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## UNIVERSITI TEKNOLOGI PETRONAS

OBJECTIVE IMAGING TECHNIQUE FOR PSORIASIS ANALYSIS

## by

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ELECTRICAL AND ELECTRONIC ENGINEERING DEPARTMENT UNIVERSITI TEKNOLOGI PETRONAS BANDAR SERI ISKANDAR, PERAK

FEBRUARY 2014

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## DEDICATION

In the Name of Allah, the Most Beneficent, the Most Merciful So, verily, with every difficulty, there is relief. Verily, with every difficulty there is relief. (Al-Inshiraḥ: 5-6)

This thesis is dedicated to my dear parents, Bapak Munadi and Ibu Suwarsih, my lovely wife, Eva Susanti, and my precious daughters, Faiza and Afiyah

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#### Abstract

Most of skin assessment methods applied in clinical practice use visual and tactile senses of dermatologists to assess features and conditions of skin diseases. However, the assessment can be subjective resulting in intra- and inter-rater variability. This research investigates the objective skin roughness measurement for psoriasis lesion assessment. Psoriasis is an incurable skin disease affecting 2-3\% of world population. The scaliness (roughness) of psoriasis lesion surface is a key assessment parameter of the Psoriasis Area and Severity Index (PASI) that subjectively assessed at four body regions. This research proposes an assessment method that incorporates 3D surface roughness algorithm with standard clustering techniques to perform objective assessment on scaliness of psoriasis lesion surfaces. High order polynomial surface fitting is implemented to estimate 3D waviness surface. The algorithm was validated on the standardised roughness and curved surfaces using 390 lesion models and a medical mannequin. The measured surface roughness is highly correlated ( 0.989 ) to the lesion models with various roughness grades and the algorithm has been validated on mannequin surface with accuracy of $94.12 \%$. The developed surface roughness algorithm is combined with an unsupervised clustering algorithm to enable objective assessment for PASI scaliness scoring using a dataset of 1,999 psoriasis lesions collected from a clinical study involving 204 patients at Department of Dermatology, Hospital Kuala Lumpur. Two unsupervised clustering methods, $k$-means and fuzzy c-means (FCM) algorithms, are used in order to form four clusters defined to represent four sets of PASI scaliness score. Qualitative statistic technique using Kappa coefficient is used in this research work to evaluate inter-rater variation of two dermatologists and two successive objective 3D scan measurements on 324 lesions. Kappa coefficient among two dermatologists is found to be 0.55 (thus cannot be considered as ground truth) whereas Kappa coefficients between two 3D scan assessments are 0.8473 ( $k$-means) and 0.8708 (FCM). These agreements are considered as a perfect agreement ( $>0.81$ ) and thus alleviated the subjectivity issues. The research work reported in this thesis shows that the developed algorithm enables to provide an objective measurement for PASI scaliness assessment on psoriasis lesions in clinical practice.


#### Abstract

ABSTRAK

Kebanyakan kaedah penilaian kulit digunakan dalam amalan klinikal menggunakan deria visual dan sentuhan pakar dermatologi untuk menilai ciri-ciri dan syarat-syarat penyakit kulit. Walau bagaimanapun, penilaian bersifat subjektif dan menyebabkan perbezaan hasil penilaian antara penilai. Kajian ini menyiasat masalah ukuran kekasaran kulit untuk penilaian luka psoriasis. Psoriasis adalah penyakit kulit yang tidak boleh diubati, menjejaskan 2-3\% penduduk dunia. Psoriasis scaliness adalah parameter penilaian utama daripada Psoriasis Area and Severity Index (PASI) yang bersifat subjektif. PASI scaliness menentukan jumlah sisik pada permukaan luka. Kajian ini mencadangkan penggabungan algoritma 3D surface roughness dan teknik clustering untuk menentukan markah PASI scaliness secara objektif. High order polynomial surface fitting digunakan untuk menganggarkan 3D waviness surface. Algoritma ini telah disahkan pada kekasaran berstandard dan permukaan melengkung menggunakan 390 model luka dan patung medik. Kekasaran permukaan yang diukur sangat bresesuaian $(0,989)$ kepada model luka dengan pelbagai gred kekasaran dan algoritma juga telah disahkan di permukaan patung medik dengan ketepatan 94,12 \%. Algoritma pengukur kekasaran digabungkan dengan algoritma unsupervised clustering bagi membolehkan penilaian objektif untuk pemarkahan PASI scaliness. Sistem pengklasifikasi ini dibuat dengan menggunakan 1,999 luka psoriasis yang dikumpulkan daripada kajian klinikal melibatkan 204 pesakit di Jabatan Dermatologi, Hospital Kuala Lumpur. Dua kaedah unsuprevised clustering, $k$-means dan fuzzy c-means (FCM) telah digunakan untuk membuat empat kelompok yang ditakrifkan untuk mewakili empat set markah PASI scaliness. Teknik statistik kualitatif menggunakan pekali Kappa digunakan dalam kerja-kerja penyelidikan ini untuk menilai variasi penilaian antara dua pakar dermatologi dan dua imbasan pengukuran 3D berturutan pada 324 luka. Pekali Kappa antara dua pakar dermatologi didapati 0.55 (dengan itu tidak boleh dianggap sebagai ground truth) manakala pekali Kappa antara dua imbasan pengukuran 3D secara berturutan adalah 0,8473 (k-means) dan 0,8708 (FCM). Kesesuaian hasil penilaian ini dianggap sempurna (> 0.81 ) dan dapat mengurangkan isu subjektiviti. Hasil kajian tesis ini menunjukkan bahawa algoritma yang dibangun membolehkan sistem pengukuran objektif untuk penilaian PASI scaliness luka psoriasis dalam amalan klinikal.


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## CHAPTER 1

## INTRODUCTION

This chapter discusses related research areas namely skin assessment and analysis, subjectivity of skin assessment, and skin roughness measurement, in formulating research problems of the thesis. The research objectives and hypotheses, and scope of work are discussed in the subsequent sections. The chapter ends with a summary of the thesis organisation.

### 1.1 Background and Motivation of Research

Various skin assessment methods are available for skin analysis in clinical practice. Most of these methods employ visual and tactile senses. Features of skin disease and their conditions such as colour, thickness, volume, or surface roughness are used as physical characteristics in these methods. These skin features which are regarded as assessment parameters are then observed, perceived and scored by dermatologists in determining severity of a particular skin condition. However, such assessments could lead to intra- and inter-rater variations due to subjectivity element in the dermatologists' assessments. Intra-rater variation is variation of assessment result that obtained from a dermatologist in different sessions. Meanwhile, inter-rater variation is result variation that given by several dermatologists to a particular object.

The need to analyse the severity of various skin diseases or disorders and their conditions (severity) through visual and tactile senses has resulted in the development of various skin disease-scoring methods. Skin disease assessment by quantifying the disease severity is common in clinical diagnosis as reported in the Medal website, listing 65 skin disorders that apply scoring method for its assessment. Many scoring methods have been developed; in the Medal website alone, a total of

313 methods are listed [1]. Disease severity is characterised by a score derived from the scores of observed parameters.

In this research, the problems of skin roughness measurement for psoriasis assessment are investigated in order to develop practical solution for daily practice. Psoriasis refers to a common chronic skin disease that affects about 2-3\% of world population [2][3]. Neimann et al. summarised the psoriasis prevalences from epidemiological studies around the world. The prevalence varies from 0.6 to $4.8 \%$ [4]. In a recent study, Chandran et al. also reported the variety of psoriasis prevalence among regions. The variation of psoriasis prevalence from several countries is $0.60-6.50 \%$ in Europe, $0.70-3.15 \%$ in North and South America and $0.08-4.00 \%$ in Africa. In Asia and Australia, the psoriasis prevalence of observed countries is reported to be $0.05-5.30 \%$ and $2.30 \%-2.57 \%$, respectively [5]. The Dermatological Society of Malaysia reported a psoriasis prevalence of $3 \%$ in Malaysia [6]. In the period of 2005-2010, the Dermatology Department of Hospital Kuala Lumpur registered 3,906 psoriasis patients of a total of 75,883 hospital's patients, giving an incidence of $5.2 \%$ [7].

Psoriasis is characterised by red plaques that can occur in a localised area of the body or it can be widespread all over the body. Psoriasis itself refers to an incurable but treatable skin disease in which the immune system sends wrong signals that accelerate the cycle of skin cells' growth. In normal conditions, skin cells grow in a cycle of about 28 days but for psoriasis new skin cells grow faster, taking about 4 days to produce as many skin cells as in normal skin [8]. This disease is distressful to the patient as the skin is itchy all over and the condition can last for a long durations [9]. Even though psoriasis is not a contagious disease, recent studies have shown that psoriasis can significantly impact on quality of life with many psoriasis patients experience social and psychological problems with their environment [10]. Figure 1.1 shows examples of plaque psoriasis lesions on several body regions.


Figure 1.1 Plaque psoriasis lesions grow on several body regions (left to right): head, upper limb, trunk, and lower limb regions.

The skin is comprised of several main skin layers. The epidermis is the outer layer and the dermis at the inner part. The layer located under the dermis is subcutaneous layer. The epidermis is made by stacking of skin cells. The living cells in the inner layer and the dead cells at the outer layer. In this layer, the skin cells are produced and maturated to replace the dead skin cells. The dead skin cells at the outermost epidermis layer form a tough, flexible, and waterproof mantle on the skin surface. These cells are shed periodically from the skin surface as keratinous scales.

The dermis forms skin characteristics such as resistance from tearing and its elasticity. This layer contains a thick network of collagen and elastic fibres. Blood vessels (artery and vein), lymphatics, nerve fibres, connective tissue cells, and immune cells are also compacted in the dermis layer. The deepest layer, subcutaneous tissue or hypodermis is not considered the real part of human skin. However, this layer has significant contributions to enable protective functions of human skin. The hypodermis comprises mainly of adipose tissue and some areolar connective tissues. This layer fixes the skin to the body, stores fat reserve, and works as thermal and mechanical insulations. Excessive amount of the adipose can thicken the hypodermis layer. The arrangement of these three main layers epidermis, dermis, and subcutaneous layer - are depicted in Figure 1.2.


Figure 1.2 Human skin is composed by several skin layers [11].
Stratum corneum is considered as the outermost layer of the skin structure and located at epidermis layer. It protects the living skin layers from the environmental harmfulness and is constructed by around twenty layers of densely packed keratinocytes, the main cell of the epidermis layer forming $95 \%$ of the skin cells [12]. The other skin cells are merkel, melanocytes, and langerhans. These cells are interlocked each other to construct some brick-like layers. In the inner part, several new keratinocytes are created to replace the older cells at the outer layers. Once created, a keratinocyte will gradually migrate from the inner to the outer layer. With this migration, the old and the dead keratinocytes at outer layer can be replaced by the keratinocytes from the inner layers [13]. The keratinocyte will transform its shape during the layer migration, from a round shape into a plate shape. This transformation process is known as differentiation. In the end of skin cell cycle, the dead keratinocyte will automatically flake away from skin surface. This final stage is known as the desquamation process. Human body can shed 30,000 to 40,000 dead skin cells from its surface every minute. This shedding amount equals 4.08 kg of cells annually [14].

The growing process of skin cell, including keratinocyte, occurs through several layers of skin epidermis. The migration process of keratinocyte through the layers within epidermis is considered as a lifetime cycle of skin cell in which a cycle
normally spends 28 days approximately. The process of skin cell regeneration is divided into four stages.

First stage, a new cell is generated from cell division in the lowest layer of the subcutis, i.e. the basal layer. In second stage, the cell shape is flattened following the cell migration to the outer layers. The cells are now located in stratum spinosum layer. This layer contains large cells with prickle contour on the surface profile. For the third stage, skin cells move to stratum granulosum layer above the stratum spinosum layer. In this stage, the cell size becomes smaller compared to its previous size. Lastly, in the fourth stage, the cells will perish at stratum corneum and fall off from the skin surface. Figure 1.3 depicts the life stages of skin cell.


Figure 1.3 The stages of skin growing at epidermis layers. The picture is reproduced from [15], [16], [17].

At higher severity of psoriasis, the lesion gets thicker with coarse white scales. The psoriasis lesions can affect any part of skin particularly on the elbows, knees, scalp, palm of the hand, chest, lower back, leg, soles and nails [18]. A periodic medical treatment for psoriatic patient is important, as the disease cannot be completely cured. Figure 1.4 shows several body parts affected by psoriasis lesions. The images are acquired from data collection sessions at Dermatology Department, Hospital Kuala Lumpur from 2007 to 2010.


Figure 1.4 Psoriasis lesions are found in various parts of the body such as trunk (1), upper limb (2), lower limb, elbow (4), and forehead (5).

### 1.2 Problem Statements

The PASI (Psoriasis Area and Severity Index) scoring method is known as gold standard for the severity assessment [19]. Six clinical severity scores i.e. PASI, BSA, PGA, LS-PGA, SPI, and SAPASI for psoriasis assessment has been systematically reviewed by Bonsard et al. [20]. Based on the review, there is no psoriasis severity scoring that meets all of the required validation criteria. Nevertheless, PASI scoring has been recommended for clinical study. Thus, PASI is considered the reference for having been widely applied in clinical studies and being the most validated among the psoriasis scoring methods [20].

PASI scoring method was introduced by Fredriksson and Pettersson in 1978. The method was proposed to evaluate a clinical efficacy of a new anti psoriatic drug [21]. To determine total PASI score, four parameters, i.e. area (ratio of lesion area to total body surface area), erythema (colour of lesion inflammation), lesion thickness, and scaliness of the lesion are required. In PASI assessment, the human body is divided into four regions namely, head, trunk, upper limbs and lower limbs. The PASI parameters of the psoriasis lesions are determined for each body region. Dermatologists use their visual and tactile senses to score the PASI parameter. The parameter scores from each region are weighted and totalled to provide a PASI score ranging from $0-72$. For the treatment efficacy, dermatologist refers at least a $75 \%$
reduction in PASI score is considered to be a clinically meaningful improvement [19]. The PASI score is calculated from following equation:

$$
\begin{align*}
\text { PASI }= & 0.1 \times\left(E_{h}+T_{h}+S_{h}\right) A_{h}+0.2 \times\left(E_{u}+T_{u}+S_{u}\right) A_{u}+  \tag{1-1}\\
& 0.3 \times\left(E_{t}+T_{t}+S_{t}\right) A_{t}+0.4 \times\left(E_{l}+T_{l}+S_{l}\right) A_{l}
\end{align*}
$$

Score ranges for PASI erythema ( $E$ ), thickness ( $T$ ), and scaliness $(S)$ are from 0 to 4 whereas the range for PASI area $(A)$ is from 0 to 6 . Four body regions namely, head, upper limb, trunk, and lower limb are denoted by subscripting $h, u, t$, and $l$ respectively for each PASI parameter. Variable $E_{h}$ for instance is used to represent PASI erythema score at the head region.

The scaliness parameter is selected as PASI parameter studied in this research. For PASI scaliness assessment, dermatologists observe several lesions that are appeared on an examined body region. The PASI scaliness scores are then given to these lesions. Dermatologists assign a scaliness score of the examined body region based on the commonest score of the lesions. To minimise tediousness in PASI scaliness assessment, a simplified procedure is performed by dermatologists by selecting a representative lesion for the examined body region. The score obtained from a representative lesion is then considered as scaliness score for the body region.

Although the PASI scoring has been accepted as the gold standard for psoriasis assessment, it is not used in daily practice. PASI scoring is tedious, time-consuming ( $\pm 30$ minutes/patient) and subjective. Four parameters must be determined at four body regions (a total of 16 assessments). The subjectivity of the scores is influenced by intra and inter-rater variation of dermatologists. Reliability and agreement of dermatologist assessments are of concerned in many studies. Assessment agreements between self assessment and dermatologist on benign melanocytic nevi patients are studied in [22]. Agreement study is also conducted to evaluate a new method for burn scar assessment [23]. A better assessment method can be decided based on the results of agreement analysis. Therefore, an objective and reliable system is required to deal with these problems.

An imaging approach to overcome the aforementioned problem is proposed to assess PASI scaliness objectively. Imaging technology is also advantageous as it can be designed to recognise skin diseases in their early stages known as the pre-disease diagnosis. As seen in chronic dermatoses, the disease is preceded by some changes. The changes may occur - even for years prior to the disorder being clinically observed [24]. This achievement can be illustrated by prevention of rosacea skin disease. This disease is characterised by reddening on facial area. The dilated blood vessels cause increasing the redness of the skin. It can occur especially in the central face region - across the cheeks, nose, and forehead. Most of rosacea cases occur in adult ages ( 30 to 50) and people with fair skin [25]. However, in early stages, the reddening symptoms on face area cannot be visually observed.

Video microscope with polarised light has been used to reveal a network of invisible dilated blood vessels underneath the skin surface. Subsurface structures can be clearly observed by applying polarised light imaging whereas finer details of blood vessels are obtained by microscope [26]. The method can detect the vessels at early stages, e.g. when the patient age is less than 10 years old. Since the method enables to detect early stages, mild topical treatments may be applied to slow down progression of rosacea [24].

Images of rosacea patterns obtained from digital camera and microscope are shown in Figure 1.5 (left) and Figure 1.5 (right) respectively. Polarising filters on both acquisitions can reveal the vessels patterns even though they are located in underneath skin surface. Microscopic magnification $100 \times$ is used to obtain the right image of Figure 1.5. Higher magnification - 200x - is applied to see the details as show in the inset image [27] [28].


Figure 1.5 Patterns of rosacea blood vessels are acquired by digital camera (left) and microscope (right).

To achieve objective assessment, the PASI scaliness visual descriptors are studied and defined in terms on surface roughness, a measurable feature that can be used to differentiate PASI scaliness scores. Abnormality of skin can be identified through several skin symptoms, such as itching skin, skin lesion, mole symptoms, acne, skin colour changes, redness, etc [29]. Skin lesions can be caused by various diseases. A total of 422 diseases that can cause skin lesions are listed in RightDiagnosis website [30]. A skin lesion is a superficial growth or patch of the skin that does not appear as surrounding normal skin [31]. Skin lesion appearance is specified by visual and tactile descriptions based on primary morphology (size, shape and thickness), secondary morphology (clustered or distributed), surface texture (roughness), location and colour [32]. A texture determination of normal and abnormal skin surface is crucial in the field of dermatology measurements, particularly in the evaluation of therapeutic or cosmetic treatments. Skin surface can be characterised by physical features, such as dryness, smoothness, thickness, and roughness.

Dermatologists conduct PASI scaliness assessment based on some descriptions of a visual appearance. These descriptions are used as a standard guidance for determining the PASI scaliness scores. In the proposed approach, surface roughness features are required to represent scaliness severity. The correspondence between surface roughness of psoriasis lesion and severity stages needs to be investigated and defined by surface roughness parameter(s) that can measured from digital surface image data.

The current problems of PASI scaliness assessment and surface roughness measurement are formulated as follows:

- Dermatologist assessment can be subjective due to intra- and inter-rater variability of human assessment. The subjectivity depends on the perception and the clinical experience of the dermatologist.
- Objective skin assessment might be performed objectively but the available methods require an invasive treatment on the measured skin. A tiny skin cut is sampled from the patient and then analysed using a scanning electron
microscopy. Trained and experienced medical personnel are required to properly prepare and analyse the skin sample.
- A non-invasive method can be applied to obtain a high accuracy measurement on surface roughness. However, several methods need a skin replica as a representation of the actual skin surface. The skin replica is required because the measuring process uses a precise profilometer (sharp needle or laser beam) to extract the profile of skin surface. Though being able to provide measurement results with high accuracy, this method in clinical practice is not so practical. In addition, well-trained personnel are required to build qualified skin replica and operate the profilometer system.

Digital image analysis of skin surface roughness for scaliness classification is proposed to resolve the aforementioned problems. 3D imaging algorithm and clustering algorithm can be applied in the proposed approach. However, there are challenges that need to be overcome in this approach. The challenges are as follows:

- Difficulty in an assessment of lesion surface roughness on curved skin surfaces. The roughness assessment should be able to measure the vertical deviation of the lesion surface at various points on human body surface.
- To determine vertical deviations on lesion surface, a reference surface is required to be a zero point of vertical distance measurement as commonly applied in industrial applications. However, there are no reference surfaces on the skin that it can be referred as the zero point. The reference surface needs to be determined before the vertical deviations can be computed.
- Dermatologists' assessment results cannot be considered as a ground truth for evaluating algorithm performance due to subjectivity influences on assessment results.


### 1.3 Research Hypothesis

In this research work, two primary hypotheses are defined. The hypotheses can be described as follows:
a. The first hypothesis sets that a lesion surface is a superimposed surface between two surfaces - curved and rough surfaces. To prepare surface roughness determination, these surfaces need to be extracted from the lesion surface. Surface roughness of lesion is calculated by averaging the vertical deviations of the rough surface. Vertical deviation due to the lesion is determined by subtracting lesion surface from an estimated waviness obtained by fitting a polynomial surface to lesion surface. Most of the estimated waviness surfaces have the form of curved surfaces.
b. The second hypothesis defines that surface roughness of the lesions can be used to build clustering algorithm for PASI scaliness scoring. With availability of large datasets, an unsupervised clustering algorithm can be applied to obtain the boundary levels of lesion surface roughness for PASI scaliness scoring.

Figure 1.6 depicts the cross-sectional view of a skin lesion on normal skin surface. Vertical deviations of lesion surface can be either positive or negative and the average of deviation magnitudes is used to represent surface roughness of the lesion.


Figure 1.6 Cross-sectional view of a skin lesion on normal skin surface.

### 1.4 Research Objectives

Two research objectives are specified for the thesis. These objectives are described as follows:
a. The first research objective is to develop a 3D imaging algorithm for accurately measuring the surface roughness of skin lesions. The algorithm must be valid for all skin lesions that appear at any body parts.
b. The second objective is to develop an objective and reliable PASI scaliness scoring using an unsupervised clustering technique on surface roughness of skin lesions.

### 1.5 Scope of Work

Three items in the scope of work have been identified that limits the coverage of the research. The limitations are defined for the type of skin disease, PASI parameter, and the size of acquired images. The type of psoriasis lesion in the research work is limited to plaque psoriasis. This is the commonest types of psoriasis accounting around $80 \%$ of psoriasis cases. Other psoriasis types are flexural, guttate, pustular, nail psoriasis, psoriatic arthritis and erythrodermic [4]. The Malaysian Psoriasis Registry found that the percentage of plaque psoriasis in Malaysia reaches $80.9 \%$ [33]. Surface roughness of psoriasis lesion is determined by applying 3D image analysis on scanned lesion surface. The PASI parameter studied in this research is limited only to PASI scaliness. The other parameters - area, erythema, thickness are not considered since the surface roughness is not involved in the scoring process of the parameters.

The measured area is limited to the maximum area of 3D scanner used in this research. The maximum measured area is $40 \times 30 \mathrm{~mm}^{2}$ and the vertical heights are stored as a matrix with $640 \times 480$ data points in size. Figure 1.7 shows an example of 3D lesion surface taken by using 3D optical scanner. This image is a second sample of psoriasis lesion. The image is acquired from upper limb region of a psoriasis patient at Dermatology Department, Hospital Kuala Lumpur.


Figure 1.7 An example of 3D lesion surface obtained from psoriasis patient.
This localised measurement is not contrary with the clinical procedure of PASI assessment. In real situations, the dermatologist scores several representative lesions one by one even though there are several lesions at the same body region (head, upper limb, trunk, or lower limb). The score given for a body region is determined by selecting the commonest PASI scores of psoriasis lesions at the same body region. The same procedure can be conducted by applying the proposed method. The surface measurement can be conducted repeatedly to another lesion located at the same region. Patient data profile, such as gender, age, ethnic, and medical history would not be considered in this study. There is no significant difference on scale surface appearances based on those parameters.

### 1.6 Organisation of the Thesis

The thesis is organised into six chapters including the first chapter. Chapter 1 introduces background of the research work, initially by illustrating the current method on skin assessment and analysis. The method can suffer from an assessment inconsistency due to human subjectivity. Skin roughness is one of the common features used in the skin assessment. The feature is intensively studied in this work for having a relationship with the scaliness characteristic of psoriasis lesion. Research problems and objectives are formulated in this chapter as well as a research hypothesis on imaging method for skin roughness analysis. Scopes of work, such as the type of skin disease, studied PASI parameter and acquired image are defined afterward. In the final section, organisation of the thesis details the thesis content started from chapters 1 to 6 .

Chapter 2 presents a medical review of the skin structure and the clinical features of skin disease. This chapter begins by describing the physical structure of human skin. Layers types and its function are mentioned as well. Psoriasis, a well-known skin disease, is studied in this research work.

Chapter 3 furthermore describes related methods that are used for surface roughness analysis. Surface roughness initially is determined by drawing a stylus on the measured surface known as mechanical surface profilometry method. Several imaging technologies for surface analysis, such as laser profilometry, speckle imaging, microscopy imaging, and structured light projection are developed to enable 3D surface acquisition. Surface roughness is determined from the extracted roughness component of surface profile. To extract roughness components from the unwanted components, surface filtering methods have been developed. This section presents the commonest filters for surface filtering i.e. Fourier transform, Gaussian filter, wavelet filtering, and polynomial surface fitting. Calculation of surface roughness parameter is then performed on filtered surface to obtain measured surface roughness. In this section, three roughness parameters; which are frequency-based parameter, amplitude-based parameter, and fractal parameter are reviewed. Applications of surface roughness measurement for manufacturing and medical purposes are provided in this chapter.

Chapter 4 describes the development of surface roughness algorithm. In the first stage, surface roughness is simulated as a computational model of lesion surface. Estimated waviness is extracted from the computed model for testing the performance of polynomial surface fitting. Validation study of surface roughness algorithm on abrasive paper, as a rough surface, and curved surface are reported in this chapter. From the validation study, accuracy and total standard deviation are determined.

Chapter 5 discusses a surface roughness analysis for a psoriasis assessment. A clinical study has been conducted to collect 3D surfaces of psoriasis lesions. Experiment set-up and procedures for data collection are detailed in the first section. Development and validation study of clustering algorithms are described in the following sections. Two unsupervised clustering algorithms - $k$-means and fuzzy $c$ -
means clustering algorithms are applied to determine PASI scaliness score based on lesion surface roughness. Objectivity of the scoring algorithm is evaluated by performing agreement analysis between the first and the second assessment.

Chapter 6 finally summarises the findings on this research work into a conclusion section. This is followed by some recommendations and suggestions for future work. Some potential applications for implementing the developed skin roughness algorithm are mentioned in the end of this chapter.

## CHAPTER 2

## MEDICAL LITERATURE REVIEW

Chapter 2 presents both general information on human skin structure comprising several skin layers and a number of related diseases including the assessment. It begins from Section 2.1 that describes the characteristics of skin layers and their functions. Section 2.2, provides a detailed discussion about psoriasis, a chronic skin disease which is the object of study in the research work. This section also includes a discussion on various psoriasis types and its treatment methods for reducing the effects of psoriasis. Following this, Section 2.3 presents the PASI (Psoriasis Area and Severity Index) scoring method as a gold standard for psoriasis assessment in order to obtain the treatment efficacy for the dermatologist in quantifying the psoriasis severity. The contents of Chapter 2 are finally summarised in Section 2.4.

### 2.1 Skin Structure

Skin is considered one of the largest organs of human body [34]; in adult, its weight is about 4 to 5 kg or $7 \%$ of total body weight covering almost $1.7 \mathrm{~m}^{2}$ of body surface area [13]. The skin area between male and female will be different in size. For male, it ranges from 1.39 to $2.10 \mathrm{~m}^{2}$ and for female the area interval is 1.19 to 1.88 $\mathrm{m}^{2}$ [35]. Based on its structure, skin is divided into two parts namely actual skin (cutis) and subcutaneous tissue (tela subcutanea) [13]. The actual skin layer comprises of two layers - epidermis and dermis. Epidermis is a stratified squamous keratinous epithelium whereas dermis is a dense network of collagen and elastic fibres. Epidermis is located in outer layer of $0.1-0.2 \mathrm{~mm}$ in thickness. This thickness range is observed at most of body areas. Exceptions are found at the hand palms and soles of the feet, the epidermis layer in these areas is thicker compared to other body areas ranging from 0.8 to 1.5 mm . The research is focused on the outer
surface of epidermis since most of the skin diseases affect and appear on this layer.

The skin organ is composed by four main tissue types - epithelial, connective, muscle, and nervous. These tissue types are also mostly found in the other organs. The tissues are combined to perform specific function of the organ. Skin organ has multiple functions [13] such as: (1) To protect the skin from mechanical, thermal and chemical damage; (2) To regulate body temperature; (3) To maintain skin moisture from dryness; (4) To sense physical stimulus trough the skin receptors; (5) To build immune system from pathogenic threats; (6) To communicate an expression by changing the skin colour of face area. These functions can be obtained for normal skin but for the diseased skin, one or more functions would not be performed normally. Skin organ, derivatives of skin (sweat glands, oil glands, hair, and nails) and accessory structures (blood vessels and nerves) are integrated to form an integumentary system. The function of this system is to protect the human body from harmful interactions with the outer substances and environment [34].

### 2.2 Clinical Features of Psoriasis

The psoriasis prevalences in several countries from 1964 to 2005 have been studied by Neimann et al. Their study found that the psoriasis prevalence around the world is in the range of 0.6 to $4.8 \%$ [4]. In a recent review, Chandran et al. reported psoriasis prevalence varies among the regions. The prevalences of psoriasis are $0.60-6.50 \%$ in Europe, $0.70-3.15 \%$ in North and South America and $0.08-4.00 \%$ in Africa. In Asia and Australia, psoriasis prevalences of $0.05-5.30 \%$ and $2.30 \%$ $2.57 \%$, respectively have been reported [5]. These prevalence data are obtained from clinical based and field survey from 1998 to 2009. However, if the worldwide population is used as the prevalence base then the psoriasis prevalence around the world is about 2 to $3 \%$ [2][3]. The Dermatological Society of Malaysia studied that the psoriasis prevalence in Malaysia is 3\% [6]. The Dermatology Department of Hospital Kuala Lumpur registered 3,906 psoriasis patients of a total of 75,883 hospital's patients for the years 2005-2010, giving a prevalence of $5.2 \%$ \% [7]. Periodical medical treatment for psoriatic patient is important, as the disease cannot be cured. The treatment for psoriasis can be grouped into three types - topical,
phototherapy, and systemic medications. The treatment is specific for each patient. It is decided based on both disease progression and medical response of treated patient. Topical method is usually used for handling psoriasis at mild levels. The second treatment is phototherapy that uses a UV light emission to control the growth of psoriasis lesions. The third treatment is systemic method applied when the two previous methods are not considered effective to deal with psoriasis.

For adults, psoriasis is believed to be the most prevalent immune-mediated skin disease, which is initiated by an activated cellular immune system. Therefore, the psoriasis is also considered as an organ-specific autoimmune disease [36]. The most common form of psoriasis is psoriasis vulgaris reaching $90 \%$ of all psoriasis cases. In psoriasis vulgaris, scaly papules and plaques are well-defined from surrounding normal skin [37].

Pathogenesis of psoriasis has not been being completely understood until recently. The immune system and T lymphocytes so far are considered as the initiator for psoriasis. In response, the epidermal cell cycle is then shortened resulting in silvery scale lesions [38]. The psoriasis scales are formed from a number of abnormally stacked cells on the stratum corneum - an outermost surface of epidermis layer. The granular layer of the epidermis is much reduced by increasing the size of cells stacking. Clinical features and severities of psoriasis are not only varying in time but also specific for each individual.

In characterizing psoriasis, four abnormalities can be used. The first abnormality is vascular changes that are shown by dilatation and tortuosity of the papillary blood vessels. At this point, there will be elongation and enlargement of epidermal blood vessels [39]. In such condition, the skin becomes reddish and this is widely known as psoriasis erythema. The second condition is inflammation in which polymorphonuclear leukocytes from the dermal vessels move into the epidermis. Here, the number of leukocyte of psoriatic lesion significantly increases and many immune-related pathways are activated [39]. In turn, lesions with a large amount of activated T helper cells $\left(\mathrm{CD}^{+}\right.$and $\left.\mathrm{CD}^{+}\right)$will release proinfalammatory cytokines. The cytokines cell is signalling cell that can activate the body immune system to create skin inflammation. The third abnormality is hyperpoliferation (high rate on
cells regeneration) of the keratinocytic layer. This layer protects the body from pathogens, heat, light radiation, and water losses. The last abnormality is the altered epidermal differentiation. Keratinocytes keep their nuclei in the protected layer (parakeratosis) and lose the granular layer. They grow excessively with different appearance compared to normal skin. Comparison between normal skin and skin affected by psoriasis is shown Figure 2.1.

(a) Histology image and corresponding scheme of normal skin

(b) Histology image and corresponding scheme of skin affected by psoriasis

Figure 2.1 Skin histology of (a) normal skin and (b) skin affected by psoriasis [36].

These progressions in the epidermis layer will introduce skin scales on the lesion locations [39]. The epidermal rete is extended and the papillary blood vessels of
psoriatic plaque are dilated. Epidermal rete is ridge profiles located at boundary area between epidermis and dermis layers. The epidermal rate enables epidermis and dermis to be interlocked into a unified skin layer. These enlarged blood vessels, as shown by psoriatic plaque cross-section in Figure 2.1 (b), make the affected skin become reddish which is defined as lesion erythema. Hyperpoliferation of keratinocyte is also depicted by changing on the stratum corneum layer. Previously, its thickness is slimmer but after affected by psoriasis, the layer goes very thick and irregular.

Psoriasis lesions typically appears in certain locations such as scalp, ears, elbows, umbilicus (belly button), buttocks (gluteal cleft) and genitalia areas, knees, soles of the feet, fingers and toes. It is possible for the lesions to grow at moist areas; for examples in armpits, under breasts, and groins [40]. Figure 2.2 describes the lesion locations that are commonly found in psoriasis cases.


Figure 2.2 Common locations of psoriasis lesions growing on human skin. The picture is created by combining graphic materials and description from [40] [41] [42].

Based on the appearance and characteristic, psoriasis could be categorised several types that are presented in the following sections.

### 2.2.1 Plaque psoriasis

Psoriasis lesion typically is covered by silvery white scales with varying thicknesses. Plaque psoriasis usually starts in early adulthood and will generally persist for a long time. Plaque psoriasis lesions frequently occur on certain areas such as elbows, knees, and the scalp [36].

The amount of scales may vary among the psoriatic patients and also at different parts of the same patient [43]. The lesion is usually clumped as a single skin patch. However, due to improper treatment, the lesion size could be enlarged. The enlarged lesion can be merged with another neighbour lesion to create a larger lesion. Some images of plaque psoriasis lesions from clinical study at Hospital Kuala Lumpur are shown in Figure 2.3. From these examples, it can be shown that the large lesions are formed by merging some smaller adjacent lesions.


Figure 2.3 Plaque psoriasis lesions appear at several body regions such as (a) elbow, (b) upper back, and (c) trunk.

### 2.2.2 Guttate psoriasis

Guttate psoriasis originated from Greek gutta meaning a tiny drop. The drop gives a description of the acute onset of unnumbered tiny psoriasis lesions with diameter 2 10 mm . A total of five to hundreds lesions can appear at certain affected regions, such as head and limbs. Guttate psoriasis occurs in $2 \%$ of total psoriasis cases. Plaque psoriasis can be initiated by suffering guttate psoriasis at younger ages [43]. Figure 2.4 shows guttate psoriasis lesions at several body regions.


Figure 2.4 Example of guttate psoriasis lesions. These lesions are spread at several body regions such as (a) chest and arm [44], (b) forearm [45] and (c) back region [45].

### 2.2.3 Flexural (inverse) psoriasis

Flexural psoriasis occurs at the flexural areas. These areas are located between two body parts such as at the folding areas of breast, genital, and armpit. The flexural lesions appear as red and shiny lesions. Its plaque boundaries are also easily to pronounce from the surrounding normal skin. However, the disease sometimes is misidentified as candidal, intertrigo, or dermatophyte infections [43]. Figure 2.5 shows the folded skin areas that are affected by psoriasis.


Figure 2.5 Flexural psoriasis growths lesions are appeared at folded skin areas. The areas such as (a) armpit [37] (b) ear [46], and (c) umbilicus.

### 2.2.4 Generalised pustular psoriasis

Pustular type is characterised by red skin on the affected area and scattered by many pustules. These pustules contain white blood cells that can dry and peel causing the
skin to be glazed. Moreover, this type also can cause a fever to the patients. In many cases, the adult patients need to be hospitalised [43]. Figure 2.6 illustrates the examples of skin affected by pustular psoriasis.


Figure 2.6 Many pustules scatter on the inflamed skin affected by generalised pustular psoriasis at (a) palm [47], (b) upper back [48], and (c) neck [49] regions.

### 2.2.5 Erythrodermic psoriasis

Erythrodermic psoriasis can affect the body skin surface, either partially or completely affected. This psoriasis type might be developed from the gradual enlargement of chronic psoriasis plaques. It is also triggered by an abrupt termination on a systemic psoriasis treatment, infections, or allergic reaction to the medicines. This psoriasis type is characterised by several symptoms, such as periodical and widespread appearance, inflaming skin surface, and lesion scales in the form of sheets. Patients with erythrodermic psoriasis will experience severe itching and pain, increasing heat rate, and unpredictable body temperature. Abnormality in skin temperature regulation can lead hypothermia, high output cardiac failure, and change in blood metabolism. Once the symptoms have been detected, hospitalised treatments need to be given immediately to the patients [43] [50]. Figure 2.7 depicts the example of erythrodermic psoriasis.


Figure 2.7 Erythrodermic psoriasis covers most of body surfaces at areas such as (a) trunk [51], (b) lower limbs [52], and (c) hand including wrist area [53].

Psoriasis treatments can be categorised into three therapeutic modalities, namely topical agents, phototherapy using appropriate ultra violet wavelength, and systemic medications. Topical agents are used for treating mild psoriasis and resistant lesions at higher severity. Phototherapy is applied for psoriasis at moderate level. Meanwhile, the systemic treatment is implemented for treating severe cases. This treatment includes photo chemotherapy, oral medicine consumption, and biological injection [54]. In topical category coal tar, vitamin D analogues, steroids, and dithranol are applied to the skin surface. On the other hand, through phototherapy, Narrow-band Ultraviolet-B (NBUVB, $315-280 \mathrm{~nm}$ ) and Psoralen Ultraviolet-A (PUVA, $400-315 \mathrm{~nm}$ ) treatments are aimed to improve the patient condition. In systemic treatment, various medicines, such as methotrexate, retinoids, sulphasalazine, cyclophosphamide, and cyclosporine can be given. All of these psoriasis treatments are discussed in the following sections [55].

### 2.3 Skin Assessment and Analysis

In the 1950s, skin assessment was relatively simple based on the direct accessibility of the skin surface using visual and tactile senses. Dermatologists are able to assess and diagnose using only their perception of the physical appearances of skin. In contrast to other medical specialists, such as internists, surgeons, and radiologists, equipments or modalities such as electroencephalography (EEG), magnetic resonance imaging (MRI), x-ray, and medical ultrasonography are used to observe
affected human body parts [24]. Nowadays, many advanced equipments have been used to examine and measure skin abnormalities. These equipments such as digital single-lens reflex camera (DSLR) [56], dermatoscopy [28][57], video dermatoscopy also known as video microscopic [28][58], optical coherence tomography [59], confocal scanning laser microscopy [60], stylus and laser profilometries on skin replicas [61][62], and tribo-acoustic system [63] have been used in skin assessment. However, most of them are only available for clinical studies and not for clinical practice.

In clinical practice, the skin diseases are commonly analysed and assessed through visual and tactile senses using skin disease-scoring methods. Many scoring methods are available to determine severity of various skin diseases. For example, eczema area and severity index (EASI) is used to grade atopic dermatitis [64]. EASI defines four parameters which are erythema, thickness, scratching and lichenification (lined skin). The intensities of these parameters are obtained from four regions of body (head and neck, upper limbs, trunk, and lower limbs). Patient age and affected body surface area are considered as the weighting factors in a final score calculation. The EASI score ranges from 0 to 72.

SCORAD (SCORing Atopic Dermatitis) is another example of well known skin disease assessment. SCORAD is used to grade atopic dermatitis severity. There are three parameters need to be assessed. The parameters are lesion existence on body regions, disease severities, and subjective symptoms. The first and second parameters are scored by dermatologists. The third parameter - subjective symptoms - is scored by patient based on his or her subjective perception and experience.

The combination of assessment results by dermatologists and patient represent the patient condition [65]. The SCORAD score ranges from 0 to 103. Both scoring methods purposively are to standardise the atopic dermatitis assessment and evaluate therapeutic response [66].

The Cutaneous Lupus Erythematosus Disease Area and Severity Index (CLASI) scoring is performed on the patient to determine the severity of cutaneous lupus erythematosus. This method assesses four parameters (erythema, scale,
dyspigmentation, and scarring) at 16 anatomical locations, such as scalp, ears, nose, back, arms, hands, legs, etc. CLASI scoring is applied to evaluate treatment efficacy on cutaneous lupus erythematosus disease [67].

Several scoring methods have also been developed for psoriasis assessment such as National Psoriasis Foundation (NPF), Psoriasis Score (NPF-PS), Physician static global assessment (PSGA), Overall lesion assessment (OLA), Lattice System Physician's Global Assessment (LS-PGA) Psoriasis Global Assessment (PGA), Copenhagen Psoriasis Severity Index (CoPSI), and Psoriasis Area and Severity Index (PASI) [19], [68], [69].

Among the existing psoriasis scoring methods, PASI is considered the gold standard for psoriasis assessment. However, new scoring methods developed are motivated to simplify assessment procedures, accommodate new parameters, and possibly minimise the score variations among the raters. However, due to the inherent subjectivity of human assessment most of these scoring methods suffer from intra- and inter-rater score variations. The subjectivity on assessment in turn can result in treatment inefficacy, as seen in psoriasis cases [24].

### 2.3.1 Subjectivity of Skin Assessment

Most assessment methods are performed by using visual and tactile senses. A standard guide based on physical characteristics of the disease is used by dermatologists to score disease severities; perception will be significant in the physical interpretation of the disease. In many cases, subjective clinical assessments by dermatologist produce intra- and inter-rater variation.

Figure 2.8 illustrates two dermatologists assessing psoriasis lesions in a session of clinical trial using visual and tactile senses. Numerous studies have reported the intra and inter-rater variation problem on scoring methods of skin diseases. Chen et al. studied inter-rater reliability in leprosy assessment in which pain sensations of the skin lesion and peripheral nerve enlargement were assessed. It was found that the agreements among the raters were less than 0.53 . In this study, leprologist, medical students, and leprosy staff were considered as the raters and involved to examine the
disease [70]. Forbes-Duchart et al. investigated an inter-rater reliability in paediatric burn scar assessment. Three raters are involved in the study.


Figure 2.8 Two dermatologists are assessing psoriasis lesions.
Modified Version of the Vancouver Scar Scale (MVSS) is used to score the scar severities. MVSS defines four parameters for scar assessment, such as pigmentation level, vascularity, pliability (skin elasticity), and height (scar thickness). In the analysis of $\kappa$ agreement between two raters, it is found that all MVSS parameters have a low inter-rater reliability. The maximum $\kappa$ coefficients for pigmentation, vascularity, pliability, and height parameters are $0.16,0.25,0.38$, and 0.58 respectively. The research suggested the use of MVSS total scores rather than that of individual parameters [71].

Berth-Jones et al. reported that intra-rater reliability of three scoring methods for psoriasis (PASI, CoPASI, and PGA) cannot be categorised as a perfect agreement. In the inter-rater reliability evaluation, agreements of PASI and CoPASI are considered only substantial and become lower for PGA assessment (moderate agreement). In this regard, CoPASI might be better than PASI and PGA, particularly for assessment of milder cases [69].

Principally, the development of objective assessments and measurements is continuously improving to reduce subjectivity and introduce objectivity in skin assessment [72]. Although the scoring parameters have been simplified, the intra and inter-rater variations problem remain. To eliminate the aforementioned
problems, it is pertinent to develop techniques to measure and quantify the parameters in order to obtain objective scores for skin assessment.

### 2.3.2 PASI Scoring for Psoriasis Assessment

Psoriasis assessment using PASI scoring method was proposed by Fredriksson and Pettersson in 1978. This new assessment is used to evaluate efficacy of Ro 10-9359, a retinoic acid derivative, for treating the psoriatic patients. In their study, more than $90 \%$ reduction of psoriatic lesions in 10 patients out of 20 after 4 to 8 weeks of medical treatment has been found. Total PASI score functionally is used to indicate the reduction of psoriasis severity [21].

As mentioned in previous chapter, there are four PASI parameters - area, erythema, thickness, and scaliness - that need to be assessed in a round PASI assessment. Dermatologists carefully select the representative lesions from each body region for scoring (head, upper limb, trunk, and lower limb). The head incorporates face, scalp, ear, and neck. Meanwhile, chest, stomach, back, armpits and genital are considered as the parts of the trunk. The upper limbs comprise the arms, hands, and palms whereas the legs, buttocks, feet, and soles are belong to the lower limbs [73]. In this research work, the patients are required to wear their underwear during the assessment sessions. It implies that the psoriasis condition on breast, buttock, and genital regions are not assessed during the clinical study.

The following description is an example of a psoriasis patient that has been involved in clinical study (Patient 052, male, 35 years old). He has been assessed by dermatologist (Dermatologist 1). Figure 2.9 shows a summary of the PASI score from the assessment.


Figure 2.9 An example of PASI scores summary obtained from dermatologist assessment to a psoriasis patient.

The total PASI score for this patient is calculated using the scores to Equation (1-1) (Chapter 1). The detail of calculation is explained as follows:

$$
\begin{array}{r}
\text { PASI }=0.1 \times\left(E_{h}+T_{h}+S_{h}\right) A_{h}+0.2 \times\left(E_{u}+T_{u}+S_{u}\right) A_{u}+  \tag{2-1}\\
0.3 \times\left(E_{t}+T_{t}+S_{t}\right) A_{t}+0.4 \times\left(E_{l}+T_{l}+S_{l}\right) A_{l}
\end{array}
$$

The result shows that the total PASI score for the observed patient is 22.3. This total value is important for dermatologist, especially to determine the treatment efficacy. In this case, the given treatment is considered significant if total PASI score after treatment can achieve at least 5.5 ( $75 \%$ reduction from initial total PASI score). According to maximum scores for each parameter and total PASI score equation, the score can achieve 72 for a maximum score. In real situation, a maximum score is uncommon since the patients are being treated to reduce the psoriasis severity. The high total PASI scores usually are not more than 40 [74].

PASI scaliness assessment is conducted by dermatologist based on certain physical descriptions as listed in Table 2.1. These descriptions are referred by dermatologists as a standard guidance for determining PASI scaliness scores.

Table 2.1 Features for PASI scaliness scoring used by dermatologists.

| PASI Score | PASI scaliness visual descriptors |
| :---: | :--- |
| 0 | No scale |
| 1 | Fine scale, some of lesion covered by scale |
| 2 | Coarse scale with most lesion partially covered by scale |
| 3 | Coarse scale almost all lesions covered by a rough surface |
| 4 | Very coarse thick scales covering all lesions, very rough surface |

Dermatologists score the lesions based on their visual and tactile perceptions. Figure 2.10 shows the psoriasis lesions for PASI scaliness scores 0 to 4 . These images are used to provide training on PASI assessment. The high score lesion is rougher compared to the lesion with low scores. It is clear that the lesion roughness is related to the scale appearance on the lesion surfaces. As shown in Figure 2.10, no scales appear on lesion score 0 and only fine scales at lesion score 1 . However, at score 4, a large amount of rough scales can spread over lesion surfaces.


Figure 2.10 Psoriasis lesions and its PASI scaliness scores given by human visual and tactile perceptions [75].

In its application, it is found that PASI assessment is subjective, time-consuming and requires trained physician. Thus, it leads the assessment to be difficult to be applied in either daily practice or a large-scale epidemiologic research. In addition, the self-assessment by the patients or clinical study participants are not included in PASI scoring but all parameter scores are obtained only from dermatologists' observation and assessment [76]. Regardless of those drawbacks, the PASI scoring, however, is still considered as the gold standard in a psoriasis assessment.

### 2.4 Summary

This chapter presents the information about human skin, psoriasis skin disease, severity assessment, and its treatments. These descriptions describe the significance of the study on skin disease assessment, especially for psoriasis skin disease. The disease brings some effects on the largest part of human body and cannot be totally cured. In response, an accurate and objective assessment is highly required to evaluate the treatment efficacy given by the dermatologist.

Psoriasis is a common skin disorder that affects about $2-3 \%$ of world population. It is known as the most prevalent immune-mediated skin disease for adults. Until recently, the initial process of psoriasis has been being in research and not totally discovered yet. The psoriasis lesions can affect at any body parts. Psoriasis is categorised into several types based on its specific appearance. The types include plaque, guttate, flexural, and erythrodermic psoriasis. Plaque psoriasis is the commonest case among the psoriasis types. Considering that psoriasis cannot be totally cured, a long term treatment for this disease is highly needed.

Psoriasis assessment using PASI (Psoriasis Area Severity Index) scoring method was proposed by Fredriksson and Pettersson in 1978. The scoring method defines four assessment parameters, namely area, erythema, thickness, and scaliness. The parameters are scored by dermatologists for each defined body regions (head, upper limb, trunk, and lower limb). The scores are given based on visual and tactile descriptions of the psoriasis lesions. Total PASI score is used to indicate the reduction of psoriasis severity.

PASI scaliness assessment is conducted by dermatologists based on their visual and tactile perceptions. The PASI scaliness scores are ranging from 0 to 4 . The scores are used to represent the amount of scale on the lesion surface. It is found that PASI assessment is subjective, time-consuming and requires trained physician. Although these disadvantages are permanently extant in PASI scoring method, the scoring method is still considered gold standard for psoriasis assessment.

## CHAPTER 3

## SURFACE MEASUREMENT METHODS

Chapter 3 describes about the background information of the methods for measuring surface roughness and analysing the measurement results using a clustering algorithm. Section 3.1 reviews the methods of the surface roughness analysis. In the initial section, a long history of the surface roughness implementation is presented. Since the beginning era, mechanical surface profilometry has been commonly used to measure the surface texture of manufacturing and metallurgical products. Here, a sharp needle is drawn along a line on the measured surface to acquire height information. Although being able to provide a high resolution, this method in fact still has the very restricted scan area and scanning time. In response to this restriction, the imaging technologies then provide a light profiling medium to extract the height profile of the measured surfaces. Three imaging methods for 3D surface measurements are presented in this subsection; those are laser profilometry, light scattering and structured light projection. To enable surface roughness measurement, surface filtering and surface roughness parameters have to be defined. Surface filtering is a method aimed to extract a roughness component from certain unsuitable components of surface profile. Such methods include Fourier transform, Gaussian filter, wavelet filtering, and polynomial surface fitting. The following subsection discusses the surface roughness parameters that dominate the current standard in a surface roughness measurement. Section 3.2, on the other hand, shows some applications of imaging methods for skin surface roughness measurement. At last, the summary of the materials that have been described in Chapter 3 is briefly described in Section 3.4.

### 3.1 Surface Roughness Analysis

The main problem of assessing surfaces is related to the small size of the machining marks. Surface roughness is an important quality parameter in many industrial applications such as manufacturing process of metals, semiconductors, ceramics, paper, and plastic products [77]. Until in the end of 1920s there was no effort to create a surface measurement system. The metallurgists at that time merely examined the surfaces by their visual and tactile perceptions. This examination might be sufficient in one factory but not for further examination outside to the factory [78]. To standardise the assessment, several samples with certain roughness were provided to be used as the references to compare with the assessed surface. By comparing with these reference samples, the metallurgists were able to classify the assessed surface according its roughness levels.

### 3.2 Mechanical Surface Profilometry

Mechanical surface profilometry is a conventional method for measuring surface roughness. It uses a sharp stylus made of diamond for sweeping along the measured surface. Here, the stylus tip, which has a few micrometers of radius size, is drawn on the surface. During the movement, the stylus body will move up and down following the profile variations of scanned surface. In common, the recommended length for stylus scanning is 5 mm [79]. The vertical variations of the stylus are automatically interpreted as the roughness profile of the surface. The scan resolution of stylus profilometer is limited by the diameter size of stylus tip [79]. To obtain a reliable measurement, ridge valleys of the surface profile should be wider than the diameter size. If the valley is narrower than the stylus tip diameter, the profilometer will not extract the actual elevation of the surface. This problem can be solved by applying a finer probe to replace the stylus tip, as applied by Atomic Force Microscopy (AFM). However, the tiny scan area and the long scanning time become the drawbacks of AFM. For example, to measure an area of $70 \times 70 \mu \mathrm{~m}^{2}$ it requires at least 1 to 2 hours for each scan [79]. This tiny scan area might not represent the actual condition of the measured surface. A larger scan area is required to cover the surface conditions accurately. The advantages of mechanical surface profilometry
are listed as follows [80]:

- Measurement is performed to the measured object without any prior treatment.
- The measurement can cover a wide range from tens of microns to ten of millimetres. In addition, the vertical sensitivity can be increased less than 10 nm .

Despite these advantages, the mechanical surface profilometry has several drawbacks:

- Contact stress of stylus tip can damage the scanned surface. The stylus also could scratch the surface especially for the soft materials. Although the stylus load is low, it can cause a high stress for being applied and centred to a very small area of stylus tip.
- A number of vertical points might not correctly be profiled by the stylus tip. It occurs when the size of stylus tip is bigger than the vertical gaps. As a consequence, the stylus tip is not able to enter deeply to the point location.
- To obtain two-dimensional topographical information, a number of the consecutive line scans are required. This scanning process can be timeconsuming. Based on this constraint, the sample area is typically limited to a small size only.

The drawbacks founded on mechanical surface profilometry motivate the researchers to develop non-contact techniques for surface roughness determination.

### 3.3 Imaging Technologies for Surface Analysis

After established for hundred years, mechanical stylus profiling has been proposed to be replaced with several optical techniques. These optical techniques give some advantages in measuring surface roughness as follows [80]:

- Scanning process is not going to damage the scanned surface. Additionally, profiling process can be performed without applying any direct contact between the sensor and the scanned surface.
- Physical stylus is not longer used as a profiling sensor. The function has
been replaced by applying a light medium that can omit the probe breakage problems.
- The scan using the light as the measurement medium can have a very fast completion.
- The light beam as the probe medium is able to penetrate the vertical profile deeper than the probe stylus. In the mechanical stylus, the probe cannot measure an actual surface that is located in small gap - smaller than the stylus.
- Measurement is able to cover a large area. The light also can be directed to perform measurement on a large area.

Beside those advantages, the optical methods have several challenges, such as:

- The light coming from a certain angle might be covered by the surface profiles in which the reflected light in this condition will not give any depth information correctly.
- The measurement results can be influenced by the optical characteristic of the surface material. In this case, the reflected light might come from the unwanted surface resulting errors for a depth measurement. As shown in Figure 3.1, the $n_{1}$ and $n_{2}$ are made of the distinct materials. Stylus profiling is able to measure the surfaces correctly since the stylus sweep is not reliant on the optical characteristics of the materials. However, for the optical measurement, it may obtain inaccurate results.


Figure 3.1 Stylus and optical measurements.
Imaging technology is applied for measuring surface roughness. This
implementation is motivated by two main objectives: to provide high resolution (depth and lateral resolutions) measurement and to perform high speed scanning. In this case, the earliest technology is laser profilometry.

### 3.3.1 Laser Profilometry Method

In laser profilometry, a laser beam is emitted and line by line directed to a scanned surface. The laser beam is used to profile the elevation points of the surface. The elevation points are determined by applying triangulation method to the reflected light. The reflected light is focused by an optical lens to a light sensitive detector. A surface topography, afterward, is interpreted from the projection of the reflected laser on light sensitive detector. Figure 3.2 describes the triangulation principle of a laser camera in extracting the elevation profiles of the scanned surface. The laser light is exactly reflected to the centre of light sensitive detector if the light is projected onto a reference plane. Once the surface is located either upper or below the reference plane, the reflected laser light would be deviated from the centre of the detector. Depth information is then determined from the deviations of reflected laser light on the light detector. A higher deviation represents a further location from the reference plane.


Figure 3.2 Laser triangulation of laser camera.
The laser profilometry has some advantages such as providing high resolution scan, being able to penetrate and scan deep surfaces and being adapted with any kinds of materials. However, this also has some drawbacks, for example the
scanning process that cannot be performed as a fast measurement. By using a laser scanner, the surface information obtained is not a grid matrix of elevation data. The information is recorded by a laser system as a point of cloud. To construct a 3D surface, these measured points need to be connected using an interconnection of small triangular meshes. In order to enable practical linear calculation, these meshes are then converted to a grid of XYZ coordinates. The data obtained from laser profilometry has been applied for measuring skin disease treatments. In this research, a laser scanner, Konica Minolta Vivid 910, has been used to scan the surface of the psoriasis lesions.

The laser scanner can properly scan the skin surface if the surface is stable. Conversely, the scanning cannot be properly performed for the regions with some vibrations on the surface such as at the chest regions. The vibrations are caused by a periodical breathing movement. For having no stability on the scanned surface, a focused surface is difficult to be obtained. Figure 3.3 shows the example of psoriasis lesions obtained from the laser scanner. Figure 3.3 (a) shows 2D image of plaque psoriasis lesions at lower back region. The surface vibrates are continuously caused by the breathing movement. Figure 3.3 (b) displays the blurriness of the scanned lesion at a vibrated surface. Some parts of the scanned surfaces are overlaid due to the surface movement during the scanning process. These overlying edges are found at the edges of tape 3 (pointed by arrow 1) and the lesion edges (shown by arrow 2).

(a)

(b)

Figure 3.3 (a) 2D image of psoriasis lesions (b). 3D surface of psoriasis lesions is scanned by a 3D laser scanner.

### 3.3.2 Light Scattering and Speckle Imaging Methods

Another technology implemented for surface roughness measurement is light scattering and speckle imaging. In these technologies, a light beam is projected to the scanned surface. The beam consists of many light paths. The reflected lights are then photographed by a 2D camera. The roughness differences are determined based on the characteristic of reflected lights. Figure 3.4 shows the reflected light of two surfaces with different roughness.


Figure 3.4 The light is uniformly reflected at (a) smooth surface whereas (b) the rough surfaces will scatter the light to various directions.

In the light scattering method, the smooth surface reflects the lights in uniform direction to the camera. In the other hand, rough surface will return back the projected lights at unordered manner. The intensity of reflected light from smooth surface will be higher than the rough surface. Photo sensor at smooth surface can receive reflected light at higher intensity. It can occur because all of the lights are reflected by smooth surface with the same reflection angles. The intensity received by photo sensor at rough surface will be lower. The reflected lights are scattered due to unordered normal direction of the rough surface. Surface roughness is then interpreted based on these reflected light characteristics.

Persson illustrates the distribution of reflected light from different surface roughness - smoother and rougher surfaces [81]. The light is emitted perpendicular onto the measured surfaces. The reflection angles and reflected light intensities are then recorded. To provide a reference value, surface roughness of the observed subject is determined by root mean square (rms)-roughness. The smoother and the rougher surfaces are represented by rms-roughness of 50 nm and 300 nm ,
respectively. The surface is made of stainless steel and it is illuminated with laser light $\lambda=700 \mathrm{~nm}$. From the experiment, it can be found that reflected light dispersion of the rough surface is higher than the smoother surface. The rough surface will scatter the reflected light to more various directions. Histogram of reflection light intensity against reflection angles are created to find the light dispersion characteristics. The intensities of reflected light from smoother surface are accumulated in a certain angle. For the rougher surface, the reflected light are scattered at various angles. Therefore a certain amount of light intensity can be found at all angles. The histograms of both surfaces are presented in Figure 3.5 (a) and (b).


Figure 3.5 Intensity distributions of reflected light from (a) smoother surface and (b) rougher surfaces.

As performed by light scattering, speckle imaging method also interprets surface roughness based on reflected illumination on the measured surface. However, in the speckle imaging, a coherent light is used to create spackle pattern of the rough surface. Average contrast of the speckle is also found related with the surface roughness [82]. Speckle is considered a random pattern region of bright and dark which is obtained when a surface is illuminated by a highly or partially coherent light beam [77]. Both methods, light scattering and speckle imaging can provide advantages on non-contact measurement and fast scanning [77]. Despite these advantages, the method has drawbacks in the 3D surface measurement. The drawbacks of this method are sophisticated system arrangement and standardised environment lighting. The measurement surface is usually limited to the small size area.

### 3.3.3 Structured Light Projection Method

Another available method for acquiring 3D surface is the structured light projection. This 3D measurement method has been used for many years in various researches and industrial applications [83]. The method utilises the projection of some particular structured patterns onto the measured surface. A series of the structured patterns can also be applied for the measurement. 2D images of the surface with several projected patterns are computed to reconstruct 3D surface. Depth information is interpreted from deviations of the projected patterns on the surface. Currently, some pixel-wise addressable projection systems on the basis of micro mirror projectors have been being used to create certain projected patterns. These projector systems in their application are able to generate various types of the pattern with high resolution. Furthermore, the application scope of the method has been expanded from measuring small size area to the large coverage area [83].

Principally, the structured light projection also uses a triangulation process as applied in the laser profilometry. This method also projects a light beam to the scanned surface, as conducted by scattering method. However, the structured projected light illuminates a structured pattern, rather than speckle imaging that illuminates random of lights. A difference between both of methods lies on the way of implementing the light profiling. Laser triangulation, as aforementioned in the previous section, uses a single beam to profile the surface. By applying this method, the surface elevations are able to be measured point by point. However, the structured light projection measures the surface elevations of the whole area at once. The area of projected pattern is composed by an equidistant of several line triangulations. The scanning is simplified by projecting the lines at the same time. Therefore, the movement of either the optical sensor and the measured object are not needed [83].

In this research, a 3D optical scanner is built by applying a structured light projection method. Thus, the details of this method are more widely elaborated. Figure 3.6 shows the basic diagram of camera and projector arrangement in the structured light projection. The projector and the camera are adjusted at fixed positions. These positions are indicated by the projection angle $\alpha$, the distance of the
projector to the camera, and the distance of the camera to the reference measured surface. The principle of 3D surface acquisition is mentioned in the following section as mentioned in [84] [85]. Phase-shifting method has been widely applied in 3D surface measurement in consideration of its scanning speed and high accuracy. Initially, three images are required to apply a phase-shifting algorithm, which therefore is called as the single three-step phase-shifting algorithm. These images are obtained from some projected fringe patterns on the measured surfaces and differentiated with a phase shift angle $\theta$. These three images can be expressed by equations (3-1), (3-2), and (3-3).

$$
\begin{gather*}
I_{1}(x, y)=I_{0}(x, y)+I_{\text {mod }}(x, y) \cos (\phi(x, y)-\theta)  \tag{3-1}\\
I_{2}(x, y)=I_{0}(x, y)+I_{\text {mod }}(x, y) \cos (\phi(x, y))  \tag{3-2}\\
I_{3}(x, y)=I_{0}(x, y)+I_{\text {mod }}(x, y) \cos (\phi(x, y)+\theta) \tag{3-3}
\end{gather*}
$$

Variables $I_{1}(x, y), I_{2}(x, y)$, and $I_{3}(x, y)$ denote the intensities of three fringe patterns. $I_{0}(x, y)$ represents the background intensity and $I_{\text {mod }}(x, y)$ represents the modulation signal amplitude. Meanwhile, variables $\phi(x, y)$ and $\theta$ are the phase and the constant phase-shift angle, respectively. The phase map $\phi(x, y)$ is calculated by applying three fringe images as written in (3-4).

$$
\begin{equation*}
\phi(x, y)=\tan ^{-1}\left[\sqrt{3} \frac{I_{1}(x, y)-I_{3}(x, y)}{2 I_{2}(x, y)-I_{1}(x, y)-I_{3}(x, y)}\right] \tag{3-4}
\end{equation*}
$$

The phase $\phi(x, y)$ calculated in (3-5) will have a range from $-\pi$ to $+\pi$ and discontinuity points at multiples of $2 \pi$. These discontinuities are located and then discarded by adding or subtracting the phase $\phi(x, y)$ with multiples of $2 \pi$. The equation for removing discontinuity is given by (3-5).

$$
\begin{equation*}
\Phi(x, y)=\phi(x, y)+2 k \pi \tag{3-5}
\end{equation*}
$$

The $\phi(x, y)$ and $\Phi(x, y)$ are known as the unwrapped and wrapped images, respectively and variable $k$ denotes the projection period. The elevation map in 3D coordinates is determined based on the phase difference between the measured
surface $\Phi(x, y)$ and reference surface $\Phi_{0}(x, y)$. The phase of reference surface $\Phi_{0}(x, y)$ is obtained from the projected pattern images on the flat surface. From Figure 3.6, it can be derived an equation for determining height $Z$ as shown in (3-6)

$$
\begin{equation*}
\frac{Z}{L-Z}=\frac{d}{B} \tag{3-6}
\end{equation*}
$$



Figure 3.6 Diagram for depth calculation based on phase differences.

Equation (3-6) is then simplified to obtain the following equations:

$$
\begin{gather*}
Z=\frac{L-Z}{B} d  \tag{3-7}\\
Z B=d L-d Z  \tag{3-8}\\
Z(B+d)=d L  \tag{3-9}\\
Z=\frac{d L}{(B+d)} \tag{3-10}
\end{gather*}
$$

Since $d \ll B$, it can be considered that $B+d \approx B$. This simplification can give

$$
\begin{equation*}
Z \approx \frac{L}{B} d \tag{3-11}
\end{equation*}
$$

It is known that $d$ is directly proportional to the phase difference, $d \propto\left(\Phi-\Phi_{0}\right)$, and then the equation (3-11) is modified to obtain a new expression (3-12).

$$
\begin{equation*}
Z \approx \frac{L}{B}\left(\Phi-\Phi_{0}\right) \tag{3-12}
\end{equation*}
$$

According to this method, the measured skin surface is exposed by a structured fringe-shaped light projection. The light is projected by applying projection angle $\alpha$ from the vertical axis. Since the scan process is quite fast, it can deal with the constant vibration of skin surface. This movement is caused by the vibration of the autonomous nervous system. Despite the method advantages, there are some limitations in the 3D surface acquisition. These limitations can be described such as measurement resolution and shadow problems. The measurement resolution depends on the projector resolution and the CCD density of the camera [86]. The method also cannot measure accurately the shadowed area and isolated surfaces [87]. These locations are usually found either as deep pit or localised peak surfaces.

### 3.4 Surface Profile Characterisation

The surface characterisation has been widely used in diverse application fields, such as road monitoring, material industry, remote sensing, and medical engineering. ISO 14460-1 defines the surface as a set of features which physically exist and separate the entire work piece from the surrounding medium [88]. A surface can be considered as a boundary layer between object and its environment. Normal profile, perpendicular to the reference surface, is used to extract the vertical variations of surface profile. The profile can be classified into three main components; namely roughness, waviness and form [88], as depicted in Figure 3.7 [89].


Figure 3.7 Surface profile is composed by three major components - roughness, waviness, and form [89] [90] [78].

These components are superimposed to form a surface profile. The surface wavelength or peak-to-peak spacing is used to differentiate a number of surface profile components [88]. Roughness has the finer texture and ridges compared to waviness, which is smoother than the roughness. Form, meanwhile, is a curved surface with a very long-range deviation.

The profile components are distinguished based on their wavelength. The roughness is the component with short wavelengths whereas the longer ones belong to the waviness and form [90]. ISO 115621, a standard for the manufacturing application, defines the separation point between roughness and waviness based on its wavelengths. Figure 3.8 depicts this component separation. The distribution of the roughness wavelength is denoted by $R$ whereas waviness wavelength is represented by $W$. The wavelengths of roughness profile range from $\lambda_{1}$ to $\lambda_{2}$ in which the shorter wavelengths $\left(\lambda<\lambda_{1}\right)$ exist in the surface. However, its quantity is not significant and can be neglected. The waviness profile has longer wavelengths ranging from $\lambda_{2}$ to $\lambda_{3}$ in which wavelength $\lambda_{2}$ is assigned to a wavelength that has the proportion of 50:50 for its roughness and waviness. The $\lambda_{3}$, meanwhile, is used to separate the waviness and form profiles [78].


Figure 3.8 Wavelength distributions of roughness and waviness [78].
The characterization and separation methods of surface profile are therefore important in a surface determination. These methods are implemented as a filtering process to the measured data. There are several filtering methods that can be applied to separate the surface components [90] such as Fourier transform, Gaussian filter, wavelet, and polynomial surface fitting.

In the earliest years, surface roughness filtering was performed manually based on a graphical plot on the 2 -dimensional coordinates. In this case, the plot is divided equally into several segments. A mean line is subsequently created for each segment. The line is drawn to represent the gradient between the segment intervals. Vertical deviations are then determined by subtracting the actual profile to the gradient line. Nowadays, some computational methods are widely applied to separate the profile components. The main objective of surface characterisation is to extract and to eliminate the waviness component. This waviness can be filtered by applying some methods such as Fourier transform, Gaussian filter, wavelet filtering, and polynomial surface fitting.

### 3.4.1 Fourier Transform

Digital Fourier transform is used to filter the components of a surface profile. The method was initially explored by Raja and Radhakrishnan [91]. In a surface engineering, the surface is considered to be formed by certain sinusoidal functions with various amplitudes and frequencies. From aforementioned discussion, it is known that the surface profiles - roughness, waviness, and form - are differentiated based on their frequencies. A number of cut-offs frequencies are defined by a user to enable a surface profiles differentiation [91]. In this case, this power spectral density analysis has been applied to determine surface roughness of metal surface [92].

### 3.4.2 Gaussian Filter

Gaussian filter is one of the surface characterisation methods widely used in recent years. A single filter here can be applied to extract roughness and waviness components [91]. However, Gaussian filter possesses some drawbacks in its implementation. The first drawback is edge distortion that can be found in the edges of the filtered data. In addition, Gaussian filter cannot properly perform to filter a surface dominated by a large form. Even, it is not robust to deal with outliers [91].

### 3.4.3 Wavelet Filter

Principally, the aim of wavelet filtering is similar with Gaussian filter. Both filters
are used to characterise the surface profile components based on their wavelengths. Compared with Gaussian filter, the wavelet filter, however, has more advance capabilities. This filter can be used to analyze data at various resolution levels. For this, its filter is also known as the multi-scale filters. Figure 3.9 depicts the decomposed signals of a surface profile [91]. Diff1 is a signal component with the shortest wavelength (the highest frequency). The advantage of wavelet filter is its ability to decompose the profile component, specifically at certain wavelength. Used in many works on surface roughness determinations, the wavelet filtering method also has a drawback in which its process requires a high computational cost. Also, it finds difficult to define the required number of level in the filtering process.


Figure 3.9 Wavelength distributions of roughness and waviness [91].

### 3.4.4 Polynomial Surface Fitting

Polynomial surface fitting has been widely used in the 3D surface characterisation of various applications. The surface fitting determines a least square mean plane. The plane here represents the nominal direction of a surface. A quadratic polynomial surface is the suitable approximation of common geometrical shapes in a small area. Further, the polynomial surface fitting is widely used to fit various surface shapes, such as partial cylindrical forms, partial spheres, and any arbitrary curved forms. Determination on a suitable polynomial order, however, can result in a problem. This problem occurs when the surface fitting is applied to a surface containing some different curvatures [93] [94]. Polynomial surface fitting is usually fitted into a rectangular mesh of XYZ coordinates. Buxton et al. have found that the whole
human body curvature can be constructed using interconnection of rectangular meshes [95]. This fact shows that polynomial surface fitting is suitable to be applied to any part of human body surfaces.

In the early roughness measurement by using stylus profilometer, there are three terms to define the sample [78] including traversing, assessment, and sampling length. Traversing length is the crossing distance of the stylus over the measured surface. Assessment length is only a part of traversing length that would be assessed. Another shorter sample is then measured from the assessment length. This shorter part is called as the sampling length. Figure 3.10 shows this sample categorisation in term of one dimensional sampling.


Figure 3.10 Sample categorisation based on the coverage area of the sample.
In the research, this categorisation is also implemented. The traversing length corresponds to a maximum scanned area by the camera. Meanwhile, assessment length is similar with the ROI of the image. In the research, ROI is defined as a lesion area in the scanned image. In this research, sampling length, additionally, can be considered as the smallest subdivided area of the developed algorithm. The next chapter of this thesis will present the determination of the optimum sampling area for surface roughness algorithm.

### 3.5 Surface Roughness Parameter

Most of 3D surface parameters are extended from the earlier 2D parameters of surface roughness. These 3D parameters were initially proposed and listed in 1990s. The parameters consist of 14 parameters leading it to be known as "Birmingham 14" [88]. New parameters were included to the list afterward. The ASME B46.1 has
also included some 3D parameters in their standard. Currently, several 3D parameters are being viewed to be adopted as the international standards [90].

Surface parameter is divided into three main groups, each of which is differentiated based on the direction of the measured unit. The groups are including amplitude, frequency, and combination of amplitude-frequency based parameters. Meanwhile, the calculation of the amplitude parameter has applied the vertical distances of the surface data points. The frequency parameter is calculated based on the distances at horizontal direction. To accommodate a measurement from both directions, these two parameters are combined in order to provide an amplitudefrequency parameter. Figure 3.11 displays the distances used for surface roughness determination.


Figure 3.11 Three main groups of surface roughness parameters.

Several roughness parameters are classified as amplitude parameters such as average roughness $\left(S_{a}\right)$, root mean square roughness $\left(S_{q}\right)$, maximum profile peak height $\left(S_{p}\right)$, maximum profile valley height $\left(S_{v}\right)$, maximum height of the profile $\left(S_{t}\right)$, and average maximum height of the profile $\left(S_{z}\right)$. Here, average roughness $\left(S_{a}\right)$ is defined as the average of the absolute vertical deviations measured from the mean surface. Root mean square roughness $\left(S_{q}\right)$, meanwhile, is determined by applying root mean square to the profile measured from the mean surface. Maximum profile peak height $\left(S_{p}\right)$ additionally refers to the distance to the highest point of the profile measured from the reference surface. Maximum profile valley height $\left(S_{v}\right)$ and
maximum height of the profile $\left(S_{t}\right)$ are determined as the distance of the lowest point to the reference surface and the distance between the highest and lowest points of the surface profile, respectively. Lastly, the average maximum height of the profile refers to the average of the successive values of $S_{t, i}$ which is calculated over the area of the profile. The $S_{t, i}$ in this case is the maximum height of the profile within the sampling length of $i$-th. Usually, the measured area is segmented into several sampling areas. If the measurement is applied to the whole areas, $S_{z}$ will be equal to $S_{t}$. Figure 3.12 and equation (3-13) to (3-18) illustrate the surface profile and the formulas based on amplitude parameters. The profiles shown in the figures are extracted from a certain column of rough surface. Variable $L$ is used to represent sampling length of surface roughness measurement. Figure 3.12 (b) depicts a measurement that computed from four successive samples.

Average roughness ( $S_{a}$ ),

$$
\begin{equation*}
S_{a}=\frac{1}{M N} \sum_{i=1}^{M} \sum_{j=1}^{N}\left|z\left(x_{i}, y_{j}\right)\right| \tag{3-13}
\end{equation*}
$$

Root mean square roughness $\left(S_{q}\right)$,

$$
\begin{equation*}
S_{q}=\sqrt{\frac{1}{M N} \sum_{i=1}^{M} \sum_{j=1}^{N} z\left(x_{i}, y_{j}\right)^{2}} \tag{3-14}
\end{equation*}
$$

Maximum profile peak height $\left(S_{p}\right)$,

$$
\begin{equation*}
S_{p}=\max \left(z\left(x_{i}, y_{j}\right)\right) \tag{3-15}
\end{equation*}
$$

Maximum profile valley height $\left(S_{v}\right)$,

$$
\begin{equation*}
S_{v}=\left|\min \left(z\left(x_{i}, y_{j}\right)\right)\right| \tag{3-16}
\end{equation*}
$$

Maximum height of the profile $\left(S_{t}\right)$, and

$$
\begin{equation*}
S_{t}=\left|S_{p}-S_{v}\right| \tag{3-17}
\end{equation*}
$$

Average maximum height of the profile $\left(S_{z}\right)$.

$$
\begin{equation*}
S_{z}=\frac{S_{t, 1}+S_{t, 2}+\cdots+S_{t, N}}{N}=\frac{\sum_{k=1}^{N} S_{t, k}}{N} \tag{3-18}
\end{equation*}
$$



Figure 3.12 (a) Surface roughness parameters $S_{p}, S_{v}$, and $S_{t}$ are determined from an evaluation length $L$. (b) $S_{t}$ is computed from four sampling length, $L_{1}, L_{2}, L_{3}$, and $L_{4}$.

Furthermore, the mean peak spacing can be used to determine the surface roughness based on a horizontal distance. This parameter is denoted by variable of $S S_{m}$, which is calculated by averaging the distance between zero crossings of the profile. At this point, two conditions needs to be satisfied to specify a crossing as zero crossing. The first condition is the greater maximum profile height between two zero crossing compared to a threshold. The threshold is usually set at $10 \%$ of $S_{Z}$. The second condition is that the spacing between two zero crossing is greater than a threshold; typically it is $1 \%$ of the sampling area. Figure 3.13 and equation (3-19) describe the profile parts and the formulas used in determination of surface roughness parameters.

$$
\begin{equation*}
S S_{m}=\frac{S_{m, 1}+S_{m, 2}+\cdots+S_{m, N}}{N}=\frac{\sum_{k=1}^{N} S_{m, k}}{N} \tag{3-19}
\end{equation*}
$$

The determination of mean peak spacing is illustrated in Figure 3.13. Two zero crossings are specified to determine the interval of peak spacing. As mentioned earlier, the zero crossings are identified by satisfying two conditions. Point 1 is firstly identified as the first zero crossing point followed by positive slope. Point 2 satisfies the first condition since the maximum profile height between point 1 and 2 is greater than the threshold. However, point 2 cannot be used as the end of peak
spacing because the distance between point 1 and 2 is too close. Point 3 cannot be considered as endpoint because it is preceded by a valley lower than the threshold. Point 4 is preceded by a valley greater than the upper threshold but it is followed a peak lower than upper threshold. The same condition as point 4 is also found at point 5. Finally, point 6 can be considered as endpoint because it is preceded by a deep valley (between point 4 and 5) and followed by a peak higher than threshold. Now, the distance between point 1 and 6 is denoted as the first peak spacing. The similar conditions are also applied to find the other peak spacing. Figure 3.13 displays the next peak spacing as $S_{m, 2}$. The interval of $S_{m, 2}$ is started from point 6 and ended at point 7. The mean peak spacing is calculated by averaging several successive peak spacings.


Figure 3.13 The mean peak spacing is obtained by averaging several peak spacings.

Furthermore, a slope is used as a hybrid parameter that combines the amplitude and frequency parameters. The value is determined as the ratio of the vertical to the horizontal distances of two consecutive data points. To increase sensitiveness on the spacing of data points, six consecutive points are used in the slope determination. The average roughness is mostly used among the amplitude parameters that can represent the actual vertical deviation of surface, minimise impulse noises, and consider all data points in roughness determination. A research on tactile perception also shows that perceived coarseness by human on several paper surfaces is correlated ( $R^{2}=0.73$ ) with the surface roughness of the papers [96].

### 3.6 Imaging Methods for Skin Surface Roughness Measurement

As mentioned in the earlier section, the skin assessment based on human senses and perception can result in a subjective assessment. To obtain objective results, a system of assessment and measurement based on imaging technologies has been developed. This research is focused on skin surface roughness as a characterisation parameter of skin lesion. Objective measurements of skin surface roughness have been attempted by several methods. The method of the skin roughness measurement can be grouped into invasive and non invasive method [97].

In invasive method, an incision procedure is required to provide sample data. For skin roughness measurement, a piece of skin sample is cut from the patient body. The measurement will be performed separately from the human body. This procedure is widely known as skin biopsy. High resolution and tiny details of skin sample surface can be acquired by microscopic camera. The skin sample preparation is conducted by a plastic surgeon or dermatologist. A pathologist will handle the analysis part of the skin sample. Both processes require well trained medical personnel and may take 3 to 10 days for the results of the analysis [98]. For instance, the skin surface topography of wrinkles on elderly people is measured by light microscopy and scanning electron microscopy [99]. Here, the measurement will detect the loss skin elasticity at wrinkled area. The skin sample is obtained by performing skin biopsy on the skin sample [99]. Scanning Electron Microscopy [100] on skin sample is considered as an invasive method. The texture impression is then observed using microscope to photograph the microscopic pattern of skin ridges and furrows. In this case, the 3D features cannot be analysed through the microscope observation.

Non-invasive method does not require skin cutting to obtain the sample. This method is preferred by most of the patient. Measurement can be performed as either non-direct or direct measurement. In non-direct measurement, the equipment is not applied directly to the skin surface. The process requires a skin surface replica to represent the actual surface. Conversely, the direct measurement does not need skin replica. The equipment can be applied directly to the skin surface of the living person. This method is also known as in-vivo measurement.

Direct measurement method is applied in early developments of surface roughness measurement. A high precision profilometer using sharp stylus is used to obtain profile lines of skin surface replica [101] [102] [103]. Since 1962 the silicon rubber plastic has been used to make an impression of skin texture by Sarkany [104] [105]. The process of replica preparation is not simple. Two mould steps using silicon materials have to be completed to obtain a skin replica. The first step is to create negative impression of skin surfaces. In the second step, the negative impression is then used to cast a positive print of the evaluated surface. Since the process is performed manually, significant errors can be introduced in both replication steps. An experienced technician is required to prepare skin replica accurately. To complete the replication process, it can take few minutes (3 to 15 minutes) for creating negative impression of the skin surface and followed by several hours (3 to 5 hours) for printing its positive impression [106]. For another impression material, it can take 24 hours to harden the skin replica [101]. Even though profilometer machines can measure with high resolution, the measurement process itself is very slow. Therefore this method cannot be adapted in daily practice that requires simple, reliable, and fast procedures. For instance, scan speed of stylus profilometer as used by Gassmueller et al. [107] ranging from 0.15 to $1.00 \mathrm{~mm} / \mathrm{s}$ [108].

Light medium is applied to improve scan speed in surface roughness determination [109]. Three-dimensional (3D) imaging and computer vision algorithms are applied to enable a fast scanning and measurement at high resolutions. By using imaging technologies, characterisation of surface components will not be limited by the dimension of the profilometer stylus; instead, surface characterisation and filtering process of imaging technologies can be performed through computational operations.

By using a laser profilometer, light profiling can speed up the scanning process. Scan speed of optical profilometer used in [110] can be increased until $250 \mathrm{~mm} / \mathrm{s}$ [111], 250× greater that scan speed of stylus profilometer. By applying light profiling, either a direct or non-direct surface measurement can be performed on skin surface [112]. Skin replica can be provided for light profiling to ensure the scanned
surface is firm and stable. The measurement of laser scanner is initially applied in cosmetic industry - in which, as mentioned in [113], it is used to measure skin surface indicators. The cosmetic efficacy is evaluated based on skin surface indicators, such as fold reduction, skin smoothness, and skin tightness. Skin surface roughness is measured to evaluate those skin indicators. Laser profilometry has been used to determine surface roughness. In the research, the laser does not scan the skin surface directly but it scans the skin impressions of the actual skin surface. These impressions are made of silicon materials and measured at size $3 \mathrm{~mm} \times 3 \mathrm{~mm}$. This size is selected to maintain the resolution and reduce scanning time [113]. Skin roughness analysis is also performed to quantify skin aging. Initially, a skin replica is required to extract the skin texture. The skin replica, made of an araldite casting resin, is scanned to provide several quantitative parameters such as $R_{a}, R_{p}, R_{t}, R_{\max }$, and others [114].

Stripe projection method has been applied to determine in vivo skin surface roughness [115]. By measuring in vivo skin surface roughness, inaccuracy caused by skin replica is investigated [116]. A 3D optical scanner using fringe projection method is used to acquire 3D surfaces from the actual skin surface and skin replica. Both scans are compared to find the 3D surface with better quality. It is found that fringe projection method on the actual skin surface can extract surface profiles more detail. Fine structures of skin surfaces cannot be represented during the process of replica making.

In the research, the skin roughness parameter of a psoriasis lesion is measured by analysing a number of digital images of lesion samples. The roughness parameter of healthy and unhealthy skin can be characterised by roughness parameters such as $R_{a}, R_{q}$, and peak-peak height ( $R_{y}$ ) [117]. These parameters are extended to measure the 3D surface roughness.

Surface roughness also can be measured from the speckle texture of the surface. The image of speckle texture is projected from scattering of coherent light in a rough surface. The contrast of the speckle is found to be related to the surface roughness [118]. A number of the texture analysis methods, such as co-occurrence matrix [119] and fractal analysis [120], have been applied to determine the roughness from the
speckle images. In a recent research, Tchvialeva et al. [109] proposed a speckle contrast method in order to determine the surface roughness. The results, afterward, have been validated with the results obtained from a fringe projection method.

Several methods are available for skin roughness determination. However, there are uncertainties in the methods that have not been defined yet. These uncertainties include the method capability to perform measurement on the skin lesion surfaces at any body parts. The methods have not been validated at various locations of the body surface. The reported studies have been selective in the location of the body surface such as cheek [121], back hand \& upper buttock [122], facial area [123], volar forearm, lower back, thigh and lower leg [124]. It is pertinent that the measurement is reliable at any location of the body surface. In addition, it is found that most of the reported researches do not focus on surface roughness of marked lesion [121] [124] [123] whereas in clinical practice, surface roughness is not only measured to normal area but also to skin lesions. Table 3.1 summarises the comparison of 3D acquisition methods - mechanical surface profilometry, laser profilometry, light scattering, and structured light projection - that currently used for surface measurement.

Table 3.1 Comparison of 3D acquisition methods for surface measurement.

| Methods | Advantages | Disadvantages |
| :--- | :--- | :--- |
| Mechanical <br> surface <br> profilometry | - Measurement is <br> performed to the <br> object without any <br> prior treatment [80] | - The scan resolution of stylus profilometer is limited by <br> the stylus size [79]. If the size of stylus tip is bigger than <br> the vertical gaps then a number of vertical points might <br> not correctly be profiled by the stylus tip [80]. |
| - The measurement |  |  |
| can cover a wide |  |  |
| range from tens of |  |  |
| microns to ten of |  |  |
| mm [80] |  |  |$\quad$| - The scanning process can be time-consuming. To obtain |
| :--- |
| topographical information, a number of the consecutive |
| line scans are required. To reduce scanning time, the |
| sample area is limited to a small size only [79] [80] |
| - Contact stress of stylus tip can damage and scratch the |
| surface especially for the soft materials [80] |

Table 3.1 Comparison of 3D acquisition methods for surface measurement (continued).

| Methods | Advantages | Disadvantages |
| :---: | :---: | :---: |
| Light <br> Scattering and Speckle <br> Imaging | - The methods can provide non-contact measurement and fast scanning [77] | The method requires sophisticated system arrangement and standardised environment lighting. <br> - The measurement surface is usually limited to the small size area |
| Structured <br> Light <br> Projection | - By performing fast scanning, the method overcomes the scan problem on constant vibration of skin surface <br> - Scanned surface is acquired as a grid matrix at high resolution | - The resolution of the measurement is limited by the resolution of the projection <br> - And the imaging CCD arrays. Multiple projections and multiple viewing cameras can be used to improve the resolution of the measurement.[86] <br> - Elevation cannot be measured accurately at local shadows and isolated surfaces[87] |

### 3.7 Summary

Chapter 3 presents a comprehensive description on surface roughness measurement methods. A surface roughness analysis in early years has been described in the beginning sections. Manufacturing and metallurgical process applies a mechanical surface profilometry to quantify the product surfaces. The profilometer system, at this point, has utilised a sharp needle to extract the profile elevation at a high resolution. However, there are drawbacks in the method related to the physical size of the needle. A small needle tip might damage the surface whereas the bigger tip might lose the profile details with narrow gaps. Some imaging technologies are proposed to overcome these problems. Here, the imaging system measures the surface directly without any physical contact. In this chapter, well-known surface imaging methods are reviewed. The methods are laser profilometry, light scattering, speckle imaging, and structured light projection.

A laser beam is emitted and directed line by line to the scanned surface. The laser triangulation is used to determine the position of a target based on the reflected light. The surface elevation is linearly proportional to the reflected light deviations. Although the laser system can provide high resolution images, the process is considered to be time-consuming. Light scattering and speckle imaging are another methods applied to measure surface roughness. In these methods, a light beam is
projected to the scanned surface. Both methods utilise its reflection to interpret the surface roughness. These methods can provide non-contact measurement and fast scanning but they require sophisticated system arrangement and standardised environment lighting. A more advanced method of 3D surface measurement is the structured light projection. The method projects some particular structured patterns onto the measured surface. The surface elevation of the scanned surface is determined from the deviated patterns on the surface. This method can perform high resolution scanning at high speed scan.

Surface filtering and surface roughness parameters are defined to enable a surface roughness determination. Surface filtering, in this case, is performed to extract roughness profile from the waviness and form components. The filtering methods are available, such as Fourier transform, Gaussian filter, wavelet filtering, and polynomial surface fitting. Digital Fourier transform is applied to identify the roughness from its frequency domains. Furthermore, Gaussian filter is applied in order to smooth the surface profile. However, Gaussian filter also has some drawbacks such as edge distortion, unable to filter a surface with a large form, and unreliable to deal with the surface outliers. Wavelet filter is used to analyze data at multi-resolution levels. However, the filtering process not only requires high computational costs but also is difficult to define the number of decomposition levels. The polynomial surface fitting as the last filtering method has been widely used in 3D surface characterisation. The surface fitting determines the mean plane that represents the nominal direction of a surface. The fitting can any arbitrary curved forms including cylindrical and spheres. Several roughness parameters are defined to determine the surface roughness. The parameters are classified into three groups: amplitude, frequency, and combination of amplitude-frequency. Average roughness - an amplitude parameter - is considered as the most reliable parameter and can adopt the characteristics of the natural surfaces.

The skin assessment based on human senses and perception can result in a subjective assessment. Imaging technologies have been developed to enable objective assessment. The method of skin roughness measurement is grouped into invasive and non invasive methods. In invasive method, an incision procedure is
required and the measurement is performed separately from the human body. High resolution and tiny details of skin sample surface can be acquired by microscopic camera. The process requires well trained medical personnel and may several days for the results of the analysis. Non-invasive method does not require skin cutting to obtain the sample. This method is preferred by most of the patient. Measurement can be performed as either non-direct or direct measurement. In non-direct measurement, the process requires a skin surface replica to represent the actual surface. Conversely, the direct measurement can be applied directly to the skin surface of the living person. Direct measurement method is applied in early developments of surface roughness measurement. A high precision profilometer using sharp stylus is used to obtain profile lines of skin surface replica. The replica making process is not simple. Significant errors can be introduced in the skin replica due to manual preparation. Light medium is applied to improve scan speed and profiling accuracy in surface roughness determination. The light profiling can be applied as either direct or non-direct surface measurement. The measurement of laser scanner is initially applied in cosmetic industry to evaluate the cosmetic efficacy. Several quantitative parameters such as $R_{a}, R_{p}, R_{t}$, and $R_{\max }$ are used to represent surface roughness. Stripe projection method has been applied to determine in vivo skin surface roughness. By applying in vivo measurement, inaccuracy caused by skin replica can be minimised. Skin surface roughness is also measured from the speckle texture of the surface. The speckle contrast is related to the surface roughness. The texture analysis method is also used to determine the surface roughness from the speckle images. These mentioned methods have been applied for skin roughness determination. However, the methods have not been validated at various locations of the body surface. It is pertinent that the measurement is reliable at any location of the body surface. In addition, it is found that most of the previous works do not focus on surface roughness of marked lesion. In the research, a surface roughness algorithm based on polynomial surface fitting has been developed to enable objective measurement. The development of this surface roughness algorithm is elaborated in the next chapter.

## CHAPTER 4

DEVELOPMENT ON SURFACE ROUGHNESS ALGORITHM

Chapter 4 is to describe the development of surface roughness algorithm. This chapter consists of several sections. The first section discusses in detail about the developed surface roughness algorithm by applying a high order polynomial surface fitting. In this case, a surface waviness is filtered out from a rough curved surface using the application of this polynomial surface fitting. The rough surface will afterward be flattened and the remaining vertical deviations are used to determine surface roughness. The following section describes a validation study on the surface roughness algorithm that is conducted to determine the accuracy and total standard deviation. In the next section, these values are used to analyse the system performance. This chapter also includes an evaluation on the rotational invariance and the sample area determination. These studies aim to evaluate and to define the algorithm limitations. The final section of this chapter summarises all of the materials presented.

### 4.1 Surface Roughness as Scaliness Parameter for PASI Scoring

As mentioned in the earlier chapter, surface roughness is used as the measurable features to grading the psoriasis scaliness. These features are selected based on the surface appearances of the lesion with different scaliness scores. Rougher surfaces, which are caused by irregular stack of dead skin cells, are found at a psoriasis lesion with higher severities. Imaging modalities i.e. 3D optical scanner is now available to perform fast surface scan at high resolution.

Figure 4.1(a)-(c) describes the 3D correspondence between skin surface roughness and scaliness severities (scores). Figure 4.1(d) illustrates the roughness profiles of normal skin; i.e., lesions with score 1 and score 4 . In lesions with score 4,
large vertical deviations have been observed that are caused by excessive coarse scales irregularly stacked on the lesion surfaces. The scores are provided based on dermatologists' visual and tactile perceptions. In this research, an imaging method is developed to assess scaliness objectively and accurately.


Figure 4.1 3D surfaces of (a) normal skin, psoriasis lesion (b) score 1, and (c) score 4; (d) Rough profiles of normal skin (solid line), lesion score 1 (dotted line), and lesion score 4 (dashed line).

### 4.2 Surface Roughness Algorithm

The 3D surface roughness is determined by averaging the vertical deviations of the lesion surface. Because lesions appear on 3D curved surface (human body), the vertical deviations due to the lesion is determined by subtracting a lesion surface from an estimated waviness surface as shown in cross-sectional view in Figure 4.2.


Figure 4.2 The cross-sectional view of a skin lesion on normal skin surface.

A 3D optical scanner with a structured light projection method, namely PRIMOS (Phase shift Rapid In vivo Measurement of Skin) portable, is used to acquire some lesion surface images. This optical scanner is designed for 3D in vivo measurements of a microscopic and macroscopic skin surface structure [125]. The structured light projection method is applied by PRIMOS camera to obtain the 3D surface image. The method provides a number of advantages such as standardised capture distance and high-speed scan (<63 ms). A high-speed capture is important for a skin surface measurement due to the inevitable movements of the subject. The PRIMOS has a high-resolution 3D surface (spatial resolution: 0.0062 mm and depth resolution: 0.0040 mm ) in which its image is used as the input to the developed algorithm to determine the surface roughness.

A higher-order polynomial surface fitting is applied to the rough lesion surface to extract an estimated 3D waviness surface from the rough lesion surface. By subtracting the rough lesion surface from an estimated waviness surface, the vertical deviations of lesion surface can be exactly determined. The vertical deviations of a lesion surface are known as the deviation surface. The second and third order polynomials are applied in this work. These polynomial orders are suitable for the small surface areas in that the vertical undulation of the waviness surface is lesser [93].

### 4.2.1 Polynomial Surface Fitting

As the lesion surface is not always flat, a polynomial surface fitting is required to fit a lesion surface. The second and third order polynomials are applied in this work. These polynomial orders are suitable for some small surface areas [93] such as lesion. In the small area, the vertical undulation of the estimated waviness is less and can be accurately fitted with the second and third order polynomials. The general form of polynomials can be written as equations (4-1) and (4-2) [126]:

The second order polynomial:

$$
\begin{align*}
z_{2}(x, y)= & \left(a_{1} x^{2}+a_{2} x+a_{3}\right) y^{2}+\left(a_{4} x^{2}+a_{5} x+a_{6}\right) y+ \\
& \left(a_{7} x^{2}+a_{8} x+a_{9}\right) \tag{4-1}
\end{align*}
$$

The third order polynomial:

$$
\begin{align*}
z_{3}(x, y)= & \left(a_{1} x^{3}+a_{2} x^{2}+a_{3} x+a_{4}\right) y^{3}+\left(a_{5} x^{3}+a_{6} x^{2}+a_{7} x+a_{8}\right) y^{2} \\
& +\left(a_{9} x^{3}+a_{10} x^{2}+a_{11} x+a_{12}\right) y  \tag{4-2}\\
& +\left(a_{13} x^{3}+a_{14} x^{2}+a_{15} x+a_{16}\right)
\end{align*}
$$

To create a surface based on a polynomial equation, 9 and 16 polynomial coefficients are required for the second and third polynomials, respectively. The coefficient of determination $\left(R^{2}\right)$ is used to measure fittingness of polynomial fitting [127]. A good fit can be obtained if $R^{2}$ is in interval [0.9, 1.0]. The best fitting result of the polynomial orders is selected based on the highest $R^{2}$. The equation of $R^{2}$ is expressed in following equation:

$$
\begin{equation*}
R^{2}=1-\frac{\sum_{i=1}^{M} \sum_{j=1}^{N}\left(z\left(x_{i}, y_{j}\right)-w_{k}\left(x_{i}, y_{j}\right)\right)^{2}}{\sum_{i=1}^{M} \sum_{j=1}^{N}\left(z\left(x_{i}, y_{j}\right)-\bar{z}\right)^{2}} \tag{4-3}
\end{equation*}
$$

where $z\left(x_{i}, y_{j}\right)$ represents the elevation of lesion surface, $\bar{z}$ represents the elevation average and $w_{k}\left(x_{i}, y_{j}\right)$ is fitted value at $\left(x_{i}, y_{j}\right)$ using $k$-th order polynomial. The surface roughness is determined by using average roughness $\left(S_{a}\right)$ equation [128]. In this equation, the surface roughness is calculated by averaging the absolute vertical deviation of all data points. The average roughness, $S_{a}$, is defined as equation (4-4). Variable $e\left(x_{i}, y_{j}\right)$ denotes the vertical deviation of lesion surface at $\left(x_{i}, y_{j}\right)$.

$$
\begin{equation*}
S_{a}=\frac{1}{M N} \sum_{i=1}^{M} \sum_{j=1}^{N}\left|z\left(x_{i}, y_{j}\right)-w_{k}\left(x_{i}, y_{j}\right)\right|=\frac{1}{M N} \sum_{i=1}^{M} \sum_{j=1}^{N}\left|e\left(x_{i}, y_{j}\right)\right| \tag{4-4}
\end{equation*}
$$

### 4.2.2 Surface Roughness Calculation

The surface roughness algorithm is described by the flow chart shown in Figure 4.3. In this flow chart, the fitting process of the second and the third orders are performed separately in order to get the best fit based on $R^{2}$.


Figure 4.3 Flowchart of the surface roughness algorithm.
The detail explanation on surface roughness algorithm is described as follows:
a) Input a 3D lesion surface matrix, $Z_{0}(x, y)$, with size $M \times N$ (see Figure 4.4). The coordinates $x, y$, and its value $z(x, y)$ are then used to calculate unknown coefficients of polynomial equations (Equations (4-1) and (4-2)). The total data point in lesion surface is $M \times N$.
b) Divide the lesion surface into $2 \times 2$ subdivided surfaces that gives four subdivided surfaces, namely $D_{1}, D_{2}, D_{3}$, and $D_{4}$, as shown in Figure 4.4. Half division is applied symmetrically to each side of the lesion area, thus the size of each subdivided surface becomes $\frac{M}{2} \times \frac{N}{2}=L$. The number of surface division is set not larger than $2 \times 2$. It aims to reduce computational cost and provide a sufficient area for surface roughness determination.
c) For each subdivided surface, $D_{1}$ to $D_{4}$, determine the polynomial coefficients of the selected order from coordinates $x, y$, and $z(x, y)$ through a matrix inversion.

This inversion is performed separately for all subdivided surfaces. As described in (4-5), matrix $V$ contains the elements of the polynomial equation, the coefficients of polynomial equation are stored in matrix $A$, and matrix $Z$ represents the subdivided lesion surface.

$$
\begin{equation*}
V A=Z \quad A=V^{-1} Z \tag{4-5}
\end{equation*}
$$



Figure 4.4 The segmented rough surface of the lesion model and its $4 \times 4$ subdivided surfaces.
d) For the first stage, the second order polynomial is selected to find polynomial coefficients $A$. Once the coefficients have been determined, the estimated waviness surface can be constructed. The waviness is obtained by applying calculated polynomial equation with polynomial coefficients $A$ at the evaluated coordinate points (x, y). Equation (4-5) can be written as $V_{2} A_{2}=Z_{D_{1}}$ to denote a second order polynomial fitting at subdivided surface $D_{1}$.
e) The matrices elements of $V_{2}, A_{2}$ and $Z_{D_{1}}$ are given by the following forms. Here, variable $L$ refers to the total number of data points at a subdivided surface.

$$
\begin{gather*}
V_{2}=\left[\begin{array}{ccccccccc}
x_{1}^{2} y_{1}^{2} & x_{1} y_{1}^{2} & y_{1}^{2} & x_{1}^{2} y_{1} & x_{1} y_{1} & y_{1} & x_{1}^{2} & x_{1} & 1 \\
x_{2}^{2} y_{2}^{2} & x_{2} y_{2}^{2} & y_{2}^{2} & x_{2}^{2} y_{2} & x_{2} y_{2} & y_{2} & x_{2}^{2} & x_{2} & 1 \\
x_{3}^{2} y_{3}^{2} & x_{3} y_{3}^{2} & y_{3}^{2} & x_{3}^{2} y_{3} & x_{3} y_{3} & y_{3} & x_{3}^{2} & x_{3} & 1 \\
x_{4}^{2} y_{4}^{2} & x_{4} y_{4}^{2} & y_{4}^{2} & x_{4}^{2} y_{4} & x_{4} y_{4} & y_{4} & x_{4}^{2} & x_{4} & 1 \\
\cdots & \cdots & \cdots & \cdots & \cdots & \cdots & \cdots & \cdots & \cdots \\
x_{L}^{2} y_{L}^{2} & x_{L} y_{L}^{2} & y_{L}^{2} & x_{L}^{2} y_{L} & x_{L} y_{L} & y_{L} & x_{L}^{2} & x_{L} & 1
\end{array}\right]  \tag{4-6}\\
A_{2}=\left[\begin{array}{llllllll}
a_{1} & a_{2} & a_{3} & \cdots & a_{8} & a_{9}
\end{array}\right]^{T} \tag{4-7}
\end{gather*}
$$

$$
Z_{D_{1}}=\left[\begin{array}{lllll}
z_{D_{1}} & \left(x_{1}, y_{1}\right) & z_{D_{1}}\left(x_{2}, y_{2}\right) & z_{D_{1}}\left(x_{3}, y_{3}\right) & \cdots  \tag{4-8}\\
z_{D_{1}} & \left(x_{L}, y_{L}\right)
\end{array}\right]^{T}
$$

f) The matrix of polynomial coefficients $A_{2}$ is unknown and inversion $A_{2}=$ $V_{2}^{-1} Z_{D_{1}}$ is applied to determine its values then. The polynomial coefficients of $A_{2}$ are determined by applying equation (4-9). Moore-Penrose pseudoinverse algorithm is used to compute $V_{2}^{-1}$. This method is able to find the solution for inverting a non-square matrix.

$$
\left[\begin{array}{l}
a_{1}  \tag{4-9}\\
a_{2} \\
a_{3} \\
\cdots \\
a_{8} \\
a_{9}
\end{array}\right]=\left[\begin{array}{ccccccc}
x_{1}^{2} y_{1}^{2} & x_{1} y_{1}^{2} & y_{1}^{2} & \cdots & x_{1}^{2} & x_{1} & 1 \\
x_{2}^{2} y_{2}^{2} & x_{2} y_{2}^{2} & y_{2}^{2} & \cdots & x_{2}^{2} & x_{2} & 1 \\
x_{3}^{2} y_{3}^{2} & x_{3} y_{3}^{2} & y_{3}^{2} & \cdots & x_{3}^{2} & x_{3} & 1 \\
x_{4}^{2} y_{4}^{2} & x_{4} y_{4}^{2} & y_{4}^{2} & \cdots & x_{4}^{2} & x_{4} & 1 \\
\cdots & \cdots & \cdots & \cdots & \cdots & \cdots & \cdots \\
x_{L}^{2} y_{L}^{2} & x_{L} y_{L}^{2} & y_{L}^{2} & \cdots & x_{L}^{2} & x_{L} & 1
\end{array}\right]^{-1} \quad\left[\begin{array}{c}
z_{D_{1}}\left(x_{1}, y_{1}\right) \\
z_{D_{1}}\left(x_{2}, y_{2}\right) \\
z_{D_{1}}\left(x_{3}, y_{3}\right) \\
z_{D_{1}}\left(x_{4}, y_{4}\right) \\
\cdots \\
z_{D_{1}}\left(x_{L}, y_{L}\right)
\end{array}\right]
$$

g) Waviness surface $W_{2}(x, y)$ (Figure 4.5-b) is estimated by applying the second order polynomial surface fitting with coefficients $A_{2}$. The estimation is limited for $(x, y)$ coordinates at $D_{1}$ area. Equations for determining $W_{2}(x, y)$ are shown by (4-10) and (4-11).

$$
\begin{gather*}
W_{2}=V_{2} A_{2}  \tag{4-10}\\
{\left[\begin{array}{c}
w_{2}\left(x_{1}, y_{1}\right) \\
w_{2}\left(x_{2}, y_{2}\right) \\
w_{2}\left(x_{3}, y_{3}\right) \\
w_{2}\left(x_{4}, y_{4}\right) \\
\ldots \\
w_{2}\left(x_{L}, y_{L}\right)
\end{array}\right]=\left[\begin{array}{ccccccc}
x_{1}^{2} y_{1}^{2} & x_{1} y_{1}^{2} & y_{1}^{2} & \cdots & x_{1}^{2} & x_{1} & 1 \\
x_{2}^{2} y_{2}^{2} & x_{2} y_{2}^{2} & y_{2}^{2} & \cdots & x_{2}^{2} & x_{2} & 1 \\
x_{3}^{2} y_{3}^{2} & x_{3} y_{3}^{2} & y_{3}^{2} & \cdots & x_{3}^{2} & x_{3} & 1 \\
x_{4}^{2} y_{4}^{2} & x_{4} y_{4}^{2} & y_{4}^{2} & \cdots & x_{4}^{2} & x_{4} & 1 \\
\cdots & \cdots & \cdots & \cdots & \cdots & \cdots & \cdots \\
x_{L}^{2} y_{L}^{2} & x_{L} y_{L}^{2} & y_{L}^{2} & \cdots & x_{L}^{2} & x_{L} & 1
\end{array}\right] \times\left[\begin{array}{c}
a_{1} \\
a_{2} \\
a_{3} \\
\cdots \\
a_{8} \\
a_{9}
\end{array}\right]} \tag{4-11}
\end{gather*}
$$

h) Deviation surface, $E_{2}(x, y)$ (Figure $4.5-\mathrm{c}$ ), is determined by subtracting the estimated waviness, $W_{2}(x, y)$, from the lesion surface $Z_{D_{1}}(x, y)$. The equation is $E_{2}(x, y)=\left|Z_{D_{1}}(x, y)-W_{2}(x, y)\right|$.
i) The coefficient of determination $\left(R^{2}\right)$ is calculated to evaluate the fitting result. Here, $W_{2}(x, y)$ is accepted if $R^{2}$ is within [0.9, 1.0]. The equation for determining $R^{2}$ is provided in (4-3). Notation $R_{2}^{2}\left(D_{1}\right)$ is used to denote the $R^{2}$ of the second order polynomial surface fitting at the subdivided surface $D_{1}$. Equation (4-3) is used to determine this coefficient.

$$
\begin{equation*}
R_{2}^{2}\left(D_{1}\right)=1-\frac{\sum_{i=1}^{\frac{M}{2}} \sum_{j=1}^{\frac{N}{2}}\left(z_{D_{1}}\left(x_{i}, y_{j}\right)-w_{2}\left(x_{i}, y_{j}\right)\right)^{2}}{\sum_{i=1}^{\frac{M}{2}} \sum_{j=1}^{\frac{N}{2}}\left(z_{D_{1}}\left(x_{i}, y_{j}\right)-\overline{z_{D_{1}}}\right)^{2}} \tag{4-12}
\end{equation*}
$$

j) Equation (4-4) is used to determine the surface roughness $S_{a}$ at the subdivided surface $D_{1}$. The input variables for this equation are deviation surface $E_{2}(x, y)$ with matrix size $\frac{M}{2} \times \frac{N}{2}$.

$$
\begin{equation*}
S_{a, 2}\left(D_{1}\right)=\frac{1}{\left(\frac{M}{2} \times \frac{N}{2}\right)} \sum_{i=1}^{\frac{M}{2}} \sum_{j=1}^{\frac{N}{2}}\left|z_{D_{1}}\left(x_{i}, y_{j}\right)-w_{2}\left(x_{i}, y_{j}\right)\right| \tag{4-13}
\end{equation*}
$$

Notation $S_{a, 2}\left(D_{1}\right)$ is used to represent the surface roughness of the subdivided surface $D_{1}$ that is determined by the second order polynomial surface fitting.
k) Repeat step (d) to (j) to compute the surface roughness values for other subdivided surfaces, $D_{2}$ to $D_{4}$. These computations are performed separately for each subdivided surface. The overall surface roughness of lesion surface is obtained by averaging the surface roughness of subdivided surfaces. Here, the surface roughness of a subdivided surface will not be included in final calculation if $R^{2}$ of polynomial surface fitting is not within an acceptable interval [ $0.9,1.0]$. Thus, the overall surface roughness of lesion surface can be expressed as

$$
\begin{equation*}
\overline{S_{a, 2}}=\frac{\sum_{i=1}^{N_{D}} S_{a, 2}\left(D_{i}\right)}{N_{D}}, \text { if } 0.9 \leq R_{2}^{2}\left(D_{i}\right) \leq 1.0, N_{D} \leq 4 \tag{4-14}
\end{equation*}
$$

The overall coefficient of determination $R^{2}$ is computed as well by using equation (4-15).

$$
\begin{equation*}
\overline{R_{2}^{2}}=\frac{\sum_{i=1}^{N_{D}} R_{2}^{2}\left(D_{i}\right)}{N_{D}} \text {, if } 0.9 \leq R_{2}^{2}\left(D_{i}\right) \leq 1.0, N_{D} \leq 4 \tag{4-15}
\end{equation*}
$$

1) Perform some similar steps as step 3 to step 11 but, instead of by fitting to a second order polynomial, use the third order polynomial surface fitting to estimate a waviness surface. For the implementation of third order polynomial to the subdivided surface $D_{1}$, equation (4-5) can be updated as $V_{3} A_{3}=Z_{D_{1}}$.
m) Matrices elements of $V_{3}$ and $A_{3}$ are arranged as in the following forms. There is no difference in the input data $Z_{D_{1}}$ compared to the previous steps. Variable $L$ here represents the total number of data points of the subdivided surface $D_{1}$.

$$
\begin{align*}
& V_{3}=\left[\begin{array}{ccccccccccccc}
x_{1}^{3} y_{1}^{3} & x_{1}^{2} y_{1}^{3} & x_{1} y_{1}^{3} & y_{1}^{3} & \ldots & x_{1}^{3} y_{1} & x_{1}^{2} y_{1} & x_{1} y_{1} & y_{1} & x_{1}^{3} & x_{1}^{2} & x_{1} & 1 \\
x_{1}^{3} y_{1}^{3} & x_{2}^{2} y_{3}^{3} & x_{2} y_{3}^{3} & y_{3}^{3} & \ldots & x_{2}^{3} y_{2} & x_{2}^{2} y_{2} & x_{2} y_{2} & y_{2} & x_{2}^{3} & x_{2}^{2} & x_{2} & 1 \\
x_{3}^{3} y_{3}^{3} & x_{3}^{2} y_{3}^{3} & x_{3} y_{3}^{3} & y_{3}^{3} & \ldots & x_{3}^{3} y_{3} & x_{3}^{2} y_{3} & x_{3} y_{3} & y_{3} & x_{3}^{3} & x_{3}^{2} & x_{3} & 1 \\
x_{4}^{3} 4_{4}^{3} & x_{4}^{2} y_{4}^{3} & x_{4} y_{4}^{3} & y_{4}^{3} & \ldots & x_{4}^{3} y_{4} & x_{4}^{2} y_{4} & x_{4} y_{4} & y_{4} & x_{4}^{3} & x_{4}^{2} & x_{4} & 1 \\
x_{1}^{3} y_{N}^{3} & x_{N}^{2} y_{N}^{3} & x_{N} y_{N}^{3} & y_{N}^{3} & \ldots & x_{N}^{3} y_{N} & x_{N}^{2} y_{N} & x_{N} y_{N} & y_{N} & x_{N}^{3} & x_{N}^{2} & x_{N} & 1
\end{array}\right]  \tag{4-16}\\
& A_{3}=\left[\begin{array}{llllll}
a_{1} & a_{2} & a_{3} & \cdots & a_{15} & a_{16}
\end{array}\right]^{T} \tag{4-17}
\end{align*}
$$

n) The Moore-Penrose inversion of $A_{3}=V_{3}^{-1} Z_{D_{1}}$ is applied to determine the polynomial coefficients $A_{3}$ and its equation is given in (4-18).

$$
\left[\begin{array}{c}
a_{1}  \tag{4-18}\\
a_{2} \\
a_{3} \\
\cdots \\
a_{15} \\
a_{16}
\end{array}\right]=\left[\begin{array}{ccccccccc}
x_{1}^{3} y_{1}^{3} & x_{1}^{2} y_{1}^{3} & x_{1} y_{1}^{3} & y_{1}^{3} & \cdots & x_{1}^{3} & x_{1}^{2} & x_{1} & 1 \\
x_{2}^{3} y_{2}^{3} & x_{2}^{2} y_{2}^{3} & x_{2} y_{2}^{3} & y_{2}^{3} & \cdots & x_{2}^{3} & x_{2}^{2} & x_{2} & 1 \\
x_{3}^{3} y_{3}^{3} & x_{3}^{2} y_{3}^{3} & x_{3} y_{3}^{3} & y_{3}^{3} & \cdots & x_{3}^{3} & x_{3}^{2} & x_{3} & 1 \\
x_{4}^{3} y_{4}^{3} & x_{4}^{2} y_{4}^{3} & x_{4} y_{4}^{3} & y_{4}^{3} & \cdots & x_{4}^{3} & x_{4}^{2} & x_{4} & 1 \\
\cdots y_{N}^{3} & x_{N}^{2} y_{N}^{3} & x_{N} y_{N}^{3} & y_{N}^{3} & \cdots & \cdots & x_{N}^{3} & x_{N}^{2} & x_{N}
\end{array}\right] \quad 1 \quad \cdots \quad\left[\begin{array}{c}
z_{D_{1}}\left(x_{1}, y_{1}\right) \\
z_{D_{1}}\left(x_{2}, y_{2}\right) \\
z_{D_{1}}\left(x_{3}, y_{3}\right) \\
z_{D_{1}}\left(x_{4}, y_{4}\right) \\
\ldots \\
z_{D_{1}}\left(x_{L}, y_{L}\right)
\end{array}\right]
$$

o) Estimated waviness surface $W_{3}(x, y)$ is obtained by the substitution of coefficients $A_{3}$ into equation (4-19). Equation (4-20) shows the expanded form of equation (4-19).

$$
\begin{align*}
& W_{3}=V_{3} A_{3}  \tag{4-19}\\
& {\left[\begin{array}{c}
w_{3}\left(x_{1}, y_{1}\right) \\
w_{3}\left(x_{2}, y_{2}\right) \\
w_{3}\left(x_{3}, y_{3}\right) \\
w_{4}\left(x_{4}, y_{4}\right) \\
\ldots \\
w_{16}\left(x_{L}, y_{L}\right)
\end{array}\right]=\left[\begin{array}{ccccccccc}
x_{1}^{3} y_{1}^{3} & x_{1}^{2} y_{1}^{3} & x_{1} y_{1}^{3} & y_{1}^{3} & \cdots & x_{1}^{3} & x_{1}^{2} & x_{1} & 1 \\
x_{2}^{3} y_{2}^{3} & x_{2}^{2} y_{2}^{3} & x_{2} y_{2}^{3} & y_{2}^{3} & \cdots & x_{2}^{3} & x_{2}^{2} & x_{2} & 1 \\
x_{3}^{3} y_{3}^{3} & x_{3}^{2} y_{3}^{3} & x_{3} y_{3}^{3} & y_{3}^{3} & \cdots & x_{3}^{3} & x_{3}^{2} & x_{3} & 1 \\
x_{4}^{3} y_{4}^{3} & x_{4}^{2} y_{4}^{3} & x_{4} y_{4}^{3} & y_{4}^{3} & \cdots & x_{4}^{3} & x_{4}^{2} & x_{4} & 1 \\
\cdots & \cdots & \cdots & \cdots & \cdots & \cdots & \cdots & \cdots & \ldots \\
\cdots & \ldots \\
x_{N}^{3} & x_{N}^{2} y_{N}^{3} & x_{N} y_{N}^{3} & y_{N}^{3} & \cdots & x_{N}^{3} & x_{N}^{2} & x_{N} & 1
\end{array}\right] \times\left[\begin{array}{c}
a_{1} \\
a_{2} \\
a_{3} \\
\cdots \\
a_{15} \\
a_{16}
\end{array}\right]} \tag{4-20}
\end{align*}
$$

p) By subtracting the estimated waviness $W_{2}(x, y)$ from the lesion surface $Z_{D_{1}}(x, y)$, the deviation surface $E_{3}(x, y)$ is determined. The equation of this subtraction is $E_{3}(x, y)=\left|Z_{D_{1}}(x, y)-W_{3}(x, y)\right|$.
q) The coefficient of determination and surface roughness of the measured subdivided surface $D_{1}$ can be given by (4-21) and (4-22).

$$
\begin{align*}
& R_{3}^{2}\left(D_{1}\right)=1-\frac{\sum_{i=1}^{\frac{M}{2}} \sum_{j=1}^{\frac{N}{2}}\left(z_{D_{1}}\left(x_{i}, y_{j}\right)-w_{3}\left(x_{i}, y_{j}\right)\right)^{2}}{\sum_{i=1}^{\frac{M}{2}} \sum_{j=1}^{\frac{N}{2}}\left(z_{D_{1}}\left(x_{i}, y_{j}\right)-\overline{z_{D_{1}}}\right)^{2}}  \tag{4-21}\\
& S_{a, 3}\left(D_{1}\right)=\frac{1}{\left(\frac{M}{2} \times \frac{N}{2}\right)} \sum_{i=1}^{\frac{M}{2}} \sum_{j=1}^{\frac{N}{2}}\left|z_{D_{1}}\left(x_{i}, y_{j}\right)-w_{3}\left(x_{i}, y_{j}\right)\right| \tag{4-22}
\end{align*}
$$

r) Determine $S_{a, 3}$ and $R_{3}^{2}$ for the remain subdivided surfaces, $D_{2}, D_{3}$, and $D_{4}$. Hence, the overall surface roughness and coefficient of determination are determined by (4-23) and (4-24).

$$
\begin{gather*}
\overline{S_{a, 3}}=\frac{\sum_{i=1}^{N_{D}} S_{a, 3}\left(D_{i}\right)}{N_{D}}, \text { if } 0.9 \leq R_{3}^{2}\left(D_{i}\right) \leq 1.0, N_{D} \leq 4  \tag{4-23}\\
\overline{R_{3}^{2}}=\frac{\sum_{i=1}^{N_{D}} R_{3}^{2}\left(D_{i}\right)}{N_{D}}, \text { if } 0.9 \leq R_{3}^{2}\left(D_{i}\right) \leq 1.0, N_{D} \leq 4 \tag{4-24}
\end{gather*}
$$

s) Coefficient $\overline{R_{2}^{2}}$ and $\overline{R_{3}^{2}}$ are compared to decide the best order of the polynomial surface fitting. The overall surface roughness $\overline{S_{a}}$ is obtained from the calculation that can give a higher $\overline{R^{2}}$ value. For example, if $\overline{R_{2}^{2}}$ is found to be higher than $\overline{R_{3}^{2}}, \overline{S_{a, 2}}$ is considered as the overall surface roughness. Consequently, the surface roughness of $\overline{S_{a, 3}}$ would be discarded. A subtraction between a lesion surface and an estimated waviness is shown in Figure 4.5. This subtraction yields deviation surface. The average roughness equation is applied to the deviation surface to compute the lesion surface roughness.


Figure 4.5 The 3D surfaces involved in surface roughness determination: (a) lesion surface, (b) estimated waviness, and (c) deviation surface.

Higher polynomial orders (higher than 3rd order) are not selected in estimating waviness surface. This limitation aims to avoid over fitting on rough profile of the lesion surface [129]. In over fitting condition, the estimated waviness tends to follow the actual profile of lesion surface. Therefore, the fitted surface does not measure the vertical deviations accurately. Figure 4.6 shows comparison of the waviness profiles that have been estimated by applying surface fitting with several orders. It can be shown that fitting lines of second (red line) and third orders (green line) are able to follow the rough profile curvature better than fourth and fifth orders.


Figure 4.6 Rough surface and fitted profiles of a lesion model. The first, second, third and fourth orders are applied to determine the fitted profiles.

These surface and fitted profiles are extracted from a lesion surface. The size of lesion surface is $75 \times 69$ pixels representing $4.710 \times 4.326 \mathrm{~mm}^{2}$. The profile line is located parallel with Y -axis at point $x$ equals 50 pixel $(3.150 \mathrm{~mm}$ from the vertical edge of the 3D surface). The fourth and the fifth orders are not able to fit properly on the lesion surface. The fitting process is trying to fit undulated profiles at centre area. Therefore, the edges of fitted surface are released freely and not set to follow the lesion curvature. Finally, the fitting error are minimised at centre area but it becomes very large at the edges. Figure 4.7 (a) and (b) demonstrate the success of the second and third order polynomial surface fitting implementations. The lesion in this figure is the same lesion that has been mentioned in previous plot (Figure 4.8).

The estimated and deviation surfaces are shown for each fitting order implementation. The fitting results of fourth and fifth orders are shown in Figure 4.7 (c) and (d). Their waviness estimation cannot be used in surface roughness determination because the the estimated waviness does not follow mean curvature of the lesion surface. The coefficient of determination $\left(R^{2}\right)$ of both calculations are less than 0.90 .


Figure 4.7 (a) 3D surface of a lesion model. (b) - (e) The estimated waviness and deviation surfaces that are determined by applying several different fitting orders.

### 4.3 Validation Study

### 4.3.1 Surface Roughness Measurement: Abrasive Paper

Several abrasive papers with several different roughness grades have been tested to validate the surface roughness measurement on various rough surfaces. There are five grades here used for the validation, those are $16,24,60,80$, and 280 . Grade value $(G)$ follows the standard of United States CAMI (Coated Abrasive Manufacturers Institute) [130]. The grade is inversely proportional with the surface
roughness. Rougher surfaces are valued by smaller grades. Notation $1 / G$ is usually used to relate linearly proportional to the surface roughness. A total of 72 3D surfaces are scanned for each grade. Since there are five grades evaluated in this study, the acquisition finally gives a total of 360 surfaces.

A surface roughness algorithm is applied to the collected 3D surfaces. Table 4.1 lists surface roughness and its standard deviation for each abrasive grade. As listed in this table, the surface roughness increases proportionally with $1 / G$. Average diameter is an average size of the abrading particles embedded in the abrasive paper. Figure 4.8, meanwhile, shows 3D surfaces of the abrasive papers. The surface images are ordered from the less rough to the very rough surfaces. Subsequently, several impulse noises appear in the rough surfaces as a result of the reflectivity of the abrasive particles. The scanner could not scan correctly on the shiny surface.


Figure 4.8 The 2D and 3D images of abrasive papers used in the surface roughness validation. The roughness grades are (a) 16, (b) 24, (c) 60, (d) 80, and (e) 280.

Table 4.1 The surface roughness of abrasive paper.

| Grade <br> Values $(G)$ | Average <br> diameter $(\mathrm{mm})$ | $1 / G$ | $\overline{S_{a}}(\mathrm{~mm})$ | $\sigma(\mathrm{mm})$ | $\overline{S_{a}}+\sigma$ <br> $(\mathrm{mm})$ | $\overline{S_{a}}-\sigma$ <br> $(\mathrm{mm})$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 280 | 0.044 | 0.004 | 0.0122 | 0.0007 | 0.0115 | 0.0128 |
| 80 | 0.192 | 0.013 | 0.0856 | 0.0090 | 0.0766 | 0.0946 |
| 60 | 0.268 | 0.017 | 0.1101 | 0.0380 | 0.0722 | 0.1481 |
| 24 | 0.715 | 0.042 | 0.2552 | 0.0336 | 0.2216 | 0.2888 |
| 16 | 1.320 | 0.063 | 0.4517 | 0.0555 | 0.3962 | 0.5072 |

The results of Table 4.1 are then plotted, as displayed in Figure 4.9. The relation between $1 / G$ and $S_{a}$ is proven by using Pearson's correlation coefficient. Its Pearson's coefficient is 0.989 showing a strong relation between them. A positive sign furthermore represents a linear relationship, which means that the smoother surface (small $S_{a}$ value) is represented by a small number of $1 / G$. This value proves that roughness algorithm can be used to determine a roughness grading. Since the surface roughness is an amplitude parameter, its value will relate to a particle size. The rougher surface is composed by particles with a bigger diameter size and results in a number of higher amplitude deviations. Therefore, the dependency of surface roughness on the size of abrading particles can be fitted as a linear line, as shown in Figure 4.9.


Figure 4.9 Plot of surface roughness vs. $1 / G$ (CAMI grade) obtained from measurement on abrasive papers.

### 4.3.2 Surface Roughness on Curve Surface: Mannequin Surface

To model a skin lesion, a medical tape that has regular and uniform texture on its surface is used. The tape has regular and uniform texture on its surface. It is made from an elastic material. Therefore, the tape can adapt the surface curvature when located onto any surfaces. However, to preserve its texture characteristics, the medical tape has to be pasted onto a smooth surface only. The average surface roughnesses $\overline{S_{a}}$ of 33 lesion models are used as the reference. These reference models are pasted at a hard paperboard flat surface. By applying lesion models on the flat surface, some vertical undulations caused by a curved surface would not contribute to the surface roughness measurement. Then, it could be ensured that only vertical deviations of the lesion model contribute to the roughness calculation. Figure 4.10 shows the lesion models at flat surface.


Figure 4.10 (a) Lesion models made of surgical tape are pasted on the flat surfaces of paperboard. (b) A 3D surface of lesion model.

In this experiment, the lesion model is scanned three times by applying a series of scans on the scanner mode. The scanning process of 33 lesion models has then given a total of 99 3D surface images. The surface roughness of a lesion model is represented by the average of surface roughness from these three consecutive scans. Additionally, the algorithm is applied separately for each 3D surface to obtain the surface roughness of the lesion models. Since the model preparation is manually performed, it results in a variety of sizes on the area of the lesion model. Table 4.2 shows the first 12 of 99 surface roughness measurements. In this case, the smallest and the largest areas are $10.06 \times 9.93 \mathrm{~mm}^{2}$ and $14.14 \times 10.69 \mathrm{~mm}^{2}$, respectively.

Table 4.2 The surface roughness of lesion models at a flat surface

| Lesion model | Scan index | $S_{a, 2}$ | $R_{2}^{2}$ | $S_{a, 3}$ | $R_{3}^{2}$ | $S_{\text {final }}$ | $R_{\text {final }}^{2}$ | $\overline{S_{\text {final }}}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | 1 | 0.0142 | 0.997 | 0.0138 | 0.997 | 0.0138 | 0.997 | 0.0139 |
|  | 2 | 0.0146 | 0.996 | 0.0142 | 0.997 | 0.0142 | 0.997 |  |
|  | 3 | 0.0140 | 0.996 | 0.0137 | 0.997 | 0.0137 | 0.997 |  |
| 2 | 1 | 0.0123 | 0.998 | 0.0120 | 0.998 | 0.0120 | 0.998 | 0.0130 |
|  | 2 | 0.0122 | 0.998 | 0.0119 | 0.998 | 0.0119 | 0.998 |  |
|  | 3 | 0.0151 | 0.997 | 0.0149 | 0.997 | 0.0149 | 0.997 |  |
| 3 | 1 | 0.0124 | 0.994 | 0.0122 | 0.994 | 0.0122 | 0.994 | 0.0122 |
|  | 2 | 0.0125 | 0.994 | 0.0122 | 0.994 | 0.0122 | 0.994 |  |
|  | 3 | 0.0123 | 0.994 | 0.0121 | 0.994 | 0.0121 | 0.994 |  |
| 4 | 1 | 0.0123 | 0.994 | 0.0119 | 0.994 | 0.0119 | 0.994 | 0.0118 |
|  | 2 | 0.0122 | 0.994 | 0.0118 | 0.994 | 0.0118 | 0.994 |  |
|  | 3 | 0.0123 | 0.994 | 0.0118 | 0.994 | 0.0118 | 0.994 |  |

The surface roughness of lesion models are summarised in a histogram as shown in Figure 4.11. From the histogram, average surface roughness $\overline{S_{a}}$ and its standard deviation are found to be $0.0122 \pm 0.0011 \mathrm{~mm}\left(\bar{x}_{R e f} \pm \sigma_{R e f}\right)$.


Figure 4.11 The histogram of lesion models pasted on flat surfaces.

To validate the surface roughness algorithm on the curved surfaces of human skin, the lesion models are placed on surfaces of mannequin that is made based on the body size of human at adult ages. In common, mannequin is used for medical purposes. The validation on mannequin uses a similar tape material with the
validation on the flat surface. As comparison, Figure 4.12 displays a lesion model in which the left figure shows a lesion model made of a surgical tape and the right figure shows the actual psoriasis lesion. Both of the objects are photographed at the same resolution.


Figure 4.12 Lesion model $\left(20 \times 11.5 \mathrm{~mm}^{2}\right)$ is made of surgical tape (left) and a psoriasis lesion (right). The images are photographed by PRIMOS camera.

A total of 390 lesion models are pasted and distributed onto several locations of the mannequin body. The lesion models are placed in the centre point of grid arrangement with size $40 \times 30 \mathrm{~mm}^{2}$. The centre point is selected to provide an accurate image because the highly focused area is located in this point. The region of the lesion models are defined following four regions (head, upper limb, trunk, and lower limb) of PASI scoring. Table 4.3 presents a list of the distribution of lesion models on body regions. Since each model requires a certain rectangular space, the number of lesions is limited by the available areas for each body region. Figure 4.13 shows the mannequin used in the validation study and a 3D surface of lesion model.


Figure 4.13 Lesion models are pasted onto life-size mannequin to simulate the lesions. A 2D image and 3D surface of lesion model are shown in the figure.

Table 4.3 The distribution of lesion models on the body regions of mannequin.

| Body region | Number of lesions |
| :--- | :---: |
| Head | 6 |
| Upper limb | 87 |
| Trunk | 159 |
| Lower limb | 138 |

The validation process determines the accuracy and total standard deviation of a roughness measurement on a 3D curved skin surface. The algorithm is considered valid if the measured surface roughness of tapes (lesion model) is constant at any locations on the skin surfaces. In this study, a surface of mannequin's skin is used to simulate the human skin surfaces. Table 4.4 presents several surface roughness values of the lesion models.

Table 4.4 The surface roughness of lesion models at a flat surface.

| Lesion <br> Index | Body <br> Region | Width <br> $(\mathrm{mm})$ | Height <br> $(\mathrm{mm})$ | $S_{a, 2}$ | $R_{2}^{2}$ | $S_{a, 3}$ | $R_{3}^{2}$ | $S_{\text {final }}$ | Pct. <br> Error <br> $(\%)$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 2 | Head | 10.05 | 9.04 | 0.014 | 0.996 | 0.015 | 0.995 | 0.014 | $14.8 \%$ |
| 6 | Head | 13.49 | 8.40 | 0.013 | 0.996 | 0.013 | 0.996 | 0.013 | $6.6 \%$ |
| 7 | Head | 9.54 | 7.64 | 0.014 | 0.998 | 0.014 | 0.998 | 0.014 | $14.8 \%$ |
| 4 | Upper Limb | 9.80 | 4.84 | 0.012 | 0.971 | 0.013 | 0.964 | 0.012 | $1.6 \%$ |
| 5 | Upper Limb | 9.67 | 16.42 | 0.012 | 0.998 | 0.012 | 0.998 | 0.012 | $1.6 \%$ |
| 6 | Upper Limb | 10.18 | 3.82 | 0.011 | 0.996 | 0.013 | 0.994 | 0.011 | $9.8 \%$ |
| 3 | Trunk | 15.01 | 7.77 | 0.012 | 0.944 | 0.015 | 0.905 | 0.012 | $1.6 \%$ |
| 4 | Trunk | 9.93 | 6.49 | 0.013 | 0.992 | 0.013 | 0.992 | 0.013 | $6.6 \%$ |
| 8 | Trunk | 12.22 | 7.51 | 0.012 | 0.977 | 0.016 | 0.962 | 0.012 | $1.6 \%$ |
| 2 | Lower Limb | 9.29 | 12.48 | 0.012 | 0.982 | 0.014 | 0.984 | 0.014 | $14.8 \%$ |
| 7 | Lower Limb | 10.18 | 5.73 | 0.013 | 0.959 | 0.014 | 0.950 | 0.013 | $6.6 \%$ |
| 8 | Lower Limb | 9.67 | 14.51 | 0.011 | 0.970 | 0.011 | 0.975 | 0.011 | $9.8 \%$ |

The surface roughness of lesion models on curve surfaces are then summarised in a histogram as depicted in Figure 4.14. The average surface roughness $\overline{S_{a}}$ and the standard deviation of lesion models are found to be $0.0130 \pm 0.0013 \mathrm{~mm}\left(\bar{x}_{\text {Curve }} \pm\right.$ $\sigma_{\text {Curve }}$ ).


Figure 4.14 Histogram of lesion models pasted on curved surfaces.

### 4.4 Performance Analysis

### 4.4.1 Determination on Measurement Accuracy

To determine accuracy and total standard deviation of a surface roughness measurement, the average $S_{a}$ of lesion models (33) on a flat surface is used as a reference $\left(S_{a R e f}\right)$. The $S_{a R e f}$ value is found to be $0.0122 \pm 0.0011 \mathrm{~mm}\left(\bar{x}_{\text {Ref }} \pm\right.$ $\sigma_{\text {Ref }}$ ). Meanwhile, the $S_{a}$ of lesion models on the mannequin surface ( $S_{a C u r v e}$ ) is found to be $0.0130 \pm 0.0013 \mathrm{~mm}\left(\bar{x}_{\text {Curve }} \pm \sigma_{\text {Curve }}\right)$. The following equations are used for the error analysis of the algorithm. The calculation results of these equations are described in Table 4.5.

$$
\begin{gather*}
\text { Error }=\left|\bar{x}_{\text {Curve }}-\bar{x}_{\text {Ref }}\right|  \tag{4-25}\\
\sigma_{\text {Total }}=\sqrt{\sigma_{\text {Ref }}^{2}+\sigma_{\text {Curve }}^{2}}  \tag{4-26}\\
\text { Accuracy }=\left[1-\frac{\left|\bar{x}_{\text {Curve }}-\bar{x}_{\text {Ref }}\right|}{\bar{x}_{\text {Ref }}}\right] \times 100 \% \tag{4-27}
\end{gather*}
$$

From the equations (4-25) and (4-26), it is found that the error is $\mid 0.0130$ $0.0122 \mid=0.0008 \mathrm{~mm}$ and the total standard deviation is $\left(0.0011^{2}+0.0013^{2}\right)^{1 / 2}=$ 0.0017 mm . Afterwards, the algorithm accuracy is computed by applying (4-27) that
gives $(1-|0.0130-0.0122| / 0.0122) \times 100 \%=94.12 \%$. In this case, the computed accuracy is the minimum accuracy. It implies that the algorithm accuracy is not less than $94.12 \%$. Based on the calculated standard deviation ( $\sigma_{\text {Total }}=0.0017 \mathrm{~mm}$ ), the 95\% confidence interval ( $Z_{\alpha / 2}=1.96$ ) for the calculated values were in the range of $\pm 0.0019 \mathrm{~mm}$ for the sample provided. It is obtained by applying the margin error equation that is written as $\left(Z_{\alpha / 2} \times \sigma_{\text {Total }}\right) / \sqrt{n}=(1.96 \times 0.0017) / \sqrt{3}=0.0019$. The variable $n$ denotes the small sample size of the trial. The margin error equation is presented more details in Section 5.1.1 This measurement range showed that it lay within the depth resolution of the PRIMOS camera ( 0.0040 mm ) [131].

Table 4.5 Performance evaluation of the surface roughness algorithm.

| Surfaces | Surface Roughness Parameter |  |  |
| :---: | :---: | :---: | :---: |
|  | $\bar{x}(\mathrm{~mm})$ | $\sigma(\mathrm{mm})$ | Accuracy (\%) |
| Reference | 0.0122 | 0.0011 |  |
| Curve | 0.0130 | 0.0013 |  |
| Absolute | $\|0.0130-0.0122\|=$ | $\sqrt{0.0011^{2}+0.0013^{2}}=$ | $\left(1-\frac{0.0008}{0.0122}\right) \times 100 \%=$ |
| Error | 0.0008 | 0.0017 | $94.12 \%$ |

### 4.4.2 Evaluation on Rotational Invariance

The rotation invariance of the algorithm with the second and third order polynomials are tested at the rotated surfaces of lesion model. Another surgical tape on a flat surface with size more than $40 \mathrm{~mm} \times 30 \mathrm{~mm}$ is used as the object. The rotation angles are varied from $0^{\circ}$ to $330^{\circ}$ with the increment of $30^{\circ}$. Thus, 12 rotation angles are applied to evaluate rotation invariance characteristics. Figure 4.15 shows the rotated surfaces of lesion model.


Figure 4.15 Rotated surfaces of lesion models by applying 12 rotation angles.

Five measurements are conducted for each angle variation. The $S_{a}$ of lesion models with various rotation angles is obtained to be $0.0128 \pm 0.0003 \mathrm{~mm}(\bar{x} \pm \sigma)$. Figure 4.16 shows the $S_{a}$ of the rotated lesions within the acceptable accuracy ( $>95 \%$ ). The results then show that the algorithm is invariant to the rotation of the measured surface. This is made possible because the matrix inversion in a polynomial coefficients determination does not depend on the element sequence of matrices $V$ and $Z$. A surface rotation will change the element sequence of these matrices.


Figure 4.16 $S_{a}$ of lesion models with a variation on rotation angles. The values of $S_{a}$ are within acceptable accuracy (>95\%).

To test the rotation invariance of a surface roughness algorithm, a lesion surface obtained in two successive scans will be measured. The surface roughness of the lesion images are then determined by a same user in separated calculation. Here, there are 465 samples of the lesions ( 930 images). The lesion surface between the first and the second scan are made slightly different. As an example (see Figure 4.17), the lesion surface of the second scan has been tilted in a clockwise direction compared to the first scan. By applying a surface roughness algorithm to the lesions (Figure 4.17), the measured $S_{a}$ for the first and the second scans are to be 0.0371 mm ( $\left.R^{2}=0.9804\right)$ and $0.0511 \mathrm{~mm}\left(R^{2}=0.9587\right)$, respectively.

Further, to prove that the algorithm is invariant to the rotation of the measured surface, the repeatability of the successive scan is evaluated. The definition of repeatability is a closeness between independent results that are obtained through the same method on an identical object, same operator, same conditions, and performed in the short time interval [132]. The absolute differences between the two measurements are used to determine the system repeatability. The repeatability itself can be accepted if $95 \%$ of the absolute of measurement differences are less than two standard deviations of the measurement difference $\left(2 \sigma_{\text {Diff }}\right)$. This rule has been presented by Bland and Altman in [133].

From the data of successive measurements, the standard deviation of measurement differences ( $\sigma_{\text {Diff }}$ ) is found to be 0.011 mm . Therefore, the system repeatability of surface roughness algorithm can be accepted since $95.27 \%$ of the measurement differences ( 443 of 465 lesions) are within the acceptable repeatability $\left(2 \sigma_{\text {Diff }}=0.021 \mathrm{~mm}\right)$ as required by Bland and Altman rule [133].


Figure 4.17 Lesion surfaces are obtained from two successive scans. Lesion (a) is from the first scan and lesion (b) is from the following scan.

The measurement differences can be due to manual segmentation of the lesion ROI as seen in an example in Figure 4.18. The ROI of lesion (a) is larger than the lesion (b). This problem might occur in a segmentation process - particularly in the case of a very large lesion. As a consequence, the lesion boundary cannot be clearly defined.


Figure 4.18 The 2D image and height map of lesion surfaces from the first (a) and the second (b) measurements.

### 4.4.3 Sample Area for Surface Roughness Determination

In the developed surface roughness, the measured lesion surface is divided into several smaller subdivided surfaces in order to minimise over fitting by a polynomial surface fitting. The overall surface roughness of lesion surface is obtained by averaging the surface roughness of subdivided surfaces, which have been separately calculated. In this algorithm, the surface division is performed without any consideration of the lesion size. The algorithm does not define minimum area of lesion surface for the optimal performance of the algorithm.

Stout and Blunt have reviewed several 3D surface measurement methods in bioengineering [134]. The review describes that the sampled area is dependent on the machine resolution, measuring time requirement and sample availability. For example, in measuring the surface roughness of tooth root surface, the sampled area is set to be $0.610 \times 0.610 \mathrm{~mm}^{2}$. This method is also applied in determining the surface roughness of blood vessel where a sample area of $6.400 \times 2.720 \mathrm{~mm}^{2}$ is used. In this case, a topography measurement on a fingerprint replica with size $1.020 \times$ $1.540 \mathrm{~mm}^{2}$ is performed using a focus detection method.

In a comparative study of two profiling methods (diamond and laser stylus methods), tracing length of 5.6 mm is defined in extracting the roughness profile of dental material [135]. In a larger scale, the landscape surface roughness varies according to the sample area of terrain images. In every landscape type (glacial, upland fluvial, and lowland fluvial), there is a threshold sample area to the extent that there is no any surface roughness variation at the sample area bigger than the
threshold [136]. By determining the surface roughness on a bigger area, it can be ensured that all roughness components from low to high frequencies are accounted in a surface roughness determination. The roughness components that have low and medium frequencies might not be accounted if the sample area is too small. As reported in [137] the surface roughness of terrain surface is varying with the spatial resolution (spatial frequency) of the acquired images. Therefore, a large enough sample area, in this case larger than certain threshold size, needs to be acquired in obtaining a representative measurement.

A study on the surface roughness similarity of a large rough surface is reported in [138]. In this study, the threshold for the sampled area in computing surface roughness is determined by plotting two parameters, i.e. fractal dimension and amplitude parameter, against the variation of sample area. Fractal dimension and amplitude parameter are found to be scale-dependent. Its values decrease along with the increase of the sample area. However, the decreasing trend stops at a particular area size. This area is considered as a stationary threshold. Fractal area and amplitude values become constant at sample area sizes larger than the stationary threshold.

The objective of this section is to determine the minimum surface area in determining skin surface roughness. To make the algorithm working properly, the lesion surface must contain all significant roughness components. The above method [137] is adopted in this thesis but only the surface roughness - not the fractal dimension - is used as an indicator. In a developed algorithm, the calculation of surface roughness is based on the vertical deviations of the rough profiles. The amplitude parameter is used to represent the actual vertical deviations of the rough surface, whereas the fractal dimension is used to describe the rough surface irregularity. In the clinical study at the Dermatology Department, Hospital Kuala Lumpur, 2,202 lesion images have been obtained from 204 patients. The lesion sizes vary according to the psoriasis severity as shown in Figure 4.19. This figure shows some lesion examples that are differentiated based on its size: (1) small, $1.66 \times 2.55$ $\mathrm{mm}^{2}$, (2) medium $10.44 \times 10.05 \mathrm{~mm}^{2}$, (3) large $26.78 \times 20.29 \mathrm{~mm}^{2}$, and (4) huge $>40 \times 30 \mathrm{~mm}^{2}$. Some of the lesions are larger than the maximum scanned area ( $40 \times$
$30 \mathrm{~mm}^{2}$ ) leading the normal skin area not to be acquired in the image. Most of the collected lesions are smaller than the maximum scanning area of the PRIMOS 3D optical scanner since normal skin is required for a lesion thickness determination (lesion thickness is another parameter studied in the clinical study).


Figure 4.19 Psoriasis lesions differentiation based on its size: (1) small, (2) medium, (3) large, and (4) huge.

To find a minimum area for a sampled area, skin surface roughness has been determined using different numbers of sampled area. In this evaluation, normal skin surfaces are used as the input data. Figure 4.20 shows the examples of 2D and 3D images of a scanned normal skin surface. The texture of normal skin at the scanned area is assumed to be uniform. Therefore, it can be used in observing the surface roughness variations on several different sampled areas.


Figure 4.20 2D image (a) and 3D surface (b) of normal skin surface.

Nine normal skin images acquired from nine subjects are evaluated in this stage. Three surface images are acquired from forearm region of three volunteers with no psoriasis and six images are obtained from psoriasis patients. The images are then cropped from the surrounding normal skin of the psoriasis lesions. These images are
divided into a number of smaller subdivided surfaces with size $n \times n \mathrm{~mm}^{2}$, as shown in Figure 4.21.


Figure 4.21 Scanned image is cropped to obtain evaluated surface. Subdivided surfaces are obtained by performing mesh division on an evaluated surface.

These subdivided surfaces are considered as the evaluated surfaces. For this evaluation, the determination of surface roughness is performed for 7 size variations. In this case, the smallest area of subdivided area is $1 \times 1 \mathrm{~mm}^{2}$ whereas the largest one is $7 \times 7 \mathrm{~mm}^{2}$.

In this study, the maximum evaluated surface is fixed to be less than $50 \%$ of the maximum scanned surface $(40 \mathrm{~mm} \times 30 \mathrm{~mm})$. The condition aims to minimise the curvedness of the evaluated surfaces. Therefore, the maximum dimension of the evaluated surface is $20 \mathrm{~mm} \times 15 \mathrm{~mm}$. According to the steps of the surface roughness algorithm (described in Figure 4.3), this surface is divided into $2 \times 2$ subdivided surfaces. This division gives a sub-divided surface with size $10 \mathrm{~mm} \times 7.5$ mm . The height size, 7.5 mm , is then rounded to 7 mm and selected as the maximum dimension of the evaluated surface. If the maximum dimension is set to 8 mm or 10 mm , it will give an evaluated surface larger than $50 \%$ of the maximum scanned surface.

The sizes are varied at increments of 1 mm on both sides. The second, third, and fourth order polynomials are applied to these subdivided surfaces. In previous section, only the second and third orders have been applied. Similarly, the fourth
order polynomial is also applied to figure out some fitting problems on a larger area. A larger curved surface may have more undulations than the smaller surface. Therefore, a polynomial with a higher order is required to fit this surface. The equation of fourth order polynomial can be expressed as in (4-28).

$$
\begin{align*}
z_{4}(x, y)= & \left(a_{1} x^{4}+a_{2} x^{3}+a_{3} x^{2}+a_{4} x+a_{5}\right) y^{4}+ \\
& \left(a_{6} x^{4}+a_{7} x^{3}+a_{8} x^{2}+a_{9} x+a_{10}\right) y^{3}+ \\
& \left(a_{11} x^{4}+a_{12} x^{3}+a_{13} x^{2}+a_{14} x+a_{15}\right) y^{2}+  \tag{4-28}\\
& \left(a_{16} x^{4}+a_{17} x^{3}+a_{18} x^{2}+a_{19} x+a_{20}\right) y+ \\
& \left(a_{21} x^{4}+a_{22} x^{3}+a_{23} x^{2}+a_{24} x+a_{25}\right)
\end{align*}
$$

A polynomial surface fitting can fit a surface properly if the fitting order is not less than the actual order of the surface. On the other hand, the actual order of the surface can be characterised by the number of peaks and valleys of the undulated surfaces. Table 4.6 shows the obtained surface roughness values. Here, the cases of the fitting results at $R^{2}<0.90$ are not included. Figure 4.22 shows the surface roughness of normal skin samples from 9 subjects with a sampling area variation. The sampling areas are varied from $1 \times 1 \mathrm{~mm}^{2}$ to $7 \times 7 \mathrm{~mm}^{2}$.

Table 4.6 The normal skin surface roughness of 9 subjects with a size variation

| Sampling <br> length (mm) | Surface roughness (mm) |  |  |  |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | $S_{1}$ | $S_{2}$ | $S_{3}$ | $S_{4}$ | $S_{5}$ | $S_{6}$ | $S_{7}$ | $S_{8}$ | $S_{9}$ |  |
| 1.0 | 0.015 | 0.008 | 0.011 | 0.014 | 0.013 | 0.015 | 0.014 | 0.013 | 0.020 |  |
| 2.0 | 0.018 | 0.011 | 0.014 | 0.019 | 0.018 | 0.024 | 0.018 | 0.018 | 0.030 |  |
| 3.0 | 0.020 | 0.011 | 0.015 | 0.022 | 0.019 | 0.032 | 0.019 | 0.022 | 0.035 |  |
| 4.0 | 0.021 | 0.012 | 0.016 | 0.024 | 0.020 | 0.035 | 0.019 | 0.023 | 0.036 |  |
| 5.0 | 0.022 | 0.012 | 0.016 | 0.025 | 0.020 | 0.034 | 0.019 | 0.024 | 0.038 |  |
| 6.0 | 0.024 | 0.013 | 0.016 | 0.031 | 0.020 | 0.036 | 0.020 | 0.024 | 0.040 |  |
| 7.0 | 0.025 | 0.013 | 0.017 | 0.031 | 0.021 | 0.034 | 0.021 | 0.026 | 0.045 |  |



Figure 4.22 The surface roughness of normal skin samples from 9 subjects with a sampling area variation. The sampling area is a square of sampling length.

Surface roughness $\left(S_{a}\right)$ versus sampling length is then fitted to a polynomial function in order to find a stationary threshold for the skin roughness determination. $R^{2}$ of these fittings is found to be greater than 0.9957 . Subjects $S_{2}$ and $S_{7}$ are then fitted with the fifth order polynomial. The sixth order polynomial, furthermore, is used to fit the subjects $S_{1}, S_{3}, S_{4}, S_{5}, S_{6}$, and $S_{8}$, while the seventh order polynomial is only applied to fit subject $S_{9}$. The polynomial equation can be written as follows.

$$
\begin{equation*}
f(n)=c_{1} n^{6}+c_{2} n^{5}+c_{3} n^{4}+c_{4} n^{3}+c_{5} n^{2}+c_{6} n+c_{7} \tag{4-29}
\end{equation*}
$$

A fitting result is considered good if $R^{2}$ is greater than 0.9 and close to 1.0 . At $R^{2} \geq 0.9$, the difference between the actual data points and the fitted data points are approaching a minimum value. All $R^{2}$ of these nine fitting equations are greater than 0.9 . The order of polynomial equation is selected from the highest $R^{2}$ that can be obtained from the polynomial fitting.

By referring the plots in Figure 4.22, it is possible to determine the threshold of the surface roughness stability. This threshold defines the minimum sampled area for measuring skin surface roughness. The threshold point can be determined from the first inflection point of the plot. To find this inflection point, the second derivative of the fitting equation (2) needs to be determined. The first inflection point occurs when the plot of the second derivative crosses the axis $Y=0$. Figure 4.23 depicts the plot of surface roughness dependency on a sampling size variation. Here, its first derivative and second derivative for the data are taken from subject $S_{3}$.


Figure 4.23 (a) Surface roughness $\left(S_{a}\right)$ depends on sampling size variations. Figure (b) and (c) are the plot of the first and the second derivative functions, respectively.

As shown in Figure 4.23 (c), the first inflection point is found at $n=4 \mathrm{~mm}$ with $d^{2} S_{a} /\left.d n^{2}\right|_{n=4}=-1.73 \times 10^{-6} \mathrm{~mm}$. It can be stated here that the roughness components are totally covered within a sampling area if its size is not less than $4 \times 4$ $\mathrm{mm}^{2}$. Surface roughness is increasing for $n<4 \mathrm{~mm}$ because the roughness components are not totally included within the sampled area. To determine a threshold value applicable for all skin surfaces, the highest value has to be determined from several skin samples. The threshold points of 9 evaluated skin samples are shown in Table 4.7.

As seen in Table 4.7, the highest threshold point is found to be 4.9 mm . This size can be considered as a minimum sampled area that can be used in a skin surface
roughness determination. The aim of this area setting is to ensure that all of roughness components are accounted in the calculation. From the findings, the surface roughness measurement on normal skin needs a sampled area not less than $3.8 \times 3.8 \mathrm{~mm}^{2}$. It is also recommended to measure a lesion surface with a sampled area greater than $4.9 \times 4.9 \mathrm{~mm}^{2}$. Moreover, its texture is less regular than the texture of normal skin. Therefore, a small lesion as shown in Figure 4.19 would not be correctly measured with the current algorithm.

Table 4.7 The threshold points of minimum sampling area.

| Subject | Threshold <br> point (mm) | Minimum <br> Sampling Area <br> $\left(\mathrm{mm}^{2}\right)$ | $d^{2} S_{a} / d n^{2}{ }_{i}$ <br> $(\mathrm{~mm})$ |
| :---: | :---: | :---: | :---: |
| $S_{1}$ | 3.4 | 11.56 | $9.75 \times 10^{-8}$ |
| $S_{2}$ | 3.9 | 15.21 | $-4.89 \times 10^{-8}$ |
| $S_{3}$ | 4.0 | 16.00 | $-1.73 \times 10^{-6}$ |
| $S_{4}$ | 3.6 | 12.96 | $-1.08 \times 10^{-6}$ |
| $S_{5}$ | 3.4 | 11.56 | $-7.81 \times 10^{-7}$ |
| $S_{6}$ | 4.3 | 18.49 | $3.57 \times 10^{-7}$ |
| $S_{7}$ | 3.6 | 12.96 | $7.96 \times 10^{-8}$ |
| $S_{8}$ | 4.9 | 24.01 | $1.91 \times 10^{-7}$ |
| $S_{9}$ | 3.5 | 12.25 | $1.91 \times 10^{-6}$ |

In the developed surface roughness algorithm, an input 3D surface is divided into several $2 \times 2$ subdivided surfaces. By using 4.9 mm as the minimum subdivided area, it can be defined that the minimum size of 3D surface is $(2 \times 4.9) \mathrm{mm} \times(2 \times 4.9) \mathrm{mm}=$ $9.8 \mathrm{~mm} \times 9.8 \mathrm{~mm}$. Generally, a surface roughness variation is less for the sample area that is higher than the threshold. However, there is a small surface roughness increment related to the sample size enlargement. As illustrated in Figure 4.23 (c), the surface roughness increment is started from sample size $(n)$ equal to 6 mm $\left(d S_{a} /\left.d n\right|_{n=6 m m}=0.0004 \mathrm{~mm}\right)$. This fact is initiated due to the polynomial surface fitting properties. In a polynomial surface fitting, the errors near the borders are increasing.

A data point can be estimated properly if the estimated data point is located in the centre of the actual data points. The fitting iteration of its neighbour data points can minimise the fitting error at the centre data point. This ideal condition cannot be obtained at the border areas where an estimated data point is fitted with the less neighbour data points. Similarly, the location of estimated data point also cannot be
considered as a centre point since only a half of neighbour data points are available. Another half of the neighbour data points cannot be obtained because the data are located outside the measured area boundaries.

Through this condition, the polynomial function cannot fit the mean curvature accurately. To fit a data point accurately, a number of surrounding data points from all sides is required in a surface fitting determination [139]. The increasing sampled size in fact can increase the error between the fitted surface and the original surface.

### 4.5 Summary

Chapter 4 details the development of surface roughness algorithm. Surface roughness is used as the measurable features to represent the severity of psoriasis scaliness. A 3D optical scanner is used to acquire some lesion surface images. The estimated 3D waviness surface is extracted from the rough lesion surface by applying a high order polynomial surface fitting. The rough lesion surface is subtracted from the estimated waviness to obtain the vertical deviations of lesion surface. Surface roughness is calculated by averaging the absolute values of the vertical deviations.

This chapter additionally presents the validation study of surface roughness algorithm on the standardised rough surface and curved surfaces. This validation is conducted in order to determine the accuracy and total standard deviation of the developed algorithm. A set of abrasive papers with different roughness grades are evaluated to perform algorithm validation on several various rough surfaces. From the validation, it can be found that the measured surface roughness of the developed algorithm is linearly related to the standardised roughness grade.

Medical tape is used to model skin lesions. The tape is elastic and can adapt with the surface curvature. To validate the surface roughness algorithm on human skin, the lesion models are placed on the curved surfaces of a life-size medical mannequin. The validation process calculates the accuracy and total standard deviation of a roughness measurement on 3D curved skin surfaces. Surface roughness of lesion models is considered valid if surface roughness of lesion model is constant at any locations. The algorithm accuracy is high and can be accepted.

The rotation invariance of the algorithm is tested at the rotated surfaces of lesion model. A lesion model is rotated at varying rotation angles. Surface roughnesses of rotated lesion models are determined. The results prove the rotation invariance characteristic of the surface roughness algorithm. The rotation invariance of a surface roughness algorithm is also tested to psoriasis lesion surface obtained from two successive scans. Due to the rotation and translation, the lesion surface between the first and the second scan is made slightly different. From the measured data, it can be found the system repeatability is within the acceptable repeatability.

Furthermore, the minimum surface area for surface roughness algorithm is determined to define the algorithm limitations. The sizes of psoriasis lesions vary according to the severity. To calculate a minimum sample area, skin surface roughness has been determined using a different number of sampled areas. These images are divided into smaller subdivided surfaces. In the evaluation, the surface roughness determination is applied to the images with several size variations. A sample area threshold defines the minimum sampled area for the surface roughness algorithm. The area threshold is determined from the plot of surface roughness against sampling length. Sample area enlargement does not change the determined surface roughness if the sample area is greater than the threshold level.

From the materials presented in this chapter, it can be shown that the developed algorithm has been validated at a significant level. In its last part, this chapter also clearly describes the algorithm performance and limitations. These remarkable results provide the high level of confidence to build a system for objective clinical assessment. The system is built by combining the developed surface roughness algorithm with the clustering algorithm. This part is elaborated in the following chapter.

## CHAPTER 5

## DEVELOPMENT ON SURFACE ROUGHNESS ALGORITHM

Chapter 5 describes the study set-up, data collection, result and analysis of the developed computerised psoriasis (PASI) scaliness assessment. For a statically representative data collection, a proper sampling size needs to be determined. Dermatologists assess the PASI scaliness of all collected data. Section 5.1 of this chapter presents the data collection requirements and procedures. The developed surface roughness algorithm (presented in Chapter 4) is applied to measure the skin roughness of psoriasis lesions. It is then followed by Section 5.2 that discusses about unsupervised clustering methods. Two clustering methods, $k$-means and fuzzy $c$ means (FCM) algorithms, are used to obtain the PASI scaliness score of psoriasis lesion. These methods are presented to classify the measurement result of skin surface roughness to be next interpreted in PASI scaliness score. The results of these clustering algorithms are described in this section. Following this, Section 5.3 describes an agreement analysis related to the PASI scaliness scoring performance that is evaluated by determining kappa agreement coefficients between the first and the second assessment of the developed algorithm. The results of agreement analysis are later used to determine the assessment objectivity. Section 5.4 recaps all findings and analyses of the Chapter 5.

### 5.1 Study Set-up and Data Collection

### 5.1.1 Sample Size Measurement

A sample data is used to represent the uncounted size of population data. The problem is about how to determine the sample size correctly. The size should be
well optimised to represent the population characteristics. A very large sample size might make the data representation better but it can be costly and time-consuming. Lowering the sample size, can reduce the amount of the resources needed. Nevertheless, this can potentially make the sample characteristics dissociated from the actual population.

The determination of the sample size can be done based on the margin error equation [140]. The margin error equation is used to determine a maximum error between the population mean $\mu$ (true value) and the sample mean $\bar{x}$ (observed from the collected data). The population data is assumed normally distributed. Only the population standard deviation can be provided to the equation whereas the population mean is unknown. The equation is given as follows,

$$
\begin{equation*}
E=z_{\alpha / 2} \frac{\sigma}{\sqrt{n}} \tag{5-1}
\end{equation*}
$$

Variable $E$ represents the margin of the error of the unknown population mean. The error is determined from the sample mean $\bar{x}$ of sample data. If the mean of sample $\bar{x}$ has the same value with the population mean, $\mu$, then $E$ will be equal to zero. Variable $Z_{\alpha / 2}$ refers to the critical value of normal distribution table (the bell curve). The term $\alpha$ is used to define the level of confidence. Any positive value less than 1 can be applied but in order to have meaningful results, the value of $\alpha$ should be close to 1 . The level of confidence is usually set at $90 \%, 95 \%$, and $99 \%$. For instance, to obtain an area with the level of confidence $95 \%, \pm Z_{\alpha / 2}$ values need to be determined. Here, it is calculated by $\alpha=1-0.95=0.05$. This remaining area ( $\alpha=$ 0.05 ) is then divided by 2 and distributed to the left ( 0.025 ) and the right $(0.025)$ tails of the normal distribution, as shown in Figure 5.1. The observed area (95\%) is located in the middle of normal distribution and bounded by $\pm Z_{\alpha / 2}$. From the standard normal distribution table, it can be found that $\pm Z_{\alpha / 2}= \pm 1.96$. The details of Z score determination is presented in Appendix A. The term $\sigma$ is the population standard deviation whereas $n$ is the sample size.


Figure 5.1 An area with a certain confidence level bounded by $\pm Z_{\alpha / 2}$.
Equation (5.1) is rewritten as follows to obtain the sample size:

$$
\begin{equation*}
n=\left[\frac{z_{\alpha / 2} \times \sigma}{E}\right]^{2} \tag{5-2}
\end{equation*}
$$

The population standard deviation $\sigma$ is considered similar to the sample standard deviation. As in the aforementioned example, the $Z_{\alpha / 2}$ is found based on the required accuracy $(1-\alpha)$ in the sampling process. For fixed $E$ and $\sigma$ values, a higher confidence level can be obtained by increasing the sample size. In the normal distribution, a high confidence level is indicated by enlargement on the interval width of confidence levels.

The required sample size of PASI scaliness is calculated by using the data of lesion scaliness from a previous study [141]. In this study, the data is acquired by a 3D laser scanner (Konica Minolta Vivid 910) with a focal length, $f=14 \mathrm{~mm}$. The object distance is set around 1 meter from the scanner and the depth accuracy of the 3D laser scanner is up to 0.10 mm to the reference surface. The scaliness data is grouped into four different scores by applying an unsupervised clustering algorithm i.e. $k$-means clustering. Table 5.1 lists the statistical values of PASI scaliness.

Table 5.1 Statistics of the scaliness value for each score.

| Scaliness <br> Score | Samples | Minimum <br> $(\mathrm{mm})$ | Maximum <br> $(\mathrm{mm})$ | Interval <br> $(\mathrm{mm})$ | Mean <br> $(\mathrm{mm})$ | Standard <br> Deviation |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | 37 | 0.024 | 0.078 | 0.054 | 0.057 | 0.014 |
| 2 | 21 | 0.083 | 0.133 | 0.050 | 0.104 | 0.014 |
| 3 | 14 | 0.140 | 0.206 | 0.066 | 0.172 | 0.021 |
| 4 | 11 | 0.221 | 0.328 | 0.108 | 0.256 | 0.038 |

From Table 5.1, score 1 is the only data that would be used for the sample size determination. This score is statistically reliable because it has more than 30 samples. The significant level $(\alpha)$ of 0.05 corresponding to confidence level $95 \%$ is assigned in the determination. To compare some options, the margin of error $E$ is varied for $5 \%, 10 \%$, and $15 \%$ and the sample mean is used as the base for determining the $E$ value. For example, a maximum difference with error $10 \%$ is determined by $10 \% \times \overline{x_{1}}=10 \% \times 0.057=0.006 \mathrm{~mm}$.

Table 5.2 Sample sizes of scaliness score 1 with $\alpha=0.05$ (confidence level 95\%) and maximum difference variations.

| Confidence <br> Level $(Z)$ | Error <br> $(\%)$ | Maximum Difference $(E)$ <br> $(\mathrm{mm})$ | Required <br> Sample Size |
| :---: | :---: | :---: | :---: |
| $95 \%$ | 5 | 0.003 | $83.66 \approx 84$ |
|  | 10 | 0.006 | $20.92 \approx 21$ |
|  | 15 | 0.009 | $9.30 \approx 10$ |

The sample size with error 5\% is used to provide sample size with acceptable mean difference from the actual value. Four scaliness scores need to be analysed. Hence, to cover all scores, the total sample size required is 4 scores $\times 84$ samples $=$ 336 samples. A scaliness sample is obtained from a scaliness calculation of a single psoriasis lesion. Based on this calculation approach, 336 psoriasis lesions are required to achieve a confidence level of $95 \%$ for all scaliness scores.

The lesion data were acquired in a clinical study involving 204 registered psoriasis patients at Department of Dermatology, Hospital Kuala Lumpur. The number of patient is defined based on a sample size determination for the PASI area parameter. Naturally, more than one lesion can be obtained from a patient. The clinical study NMRR-09-1098-4863 for data collection has been approved by the Clinical Research Centre, Ministry of Health, Malaysia. In this study, the data collection for scaliness parameter is part of the data collection for other PASI parameters (area, erythema, and thickness).

### 5.1.2 Data Collection Procedure

The patients involved are referred as participants in the study. Two criteria inclusion and exclusion - are applied in recruiting participants in the clinical study.

The details for these criteria are described in the following lists.
Inclusion criteria:

- Participant has been informed regarding the nature and the aims of the study and has given their written informed consent.
- Male and female participants can be involved in the study.
- The age of participant is not less than 18 years.
- Participant has been clinically diagnosed to have plaque psoriasis.

Exclusion criteria:

- Participant refuses consent to be involved in the study.
- Participant with other forms of psoriasis, such as erythrodermic, pustular, flexural, palmoplantar, or guttate psoriasis.
- Presence of other dermatological diseases that can interfere with the interpretation of the photographs (e.g. eczema, vitiligo).
- Participant who is unfit and is unable to stand for a long time.

A PRIMOS 3D optical scanner in this procedure is also used to acquire 3D surface images of the plaque psoriasis lesion of patients. The scanner gives the maximum depth resolution of $4 \mu \mathrm{~m}$ and the lateral resolution of $63 \mu \mathrm{~m}$. The image size obtained at this resolution is 640 by 480 pixels. The dermatologists subsequently select a representative lesion for each body region (head, trunk, upper and lower limbs). All four representative lesions are scanned in order to obtain 3D surfaces. The collected lesions are also being applied for another measurement (lesion thickness). Hence, the selection requires a lesion, which has an area less than the total scanned area $\left(40 \times 30 \mathrm{~mm}^{2}\right)$. For patients with severe psoriasis condition, additional lesions are collected from the patient without lesion area limitation.

The 3D scanning of psoriasis lesions requires certain conditions and room settings. These requirements are described as follows:
a. The illumination of the projected pattern of PRIMOS 3D optical scanner should be higher than the ambient illumination, as shown in Figure 5.2 (a). The illumination of the projected pattern produced by the optical scanner is measured to be $1,435 \pm 29$ Lux. The lights in the room are switched off to lower the ambient illumination below the projector illumination. The
ambient illumination during data collection is $249 \pm 11$ Lux.
b. The distance between the optical scanner and the object (lesion surface) is fixed at 220 mm . The measured surface must be parallel with the surface of the scanner lens. On the other hand, a fixed distance is required to scale a surface pixel to an actual size in length unit. A parallel and fixed distance arrangement is to ensure the surface is within the area of depth of field.
c. Two persons are required during the scanning process. The first person handles the camera and scans the lesion. The second person operates the PRIMOS software running on the laptop. Figure 5.2 (b) shows the data acquisition method for 3D surface of skin lesion.

(a)

(b)

Figure 5.2 (a) A projected pattern of PRIMOS 3D optical scanner. (b) A scanning process for acquiring the data for a scaliness measurement.

The process flow during the data collection involves several steps such as patient's preparation, PASI assessment by dermatologists, and data acquisition. It takes 28 minutes to complete the whole process; 12 minutes are required by dermatologists to assess PASI scores and another 16 minutes for the computerised PASI assessment. The detailed steps are as follows:
a. Dermatologist briefs patient on the procedures of data collection.
b. Patient takes off clothes and then put on underwear. The underwear is used to cover genital area for all patients and chest area for female patient.
c. Dermatologist performs the PASI assessment to score area, erythema, thickness, and scaliness. The assessment details performed by the dermatologists are presented in the section 5.1.3
d. Patient is asked to move from the medical examination room to the photography room. In this room, some investigators will perform a computerised PASI assessment. In this assessment, three imaging modalities - 2D camera, chromameter, and 3D optical scanner - are integrated to the computer with PASI software. Male patients are handled by male investigators. For handling the female patients, only female investigator is allowed to enter the photography room.
e. PASI area assessment is carried out by acquiring 2D images of the whole body of the patient. The images are photographed from four different views - front, back, right, and left. PASI area algorithm uses these 2D images to determine PASI area score.
f. PASI erythema assessment is performed by measuring the CIE Lab data of selected lesions and surrounding normal skin. A chromameter CR-400 is used to measure the colour data.
g. The last step of data collection is 3D surface measurement. The optical scanner is used to acquire the lesion surfaces. The 3D surfaces obtained in this step are used to determine PASI thickness and scaliness scores. Figure 5.3 displays acquisition processes for PASI area, erythema, and thicknessscaliness parameters, as mentioned in step 5, 6, and 7 .


Figure 5.3 Data acquisition for determining PASI parameters: (a) area (b) erythema, and (c) lesion thickness and scaliness.
h. In performing a double assessment, the acquisition steps are repeated from step 4 to step 6 . This session is performed after step 6 has been completed in the earlier session. In this research, the double assessment is applied to the last 43 patients. 161 patients earlier are assessed in one session only. Not all of patients are assessed twice because double assessment was only proposed during the later stages of the study.

### 5.1.3 Procedure of Dermatologist Assessment

There are two dermatologists involved in the clinical study of the research. These dermatologists independently assess the patient. The dermatologists perform the PASI assessment to score area, erythema, thickness, and scaliness, as depicted in Chapter 2 by Figure 2.8. The steps of dermatologist assessment are as follows:
a. For erythema, thickness, and scaliness assessment, the first dermatologist selects a representative lesion for each body region (head, upper limbs, trunk, and lower limbs). After being labelled using a pen marker, the lesions are scored by the first and the second dermatologists consecutively. Figure 5.4 (a) shows the dermatologist gives label on a representative lesion. An example of a labelled lesion (Lesion 2268 obtained from lower limb region of Patient 184) is shown in Figure 5.4 (b). Number 2 indicates the lesion is the second lesion to be assessed at particular body region. This label is written on the surrounding normal skin surfaces and it would not affect the process of surface measurement. The labels are used to help the dermatologists to differentiate the assessed lesions. Another one or two additional lesions at four body regions can be collected and assessed by the dermatologist. Therefore, for each patient, the maximum number of lesions assessed by dermatologist is 12 lesions ( 3 lesions $\times 4$ body regions).
b. In the PASI scaliness assessment, the dermatologists give the scores based on their visual and tactile perceptions. The dermatologists sense the skin scaliness by visually observing the lesion appearance and repetitively sliding their pointer finger on the lesion surface. The first dermatologist performs the assessment and then records the PASI scores on the
assessment sheet.
c. The second dermatologist conducts PASI assessment on the representative lesions selected by the first dermatologist. This second assessment is conducted after the first dermatologist completed the assessment for all lesions. Therefore, all of scores are obtained independently and there are no mutual influences among the dermatologists. The second dermatologist records the PASI scores on the different assessment sheet.


Figure 5.4 (a) Dermatologist selects and labels the lesion. (b) An example of a labelled lesion on lower limb region of Patient 184.
d. The second dermatologist conducts PASI assessment on the representative lesions selected by the first dermatologist. This second assessment is conducted after the first dermatologist completed the assessment for all lesions. Therefore, all of scores are obtained independently and there are no mutual influences among the dermatologists. The second dermatologist records the PASI scores on the different assessment sheet.
e. Finally, the investigators collect the assessment results given by the dermatologists. The scores are recorded and summarised to be used in the analysis stage.

### 5.1.4 Data Profile of Recruited Patients in Clinical Study

In this research, the psoriasis lesions dataset is applied to the surface roughness algorithm. The data have been acquired from a number of the registered psoriasis patients of Dermatology Department, Hospital Kuala Lumpur. The data collection
was conducted from 10 March to 5 October 2010 (10 months) involving 204 patients ( 163 males and 41 females) with ages ranging from 19 to 82 years. The average psoriasis duration of the recruited patients is 14.05 years. Therefore, most of the patients have been affected by psoriasis more than 10 years.

The demographic data might not complete enough to represent the actual condition of psoriasis - especially at gender category. The number of male patients is higher than that of female patients. A population-based study has shown that the male and female patients can suffer psoriasis with a male to female prevalence ratio almost 1:1, except for younger females (i.e. age <20 years) whereas the female prevalence is higher with ratio of $1: 1.3$ [142]. In this study however, more male patients were recruited due the lack consenting female patients. They worry about the assessment procedure, especially in taking 2D images of their body. The patient demography based on gender, race, age, and psoriasis duration are shown as charts in Figure 5.5.


Figure 5.5 Charts of patient demography based on (a) gender, (b) race, (c) age, and (d) psoriasis duration.

The sample size of roughness measurement is 1,999 psoriasis lesion images collected from 204 psoriasis patients. The images consist of 1,351 images scanned in
a single measurement and 648 scanned images in double assessments on 324 lesions. 1,892 psoriasis lesions have been scored $1,2,3$, or 4 by dermatologists. The score distribution of the lesions is described in Table 5.3.

Table 5.3 Distribution of PASI Scaliness Score ( $N=1,892$ ).

| PASI Scaliness Score | Number of Lesion | Percentage (\%) |
| :---: | :---: | :---: |
| 1 | 1,266 | 66.91 |
| 2 | 409 | 21.62 |
| 3 | 153 | 8.09 |
| 4 | 64 | 3.38 |
| Total | 1,892 | 100.00 |

Figure 5.6 shows the histogram of the surface roughness of 1,999 psoriasis lesion images. Based on the histogram, the minimum and maximum roughness values are 0.010 mm and 0.187 mm , respectively. A roughness algorithm has been tested at these limit values. The lowest limit of surface roughness has been determined in the measurement of lesion model at flat surface. The value of the lowest limit is 0.0100 $\mathrm{mm}\left(\bar{x}_{\text {Ref }}-2 \sigma_{\text {Ref }}\right)$.


Figure 5.6 Histogram of surface roughness of collected lesions.

For the highest limit, it can be obtained from validation results of surface roughness algorithm on abrasive papers. The highest surface roughness is found to be $0.3962 \mathrm{~mm}\left(\overline{S_{a}}+\sigma\right)$. This roughness is found from measurement of the roughest abrasive paper evaluated in the research. Surface roughness values of abrasive paper and lesion model are mentioned in section 4.3.1 and 4.3.2. The histogram is
normally distributed but is skewed to the left side with positive skewness of 2.002 and with mean $(0.043 \mathrm{~mm})$ and median $(0.038 \mathrm{~mm})$. The proportion of lesions with low roughness is greater since most of psoriasis lesions are being treated. Therefore, the number of lesion at a severe stage is lesser than the milder stages.

The collected lesion images are categorised into two datasets, namely training dataset and testing dataset. This categorisation is based on following conditions:
a) Training dataset:

- Images are obtained from a single measurement with $R^{2} \geq 0.9$ on a surface roughness algorithm.
- Images are scanned from a double measurement but one of the images gives $R^{2}<0.9$. The image with $R^{2} \geq 0.9$ can be included as training.
- Images are acquired from a double assessment but $S_{a}$ difference between the first and the second measurement is greater than $3 \sigma_{\text {Total }}(0.007 \mathrm{~mm})$. The images with $R^{2} \geq 0.9$ are selected as training data.

Total number of training dataset is 1,351 lesions.
b) Testing dataset:

- Images are obtained from a double assessment whereas $R^{2} \geq 0.9$ and $S_{a}$ difference of a double measurement is less than $3 \sigma_{\text {Total }}$.
Total number of testing dataset is 324 lesions. These lesions are imaged two times therefore it gives 628 lesion images.


### 5.2 Clustering Algorithm Classifying Surface Roughness

Data classification is differentiated into two approaches, namely supervised and unsupervised classifications. As mentioned by Jain et al. [143], the supervised classification is referred as discriminant analysis whereas the clustering is considered to be an unsupervised classification. The different between these two approaches is in the training data initialisation. For the supervised classification, the training data is labelled and used to define the cluster descriptions. These descriptions are then used to classify the inputted data. In contrast, the unsupervised classification uses
unlabelled data for the training. The data are clustered based on its similarity criteria. The process is performed iteratively until the centroids are maximally separated [143]. In data management, the clustering algorithm is required since the volume and the variety of data are rapidly increasing nowadays. The clustering algorithm here can enable the user to understand, to process, and to summarise this abundant data [144]. By observing at the clustered data, the analysis and the decision can be carried out easier and more organised. The clustering algorithm can be applied in various applications including medical fields.

The unsupervised clustering algorithms have been widely applied for the image analyses of medical applications. Several applications are mentioned in the following section. Li and Yuen have applied a regularised colour clustering algorithm in order to find the significant colours for tongue colour diagnosis. These colour groups are used to classify eight medical categories [145]. Masulli and Schenone developed the possibilistic neuro fuzzy c-means algorithm (PNFCM) to segment the brain features of MRI images [146]. The algorithm is able to segment the brain structures of normal people and to locate a tumour area of the brain images with meningioma [146].

The $k$-means and the fuzzy $c$-means (FCM) clustering algorithms are widely applied in clustering implementation. These algorithms are able to recognise and classify the data into a number of clusters. The main difference between $k$-means and FCM is on the probability of the data of a certain cluster. In $k$-means, a data belongs to a cluster only. Thus, there are only two options for its probability; those are probability equal to one at its cluster and zeros for the other clusters. In contrast to $k$-means, the FCM algorithm defines the data probabilities for each defined cluster. These probabilities are known as membership degrees. The function used to determine the degrees is referred to the membership function. The FCM defines some different membership degrees for each data to the existing clusters. The degrees might be calculated by applying the input data to the available membership functions.

Figure 5.7 shows the principle difference of classification methods in hard clustering ( $k$-means) and soft clustering (FCM). For the hard clustering, an input $x_{1}$
is classified as cluster 2 since its value is located inside two boundary levels (from $T_{1}$ to $T_{2}$ ). There are only two options for the membership degree -0 or 1. Figure 5.7 (a) shows that membership degree of $x_{1}$ of cluster 2 is 1 whereas the membership degrees of $x_{1}$ for the other clusters are 0 . In soft clustering (FCM), as shown by Figure 5.7 (b), an input $x_{1}$ can be considered as the member of three clusters (cluster 1, 2 and 3). However, it has several different membership degrees. For example, the degree of $x_{1}$ belongs to cluster 1,2 , and 3 are $0.8,0.4$, and 0.0 , respectively. The classification system decides $x_{1}$ as a member of cluster 1 . The decision is taken since the membership degree at cluster 1 is the highest one among the clusters.


Figure 5.7 Comparison of classification methods used in (a) hard and (b) soft clustering algorithms.

Both unsupervised clustering algorithms are used in this research. The term of $k$ means was proposed by James MacQueen in 1967 based on the idea of a Polish mathematician, Hugo Steinhaus in 1956. The algorithm was then standardised by Stuart Lloyd in 1957 but not published until 1982. At that time, the algorithm was implemented for a pulse-code modulation. Meanwhile, Forgy proposed a similar clustering method in 1965. For this, $k$-means algorithm is also recognised as LloydForgy method. The $k$-means algorithm is simple, efficient, and easy to implement [144]. $K$-means groups the dataset into a predefined number of clusters. Initial centroids are assigned randomly or by applying certain algorithms to the clusters. Similarities of the dataset and the centroids are examined for every single data point. The data will be considered to belong to a cluster that has the highest similarity. After all of the dataset are completely clustered, some new centroids are recalculated by finding the mean values of the clusters. The process is iterated until the centroids
are considered constant. This consistency is indicated by the minimum changes at all centroids. A certain number of iterations can be also used to define the termination point of the iteration process. A classification system is built based on the final centroids obtained from the clustering process. The system classifies an input data based on the distance between the input data and the cluster centroid. The data will belong to a cluster if it has the highest similarity to the cluster, which is represented by the cluster centroid.

FCM clustering algorithms is also applied in this research to improve scoring performance. Soft decision of FCM is more similar with the fuzziness of human being. Fuzzy $c$-means clustering is initially proposed by Bezdek et al. in 1984 [147]. The algorithm has been developed based on a fuzzy set proposed by Zadeh [148]. Bezdek introduced this algorithm in order to improve the $k$-means algorithm. FCM minimises the limitation of hard clustering algorithm, e.g. $k$-means clustering, by associating each data to all existing clusters with a membership degree [149]. The membership degree is represented by a membership matrix $U=\left[u_{i k}\right]$. The membership degree of a data point in a particular cluster can be considered as a probability of this data point that belongs to the cluster. Subscripts $i$ and $k$ are used to index the data point and the cluster, respectively. For instance, if there are a hundred data points ( $i=1, \cdots, 100$ ) and three clusters $(k=1,2,3)$, the matrix size of $U$ will become $100 \times 3$. The membership degrees can be any real values but it must be within 0 to 1 . A data point will be associated to each cluster and determined by a membership degree. The accumulation of these degrees is always equal to 1 . The following equation describes the constraints of the membership degree [150].

$$
\begin{equation*}
\sum_{k=1}^{K} u_{i k}=1, \quad \forall i=1, \cdots, N ; 0 \leq u_{i k} \leq 1 \tag{5-3}
\end{equation*}
$$

The iteration process of FCM algorithm is performed to achieve membership degrees stability. This condition indicates that the data points have been clustered and maximally separated. As applied in $k$-means algorithm, the FCM algorithm also has an objective function for the iteration process, as written in (5-4). A new variable, fuzziness coefficient $(m)$, is introduced to the equation. The aim of this coefficient is used to symbol the fuzziness of the membership function. This
fuzziness represents the softness of the cluster boundaries. As a comparison, the fuzziness coefficient will be zero in a hard clustering. It denotes firmness of the cluster boundaries for the hard clustering. Variable $(K)$ is the number of cluster whereas $(N)$ is the number of data point. Centroids initialisation can be executed either randomly or based on prior knowledge.

$$
\begin{equation*}
J(C, U, X)=\sum_{k=1}^{K} \sum_{i=1}^{N} u_{i k}^{m}\left\|x_{i}-c_{k}\right\|^{2} \tag{5-4}
\end{equation*}
$$

The iteration process updates the centroids and the membership matrix continuously. The equation for calculating the new centroids is given by (5-5).

$$
\begin{equation*}
c_{k}=\frac{\sum_{i=1}^{N} u_{i k}^{m} x_{i}}{\sum_{i=1}^{N} u_{i k}^{m}} \tag{5-5}
\end{equation*}
$$

The membership matrix is determined by applying equation (5-6).

$$
\begin{equation*}
u_{i k}=\frac{1}{\sum_{l=1}^{K}\left[\frac{\left\|x_{i}-c_{k}\right\|}{\left\|x_{i}-c_{l}\right\|}\right]^{2 /(m-1)}} \quad i=1,2, \cdots, N \quad k=1,2, \cdots, K \tag{5-6}
\end{equation*}
$$

The objective function is going to be stable when some high membership degrees are assigned near the cluster centroids. While, some low membership degrees are obtained at the data far from the centroids [151]. The iteration process to update matrixes U and M is repeated until no changes in the cluster centroids are found. It is indicated by $\Delta J<\varepsilon$ where $\varepsilon$ is an acceptable difference to terminate the iteration. The membership degree of a data point in a particular cluster can be considered as a probability of this data point that belongs to the cluster.

Algorithm 5.1: The FCM algorithm
i) Define the number of cluster $(K)$, membership matrix $\left(u_{i, k}\right)$, and fuzziness coefficient ( $m$ ). Use a number of small random numbers to the membership matrix $u_{i, k}$ and a real value between 1 and 2 for the fuzziness coefficient. Fix the objective function $J(C)$ equal to 0 . The acceptable $\operatorname{error}(\varepsilon)$ and the maximum number of iteration are also initialised in this step. These values are used to indicate termination point of the iteration.
ii) Compute new centroids by applying $u_{i, k}$ and $x_{i}$ to equation (5-19).
iii) Update the new membership matrix based on new centroids at step 2 and equation (5-20).
iv) Determine the objective function $J(C)$ by applying the centroids and the membership matrix to equation (5-4).
v) Determine the difference between the latest to the previous objective function. The difference is denoted as $\Delta J$. If the difference is larger than the predefined acceptable error $(\varepsilon)$, continue the iteration to step 3 . The iteration could be terminated if $\Delta J<\varepsilon$ or the iteration number is over the maximum value. Here, the value of $\varepsilon$ is $10^{-5}$.

## End.

From the clustering result, it is found that the membership degrees are not calculated at every data point values outside the values given in the training stage. Therefore, a membership function is required to determine the membership degrees at any data points. The membership degrees of the cluster are used to create a membership function. In this case, the curve fitting algorithms can be applied to find the best membership function. Gaussian functions might also be used to fit the membership degrees of the clustered data points. The classification of any input data is decided by comparing the membership degrees of the input. The data will be included to a cluster if its membership degree is higher than the membership degrees to the other clusters.

Two unsupervised clustering methods, $k$-means and FCM algorithms are investigated. $K$-means clustering is used to classify $n$ observed pattern into $k$ clusters. Each observation data point is classified into a cluster with the nearest mean. In this research, psoriasis lesions are classified into four score groups by applying $k$-means clustering to the surface roughness dataset. The objective of the $k$ means algorithm is to divide the dataset into $k$ clusters to achieve the minimum within-group sum of squares. The basic form of the $k$-means algorithm is based on the iteration of two processes. The first process is the assignment of data points into score groups. A data point is assigned to the closest group in term of Euclidean distance. The second process is the calculation of new group means, namely group
centroid, based on the new assignments. The processes will be terminated when there is no any new assignment required. In this stage, the centroids of score groups are maximally separated [144]. A boundary level between two adjacent score groups is then determined by finding a middle point between the centroids of adjacent groups. 1,351 lesions are used in the training. The dataset is selected based on the criteria as mentioned in the previous section. Table 5.4 describes the score group centroids of PASI scaliness scores.

Table 5.4 Centroid of PASI scaliness scores.

| PASI scaliness score | Score Centroid (mm) | $N$ | Percentage (\%) |
| :---: | :---: | :---: | :---: |
| 1 | 0.024 | 486 | 35.97 |
| 2 | 0.040 | 523 | 38.71 |
| 3 | 0.061 | 285 | 21.10 |
| 4 | 0.106 | 57 | 4.22 |
| Total |  | 1,351 | 100.00 |

Furthermore, the $k$-means clustering algorithm ability to classify scaliness of psoriasis lesion based on surface roughness is validated by applying the algorithm to divided datasets. Here, the dataset is randomly partitioned into two equal sized datasets (Subset 1 and Subset 2). The scores are properly clustered (maximally separated) if the score centroids for all datasets are consistent. In other words, the score centroids are not influenced by a dataset partition since the sample size of each score is statistically accepted (sample size is greater than 30 ). $K$-means clustering algorithm, thus, requires a large dataset for each score to achieve centroid stability.

Table 5.5 and Table 5.6 depict the stability of the score centroids of partitioned datasets. The centroid differences of each score between Subset 1 and Subset 2 are smaller than 0.004 mm . Meanwhile, to observe centroid consistency on a smaller sample size, the dataset is randomly partitioned into three datasets (Subset 1, Subset 2, and Subset 3). The centroid consistency of subsets can be observed in two and three partitions. The differences of centroid are not more than $3 \sigma_{\text {Total }}(0.005 \mathrm{~mm})$ except for the centroid difference of score 4 in three partitions ( 0.008 mm ). The centroid difference is greater because the size of clustered dataset is insufficient ( $<30$ samples).

Table 5.5 Score centroids of subset 1 and 2.

| PASI <br> Score | Subset $1(N=676)$ |  | Subset 2 $(N=675)$ |  | Centroid <br> difference $(\mathrm{mm})$ |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  | Centroid (mm) | $N$ | Centroid (mm) | $N$ |  |
| 1 | 0.024 | 226 | 0.024 | 227 | 0.000 |
| 2 | 0.038 | 247 | 0.038 | 264 | 0.004 |
| 3 | 0.057 | 169 | 0.059 | 155 | 0.000 |
| 4 | 0.104 | 34 | 0.104 | 29 |  |

Table 5.6 Score centroids of subset 1, 2, and 3 .

| PASI <br> Score | Subset $1(N=451)$ |  | Subset 2 $(N=450)$ |  | Subset 3 $(N=450)$ | Average <br> centroid |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | Centroid <br> $(\mathrm{mm})$ | $N$ | Centroid <br> $(\mathrm{mm})$ | $N$ | Centroid <br> $(\mathrm{mm})$ | $N$ | difference <br> $(\mathrm{mm})$ |
| 1 | 0.024 | 156 | 0.023 | 164 | 0.025 | 158 | 0.001 |
| 2 | 0.040 | 180 | 0.038 | 155 | 0.039 | 164 | 0.001 |
| 3 | 0.061 | 90 | 0.058 | 117 | 0.058 | 96 | 0.002 |
| 4 | 0.102 | 25 | 0.109 | 14 | 0.097 | 32 | 0.008 |

The boundary levels of surface roughness for PASI Scaliness scores are calculated from Table 5.4 . A boundary level between Score 1 and Score 2 is the midpoint between score centroids. For instance, the boundary level between score 1 and 2 is 0.032 mm . It is obtained from (Centroid $1+$ Centroid 2$) / 2=(0.024+$ $0.040) / 2=0.032 \mathrm{~mm}$. This method is also applied to determine the next boundary levels. If there is a lesion with surface roughness equal to the boundary level, the lesion will be classified to have a higher score. Finally, the boundary levels of surface roughness for PASI Scaliness score are given as described in Table 5.7

Table 5.7 Boundary levels of surface roughness for PASI scaliness score.

| Score | Boundary Levels (mm) |
| :---: | :---: |
| 1 | $S_{a} \leq 0.032$ |
| 2 | $0.032<S_{a} \leq 0.051$ |
| 3 | $0.051<S_{a} \leq 0.084$ |
| 4 | $S_{a}>0.084$ |

PASI scaliness score is then determined by applying the score rules to the surface roughness of psoriasis lesions. The clustering results on the test dataset is summarised as shown in Figure 5.8. Assessment consistency of first and second assessment is symbolised by coincident markers of both assessments.


Figure 5.8 Clustering results of $k$-means algorithm on test dataset.
324 data points are used as test dataset. The detailed clustering result is described in the appendix section. $289(89.2 \%)$ data points are successfully classified as the same cluster in the first and second assessments. However, 35 (10.8 \%) data points of double assessments are classified as different cluster. The misclassifications can be categorised into three types based on location of the occurrences. The first type is misclassification in determining either score 1 or score 2. The score 1 might be found in the first assessment and then followed by score 2 in the second assessment otherwise score 2 precedes the score 1 . The second type is the classification uncertainty of score 2 or 3 . The last type is the scoring vagueness between score 3 and 4. Two different scores can also be obtained in the second and the third types as exemplified by the first type. These measurement inconsistencies occur when the surface roughness data points are located too close to the score
boundary levels.

As described in Table 5.7, there are three boundary levels can be defined. These boundary levels distinguish a score cluster from the other clusters. The levels are labelled as $B_{1}, B_{2}$, and $B_{3}$. $B_{1}$ is used to separate score 1 from score 2 . The second boundary is $B_{2}$ that divides score 2 from score 3 . The last boundary is $B_{3}$. This boundary splits score 3 from score 4 . The values of $B_{1}, B_{2}$, and $B_{3}$ are 0.032 mm , 0.051 mm , and 0.084 respectively. Closeness of the measured surface roughness to each boundary level is computed by applying Euclidean distance equation. The equation is given by

$$
\begin{equation*}
D_{B_{i}}\left(S_{a}\right)=\sqrt{\left(S_{a_{1}}-B_{i}\right)^{2}+\left(S_{a_{2}}-B_{i}\right)^{2}} \quad i=1,2,3 \tag{5-7}
\end{equation*}
$$

$S_{a_{1}}$ and $S_{a_{2}}$ are measured surface roughness from the first and the second assessments. $B_{i}$ is used to represent the boundary level at $i$, as mentioned in previous paragraph. The Euclidean distance equation is applied to the test dataset that have been misclassified in the score clustering. All of the misclassification types are found in this work. There are 21 misclassification cases of type 1 (score 1 and 2), 10 cases of type 2 (score 2 and 3 ), and only 4 cases for the type 3 (score 3 and 4). The closeness of the measured surface roughness to the existing boundary levels are determined by applying equation (5-7). Furthermore, an average of the Euclidean distance is calculated to indicate closeness of the misclassification type with a certain boundary level.

The closeness measurement results are described in Table 5.8. The distances are determined from misclassification cases on test dataset. Misclassification type 1 occurs when the measured surface roughness values are close to the boundary level $B_{1}$. This closeness is shown by the $\overline{D_{B_{1}}}$ value $(0.003 \mathrm{~mm})$ which is $11 \%$ smaller than $D_{B_{2}}(0.028 \mathrm{~mm})$ and $D_{B_{3}}(0.075 \mathrm{~mm})$. Misclassification type 2 is indicated by the smallest value of $D_{B_{2}}(0.003 \mathrm{~mm})$ compared to $D_{B_{1}}(0.026 \mathrm{~mm})$ and $D_{B_{3}}(0.048 \mathrm{~mm})$. For the type 3, the trend still follows the previous types. The smallest Euclidean distance is obtained at the $D_{B_{3}}(0.004 \mathrm{~mm})$ value. The $D_{B_{1}}(0.072 \mathrm{~mm})$ and $D_{B_{2}}$ (0.046) are found much larger than $D_{B_{3}}$. The closeness distances of all
misclassification types are less than 0.005 mm .
Table 5.8 Average of Euclidean distances of surface roughness and boundary levels of $k$-means classification algorithm.

| Misclassification Type | $\overline{D_{B_{1}}}(\mathrm{~mm})$ | $\overline{D_{B_{2}}}(\mathrm{~mm})$ | $\overline{D_{B_{3}}}(\mathrm{~mm})$ | $N$ |
| :---: | :---: | :---: | :---: | :---: |
| Type 1, score 1 and 2 | 0.003 | 0.028 | 0.075 | 21 |
| Type 2, score 2 and 3 | 0.026 | 0.003 | 0.048 | 10 |
| Type 3, score 3 and 4 | 0.072 | 0.046 | 0.004 | 4 |

A small Euclidean distance at a certain boundary level is only found in the misclassification cases. Small Euclidean distances cannot be found in the data points that correctly classified. All of surface roughness data points are located beyond from the boundary levels. Thus, the misclassification cases can be minimised. The average of Euclidean distance is much higher than 0.005 mm for all score groups, as listed in Table 5.9.

Table 5.9 Average of Euclidean distances of surface roughness and boundary levels determined from correct classification cases of $k$-means algorithm.

| Score Group | $\overline{D_{B_{1}}}(\mathrm{~mm})$ | $\overline{D_{B_{2}}}(\mathrm{~mm})$ | $\overline{D_{B_{3}}}(\mathrm{~mm})$ | $N$ |
| :---: | :---: | :---: | :---: | :---: |
| Score 1 | 0.012 | 0.039 | 0.085 | 79 |
| Score 2 | 0.011 | 0.016 | 0.063 | 110 |
| Score 3 | 0.046 | 0.019 | 0.028 | 70 |
| Score 4 | 0.120 | 0.093 | 0.046 | 30 |

To avoid incorrect classification, it is recommended to collect more than a single data point if the surface roughness measured near the boundary levels. The surface roughness input is then determined by averaging the surface roughness values of multiple measurements. The surface roughness ranges that require multiple measurements are defined in Table 5.10.

Table 5.10 The surface roughness ranges near the boundary levels of $k$-means
classification algorithm.

| Misclassification Type | Surface Roughness Interval (mm) <br> $S_{a}-3 \sigma_{\text {Total }} \leq B_{i} \leq S_{a}+3 \sigma_{\text {Total }}$ |
| :---: | :---: |
| Type 1, score 1 and 2 | $0.027 \leq S_{a} \leq 0.037$ |
| Type 2, score 2 and 3 | $0.046 \leq S_{a} \leq 0.056$ |
| Type 3, score 3 and 4 | $0.079 \leq S_{a} \leq 0.089$ |

The second clustering method applied in the research is FCM (fuzzy $c$-means) clustering. This algorithm aims to improve the classification reliability especially at
a border region between two adjacent clusters. Rigidity at the border region in the $k$ means clustering is reduced by implementing the fuzziness of FCM clustering.

An example that shows $k$-means rigidity is when the results of two consecutive measurements are classified into two different clusters. It can be occurred when the boundary level is located in the middle area of the first and the second assessments. The first measurement is classified belong to the first cluster whereas the second measurement is considered to be the second cluster. Here, the similar dataset that has been applied to $k$-means clustering is used by FCM clustering. The size of training dataset is 1,351 . The clustering iteration gives the membership degrees of each data point. For this dataset, 67 iteration steps are required to achieve its stability at the objective function $J(C, U, X)$ equals 0.0503 . The function $J(C, U, X)$ is determined by applying equation (5-4). This objective function determines summation of the cluster member aggregation and the separability among the existing clusters. In perfect condition the value of the final objective function should be nearly zero. This condition is achieved when all of the cluster members are located close to its cluster centroids and there are no intersections among the membership functions, as found in hard clustering.

To reduce the iteration number, the predefined membership degrees are assigned before the iteration process. Four functions are required to determine predefined membership degrees of the score groups. To build the functions, the maximum interval of surface roughness is then divided into four equally score intervals. The cluster centroid is computed from the middle point of the score interval. Figure 5.9 show the score intervals that divide surface roughness interval into four score groups. Each score group is bordered with two boundary levels. For instance, score 1 is bordered by $L_{1}$ and $L_{3}$. The centroid of score 1 is then calculated from the middle point of $L_{1}$ and $L_{3}$. Here, the centroid is represented by variable $L_{2}$. This similar method is also applied to determine the centroids for cluster 2,3 , and 4.


Figure 5.9 Surface roughness intervals of score groups.
A triangular function is used to calculate the membership degrees of the training dataset. This triangular shape is selected because the shape will have the maximum value only at a data point. By using this shape, the membership degrees of another data points can be also linearly determined. The edges of the triangle legs are fixed at minimum and maximum values of training dataset. Here, the minimum and maximum values of abscissae are 0.0120 mm and 0.1780 mm , respectively. To create triangle shape, the highest membership degree is assigned to the cluster centroid. The predefined membership degrees of score clusters are determined by applying equations (5-8) to (5-15). These predefined membership degrees are used to initiate the clustering iteration. Variables $c_{1}, c_{2}, c_{3}$, and $c_{4}$ are the cluster centroids that assigned from $L_{2}, L_{4}, L_{6}$, and $L_{8}$ respectively, as described in Figure 5.9. $L_{1}$ and $L_{9}$ are the minimum and maximum values of the training dataset, $S_{a}$. The matrix size of of the membership degree is $N \times 4$. The $N$ is the size of training dataset. The details of membership degree equations are formulated as follows:

Membership degrees of Score 1, $c_{1}=L_{2}$

$$
\begin{equation*}
\text { if } S_{a}(i)<c_{1} ; \quad u_{i, 1}=\frac{1}{\left(c_{1}-L_{1}\right)} \times\left(S_{a}(i)-L_{1}\right) \tag{5-8}
\end{equation*}
$$

$$
\begin{equation*}
\text { if } S_{a}(i) \geq c_{1} ; \quad u_{i, 1}=\frac{-1}{\left(L_{9}-c_{1}\right)} \times\left(S_{a}(i)-c_{1}\right)+1 \tag{5-9}
\end{equation*}
$$

Membership degrees of Score 2, $c_{2}=L_{4}$

$$
\begin{align*}
& \text { if } S_{a}(i)<c_{2} ; \quad u_{i, 2}=\frac{1}{\left(c_{2}-L_{1}\right)} \times\left(S_{a}(i)-L_{1}\right)  \tag{5-10}\\
& \text { if } S_{a}(i) \geq c_{2} ; \quad u_{i, 2}=\frac{-1}{\left(L_{9}-c_{2}\right)} \times\left(S_{a}(i)-c_{2}\right)+1 \tag{5-11}
\end{align*}
$$

Membership degrees of Score 3, $c_{3}=L_{6}$

$$
\begin{align*}
& \quad \text { if } S_{a}(i)<c_{3} ; \quad u_{i, 3}=\frac{1}{\left(c_{3}-L_{1}\right)} \times\left(S_{a}(i)-L_{1}\right)  \tag{5-12}\\
& \text { if } S_{a}(i) \geq c_{3} ; \quad u_{i, 3}=\frac{-1}{\left(L_{9}-c_{3}\right)} \times\left(S_{a}(i)-c_{3}\right)+1 \tag{5-13}
\end{align*}
$$

Membership degrees of Score 4, $c_{4}=L_{8}$

$$
\begin{align*}
& \text { if } S_{a}(i)<c_{2} ; \quad u_{i, 4}=\frac{1}{\left(c_{4}-L_{1}\right)} \times\left(S_{a}(i)-L_{1}\right)  \tag{5-14}\\
& \text { if } S_{a}(i) \geq c_{2} ; \quad u_{i, 4}=\frac{-1}{\left(L_{9}-c_{4}\right)} \times\left(S_{a}(i)-c_{4}\right)+1 \tag{5-15}
\end{align*}
$$

The predefined membership degrees of the score groups are plotted to create membership degree distribution. The degrees obtained from interval division are plotted as shown in Figure 5.10.


Figure 5.10 Predefined membership degrees of score groups.

Comparison of iteration process that uses predefined and random membership degrees is shown in Figure 5.11. By applying predefined membership degrees, the iteration number has been reduced from 67 steps to 31 steps.


Figure 5.11 Iteration of FCM clustering with random (dotted line) and predefined (solid line) membership degrees.

Membership degrees of the training dataset are obtained from the iteration of FCM clustering. The degrees are clustered and scattered according to the existing clusters as depicted in Figure 5.12.


Figure 5.12 Membership degrees scattering of clustered dataset.
Gaussian fitting function is used to obtain the membership degrees or the probabilities of surface roughness at any input values. The general form of the Gaussian equation for representing the membership function of scaliness score is given by equation (5-16). The coefficients of scaliness score functions are listed in Table 5.11. The membership degrees of clustered data have been fitted to the Gaussian function with $R^{2} \approx 1$. The membership degree of a certain surface roughness $S_{a}$ is calculated by applying all membership functions, $P_{n}\left(S_{a}\right)$, from $n=1$ to $n=4$. The subscript $n$ on variable $P_{n}\left(S_{a}\right)$ represents the group of scaliness score. Equation (5-16) is constructed from the summation of two Gaussian fitting functions, which have different means and standard deviations. For the first Gaussian function, coefficient $a_{1}$ is the maximum height of the distribution, $b_{1}$ is the mean of Gaussian distribution, and $c_{1}$ is the standard deviation. The same meanings are also applied for the second Gaussian function.

$$
\begin{equation*}
P_{n}\left(S_{a}\right)=a_{1} \exp \left(-\left(\frac{S_{a}-b_{1}}{c_{1}}\right)^{2}\right)+a_{2} \exp \left(-\left(\frac{S_{a}-b_{2}}{c_{2}}\right)^{2}\right) \tag{5-16}
\end{equation*}
$$

Table 5.11 Coefficients of Gaussian functions for the roughness classification of PASI scaliness scores.

| Scaliness score | $a_{1}$ | $b_{1}$ | $c_{1}$ | $a_{2}$ | $b_{2}$ | $c_{2}$ | $R^{2}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Score 1 (S1) | 0.609 | 0.029 | 0.007 | 0.888 | 0.018 | 0.011 | 0.994 |
| Score 2 (S2) | 0.514 | 0.038 | 0.006 | 0.858 | 0.048 | 0.010 | 0.992 |
| Score 3 (S3) | 0.635 | 0.064 | 0.010 | 0.784 | 0.081 | 0.018 | 0.995 |
| Score 4 (S4) | 0.708 | 0.118 | 0.023 | 0.792 | 0.160 | 0.039 | 0.997 |

The membership functions of PASI scaliness scores are plotted in Figure 5.13. PASI scaliness score is determined by applying the rule of scaliness score as written in (5-17). An input of surface roughness is classified into a particular score if the input gives the highest membership degree or probability among the scaliness scores.


Figure 5.13 Membership functions of PASI scaliness scores.

$$
S\left(S_{a}\right)=\left\{\begin{array}{l}
1, \max \left(P_{1}\left(S_{a}\right), P_{2}\left(S_{a}\right), P_{3}\left(S_{a}\right), P_{4}\left(S_{a}\right)\right)=P_{1}\left(S_{a}\right)  \tag{5-17}\\
2, \max \left(P_{1}\left(S_{a}\right), P_{2}\left(S_{a}\right), P_{3}\left(S_{a}\right), P_{4}\left(S_{a}\right)\right)=P_{2}\left(S_{a}\right) \\
3, \max \left(P_{1}\left(S_{a}\right), P_{2}\left(S_{a}\right), P_{3}\left(S_{a}\right), P_{4}\left(S_{a}\right)\right)=P_{3}\left(S_{a}\right) \\
4, \max \left(P_{1}\left(S_{a}\right), P_{2}\left(S_{a}\right), P_{3}\left(S_{a}\right), P_{4}\left(S_{a}\right)\right)=P_{4}\left(S_{a}\right)
\end{array}\right.
$$

Some examples of score calculations are listed in Table 5.12. Sample 1, a lesion with surface roughness 0.039 mm would be classified into a PASI scaliness score. Membership degrees of the sample for score clusters (score 1 to 4 ) are $0.100,0.913$, 0,005 , and 0.000 . The highest degree is obtained at second cluster $P_{2}\left(\overline{S_{a}}\right)=0.913$ therefore the sample can be categorised as score 2 . This is similarly applied to the
subsequent lesion samples in Table 5.12.

Table 5.12 Calculations of PASI scaliness scores by applying FCM algorithm

| No | $\overline{S_{a}}(\mathrm{~mm})$ | $P_{1}\left(\overline{S_{a}}\right)$ | $P_{2}\left(\overline{S_{a}}\right)$ | $P_{3}\left(\overline{S_{a}}\right)$ | $P_{4}\left(\overline{S_{a}}\right)$ | $P_{\max }$ | Score |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | 0.039 | 0.100 | 0.913 | 0.005 | 0.000 | 0.913 | 2 |
| 2 | 0.017 | 0.908 | 0.000 | 0.000 | 0.000 | 0.908 | 1 |
| 3 | 0.063 | 0.000 | 0.085 | 0.911 | 0.004 | 0.911 | 3 |
| 4 | 0.112 | 0.000 | 0.000 | 0.044 | 0.837 | 0.837 | 4 |

Figure 5.14 describes the complete scaliness scores that have been classified by applying FCM algorithm. Several lesions that have been scored by FCM are shown in Figure 5.15. The complete results of FCM are listed in appendix section.


Figure 5.14 Clustering results of FCM algorithm on test dataset.


Figure 5.15 Deviation surfaces of psoriasis lesions: (a) score 1, (b) score 2, (c) score 3 , and (d) score 4 , determined by the surface roughness algorithm and classified by the FCM algorithm.

The same test dataset is applied to FCM algorithm. From 324 data points tested, a total of $295(91.0 \%)$ data points are correctly classified in the double assessment sessions. Misclassified data points are found at $29(9.0 \%)$ data points. As observed in $k$-means algorithm, similar misclassifications also happen in FCM implementation. It occurs in the locations near the boundary levels of the score groups. Here, the boundary levels of FCM algorithm would not be considered in classification process. These boundary levels are determined in order to find the transition boundary from a certain score to another score group.

A boundary level is defined as intersection point of two overlapping membership functions. Figure 5.16 shows an example of boundary level between score 1 and score 2. The boundary level is depicted as a green circle. In this point, both overlapping membership functions (score 1 and score 2) give a same membership degree for a certain surface roughness. This membership degree relation can be expressed by $P_{1}(0.034)=P_{2}(0.034)=0.4564$. A misclassification case appears when the measured surface roughness values are located in the left and the right sides
of boundary level, as illustrated in Figure 5.16. The first assessment, $S_{a 1}$, is classified as score 1 because membership function of score 1 gives the largest membership degree at $P_{1}(0.030)=0.8684$. Conversely, in the second assessment, the highest membership degree, $P_{2}(0.036)=0.6781$, is given by membership function of score 2. Thus, the measured surface roughness $S_{a 2}$ is considered as score 2. Finally, all boundary levels of FCM algorithm are obtained from the intersection point of membership functions. The boundary levels of misclassification type 1, 2, and 3 are $0.034 \mathrm{~mm}, 0.056 \mathrm{~mm}$, and 0.097 mm respectively. These levels are not equal to the boundary levels of $k$-means algorithm. In the FCM algorithm, the levels depend on the curve shape of the membership functions. Therefore, the levels are not exactly located in the middle point of two cluster centroids, as found in $k$-means algorithm.


Figure 5.16 A boundary level (white circle) splits membership functions of score 1 (dot-dashed line) and score 2 (dashed line).

As conducted in previous section, the average of Euclidean distance is computed to find closeness of the measured surface roughness with the boundary levels. Equation (5-7) is applied to compute the Euclidean distance for each misclassified data point. Table 5.13 shows the closeness of measured data to the boundary levels of FCM algorithm. The data is summarised from misclassified data points. Misclassification type 1 exists when the measurement results are located near to the first boundary level, $B_{1}$. The average distance $\overline{D_{B_{1}}}(0.003 \mathrm{~mm})$ is small and not more
then $10 \%$ of $D_{B_{2}}(0.030 \mathrm{~mm})$ and $D_{B_{3}}(0.088 \mathrm{~mm})$. Misclassification type 2 and 3 are also indicated by the closeness of the measured surface roughness to the boundary levels. A small value of $D_{B_{2}}(0.004 \mathrm{~mm})$ shows the closeness of measured surface roughness to the boundary level of misclassification type 2 . The occurrence of misclassification type 3 is explained by a small distance value of $D_{B_{3}}(0.004 \mathrm{~mm})$.

Table 5.13 Average of Euclidean distances of surface roughness and boundary levels of FCM classification algorithm

| Misclassification Type | $\overline{D_{B_{1}}}(\mathrm{~mm})$ | $\overline{D_{B_{2}}}(\mathrm{~mm})$ | $\overline{D_{B_{3}}}(\mathrm{~mm})$ | $N$ |
| :---: | :---: | :---: | :---: | :---: |
| Type 1, score 1 and 2 | 0.003 | 0.030 | 0.088 | 17 |
| Type 2, score 2 and 3 | 0.033 | 0.004 | 0.056 | 8 |
| Type 3, score 3 and 4 | 0.091 | 0.060 | 0.004 | 4 |

These small distances do not occur for the cases with correct classifications. The measured surface roughnesses are located far from the boundary levels. Therefore, misclassification cases could be avoided in these measurements. Here, the averages of Euclidean distances are also found higher than 0.005 mm , as shown in correct classification cases of $k$-means algorithm. The distances for all score groups are summarised in Table 5.17.

Table 5.14 Average of Euclidean distances of surface roughness and boundary levels determined from correct classification cases of FCM algorithm.

| Score Group | $\overline{D_{B_{1}}}(\mathrm{~mm})$ | $\overline{D_{B_{2}}}(\mathrm{~mm})$ | $\overline{D_{B_{3}}}(\mathrm{~mm})$ | $N$ |
| :---: | :---: | :---: | :---: | :---: |
| Score 1 | 0.012 | 0.043 | 0.101 | 104 |
| Score 2 | 0.014 | 0.018 | 0.076 | 114 |
| Score 3 | 0.053 | 0.022 | 0.036 | 55 |
| Score 4 | 0.128 | 0.097 | 0.039 | 22 |

Two methods are recommended to minimise misclassification cases. The first method is performed by acquiring more data points if the surface roughness is lie close to the boundary levels. An average value of this multiple measurement is then applied to score the surface roughness. This first method is also suggested to resolve the boundary problems of $k$-means algorithm. Table 5.18 summarises the surface roughness ranges that need to be considered for performing multiple measurements.

Table 5.15 The surface roughness ranges near the boundary levels of FCM classification algorithm.

| Misclassification Type | Surface Roughness Interval (mm) <br> $S_{a}-3 \sigma_{\text {Total }} \leq B_{i} \leq S_{a}+3 \sigma_{\text {Total }}$ |
| :---: | :---: |
| Type 1, score 1 and 2 | $0.029 \leq S_{a} \leq 0.084$ |
| Type 2, score 2 and 3 | $0.051 \leq S_{a} \leq 0.106$ |
| Type 3, score 3 and 4 | $0.092 \leq S_{a} \leq 0.147$ |

In the second method, the score group is determined by comparing the membership degrees of the decided score groups. As exemplified in Figure 5.16, the final score cannot be determined because there are two different scores from two measurement sessions. The final score can be obtained by comparing the maximum membership degrees of the first $\left(P_{1}(0.030)=0.8684\right)$ and the second $\left(P_{2}(0.036)=\right.$ 0.6781 ) assessments. From the comparison, it is known that $0.8684>0.6781$ therefore the final score for this data point is score 1. A mathematical expression is formulated to determine final score $S\left(S_{a}\right)$ from $n$-times measurements. Let $m_{i}\left(S_{i}, P_{\max , i}\right)$ represents $i$-th measurement which has final score $S_{i}$ at maximum membership degree $P_{\max , i}$. Then, the equation can be given by

$$
\begin{equation*}
S\left(S_{a}\right)=S_{k}, \max \left(P_{\max , 1}, P_{\max , 2}, \cdots, P_{\max , k}, \cdots, P_{\max , n}\right)=P_{\max , k}\left(S_{a}\right) \tag{5-18}
\end{equation*}
$$

### 5.3 The Agreement Analysis of Psoriasis Assessment

The inter-rater variation of two independent observers can be evaluated by using the kappa coefficient analysis. The Kappa coefficient has been proposed by Cohen in 1960 and has been widely used to measure an agreement among clinicians on the scores of a medical assessment. Kappa coefficient is expressed by following equation [152]:

$$
\begin{equation*}
\kappa=\frac{p_{0}-p_{c}}{1-p_{c}} \tag{5-19}
\end{equation*}
$$

The probability variable, $p_{0}$ is the proportion of units in which the observers agreed with and $p_{c}$ is the proportion of units for which agreement is expected by chance. Kappa's possible values are limited from -1 up to 1 . As an example, suppose there are two observers, $O_{1}$ and $O_{2}$. As an example, suppose there are two observers, $O_{1}$
and $O_{2}$. The observers are asked to examine and score 20 subjects. They may give a score for the subjects, either score 1 or 2 . Scoring results and summary of agreement for this scoring are given by Table 5.16 and Table 5.17, respectively.

Table 5.16 Scoring results of two observers, $O_{1}$ and $O_{2}$.

| $O_{1}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| vs |
| $O_{2}$ | $1_{1}$

Table 5.17 Summary of agreement from scoring by two observers.

| $O_{1}$ Assessment vs. | Scores of $O_{2}$ Assessment |  | Total |  |
| :---: | :---: | :---: | :---: | :---: |
|  | 1 | 2 |  |  |
| Scores of | 1 | 10 | 2 | 12 |
| S Assessment | 2 | 0 | 8 | 8 |
| Total |  | 10 | 10 | 18 |

To obtain kappa coefficient as formulated in (5-19), variables $p_{0}$ and $p_{c}$ need to be determined. Variable $p_{0}$ is determined from a ratio of total agreement of both observers to the total of sample. The calculations of $p_{0}$ is given as follows:

$$
\begin{equation*}
p_{0}=\frac{\left(N_{O_{1} O_{2}, \text { Score } 1}+N_{O_{1} O_{2}, \text { score } 2}\right)}{N_{\text {Sample }}}=\frac{(10+8)}{20}=0.90 \tag{5-20}
\end{equation*}
$$

Variable $p_{c}$ is computed by adding between probability both observers give score 1 and probability both observers give score 2 . The solution can be found as:

$$
\begin{align*}
& p_{c}=\left(\frac{N_{O_{1}, \text { Score } 1}}{N_{\text {Sample }}} \times \frac{N_{O_{2}, \text { Score } 1}}{N_{\text {Sample }}}\right)+\left(\frac{N_{O_{1}, \text { Score } 2}}{N_{\text {Sample }}} \times \frac{N_{O_{2}, \text { Score } 2}}{N_{\text {Sample }}}\right)  \tag{5-21}\\
& p_{c}=\left(\frac{10}{20} \times \frac{12}{20}\right)+\left(\frac{10}{20} \times \frac{8}{20}\right)=(0.5 \times 0.6)+(0.5 \times 0.4)  \tag{5-22}\\
& p_{c}=0.3+0.2=0.50 \tag{5-23}
\end{align*}
$$

Finally, values of $p_{0}$ and $p_{c}$ are substituted into (5-19) to calculate the kappa coefficient,

$$
\begin{equation*}
\kappa=\frac{0.90-0.50}{1-0.50}=0.80 \tag{5-24}
\end{equation*}
$$

The maximum value $(\kappa=1.0)$ represents that the observers agree in all of examination samples. Landis interpretation on Kappa coefficient values can be summarised as shown in Table 5.18 [153].

Table 5.18 Agreement interpretation of Kappa coefficient.

| Kappa | Agreement interpretation |
| :---: | :---: |
| $<0$ | Less than chance agreement |
| $0.01-0.20$ | Slight agreement |
| $0.21-0.40$ | Fair agreement |
| $0.41-0.60$ | Moderate agreement |
| $0.61-0.80$ | Substantial agreement |
| $0.81-0.99$ | Almost perfect agreement |

To evaluate an agreement between two dermatologists, the Kappa coefficient value of PASI Scaliness scores of 1,283 lesions are determined. The Kappa coefficient value between Dermatologist 1 and Dermatologist 2 is 0.55 and is categorised as a moderate agreement. The Kappa coefficient of 0.55 indicates that the dermatologists have achieved $55 \%$ agreement of total assessment and $45 \%$ agreement will be expected by chance. A perfect agreement is achieved if the Kappa coefficient is greater than 0.80 . As a result, the dermatologists' scores cannot be considered as a ground truth in evaluating the algorithm performance since the Kappa coefficient among the dermatologists is only 0.55 .

To evaluate the performance of PASI Scaliness algorithm, several lesion samples are imaged in two successive scans. PASI scaliness scores of lesion images are then analysed by the same user in separated calculations. The number of samples is 324 lesions (648 images). The number of tested lesions is less than the number of scored lesions by dermatologists because not all lesions are scanned twice. Kappa coefficients between the first assessment and the second assessment that are obtained from $k$-means and FCM algorithms are evaluated. Table 5.19, Table 5.20, and Table 5.21 show the agreement summary of dermatologists, $k$-means clustering, and FCM clustering, respectively. Table 5.22 lists the comparisons on Kappa coefficients obtained from three scoring methods. The Kappa coefficients are computed through the same way as exemplified in the previous section. It can be seen that the FCM
gives better Kappa coefficient agreement than $k$-means clustering.
Table 5.19 Summary of agreement from scoring by dermatologists.

| First Assessment vs. | Scores of Dermatologist 2 Assessment |  |  |  | Total |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 1 | 2 | 3 | 4 |  |  |
| Scores of | 1 | 634 | 200 | 4 | 0 | 838 |
|  | 2 | 37 | 204 | 39 | 0 | 280 |
|  | 3 | 4 | 24 | 75 | 5 | 108 |
|  | 4 | 1 | 0 | 16 | 40 | 57 |
| Total |  | 676 | 428 | 134 | 45 | 1,283 |

Table 5.20 Summary of agreement from scoring by applying $k$-means algorithm.

| First Assessment <br> vs. Second <br> Assessment | Scores of Second Assessment |  |  |  | Total |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 1 | 2 | 3 | 4 |  |  |
| Scores of <br> First <br> Assessment | 1 | 79 | 10 | 0 | 0 | 89 |
|  | 2 | 11 | 110 | 7 | 0 | 128 |
|  | 3 | 0 | 3 | 70 | 2 | 75 |
| Total |  | 4 | 0 | 0 | 2 | 30 |
| 32 |  |  |  |  |  |  |

Table 5.21 Summary of agreement from scoring by applying FCM algorithm.

| First Assessment <br> vs. Second <br> Assessment | Scores of Second Assessment |  |  |  | Total |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 1 | 2 | 3 | 4 |  |  |
| Scores of <br> First <br> Assessment | 1 | 104 | 12 | 0 | 0 | 116 |
|  | 2 | 5 | 108 | 4 | 0 | 117 |
| Total |  |  |  |  |  |  |
|  | 4 | 0 | 6 | 59 | 1 | 66 |

Table 5.22 Comparison of Kappa coefficients using $k$-means and FCM.

| Clustering algorithm | Kappa coefficient | N |
| :---: | :---: | :---: |
| Dermatologist | 0.5500 | 1,283 |
| $K$-means | 0.8473 | 324 |
| FCM | 0.8708 | 324 |

Perfect agreement between the first assessment and the second assessment for $k$ means and FCM algorithms are achieved since Kappa coefficients are found to be greater than 0.81 . The FCM performance ( 0.8708 ) of surface roughness classification is slightly better than $k$-means ( 0.8473 ). FCM clustering, as a soft clustering, can better solve the boundary problems compare to $k$-means clustering. Figure 5.17 depicts an example of the double measurement result. These measurements are taken from a lesion at an upper limb region (lesion number: 1600
and 1601). The first and second measurements give $S_{a}=0.030 \mathrm{~mm}$ and 0.033 mm , respectively. In $k$-means clustering, the measurement results will be classified as the different score clusters (score 1 and score 2 ) because the boundary level of score 1 and 2 is located in the middle of measurement results. These measurement results are sharply classified into different scores by $k$-means clustering even the images are acquired from the same lesion. FCM however, is able to accommodate this condition by giving the membership degrees or the probabilities of the measured data to cluster 1 and 2. Although the membership degrees of the first and second measurements are changing, the final decision is equal, both of measurements are classified as score 1 .


Figure 5.17 Clustering of the $k$-means and FCM at the boundary of two clusters score 1 and score 2. A boundary level of the $k$-means clustering (dotted-line) crosses the membership functions of score 1 (dot-dashed line) and score 2 (solid line).

### 5.4 Summary

Sampling sizes are determined to ensure that the collected data is statically representative. A margin error equation in this case is rearranged to provide a sample size equation. $Z$ scores at a certain confidence level, population standard
deviation, and maximum allowable error are used to compute the sample size. To obtain the confidence level $95 \%$ and error $10 \%$, the sample size of scaliness scores are differently found among the scores. A minimum required sample size is found to be 25 samples. To cover the data from all scores (four score groups), the total required sample size is 100 samples. A scaliness sample is calculated from surface roughness of a single psoriasis lesion. Based on the sample size calculation, it is found that a total amount of 100 psoriasis lesions are required to achieve the confidence level of $95 \%$ for all scaliness scores. The lesion data themselves have been collected from registered psoriasis patients at Department of Dermatology, Hospital Kuala Lumpur.

A total of 204 patients are recruited in this clinical study - consisting of 163 males and 41 females. The study has been approved by the Clinical Research Centre, Ministry of Health, Malaysia and registered at NMRR-09-1098-4863. A total of 1,999 psoriasis lesions are collected in the study. From this total number, 1,892 lesions are assessed by dermatologists to obtain PASI scaliness scores. The percentage of the lesion with score $1,2,3$, and 4 are $66.91 \%, 21.62 \%, 8.09 \%$, and $3.38 \%$, respectively. The proportion of lesions at a mild severity is greater since most of the lesions are still under a continuous treatment.

Two unsupervised clustering methods, $k$-means and fuzzy $c$-means (FCM) algorithms are studied in the research work. These clustering algorithms are widely applied in many medical applications. $K$-means clustering is used to classify $n$ observed pattern into $k$ clusters. Here, each data point is classified into the cluster with the nearest mean. There are four clusters representing four sets of PASI scaliness score. $K$-means clustering algorithm is validated by applying a clustering algorithm towards the divided dataset. The dataset of lesion roughness is randomly partitioned into two equal sized datasets. The consistency of score centroids for all dataset then shows that the score clusters are maximally separated. Centroid consistency is additionally observed on a smaller sample size and the dataset is randomly partitioned into three datasets. It is found that centroid differences are not more than $3 \sigma_{\text {Total }}(0.005 \mathrm{~mm})$ - except for one-third partitions of score $4(0.008 \mathrm{~mm})$. The greater difference is a result of the size of clustered dataset that is less than 30
samples.
In addition, FCM clustering is applied to improve the classification reliability. The improvement is given at a border area between two adjacent clusters. The rigorousness of $k$-means clustering is reduced by applying the softness of FCM clustering. The clustering iteration provides the membership degrees of each data point. Gaussian fitting function furthermore is fitted to the clustered dataset in order to obtain the membership degrees of the dataset. Meanwhile, the PASI scaliness score is determined by comparing the membership degrees of an input data to the existing clusters. The input surface roughness is classified into a particular score cluster if it has the highest membership degree among the clusters of scaliness scores.

Kappa coefficient analysis is used to evaluate Inter-rater variation of two independent observers. To evaluate a dermatologist agreement, the assessments of two dermatologists on 1,283 lesions are determined. Kappa coefficient of dermatologist is found to be 0.55 that is less than the required agreement, namely 0.81 . Therefore, the dermatologist agreement is not considered as the ground truth for evaluating algorithm performance.

The agreement analysis of PASI scaliness algorithm is performed by comparing two measurements obtained from two successive scans. The number of samples is 324 lesions ( 648 two successive images). Kappa coefficients between the first assessment and the second assessment obtained from $k$-means and FCM clustering algorithms are evaluated. The Kappa coefficients for $k$-means and FCM are 0.8473 and 0.8708 , respectively. Both of agreements are considered as a perfect agreement since the coefficients are found to be greater than 0.81 . In this agreement analysis, FCM classification is better than $k$-means.

FCM can solve the boundary problems better than $k$-means clustering. In the classification of $k$-means clustering, a surface roughness can be classified as different groups since the group separation is very strict. The FCM algorithm determines membership degrees for each measured data point. Therefore, the score classifications are decided based on similarity quantification which is represented by these degrees. In several boundary cases, the $k$-means algorithm can give several
different scores for the same lesions that are successively acquired in double measurements. FCM similarly can give several different membership degrees for the lesions. However, the final classification decides that the lesions are still in the same cluster. This decision is taken because the membership degrees are considered maximum in the first and second measurements.

## CHAPTER 6

## CONCLUSION AND RECOMMENDATION

### 6.1 Conclusion

In this research, the problems of skin roughness measurement for psoriasis assessment are investigated. The investigation aims to provide a practical solution for clinical treatment in daily practice. Psoriasis appears as a red plaque lesion and covers a localised area of the body. In severe stages, the lesion can be widespread all over the body. The disease is caused by the wrong signals of the immune system that accelerates the cycle of skin cells' growth.

Psoriasis patients need periodic medical treatment continuously as the disease cannot be completely cured. The PASI (Psoriasis Area and Severity Index) scoring method is considered as gold standard for the severity assessment. To determine the psoriasis severity, the PASI scoring system defines four parameters, i.e. area (ratio of lesion area to total body surface area), erythema (colour of lesion inflammation), lesion thickness and scaliness. These parameters are scored at four body regions, namely head, trunk, upper limbs and lower limbs. In practice, an assessment of dermatologist can be subjective due to intra- and inter-rater variability of human assessment. The subjectivity is influenced either by the perception or by the clinical experience of the dermatologist.

An imaging approach is proposed to provide an objective PASI scaliness assessment based on skin surface roughness measurement. Several imaging solutions have been developed for measuring skin surface roughness. The earlier methods including mechanical surface profilometry, laser profilometry, light scattering, and speckle imaging have been applied in skin surface roughness measurement. However, these methods have some disadvantages, such as limited
resolution of profiler size and time-consuming process that occur in mechanical surface profilometry. The fast scanning also could not be performed by applying laser profilometry tough it can provide high scan resolution. Light scattering and speckle imaging methods are able to provide non-contact measurement and fast scanning. However, both methods require sophisticated system arrangement and standardised environment lighting. Moreover, the measured area is limited to a small dimension. Lastly, 3D surface acquired by structured light projection method is applied in the research. This method has advantages more than the aforementioned methods. The method can perform in vivo high speed scan at high resolution. Therefore it can overcome the scan problem on constant vibration of skin surface.

To achieve this objective, the PASI scaliness visual descriptors are studied and defined in terms on surface roughness. This feature can be measured by an imaging modality and its values are related to the PASI scaliness scores. An unsupervised clustering algorithm is applied to classify the lesion surface into the PASI scaliness scores. In this research work, the clustering algorithm is built based on the collected data from a clinical study at Department of Dermatology, Hospital Kuala Lumpur.

As stated in the Chapter 1, there are two research objectives of this thesis. The first objective is to develop a 3D imaging algorithm for accurately measuring the surface roughness of skin lesions. This first objective has been achieved in the research in which the algorithm has been validated for all skin lesions that appear at any body parts. A validation study furthermore is conducted on a set of lesion models. A total of 390 lesion models are pasted on a life-size mannequin to simulate a number of psoriasis lesions on human body. The life-size mannequin is suitable to model the human body curvature. The mannequin is commonly used in the research related with the surface determination such as body surface area measurement. The measurement methods are mostly applied in clothes designing and medical applications. Wurong and Bugao used life-size mannequin to validate a 3D reconstruction method for whole body surface imaging [154]. Wang et al. conducted a research on measurement of human body volumes. A laser scanner is applied to acquire 3D rigid object of the life-size mannequin. The body volume is computed from the 3D volume of the scanned mannequin. To validate the result, the mannequin is then immersed into a water tank to obtain body volume based on water
displacement [155]. The life-size mannequin is usually constructed based on human body modelling. This model is determined and developed from the real body proportions. As reported in [156], Jin Gu et al. have developed a human body model by stacking a set of skin contours. The contours are extracted from the edge images of the real human body.

Douros and Buxton have conducted a research on the construction of curvature maps for 3D human body surfaces. In this research they defined that the body skin surface is constructed by many skin patches. There are four basic shapes of the skin patches. The shapes are flat, elliptic, parabolic, and saddle surfaces. All of these basic shapes can be properly modelled by applying second order polynomial [157]. In this research smaller lesion surface areas are fitted with the second and third order polynomial. Moreover, the lesion area is also divided into four subdivided surfaces to reduce the surface curvedness. By applying this method, the lesion surface can be flattened and its vertical deviations are determined accurately.

From the validation study, it can be found that the developed surface roughness has an error of $0.0008 \pm 0.0017 \mathrm{~mm}\left(\right.$ Error $\left.\pm \sigma_{\text {Total }}\right)$ and gives accuracy of $94.12 \%$. This accuracy is quite high ( $>90 \%$ ) and therefore the developed algorithm can be implemented to measure real lesions. The measurement errors mostly are caused by the limitation of the scanning system. The system could not measure the lesion model surface accurately if a part of the lesion surface is in a deep concave area. In this area, the scanned surface might be located outside the depth of field interval ( $\pm 50 \mathrm{~mm}$ ).

There is no previous works on surface roughness measurement of psoriasis lesion. Most of the works were conducted on normal skin surfaces. Tchvialeva et al. [109] sumarised the skin surface roughness determinations using two different acqusition methods, structured light projection and speckle imaging [116] [158] [159]. The structured light projection method is evaluated for measuring skin replica and the actual skin surfaces. Jacobi et al. have conducted a research on skin surface measurement. The samples were taken from several regions such as forearm, back, and forehead with size $18 \mathrm{~mm} \times 13 \mathrm{~mm}$. A 3D scanner using structured light projection is applied to measure the skin surfaces. From the experiment, the average roughness of skin surfaces at back region is $0.0368 \pm 0.0094 \mathrm{~mm}$ whereas the surface
roughness of forearm region is $0.0302 \pm 0.0038 \mathrm{~mm}$ [160]. Table 6.1 summarises the comparison of surface roughness measurements on skin surfaces.

For the arm and hand regions, it can be shown that the surface roughness values are increasing from the structured light projection on skin replica to speckle imaging methods. The printing process of skin replica cannot completely duplicate the skin profile details with high frequencies. Only the skin profiles with lower frequencies and less variance are preserved by skin replica. Therefore its surface roughness is lower than in vivo structured light projection and speckle imaging methods. In 2D calculation, the surface roughness is determined higher than the other methods. This result can be caused by incompleteness of measured profiles. Here, the vertical deviations of the surface profile are extracted only from a single axis. The maximum surface roughness measured from psoriasis lesion $(0.1870 \mathrm{~mm})$ is higher than all surface roughness values presented in Table 6.1. This value shows that the psoriasis lesions surfaces are rougher than the normal skin surfaces.

Table 6.1 Comparison of skin surface roughness measurements.

|  | Average roughness of skin surfaces at several body regions (mm) |  |  |  |
| :---: | :---: | :---: | :---: | :---: |
| Method | Arm | Hand | Cheek | Back |
| Sturctured light projection on skin replica | $\begin{gathered} 0.0050-0.0078 \\ {[116][158][159]} \end{gathered}$ | 0.0122 [116] | 0.0110 [121] |  |
| In vivo sturctured light projection | $\begin{gathered} 0.0074-0.0184 \\ {[116][158][159]} \end{gathered}$ | 0.0134 [116] | 0.0430 [121] |  |
| Speckle imaging | $\begin{gathered} 0.0276 \pm 0.0076 \\ {[109]} \end{gathered}$ | $\begin{gathered} 0.0386 \pm 0.0056 \\ {[109]} \end{gathered}$ | $\begin{gathered} 0.0354 \pm 0.0064 \\ {[109]} \end{gathered}$ | $\begin{gathered} 0.0298 \pm 0.0092 \\ {[109]} \end{gathered}$ |
| Sturctured light projection, 2D calculation | $\begin{gathered} 0.0302 \pm 0.0038 \\ {[160]} \end{gathered}$ |  |  | $\begin{gathered} 0.0368 \pm 0.0094 \\ {[160]} \end{gathered}$ |
| Developed method on the surfaces of psoriasis lesion | 0.0100-0.1870 |  |  |  |

The second objective is to develop an objective and reliable PASI scaliness scoring. An unsupervised clustering is also applied to score PASI scaliness based on lesion surface roughness. $K$-means and fuzzy $c$-means (FCM) algorithms furthermore are studied in the research work. The implementation of $k$-means clustering algorithm has been validated by applying a clustering algorithm towards the divided dataset. The dataset of lesion roughness is randomly split into two equal
sized datasets. The clustering reliability is shown by the consistency of score centroids for all datasets. It proves that the score clusters are maximally separated. Centroid differences are not more than $3 \sigma_{\text {Total }}(0.005 \mathrm{~mm})$, except for a clustered data that have size less than 30 samples. Stability of FCM algorithms is shown by fittingness of Gaussian function to the membership functions of scaliness scores. All of the Gaussian function can be fitted with $R^{2} \approx 1$. The objectives of these clustering algorithms are indicated from the kappa agreement analysis. Since the scoring agreement between the dermatologists (0.55) is less than 0.81 , their assessment cannot be considered as a ground truth for evaluating the algorithm. 3D images from two successive scans are used to evaluate the scoring objectivity of the developed algorithm. The objectives of both clustering algorithms have been proven by the kappa coefficients of $k$-means and FCM algorithms. The coefficient value of $k$ means is 0.8473 whereas the coefficient of FCM is 0.8708 . FCM clustering, as the soft clustering, can solve the boundary problems better than $k$-means clustering and it therefore can give higher kappa agreement coefficient.

The research works in this thesis show that the developed algorithm enables to provide an objective measurement for PASI scaliness assessment. The developed algorithm resolves the problems of surface roughness determination on curved surfaces of human skin. The algorithm has been validated for accurately measuring the surface roughness of skin lesions at any body parts. These methods have been validated and accepted to be used in a clinical study involving registered 204 psoriasis patients of Hospital Kuala Lumpur. Furthermore, unsupervised clustering algorithm for scoring PASI scaliness has been developed. The algorithm is developed based on surface roughness data of the collected psoriasis lesions. High agreement between the first and the second assessments of psoriasis lesions demonstrates objectivity and reliability of the developed clustering algorithm.

### 6.2 Recommendations for Further Work

Currently, many scoring systems are still manually performed by dermatologist. As explained from this research, the assessment given by dermatologist might be subjective and can result in treatment inefficacy for the patient. Therefore, an objective assessment is still required in a clinical practice. Surface roughness is
known as one of the important parameters on skin analysis and assessment. The developed algorithm is potentially applied to another skin surface analysis that considers skin surface roughness as its parameter. The example of these skin problems, beside psoriasis, might be atopic dermatitis (atopic eczema), wrinkle analysis, and burnt skin. The surface roughness measurement either can subsequently be intended for monitoring on skin disease treatment or for any applications related to the cosmetics purposes.

Polynomial surface fitting has been validated to fit most of body surface areas. From this result, it can be shown that the algorithm is suitable for determining the curvature of human body surfaces. The maximum coverage area of the algorithm is limited by the scanning frame of 3D optical scanner. The size of this frame is $40 \times 30$ $\mathrm{mm}^{2}$. Lesions could be widely distributed on the body regions or appear as a single lesion covering a large area. To scan a larger area ( $>40 \times 30 \mathrm{~mm}^{2}$ ), multiple scans and assessments on adjacent areas are required, as shown in Figure 6.1. The lesions can be represented by several sampled areas. The average surface roughness for the assessed region would then be determined by averaging the surface roughness of the sampled areas.


Figure 6.1 Multiple scans for a single large lesion: (a) Large lesion covering part of the region and (b) Single large lesion covering the whole region.

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## APPENDIX A: DETERMINATION OF Z-SCORE

The Z value (Z-score) is determined to find normal distribution boundaries at a certain confidence level. As mentioned in previous section, the confidence level is usually set at $90 \%, 95 \%$, and $99 \%$. This appendix describes Z-score determination based on standard normal distribution table. The table maps the probabilities according to the Z-scores up to two decimal places. For instance, a confidence level is set at $95 \%$ ( $\alpha=0.05$ ). This level is then considered as a probability value at the normal distribution. The Z-score is determined by finding the probability listed in the cells. According to the table, probability 0.95 is located between $Z_{\alpha}=1.64$ and $Z_{\alpha}$ $=1.65$. Here, the probability covers the area to the left of $Z_{\alpha}$ and excludes the area to the right of $Z_{\alpha}$. This remaining area is considered as a single tail of the normal distribution with size of 0.05 . To provide two tails at left and right sides of the normal distribution, the remaining area $(\alpha=0.05)$ is then divided by two. For each tail, it gives an area of 0.025 . A new Z -score needs to be defined to represent these two-tailed areas.
$Z_{\alpha / 2}$ is used to indicate the boundary of two-tailed areas at the standard normal distribution. The $Z_{\alpha / 2}$ value is determined by searching the Z that provides probability $1-0.025=0.975$. By referring Table A.1, probability 0.975 is exactly located at $Z=1.9+0.06=1.96$. The rows and the columns are used to represent the tenth and the hundredth of Z respectively. Finally, it can be found that the two-tailed normal distribution with confidence level $95 \%$ is bounded by $Z_{\alpha / 2}=1.96$.

Table A. 1 Z-score of the standard normal distribution.

| $\mathbf{Z}$ | $\mathbf{0 . 0 0}$ | $\mathbf{0 . 0 1}$ | $\mathbf{0 . 0 2}$ | $\mathbf{0 . 0 3}$ | $\mathbf{0 . 0 4}$ | $\mathbf{0 . 0 5}$ | $\mathbf{0 . 0 6}$ | $\mathbf{0 . 0 7}$ | $\mathbf{0 . 0 8}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| $\mathbf{0 . 0}$ | 0.5000 | 0.5040 | 0.5080 | 0.5120 | 0.5160 | 0.5199 | 0.5239 | 0.5279 | 0.5319 |
| $\mathbf{0 . 1}$ | 0.5398 | 0.5438 | 0.5478 | 0.5517 | 0.5557 | 0.5596 | 0.5636 | 0.5675 | 0.5714 |
| $\ldots$ | $\ldots$ | $\ldots$ | $\ldots$ | $\ldots$ | $\ldots$ | $\ldots$ | $\ldots$ | $\ldots$ | $\ldots$ |
| $\mathbf{1 . 5}$ | 0.9332 | 0.9345 | 0.9357 | 0.9370 | 0.9382 | 0.9394 | 0.9406 | 0.9418 | 0.9429 |
| $\mathbf{1 . 6}$ | 0.9452 | 0.9463 | 0.9474 | 0.9484 | 0.9495 | 0.9505 | 0.9515 | 0.9525 | 0.9535 |
| $\mathbf{1 . 7}$ | 0.9554 | 0.9564 | 0.9573 | 0.9582 | 0.9591 | 0.9599 | 0.9608 | 0.9616 | 0.9625 |
| $\mathbf{1 . 8}$ | 0.9641 | 0.9649 | 0.9656 | 0.9664 | 0.9671 | 0.9678 | 0.9686 | 0.9693 | 0.9699 |
| $\mathbf{1 . 9}$ | 0.9713 | 0.9719 | 0.9726 | 0.9732 | 0.9738 | 0.9744 | $\mathbf{0 . 9 7 5 0}$ | 0.9756 | 0.9761 |
| $\mathbf{2 . 0}$ | 0.9772 | 0.9778 | 0.9783 | 0.9788 | 0.9793 | 0.9798 | 0.9803 | 0.9808 | 0.9812 |

## APPENDIX B: SURFACE ROUGHNESS OF LESION MODEL AT FLAT SURFACES

| Lesion model | Assessment | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $S_{\text {final }}$ <br> (mm) | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | 1 | 12.23 | 10.44 | 0.0142 | 0.997 | 0.0138 | 0.997 | 0.0138 | 0.997 | 0.0139 |
|  | 2 | 12.23 | 10.44 | 0.0146 | 0.996 | 0.0142 | 0.997 | 0.0142 | 0.997 |  |
|  | 3 | 12.23 | 10.44 | 0.0140 | 0.996 | 0.0137 | 0.997 | 0.0137 | 0.997 |  |
| 2 | 1 | 14.65 | 10.31 | 0.0123 | 0.998 | 0.0120 | 0.998 | 0.0120 | 0.998 | 0.0130 |
|  | 2 | 14.65 | 10.31 | 0.0122 | 0.998 | 0.0119 | 0.998 | 0.0119 | 0.998 |  |
|  | 3 | 14.65 | 10.31 | 0.0151 | 0.997 | 0.0149 | 0.997 | 0.0149 | 0.997 |  |
| 3 | 1 | 14.65 | 10.31 | 0.0124 | 0.994 | 0.0122 | 0.994 | 0.0122 | 0.994 | 0.0122 |
|  | 2 | 14.65 | 10.31 | 0.0125 | 0.994 | 0.0122 | 0.994 | 0.0122 | 0.994 |  |
|  | 3 | 14.65 | 10.31 | 0.0123 | 0.994 | 0.0121 | 0.994 | 0.0121 | 0.994 |  |
| 4 | 1 | 13.88 | 9.80 | 0.0123 | 0.994 | 0.0119 | 0.994 | 0.0119 | 0.994 | 0.0118 |
|  | 2 | 13.88 | 9.80 | 0.0122 | 0.994 | 0.0118 | 0.994 | 0.0118 | 0.994 |  |
|  | 3 | 13.88 | 9.80 | 0.0123 | 0.994 | 0.0118 | 0.994 | 0.0118 | 0.994 |  |
| 5 | 1 | 12.86 | 9.67 | 0.0134 | 0.995 | 0.0132 | 0.995 | 0.0132 | 0.995 | 0.0149 |
|  | 2 | 12.86 | 9.67 | 0.0187 | 0.992 | 0.0186 | 0.992 | 0.0186 | 0.992 |  |
|  | 3 | 12.86 | 9.67 | 0.0130 | 0.996 | 0.0129 | 0.996 | 0.0129 | 0.996 |  |
| 6 | 1 | 12.10 | 9.93 | 0.0146 | 0.950 | 0.0139 | 0.954 | 0.0139 | 0.954 | 0.0132 |
|  | 2 | 12.10 | 9.93 | 0.0129 | 0.952 | 0.0122 | 0.957 | 0.0122 | 0.957 |  |
|  | 3 | 12.10 | 9.93 | 0.0141 | 0.921 | 0.0135 | 0.928 | 0.0135 | 0.928 |  |
| 7 | 1 | 11.21 | 9.67 | 0.0112 | 0.997 | 0.0111 | 0.997 | 0.0111 | 0.997 | 0.0112 |
|  | 2 | 11.21 | 9.67 | 0.0115 | 0.997 | 0.0114 | 0.997 | 0.0114 | 0.997 |  |
|  | 3 | 11.21 | 9.67 | 0.0112 | 0.997 | 0.0112 | 0.997 | 0.0112 | 0.997 |  |
| 8 | 1 | 12.35 | 9.67 | 0.0128 | 0.993 | 0.0122 | 0.994 | 0.0122 | 0.994 | 0.0114 |
|  | 2 | 12.35 | 9.67 | 0.0118 | 0.994 | 0.0110 | 0.995 | 0.0110 | 0.995 |  |
|  | 3 | 12.35 | 9.67 | 0.0117 | 0.994 | 0.0109 | 0.995 | 0.0109 | 0.995 |  |
| 9 | 1 | 14.14 | 9.55 | 0.0125 | 0.980 | 0.0122 | 0.981 | 0.0122 | 0.981 | 0.0122 |
|  | 2 | 14.14 | 9.55 | 0.0124 | 0.981 | 0.0121 | 0.982 | 0.0121 | 0.982 |  |
|  | 3 | 14.14 | 9.55 | 0.0125 | 0.982 | 0.0122 | 0.983 | 0.0122 | 0.983 |  |
| 10 | 1 | 13.25 | 9.29 | 0.0113 | 0.993 | 0.0110 | 0.994 | 0.0110 | 0.994 | 0.0109 |
|  | 2 | 13.25 | 9.29 | 0.0109 | 0.994 | 0.0105 | 0.994 | 0.0105 | 0.994 |  |
|  | 3 | 13.25 | 9.29 | 0.0116 | 0.993 | 0.0112 | 0.993 | 0.0112 | 0.993 |  |
| 11 | 1 | 12.23 | 9.93 | 0.0133 | 0.987 | 0.0129 | 0.987 | 0.0129 | 0.987 | 0.0124 |
|  | 2 | 12.23 | 9.93 | 0.0119 | 0.990 | 0.0116 | 0.990 | 0.0116 | 0.990 |  |
|  | 3 | 12.23 | 9.93 | 0.0131 | 0.988 | 0.0128 | 0.989 | 0.0128 | 0.989 |  |
| 12 | 1 | 11.59 | 9.67 | 0.0119 | 0.985 | 0.0115 | 0.986 | 0.0115 | 0.986 | 0.0118 |
|  | 2 | 11.59 | 9.67 | 0.0127 | 0.983 | 0.0124 | 0.985 | 0.0124 | 0.985 |  |
|  | 3 | 11.59 | 9.67 | 0.0120 | 0.985 | 0.0116 | 0.986 | 0.0116 | 0.986 |  |


| Lesion model | Assessment | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $S_{\text {final }}$ <br> (mm) | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 13 | 1 | 12.99 | 10.05 | 0.0113 | 0.990 | 0.0109 | 0.991 | 0.0109 | 0.991 | 0.0108 |
|  | 2 | 12.99 | 10.05 | 0.0110 | 0.991 | 0.0107 | 0.991 | 0.0107 | 0.991 |  |
|  | 3 | 12.99 | 10.05 | 0.0113 | 0.991 | 0.0109 | 0.991 | 0.0109 | 0.991 |  |
| 14 | 1 | 10.06 | 9.93 | 0.0134 | 0.972 | 0.0127 | 0.975 | 0.0127 | 0.975 | 0.0117 |
|  | 2 | 10.06 | 9.93 | 0.0117 | 0.980 | 0.0109 | 0.983 | 0.0109 | 0.983 |  |
|  | 3 | 10.06 | 9.93 | 0.0122 | 0.976 | 0.0115 | 0.979 | 0.0115 | 0.979 |  |
| 15 | 1 | 10.44 | 9.93 | 0.0124 | 0.991 | 0.0116 | 0.992 | 0.0116 | 0.992 | 0.0115 |
|  | 2 | 10.44 | 9.93 | 0.0121 | 0.992 | 0.0115 | 0.993 | 0.0115 | 0.993 |  |
|  | 3 | 10.44 | 9.93 | 0.0121 | 0.992 | 0.0114 | 0.992 | 0.0114 | 0.992 |  |
| 16 | 1 | 11.59 | 10.56 | 0.0114 | 0.993 | 0.0111 | 0.994 | 0.0111 | 0.994 | 0.0110 |
|  | 2 | 11.59 | 10.56 | 0.0113 | 0.993 | 0.0110 | 0.993 | 0.0110 | 0.993 |  |
|  | 3 | 11.59 | 10.56 | 0.0112 | 0.993 | 0.0109 | 0.993 | 0.0109 | 0.993 |  |
| 17 | 1 | 11.21 | 9.93 | 0.0161 | 0.988 | 0.0152 | 0.989 | 0.0152 | 0.989 | 0.0131 |
|  | 2 | 11.21 | 9.93 | 0.0130 | 0.992 | 0.0119 | 0.993 | 0.0119 | 0.993 |  |
|  | 3 | 11.21 | 9.93 | 0.0135 | 0.992 | 0.0123 | 0.993 | 0.0123 | 0.993 |  |
| 18 | 1 | 11.21 | 10.18 | 0.0122 | 0.997 | 0.0113 | 0.998 | 0.0113 | 0.998 | 0.0114 |
|  | 2 | 11.21 | 10.18 | 0.0126 | 0.997 | 0.0118 | 0.997 | 0.0118 | 0.997 |  |
|  | 3 | 11.21 | 10.18 | 0.0121 | 0.997 | 0.0112 | 0.998 | 0.0112 | 0.998 |  |
| 19 | 1 | 11.59 | 10.05 | 0.0169 | 0.992 | 0.0166 | 0.992 | 0.0166 | 0.992 | 0.0137 |
|  | 2 | 11.59 | 10.05 | 0.0128 | 0.995 | 0.0125 | 0.995 | 0.0125 | 0.995 |  |
|  | 3 | 11.59 | 10.05 | 0.0120 | 0.996 | 0.0118 | 0.996 | 0.0118 | 0.996 |  |
| 20 | 1 | 12.61 | 10.56 | 0.0117 | 0.998 | 0.0106 | 0.998 | 0.0106 | 0.998 | 0.0108 |
|  | 2 | 12.61 | 10.56 | 0.0123 | 0.997 | 0.0111 | 0.998 | 0.0111 | 0.998 |  |
|  | 3 | 12.61 | 10.56 | 0.0116 | 0.998 | 0.0106 | 0.998 | 0.0106 | 0.998 |  |
| 21 | 1 | 13.50 | 10.82 | 0.0134 | 0.972 | 0.0128 | 0.975 | 0.0128 | 0.975 | 0.0118 |
|  | 2 | 13.50 | 10.82 | 0.0124 | 0.977 | 0.0116 | 0.979 | 0.0116 | 0.979 |  |
|  | 3 | 13.50 | 10.82 | 0.0119 | 0.975 | 0.0111 | 0.979 | 0.0111 | 0.979 |  |
| 22 | 1 | 13.76 | 10.44 | 0.0127 | 0.989 | 0.0123 | 0.990 | 0.0123 | 0.990 | 0.0124 |
|  | 2 | 13.76 | 10.44 | 0.0134 | 0.987 | 0.0130 | 0.988 | 0.0130 | 0.988 |  |
|  | 3 | 13.76 | 10.44 | 0.0123 | 0.990 | 0.0120 | 0.990 | 0.0120 | 0.990 |  |
| 23 | 1 | 11.72 | 10.56 | 0.0133 | 0.971 | 0.0126 | 0.974 | 0.0126 | 0.974 | 0.0125 |
|  | 2 | 11.72 | 10.56 | 0.0128 | 0.973 | 0.0123 | 0.976 | 0.0123 | 0.976 |  |
|  | 3 | 11.72 | 10.56 | 0.0133 | 0.969 | 0.0126 | 0.972 | 0.0126 | 0.972 |  |
| 24 | 1 | 13.12 | 10.05 | 0.0129 | 0.958 | 0.0128 | 0.959 | 0.0128 | 0.959 | 0.0129 |
|  | 2 | 13.12 | 10.05 | 0.0126 | 0.966 | 0.0125 | 0.966 | 0.0125 | 0.966 |  |
|  | 3 | 13.12 | 10.05 | 0.0136 | 0.962 | 0.0134 | 0.963 | 0.0134 | 0.963 |  |
| 25 | 1 | 12.48 | 10.05 | 0.0122 | 0.996 | 0.0113 | 0.997 | 0.0113 | 0.997 | 0.0109 |
|  | 2 | 12.48 | 10.05 | 0.0116 | 0.996 | 0.0108 | 0.997 | 0.0108 | 0.997 |  |
|  | 3 | 12.48 | 10.05 | 0.0114 | 0.997 | 0.0106 | 0.997 | 0.0106 | 0.997 |  |
| 26 | 1 | 12.10 | 10.44 | 0.0127 | 0.990 | 0.0124 | 0.991 | 0.0124 | 0.991 | 0.0127 |
|  | 2 | 12.10 | 10.44 | 0.0131 | 0.986 | 0.0126 | 0.989 | 0.0126 | 0.989 |  |
|  | 3 | 12.10 | 10.44 | 0.0136 | 0.984 | 0.0131 | 0.987 | 0.0131 | 0.987 |  |


| Lesion model | Assessment | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $\begin{aligned} & S_{\text {final }} \\ & (\mathrm{mm}) \end{aligned}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 27 | 1 | 12.23 | 10.18 | 0.0129 | 0.997 | 0.0124 | 0.997 | 0.0124 | 0.997 | 0.0120 |
|  | 2 | 12.23 | 10.18 | 0.0123 | 0.998 | 0.0118 | 0.998 | 0.0118 | 0.998 |  |
|  | 3 | 12.23 | 10.18 | 0.0124 | 0.998 | 0.0119 | 0.998 | 0.0119 | 0.998 |  |
| 28 | 1 | 14.14 | 10.69 | 0.0133 | 0.997 | 0.0130 | 0.997 | 0.0130 | 0.997 | 0.0130 |
|  | 2 | 14.14 | 10.69 | 0.0139 | 0.996 | 0.0135 | 0.997 | 0.0135 | 0.997 |  |
|  | 3 | 14.14 | 10.69 | 0.0129 | 0.997 | 0.0125 | 0.997 | 0.0125 | 0.997 |  |
| 29 | 1 | 14.65 | 9.5450 | 0.0116 | 0.985 | 0.0115 | 0.985 | 0.0115 | 0.985 | 0.0116 |
|  | 2 | 15.16 | 8.7820 | 0.0119 | 0.984 | 0.0116 | 0.985 | 0.0116 | 0.985 |  |
|  | 3 | 15.92 | 9.2910 | 0.0117 | 0.986 | 0.0116 | 0.986 | 0.0116 | 0.986 |  |
| 30 | 1 | 14.52 | 8.5280 | 0.0113 | 0.989 | 0.0109 | 0.989 | 0.0109 | 0.989 | 0.0109 |
|  | 2 | 13.63 | 8.9090 | 0.0112 | 0.988 | 0.0109 | 0.988 | 0.0109 | 0.988 |  |
|  | 3 | 14.65 | 9.1640 | 0.0112 | 0.989 | 0.0108 | 0.990 | 0.0108 | 0.990 |  |
| 31 | 1 | 15.41 | 8.6550 | 0.0112 | 0.986 | 0.0111 | 0.986 | 0.0111 | 0.986 | 0.0112 |
|  | 2 | 14.65 | 8.5270 | 0.0114 | 0.984 | 0.0113 | 0.984 | 0.0113 | 0.984 |  |
|  | 3 | 15.03 | 8.5270 | 0.0113 | 0.985 | 0.0111 | 0.985 | 0.0111 | 0.985 |  |
| 32 | 1 | 0.0146 | 0.965 | 0.0147 | 0.964 | 0.0146 | 0.965 | 0.0146 | 0.965 | 0.0145 |
|  | 2 | 0.0117 | 0.977 | 0.0119 | 0.976 | 0.0117 | 0.977 | 0.0117 | 0.977 |  |
|  | 3 | 0.0172 | 0.954 | 0.0175 | 0.952 | 0.0172 | 0.954 | 0.0172 | 0.954 |  |
| 33 | 1 | 0.0096 | 0.977 | 0.0095 | 0.978 | 0.0095 | 0.978 | 0.0095 | 0.978 | 0.0121 |
|  | 2 | 0.0124 | 0.965 | 0.0123 | 0.965 | 0.0123 | 0.965 | 0.0123 | 0.965 |  |
|  | 3 | 0.0147 | 0.948 | 0.0146 | 0.949 | 0.0146 | 0.949 | 0.0146 | 0.949 |  |

## APPENDIX C: SURFACE ROUGHNESS OF LESION MODEL AT

 CURVE SURFACES| No | Lesion ID | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | H-2 | 10.05 | 9.04 | 0.014 | 0.996 | 0.015 | 0.995 | 0.996 | 0.014 |
| 2 | H-4 | 11.58 | 9.42 | 0.015 | 0.991 | 0.015 | 0.991 | 0.991 | 0.015 |
| 3 | H-5 | 11.96 | 7.77 | 0.015 | 0.999 | 0.015 | 0.999 | 0.999 | 0.015 |
| 4 | H-6 | 13.49 | 8.40 | 0.013 | 0.996 | 0.013 | 0.996 | 0.996 | 0.013 |
| 5 | H-7 | 9.54 | 7.64 | 0.014 | 0.998 | 0.014 | 0.998 | 0.998 | 0.014 |
| 6 | H-10 | 11.71 | 6.87 | 0.014 | 0.981 | 0.014 | 0.982 | 0.982 | 0.014 |
| 7 | U-2 | 8.53 | 5.09 | 0.015 | 0.987 | 0.015 | 0.987 | 0.987 | 0.015 |
| 8 | U-4 | 9.80 | 4.84 | 0.012 | 0.971 | 0.013 | 0.964 | 0.971 | 0.012 |
| 9 | U-5 | 9.67 | 16.42 | 0.012 | 0.998 | 0.012 | 0.998 | 0.998 | 0.012 |
| 10 | U-6 | 10.18 | 3.82 | 0.011 | 0.996 | 0.013 | 0.994 | 0.996 | 0.011 |
| 11 | U-7 | 8.53 | 4.07 | 0.012 | 0.970 | 0.012 | 0.964 | 0.970 | 0.012 |
| 12 | U-8 | 10.56 | 4.07 | 0.012 | 0.995 | 0.013 | 0.994 | 0.995 | 0.012 |
| 13 | U-9 | 7.51 | 4.33 | 0.012 | 0.995 | 0.012 | 0.995 | 0.995 | 0.012 |
| 14 | U-10 | 8.53 | 4.20 | 0.012 | 0.963 | 0.012 | 0.961 | 0.963 | 0.012 |
| 15 | U-11 | 10.18 | 16.42 | 0.013 | 0.997 | 0.015 | 0.997 | 0.997 | 0.013 |
| 16 | U-12 | 9.54 | 5.47 | 0.012 | 0.994 | 0.014 | 0.992 | 0.994 | 0.012 |
| 17 | U-13 | 9.29 | 3.95 | 0.015 | 0.985 | 0.015 | 0.984 | 0.985 | 0.015 |
| 18 | U-14 | 9.54 | 4.33 | 0.011 | 0.991 | 0.013 | 0.988 | 0.991 | 0.011 |
| 19 | U-15 | 9.03 | 4.58 | 0.012 | 0.960 | 0.012 | 0.961 | 0.961 | 0.012 |
| 20 | U-16 | 8.91 | 3.69 | 0.012 | 0.998 | 0.015 | 0.998 | 0.998 | 0.012 |
| 21 | U-17 | 9.80 | 15.78 | 0.011 | 0.988 | 0.013 | 0.989 | 0.989 | 0.013 |
| 22 | U-18 | 10.43 | 16.42 | 0.012 | 0.996 | 0.016 | 0.993 | 0.996 | 0.012 |
| 23 | U-19 | 8.91 | 4.46 | 0.013 | 0.993 | 0.016 | 0.988 | 0.993 | 0.013 |
| 24 | U-20 | 15.27 | 7.26 | 0.014 | 0.989 | 0.014 | 0.990 | 0.990 | 0.014 |
| 25 | U-23 | 9.67 | 11.08 | 0.013 | 0.986 | 0.014 | 0.982 | 0.986 | 0.013 |
| 26 | U-24 | 9.67 | 5.35 | 0.011 | 0.994 | 0.012 | 0.993 | 0.994 | 0.011 |
| 27 | U-25 | 7.76 | 5.35 | 0.012 | 0.991 | 0.013 | 0.989 | 0.991 | 0.012 |
| 28 | U-26 | 9.54 | 5.09 | 0.011 | 0.992 | 0.011 | 0.991 | 0.992 | 0.011 |
| 29 | U-27 | 9.93 | 16.93 | 0.011 | 0.990 | 0.013 | 0.988 | 0.990 | 0.011 |
| 30 | U-28 | 10.05 | 4.84 | 0.012 | 0.993 | 0.012 | 0.992 | 0.993 | 0.012 |
| 31 | U-30 | 8.40 | 15.40 | 0.015 | 0.996 | 0.011 | 0.998 | 0.998 | 0.011 |
| 32 | U-31 | 10.05 | 4.58 | 0.012 | 0.989 | 0.012 | 0.989 | 0.989 | 0.012 |
| 33 | U-32 | 9.92 | 15.66 | 0.014 | 0.998 | 0.020 | 0.996 | 0.998 | 0.014 |
| 34 | U-33 | 9.54 | 4.20 | 0.013 | 0.984 | 0.013 | 0.985 | 0.985 | 0.013 |
| 35 | U-34 | 9.92 | 16.55 | 0.012 | 0.994 | 0.012 | 0.994 | 0.994 | 0.012 |
| 36 | U-35 | 9.67 | 15.78 | 0.011 | 0.989 | 0.013 | 0.985 | 0.989 | 0.011 |
| 37 | U-36 | 9.16 | 5.22 | 0.012 | 0.956 | 0.012 | 0.956 | 0.956 | 0.012 |


| No | Lesion ID | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{gathered} \overline{S_{\text {final }}} \\ (\mathrm{mm}) \end{gathered}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 38 | U-38 | 9.67 | 14.51 | 0.012 | 0.998 | 0.014 | 0.997 | 0.998 | 0.012 |
| 39 | U-39 | 9.16 | 5.73 | 0.011 | 0.991 | 0.010 | 0.991 | 0.991 | 0.011 |
| 40 | U-40 | 9.54 | 15.40 | 0.012 | 0.996 | 0.013 | 0.995 | 0.996 | 0.012 |
| 41 | U-41 | 9.67 | 3.95 | 0.011 | 0.973 | 0.012 | 0.967 | 0.973 | 0.011 |
| 42 | U-42 | 9.42 | 15.66 | 0.011 | 0.993 | 0.013 | 0.990 | 0.993 | 0.011 |
| 43 | U-43 | 10.05 | 5.35 | 0.012 | 0.984 | 0.012 | 0.984 | 0.984 | 0.012 |
| 44 | U-44 | 6.87 | 4.58 | 0.014 | 0.975 | 0.013 | 0.978 | 0.978 | 0.013 |
| 45 | U-45 | 9.42 | 3.69 | 0.011 | 0.995 | 0.011 | 0.996 | 0.996 | 0.011 |
| 46 | U-46 | 9.54 | 4.33 | 0.010 | 0.928 | 0.011 | 0.945 | 0.945 | 0.011 |
| 47 | U-47 | 10.56 | 5.73 | 0.011 | 0.992 | 0.011 | 0.992 | 0.992 | 0.011 |
| 48 | U-48 | 10.82 | 15.15 | 0.011 | 0.993 | 0.012 | 0.993 | 0.993 | 0.011 |
| 49 | U-49 | 10.56 | 4.58 | 0.012 | 0.980 | 0.013 | 0.979 | 0.980 | 0.012 |
| 50 | U-50 | 9.03 | 4.71 | 0.013 | 0.948 | 0.013 | 0.948 | 0.948 | 0.013 |
| 51 | U-52 | 8.53 | 4.07 | 0.015 | 0.968 | 0.017 | 0.961 | 0.968 | 0.015 |
| 52 | U-53 | 9.03 | 3.95 | 0.011 | 0.974 | 0.012 | 0.973 | 0.974 | 0.011 |
| 53 | U-54 | 9.67 | 5.86 | 0.012 | 0.969 | 0.013 | 0.962 | 0.969 | 0.012 |
| 54 | U-55 | 10.18 | 4.97 | 0.014 | 0.983 | 0.014 | 0.983 | 0.983 | 0.014 |
| 55 | U-56 | 9.03 | 6.11 | 0.011 | 0.984 | 0.011 | 0.982 | 0.984 | 0.011 |
| 56 | U-57 | 9.16 | 4.84 | 0.011 | 0.910 | 0.012 | 0.920 | 0.920 | 0.012 |
| 57 | U-58 | 9.92 | 5.35 | 0.011 | 0.963 | 0.012 | 0.956 | 0.963 | 0.011 |
| 58 | U-59 | 9.93 | 4.71 | 0.015 | 0.984 | 0.014 | 0.986 | 0.986 | 0.014 |
| 59 | U-60 | 9.16 | 3.57 | 0.013 | 0.936 | 0.013 | 0.934 | 0.936 | 0.013 |
| 60 | U-62 | 9.80 | 4.46 | 0.011 | 0.997 | 0.012 | 0.997 | 0.997 | 0.011 |
| 61 | U-63 | 9.29 | 5.09 | 0.011 | 0.986 | 0.012 | 0.983 | 0.986 | 0.011 |
| 62 | U-64 | 9.16 | 4.71 | 0.011 | 0.926 | 0.011 | 0.939 | 0.939 | 0.011 |
| 63 | U-65 | 9.29 | 5.09 | 0.011 | 0.984 | 0.011 | 0.983 | 0.984 | 0.011 |
| 64 | U-66 | 10.56 | 17.69 | 0.011 | 0.990 | 0.013 | 0.988 | 0.990 | 0.011 |
| 65 | U-67 | 9.42 | 5.22 | 0.014 | 0.958 | 0.014 | 0.954 | 0.958 | 0.014 |
| 66 | U-68 | 9.03 | 4.71 | 0.014 | 0.991 | 0.015 | 0.991 | 0.991 | 0.014 |
| 67 | U-69 | 9.03 | 3.31 | 0.012 | 0.996 | 0.012 | 0.996 | 0.996 | 0.012 |
| 68 | U-70 | 10.18 | 4.20 | 0.012 | 0.972 | 0.012 | 0.969 | 0.972 | 0.012 |
| 69 | U-71 | 10.82 | 17.06 | 0.015 | 0.996 | 0.013 | 0.996 | 0.996 | 0.015 |
| 70 | U-72 | 9.67 | 4.71 | 0.013 | 0.985 | 0.013 | 0.985 | 0.985 | 0.013 |
| 71 | U-73 | 10.05 | 4.58 | 0.015 | 0.992 | 0.015 | 0.992 | 0.992 | 0.015 |
| 72 | U-75 | 9.54 | 5.73 | 0.012 | 0.991 | 0.012 | 0.990 | 0.991 | 0.012 |
| 73 | U-76 | 8.27 | 5.86 | 0.014 | 0.984 | 0.014 | 0.983 | 0.984 | 0.014 |
| 74 | U-77 | 9.54 | 16.04 | 0.012 | 0.999 | 0.015 | 0.998 | 0.999 | 0.012 |
| 75 | U-78 | 10.69 | 14.77 | 0.011 | 0.997 | 0.011 | 0.997 | 0.997 | 0.011 |
| 76 | U-79 | 10.56 | 16.42 | 0.012 | 0.997 | 0.011 | 0.997 | 0.997 | 0.012 |
| 77 | U-80 | 9.93 | 15.79 | 0.011 | 0.996 | 0.012 | 0.995 | 0.996 | 0.011 |
| 78 | U-81 | 10.43 | 17.44 | 0.011 | 0.995 | 0.011 | 0.995 | 0.995 | 0.011 |
| 79 | U-82 | 10.69 | 5.09 | 0.012 | 0.984 | 0.013 | 0.983 | 0.984 | 0.012 |
| 80 | U-83 | 10.05 | 16.04 | 0.013 | 0.994 | 0.014 | 0.994 | 0.994 | 0.013 |


| No | Lesion ID | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 81 | U-84 | 10.43 | 5.09 | 0.012 | 0.997 | 0.014 | 0.996 | 0.997 | 0.012 |
| 82 | U-87 | 10.31 | 5.98 | 0.012 | 0.998 | 0.012 | 0.998 | 0.998 | 0.012 |
| 83 | U-88 | 10.05 | 6.24 | 0.014 | 0.978 | 0.014 | 0.980 | 0.980 | 0.014 |
| 84 | U-89 | 10.69 | 5.47 | 0.011 | 0.992 | 0.011 | 0.992 | 0.992 | 0.011 |
| 85 | U-90 | 9.93 | 5.22 | 0.013 | 0.997 | 0.014 | 0.997 | 0.997 | 0.013 |
| 86 | U-91 | 10.43 | 4.71 | 0.012 | 0.993 | 0.013 | 0.993 | 0.993 | 0.012 |
| 87 | U-92 | 8.02 | 17.31 | 0.012 | 0.988 | 0.012 | 0.987 | 0.988 | 0.012 |
| 88 | U-93 | 10.18 | 4.84 | 0.011 | 0.987 | 0.012 | 0.984 | 0.987 | 0.011 |
| 89 | U-94 | 8.40 | 17.70 | 0.013 | 0.991 | 0.014 | 0.990 | 0.991 | 0.013 |
| 90 | U-95 | 10.18 | 5.22 | 0.012 | 0.930 | 0.014 | 0.914 | 0.930 | 0.012 |
| 91 | U-96 | 10.82 | 5.09 | 0.011 | 0.993 | 0.011 | 0.993 | 0.993 | 0.011 |
| 92 | U-97 | 9.80 | 5.35 | 0.013 | 0.969 | 0.013 | 0.968 | 0.969 | 0.013 |
| 93 | U-98 | 10.18 | 16.55 | 0.012 | 0.988 | 0.014 | 0.986 | 0.988 | 0.012 |
| 94 | T-1 | 14.12 | 8.02 | 0.018 | 0.977 | 0.011 | 0.992 | 0.992 | 0.011 |
| 95 | T-2 | 13.23 | 7.00 | 0.011 | 0.949 | 0.012 | 0.940 | 0.949 | 0.011 |
| 96 | T-3 | 15.01 | 7.77 | 0.012 | 0.944 | 0.015 | 0.905 | 0.944 | 0.012 |
| 97 | T-4 | 9.93 | 6.49 | 0.013 | 0.992 | 0.013 | 0.992 | 0.992 | 0.013 |
| 98 | T-5 | 12.34 | 7.13 | 0.014 | 0.990 | 0.014 | 0.990 | 0.990 | 0.014 |
| 99 | T-6 | 11.71 | 6.24 | 0.015 | 0.999 | 0.018 | 0.999 | 0.999 | 0.015 |
| 100 | T-7 | 10.82 | 7.26 | 0.011 | 0.988 | 0.012 | 0.985 | 0.988 | 0.011 |
| 101 | T-8 | 12.22 | 7.51 | 0.012 | 0.977 | 0.016 | 0.962 | 0.977 | 0.012 |
| 102 | T-9 | 12.85 | 6.49 | 0.013 | 0.988 | 0.013 | 0.989 | 0.989 | 0.013 |
| 103 | T-10 | 12.98 | 10.18 | 0.016 | 0.989 | 0.017 | 0.988 | 0.989 | 0.016 |
| 104 | T-11 | 11.20 | 6.62 | 0.012 | 0.968 | 0.011 | 0.971 | 0.971 | 0.011 |
| 105 | T-12 | 13.23 | 8.40 | 0.012 | 0.986 | 0.013 | 0.985 | 0.986 | 0.012 |
| 106 | T-13 | 11.83 | 7.51 | 0.012 | 0.989 | 0.014 | 0.974 | 0.989 | 0.012 |
| 107 | T-14 | 11.20 | 9.55 | 0.014 | 0.988 | 0.013 | 0.988 | 0.988 | 0.014 |
| 108 | T-16 | 10.56 | 8.02 | 0.013 | 0.983 | 0.014 | 0.977 | 0.983 | 0.013 |
| 109 | T-17 | 9.67 | 8.40 | 0.015 | 0.983 | 0.015 | 0.983 | 0.983 | 0.015 |
| 110 | T-18 | 10.31 | 7.77 | 0.014 | 0.996 | 0.016 | 0.995 | 0.996 | 0.014 |
| 111 | T-19 | 13.11 | 7.89 | 0.012 | 0.962 | 0.012 | 0.981 | 0.981 | 0.012 |
| 112 | T-20 | 9.42 | 8.28 | 0.014 | 0.997 | 0.014 | 0.997 | 0.997 | 0.014 |
| 113 | T-21 | 10.43 | 7.26 | 0.012 | 0.993 | 0.013 | 0.993 | 0.993 | 0.012 |
| 114 | T-22 | 13.11 | 9.80 | 0.011 | 0.991 | 0.011 | 0.991 | 0.991 | 0.011 |
| 115 | T-23 | 12.72 | 8.15 | 0.011 | 0.974 | 0.011 | 0.975 | 0.975 | 0.011 |
| 116 | T-24 | 17.56 | 8.40 | 0.012 | 0.933 | 0.012 | 0.933 | 0.933 | 0.012 |
| 117 | T-25 | 12.09 | 7.77 | 0.012 | 0.963 | 0.012 | 0.964 | 0.964 | 0.012 |
| 118 | T-26 | 12.72 | 7.26 | 0.012 | 0.995 | 0.012 | 0.995 | 0.995 | 0.012 |
| 119 | T-28 | 12.72 | 9.42 | 0.014 | 0.963 | 0.014 | 0.964 | 0.964 | 0.014 |
| 120 | T-29 | 13.49 | 7.13 | 0.013 | 0.946 | 0.014 | 0.958 | 0.958 | 0.014 |
| 121 | T-30 | 12.34 | 7.26 | 0.011 | 0.950 | 0.012 | 0.942 | 0.950 | 0.011 |
| 122 | T-31 | 11.83 | 7.89 | 0.014 | 0.924 | 0.014 | 0.930 | 0.930 | 0.014 |
| 123 | T-32 | 15.01 | 9.29 | 0.013 | 0.979 | 0.017 | 0.964 | 0.979 | 0.013 |


| No | Lesion ID | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 124 | T-33 | 10.56 | 7.13 | 0.014 | 0.998 | 0.014 | 0.998 | 0.998 | 0.014 |
| 125 | T-34 | 12.60 | 9.04 | 0.013 | 0.985 | 0.015 | 0.978 | 0.985 | 0.013 |
| 126 | T-35 | 12.22 | 7.51 | 0.012 | 0.985 | 0.013 | 0.984 | 0.985 | 0.012 |
| 127 | T-36 | 15.01 | 7.26 | 0.013 | 0.980 | 0.013 | 0.981 | 0.981 | 0.013 |
| 128 | T-37 | 10.56 | 7.89 | 0.013 | 0.943 | 0.012 | 0.950 | 0.950 | 0.012 |
| 129 | T-38 | 13.23 | 8.91 | 0.013 | 0.996 | 0.014 | 0.995 | 0.996 | 0.013 |
| 130 | T-39 | 14.38 | 9.93 | 0.013 | 0.996 | 0.015 | 0.995 | 0.996 | 0.013 |
| 131 | T-40 | 15.65 | 8.78 | 0.011 | 0.993 | 0.012 | 0.992 | 0.993 | 0.011 |
| 132 | T-41 | 13.23 | 9.55 | 0.013 | 0.997 | 0.013 | 0.997 | 0.997 | 0.013 |
| 133 | T-42 | 14.51 | 7.13 | 0.013 | 0.991 | 0.012 | 0.992 | 0.992 | 0.012 |
| 134 | T-43 | 13.11 | 8.02 | 0.013 | 0.990 | 0.012 | 0.990 | 0.990 | 0.013 |
| 135 | T-45 | 14.12 | 9.80 | 0.013 | 0.988 | 0.013 | 0.986 | 0.988 | 0.013 |
| 136 | T-46 | 16.80 | 9.29 | 0.014 | 0.983 | 0.017 | 0.976 | 0.983 | 0.014 |
| 137 | T-47 | 11.96 | 8.66 | 0.014 | 0.957 | 0.014 | 0.948 | 0.957 | 0.014 |
| 138 | T-48 | 13.23 | 9.80 | 0.013 | 0.982 | 0.013 | 0.960 | 0.982 | 0.013 |
| 139 | T-49 | 15.40 | 10.57 | 0.016 | 0.975 | 0.020 | 0.954 | 0.975 | 0.016 |
| 140 | T-50 | 11.45 | 7.26 | 0.012 | 0.979 | 0.012 | 0.980 | 0.980 | 0.012 |
| 141 | T-51 | 13.36 | 8.40 | 0.013 | 0.977 | 0.013 | 0.978 | 0.978 | 0.013 |
| 142 | T-52 | 11.45 | 8.66 | 0.012 | 0.924 | 0.011 | 0.929 | 0.929 | 0.011 |
| 143 | T-53 | 12.60 | 8.15 | 0.011 | 0.981 | 0.012 | 0.975 | 0.981 | 0.011 |
| 144 | T-58 | 9.29 | 14.51 | 0.014 | 0.999 | 0.017 | 0.998 | 0.999 | 0.014 |
| 145 | T-60 | 17.56 | 9.29 | 0.015 | 0.932 | 0.015 | 0.933 | 0.933 | 0.015 |
| 146 | T-61 | 9.93 | 8.91 | 0.011 | 0.980 | 0.012 | 0.978 | 0.980 | 0.011 |
| 147 | T-62 | 12.85 | 8.53 | 0.014 | 0.972 | 0.015 | 0.968 | 0.972 | 0.014 |
| 148 | T-63 | 15.27 | 8.15 | 0.015 | 0.973 | 0.018 | 0.961 | 0.973 | 0.015 |
| 149 | T-64 | 15.65 | 8.91 | 0.012 | 0.997 | 0.026 | 0.984 | 0.997 | 0.012 |
| 150 | T-65 | 14.51 | 8.53 | 0.012 | 0.991 | 0.014 | 0.987 | 0.991 | 0.012 |
| 151 | T-67 | 16.54 | 10.44 | 0.015 | 0.991 | 0.015 | 0.992 | 0.992 | 0.015 |
| 152 | T-68 | 11.58 | 7.77 | 0.013 | 0.953 | 0.012 | 0.975 | 0.975 | 0.012 |
| 153 | T-69 | 13.11 | 8.91 | 0.013 | 0.989 | 0.013 | 0.988 | 0.989 | 0.013 |
| 154 | T-72 | 11.83 | 7.13 | 0.012 | 0.991 | 0.012 | 0.992 | 0.992 | 0.012 |
| 155 | T-73 | 15.40 | 10.31 | 0.011 | 0.958 | 0.011 | 0.960 | 0.960 | 0.011 |
| 156 | T-78 | 14.25 | 9.67 | 0.015 | 0.990 | 0.015 | 0.990 | 0.990 | 0.015 |
| 157 | T-79 | 11.96 | 8.02 | 0.014 | 0.999 | 0.015 | 0.998 | 0.999 | 0.014 |
| 158 | T-80 | 13.36 | 5.98 | 0.011 | 0.997 | 0.011 | 0.997 | 0.997 | 0.011 |
| 159 | T-81 | 14.25 | 7.89 | 0.013 | 0.977 | 0.013 | 0.974 | 0.977 | 0.013 |
| 160 | T-82 | 13.74 | 7.26 | 0.013 | 0.974 | 0.014 | 0.963 | 0.974 | 0.013 |
| 161 | T-83 | 13.87 | 9.17 | 0.013 | 0.985 | 0.013 | 0.982 | 0.985 | 0.013 |
| 162 | T-85 | 12.09 | 8.53 | 0.012 | 0.971 | 0.011 | 0.974 | 0.974 | 0.011 |
| 163 | T-86 | 14.00 | 8.40 | 0.012 | 0.993 | 0.011 | 0.993 | 0.993 | 0.012 |
| 164 | T-88 | 15.14 | 9.29 | 0.013 | 0.993 | 0.013 | 0.993 | 0.993 | 0.013 |
| 165 | T-89 | 12.72 | 8.15 | 0.012 | 0.965 | 0.013 | 0.960 | 0.965 | 0.012 |
| 166 | T-90 | 13.62 | 7.26 | 0.012 | 0.997 | 0.011 | 0.998 | 0.998 | 0.011 |


| No | Lesion ID | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{gathered} \overline{S_{\text {final }}} \\ (\mathrm{mm}) \end{gathered}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 167 | T-91 | 15.01 | 9.04 | 0.012 | 0.996 | 0.013 | 0.996 | 0.996 | 0.012 |
| 168 | T-92 | 15.14 | 8.27 | 0.013 | 0.995 | 0.014 | 0.995 | 0.995 | 0.013 |
| 169 | T-93 | 14.38 | 9.80 | 0.014 | 0.977 | 0.014 | 0.977 | 0.977 | 0.014 |
| 170 | T-94 | 14.38 | 8.53 | 0.013 | 0.955 | 0.013 | 0.953 | 0.955 | 0.013 |
| 171 | T-95 | 10.82 | 7.89 | 0.015 | 0.988 | 0.014 | 0.989 | 0.989 | 0.014 |
| 172 | T-96 | 12.22 | 7.64 | 0.012 | 0.947 | 0.013 | 0.954 | 0.954 | 0.013 |
| 173 | T-98 | 11.83 | 7.89 | 0.013 | 0.998 | 0.018 | 0.997 | 0.998 | 0.013 |
| 174 | T-99 | 9.42 | 7.26 | 0.014 | 0.984 | 0.013 | 0.986 | 0.986 | 0.013 |
| 175 | T-101 | 9.03 | 5.47 | 0.011 | 0.991 | 0.012 | 0.991 | 0.991 | 0.011 |
| 176 | T-102 | 11.45 | 8.27 | 0.014 | 0.996 | 0.014 | 0.996 | 0.996 | 0.014 |
| 177 | T-104 | 13.23 | 8.91 | 0.012 | 0.988 | 0.013 | 0.987 | 0.988 | 0.012 |
| 178 | T-105 | 11.32 | 7.38 | 0.015 | 0.975 | 0.016 | 0.970 | 0.975 | 0.015 |
| 179 | T-107 | 10.05 | 6.24 | 0.015 | 0.966 | 0.016 | 0.959 | 0.966 | 0.015 |
| 180 | T-108 | 14.38 | 7.13 | 0.014 | 0.970 | 0.014 | 0.967 | 0.970 | 0.014 |
| 181 | T-109 | 12.34 | 6.62 | 0.013 | 0.993 | 0.014 | 0.992 | 0.993 | 0.013 |
| 182 | T-110 | 16.80 | 7.38 | 0.015 | 0.996 | 0.016 | 0.996 | 0.996 | 0.015 |
| 183 | T-111 | 10.82 | 5.22 | 0.015 | 0.994 | 0.016 | 0.994 | 0.994 | 0.015 |
| 184 | T-114 | 13.23 | 7.38 | 0.014 | 0.962 | 0.015 | 0.955 | 0.962 | 0.014 |
| 185 | T-118 | 13.49 | 8.53 | 0.014 | 0.982 | 0.016 | 0.975 | 0.982 | 0.014 |
| 186 | T-124 | 9.03 | 4.58 | 0.014 | 0.999 | 0.016 | 0.999 | 0.999 | 0.014 |
| 187 | T-127 | 9.54 | 6.36 | 0.014 | 0.992 | 0.014 | 0.992 | 0.992 | 0.014 |
| 188 | T-128 | 6.74 | 8.02 | 0.014 | 0.998 | 0.014 | 0.998 | 0.998 | 0.014 |
| 189 | T-131 | 12.34 | 5.98 | 0.015 | 0.998 | 0.015 | 0.998 | 0.998 | 0.015 |
| 190 | T-132 | 15.14 | 8.40 | 0.015 | 0.969 | 0.014 | 0.984 | 0.984 | 0.014 |
| 191 | T-133 | 10.43 | 8.02 | 0.016 | 0.967 | 0.016 | 0.967 | 0.967 | 0.016 |
| 192 | T-134 | 11.07 | 7.13 | 0.014 | 0.995 | 0.014 | 0.995 | 0.995 | 0.014 |
| 193 | T-135 | 7.00 | 8.40 | 0.015 | 0.981 | 0.015 | 0.980 | 0.981 | 0.015 |
| 194 | T-136 | 13.36 | 6.87 | 0.013 | 0.993 | 0.013 | 0.993 | 0.993 | 0.013 |
| 195 | T-137 | 13.11 | 7.38 | 0.014 | 0.999 | 0.014 | 0.999 | 0.999 | 0.014 |
| 196 | T-138 | 13.36 | 8.53 | 0.015 | 0.998 | 0.015 | 0.998 | 0.998 | 0.015 |
| 197 | T-139 | 11.32 | 7.13 | 0.015 | 0.998 | 0.016 | 0.998 | 0.998 | 0.015 |
| 198 | T-143 | 10.05 | 6.75 | 0.013 | 0.995 | 0.013 | 0.995 | 0.995 | 0.013 |
| 199 | T-146 | 9.29 | 9.29 | 0.015 | 0.993 | 0.015 | 0.993 | 0.993 | 0.015 |
| 200 | T-147 | 12.85 | 6.49 | 0.013 | 0.993 | 0.014 | 0.991 | 0.993 | 0.013 |
| 201 | T-150 | 10.18 | 5.22 | 0.013 | 0.990 | 0.014 | 0.990 | 0.990 | 0.013 |
| 202 | T-151 | 9.80 | 5.73 | 0.015 | 0.976 | 0.015 | 0.974 | 0.976 | 0.015 |
| 203 | T-156 | 7.89 | 4.07 | 0.014 | 0.992 | 0.014 | 0.992 | 0.992 | 0.014 |
| 204 | T-158 | 8.78 | 3.95 | 0.014 | 0.974 | 0.014 | 0.973 | 0.974 | 0.014 |
| 205 | T-159 | 10.05 | 4.33 | 0.011 | 0.936 | 0.012 | 0.923 | 0.936 | 0.011 |
| 206 | T-160 | 9.67 | 4.96 | 0.013 | 0.964 | 0.013 | 0.962 | 0.964 | 0.013 |
| 207 | T-161 | 9.03 | 4.71 | 0.014 | 0.924 | 0.014 | 0.927 | 0.927 | 0.014 |
| 208 | T-163 | 10.18 | 4.97 | 0.012 | 0.976 | 0.014 | 0.970 | 0.976 | 0.012 |
| 209 | T-165 | 10.69 | 5.98 | 0.015 | 0.985 | 0.015 | 0.985 | 0.985 | 0.015 |


| No | Lesion ID | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{gathered} \overline{S_{\text {final }}} \\ (\mathrm{mm}) \end{gathered}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 210 | T-166 | 8.27 | 3.82 | 0.013 | 0.994 | 0.013 | 0.994 | 0.994 | 0.013 |
| 211 | T-170 | 8.53 | 5.09 | 0.011 | 0.994 | 0.011 | 0.994 | 0.994 | 0.011 |
| 212 | T-171 | 8.40 | 5.98 | 0.012 | 0.997 | 0.017 | 0.994 | 0.997 | 0.012 |
| 213 | T-172 | 8.53 | 4.07 | 0.012 | 0.995 | 0.012 | 0.995 | 0.995 | 0.012 |
| 214 | T-173 | 9.42 | 4.33 | 0.011 | 0.968 | 0.011 | 0.967 | 0.968 | 0.011 |
| 215 | T-174 | 7.64 | 12.48 | 0.015 | 0.993 | 0.015 | 0.993 | 0.993 | 0.015 |
| 216 | T-175 | 7.25 | 4.07 | 0.011 | 0.971 | 0.012 | 0.967 | 0.971 | 0.011 |
| 217 | T-176 | 8.65 | 3.95 | 0.012 | 0.939 | 0.013 | 0.932 | 0.939 | 0.012 |
| 218 | T-178 | 10.05 | 4.97 | 0.012 | 0.942 | 0.014 | 0.918 | 0.942 | 0.012 |
| 219 | T-182 | 8.65 | 4.71 | 0.013 | 0.961 | 0.013 | 0.957 | 0.961 | 0.013 |
| 220 | T-183 | 9.16 | 8.40 | 0.013 | 0.987 | 0.015 | 0.983 | 0.987 | 0.013 |
| 221 | T-185 | 8.14 | 3.82 | 0.014 | 0.984 | 0.014 | 0.983 | 0.984 | 0.014 |
| 222 | T-187 | 8.14 | 3.95 | 0.013 | 0.997 | 0.017 | 0.995 | 0.997 | 0.013 |
| 223 | T-188 | 7.51 | 4.07 | 0.013 | 0.993 | 0.016 | 0.990 | 0.993 | 0.013 |
| 224 | T-189 | 7.51 | 4.96 | 0.011 | 0.968 | 0.012 | 0.964 | 0.968 | 0.011 |
| 225 | T-190 | 11.45 | 6.62 | 0.013 | 0.990 | 0.013 | 0.990 | 0.990 | 0.013 |
| 226 | T-191 | 8.53 | 5.09 | 0.014 | 0.933 | 0.014 | 0.926 | 0.933 | 0.014 |
| 227 | T-193 | 10.69 | 3.69 | 0.013 | 0.953 | 0.013 | 0.951 | 0.953 | 0.013 |
| 228 | T-197 | 10.31 | 4.46 | 0.014 | 0.993 | 0.014 | 0.993 | 0.993 | 0.014 |
| 229 | T-199 | 15.40 | 6.49 | 0.013 | 0.999 | 0.014 | 0.998 | 0.999 | 0.013 |
| 230 | T-200 | 17.18 | 6.49 | 0.014 | 0.998 | 0.013 | 0.998 | 0.998 | 0.014 |
| 231 | T-201 | 13.49 | 7.13 | 0.013 | 0.995 | 0.015 | 0.994 | 0.995 | 0.013 |
| 232 | T-202 | 12.60 | 6.37 | 0.011 | 0.997 | 0.013 | 0.996 | 0.997 | 0.011 |
| 233 | T-203 | 13.23 | 6.24 | 0.011 | 0.998 | 0.013 | 0.997 | 0.998 | 0.011 |
| 234 | T-204 | 15.14 | 6.11 | 0.014 | 0.998 | 0.014 | 0.998 | 0.998 | 0.014 |
| 235 | T-205 | 16.16 | 7.00 | 0.014 | 0.991 | 0.014 | 0.990 | 0.991 | 0.014 |
| 236 | T-207 | 14.89 | 7.00 | 0.013 | 0.998 | 0.013 | 0.998 | 0.998 | 0.013 |
| 237 | T-208 | 12.72 | 6.24 | 0.014 | 0.996 | 0.013 | 0.997 | 0.997 | 0.013 |
| 238 | T-209 | 14.00 | 6.87 | 0.013 | 0.994 | 0.014 | 0.993 | 0.994 | 0.013 |
| 239 | T-210 | 13.74 | 6.62 | 0.012 | 0.950 | 0.012 | 0.951 | 0.951 | 0.012 |
| 240 | T-211 | 11.45 | 5.60 | 0.012 | 0.993 | 0.012 | 0.993 | 0.993 | 0.012 |
| 241 | T-212 | 13.36 | 7.13 | 0.015 | 0.989 | 0.014 | 0.990 | 0.990 | 0.014 |
| 242 | T-213 | 12.72 | 6.87 | 0.014 | 0.998 | 0.014 | 0.998 | 0.998 | 0.014 |
| 243 | T-214 | 11.07 | 10.31 | 0.014 | 0.997 | 0.013 | 0.997 | 0.997 | 0.014 |
| 244 | T-215 | 14.12 | 7.51 | 0.015 | 0.997 | 0.014 | 0.997 | 0.997 | 0.015 |
| 245 | T-216 | 12.85 | 6.62 | 0.013 | 0.995 | 0.013 | 0.995 | 0.995 | 0.013 |
| 246 | T-218 | 10.56 | 3.82 | 0.013 | 0.922 | 0.012 | 0.928 | 0.928 | 0.012 |
| 247 | T-219 | 9.80 | 4.20 | 0.013 | 0.990 | 0.013 | 0.990 | 0.990 | 0.013 |
| 248 | T-220 | 5.47 | 5.98 | 0.014 | 0.968 | 0.014 | 0.968 | 0.968 | 0.014 |
| 249 | T-221 | 10.69 | 5.86 | 0.014 | 0.987 | 0.015 | 0.985 | 0.987 | 0.014 |
| 250 | T-222 | 10.05 | 5.86 | 0.014 | 0.994 | 0.014 | 0.994 | 0.994 | 0.014 |
| 251 | T-223 | 5.85 | 6.11 | 0.014 | 0.992 | 0.014 | 0.992 | 0.992 | 0.014 |
| 252 | T-224 | 10.18 | 5.47 | 0.013 | 0.998 | 0.013 | 0.998 | 0.998 | 0.013 |


| No | Lesion ID | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 253 | L-2 | 9.29 | 12.48 | 0.012 | 0.982 | 0.014 | 0.984 | 0.984 | 0.014 |
| 254 | L-3 | 9.54 | 4.46 | 0.014 | 0.967 | 0.015 | 0.959 | 0.967 | 0.014 |
| 255 | L-4 | 9.42 | 9.42 | 0.016 | 0.970 | 0.018 | 0.965 | 0.970 | 0.016 |
| 256 | L-7 | 10.18 | 5.73 | 0.013 | 0.959 | 0.014 | 0.950 | 0.959 | 0.013 |
| 257 | L-8 | 9.67 | 14.51 | 0.011 | 0.970 | 0.011 | 0.975 | 0.975 | 0.011 |
| 258 | L-10 | 9.42 | 4.07 | 0.013 | 0.960 | 0.014 | 0.956 | 0.960 | 0.013 |
| 259 | L-11 | 9.42 | 4.46 | 0.014 | 0.949 | 0.015 | 0.942 | 0.949 | 0.014 |
| 260 | L-13 | 9.54 | 5.60 | 0.014 | 0.986 | 0.014 | 0.986 | 0.986 | 0.014 |
| 261 | L-14 | 9.67 | 5.35 | 0.015 | 0.918 | 0.015 | 0.913 | 0.918 | 0.015 |
| 262 | L-18 | 8.40 | 15.53 | 0.015 | 0.983 | 0.020 | 0.970 | 0.983 | 0.015 |
| 263 | L-19 | 10.05 | 5.73 | 0.012 | 0.952 | 0.013 | 0.954 | 0.954 | 0.013 |
| 264 | L-20 | 9.16 | 5.35 | 0.012 | 0.920 | 0.012 | 0.913 | 0.920 | 0.012 |
| 265 | L-21 | 9.54 | 4.96 | 0.014 | 0.954 | 0.016 | 0.946 | 0.954 | 0.014 |
| 266 | L-22 | 8.40 | 4.97 | 0.013 | 0.978 | 0.015 | 0.970 | 0.978 | 0.013 |
| 267 | L-23 | 9.92 | 4.33 | 0.012 | 0.963 | 0.014 | 0.966 | 0.966 | 0.014 |
| 268 | L-24 | 8.91 | 5.22 | 0.011 | 0.988 | 0.014 | 0.980 | 0.988 | 0.011 |
| 269 | L-25 | 8.91 | 4.96 | 0.012 | 0.987 | 0.013 | 0.983 | 0.987 | 0.012 |
| 270 | L-28 | 9.54 | 16.80 | 0.016 | 0.954 | 0.019 | 0.934 | 0.954 | 0.016 |
| 271 | L-29 | 9.93 | 16.68 | 0.016 | 0.971 | 0.016 | 0.971 | 0.971 | 0.016 |
| 272 | L-30 | 9.42 | 5.60 | 0.014 | 0.955 | 0.019 | 0.913 | 0.955 | 0.014 |
| 273 | L-31 | 9.42 | 14.13 | 0.015 | 0.965 | 0.015 | 0.973 | 0.973 | 0.015 |
| 274 | L-32 | 10.05 | 5.09 | 0.014 | 0.958 | 0.015 | 0.947 | 0.958 | 0.014 |
| 275 | L-36 | 9.03 | 4.46 | 0.015 | 0.953 | 0.016 | 0.943 | 0.953 | 0.015 |
| 276 | L-37 | 9.29 | 4.33 | 0.015 | 0.965 | 0.017 | 0.952 | 0.965 | 0.015 |
| 277 | L-38 | 9.67 | 4.33 | 0.011 | 0.994 | 0.015 | 0.987 | 0.994 | 0.011 |
| 278 | L-39 | 9.80 | 4.96 | 0.013 | 0.978 | 0.014 | 0.974 | 0.978 | 0.013 |
| 279 | L-41 | 10.05 | 5.22 | 0.015 | 0.944 | 0.017 | 0.932 | 0.944 | 0.015 |
| 280 | L-42 | 10.43 | 4.07 | 0.013 | 0.962 | 0.013 | 0.961 | 0.962 | 0.013 |
| 281 | L-43 | 9.29 | 5.09 | 0.013 | 0.943 | 0.013 | 0.937 | 0.943 | 0.013 |
| 282 | L-44 | 10.05 | 5.47 | 0.012 | 0.960 | 0.014 | 0.945 | 0.960 | 0.012 |
| 283 | L-45 | 10.18 | 4.84 | 0.013 | 0.941 | 0.015 | 0.938 | 0.941 | 0.013 |
| 284 | L-46 | 8.53 | 4.46 | 0.014 | 0.976 | 0.014 | 0.975 | 0.976 | 0.014 |
| 285 | L-51 | 9.80 | 5.22 | 0.011 | 0.974 | 0.012 | 0.970 | 0.974 | 0.011 |
| 286 | L-52 | 9.42 | 5.22 | 0.015 | 0.942 | 0.017 | 0.929 | 0.942 | 0.015 |
| 287 | L-53 | 9.16 | 5.22 | 0.012 | 0.944 | 0.012 | 0.950 | 0.950 | 0.012 |
| 288 | L-58 | 9.03 | 5.22 | 0.014 | 0.921 | 0.014 | 0.923 | 0.923 | 0.014 |
| 289 | L-59 | 8.40 | 5.35 | 0.015 | 0.934 | 0.016 | 0.928 | 0.934 | 0.015 |
| 290 | L-69 | 9.16 | 5.98 | 0.012 | 0.918 | 0.012 | 0.917 | 0.918 | 0.012 |
| 291 | L-70 | 10.18 | 6.11 | 0.015 | 0.935 | 0.015 | 0.930 | 0.935 | 0.015 |
| 292 | L-73 | 8.65 | 4.58 | 0.012 | 0.939 | 0.012 | 0.944 | 0.944 | 0.012 |
| 293 | L-74 | 8.91 | 4.71 | 0.012 | 0.989 | 0.012 | 0.990 | 0.990 | 0.012 |
| 294 | L-75 | 9.03 | 4.58 | 0.012 | 0.996 | 0.012 | 0.996 | 0.996 | 0.012 |
| 295 | L-76 | 9.03 | 5.47 | 0.013 | 0.918 | 0.014 | 0.903 | 0.918 | 0.013 |


| No | Lesion ID | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 296 | L-78 | 8.02 | 4.46 | 0.013 | 0.974 | 0.014 | 0.969 | 0.974 | 0.013 |
| 297 | L-80 | 8.65 | 5.86 | 0.015 | 0.930 | 0.016 | 0.919 | 0.930 | 0.015 |
| 298 | L-85 | 8.91 | 4.46 | 0.014 | 0.936 | 0.015 | 0.958 | 0.958 | 0.015 |
| 299 | L-88 | 9.29 | 4.58 | 0.013 | 0.983 | 0.015 | 0.977 | 0.983 | 0.013 |
| 300 | L-89 | 11.83 | 5.60 | 0.013 | 0.982 | 0.013 | 0.982 | 0.982 | 0.013 |
| 301 | L-90 | 11.58 | 5.73 | 0.014 | 0.995 | 0.014 | 0.995 | 0.995 | 0.014 |
| 302 | L-92 | 10.69 | 4.97 | 0.014 | 0.978 | 0.014 | 0.977 | 0.978 | 0.014 |
| 303 | L-93 | 9.93 | 4.33 | 0.013 | 0.988 | 0.013 | 0.987 | 0.988 | 0.013 |
| 304 | L-94 | 8.65 | 4.84 | 0.012 | 0.972 | 0.013 | 0.964 | 0.972 | 0.012 |
| 305 | L-95 | 10.05 | 4.97 | 0.013 | 0.966 | 0.013 | 0.966 | 0.966 | 0.013 |
| 306 | L-96 | 8.40 | 4.84 | 0.015 | 0.932 | 0.015 | 0.932 | 0.932 | 0.015 |
| 307 | L-97 | 8.53 | 5.22 | 0.012 | 0.947 | 0.012 | 0.961 | 0.961 | 0.012 |
| 308 | L-98 | 9.16 | 5.22 | 0.014 | 0.981 | 0.014 | 0.981 | 0.981 | 0.014 |
| 309 | L-99 | 9.42 | 4.46 | 0.012 | 0.963 | 0.012 | 0.963 | 0.963 | 0.012 |
| 310 | L-100 | 10.56 | 6.11 | 0.015 | 0.903 | 0.014 | 0.910 | 0.910 | 0.014 |
| 311 | L-101 | 8.40 | 5.47 | 0.016 | 0.945 | 0.016 | 0.946 | 0.946 | 0.016 |
| 312 | L-103 | 6.49 | 4.07 | 0.011 | 0.968 | 0.011 | 0.968 | 0.968 | 0.011 |
| 313 | L-104 | 8.65 | 5.98 | 0.013 | 0.962 | 0.013 | 0.962 | 0.962 | 0.013 |
| 314 | L-105 | 8.27 | 5.22 | 0.013 | 0.986 | 0.013 | 0.987 | 0.987 | 0.013 |
| 315 | L-106 | 9.03 | 12.86 | 0.013 | 0.958 | 0.013 | 0.959 | 0.959 | 0.013 |
| 316 | L-107 | 10.05 | 5.60 | 0.013 | 0.935 | 0.014 | 0.917 | 0.935 | 0.013 |
| 317 | L-108 | 9.16 | 6.11 | 0.011 | 0.944 | 0.011 | 0.946 | 0.946 | 0.011 |
| 318 | L-109 | 9.93 | 6.37 | 0.012 | 0.924 | 0.012 | 0.926 | 0.926 | 0.012 |
| 319 | L-111 | 9.16 | 5.47 | 0.012 | 0.929 | 0.013 | 0.924 | 0.929 | 0.012 |
| 320 | L-117 | 9.54 | 6.37 | 0.014 | 0.996 | 0.014 | 0.996 | 0.996 | 0.014 |
| 321 | L-118 | 9.29 | 6.75 | 0.014 | 0.983 | 0.015 | 0.982 | 0.983 | 0.014 |
| 322 | L-123 | 10.05 | 5.47 | 0.014 | 0.987 | 0.013 | 0.988 | 0.988 | 0.013 |
| 323 | L-125 | 8.91 | 6.24 | 0.015 | 0.989 | 0.015 | 0.988 | 0.989 | 0.015 |
| 324 | L-127 | 9.29 | 4.96 | 0.012 | 0.971 | 0.011 | 0.972 | 0.972 | 0.011 |
| 325 | L-128 | 9.80 | 5.86 | 0.014 | 0.998 | 0.015 | 0.998 | 0.998 | 0.014 |
| 326 | L-130 | 9.54 | 6.24 | 0.012 | 0.983 | 0.013 | 0.977 | 0.983 | 0.012 |
| 327 | L-131 | 9.67 | 5.86 | 0.013 | 0.934 | 0.015 | 0.924 | 0.934 | 0.013 |
| 328 | L-135 | 10.43 | 5.09 | 0.014 | 0.946 | 0.013 | 0.966 | 0.966 | 0.013 |
| 329 | L-137 | 6.62 | 2.80 | 0.012 | 0.990 | 0.012 | 0.990 | 0.990 | 0.012 |
| 330 | L-139 | 9.80 | 4.46 | 0.013 | 0.993 | 0.013 | 0.993 | 0.993 | 0.013 |
| 331 | L-140 | 9.80 | 4.84 | 0.012 | 0.990 | 0.012 | 0.989 | 0.990 | 0.012 |
| 332 | L-143 | 9.67 | 6.36 | 0.015 | 0.928 | 0.015 | 0.915 | 0.928 | 0.015 |
| 333 | L-144 | 10.18 | 16.04 | 0.016 | 0.996 | 0.023 | 0.991 | 0.996 | 0.016 |
| 334 | L-145 | 10.18 | 5.22 | 0.014 | 0.948 | 0.015 | 0.941 | 0.948 | 0.014 |
| 335 | L-147 | 10.56 | 5.09 | 0.013 | 0.953 | 0.013 | 0.963 | 0.963 | 0.013 |
| 336 | L-151 | 9.80 | 5.60 | 0.013 | 0.965 | 0.016 | 0.946 | 0.965 | 0.013 |
| 337 | L-152 | 8.65 | 6.62 | 0.014 | 0.982 | 0.016 | 0.978 | 0.982 | 0.014 |
| 338 | L-153 | 9.16 | 6.24 | 0.012 | 0.975 | 0.015 | 0.963 | 0.975 | 0.012 |


| No | Lesion ID | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 339 | L-156 | 9.42 | 5.98 | 0.014 | 0.942 | 0.016 | 0.929 | 0.942 | 0.014 |
| 340 | L-158 | 8.91 | 4.33 | 0.013 | 0.988 | 0.017 | 0.982 | 0.988 | 0.013 |
| 341 | L-160 | 10.82 | 5.60 | 0.015 | 0.996 | 0.015 | 0.996 | 0.996 | 0.015 |
| 342 | L-162 | 10.43 | 4.96 | 0.013 | 0.992 | 0.015 | 0.989 | 0.992 | 0.013 |
| 343 | L-164 | 10.05 | 5.22 | 0.012 | 0.971 | 0.015 | 0.958 | 0.971 | 0.012 |
| 344 | L-165 | 9.54 | 4.71 | 0.015 | 0.954 | 0.016 | 0.948 | 0.954 | 0.015 |
| 345 | L-166 | 9.67 | 5.86 | 0.012 | 0.982 | 0.014 | 0.975 | 0.982 | 0.012 |
| 346 | L-172 | 8.91 | 5.73 | 0.011 | 0.986 | 0.011 | 0.986 | 0.986 | 0.011 |
| 347 | L-173 | 10.56 | 5.86 | 0.013 | 0.957 | 0.013 | 0.957 | 0.957 | 0.013 |
| 348 | L-174 | 11.71 | 17.31 | 0.015 | 0.987 | 0.015 | 0.988 | 0.988 | 0.015 |
| 349 | L-175 | 16.29 | 6.87 | 0.012 | 0.985 | 0.019 | 0.963 | 0.985 | 0.012 |
| 350 | L-176 | 11.07 | 5.35 | 0.014 | 0.948 | 0.014 | 0.951 | 0.951 | 0.014 |
| 351 | L-180 | 9.54 | 5.60 | 0.015 | 0.994 | 0.015 | 0.994 | 0.994 | 0.015 |
| 352 | L-181 | 9.03 | 5.47 | 0.014 | 0.992 | 0.014 | 0.991 | 0.992 | 0.014 |
| 353 | L-183 | 9.29 | 5.35 | 0.013 | 0.974 | 0.015 | 0.963 | 0.974 | 0.013 |
| 354 | L-184 | 9.93 | 5.98 | 0.014 | 0.974 | 0.016 | 0.964 | 0.974 | 0.014 |
| 355 | L-186 | 9.54 | 5.47 | 0.013 | 0.981 | 0.014 | 0.977 | 0.981 | 0.013 |
| 356 | L-187 | 9.80 | 16.80 | 0.014 | 0.987 | 0.018 | 0.970 | 0.987 | 0.014 |
| 357 | L-188 | 10.05 | 5.73 | 0.012 | 0.977 | 0.011 | 0.978 | 0.978 | 0.011 |
| 358 | L-190 | 9.42 | 5.73 | 0.011 | 0.984 | 0.012 | 0.979 | 0.984 | 0.011 |
| 359 | L-191 | 9.42 | 4.71 | 0.012 | 0.987 | 0.013 | 0.984 | 0.987 | 0.012 |
| 360 | L-192 | 8.02 | 6.62 | 0.013 | 0.937 | 0.014 | 0.945 | 0.945 | 0.014 |
| 361 | L-193 | 10.05 | 5.98 | 0.013 | 0.969 | 0.014 | 0.964 | 0.969 | 0.013 |
| 362 | L-194 | 10.18 | 16.55 | 0.012 | 0.987 | 0.016 | 0.980 | 0.987 | 0.012 |
| 363 | L-195 | 9.67 | 5.09 | 0.014 | 0.926 | 0.014 | 0.926 | 0.926 | 0.014 |
| 364 | L-196 | 9.54 | 4.33 | 0.015 | 0.927 | 0.014 | 0.930 | 0.930 | 0.014 |
| 365 | L-197 | 9.93 | 4.97 | 0.012 | 0.979 | 0.014 | 0.971 | 0.979 | 0.012 |
| 366 | L-198 | 8.78 | 4.96 | 0.011 | 0.959 | 0.011 | 0.961 | 0.961 | 0.011 |
| 367 | L-199 | 9.93 | 4.96 | 0.011 | 0.992 | 0.015 | 0.981 | 0.992 | 0.011 |
| 368 | L-201 | 7.89 | 3.69 | 0.015 | 0.906 | 0.015 | 0.905 | 0.906 | 0.015 |
| 369 | L-202 | 9.42 | 14.38 | 0.015 | 0.969 | 0.015 | 0.965 | 0.969 | 0.015 |
| 370 | L-203 | 8.14 | 4.96 | 0.012 | 0.918 | 0.012 | 0.916 | 0.918 | 0.012 |
| 371 | L-204 | 8.53 | 5.73 | 0.013 | 0.952 | 0.013 | 0.952 | 0.952 | 0.013 |
| 372 | L-205 | 9.93 | 5.98 | 0.015 | 0.961 | 0.015 | 0.960 | 0.961 | 0.015 |
| 373 | L-211 | 7.76 | 3.44 | 0.013 | 0.998 | 0.014 | 0.998 | 0.998 | 0.013 |
| 374 | L-213 | 8.91 | 5.09 | 0.015 | 0.974 | 0.016 | 0.969 | 0.974 | 0.015 |
| 375 | L-215 | 9.16 | 6.24 | 0.011 | 0.903 | 0.011 | 0.903 | 0.903 | 0.011 |
| 376 | L-217 | 9.42 | 14.51 | 0.011 | 0.940 | 0.012 | 0.944 | 0.944 | 0.012 |
| 377 | L-218 | 10.56 | 5.09 | 0.013 | 0.959 | 0.013 | 0.959 | 0.959 | 0.013 |
| 378 | L-219 | 8.78 | 4.20 | 0.014 | 0.941 | 0.015 | 0.936 | 0.941 | 0.014 |
| 379 | L-221 | 10.82 | 4.71 | 0.012 | 0.942 | 0.013 | 0.931 | 0.942 | 0.012 |
| 380 | L-222 | 10.56 | 4.20 | 0.011 | 0.956 | 0.011 | 0.957 | 0.957 | 0.011 |
| 381 | L-223 | 8.91 | 4.71 | 0.013 | 0.965 | 0.013 | 0.963 | 0.965 | 0.013 |


| No | Lesion <br> ID | Width <br> $(\mathrm{mm})$ | Height <br> $(\mathrm{mm})$ | $S_{a, 2}$ <br> $(\mathrm{~mm})$ | $R_{2}^{2}$ | $S_{a, 3}$ <br> $(\mathrm{~mm})$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $S_{\text {final }}$ <br> $(\mathrm{mm})$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 382 | L-224 | 10.56 | 4.58 | 0.015 | 0.979 | 0.015 | 0.979 | 0.979 | 0.015 |
| 383 | L-226 | 9.29 | 3.82 | 0.013 | 0.998 | 0.017 | 0.996 | 0.998 | 0.013 |
| 384 | L-227 | 9.93 | 4.07 | 0.013 | 0.987 | 0.012 | 0.989 | 0.989 | 0.012 |
| 385 | L-228 | 8.02 | 3.95 | 0.012 | 0.998 | 0.012 | 0.998 | 0.998 | 0.012 |
| 386 | L-229 | 8.53 | 4.58 | 0.014 | 0.996 | 0.014 | 0.996 | 0.996 | 0.014 |
| 387 | L-231 | 9.54 | 5.22 | 0.012 | 0.991 | 0.012 | 0.991 | 0.991 | 0.012 |
| 388 | L-233 | 10.94 | 7.64 | 0.014 | 0.998 | 0.014 | 0.998 | 0.998 | 0.014 |
| 389 | L-236 | 11.32 | 3.69 | 0.014 | 0.992 | 0.015 | 0.989 | 0.992 | 0.014 |
| 390 | L-237 | 14.51 | 6.87 | 0.014 | 0.996 | 0.014 | 0.996 | 0.996 | 0.014 |

APPENDIX D: TRAINING DATASET OF PSORIASIS LESION

| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | P-4-MOH | 36 | M | H1 | 12.86 | 17.69 | 0.031 | 0.967 | 0.033 | 0.967 | 0.967 | 0.033 |
| 2 | P-4-MOH | 36 | M | T1 | 16.43 | 19.22 | 0.036 | 0.939 | 0.038 | 0.930 | 0.939 | 0.036 |
| 3 | P-4-MOH | 36 | M | T2 | 9.17 | 10.44 | 0.041 | 0.960 | 0.041 | 0.961 | 0.961 | 0.041 |
| 4 | P-4-MOH | 36 | M | U2 | 6.50 | 10.05 | 0.018 | 0.997 | 0.016 | 0.998 | 0.998 | 0.016 |
| 5 | P-5-IBR | 69 | M | U3 | 6.75 | 7.13 | 0.026 | 0.926 | 0.028 | 0.918 | 0.926 | 0.026 |
| 6 | P-6-AZI | 32 | M | H1 | 9.30 | 7.38 | 0.049 | 0.978 | 0.049 | 0.978 | 0.978 | 0.049 |
| 7 | P-6-AZI | 32 | M | L1 | 12.86 | 12.09 | 0.052 | 0.964 | 0.052 | 0.963 | 0.964 | 0.052 |
| 8 | P-6-AZI | 32 | M | T1 | 13.88 | 7.51 | 0.058 | 0.950 | 0.052 | 0.941 | 0.950 | 0.058 |
| 9 | P-6-AZI | 32 | M | U1 | 11.72 | 7.76 | 0.035 | 0.977 | 0.037 | 0.975 | 0.977 | 0.035 |
| 10 | P-6-AZI | 32 | M | U2 | 13.50 | 12.60 | 0.054 | 0.779 | 0.043 | 0.915 | 0.915 | 0.043 |
| 11 | P-6-AZI | 32 | M | U3 | 8.79 | 7.51 | 0.046 | 0.954 | 0.045 | 0.954 | 0.954 | 0.045 |
| 12 | P-7-SIV | 40 | M | L2 | 10.44 | 10.31 | 0.030 | 0.904 | 0.027 | 0.920 | 0.920 | 0.027 |
| 13 | P-7-SIV | 40 | M | L3 | 6.37 | 6.75 | 0.036 | 0.916 | 0.037 | 0.914 | 0.916 | 0.036 |
| 14 | P-7-SIV | 40 | M | T1 | 6.62 | 6.87 | 0.029 | 0.969 | 0.029 | 0.968 | 0.969 | 0.029 |
| 15 | P-7-SIV | 40 | M | T2 | 7.39 | 7.89 | 0.044 | 0.901 | 0.046 | 0.857 | 0.901 | 0.044 |
| 16 | P-7-SIV | 40 | M | T3 | 6.88 | 7.13 | 0.024 | 0.964 | 0.020 | 0.976 | 0.976 | 0.020 |
| 17 | P-8-SUN | 62 | M | H1 | 4.97 | 5.85 | 0.020 | 0.924 | 0.020 | 0.924 | 0.924 | 0.020 |
| 18 | P-8-SUN | 62 | M | L2 | 7.52 | 7.25 | 0.025 | 0.964 | 0.027 | 0.954 | 0.964 | 0.025 |
| 19 | P-8-SUN | 62 | M | L3 | 6.88 | 4.96 | 0.025 | 0.928 | 0.025 | 0.928 | 0.928 | 0.025 |
| 20 | P-8-SUN | 62 | M | T1 | 14.14 | 13.11 | 0.035 | 0.984 | 0.036 | 0.984 | 0.984 | 0.036 |
| 21 | P-8-SUN | 62 | M | T3 | 5.73 | 4.71 | 0.020 | 0.949 | 0.020 | 0.949 | 0.949 | 0.020 |
| 22 | P-8-SUN | 62 | M | U1 | 2.93 | 3.44 | 0.020 | 0.908 | 0.022 | 0.911 | 0.911 | 0.022 |
| 23 | P-8-SUN | 62 | M | U3 | 3.95 | 3.69 | 0.019 | 0.932 | 0.019 | 0.932 | 0.932 | 0.019 |
| 24 | P-9-MOH | 60 | M | L12 | 12.74 | 14.25 | 0.065 | 0.955 | 0.061 | 0.961 | 0.961 | 0.061 |
| 25 | P-9-MOH | 60 | M | L2 | 16.05 | 17.05 | 0.041 | 0.985 | 0.038 | 0.987 | 0.987 | 0.038 |
| 26 | P-9-MOH | 60 | M | T1 | 16.56 | 15.53 | 0.045 | 0.976 | 0.045 | 0.975 | 0.976 | 0.045 |
| 27 | P-9-MOH | 60 | M | T2 | 12.23 | 12.34 | 0.040 | 0.967 | 0.042 | 0.951 | 0.967 | 0.040 |
| 28 | P-9-MOH | 60 | M | T3 | 16.17 | 14.38 | 0.096 | 0.945 | 0.092 | 0.950 | 0.950 | 0.092 |
| 29 | P-9-MOH | 60 | M | U1 | 7.26 | 13.87 | 0.032 | 0.969 | 0.029 | 0.955 | 0.969 | 0.032 |
| 30 | P-9-MOH | 60 | M | U2 | 7.90 | 13.75 | 0.038 | 0.980 | 0.039 | 0.978 | 0.980 | 0.038 |
| 31 | P-9-MOH | 60 | M | U3 | 11.85 | 11.45 | 0.042 | 0.983 | 0.040 | 0.985 | 0.985 | 0.040 |
| 32 | P-10-MUH | 28 | M | H1 | 9.55 | 9.29 | 0.038 | 0.950 | 0.037 | 0.948 | 0.950 | 0.038 |
| 33 | P-10-MUH | 28 | M | L12 | 6.24 | 11.96 | 0.072 | 0.969 | 0.069 | 0.972 | 0.972 | 0.069 |
| 34 | P-10-MUH | 28 | M | L3 | 9.55 | 12.85 | 0.051 | 0.976 | 0.051 | 0.975 | 0.976 | 0.051 |
| 35 | P-10-MUH | 28 | M | T2 | 5.48 | 4.20 | 0.037 | 0.908 | 0.037 | 0.907 | 0.908 | 0.037 |
| 36 | P-10-MUH | 28 | M | T4 | 16.05 | 15.40 | 0.077 | 0.967 | 0.070 | 0.972 | 0.972 | 0.070 |
| 37 | P-10-MUH | 28 | M | U2 | 4.71 | 8.53 | 0.026 | 0.920 | 0.025 | 0.928 | 0.928 | 0.025 |
| 38 | P-11-THA | 43 | F | H | 10.32 | 13.24 | 0.023 | 0.977 | 0.024 | 0.975 | 0.977 | 0.023 |
| 39 | P-11-THA | 43 | F | L1 | 4.20 | 5.60 | 0.016 | 0.933 | 0.016 | 0.936 | 0.936 | 0.016 |
| 40 | P-11-THA | 43 | F | L3 | 4.84 | 5.60 | 0.024 | 0.959 | 0.024 | 0.959 | 0.959 | 0.024 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 41 | P-11-THA | 43 | F | T1 | 11.08 | 8.66 | 0.058 | 0.912 | 0.052 | 0.907 | 0.912 | 0.058 |
| 42 | P-11-THA | 43 | F | T2 | 7.51 | 5.09 | 0.026 | 0.965 | 0.025 | 0.966 | 0.966 | 0.025 |
| 43 | P-11-THA | 43 | F | T3 | 6.62 | 10.05 | 0.026 | 0.950 | 0.024 | 0.956 | 0.956 | 0.024 |
| 44 | P-11-THA | 43 | F | U1 | 8.28 | 9.93 | 0.030 | 0.958 | 0.029 | 0.960 | 0.960 | 0.029 |
| 45 | P-11-THA | 43 | F | U2 | 5.09 | 5.47 | 0.028 | 0.974 | 0.026 | 0.977 | 0.977 | 0.026 |
| 46 | P-11-THA | 43 | F | U3 | 7.90 | 11.58 | 0.029 | 0.986 | 0.030 | 0.986 | 0.986 | 0.030 |
| 47 | P-12-YEE | 57 | M | L2 | 10.95 | 7.26 | 0.043 | 0.952 | 0.042 | 0.953 | 0.953 | 0.042 |
| 48 | P-12-YEE | 57 | M | L3 | 7.39 | 11.96 | 0.035 | 0.956 | 0.033 | 0.962 | 0.962 | 0.033 |
| 49 | P-12-YEE | 57 | M | T2 | 9.17 | 6.62 | 0.026 | 0.978 | 0.027 | 0.976 | 0.978 | 0.026 |
| 50 | P-12-YEE | 57 | M | T3 | 7.64 | 5.60 | 0.035 | 0.909 | 0.046 | 0.786 | 0.909 | 0.035 |
| 51 | P-12-YEE | 57 | M | U3 | 5.09 | 5.09 | 0.016 | 0.981 | 0.015 | 0.982 | 0.982 | 0.015 |
| 52 | P-13-ABD | 60 | M | T1 | 6.24 | 8.27 | 0.039 | 0.938 | 0.039 | 0.938 | 0.938 | 0.039 |
| 53 | P-13-ABD | 60 | M | T2 | 5.60 | 6.11 | 0.015 | 0.943 | 0.015 | 0.943 | 0.943 | 0.015 |
| 54 | P-13-ABD | 60 | M | T4 | 21.14 | 26.98 | 0.059 | 0.970 | 0.065 | 0.962 | 0.970 | 0.059 |
| 55 | P-13-ABD | 60 | M | U3 | 8.15 | 5.34 | 0.041 | 0.850 | 0.034 | 0.921 | 0.921 | 0.034 |
| 56 | P-14-FAT | 60 | F | H2 | 5.73 | 5.98 | 0.028 | 0.933 | 0.028 | 0.934 | 0.934 | 0.028 |
| 57 | P-14-FAT | 60 | F | L2 | 7.00 | 4.33 | 0.027 | 0.959 | 0.026 | 0.961 | 0.961 | 0.026 |
| 58 | P-14-FAT | 60 | F | T2 | 5.35 | 5.09 | 0.028 | 0.930 | 0.027 | 0.933 | 0.933 | 0.027 |
| 59 | P-14-FAT | 60 | F | U1 | 3.82 | 4.46 | 0.019 | 0.947 | 0.018 | 0.952 | 0.952 | 0.018 |
| 60 | P-14-FAT | 60 | F | U2 | 4.97 | 4.45 | 0.021 | 0.964 | 0.021 | 0.964 | 0.964 | 0.021 |
| 61 | P-14-FAT | 60 | F | U3 | 4.97 | 4.20 | 0.020 | 0.960 | 0.018 | 0.966 | 0.966 | 0.018 |
| 62 | P-15-AIZ | 37 | M | L1 | 6.37 | 10.05 | 0.029 | 0.926 | 0.029 | 0.923 | 0.926 | 0.029 |
| 63 | P-15-AIZ | 37 | M | T1 | 5.48 | 12.60 | 0.032 | 0.983 | 0.031 | 0.984 | 0.984 | 0.031 |
| 64 | P-15-AIZ | 37 | M | T2 | 11.21 | 7.64 | 0.035 | 0.944 | 0.035 | 0.943 | 0.944 | 0.035 |
| 65 | P-15-AIZ | 37 | M | U2 | 4.59 | 5.22 | 0.019 | 0.912 | 0.018 | 0.915 | 0.915 | 0.018 |
| 66 | P-15-AIZ | 37 | M | U3 | 5.09 | 5.85 | 0.030 | 0.906 | 0.030 | 0.903 | 0.906 | 0.030 |
| 67 | P-16-AZH | 47 | M | U2 | 5.22 | 6.87 | 0.048 | 0.906 | 0.049 | 0.904 | 0.906 | 0.048 |
| 68 | P-16-AZH | 47 | M | T1 | 4.46 | 6.24 | 0.017 | 0.945 | 0.017 | 0.945 | 0.945 | 0.017 |
| 69 | P-16-AZH | 47 | M | L1 | 5.86 | 8.40 | 0.032 | 0.926 | 0.034 | 0.933 | 0.933 | 0.034 |
| 70 | P-16-AZH | 47 | M | L2 | 4.46 | 5.85 | 0.019 | 0.973 | 0.020 | 0.973 | 0.973 | 0.020 |
| 71 | P-17-MUH | 22 | M | U1 | 11.84 | 12.86 | 0.028 | 0.981 | 0.028 | 0.981 | 0.981 | 0.028 |
| 72 | P-17-MUH | 22 | M | U2 | 12.48 | 14.25 | 0.022 | 0.958 | 0.021 | 0.960 | 0.960 | 0.021 |
| 73 | P-17-MUH | 22 | M | U3 | 10.44 | 10.05 | 0.036 | 0.969 | 0.037 | 0.968 | 0.969 | 0.036 |
| 74 | P-17-MUH | 22 | M | T3 | 8.41 | 8.53 | 0.031 | 0.925 | 0.030 | 0.930 | 0.930 | 0.030 |
| 75 | P-17-MUH | 22 | M | L1 | 10.19 | 16.80 | 0.034 | 0.987 | 0.034 | 0.984 | 0.987 | 0.034 |
| 76 | P-17-MUH | 22 | M | L2 | 15.03 | 13.75 | 0.080 | 0.960 | 0.078 | 0.961 | 0.961 | 0.078 |
| 77 | P-17-MUH | 22 | M | L3 | 11.84 | 16.55 | 0.035 | 0.988 | 0.035 | 0.988 | 0.988 | 0.035 |
| 78 | P-17-MUH | 22 | M | L4 | 29.55 | 22.91 | 0.040 | 0.977 | 0.042 | 0.991 | 0.991 | 0.042 |
| 79 | P-17-MUH | 22 | M | L5 | 18.98 | 21.38 | 0.038 | 0.987 | 0.038 | 0.987 | 0.987 | 0.038 |
| 80 | P-18-SID | 41 | M | U2 | 5.22 | 5.85 | 0.028 | 0.951 | 0.028 | 0.952 | 0.952 | 0.028 |
| 81 | P-18-SID | 41 | M | U3 | 6.62 | 6.87 | 0.031 | 0.921 | 0.031 | 0.918 | 0.921 | 0.031 |
| 82 | P-18-SID | 41 | M | T3 | 5.73 | 7.89 | 0.034 | 0.934 | 0.033 | 0.936 | 0.936 | 0.033 |
| 83 | P-18-SID | 41 | M | L1 | 11.59 | 10.06 | 0.025 | 0.967 | 0.025 | 0.968 | 0.968 | 0.025 |
| 84 | P-18-SID | 41 | M | L2 | 8.66 | 11.20 | 0.029 | 0.969 | 0.028 | 0.970 | 0.970 | 0.028 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{\overline{S_{\text {final }}}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 85 | P-19-NUR | 19 | F | H1 | 6.88 | 12.35 | 0.027 | 0.953 | 0.029 | 0.947 | 0.953 | 0.027 |
| 86 | P-19-NUR | 19 | F | U1 | 7.77 | 10.82 | 0.039 | 0.946 | 0.043 | 0.937 | 0.946 | 0.039 |
| 87 | P-19-NUR | 19 | F | U2 | 8.79 | 11.96 | 0.040 | 0.949 | 0.039 | 0.951 | 0.951 | 0.039 |
| 88 | P-19-NUR | 19 | F | U3 | 11.85 | 15.40 | 0.057 | 0.951 | 0.059 | 0.947 | 0.951 | 0.057 |
| 89 | P-19-NUR | 19 | F | T1 | 11.84 | 10.82 | 0.041 | 0.980 | 0.041 | 0.980 | 0.980 | 0.041 |
| 90 | P-19-NUR | 19 | F | T2 | 11.08 | 9.42 | 0.040 | 0.981 | 0.040 | 0.981 | 0.981 | 0.040 |
| 91 | P-19-NUR | 19 | F | T3 | 8.79 | 8.02 | 0.042 | 0.950 | 0.041 | 0.954 | 0.954 | 0.041 |
| 92 | P-19-NUR | 19 | F | L1 | 9.30 | 13.11 | 0.036 | 0.993 | 0.036 | 0.993 | 0.993 | 0.036 |
| 93 | P-19-NUR | 19 | F | L2 | 8.79 | 11.20 | 0.035 | 0.956 | 0.034 | 0.958 | 0.958 | 0.034 |
| 94 | P-19-NUR | 19 | F | L3 | 6.88 | 8.65 | 0.022 | 0.970 | 0.022 | 0.973 | 0.973 | 0.022 |
| 95 | P-20-MOH | 34 | M | U3 | 34.64 | 13.75 | 0.079 | 0.961 | 0.078 | 0.962 | 0.962 | 0.078 |
| 96 | $\mathrm{P}-20-\mathrm{MOH}$ | 34 | M | T1 | 20.89 | 16.80 | 0.078 | 0.948 | 0.076 | 0.950 | 0.950 | 0.076 |
| 97 | P-20-MOH | 34 | M | T3 | 13.12 | 10.82 | 0.051 | 0.967 | 0.048 | 0.970 | 0.970 | 0.048 |
| 98 | P-20-MOH | 34 | M | T5 | 23.94 | 21.00 | 0.115 | 0.967 | 0.120 | 0.964 | 0.967 | 0.115 |
| 99 | P-20-MOH | 34 | M | T6 | 37.95 | 28.38 | 0.136 | 0.964 | 0.148 | 0.954 | 0.964 | 0.136 |
| 100 | $\mathrm{P}-20-\mathrm{MOH}$ | 34 | M | T7 | 9.93 | 7.25 | 0.076 | 0.958 | 0.063 | 0.953 | 0.958 | 0.076 |
| 101 | P-20-MOH | 34 | M | L1 | 19.10 | 16.04 | 0.131 | 0.931 | 0.131 | 0.933 | 0.933 | 0.131 |
| 102 | P-20-MOH | 34 | M | L2 | 37.19 | 13.24 | 0.125 | 0.941 | 0.116 | 0.923 | 0.941 | 0.125 |
| 103 | $\mathrm{P}-20-\mathrm{MOH}$ | 34 | M | L3 | 7.00 | 8.78 | 0.030 | 0.980 | 0.027 | 0.966 | 0.980 | 0.030 |
| 104 | P-21-ZAI | 62 | M | U1 | 13.25 | 11.58 | 0.049 | 0.941 | 0.044 | 0.956 | 0.956 | 0.044 |
| 105 | P-21-ZAI | 62 | M | U3 | 6.11 | 6.74 | 0.029 | 0.948 | 0.030 | 0.947 | 0.948 | 0.029 |
| 106 | P-21-ZAI | 62 | M | T1 | 25.47 | 21.89 | 0.036 | 0.996 | 0.032 | 0.997 | 0.997 | 0.032 |
| 107 | P-21-ZAI | 62 | M | T2 | 13.25 | 12.85 | 0.038 | 0.960 | 0.035 | 0.963 | 0.963 | 0.035 |
| 108 | P-21-ZAI | 62 | M | T3 | 25.35 | 7.51 | 0.067 | 0.962 | 0.070 | 0.960 | 0.962 | 0.067 |
| 109 | P-21-ZAI | 62 | M | L1 | 6.62 | 11.20 | 0.028 | 0.985 | 0.028 | 0.985 | 0.985 | 0.028 |
| 110 | P-21-ZAI | 62 | M | L2 | 20.38 | 17.82 | 0.041 | 0.990 | 0.042 | 0.990 | 0.990 | 0.042 |
| 111 | P-21-ZAI | 62 | M | L3 | 15.67 | 19.22 | 0.029 | 0.963 | 0.027 | 0.968 | 0.968 | 0.027 |
| 112 | P-22-MOH | 38 | M | U1 | 20.76 | 17.44 | 0.050 | 0.968 | 0.051 | 0.968 | 0.968 | 0.051 |
| 113 | $\mathrm{P}-22-\mathrm{MOH}$ | 38 | M | L4 | 34.26 | 27.74 | 0.045 | 0.975 | 0.050 | 0.972 | 0.975 | 0.045 |
| 114 | P-22-MOH | 38 | M | L5 | 30.44 | 27.49 | 0.032 | 0.976 | 0.046 | 0.987 | 0.987 | 0.046 |
| 115 | P-22-MOH | 38 | M | L6 | 31.08 | 26.35 | 0.032 | 0.982 | 0.042 | 0.990 | 0.990 | 0.042 |
| 116 | P-24-KAM | 60 | M | U1 | 8.28 | 12.22 | 0.036 | 0.965 | 0.036 | 0.965 | 0.965 | 0.036 |
| 117 | P-24-KAM | 60 | M | T1 | 9.43 | 10.56 | 0.026 | 0.965 | 0.026 | 0.963 | 0.965 | 0.026 |
| 118 | P-24-KAM | 60 | M | T2 | 9.17 | 12.98 | 0.026 | 0.978 | 0.025 | 0.979 | 0.979 | 0.025 |
| 119 | P-24-KAM | 60 | M | T3 | 17.83 | 13.75 | 0.035 | 0.971 | 0.034 | 0.973 | 0.973 | 0.034 |
| 120 | P-25-RAB | 74 | F | U1 | 8.66 | 5.98 | 0.019 | 0.985 | 0.019 | 0.985 | 0.985 | 0.019 |
| 121 | P-25-RAB | 74 | F | T4 | 35.79 | 28.25 | 0.051 | 0.913 | 0.056 | 0.736 | 0.913 | 0.051 |
| 122 | P-25-RAB | 74 | F | T2 | 18.09 | 11.58 | 0.026 | 0.991 | 0.025 | 0.992 | 0.992 | 0.025 |
| 123 | P-25-RAB | 74 | F | L2 | 10.06 | 5.73 | 0.031 | 0.924 | 0.030 | 0.928 | 0.928 | 0.030 |
| 124 | P-25-RAB | 74 | F | L3 | 12.35 | 12.34 | 0.071 | 0.917 | 0.071 | 0.917 | 0.917 | 0.071 |
| 125 | P-26-SAM | 44 | M | U1 | 4.33 | 3.95 | 0.033 | 0.953 | 0.032 | 0.954 | 0.954 | 0.032 |
| 126 | P-26-SAM | 44 | M | U22 | 2.29 | 2.93 | 0.018 | 0.923 | 0.018 | 0.923 | 0.923 | 0.018 |
| 127 | P-26-SAM | 44 | M | U3 | 4.33 | 4.45 | 0.015 | 0.925 | 0.015 | 0.928 | 0.928 | 0.015 |
| 128 | P-26-SAM | 44 | M | T2 | 5.22 | 5.73 | 0.025 | 0.917 | 0.027 | 0.931 | 0.931 | 0.027 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 129 | P-26-SAM | 44 | M | L2 | 23.05 | 24.18 | 0.051 | 0.976 | 0.055 | 0.985 | 0.985 | 0.055 |
| 130 | P-26-SAM | 44 | M | L3 | 7.26 | 8.15 | 0.023 | 0.941 | 0.023 | 0.942 | 0.942 | 0.023 |
| 131 | P-27-ROH | 27 | F | H1 | 6.88 | 9.29 | 0.025 | 0.984 | 0.024 | 0.985 | 0.985 | 0.024 |
| 132 | P-27-ROH | 27 | F | U1 | 4.84 | 5.73 | 0.033 | 0.964 | 0.034 | 0.963 | 0.964 | 0.033 |
| 133 | P-27-ROH | 27 | F | U2 | 3.06 | 3.56 | 0.024 | 0.920 | 0.023 | 0.926 | 0.926 | 0.023 |
| 134 | P-27-ROH | 27 | F | U3 | 4.58 | 5.47 | 0.026 | 0.946 | 0.026 | 0.946 | 0.946 | 0.026 |
| 135 | P-27-ROH | 27 | F | T1 | 3.44 | 3.95 | 0.026 | 0.952 | 0.026 | 0.933 | 0.952 | 0.026 |
| 136 | P-27-ROH | 27 | F | L1 | 8.02 | 8.65 | 0.025 | 0.960 | 0.023 | 0.965 | 0.965 | 0.023 |
| 137 | P-27-ROH | 27 | F | L2 | 3.44 | 3.82 | 0.016 | 0.951 | 0.015 | 0.956 | 0.956 | 0.015 |
| 138 | P-27-ROH | 27 | F | L3 | 6.62 | 2.93 | 0.031 | 0.937 | 0.031 | 0.937 | 0.937 | 0.031 |
| 139 | P-28-MAR | 52 | M | U2 | 12.86 | 13.24 | 0.068 | 0.921 | 0.068 | 0.921 | 0.921 | 0.068 |
| 140 | P-28-MAR | 52 | M | T1 | 9.93 | 9.42 | 0.047 | 0.911 | 0.047 | 0.913 | 0.913 | 0.047 |
| 141 | P-28-MAR | 52 | M | T2 | 3.69 | 4.84 | 0.025 | 0.924 | 0.025 | 0.925 | 0.925 | 0.025 |
| 142 | P-28-MAR | 52 | M | L2 | 20.63 | 16.67 | 0.036 | 0.978 | 0.033 | 0.979 | 0.979 | 0.033 |
| 143 | P-28-MAR | 52 | M | L3 | 18.60 | 18.84 | 0.049 | 0.965 | 0.051 | 0.961 | 0.965 | 0.049 |
| 144 | P-28-MAR | 52 | M | L5 | 24.84 | 27.87 | 0.044 | 0.989 | 0.046 | 0.986 | 0.989 | 0.044 |
| 145 | P-29-BAK | 39 | M | U3 | 10.44 | 10.05 | 0.031 | 0.955 | 0.030 | 0.955 | 0.955 | 0.030 |
| 146 | P-29-BAK | 39 | M | L1 | 9.93 | 9.29 | 0.044 | 0.938 | 0.044 | 0.939 | 0.939 | 0.044 |
| 147 | P-29-BAK | 39 | M | L2 | 17.96 | 8.02 | 0.040 | 0.950 | 0.042 | 0.935 | 0.950 | 0.040 |
| 148 | P-29-BAK | 39 | M | L4 | 11.72 | 14.89 | 0.032 | 0.960 | 0.032 | 0.960 | 0.960 | 0.032 |
| 149 | P-30-CHE | 27 | M | H1 | 4.97 | 6.24 | 0.023 | 0.984 | 0.023 | 0.984 | 0.984 | 0.023 |
| 150 | P-30-CHE | 27 | M | U1 | 14.65 | 8.15 | 0.052 | 0.939 | 0.045 | 0.920 | 0.939 | 0.052 |
| 151 | P-30-CHE | 27 | M | T1 | 18.21 | 13.87 | 0.054 | 0.975 | 0.053 | 0.975 | 0.975 | 0.053 |
| 152 | P-30-CHE | 27 | M | T2 | 17.96 | 11.84 | 0.037 | 0.936 | 0.033 | 0.948 | 0.948 | 0.033 |
| 153 | P-30-CHE | 27 | M | T3 | 25.22 | 15.27 | 0.077 | 0.923 | 0.077 | 0.924 | 0.924 | 0.077 |
| 154 | P-30-CHE | 27 | M | T4 | 14.14 | 26.22 | 0.048 | 0.910 | 0.049 | 0.906 | 0.910 | 0.048 |
| 155 | P-30-CHE | 27 | M | T5 | 24.07 | 13.24 | 0.059 | 0.950 | 0.059 | 0.951 | 0.951 | 0.059 |
| 156 | P-30-CHE | 27 | M | L1 | 22.42 | 23.80 | 0.040 | 0.986 | 0.042 | 0.984 | 0.986 | 0.040 |
| 157 | P-30-CHE | 27 | M | L32 | 16.30 | 26.73 | 0.051 | 0.960 | 0.050 | 0.961 | 0.961 | 0.050 |
| 158 | P-31-CHE | 32 | F | H1 | 11.08 | 10.18 | 0.061 | 0.920 | 0.061 | 0.921 | 0.921 | 0.061 |
| 159 | P-31-CHE | 32 | F | U2 | 17.19 | 17.95 | 0.057 | 0.940 | 0.053 | 0.946 | 0.946 | 0.053 |
| 160 | P-31-CHE | 32 | F | U3 | 20.76 | 12.22 | 0.060 | 0.917 | 0.059 | 0.918 | 0.918 | 0.059 |
| 161 | P-31-CHE | 32 | F | T1 | 16.17 | 21.38 | 0.055 | 0.942 | 0.053 | 0.948 | 0.948 | 0.053 |
| 162 | P-31-CHE | 32 | F | T4 | 12.10 | 9.93 | 0.055 | 0.934 | 0.054 | 0.935 | 0.935 | 0.054 |
| 163 | P-31-CHE | 32 | F | L1 | 13.37 | 9.54 | 0.044 | 0.953 | 0.043 | 0.943 | 0.953 | 0.044 |
| 164 | P-31-CHE | 32 | F | L3 | 18.98 | 16.04 | 0.029 | 0.966 | 0.029 | 0.966 | 0.966 | 0.029 |
| 165 | P-31-CHE | 32 | F | L4 | 16.30 | 15.65 | 0.042 | 0.978 | 0.042 | 0.978 | 0.978 | 0.042 |
| 166 | P-31-CHE | 32 | F | L5 | 8.79 | 10.05 | 0.034 | 0.949 | 0.033 | 0.951 | 0.951 | 0.033 |
| 167 | P-32-LEA | 48 | F | U3 | 14.52 | 14.26 | 0.040 | 0.967 | 0.040 | 0.967 | 0.967 | 0.040 |
| 168 | P-32-LEA | 48 | F | T1 | 15.41 | 14.00 | 0.033 | 0.951 | 0.033 | 0.952 | 0.952 | 0.033 |
| 169 | P-32-LEA | 48 | F | T2 | 11.46 | 14.13 | 0.031 | 0.948 | 0.030 | 0.950 | 0.950 | 0.030 |
| 170 | P-32-LEA | 48 | F | T3 | 16.05 | 11.33 | 0.049 | 0.970 | 0.048 | 0.971 | 0.971 | 0.048 |
| 171 | P-32-LEA | 48 | F | L3 | 16.17 | 11.58 | 0.089 | 0.950 | 0.089 | 0.951 | 0.951 | 0.089 |
| 172 | P-33-LAW | 59 | M | H1 | 8.92 | 9.04 | 0.038 | 0.940 | 0.038 | 0.941 | 0.941 | 0.038 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 173 | P-33-LAW | 59 | M | U1 | 11.59 | 12.98 | 0.034 | 0.980 | 0.034 | 0.980 | 0.980 | 0.034 |
| 174 | P-33-LAW | 59 | M | U3 | 10.70 | 10.82 | 0.034 | 0.951 | 0.033 | 0.954 | 0.954 | 0.033 |
| 175 | P-33-LAW | 59 | M | T1 | 21.27 | 16.29 | 0.069 | 0.967 | 0.073 | 0.963 | 0.967 | 0.069 |
| 176 | P-33-LAW | 59 | M | T2 | 25.73 | 14.00 | 0.058 | 0.970 | 0.056 | 0.972 | 0.972 | 0.056 |
| 177 | P-33-LAW | 59 | M | T3 | 18.34 | 13.36 | 0.066 | 0.959 | 0.064 | 0.960 | 0.960 | 0.064 |
| 178 | P-33-LAW | 59 | M | T4 | 34.26 | 27.36 | 0.065 | 0.953 | 0.072 | 0.952 | 0.953 | 0.065 |
| 179 | P-33-LAW | 59 | M | T5 | 19.23 | 12.47 | 0.053 | 0.967 | 0.051 | 0.968 | 0.968 | 0.051 |
| 180 | P-33-LAW | 59 | M | L1 | 8.28 | 10.82 | 0.031 | 0.945 | 0.030 | 0.947 | 0.947 | 0.030 |
| 181 | P-33-LAW | 59 | M | L3 | 8.79 | 10.94 | 0.046 | 0.963 | 0.046 | 0.964 | 0.964 | 0.046 |
| 182 | P-34-RIZ | 23 | F | H1 | 5.48 | 8.27 | 0.021 | 0.989 | 0.020 | 0.990 | 0.990 | 0.020 |
| 183 | P-34-RIZ | 23 | F | U1 | 10.83 | 24.82 | 0.040 | 0.984 | 0.039 | 0.984 | 0.984 | 0.039 |
| 184 | P-34-RIZ | 23 | F | T1 | 20.00 | 13.36 | 0.038 | 0.951 | 0.040 | 0.942 | 0.951 | 0.038 |
| 185 | P-34-RIZ | 23 | F | T2 | 12.99 | 7.38 | 0.047 | 0.901 | 0.044 | 0.911 | 0.911 | 0.044 |
| 186 | P-34-RIZ | 23 | F | T3 | 12.10 | 10.44 | 0.041 | 0.964 | 0.041 | 0.964 | 0.964 | 0.041 |
| 187 | P-34-RIZ | 23 | F | L1 | 18.34 | 10.95 | 0.031 | 0.960 | 0.031 | 0.958 | 0.960 | 0.031 |
| 188 | P-34-RIZ | 23 | F | L2 | 8.53 | 16.16 | 0.035 | 0.970 | 0.034 | 0.971 | 0.971 | 0.034 |
| 189 | P-34-RIZ | 23 | F | L3 | 13.63 | 8.15 | 0.041 | 0.959 | 0.041 | 0.940 | 0.959 | 0.041 |
| 190 | P-34-RIZ | 23 | F | L4 | 10.06 | 7.64 | 0.031 | 0.955 | 0.029 | 0.962 | 0.962 | 0.029 |
| 191 | P-34-RIZ | 23 | F | L5 | 29.04 | 15.65 | 0.068 | 0.960 | 0.069 | 0.957 | 0.960 | 0.068 |
| 192 | P-35-RAJ | 67 | M | H1 | 11.08 | 12.60 | 0.029 | 0.908 | 0.029 | 0.910 | 0.910 | 0.029 |
| 193 | P-35-RAJ | 67 | M | U1 | 20.00 | 12.22 | 0.044 | 0.950 | 0.043 | 0.953 | 0.953 | 0.043 |
| 194 | P-35-RAJ | 67 | M | U2 | 13.88 | 10.82 | 0.022 | 0.978 | 0.021 | 0.979 | 0.979 | 0.021 |
| 195 | P-35-RAJ | 67 | M | U3 | 10.06 | 8.78 | 0.037 | 0.925 | 0.035 | 0.921 | 0.925 | 0.037 |
| 196 | P-35-RAJ | 67 | M | T1 | 9.93 | 14.13 | 0.046 | 0.843 | 0.040 | 0.907 | 0.907 | 0.040 |
| 197 | P-35-RAJ | 67 | M | T2 | 21.01 | 17.95 | 0.044 | 0.972 | 0.045 | 0.968 | 0.972 | 0.044 |
| 198 | P-35-RAJ | 67 | M | T4 | 5.60 | 5.22 | 0.045 | 0.914 | 0.045 | 0.914 | 0.914 | 0.045 |
| 199 | P-35-RAJ | 67 | M | L1 | 7.64 | 15.40 | 0.037 | 0.985 | 0.035 | 0.986 | 0.986 | 0.035 |
| 200 | P-35-RAJ | 67 | M | L2 | 8.15 | 15.91 | 0.027 | 0.996 | 0.025 | 0.996 | 0.996 | 0.025 |
| 201 | P-36-NUR | 21 | F | H1 | 14.77 | 16.93 | 0.065 | 0.948 | 0.063 | 0.952 | 0.952 | 0.063 |
| 202 | P-36-NUR | 21 | F | U3 | 31.97 | 8.91 | 0.042 | 0.988 | 0.036 | 0.991 | 0.991 | 0.036 |
| 203 | P-36-NUR | 21 | F | T3 | 26.75 | 15.53 | 0.053 | 0.941 | 0.052 | 0.944 | 0.944 | 0.052 |
| 204 | P-36-NUR | 21 | F | L2 | 14.27 | 13.75 | 0.036 | 0.990 | 0.038 | 0.989 | 0.990 | 0.036 |
| 205 | P-36-NUR | 21 | F | L3 | 16.30 | 21.38 | 0.040 | 0.991 | 0.043 | 0.990 | 0.991 | 0.040 |
| 206 | P-36-NUR | 21 | F | L4 | 31.46 | 26.47 | 0.057 | 0.990 | 0.063 | 0.987 | 0.990 | 0.057 |
| 207 | P-37-VAD | 37 | F | H12 | 4.59 | 4.07 | 0.032 | 0.951 | 0.031 | 0.955 | 0.955 | 0.031 |
| 208 | P-37-VAD | 37 | F | U1 | 2.55 | 3.06 | 0.016 | 0.959 | 0.016 | 0.959 | 0.959 | 0.016 |
| 209 | P-37-VAD | 37 | F | U2 | 2.55 | 3.18 | 0.016 | 0.953 | 0.016 | 0.954 | 0.954 | 0.016 |
| 210 | P-37-VAD | 37 | F | T1 | 11.08 | 11.33 | 0.022 | 0.973 | 0.021 | 0.975 | 0.975 | 0.021 |
| 211 | P-37-VAD | 37 | F | T2 | 5.10 | 6.11 | 0.019 | 0.973 | 0.019 | 0.973 | 0.973 | 0.019 |
| 212 | P-37-VAD | 37 | F | T3 | 4.33 | 6.11 | 0.048 | 0.936 | 0.047 | 0.939 | 0.939 | 0.047 |
| 213 | P-38-ROM | 43 | F | H1 | 3.31 | 4.07 | 0.028 | 0.904 | 0.034 | 0.692 | 0.904 | 0.028 |
| 214 | P-38-ROM | 43 | F | U1 | 10.19 | 12.22 | 0.022 | 0.952 | 0.019 | 0.947 | 0.952 | 0.022 |
| 215 | P-38-ROM | 43 | F | U2 | 4.20 | 5.85 | 0.017 | 0.957 | 0.017 | 0.957 | 0.957 | 0.017 |
| 216 | P-38-ROM | 43 | F | U3 | 3.44 | 4.71 | 0.016 | 0.943 | 0.017 | 0.943 | 0.943 | 0.017 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 217 | P-38-ROM | 43 | F | T2 | 8.53 | 7.76 | 0.016 | 0.938 | 0.016 | 0.936 | 0.938 | 0.016 |
| 218 | P-38-ROM | 43 | F | T3 | 8.02 | 6.62 | 0.022 | 0.977 | 0.022 | 0.977 | 0.977 | 0.022 |
| 219 | P-38-ROM | 43 | F | L1 | 5.09 | 7.89 | 0.018 | 0.936 | 0.018 | 0.938 | 0.938 | 0.018 |
| 220 | P-38-ROM | 43 | F | L2 | 9.04 | 7.00 | 0.018 | 0.963 | 0.017 | 0.964 | 0.964 | 0.017 |
| 221 | P-38-ROM | 43 | F | L3 | 3.06 | 3.69 | 0.016 | 0.957 | 0.016 | 0.959 | 0.959 | 0.016 |
| 222 | P-39-MOH | 44 | M | U1 | 7.13 | 9.80 | 0.048 | 0.949 | 0.048 | 0.949 | 0.949 | 0.048 |
| 223 | P-39-MOH | 44 | M | U2 | 2.55 | 4.46 | 0.023 | 0.914 | 0.023 | 0.913 | 0.914 | 0.023 |
| 224 | P-39-MOH | 44 | M | U3 | 5.35 | 10.18 | 0.022 | 0.985 | 0.020 | 0.988 | 0.988 | 0.020 |
| 225 | P-39-MOH | 44 | M | T1 | 11.08 | 7.76 | 0.030 | 0.964 | 0.028 | 0.970 | 0.970 | 0.028 |
| 226 | P-39-MOH | 44 | M | T2 | 10.83 | 14.00 | 0.023 | 0.933 | 0.022 | 0.937 | 0.937 | 0.022 |
| 227 | P-39-MOH | 44 | M | T3 | 18.85 | 6.75 | 0.033 | 0.981 | 0.032 | 0.982 | 0.982 | 0.032 |
| 228 | P-39-MOH | 44 | M | L1 | 8.41 | 10.69 | 0.054 | 0.940 | 0.052 | 0.946 | 0.946 | 0.052 |
| 229 | P-39-MOH | 44 | M | L3 | 8.79 | 7.51 | 0.069 | 0.925 | 0.068 | 0.927 | 0.927 | 0.068 |
| 230 | P-40-MOH | 46 | M | H1 | 5.60 | 9.67 | 0.018 | 0.970 | 0.017 | 0.971 | 0.971 | 0.017 |
| 231 | P-40-MOH | 46 | M | U1 | 9.42 | 10.44 | 0.044 | 0.940 | 0.041 | 0.946 | 0.946 | 0.041 |
| 232 | P-40-MOH | 46 | M | U3 | 7.26 | 9.42 | 0.030 | 0.949 | 0.030 | 0.949 | 0.949 | 0.030 |
| 233 | P-40-MOH | 46 | M | L1 | 14.90 | 15.65 | 0.053 | 0.968 | 0.058 | 0.961 | 0.968 | 0.053 |
| 234 | P-40-MOH | 46 | M | L2 | 7.77 | 8.91 | 0.034 | 0.952 | 0.034 | 0.954 | 0.954 | 0.034 |
| 235 | P-41-SHA | 59 | M | H1 | 11.97 | 15.91 | 0.059 | 0.919 | 0.059 | 0.921 | 0.921 | 0.059 |
| 236 | P-41-SHA | 59 | M | U1 | 4.59 | 9.16 | 0.024 | 0.952 | 0.023 | 0.955 | 0.955 | 0.023 |
| 237 | P-41-SHA | 59 | M | U2 | 3.69 | 4.07 | 0.016 | 0.915 | 0.016 | 0.927 | 0.927 | 0.016 |
| 238 | P-41-SHA | 59 | M | U3 | 4.46 | 4.58 | 0.015 | 0.926 | 0.015 | 0.926 | 0.926 | 0.015 |
| 239 | P-41-SHA | 59 | M | T1 | 4.33 | 4.33 | 0.023 | 0.939 | 0.024 | 0.938 | 0.939 | 0.023 |
| 240 | P-41-SHA | 59 | M | T2 | 4.71 | 6.49 | 0.018 | 0.968 | 0.018 | 0.969 | 0.969 | 0.018 |
| 241 | P-41-SHA | 59 | M | T3 | 3.57 | 6.11 | 0.022 | 0.929 | 0.021 | 0.936 | 0.936 | 0.021 |
| 242 | P-41-SHA | 59 | M | L2 | 14.39 | 28.00 | 0.069 | 0.957 | 0.066 | 0.964 | 0.964 | 0.066 |
| 243 | P-41-SHA | 59 | M | L3 | 5.60 | 6.11 | 0.027 | 0.909 | 0.021 | 0.907 | 0.909 | 0.027 |
| 244 | P-42-SEL | 34 | M | H1 | 4.20 | 3.69 | 0.020 | 0.952 | 0.018 | 0.963 | 0.963 | 0.018 |
| 245 | P-42-SEL | 34 | M | U1 | 6.37 | 5.47 | 0.021 | 0.987 | 0.021 | 0.985 | 0.987 | 0.021 |
| 246 | P-42-SEL | 34 | M | U2 | 3.31 | 4.58 | 0.023 | 0.946 | 0.022 | 0.946 | 0.946 | 0.022 |
| 247 | P-42-SEL | 34 | M | T1 | 7.00 | 4.96 | 0.032 | 0.964 | 0.032 | 0.964 | 0.964 | 0.032 |
| 248 | P-42-SEL | 34 | M | T3 | 3.44 | 3.31 | 0.032 | 0.944 | 0.033 | 0.943 | 0.944 | 0.032 |
| 249 | P-42-SEL | 34 | M | L1 | 2.55 | 3.18 | 0.016 | 0.962 | 0.016 | 0.962 | 0.962 | 0.016 |
| 250 | P-42-SEL | 34 | M | L2 | 5.48 | 4.46 | 0.031 | 0.906 | 0.031 | 0.906 | 0.906 | 0.031 |
| 251 | P-42-SEL | 34 | M | L3 | 8.53 | 9.55 | 0.040 | 0.904 | 0.037 | 0.918 | 0.918 | 0.037 |
| 252 | P-43-MUK | 56 | M | H1 | 9.04 | 9.54 | 0.035 | 0.975 | 0.037 | 0.973 | 0.975 | 0.035 |
| 253 | P-43-MUK | 56 | M | U1 | 9.17 | 8.02 | 0.032 | 0.980 | 0.033 | 0.979 | 0.980 | 0.032 |
| 254 | P-43-MUK | 56 | M | U2 | 5.22 | 6.36 | 0.021 | 0.987 | 0.021 | 0.987 | 0.987 | 0.021 |
| 255 | P-43-MUK | 56 | M | U3 | 4.71 | 7.51 | 0.072 | 0.906 | 0.072 | 0.908 | 0.908 | 0.072 |
| 256 | P-43-MUK | 56 | M | T2 | 6.62 | 4.96 | 0.019 | 0.975 | 0.019 | 0.976 | 0.976 | 0.019 |
| 257 | P-43-MUK | 56 | M | T3 | 5.60 | 4.84 | 0.034 | 0.959 | 0.034 | 0.959 | 0.959 | 0.034 |
| 258 | P-43-MUK | 56 | M | L1 | 6.11 | 7.00 | 0.020 | 0.958 | 0.021 | 0.947 | 0.958 | 0.020 |
| 259 | P-43-MUK | 56 | M | L2 | 3.82 | 5.09 | 0.028 | 0.938 | 0.027 | 0.925 | 0.938 | 0.028 |
| 260 | P-43-MUK | 56 | M | L3 | 4.58 | 5.73 | 0.014 | 0.984 | 0.014 | 0.984 | 0.984 | 0.014 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 261 | P-44-RAM | 38 | M | H1 | 14.14 | 10.95 | 0.037 | 0.976 | 0.041 | 0.972 | 0.976 | 0.037 |
| 262 | P-44-RAM | 38 | M | U1 | 8.40 | 8.53 | 0.043 | 0.906 | 0.043 | 0.906 | 0.906 | 0.043 |
| 263 | P-44-RAM | 38 | M | U2 | 5.99 | 8.02 | 0.042 | 0.962 | 0.042 | 0.961 | 0.962 | 0.042 |
| 264 | P-44-RAM | 38 | M | T2 | 10.44 | 9.54 | 0.016 | 0.984 | 0.015 | 0.985 | 0.985 | 0.015 |
| 265 | P-44-RAM | 38 | M | T3 | 6.11 | 10.05 | 0.028 | 0.963 | 0.026 | 0.968 | 0.968 | 0.026 |
| 266 | P-44-RAM | 38 | M | L1 | 7.77 | 8.78 | 0.026 | 0.983 | 0.028 | 0.981 | 0.983 | 0.026 |
| 267 | P-44-RAM | 38 | M | L2 | 7.64 | 6.49 | 0.020 | 0.952 | 0.020 | 0.950 | 0.952 | 0.020 |
| 268 | P-44-RAM | 38 | M | L3 | 13.50 | 11.07 | 0.036 | 0.950 | 0.036 | 0.950 | 0.950 | 0.036 |
| 269 | P-45-CHA | 67 | M | U3 | 4.08 | 6.11 | 0.035 | 0.930 | 0.035 | 0.931 | 0.931 | 0.035 |
| 270 | P-45-CHA | 67 | M | T1 | 17.07 | 18.07 | 0.043 | 0.950 | 0.040 | 0.957 | 0.957 | 0.040 |
| 271 | P-45-CHA | 67 | M | T2 | 7.13 | 8.15 | 0.038 | 0.944 | 0.039 | 0.942 | 0.944 | 0.038 |
| 272 | P-45-CHA | 67 | M | T3 | 25.22 | 20.49 | 0.036 | 0.988 | 0.045 | 0.969 | 0.988 | 0.036 |
| 273 | P-45-CHA | 67 | M | L1 | 19.10 | 13.75 | 0.048 | 0.954 | 0.048 | 0.954 | 0.954 | 0.048 |
| 274 | P-45-CHA | 67 | M | L2 | 18.21 | 12.73 | 0.036 | 0.981 | 0.035 | 0.982 | 0.982 | 0.035 |
| 275 | P-45-CHA | 67 | M | L3 | 18.98 | 14.38 | 0.035 | 0.971 | 0.034 | 0.972 | 0.972 | 0.034 |
| 276 | P-46-SUB | 61 | M | T1 | 5.48 | 4.33 | 0.027 | 0.920 | 0.027 | 0.918 | 0.920 | 0.027 |
| 277 | P-46-SUB | 61 | M | T2 | 2.42 | 1.78 | 0.012 | 0.971 | 0.012 | 0.971 | 0.971 | 0.012 |
| 278 | P-46-SUB | 61 | M | T3 | 2.80 | 4.20 | 0.021 | 0.961 | 0.020 | 0.961 | 0.961 | 0.020 |
| 279 | P-46-SUB | 61 | M | L1 | 28.02 | 9.29 | 0.023 | 0.983 | 0.021 | 0.985 | 0.985 | 0.021 |
| 280 | P-46-SUB | 61 | M | L2 | 9.68 | 10.44 | 0.019 | 0.967 | 0.019 | 0.968 | 0.968 | 0.019 |
| 281 | P-46-SUB | 61 | M | L3 | 7.26 | 9.04 | 0.017 | 0.931 | 0.017 | 0.932 | 0.932 | 0.017 |
| 282 | P-47-MUH | 34 | M | U1 | 17.32 | 10.06 | 0.039 | 0.950 | 0.039 | 0.931 | 0.950 | 0.039 |
| 283 | P-47-MUH | 34 | M | U2 | 6.24 | 7.26 | 0.017 | 0.979 | 0.017 | 0.979 | 0.979 | 0.017 |
| 284 | P-47-MUH | 34 | M | U3 | 6.37 | 7.13 | 0.016 | 0.987 | 0.016 | 0.988 | 0.988 | 0.016 |
| 285 | P-47-MUH | 34 | M | T1 | 8.53 | 8.15 | 0.033 | 0.976 | 0.032 | 0.978 | 0.978 | 0.032 |
| 286 | P-47-MUH | 34 | M | T2 | 5.60 | 7.00 | 0.019 | 0.956 | 0.017 | 0.941 | 0.956 | 0.019 |
| 287 | P-47-MUH | 34 | M | L1 | 15.28 | 16.42 | 0.057 | 0.991 | 0.056 | 0.991 | 0.991 | 0.056 |
| 288 | P-47-MUH | 34 | M | L2 | 12.86 | 17.56 | 0.038 | 0.952 | 0.039 | 0.950 | 0.952 | 0.038 |
| 289 | P-48-WON | 49 | M | U1 | 8.66 | 10.18 | 0.028 | 0.968 | 0.028 | 0.969 | 0.969 | 0.028 |
| 290 | P-48-WON | 49 | M | U2 | 6.88 | 8.02 | 0.038 | 0.967 | 0.038 | 0.967 | 0.967 | 0.038 |
| 291 | P-48-WON | 49 | M | T1 | 10.06 | 12.60 | 0.041 | 0.968 | 0.041 | 0.967 | 0.968 | 0.041 |
| 292 | P-48-WON | 49 | M | T2 | 8.28 | 14.25 | 0.037 | 0.964 | 0.040 | 0.959 | 0.964 | 0.037 |
| 293 | P-48-WON | 49 | M | L1 | 24.07 | 15.02 | 0.040 | 0.991 | 0.034 | 0.993 | 0.993 | 0.034 |
| 294 | P-48-WON | 49 | M | L2 | 5.86 | 6.24 | 0.024 | 0.957 | 0.024 | 0.954 | 0.957 | 0.024 |
| 295 | P-48-WON | 49 | M | L3 | 2.80 | 3.31 | 0.017 | 0.961 | 0.017 | 0.961 | 0.961 | 0.017 |
| 296 | P-49-MUN | 47 | M | H1 | 16.30 | 12.35 | 0.052 | 0.987 | 0.048 | 0.990 | 0.990 | 0.048 |
| 297 | P-49-MUN | 47 | M | U1 | 6.50 | 7.51 | 0.038 | 0.952 | 0.036 | 0.952 | 0.952 | 0.036 |
| 298 | P-49-MUN | 47 | M | U2 | 5.60 | 6.62 | 0.020 | 0.976 | 0.019 | 0.978 | 0.978 | 0.019 |
| 299 | P-49-MUN | 47 | M | T1 | 10.44 | 8.65 | 0.046 | 0.973 | 0.045 | 0.973 | 0.973 | 0.045 |
| 300 | P-49-MUN | 47 | M | T3 | 7.01 | 6.24 | 0.042 | 0.929 | 0.040 | 0.934 | 0.934 | 0.040 |
| 301 | P-49-MUN | 47 | M | L1 | 6.75 | 12.60 | 0.031 | 0.904 | 0.030 | 0.914 | 0.914 | 0.030 |
| 302 | P-49-MUN | 47 | M | L3B | 9.42 | 10.44 | 0.039 | 0.920 | 0.040 | 0.919 | 0.920 | 0.039 |
| 303 | P-50-LIM | 57 | M | H1 | 10.06 | 9.42 | 0.039 | 0.993 | 0.037 | 0.994 | 0.994 | 0.037 |
| 304 | P-50-LIM | 57 | M | U3 | 4.84 | 7.26 | 0.029 | 0.952 | 0.028 | 0.952 | 0.952 | 0.028 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 305 | P-50-LIM | 57 | M | T1 | 9.81 | 6.75 | 0.030 | 0.947 | 0.029 | 0.949 | 0.949 | 0.029 |
| 306 | P-50-LIM | 57 | M | T2 | 11.46 | 5.60 | 0.027 | 0.911 | 0.026 | 0.914 | 0.914 | 0.026 |
| 307 | P-50-LIM | 57 | M | T3 | 6.24 | 12.09 | 0.067 | 0.924 | 0.074 | 0.921 | 0.924 | 0.067 |
| 308 | P-50-LIM | 57 | M | T4 | 36.43 | 25.58 | 0.076 | 0.974 | 0.080 | 0.992 | 0.992 | 0.080 |
| 309 | P-50-LIM | 57 | M | L1 | 7.64 | 7.76 | 0.020 | 0.972 | 0.020 | 0.972 | 0.972 | 0.020 |
| 310 | P-50-LIM | 57 | M | L3 | 5.99 | 7.76 | 0.032 | 0.960 | 0.032 | 0.961 | 0.961 | 0.032 |
| 311 | P-51-TEN | 32 | M | U1 | 14.77 | 17.31 | 0.027 | 0.989 | 0.027 | 0.990 | 0.990 | 0.027 |
| 312 | P-51-TEN | 32 | M | U3 | 6.11 | 5.85 | 0.033 | 0.939 | 0.033 | 0.938 | 0.939 | 0.033 |
| 313 | P-51-TEN | 32 | M | T1 | 7.51 | 7.38 | 0.035 | 0.901 | 0.030 | 0.918 | 0.918 | 0.030 |
| 314 | P-51-TEN | 32 | M | T2 | 7.51 | 10.95 | 0.029 | 0.918 | 0.026 | 0.933 | 0.933 | 0.026 |
| 315 | P-51-TEN | 32 | M | T3 | 9.55 | 11.96 | 0.027 | 0.956 | 0.028 | 0.953 | 0.956 | 0.027 |
| 316 | P-51-TEN | 32 | M | L1 | 11.08 | 14.51 | 0.034 | 0.992 | 0.034 | 0.992 | 0.992 | 0.034 |
| 317 | P-51-TEN | 32 | M | L2 | 20.00 | 14.13 | 0.054 | 0.955 | 0.052 | 0.957 | 0.957 | 0.052 |
| 318 | P-51-TEN | 32 | M | L3 | 17.45 | 21.38 | 0.033 | 0.993 | 0.036 | 0.992 | 0.993 | 0.033 |
| 319 | P-52-VIJ | 35 | M | H1 | 5.86 | 7.89 | 0.026 | 0.914 | 0.026 | 0.917 | 0.917 | 0.026 |
| 320 | P-52-VIJ | 35 | M | U1 | 25.47 | 20.11 | 0.072 | 0.981 | 0.071 | 0.982 | 0.982 | 0.071 |
| 321 | P-52-VIJ | 35 | M | U2 | 14.39 | 19.98 | 0.038 | 0.982 | 0.039 | 0.981 | 0.982 | 0.038 |
| 322 | P-52-VIJ | 35 | M | U3 | 16.81 | 23.67 | 0.055 | 0.985 | 0.053 | 0.985 | 0.985 | 0.053 |
| 323 | P-52-VIJ | 35 | M | T1 | 18.98 | 23.80 | 0.052 | 0.925 | 0.048 | 0.940 | 0.940 | 0.048 |
| 324 | P-52-VIJ | 35 | M | T2 | 12.99 | 10.44 | 0.039 | 0.921 | 0.042 | 0.941 | 0.941 | 0.042 |
| 325 | P-52-VIJ | 35 | M | T3 | 10.06 | 10.18 | 0.039 | 0.984 | 0.039 | 0.984 | 0.984 | 0.039 |
| 326 | P-52-VIJ | 35 | M | T5 | 18.34 | 25.33 | 0.082 | 0.970 | 0.081 | 0.971 | 0.971 | 0.081 |
| 327 | P-52-VIJ | 35 | M | L1 | 18.60 | 9.29 | 0.026 | 0.979 | 0.025 | 0.981 | 0.981 | 0.025 |
| 328 | P-52-VIJ | 35 | M | L2 | 27.51 | 23.80 | 0.073 | 0.977 | 0.081 | 0.969 | 0.977 | 0.073 |
| 329 | P-52-VIJ | 35 | M | L3 | 13.63 | 11.46 | 0.030 | 0.988 | 0.029 | 0.988 | 0.988 | 0.029 |
| 330 | P-53-KAM | 63 | M | U1 | 4.84 | 6.36 | 0.017 | 0.968 | 0.018 | 0.967 | 0.968 | 0.017 |
| 331 | P-53-KAM | 63 | M | U2 | 4.46 | 6.24 | 0.029 | 0.961 | 0.031 | 0.948 | 0.961 | 0.029 |
| 332 | P-53-KAM | 63 | M | T1 | 5.86 | 12.22 | 0.031 | 0.953 | 0.030 | 0.946 | 0.953 | 0.031 |
| 333 | P-53-KAM | 63 | M | T2 | 6.24 | 7.38 | 0.017 | 0.946 | 0.017 | 0.947 | 0.947 | 0.017 |
| 334 | P-53-KAM | 63 | M | T3 | 20.63 | 19.85 | 0.067 | 0.953 | 0.066 | 0.954 | 0.954 | 0.066 |
| 335 | P-53-KAM | 63 | M | T4 | 3.44 | 3.44 | 0.038 | 0.919 | 0.038 | 0.919 | 0.919 | 0.038 |
| 336 | P-53-KAM | 63 | M | L1 | 6.49 | 5.73 | 0.023 | 0.933 | 0.022 | 0.936 | 0.936 | 0.022 |
| 337 | P-53-KAM | 63 | M | L3 | 9.68 | 7.76 | 0.025 | 0.945 | 0.028 | 0.930 | 0.945 | 0.025 |
| 338 | P-54-LIE | 52 | M | U1 | 5.73 | 5.86 | 0.045 | 0.938 | 0.045 | 0.939 | 0.939 | 0.045 |
| 339 | P-54-LIE | 52 | M | U2 | 5.48 | 4.96 | 0.031 | 0.976 | 0.031 | 0.976 | 0.976 | 0.031 |
| 340 | P-54-LIE | 52 | M | U3 | 5.60 | 5.86 | 0.030 | 0.906 | 0.029 | 0.907 | 0.907 | 0.029 |
| 341 | P-54-LIE | 52 | M | T1 | 15.28 | 11.84 | 0.042 | 0.962 | 0.041 | 0.964 | 0.964 | 0.041 |
| 342 | P-54-LIE | 52 | M | T5 | 12.35 | 8.27 | 0.025 | 0.947 | 0.024 | 0.949 | 0.949 | 0.024 |
| 343 | P-54-LIE | 52 | M | L1 | 4.46 | 5.73 | 0.027 | 0.962 | 0.028 | 0.961 | 0.962 | 0.027 |
| 344 | P-54-LIE | 52 | M | L3 | 4.97 | 8.27 | 0.039 | 0.923 | 0.037 | 0.931 | 0.931 | 0.037 |
| 345 | P-55-NED | 42 | M | H1 | 8.28 | 5.85 | 0.018 | 0.991 | 0.016 | 0.993 | 0.993 | 0.016 |
| 346 | P-55-NED | 42 | M | U1 | 11.59 | 10.82 | 0.043 | 0.975 | 0.041 | 0.978 | 0.978 | 0.041 |
| 347 | P-55-NED | 42 | M | U3 | 16.17 | 14.51 | 0.086 | 0.981 | 0.086 | 0.981 | 0.981 | 0.086 |
| 348 | P-55-NED | 42 | M | T1 | 13.12 | 14.51 | 0.054 | 0.982 | 0.055 | 0.981 | 0.982 | 0.054 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 349 | P-55-NED | 42 | M | T2 | 11.34 | 11.84 | 0.035 | 0.923 | 0.034 | 0.928 | 0.928 | 0.034 |
| 350 | P-55-NED | 42 | M | T3 | 3.95 | 6.87 | 0.019 | 0.968 | 0.017 | 0.976 | 0.976 | 0.017 |
| 351 | P-55-NED | 42 | M | L1 | 32.60 | 10.18 | 0.057 | 0.962 | 0.056 | 0.967 | 0.967 | 0.056 |
| 352 | P-55-NED | 42 | M | L2 | 7.64 | 6.62 | 0.027 | 0.965 | 0.027 | 0.963 | 0.965 | 0.027 |
| 353 | P-55-NED | 42 | M | L3 | 9.93 | 14.38 | 0.027 | 0.954 | 0.031 | 0.945 | 0.954 | 0.027 |
| 354 | P-55-NED | 42 | M | L4 | 32.22 | 27.11 | 0.084 | 0.971 | 0.103 | 0.946 | 0.971 | 0.084 |
| 355 | P-56-ROS | 53 | M | U1 | 13.63 | 11.45 | 0.044 | 0.983 | 0.042 | 0.985 | 0.985 | 0.042 |
| 356 | P-56-ROS | 53 | M | U2 | 8.79 | 11.07 | 0.052 | 0.919 | 0.052 | 0.914 | 0.919 | 0.052 |
| 357 | P-56-ROS | 53 | M | U3 | 9.04 | 9.29 | 0.041 | 0.903 | 0.042 | 0.901 | 0.903 | 0.041 |
| 358 | P-56-ROS | 53 | M | T12 | 12.99 | 17.82 | 0.074 | 0.929 | 0.066 | 0.947 | 0.947 | 0.066 |
| 359 | P-56-ROS | 53 | M | T2 | 5.99 | 4.96 | 0.023 | 0.928 | 0.021 | 0.941 | 0.941 | 0.021 |
| 360 | P-56-ROS | 53 | M | L1 | 8.92 | 7.38 | 0.020 | 0.938 | 0.022 | 0.929 | 0.938 | 0.020 |
| 361 | P-56-ROS | 53 | M | L2 | 13.76 | 18.96 | 0.095 | 0.946 | 0.093 | 0.949 | 0.949 | 0.093 |
| 362 | P-56-ROS | 53 | M | L3 | 13.63 | 12.09 | 0.034 | 0.931 | 0.034 | 0.932 | 0.932 | 0.034 |
| 363 | P-56-ROS | 53 | M | L4 | 30.18 | 22.15 | 0.098 | 0.951 | 0.106 | 0.952 | 0.952 | 0.106 |
| 364 | P-56-ROS | 53 | M | L5 | 27.89 | 25.33 | 0.071 | 0.944 | 0.072 | 0.949 | 0.949 | 0.072 |
| 365 | P-56-ROS | 53 | M | L6 | 12.48 | 10.94 | 0.090 | 0.914 | 0.089 | 0.908 | 0.914 | 0.090 |
| 366 | P-57-RAM | 56 | F | H | 5.99 | 7.89 | 0.026 | 0.980 | 0.025 | 0.982 | 0.982 | 0.025 |
| 367 | P-57-RAM | 56 | F | U1 | 4.58 | 5.86 | 0.023 | 0.967 | 0.023 | 0.966 | 0.967 | 0.023 |
| 368 | P-57-RAM | 56 | F | U2 | 9.68 | 13.24 | 0.042 | 0.981 | 0.042 | 0.981 | 0.981 | 0.042 |
| 369 | P-57-RAM | 56 | F | U3 | 3.95 | 5.35 | 0.024 | 0.956 | 0.022 | 0.961 | 0.961 | 0.022 |
| 370 | P-57-RAM | 56 | F | T1 | 13.37 | 20.49 | 0.041 | 0.908 | 0.039 | 0.911 | 0.911 | 0.039 |
| 371 | P-57-RAM | 56 | F | T2 | 10.06 | 10.44 | 0.041 | 0.914 | 0.041 | 0.914 | 0.914 | 0.041 |
| 372 | P-57-RAM | 56 | F | T3 | 7.77 | 8.78 | 0.022 | 0.932 | 0.022 | 0.934 | 0.934 | 0.022 |
| 373 | P-57-RAM | 56 | F | L1 | 7.51 | 9.67 | 0.037 | 0.941 | 0.036 | 0.944 | 0.944 | 0.036 |
| 374 | P-57-RAM | 56 | F | L2 | 6.50 | 9.42 | 0.029 | 0.970 | 0.027 | 0.974 | 0.974 | 0.027 |
| 375 | P-58-MOH | 31 | M | H1 | 7.00 | 5.73 | 0.022 | 0.969 | 0.022 | 0.970 | 0.970 | 0.022 |
| 376 | P-58-MOH | 31 | M | U1 | 3.06 | 3.06 | 0.034 | 0.917 | 0.034 | 0.917 | 0.917 | 0.034 |
| 377 | P-58-MOH | 31 | M | U2 | 2.42 | 3.31 | 0.023 | 0.968 | 0.023 | 0.967 | 0.968 | 0.023 |
| 378 | P-58-MOH | 31 | M | U3 | 7.51 | 9.55 | 0.044 | 0.919 | 0.044 | 0.920 | 0.920 | 0.044 |
| 379 | P-58-MOH | 31 | M | T2 | 2.29 | 2.93 | 0.014 | 0.917 | 0.015 | 0.915 | 0.917 | 0.014 |
| 380 | P-58-MOH | 31 | M | L3 | 11.84 | 12.47 | 0.030 | 0.987 | 0.030 | 0.988 | 0.988 | 0.030 |
| 381 | P-59-TAM | 47 | M | U2 | 12.86 | 22.53 | 0.028 | 0.978 | 0.026 | 0.969 | 0.978 | 0.028 |
| 382 | P-59-TAM | 47 | M | T1 | 17.58 | 18.84 | 0.026 | 0.995 | 0.031 | 0.993 | 0.995 | 0.026 |
| 383 | P-59-TAM | 47 | M | L11 | 12.99 | 21.13 | 0.028 | 0.957 | 0.028 | 0.958 | 0.958 | 0.028 |
| 384 | P-60-JAM | 59 | M | H1 | 12.99 | 15.27 | 0.040 | 0.988 | 0.039 | 0.989 | 0.989 | 0.039 |
| 385 | P-60-JAM | 59 | M | L12 | 36.43 | 28.25 | 0.053 | 0.989 | 0.058 | 0.987 | 0.989 | 0.053 |
| 386 | P-61-DUR | 28 | M | H1 | 5.60 | 6.49 | 0.051 | 0.939 | 0.051 | 0.938 | 0.939 | 0.051 |
| 387 | P-61-DUR | 28 | M | U1 | 4.58 | 5.09 | 0.015 | 0.964 | 0.015 | 0.965 | 0.965 | 0.015 |
| 388 | P-61-DUR | 28 | M | U3 | 2.80 | 3.44 | 0.015 | 0.938 | 0.015 | 0.938 | 0.938 | 0.015 |
| 389 | P-61-DUR | 28 | M | T1 | 2.80 | 4.20 | 0.015 | 0.933 | 0.015 | 0.932 | 0.933 | 0.015 |
| 390 | P-61-DUR | 28 | M | T2 | 4.07 | 4.20 | 0.019 | 0.921 | 0.019 | 0.918 | 0.921 | 0.019 |
| 391 | P-61-DUR | 28 | M | T3 | 5.99 | 6.87 | 0.015 | 0.970 | 0.014 | 0.971 | 0.971 | 0.014 |
| 392 | P-61-DUR | 28 | M | L1 | 5.98 | 7.64 | 0.020 | 0.957 | 0.019 | 0.958 | 0.958 | 0.019 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 393 | P-61-DUR | 28 | M | L2 | 5.60 | 5.22 | 0.022 | 0.945 | 0.023 | 0.945 | 0.945 | 0.023 |
| 394 | P-61-DUR | 28 | M | L3 | 3.44 | 4.20 | 0.038 | 0.930 | 0.038 | 0.930 | 0.930 | 0.038 |
| 395 | P-62-VEE | 50 | M | H1 | 11.08 | 16.42 | 0.101 | 0.926 | 0.084 | 0.946 | 0.946 | 0.084 |
| 396 | P-62-VEE | 50 | M | U1 | 3.95 | 3.31 | 0.026 | 0.945 | 0.026 | 0.946 | 0.946 | 0.026 |
| 397 | P-62-VEE | 50 | M | U2 | 13.88 | 14.26 | 0.052 | 0.909 | 0.052 | 0.911 | 0.911 | 0.052 |
| 398 | P-62-VEE | 50 | M | U3 | 22.16 | 11.20 | 0.053 | 0.968 | 0.053 | 0.968 | 0.968 | 0.053 |
| 399 | P-62-VEE | 50 | M | T2 | 8.92 | 10.69 | 0.036 | 0.961 | 0.037 | 0.946 | 0.961 | 0.036 |
| 400 | P-62-VEE | 50 | M | T3 | 11.34 | 11.07 | 0.040 | 0.954 | 0.035 | 0.962 | 0.962 | 0.035 |
| 401 | P-62-VEE | 50 | M | T4 | 37.19 | 29.02 | 0.052 | 0.978 | 0.052 | 0.976 | 0.978 | 0.052 |
| 402 | P-62-VEE | 50 | M | T5 | 36.93 | 28.76 | 0.089 | 0.931 | 0.094 | 0.926 | 0.931 | 0.089 |
| 403 | P-62-VEE | 50 | M | L1 | 15.67 | 14.89 | 0.038 | 0.982 | 0.037 | 0.983 | 0.983 | 0.037 |
| 404 | P-62-VEE | 50 | M | L2 | 18.09 | 16.42 | 0.028 | 0.981 | 0.030 | 0.980 | 0.981 | 0.028 |
| 405 | P-62-VEE | 50 | M | L3 | 16.56 | 14.13 | 0.047 | 0.933 | 0.045 | 0.939 | 0.939 | 0.045 |
| 406 | P-63-MUN | 32 | M | H1 | 7.90 | 14.38 | 0.048 | 0.929 | 0.046 | 0.934 | 0.934 | 0.046 |
| 407 | P-63-MUN | 32 | M | U3 | 5.48 | 5.98 | 0.038 | 0.924 | 0.036 | 0.930 | 0.930 | 0.036 |
| 408 | P-63-MUN | 32 | M | T2 | 10.32 | 9.55 | 0.035 | 0.936 | 0.035 | 0.937 | 0.937 | 0.035 |
| 409 | P-63-MUN | 32 | M | T3 | 13.63 | 14.89 | 0.066 | 0.947 | 0.064 | 0.947 | 0.947 | 0.064 |
| 410 | P-63-MUN | 32 | M | L1 | 7.00 | 7.89 | 0.030 | 0.975 | 0.030 | 0.975 | 0.975 | 0.030 |
| 411 | P-63-MUN | 32 | M | L2 | 7.00 | 8.65 | 0.026 | 0.969 | 0.026 | 0.969 | 0.969 | 0.026 |
| 412 | P-64-PAL | 55 | M | H1 | 7.39 | 9.80 | 0.044 | 0.925 | 0.044 | 0.926 | 0.926 | 0.044 |
| 413 | P-64-PAL | 55 | M | U1 | 12.35 | 14.76 | 0.030 | 0.986 | 0.028 | 0.988 | 0.988 | 0.028 |
| 414 | P-64-PAL | 55 | M | U2 | 6.37 | 8.27 | 0.025 | 0.951 | 0.024 | 0.953 | 0.953 | 0.024 |
| 415 | P-64-PAL | 55 | M | U3 | 11.84 | 13.36 | 0.038 | 0.968 | 0.036 | 0.952 | 0.968 | 0.038 |
| 416 | P-64-PAL | 55 | M | T1 | 11.08 | 13.49 | 0.048 | 0.934 | 0.048 | 0.933 | 0.934 | 0.048 |
| 417 | P-64-PAL | 55 | M | T2 | 10.83 | 9.55 | 0.042 | 0.972 | 0.041 | 0.973 | 0.973 | 0.041 |
| 418 | P-64-PAL | 55 | M | T3 | 9.81 | 10.31 | 0.040 | 0.959 | 0.038 | 0.963 | 0.963 | 0.038 |
| 419 | P-64-PAL | 55 | M | L1 | 14.90 | 17.82 | 0.046 | 0.975 | 0.043 | 0.978 | 0.978 | 0.043 |
| 420 | P-64-PAL | 55 | M | L2 | 23.43 | 15.02 | 0.046 | 0.992 | 0.041 | 0.993 | 0.993 | 0.041 |
| 421 | P-64-PAL | 55 | M | L3 | 12.10 | 10.82 | 0.052 | 0.946 | 0.052 | 0.947 | 0.947 | 0.052 |
| 422 | P-65-ONG | 56 | M | H1 | 10.57 | 7.76 | 0.033 | 0.973 | 0.028 | 0.980 | 0.980 | 0.028 |
| 423 | P-65-ONG | 56 | M | U3 | 3.44 | 4.71 | 0.037 | 0.929 | 0.037 | 0.929 | 0.929 | 0.037 |
| 424 | P-65-ONG | 56 | M | T1 | 34.01 | 10.31 | 0.052 | 0.987 | 0.054 | 0.983 | 0.987 | 0.052 |
| 425 | P-65-ONG | 56 | M | T2 | 14.65 | 13.11 | 0.024 | 0.959 | 0.029 | 0.944 | 0.959 | 0.024 |
| 426 | P-65-ONG | 56 | M | T3 | 11.84 | 8.91 | 0.018 | 0.952 | 0.018 | 0.953 | 0.953 | 0.018 |
| 427 | P-65-ONG | 56 | M | T4 | 9.42 | 8.02 | 0.029 | 0.931 | 0.029 | 0.931 | 0.931 | 0.029 |
| 428 | P-65-ONG | 56 | M | L1 | 6.50 | 7.64 | 0.035 | 0.944 | 0.033 | 0.950 | 0.950 | 0.033 |
| 429 | P-65-ONG | 56 | M | L2 | 8.53 | 7.13 | 0.053 | 0.941 | 0.053 | 0.942 | 0.942 | 0.053 |
| 430 | P-65-ONG | 56 | M | L3 | 8.41 | 10.95 | 0.058 | 0.906 | 0.056 | 0.907 | 0.907 | 0.056 |
| 431 | P-66-AZR | 49 | M | H1 | 4.71 | 6.49 | 0.035 | 0.965 | 0.034 | 0.966 | 0.966 | 0.034 |
| 432 | P-66-AZR | 49 | M | U1 | 3.57 | 4.46 | 0.029 | 0.759 | 0.027 | 0.907 | 0.907 | 0.027 |
| 433 | P-66-AZR | 49 | M | U2 | 3.82 | 4.58 | 0.027 | 0.955 | 0.026 | 0.959 | 0.959 | 0.026 |
| 434 | P-66-AZR | 49 | M | U3 | 2.93 | 3.18 | 0.027 | 0.939 | 0.025 | 0.950 | 0.950 | 0.025 |
| 435 | P-66-AZR | 49 | M | T1 | 18.72 | 13.75 | 0.050 | 0.952 | 0.049 | 0.953 | 0.953 | 0.049 |
| 436 | P-66-AZR | 49 | M | T2 | 8.41 | 14.13 | 0.045 | 0.929 | 0.045 | 0.933 | 0.933 | 0.045 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 437 | P-66-AZR | 49 | M | T3 | 17.32 | 9.16 | 0.057 | 0.937 | 0.059 | 0.934 | 0.937 | 0.057 |
| 438 | P-66-AZR | 49 | M | L1 | 24.84 | 16.16 | 0.047 | 0.971 | 0.048 | 0.969 | 0.971 | 0.047 |
| 439 | P-66-AZR | 49 | M | L2 | 9.93 | 21.51 | 0.049 | 0.945 | 0.059 | 0.930 | 0.945 | 0.049 |
| 440 | P-66-AZR | 49 | M | L3 | 9.93 | 11.71 | 0.038 | 0.984 | 0.039 | 0.984 | 0.984 | 0.039 |
| 441 | P-67-CHE | 60 | M | U1 | 6.11 | 7.38 | 0.021 | 0.965 | 0.020 | 0.969 | 0.969 | 0.020 |
| 442 | P-67-CHE | 60 | M | U2 | 5.99 | 9.04 | 0.024 | 0.937 | 0.022 | 0.946 | 0.946 | 0.022 |
| 443 | P-67-CHE | 60 | M | U3 | 9.81 | 5.85 | 0.028 | 0.951 | 0.028 | 0.951 | 0.951 | 0.028 |
| 444 | P-67-CHE | 60 | M | T1 | 15.03 | 12.09 | 0.036 | 0.955 | 0.035 | 0.955 | 0.955 | 0.035 |
| 445 | P-67-CHE | 60 | M | L1 | 21.27 | 9.42 | 0.033 | 0.994 | 0.035 | 0.994 | 0.994 | 0.035 |
| 446 | P-67-CHE | 60 | M | L2 | 19.10 | 15.78 | 0.035 | 0.965 | 0.036 | 0.965 | 0.965 | 0.036 |
| 447 | P-67-CHE | 60 | M | L3 | 17.45 | 12.35 | 0.035 | 0.984 | 0.036 | 0.983 | 0.984 | 0.035 |
| 448 | P-68-CHO | 61 | M | U1 | 6.24 | 5.98 | 0.032 | 0.978 | 0.033 | 0.975 | 0.978 | 0.032 |
| 449 | P-68-CHO | 61 | M | U2 | 5.60 | 5.85 | 0.058 | 0.911 | 0.058 | 0.910 | 0.911 | 0.058 |
| 450 | P-68-CHO | 61 | M | U3 | 17.32 | 13.11 | 0.036 | 0.961 | 0.038 | 0.958 | 0.961 | 0.036 |
| 451 | P-68-CHO | 61 | M | T1 | 5.86 | 4.46 | 0.030 | 0.945 | 0.030 | 0.945 | 0.945 | 0.030 |
| 452 | P-68-CHO | 61 | M | T2 | 8.02 | 4.46 | 0.020 | 0.939 | 0.020 | 0.936 | 0.939 | 0.020 |
| 453 | P-68-CHO | 61 | M | T3 | 4.07 | 4.07 | 0.027 | 0.955 | 0.027 | 0.955 | 0.955 | 0.027 |
| 454 | P-68-CHO | 61 | M | L1 | 5.86 | 5.98 | 0.027 | 0.945 | 0.027 | 0.949 | 0.949 | 0.027 |
| 455 | P-68-CHO | 61 | M | L3 | 34.90 | 12.85 | 0.035 | 0.979 | 0.034 | 0.981 | 0.981 | 0.034 |
| 456 | P-69-MAL | 32 | M | H1 | 2.80 | 4.07 | 0.037 | 0.916 | 0.038 | 0.915 | 0.916 | 0.037 |
| 457 | P-69-MAL | 32 | M | U1 | 9.42 | 5.09 | 0.039 | 0.978 | 0.040 | 0.978 | 0.978 | 0.040 |
| 458 | P-69-MAL | 32 | M | U2 | 8.15 | 5.85 | 0.023 | 0.936 | 0.023 | 0.940 | 0.940 | 0.023 |
| 459 | P-69-MAL | 32 | M | U3 | 4.71 | 3.18 | 0.022 | 0.943 | 0.022 | 0.944 | 0.944 | 0.022 |
| 460 | P-69-MAL | 32 | M | T12 | 3.57 | 3.69 | 0.026 | 0.931 | 0.027 | 0.930 | 0.931 | 0.026 |
| 461 | P-69-MAL | 32 | M | T2 | 3.31 | 2.93 | 0.017 | 0.909 | 0.017 | 0.908 | 0.909 | 0.017 |
| 462 | P-69-MAL | 32 | M | T3 | 3.57 | 3.31 | 0.026 | 0.941 | 0.026 | 0.942 | 0.942 | 0.026 |
| 463 | P-69-MAL | 32 | M | 12 | 15.67 | 27.36 | 0.050 | 0.975 | 0.046 | 0.979 | 0.979 | 0.046 |
| 464 | P-69-MAL | 32 | M | L3 | 27.77 | 13.62 | 0.030 | 0.976 | 0.039 | 0.974 | 0.976 | 0.030 |
| 465 | P-70-GAN | 51 | M | H1 | 27.38 | 26.73 | 0.081 | 0.973 | 0.080 | 0.973 | 0.973 | 0.080 |
| 466 | P-70-GAN | 51 | M | U1 | 25.98 | 10.82 | 0.065 | 0.964 | 0.060 | 0.968 | 0.968 | 0.060 |
| 467 | P-70-GAN | 51 | M | U2 | 35.53 | 9.55 | 0.070 | 0.966 | 0.064 | 0.967 | 0.967 | 0.064 |
| 468 | P-70-GAN | 51 | M | T1 | 10.19 | 11.96 | 0.059 | 0.909 | 0.058 | 0.910 | 0.910 | 0.058 |
| 469 | P-70-GAN | 51 | M | T4 | 36.94 | 28.13 | 0.094 | 0.984 | 0.116 | 0.972 | 0.984 | 0.094 |
| 470 | P-70-GAN | 51 | M | T7 | 37.95 | 28.13 | 0.125 | 0.942 | 0.131 | 0.940 | 0.942 | 0.125 |
| 471 | P-70-GAN | 51 | M | L1 | 19.10 | 26.35 | 0.074 | 0.955 | 0.071 | 0.956 | 0.956 | 0.071 |
| 472 | P-70-GAN | 51 | M | L2 | 19.49 | 16.80 | 0.090 | 0.972 | 0.082 | 0.977 | 0.977 | 0.082 |
| 473 | P-70-GAN | 51 | M | L3 | 31.97 | 9.67 | 0.051 | 0.985 | 0.049 | 0.987 | 0.987 | 0.049 |
| 474 | P-71-BUR | 34 | M | H1 | 15.41 | 15.27 | 0.088 | 0.928 | 0.078 | 0.944 | 0.944 | 0.078 |
| 475 | P-71-BUR | 34 | M | U1 | 16.81 | 18.58 | 0.044 | 0.988 | 0.044 | 0.988 | 0.988 | 0.044 |
| 476 | P-71-BUR | 34 | M | U3 | 16.81 | 11.71 | 0.029 | 0.981 | 0.028 | 0.983 | 0.983 | 0.028 |
| 477 | P-71-BUR | 34 | M | T1 | 16.05 | 13.49 | 0.043 | 0.964 | 0.045 | 0.963 | 0.964 | 0.043 |
| 478 | P-71-BUR | 34 | M | T3 | 12.10 | 15.65 | 0.050 | 0.956 | 0.050 | 0.956 | 0.956 | 0.050 |
| 479 | P-71-BUR | 34 | M | T4 | 17.07 | 20.49 | 0.044 | 0.932 | 0.043 | 0.936 | 0.936 | 0.043 |
| 480 | P-71-BUR | 34 | M | T5 | 10.70 | 11.58 | 0.036 | 0.929 | 0.035 | 0.931 | 0.931 | 0.035 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 481 | P-71-BUR | 34 | M | T7 | 7.90 | 11.20 | 0.023 | 0.968 | 0.024 | 0.943 | 0.968 | 0.023 |
| 482 | P-71-BUR | 34 | M | L1 | 16.17 | 13.11 | 0.035 | 0.966 | 0.032 | 0.969 | 0.969 | 0.032 |
| 483 | P-71-BUR | 34 | M | L2 | 15.54 | 15.02 | 0.035 | 0.990 | 0.033 | 0.991 | 0.991 | 0.033 |
| 484 | P-72-CHE | 47 | M | H1 | 3.57 | 5.98 | 0.025 | 0.923 | 0.025 | 0.922 | 0.923 | 0.025 |
| 485 | P-72-CHE | 47 | M | U1 | 12.99 | 17.82 | 0.051 | 0.954 | 0.048 | 0.959 | 0.959 | 0.048 |
| 486 | P-72-CHE | 47 | M | U2 | 11.59 | 15.40 | 0.033 | 0.945 | 0.031 | 0.953 | 0.953 | 0.031 |
| 487 | P-72-CHE | 47 | M | U3 | 20.76 | 15.27 | 0.064 | 0.982 | 0.071 | 0.980 | 0.982 | 0.064 |
| 488 | P-72-CHE | 47 | M | T1 | 20.38 | 7.25 | 0.087 | 0.982 | 0.088 | 0.981 | 0.982 | 0.087 |
| 489 | P-72-CHE | 47 | M | T2 | 18.34 | 16.55 | 0.053 | 0.953 | 0.051 | 0.957 | 0.957 | 0.051 |
| 490 | P-72-CHE | 47 | M | T3 | 19.36 | 29.53 | 0.084 | 0.933 | 0.083 | 0.933 | 0.933 | 0.083 |
| 491 | P-72-CHE | 47 | M | T4 | 19.36 | 18.20 | 0.052 | 0.932 | 0.055 | 0.919 | 0.932 | 0.052 |
| 492 | P-72-CHE | 47 | M | T6 | 37.83 | 28.89 | 0.077 | 0.985 | 0.091 | 0.976 | 0.985 | 0.077 |
| 493 | P-72-CHE | 47 | M | L1 | 18.34 | 16.55 | 0.028 | 0.957 | 0.028 | 0.963 | 0.963 | 0.028 |
| 494 | P-72-CHE | 47 | M | L2 | 11.84 | 12.98 | 0.022 | 0.963 | 0.021 | 0.988 | 0.988 | 0.021 |
| 495 | P-72-CHE | 47 | M | L3 | 23.05 | 17.82 | 0.049 | 0.968 | 0.045 | 0.973 | 0.973 | 0.045 |
| 496 | P-72-CHE | 47 | M | L4 | 12.23 | 10.05 | 0.055 | 0.930 | 0.060 | 0.920 | 0.930 | 0.055 |
| 497 | P-73-NUR | 26 | F | U1 | 5.48 | 4.71 | 0.032 | 0.962 | 0.031 | 0.964 | 0.964 | 0.031 |
| 498 | P-73-NUR | 26 | F | U2 | 7.01 | 6.11 | 0.034 | 0.920 | 0.033 | 0.924 | 0.924 | 0.033 |
| 499 | P-73-NUR | 26 | F | U3 | 8.02 | 16.42 | 0.039 | 0.978 | 0.041 | 0.977 | 0.978 | 0.039 |
| 500 | P-73-NUR | 26 | F | T1 | 11.21 | 11.33 | 0.043 | 0.962 | 0.041 | 0.964 | 0.964 | 0.041 |
| 501 | P-73-NUR | 26 | F | T2 | 9.68 | 9.29 | 0.040 | 0.959 | 0.041 | 0.952 | 0.959 | 0.040 |
| 502 | P-73-NUR | 26 | F | T3 | 6.62 | 9.67 | 0.042 | 0.953 | 0.041 | 0.955 | 0.955 | 0.041 |
| 503 | P-73-NUR | 26 | F | L1 | 6.62 | 5.98 | 0.037 | 0.928 | 0.037 | 0.926 | 0.928 | 0.037 |
| 504 | P-73-NUR | 26 | F | L2 | 5.35 | 5.09 | 0.036 | 0.930 | 0.036 | 0.930 | 0.930 | 0.036 |
| 505 | P-95-MOH | 53 | M | H1 | 5.22 | 10.18 | 0.051 | 0.917 | 0.052 | 0.917 | 0.917 | 0.052 |
| 506 | P-95-MOH | 53 | M | U1 | 23.82 | 13.49 | 0.051 | 0.968 | 0.051 | 0.966 | 0.968 | 0.051 |
| 507 | P-95-MOH | 53 | M | U2 | 11.33 | 8.65 | 0.025 | 0.954 | 0.025 | 0.954 | 0.954 | 0.025 |
| 508 | P-95-MOH | 53 | M | U3 | 9.81 | 13.11 | 0.048 | 0.950 | 0.048 | 0.949 | 0.950 | 0.048 |
| 509 | P-95-MOH | 53 | M | T1 | 18.72 | 21.38 | 0.036 | 0.972 | 0.039 | 0.966 | 0.972 | 0.036 |
| 510 | P-95-MOH | 53 | M | T2 | 30.06 | 20.24 | 0.038 | 0.983 | 0.050 | 0.969 | 0.983 | 0.038 |
| 511 | P-96-ZAI | 62 | M | H1 | 7.00 | 10.82 | 0.028 | 0.966 | 0.029 | 0.963 | 0.966 | 0.028 |
| 512 | P-96-ZAI | 62 | M | U1 | 17.07 | 23.80 | 0.067 | 0.950 | 0.064 | 0.951 | 0.951 | 0.064 |
| 513 | P-96-ZAI | 62 | M | U2 | 14.14 | 13.87 | 0.049 | 0.971 | 0.050 | 0.957 | 0.971 | 0.049 |
| 514 | P-96-ZAI | 62 | M | T2 | 6.11 | 8.27 | 0.032 | 0.936 | 0.031 | 0.937 | 0.937 | 0.031 |
| 515 | P-96-ZAI | 62 | M | T3 | 15.41 | 10.82 | 0.054 | 0.946 | 0.040 | 0.970 | 0.970 | 0.040 |
| 516 | P-96-ZAI | 62 | M | T4 | 30.18 | 26.34 | 0.155 | 0.932 | 0.141 | 0.921 | 0.932 | 0.155 |
| 517 | P-96-ZAI | 62 | M | T5 | 35.02 | 28.64 | 0.123 | 0.973 | 0.139 | 0.964 | 0.973 | 0.123 |
| 518 | P-96-ZAI | 62 | M | T6 | 39.61 | 24.31 | 0.186 | 0.941 | 0.178 | 0.943 | 0.943 | 0.178 |
| 519 | P-96-ZAI | 62 | M | L1 | 32.48 | 6.49 | 0.069 | 0.964 | 0.064 | 0.969 | 0.969 | 0.064 |
| 520 | P-96-ZAI | 62 | M | L2 | 8.53 | 21.51 | 0.063 | 0.968 | 0.060 | 0.971 | 0.971 | 0.060 |
| 521 | P-96-ZAI | 62 | M | L4 | 5.60 | 14.64 | 0.044 | 0.984 | 0.044 | 0.983 | 0.984 | 0.044 |
| 522 | P-96-ZAI | 62 | M | L5 | 11.84 | 17.31 | 0.070 | 0.952 | 0.071 | 0.951 | 0.952 | 0.070 |
| 523 | P-97-SAT | 54 | M | U1 | 8.53 | 11.45 | 0.022 | 0.970 | 0.021 | 0.974 | 0.974 | 0.021 |
| 524 | P-97-SAT | 54 | M | U2 | 16.81 | 23.67 | 0.034 | 0.984 | 0.041 | 0.978 | 0.984 | 0.034 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 525 | P-97-SAT | 54 | M | U3 | 8.02 | 14.25 | 0.023 | 0.984 | 0.024 | 0.983 | 0.984 | 0.023 |
| 526 | P-97-SAT | 54 | M | T1 | 6.62 | 13.49 | 0.021 | 0.950 | 0.023 | 0.941 | 0.950 | 0.021 |
| 527 | P-97-SAT | 54 | M | T2 | 7.51 | 6.24 | 0.027 | 0.940 | 0.027 | 0.940 | 0.940 | 0.027 |
| 528 | P-97-SAT | 54 | M | L1 | 16.18 | 23.04 | 0.050 | 0.933 | 0.049 | 0.935 | 0.935 | 0.049 |
| 529 | P-97-SAT | 54 | M | L2 | 14.90 | 12.73 | 0.032 | 0.961 | 0.031 | 0.962 | 0.962 | 0.031 |
| 530 | P-98-MOH | 35 | M | H1 | 7.39 | 7.76 | 0.044 | 0.963 | 0.044 | 0.963 | 0.963 | 0.044 |
| 531 | P-98-MOH | 35 | M | U1 | 7.13 | 7.00 | 0.031 | 0.974 | 0.031 | 0.974 | 0.974 | 0.031 |
| 532 | P-98-MOH | 35 | M | U2 | 8.15 | 10.56 | 0.039 | 0.931 | 0.037 | 0.937 | 0.937 | 0.037 |
| 533 | P-98-MOH | 35 | M | U3 | 7.13 | 5.85 | 0.060 | 0.982 | 0.060 | 0.981 | 0.982 | 0.060 |
| 534 | P-98-MOH | 35 | M | T1 | 3.82 | 4.58 | 0.022 | 0.946 | 0.023 | 0.944 | 0.946 | 0.022 |
| 535 | P-98-MOH | 35 | M | T2 | 3.18 | 4.07 | 0.019 | 0.938 | 0.019 | 0.938 | 0.938 | 0.019 |
| 536 | P-98-MOH | 35 | M | T3 | 3.06 | 3.44 | 0.018 | 0.917 | 0.018 | 0.915 | 0.917 | 0.018 |
| 537 | P-98-MOH | 35 | M | L1 | 2.93 | 2.80 | 0.023 | 0.953 | 0.022 | 0.958 | 0.958 | 0.022 |
| 538 | P-99-ROH | 39 | M | H1 | 8.28 | 8.53 | 0.024 | 0.964 | 0.026 | 0.958 | 0.964 | 0.024 |
| 539 | P-99-ROH | 39 | M | U1 | 15.54 | 13.11 | 0.036 | 0.980 | 0.035 | 0.982 | 0.982 | 0.035 |
| 540 | P-99-ROH | 39 | M | U2 | 9.17 | 7.13 | 0.036 | 0.988 | 0.037 | 0.988 | 0.988 | 0.037 |
| 541 | P-99-ROH | 39 | M | U3 | 11.72 | 11.84 | 0.022 | 0.989 | 0.022 | 0.989 | 0.989 | 0.022 |
| 542 | P-99-ROH | 39 | M | T3 | 6.75 | 4.07 | 0.024 | 0.949 | 0.025 | 0.944 | 0.949 | 0.024 |
| 543 | P-99-ROH | 39 | M | L2 | 11.34 | 8.53 | 0.032 | 0.955 | 0.031 | 0.956 | 0.956 | 0.031 |
| 544 | P-99-ROH | 39 | M | L3 | 7.77 | 8.15 | 0.028 | 0.962 | 0.028 | 0.962 | 0.962 | 0.028 |
| 545 | P-100-MAT | 55 | M | H1 | 10.19 | 8.40 | 0.048 | 0.939 | 0.048 | 0.940 | 0.940 | 0.048 |
| 546 | P-100-MAT | 55 | M | U1 | 16.56 | 25.07 | 0.079 | 0.965 | 0.075 | 0.969 | 0.969 | 0.075 |
| 547 | P-100-MAT | 55 | M | U3 | 11.59 | 18.33 | 0.035 | 0.986 | 0.036 | 0.986 | 0.986 | 0.036 |
| 548 | P-100-MAT | 55 | M | T1 | 31.33 | 7.76 | 0.058 | 0.927 | 0.056 | 0.932 | 0.932 | 0.056 |
| 549 | P-100-MAT | 55 | M | T2 | 27.51 | 11.71 | 0.057 | 0.963 | 0.065 | 0.952 | 0.963 | 0.057 |
| 550 | P-100-MAT | 55 | M | T4 | 36.68 | 28.13 | 0.105 | 0.964 | 0.125 | 0.950 | 0.964 | 0.105 |
| 551 | P-100-MAT | 55 | M | T5 | 9.43 | 11.20 | 0.037 | 0.952 | 0.033 | 0.961 | 0.961 | 0.033 |
| 552 | P-100-MAT | 55 | M | L1 | 13.88 | 27.24 | 0.073 | 0.971 | 0.069 | 0.973 | 0.973 | 0.069 |
| 553 | P-100-MAT | 55 | M | L2 | 24.45 | 16.42 | 0.026 | 0.987 | 0.034 | 0.977 | 0.987 | 0.026 |
| 554 | P-100-MAT | 55 | M | L3 | 11.72 | 27.87 | 0.037 | 0.970 | 0.037 | 0.970 | 0.970 | 0.037 |
| 555 | P-101-NAN | 31 | M | H1 | 11.33 | 9.67 | 0.042 | 0.950 | 0.040 | 0.956 | 0.956 | 0.040 |
| 556 | P-101-NAN | 31 | M | U1 | 25.98 | 11.33 | 0.059 | 0.945 | 0.057 | 0.948 | 0.948 | 0.057 |
| 557 | P-101-NAN | 31 | M | U3 | 15.54 | 14.00 | 0.038 | 0.974 | 0.038 | 0.975 | 0.975 | 0.038 |
| 558 | P-101-NAN | 31 | M | T2 | 16.43 | 21.13 | 0.043 | 0.939 | 0.043 | 0.941 | 0.941 | 0.043 |
| 559 | P-101-NAN | 31 | M | T3 | 27.51 | 11.58 | 0.054 | 0.990 | 0.054 | 0.990 | 0.990 | 0.054 |
| 560 | P-101-NAN | 31 | M | L1 | 25.22 | 13.49 | 0.047 | 0.988 | 0.047 | 0.988 | 0.988 | 0.047 |
| 561 | P-101-NAN | 31 | M | L2 | 15.67 | 20.11 | 0.057 | 0.983 | 0.056 | 0.984 | 0.984 | 0.056 |
| 562 | P-102-MUH | 56 | M | U1 | 13.88 | 13.75 | 0.042 | 0.970 | 0.039 | 0.975 | 0.975 | 0.039 |
| 563 | P-102-MUH | 56 | M | U2 | 15.79 | 8.78 | 0.053 | 0.927 | 0.051 | 0.927 | 0.927 | 0.051 |
| 564 | P-102-MUH | 56 | M | U3 | 11.46 | 7.64 | 0.028 | 0.984 | 0.029 | 0.984 | 0.984 | 0.029 |
| 565 | P-102-MUH | 56 | M | T1 | 12.35 | 13.75 | 0.043 | 0.930 | 0.042 | 0.930 | 0.930 | 0.042 |
| 566 | P-102-MUH | 56 | M | 9T2 | 10.06 | 8.53 | 0.048 | 0.981 | 0.048 | 0.982 | 0.982 | 0.048 |
| 567 | P-102-MUH | 56 | M | T3 | 15.54 | 18.07 | 0.083 | 0.964 | 0.083 | 0.966 | 0.966 | 0.083 |
| 568 | P-102-MUH | 56 | M | T4 | 36.17 | 28.76 | 0.037 | 0.994 | 0.044 | 0.990 | 0.994 | 0.037 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 569 | P-102-MUH | 56 | M | L1A | 15.54 | 15.53 | 0.049 | 0.988 | 0.053 | 0.985 | 0.988 | 0.049 |
| 570 | P-102-MUH | 56 | M | L2 | 13.76 | 8.27 | 0.038 | 0.962 | 0.036 | 0.967 | 0.967 | 0.036 |
| 571 | P-102-MUH | 56 | M | L3 | 9.30 | 7.13 | 0.032 | 0.951 | 0.031 | 0.955 | 0.955 | 0.031 |
| 572 | P-103-TAN | 56 | M | H1 | 7.90 | 7.51 | 0.040 | 0.936 | 0.040 | 0.936 | 0.936 | 0.040 |
| 573 | P-103-TAN | 56 | M | U2 | 7.39 | 8.53 | 0.036 | 0.954 | 0.036 | 0.955 | 0.955 | 0.036 |
| 574 | P-103-TAN | 56 | M | U3 | 17.70 | 26.09 | 0.053 | 0.968 | 0.048 | 0.954 | 0.968 | 0.053 |
| 575 | P-103-TAN | 56 | M | T1 | 12.48 | 16.80 | 0.037 | 0.845 | 0.038 | 0.906 | 0.906 | 0.038 |
| 576 | P-103-TAN | 56 | M | T2 | 25.09 | 19.47 | 0.043 | 0.957 | 0.054 | 0.954 | 0.957 | 0.043 |
| 577 | P-103-TAN | 56 | M | L1 | 14.26 | 16.55 | 0.033 | 0.982 | 0.038 | 0.978 | 0.982 | 0.033 |
| 578 | P-103-TAN | 56 | M | L3 | 13.12 | 12.98 | 0.034 | 0.962 | 0.035 | 0.962 | 0.962 | 0.035 |
| 579 | P-104-AHM | 21 | M | H1 | 5.86 | 5.98 | 0.021 | 0.979 | 0.021 | 0.978 | 0.979 | 0.021 |
| 580 | P-104-AHM | 21 | M | U1 | 3.31 | 5.09 | 0.030 | 0.943 | 0.029 | 0.946 | 0.946 | 0.029 |
| 581 | P-104-AHM | 21 | M | U2 | 3.69 | 4.71 | 0.025 | 0.938 | 0.024 | 0.941 | 0.941 | 0.024 |
| 582 | P-104-AHM | 21 | M | U3 | 2.17 | 2.42 | 0.015 | 0.928 | 0.015 | 0.927 | 0.928 | 0.015 |
| 583 | P-104-AHM | 21 | M | T1 | 10.95 | 9.93 | 0.044 | 0.973 | 0.046 | 0.972 | 0.973 | 0.044 |
| 584 | P-104-AHM | 21 | M | T2 | 7.77 | 8.65 | 0.030 | 0.935 | 0.029 | 0.937 | 0.937 | 0.029 |
| 585 | P-104-AHM | 21 | M | T3 | 9.68 | 5.60 | 0.063 | 0.945 | 0.064 | 0.941 | 0.945 | 0.063 |
| 586 | P-104-AHM | 21 | M | L1 | 5.10 | 5.60 | 0.025 | 0.945 | 0.025 | 0.945 | 0.945 | 0.025 |
| 587 | P-104-AHM | 21 | M | L2 | 3.82 | 3.44 | 0.034 | 0.920 | 0.034 | 0.921 | 0.921 | 0.034 |
| 588 | P-105-SHA | 30 | M | T1 | 34.39 | 10.82 | 0.065 | 0.955 | 0.066 | 0.953 | 0.955 | 0.065 |
| 589 | P-105-SHA | 30 | M | T2 | 22.67 | 13.36 | 0.050 | 0.957 | 0.049 | 0.958 | 0.958 | 0.049 |
| 590 | P-105-SHA | 30 | M | T3 | 12.86 | 8.78 | 0.034 | 0.926 | 0.034 | 0.928 | 0.928 | 0.034 |
| 591 | P-105-SHA | 30 | M | L1 | 31.33 | 21.00 | 0.053 | 0.992 | 0.058 | 0.990 | 0.992 | 0.053 |
| 592 | P-105-SHA | 30 | M | L1B | 23.94 | 15.65 | 0.045 | 0.985 | 0.046 | 0.985 | 0.985 | 0.046 |
| 593 | P-105-SHA | 30 | M | L2 | 11.46 | 17.31 | 0.060 | 0.956 | 0.061 | 0.956 | 0.956 | 0.061 |
| 594 | P-105-SHA | 30 | M | L3 | 11.84 | 7.64 | 0.054 | 0.917 | 0.052 | 0.922 | 0.922 | 0.052 |
| 595 | P-106-HUS | 38 | M | H1 | 3.82 | 4.58 | 0.026 | 0.976 | 0.026 | 0.976 | 0.976 | 0.026 |
| 596 | P-106-HUS | 38 | M | U1 | 6.62 | 11.45 | 0.038 | 0.958 | 0.033 | 0.968 | 0.968 | 0.033 |
| 597 | P-106-HUS | 38 | M | U2 | 3.57 | 3.56 | 0.016 | 0.944 | 0.016 | 0.944 | 0.944 | 0.016 |
| 598 | P-106-HUS | 38 | M | U3 | 5.99 | 7.64 | 0.021 | 0.993 | 0.019 | 0.967 | 0.993 | 0.021 |
| 599 | P-106-HUS | 38 | M | T1 | 6.88 | 8.27 | 0.016 | 0.991 | 0.016 | 0.991 | 0.991 | 0.016 |
| 600 | P-106-HUS | 38 | M | T2 | 7.13 | 10.56 | 0.022 | 0.955 | 0.019 | 0.967 | 0.967 | 0.019 |
| 601 | P-106-HUS | 38 | M | T3 | 9.17 | 4.45 | 0.019 | 0.976 | 0.019 | 0.976 | 0.976 | 0.019 |
| 602 | P-106-HUS | 38 | M | T4 | 4.71 | 4.20 | 0.021 | 0.989 | 0.021 | 0.989 | 0.989 | 0.021 |
| 603 | P-106-HUS | 38 | M | T5 | 3.31 | 5.09 | 0.016 | 0.943 | 0.016 | 0.943 | 0.943 | 0.016 |
| 604 | P-106-HUS | 38 | M | L1 | 7.90 | 13.75 | 0.024 | 0.938 | 0.021 | 0.945 | 0.945 | 0.021 |
| 605 | P-106-HUS | 38 | M | L2 | 8.92 | 12.47 | 0.030 | 0.983 | 0.030 | 0.984 | 0.984 | 0.030 |
| 606 | P-106-HUS | 38 | M | L3 | 6.75 | 10.94 | 0.022 | 0.972 | 0.020 | 0.976 | 0.976 | 0.020 |
| 607 | P-107-ABD | 54 | M | U1 | 8.15 | 8.78 | 0.040 | 0.962 | 0.040 | 0.961 | 0.962 | 0.040 |
| 608 | P-107-ABD | 54 | M | U2 | 13.25 | 9.93 | 0.028 | 0.960 | 0.028 | 0.959 | 0.960 | 0.028 |
| 609 | P-107-ABD | 54 | M | T1 | 10.19 | 7.89 | 0.038 | 0.918 | 0.038 | 0.919 | 0.919 | 0.038 |
| 610 | P-107-ABD | 54 | M | T2 | 3.57 | 4.71 | 0.029 | 0.936 | 0.028 | 0.937 | 0.937 | 0.028 |
| 611 | P-107-ABD | 54 | M | T3 | 5.99 | 6.24 | 0.032 | 0.945 | 0.032 | 0.945 | 0.945 | 0.032 |
| 612 | P-107-ABD | 54 | M | L1 | 8.41 | 9.93 | 0.043 | 0.934 | 0.043 | 0.933 | 0.934 | 0.043 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 613 | P-108-RAH | 56 | M | H1 | 9.30 | 11.58 | 0.067 | 0.938 | 0.063 | 0.942 | 0.942 | 0.063 |
| 614 | P-108-RAH | 56 | M | U1 | 13.88 | 10.82 | 0.041 | 0.942 | 0.038 | 0.928 | 0.942 | 0.041 |
| 615 | P-108-RAH | 56 | M | U2 | 8.02 | 8.91 | 0.043 | 0.916 | 0.042 | 0.931 | 0.931 | 0.042 |
| 616 | P-108-RAH | 56 | M | U3 | 5.22 | 5.85 | 0.020 | 0.943 | 0.020 | 0.943 | 0.943 | 0.020 |
| 617 | P-108-RAH | 56 | M | T1 | 24.45 | 19.60 | 0.050 | 0.973 | 0.049 | 0.973 | 0.973 | 0.049 |
| 618 | P-108-RAH | 56 | M | T2 | 25.34 | 15.40 | 0.074 | 0.978 | 0.074 | 0.978 | 0.978 | 0.074 |
| 619 | P-108-RAH | 56 | M | L1 | 18.98 | 23.29 | 0.059 | 0.983 | 0.059 | 0.983 | 0.983 | 0.059 |
| 620 | P-108-RAH | 56 | M | L2 | 23.69 | 26.60 | 0.039 | 0.996 | 0.038 | 0.996 | 0.996 | 0.038 |
| 621 | P-108-RAH | 56 | M | L3 | 29.55 | 24.95 | 0.046 | 0.996 | 0.046 | 0.996 | 0.996 | 0.046 |
| 622 | P-109-BAF | 60 | M | H1 | 7.77 | 8.66 | 0.051 | 0.946 | 0.051 | 0.946 | 0.946 | 0.051 |
| 623 | P-109-BAF | 60 | M | U3 | 18.98 | 28.51 | 0.043 | 0.974 | 0.063 | 0.952 | 0.974 | 0.043 |
| 624 | P-109-BAF | 60 | M | T1 | 27.00 | 16.80 | 0.052 | 0.949 | 0.074 | 0.928 | 0.949 | 0.052 |
| 625 | P-109-BAF | 60 | M | T2 | 24.33 | 8.78 | 0.047 | 0.955 | 0.045 | 0.959 | 0.959 | 0.045 |
| 626 | P-109-BAF | 60 | M | T3 | 8.41 | 11.96 | 0.038 | 0.956 | 0.038 | 0.956 | 0.956 | 0.038 |
| 627 | P-109-BAF | 60 | M | T4 | 25.73 | 13.75 | 0.064 | 0.961 | 0.058 | 0.967 | 0.967 | 0.058 |
| 628 | P-109-BAF | 60 | M | L1 | 18.47 | 20.62 | 0.063 | 0.969 | 0.066 | 0.968 | 0.969 | 0.063 |
| 629 | P-109-BAF | 60 | M | L2 | 23.94 | 12.35 | 0.045 | 0.951 | 0.035 | 0.968 | 0.968 | 0.035 |
| 630 | P-109-BAF | 60 | M | L3 | 7.90 | 6.87 | 0.026 | 0.962 | 0.025 | 0.962 | 0.962 | 0.025 |
| 631 | P-110-TAN | 38 | M | U1 | 16.05 | 14.00 | 0.046 | 0.977 | 0.041 | 0.980 | 0.980 | 0.041 |
| 632 | P-110-TAN | 38 | M | U3 | 12.23 | 11.96 | 0.047 | 0.966 | 0.046 | 0.967 | 0.967 | 0.046 |
| 633 | P-110-TAN | 38 | M | T2 | 9.55 | 16.16 | 0.038 | 0.959 | 0.038 | 0.959 | 0.959 | 0.038 |
| 634 | P-110-TAN | 38 | M | T3 | 19.61 | 19.60 | 0.091 | 0.954 | 0.089 | 0.957 | 0.957 | 0.089 |
| 635 | P-110-TAN | 38 | M | T5 | 36.94 | 28.76 | 0.124 | 0.957 | 0.156 | 0.930 | 0.957 | 0.124 |
| 636 | P-110-TAN | 38 | M | L1 | 24.07 | 17.56 | 0.082 | 0.946 | 0.068 | 0.948 | 0.948 | 0.068 |
| 637 | P-110-TAN | 38 | M | L2 | 15.16 | 29.02 | 0.077 | 0.978 | 0.072 | 0.982 | 0.982 | 0.072 |
| 638 | P-110-TAN | 38 | M | L3 | 24.58 | 26.73 | 0.067 | 0.961 | 0.061 | 0.963 | 0.963 | 0.061 |
| 639 | P-110-TAN | 38 | M | L4 | 38.34 | 29.02 | 0.101 | 0.962 | 0.126 | 0.990 | 0.990 | 0.126 |
| 640 | P-110-TAN | 38 | M | L5 | 38.72 | 28.25 | 0.095 | 0.971 | 0.089 | 0.973 | 0.973 | 0.089 |
| 641 | P-111-ONG | 64 | M | H1 | 10.95 | 10.56 | 0.035 | 0.936 | 0.032 | 0.943 | 0.943 | 0.032 |
| 642 | P-111-ONG | 64 | M | U1 | 7.39 | 10.18 | 0.045 | 0.962 | 0.044 | 0.963 | 0.963 | 0.044 |
| 643 | P-111-ONG | 64 | M | U2 | 6.49 | 7.76 | 0.035 | 0.971 | 0.033 | 0.973 | 0.973 | 0.033 |
| 644 | P-111-ONG | 64 | M | U3 | 7.13 | 7.13 | 0.049 | 0.932 | 0.048 | 0.932 | 0.932 | 0.048 |
| 645 | P-111-ONG | 64 | M | T3 | 5.60 | 7.38 | 0.031 | 0.962 | 0.030 | 0.963 | 0.963 | 0.030 |
| 646 | P-111-ONG | 64 | M | L1 | 30.06 | 16.16 | 0.067 | 0.941 | 0.064 | 0.943 | 0.943 | 0.064 |
| 647 | P-111-ONG | 64 | M | L1B | 20.63 | 18.71 | 0.037 | 0.983 | 0.035 | 0.985 | 0.985 | 0.035 |
| 648 | P-111-ONG | 64 | M | L2 | 17.19 | 17.82 | 0.053 | 0.979 | 0.051 | 0.976 | 0.979 | 0.053 |
| 649 | P-111-ONG | 64 | M | L3 | 22.03 | 19.09 | 0.044 | 0.967 | 0.039 | 0.974 | 0.974 | 0.039 |
| 650 | P-112-SEL | 42 | M | U3 | 16.68 | 17.44 | 0.067 | 0.969 | 0.068 | 0.968 | 0.969 | 0.067 |
| 651 | P-112-SEL | 42 | M | T1 | 10.70 | 20.74 | 0.039 | 0.988 | 0.034 | 0.967 | 0.988 | 0.039 |
| 652 | P-112-SEL | 42 | M | T2 | 20.50 | 13.62 | 0.049 | 0.940 | 0.046 | 0.953 | 0.953 | 0.046 |
| 653 | P-112-SEL | 42 | M | T3 | 29.55 | 23.04 | 0.037 | 0.992 | 0.043 | 0.981 | 0.992 | 0.037 |
| 654 | P-112-SEL | 42 | M | L1 | 23.82 | 28.64 | 0.049 | 0.981 | 0.070 | 0.962 | 0.981 | 0.049 |
| 655 | P-112-SEL | 42 | M | L2 | 20.89 | 16.93 | 0.062 | 0.985 | 0.064 | 0.984 | 0.985 | 0.062 |
| 656 | P-112-SEL | 42 | M | L3 | 30.57 | 13.24 | 0.059 | 0.968 | 0.051 | 0.982 | 0.982 | 0.051 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 657 | P-113-NOR | 21 | F | H1 | 5.22 | 5.85 | 0.037 | 0.929 | 0.037 | 0.930 | 0.930 | 0.037 |
| 658 | P-113-NOR | 21 | F | U1 | 8.41 | 6.24 | 0.030 | 0.940 | 0.032 | 0.931 | 0.940 | 0.030 |
| 659 | P-113-NOR | 21 | F | U2 | 3.44 | 3.31 | 0.025 | 0.924 | 0.025 | 0.924 | 0.924 | 0.025 |
| 660 | P-113-NOR | 21 | F | U3 | 3.18 | 2.55 | 0.023 | 0.915 | 0.022 | 0.916 | 0.916 | 0.022 |
| 661 | P-113-NOR | 21 | F | T3 | 7.00 | 7.00 | 0.032 | 0.919 | 0.032 | 0.918 | 0.919 | 0.032 |
| 662 | P-113-NOR | 21 | F | L1 | 11.59 | 8.40 | 0.034 | 0.952 | 0.029 | 0.962 | 0.962 | 0.029 |
| 663 | P-113-NOR | 21 | F | L2 | 3.95 | 3.69 | 0.018 | 0.929 | 0.021 | 0.916 | 0.929 | 0.018 |
| 664 | P-113-NOR | 21 | F | L3 | 5.09 | 4.58 | 0.022 | 0.930 | 0.021 | 0.937 | 0.937 | 0.021 |
| 665 | P-114-MOH | 29 | M | H1 | 5.48 | 6.24 | 0.038 | 0.927 | 0.037 | 0.929 | 0.929 | 0.037 |
| 666 | P-114-MOH | 29 | M | U1 | 5.99 | 11.71 | 0.037 | 0.942 | 0.030 | 0.946 | 0.946 | 0.030 |
| 667 | $\mathrm{P}-114-\mathrm{MOH}$ | 29 | M | U2 | 7.39 | 7.76 | 0.031 | 0.971 | 0.028 | 0.974 | 0.974 | 0.028 |
| 668 | P-114-MOH | 29 | M | T1 | 31.08 | 15.02 | 0.064 | 0.907 | 0.062 | 0.911 | 0.911 | 0.062 |
| 669 | $\mathrm{P}-114-\mathrm{MOH}$ | 29 | M | T2 | 29.17 | 17.18 | 0.075 | 0.970 | 0.065 | 0.960 | 0.970 | 0.075 |
| 670 | $\mathrm{P}-114-\mathrm{MOH}$ | 29 | M | T3 | 25.35 | 12.86 | 0.086 | 0.969 | 0.081 | 0.973 | 0.973 | 0.081 |
| 671 | $\mathrm{P}-114-\mathrm{MOH}$ | 29 | M | T4 | 38.21 | 28.89 | 0.088 | 0.978 | 0.126 | 0.950 | 0.978 | 0.088 |
| 672 | $\mathrm{P}-114-\mathrm{MOH}$ | 29 | M | L1 | 13.37 | 17.56 | 0.046 | 0.973 | 0.045 | 0.955 | 0.973 | 0.046 |
| 673 | $\mathrm{P}-114-\mathrm{MOH}$ | 29 | M | L2 | 30.31 | 12.60 | 0.061 | 0.973 | 0.054 | 0.977 | 0.977 | 0.054 |
| 674 | P-114-MOH | 29 | M | L3 | 9.55 | 9.93 | 0.042 | 0.948 | 0.041 | 0.949 | 0.949 | 0.041 |
| 675 | P-114-MOH | 29 | M | L4 | 9.04 | 11.20 | 0.098 | 0.929 | 0.093 | 0.937 | 0.937 | 0.093 |
| 676 | P-114-MOH | 29 | M | L5 | 38.34 | 28.76 | 0.088 | 0.948 | 0.088 | 0.949 | 0.949 | 0.088 |
| 677 | P-115-RAJ | 32 | M | U1 | 37.57 | 13.36 | 0.090 | 0.923 | 0.067 | 0.931 | 0.931 | 0.067 |
| 678 | P-115-RAJ | 32 | M | U2 | 39.61 | 10.56 | 0.070 | 0.951 | 0.070 | 0.950 | 0.951 | 0.070 |
| 679 | P-115-RAJ | 32 | M | U3 | 22.29 | 22.15 | 0.054 | 0.971 | 0.053 | 0.972 | 0.972 | 0.053 |
| 680 | P-115-RAJ | 32 | M | T1 | 19.87 | 18.20 | 0.078 | 0.966 | 0.078 | 0.967 | 0.967 | 0.078 |
| 681 | P-115-RAJ | 32 | M | T2 | 28.02 | 14.76 | 0.064 | 0.964 | 0.067 | 0.961 | 0.964 | 0.064 |
| 682 | P-115-RAJ | 32 | M | T3 | 27.13 | 19.73 | 0.068 | 0.955 | 0.066 | 0.958 | 0.958 | 0.066 |
| 683 | P-115-RAJ | 32 | M | T4 | 36.17 | 26.73 | 0.051 | 0.932 | 0.051 | 0.932 | 0.932 | 0.051 |
| 684 | P-115-RAJ | 32 | M | L1 | 17.19 | 18.33 | 0.047 | 0.951 | 0.045 | 0.954 | 0.954 | 0.045 |
| 685 | P-115-RAJ | 32 | M | L2 | 22.67 | 24.56 | 0.051 | 0.970 | 0.050 | 0.971 | 0.971 | 0.050 |
| 686 | P-115-RAJ | 32 | M | L3 | 15.54 | 28.38 | 0.052 | 0.987 | 0.051 | 0.988 | 0.988 | 0.051 |
| 687 | P-116-SUP | 54 | M | H1 | 21.91 | 26.09 | 0.045 | 0.993 | 0.044 | 0.993 | 0.993 | 0.044 |
| 688 | P-116-SUP | 54 | M | U1 | 6.37 | 6.49 | 0.032 | 0.960 | 0.031 | 0.961 | 0.961 | 0.031 |
| 689 | P-116-SUP | 54 | M | U3 | 4.97 | 5.35 | 0.020 | 0.953 | 0.020 | 0.953 | 0.953 | 0.020 |
| 690 | P-116-SUP | 54 | M | T1 | 4.07 | 5.09 | 0.025 | 0.944 | 0.026 | 0.943 | 0.944 | 0.025 |
| 691 | P-116-SUP | 54 | M | T2 | 4.07 | 2.93 | 0.024 | 0.914 | 0.023 | 0.917 | 0.917 | 0.023 |
| 692 | P-116-SUP | 54 | M | T3 | 5.35 | 6.24 | 0.034 | 0.928 | 0.034 | 0.928 | 0.928 | 0.034 |
| 693 | P-116-SUP | 54 | M | L1 | 28.91 | 16.29 | 0.029 | 0.989 | 0.028 | 0.989 | 0.989 | 0.028 |
| 694 | P-116-SUP | 54 | M | L2 | 33.11 | 15.53 | 0.035 | 0.993 | 0.037 | 0.992 | 0.993 | 0.035 |
| 695 | P-117-ROS | 45 | M | U1 | 4.46 | 4.45 | 0.022 | 0.935 | 0.022 | 0.935 | 0.935 | 0.022 |
| 696 | P-117-ROS | 45 | M | T1 | 12.74 | 15.27 | 0.050 | 0.921 | 0.040 | 0.947 | 0.947 | 0.040 |
| 697 | P-117-ROS | 45 | M | T2 | 10.06 | 10.82 | 0.053 | 0.916 | 0.053 | 0.918 | 0.918 | 0.053 |
| 698 | P-117-ROS | 45 | M | L1 | 19.87 | 12.35 | 0.050 | 0.974 | 0.048 | 0.975 | 0.975 | 0.048 |
| 699 | P-117-ROS | 45 | M | L2 | 4.97 | 5.60 | 0.031 | 0.952 | 0.031 | 0.952 | 0.952 | 0.031 |
| 700 | P-117-ROS | 45 | M | L3 | 13.88 | 11.71 | 0.066 | 0.930 | 0.065 | 0.931 | 0.931 | 0.065 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 701 | P-118-RAJ | 47 | M | U1 | 8.92 | 15.91 | 0.046 | 0.915 | 0.043 | 0.926 | 0.926 | 0.043 |
| 702 | P-118-RAJ | 47 | M | U3 | 6.75 | 7.38 | 0.033 | 0.943 | 0.033 | 0.945 | 0.945 | 0.033 |
| 703 | P-118-RAJ | 47 | M | T3 | 7.90 | 8.65 | 0.060 | 0.941 | 0.062 | 0.953 | 0.953 | 0.062 |
| 704 | P-118-RAJ | 47 | M | L1B | 33.88 | 12.85 | 0.052 | 0.972 | 0.049 | 0.973 | 0.973 | 0.049 |
| 705 | P-118-RAJ | 47 | M | L2C | 15.67 | 27.49 | 0.054 | 0.972 | 0.051 | 0.973 | 0.973 | 0.051 |
| 706 | P-118-RAJ | 47 | M | L3a | 8.02 | 10.56 | 0.054 | 0.956 | 0.056 | 0.953 | 0.956 | 0.054 |
| 707 | P-119-RAT | 45 | M | H1 | 7.13 | 6.62 | 0.021 | 0.987 | 0.021 | 0.987 | 0.987 | 0.021 |
| 708 | P-119-RAT | 45 | M | U1 | 20.00 | 16.42 | 0.035 | 0.990 | 0.034 | 0.991 | 0.991 | 0.034 |
| 709 | P-119-RAT | 45 | M | U2 | 13.63 | 15.53 | 0.040 | 0.971 | 0.041 | 0.971 | 0.971 | 0.041 |
| 710 | P-119-RAT | 45 | M | U4 | 39.74 | 22.91 | 0.045 | 0.980 | 0.048 | 0.978 | 0.980 | 0.045 |
| 711 | P-119-RAT | 45 | M | T1 | 25.47 | 16.42 | 0.040 | 0.960 | 0.044 | 0.976 | 0.976 | 0.044 |
| 712 | P-119-RAT | 45 | M | T2 | 27.77 | 16.67 | 0.045 | 0.929 | 0.043 | 0.937 | 0.937 | 0.043 |
| 713 | P-119-RAT | 45 | M | T3 | 15.79 | 16.54 | 0.048 | 0.951 | 0.048 | 0.953 | 0.953 | 0.048 |
| 714 | P-119-RAT | 45 | M | L1 | 38.72 | 19.22 | 0.032 | 0.991 | 0.050 | 0.981 | 0.991 | 0.032 |
| 715 | P-119-RAT | 45 | M | L2 | 13.75 | 16.04 | 0.033 | 0.992 | 0.033 | 0.992 | 0.992 | 0.033 |
| 716 | P-120-ROS | 40 | M | U1 | 7.26 | 6.11 | 0.034 | 0.942 | 0.034 | 0.943 | 0.943 | 0.034 |
| 717 | P-120-ROS | 40 | M | U2 | 5.22 | 5.60 | 0.029 | 0.949 | 0.028 | 0.950 | 0.950 | 0.028 |
| 718 | P-120-ROS | 40 | M | U3 | 2.68 | 3.05 | 0.018 | 0.959 | 0.018 | 0.958 | 0.959 | 0.018 |
| 719 | P-120-ROS | 40 | M | T1 | 9.30 | 11.07 | 0.029 | 0.954 | 0.029 | 0.956 | 0.956 | 0.029 |
| 720 | P-120-ROS | 40 | M | T2 | 7.64 | 4.20 | 0.026 | 0.964 | 0.024 | 0.969 | 0.969 | 0.024 |
| 721 | P-120-ROS | 40 | M | L1 | 6.11 | 9.93 | 0.028 | 0.953 | 0.028 | 0.953 | 0.953 | 0.028 |
| 722 | P-120-ROS | 40 | M | L2 | 6.37 | 9.55 | 0.029 | 0.957 | 0.029 | 0.957 | 0.957 | 0.029 |
| 723 | P-121-PAR | 52 | M | U1 | 16.43 | 17.31 | 0.065 | 0.933 | 0.063 | 0.936 | 0.936 | 0.063 |
| 724 | P-121-PAR | 52 | M | U2 | 20.38 | 19.09 | 0.046 | 0.985 | 0.044 | 0.986 | 0.986 | 0.044 |
| 725 | P-121-PAR | 52 | M | U3 | 15.67 | 19.22 | 0.037 | 0.939 | 0.037 | 0.940 | 0.940 | 0.037 |
| 726 | P-121-PAR | 52 | M | T1 | 12.74 | 11.07 | 0.029 | 0.965 | 0.028 | 0.968 | 0.968 | 0.028 |
| 727 | P-121-PAR | 52 | M | T2 | 10.70 | 16.16 | 0.035 | 0.944 | 0.033 | 0.948 | 0.948 | 0.033 |
| 728 | P-121-PAR | 52 | M | T3 | 12.23 | 15.40 | 0.022 | 0.958 | 0.024 | 0.954 | 0.958 | 0.022 |
| 729 | P-121-PAR | 52 | M | L2 | 19.36 | 18.71 | 0.028 | 0.947 | 0.030 | 0.946 | 0.947 | 0.028 |
| 730 | P-121-PAR | 52 | M | L3 | 17.58 | 15.53 | 0.061 | 0.953 | 0.058 | 0.958 | 0.958 | 0.058 |
| 731 | P-121-PAR | 52 | M | L4 | 25.35 | 24.56 | 0.026 | 0.965 | 0.032 | 0.992 | 0.992 | 0.032 |
| 732 | P-122-RIZ | 38 | M | H1 | 12.86 | 12.98 | 0.027 | 0.996 | 0.026 | 0.996 | 0.996 | 0.026 |
| 733 | P-122-RIZ | 38 | M | U1 | 10.19 | 8.78 | 0.024 | 0.980 | 0.024 | 0.981 | 0.981 | 0.024 |
| 734 | P-122-RIZ | 38 | M | U2 | 9.81 | 8.91 | 0.054 | 0.947 | 0.053 | 0.948 | 0.948 | 0.053 |
| 735 | P-122-RIZ | 38 | M | U3 | 5.99 | 8.15 | 0.033 | 0.966 | 0.032 | 0.968 | 0.968 | 0.032 |
| 736 | P-122-RIZ | 38 | M | T1 | 20.76 | 14.89 | 0.049 | 0.968 | 0.049 | 0.967 | 0.968 | 0.049 |
| 737 | P-122-RIZ | 38 | M | T2 | 18.60 | 10.95 | 0.039 | 0.914 | 0.046 | 0.905 | 0.914 | 0.039 |
| 738 | P-122-RIZ | 38 | M | T3 | 8.92 | 12.09 | 0.042 | 0.947 | 0.042 | 0.933 | 0.947 | 0.042 |
| 739 | P-122-RIZ | 38 | M | L1 | 15.28 | 25.33 | 0.043 | 0.964 | 0.042 | 0.964 | 0.964 | 0.042 |
| 740 | P-122-RIZ | 38 | M | L2 | 16.94 | 15.02 | 0.034 | 0.980 | 0.033 | 0.981 | 0.981 | 0.033 |
| 741 | P-122-RIZ | 38 | M | L5 | 8.28 | 11.45 | 0.030 | 0.964 | 0.030 | 0.963 | 0.964 | 0.030 |
| 742 | P-123-NUR | 23 | F | U1 | 17.58 | 12.85 | 0.022 | 0.988 | 0.022 | 0.988 | 0.988 | 0.022 |
| 743 | P-123-NUR | 23 | F | U2B | 7.13 | 8.91 | 0.051 | 0.970 | 0.050 | 0.971 | 0.971 | 0.050 |
| 744 | P-123-NUR | 23 | F | T2 | 8.79 | 7.13 | 0.027 | 0.943 | 0.025 | 0.947 | 0.947 | 0.025 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 745 | P-123-NUR | 23 | F | T3 | 6.37 | 5.22 | 0.028 | 0.924 | 0.026 | 0.933 | 0.933 | 0.026 |
| 746 | P-123-NUR | 23 | F | L1 | 28.78 | 19.98 | 0.030 | 0.990 | 0.035 | 0.988 | 0.990 | 0.030 |
| 747 | P-123-NUR | 23 | F | L2 | 20.25 | 19.47 | 0.023 | 0.959 | 0.022 | 0.964 | 0.964 | 0.022 |
| 748 | P-123-NUR | 23 | F | L3B | 10.06 | 8.02 | 0.050 | 0.919 | 0.050 | 0.918 | 0.919 | 0.050 |
| 749 | P-124-SAN | 38 | M | H1 | 32.73 | 13.11 | 0.054 | 0.989 | 0.051 | 0.990 | 0.990 | 0.051 |
| 750 | P-124-SAN | 38 | M | U1 | 10.83 | 12.47 | 0.028 | 0.970 | 0.027 | 0.972 | 0.972 | 0.027 |
| 751 | P-124-SAN | 38 | M | U2 | 15.28 | 27.87 | 0.047 | 0.987 | 0.046 | 0.987 | 0.987 | 0.046 |
| 752 | P-124-SAN | 38 | M | U3 | 13.75 | 13.36 | 0.034 | 0.968 | 0.033 | 0.968 | 0.968 | 0.033 |
| 753 | P-124-SAN | 38 | M | T1 | 17.07 | 25.07 | 0.058 | 0.959 | 0.057 | 0.958 | 0.959 | 0.058 |
| 754 | P-124-SAN | 38 | M | T3 | 21.14 | 12.34 | 0.083 | 0.934 | 0.082 | 0.936 | 0.936 | 0.082 |
| 755 | P-124-SAN | 38 | M | L2 | 17.83 | 17.82 | 0.021 | 0.994 | 0.020 | 0.994 | 0.994 | 0.020 |
| 756 | P-124-SAN | 38 | M | L4 | 33.37 | 26.60 | 0.032 | 0.996 | 0.044 | 0.990 | 0.996 | 0.032 |
| 757 | P-125-LIM | 71 | M | U2 | 16.94 | 14.13 | 0.034 | 0.962 | 0.035 | 0.962 | 0.962 | 0.035 |
| 758 | P-125-LIM | 71 | M | U3 | 12.61 | 15.53 | 0.049 | 0.917 | 0.048 | 0.918 | 0.918 | 0.048 |
| 759 | P-125-LIM | 71 | M | T1 | 23.69 | 14.76 | 0.055 | 0.941 | 0.054 | 0.942 | 0.942 | 0.054 |
| 760 | P-125-LIM | 71 | M | T2 | 16.94 | 23.16 | 0.056 | 0.977 | 0.052 | 0.980 | 0.980 | 0.052 |
| 761 | P-125-LIM | 71 | M | T3 | 14.01 | 20.11 | 0.033 | 0.964 | 0.033 | 0.964 | 0.964 | 0.033 |
| 762 | P-125-LIM | 71 | M | L1B | 22.54 | 22.78 | 0.054 | 0.988 | 0.051 | 0.988 | 0.988 | 0.051 |
| 763 | P-125-LIM | 71 | M | L2 | 13.88 | 12.22 | 0.045 | 0.983 | 0.044 | 0.983 | 0.983 | 0.044 |
| 764 | P-125-LIM | 71 | M | L3 | 24.20 | 13.49 | 0.064 | 0.973 | 0.058 | 0.977 | 0.977 | 0.058 |
| 765 | P-126-WAN | 23 | M | H1 | 5.35 | 7.64 | 0.025 | 0.967 | 0.025 | 0.967 | 0.967 | 0.025 |
| 766 | P-126-WAN | 23 | M | U1 | 4.46 | 4.96 | 0.020 | 0.970 | 0.020 | 0.971 | 0.971 | 0.020 |
| 767 | P-126-WAN | 23 | M | U2 | 3.82 | 4.96 | 0.025 | 0.932 | 0.025 | 0.935 | 0.935 | 0.025 |
| 768 | P-126-WAN | 23 | M | U3 | 3.18 | 2.80 | 0.020 | 0.948 | 0.020 | 0.948 | 0.948 | 0.020 |
| 769 | P-126-WAN | 23 | M | T1 | 5.99 | 9.04 | 0.014 | 0.979 | 0.015 | 0.962 | 0.979 | 0.014 |
| 770 | P-126-WAN | 23 | M | T2 | 3.18 | 4.84 | 0.014 | 0.988 | 0.014 | 0.988 | 0.988 | 0.014 |
| 771 | P-126-WAN | 23 | M | T3 | 3.82 | 3.69 | 0.018 | 0.936 | 0.018 | 0.936 | 0.936 | 0.018 |
| 772 | P-126-WAN | 23 | M | L1 | 11.46 | 19.09 | 0.040 | 0.986 | 0.037 | 0.988 | 0.988 | 0.037 |
| 773 | P-126-WAN | 23 | M | L2 | 13.50 | 16.04 | 0.039 | 0.977 | 0.038 | 0.979 | 0.979 | 0.038 |
| 774 | P-126-WAN | 23 | M | L3B | 7.51 | 6.11 | 0.020 | 0.932 | 0.020 | 0.929 | 0.932 | 0.020 |
| 775 | P-127-EMA | 60 | F | T1 | 9.81 | 5.47 | 0.023 | 0.955 | 0.022 | 0.958 | 0.958 | 0.022 |
| 776 | P-127-EMA | 60 | F | T2 | 12.10 | 11.71 | 0.055 | 0.948 | 0.052 | 0.952 | 0.952 | 0.052 |
| 777 | P-127-EMA | 60 | F | T3 | 4.84 | 8.15 | 0.032 | 0.960 | 0.031 | 0.963 | 0.963 | 0.031 |
| 778 | P-127-EMA | 60 | F | L1 | 12.74 | 17.95 | 0.048 | 0.967 | 0.047 | 0.969 | 0.969 | 0.047 |
| 779 | P-127-EMA | 60 | F | L2B | 4.20 | 4.84 | 0.020 | 0.934 | 0.020 | 0.934 | 0.934 | 0.020 |
| 780 | P-127-EMA | 60 | F | L3 | 5.35 | 12.98 | 0.049 | 0.971 | 0.049 | 0.970 | 0.971 | 0.049 |
| 781 | P-128-SHA | 41 | M | H1 | 9.93 | 10.82 | 0.026 | 0.986 | 0.024 | 0.988 | 0.988 | 0.024 |
| 782 | P-128-SHA | 41 | M | U1 | 10.83 | 15.15 | 0.033 | 0.982 | 0.036 | 0.981 | 0.982 | 0.033 |
| 783 | P-128-SHA | 41 | M | U3 | 20.76 | 12.22 | 0.047 | 0.971 | 0.045 | 0.973 | 0.973 | 0.045 |
| 784 | P-128-SHA | 41 | M | T2 | 7.51 | 7.64 | 0.028 | 0.977 | 0.026 | 0.978 | 0.978 | 0.026 |
| 785 | P-128-SHA | 41 | M | T3 | 10.32 | 9.67 | 0.051 | 0.946 | 0.052 | 0.944 | 0.946 | 0.051 |
| 786 | P-128-SHA | 41 | M | T4 | 37.44 | 28.76 | 0.103 | 0.971 | 0.111 | 0.986 | 0.986 | 0.111 |
| 787 | P-128-SHA | 41 | M | L1 | 20.50 | 16.93 | 0.038 | 0.992 | 0.039 | 0.992 | 0.992 | 0.039 |
| 788 | P-128-SHA | 41 | M | L2 | 18.60 | 13.62 | 0.059 | 0.987 | 0.047 | 0.992 | 0.992 | 0.047 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 789 | P-128-SHA | 41 | M | L3 | 14.14 | 11.07 | 0.027 | 0.979 | 0.027 | 0.979 | 0.979 | 0.027 |
| 790 | P-128-SHA | 41 | M | L4 | 36.68 | 29.15 | 0.042 | 0.994 | 0.077 | 0.972 | 0.994 | 0.042 |
| 791 | P-74-LAT | 50 | M | H1 | 4.075 | 3.691 | 0.021 | 0.936 | 0.021 | 0.935 | 0.936 | 0.021 |
| 792 | P-74-LAT | 50 | M | U1 | 12.864 | 20.491 | 0.095 | 0.958 | 0.101 | 0.955 | 0.958 | 0.095 |
| 793 | P-74-LAT | 50 | M | U2 | 3.311 | 3.437 | 0.028 | 0.931 | 0.028 | 0.931 | 0.931 | 0.028 |
| 794 | P-74-LAT | 50 | M | U3 | 9.552 | 12.345 | 0.045 | 0.948 | 0.041 | 0.955 | 0.955 | 0.041 |
| 795 | P-74-LAT | 50 | M | T1 | 4.203 | 4.963 | 0.035 | 0.929 | 0.035 | 0.928 | 0.929 | 0.035 |
| 796 | P-74-LAT | 50 | M | T2 | 4.203 | 5.855 | 0.027 | 0.912 | 0.026 | 0.917 | 0.917 | 0.026 |
| 797 | P-74-LAT | 50 | M | T3 | 11.717 | 11.582 | 0.044 | 0.962 | 0.042 | 0.965 | 0.965 | 0.042 |
| 798 | P-74-LAT | 50 | M | L1 | 12.226 | 15.654 | 0.063 | 0.950 | 0.057 | 0.957 | 0.957 | 0.057 |
| 799 | P-74-LAT | 50 | M | L2 | 16.811 | 14.890 | 0.057 | 0.967 | 0.057 | 0.966 | 0.967 | 0.057 |
| 800 | P-74-LAT | 50 | M | L3 | 15.156 | 16.672 | 0.069 | 0.953 | 0.066 | 0.956 | 0.956 | 0.066 |
| 801 | P-75-PON | 51 | M | H1 | 3.821 | 3.309 | 0.016 | 0.947 | 0.019 | 0.924 | 0.947 | 0.016 |
| 802 | P-75-PON | 51 | M | U1 | 7.259 | 5.981 | 0.037 | 0.945 | 0.037 | 0.945 | 0.945 | 0.037 |
| 803 | P-75-PON | 51 | M | U2 | 10.189 | 11.327 | 0.056 | 0.979 | 0.054 | 0.980 | 0.980 | 0.054 |
| 804 | P-75-PON | 51 | M | U3 | 9.170 | 6.109 | 0.045 | 0.962 | 0.047 | 0.958 | 0.962 | 0.045 |
| 805 | P-75-PON | 51 | M | T2 | 11.462 | 16.036 | 0.036 | 0.977 | 0.034 | 0.979 | 0.979 | 0.034 |
| 806 | P-75-PON | 51 | M | T1 | 11.080 | 13.490 | 0.039 | 0.947 | 0.040 | 0.946 | 0.947 | 0.039 |
| 807 | P-75-PON | 51 | M | L1 | 21.142 | 12.599 | 0.023 | 0.976 | 0.028 | 0.984 | 0.984 | 0.028 |
| 808 | P-75-PON | 51 | M | L2 | 16.684 | 16.163 | 0.087 | 0.917 | 0.091 | 0.911 | 0.917 | 0.087 |
| 809 | P-75-PON | 51 | M | L3 | 12.354 | 11.200 | 0.029 | 0.984 | 0.029 | 0.985 | 0.985 | 0.029 |
| 810 | $\mathrm{P}-76-\mathrm{MOH}$ | 47 | M | H1 | 6.113 | 7.509 | 0.018 | 0.985 | 0.018 | 0.986 | 0.986 | 0.018 |
| 811 | P-76-MOH | 47 | M | U1 | 17.194 | 15.654 | 0.032 | 0.963 | 0.033 | 0.962 | 0.963 | 0.032 |
| 812 | P-76-MOH | 47 | M | U2 | 15.028 | 12.091 | 0.029 | 0.972 | 0.030 | 0.971 | 0.972 | 0.029 |
| 813 | P-76-MOH | 47 | M | U3 | 7.259 | 11.454 | 0.028 | 0.980 | 0.027 | 0.981 | 0.981 | 0.027 |
| 814 | P-76-MOH | 47 | M | T1 | 21.524 | 19.599 | 0.033 | 0.975 | 0.030 | 0.978 | 0.978 | 0.030 |
| 815 | P-76-MOH | 47 | M | T2 | 19.359 | 17.181 | 0.062 | 0.945 | 0.061 | 0.948 | 0.948 | 0.061 |
| 816 | P-76-MOH | 47 | M | T3 | 11.844 | 9.163 | 0.025 | 0.961 | 0.026 | 0.957 | 0.961 | 0.025 |
| 817 | P-76-MOH | 47 | M | T4 | 11.081 | 8.654 | 0.049 | 0.975 | 0.048 | 0.975 | 0.975 | 0.048 |
| 818 | P-76-MOH | 47 | M | L1 | 10.316 | 7.382 | 0.031 | 0.987 | 0.031 | 0.986 | 0.987 | 0.031 |
| 819 | P-76-MOH | 47 | M | L2 | 4.202 | 7.509 | 0.030 | 0.967 | 0.029 | 0.969 | 0.969 | 0.029 |
| 820 | P-76-MOH | 47 | M | L3 | 9.552 | 9.036 | 0.035 | 0.989 | 0.035 | 0.990 | 0.990 | 0.035 |
| 821 | P-77-AZI | 46 | M | H1 | 14.647 | 13.745 | 0.022 | 0.979 | 0.022 | 0.979 | 0.979 | 0.022 |
| 822 | P-77-AZI | 46 | M | T1 | 21.269 | 28.254 | 0.040 | 0.937 | 0.045 | 0.928 | 0.937 | 0.040 |
| 823 | P-77-AZI | 46 | M | T4 | 20.123 | 18.327 | 0.037 | 0.994 | 0.036 | 0.994 | 0.994 | 0.036 |
| 824 | P-77-AZI | 46 | M | L2 | 14.646 | 13.363 | 0.051 | 0.977 | 0.052 | 0.978 | 0.978 | 0.052 |
| 825 | P-77-AZI | 46 | M | L3 | 9.551 | 9.927 | 0.053 | 0.983 | 0.053 | 0.983 | 0.983 | 0.053 |
| 826 | P-78-NUR | 25 | F | H1 | 30.948 | 10.055 | 0.039 | 0.984 | 0.038 | 0.987 | 0.987 | 0.038 |
| 827 | P-78-NUR | 25 | F | U1 | 19.996 | 20.108 | 0.058 | 0.984 | 0.059 | 0.983 | 0.984 | 0.058 |
| 828 | P-78-NUR | 25 | F | U2 | 19.358 | 28.890 | 0.055 | 0.958 | 0.057 | 0.952 | 0.958 | 0.055 |
| 829 | P-78-NUR | 25 | F | U3 | 16.684 | 15.273 | 0.036 | 0.996 | 0.035 | 0.996 | 0.996 | 0.035 |
| 830 | P-78-NUR | 25 | F | T1 | 18.085 | 10.563 | 0.076 | 0.980 | 0.075 | 0.980 | 0.980 | 0.075 |
| 831 | P-78-NUR | 25 | F | T2 | 16.557 | 29.272 | 0.050 | 0.932 | 0.049 | 0.933 | 0.933 | 0.049 |
| 832 | P-78-NUR | 25 | F | T3 | 14.137 | 21.891 | 0.073 | 0.925 | 0.073 | 0.925 | 0.925 | 0.073 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 833 | P-78-NUR | 25 | F | L2 | 17.958 | 20.490 | 0.055 | 0.965 | 0.062 | 0.957 | 0.965 | 0.055 |
| 834 | P-78-NUR | 25 | F | L3 | 12.481 | 11.327 | 0.066 | 0.967 | 0.064 | 0.969 | 0.969 | 0.064 |
| 835 | P-79-AZM | 27 | M | H1 | 4.967 | 4.964 | 0.036 | 0.945 | 0.036 | 0.944 | 0.945 | 0.036 |
| 836 | P-79-AZM | 27 | M | U1 | 20.887 | 20.999 | 0.041 | 0.971 | 0.044 | 0.978 | 0.978 | 0.044 |
| 837 | P-79-AZM | 27 | M | U2 | 10.953 | 14.254 | 0.048 | 0.928 | 0.047 | 0.933 | 0.933 | 0.047 |
| 838 | P-79-AZM | 27 | M | U3 | 11.462 | 11.072 | 0.034 | 0.979 | 0.033 | 0.980 | 0.980 | 0.033 |
| 839 | P-79-AZM | 27 | M | T1 | 11.972 | 11.581 | 0.047 | 0.950 | 0.046 | 0.951 | 0.951 | 0.046 |
| 840 | P-79-AZM | 27 | M | T2 | 13.245 | 17.436 | 0.057 | 0.930 | 0.057 | 0.928 | 0.930 | 0.057 |
| 841 | P-79-AZM | 27 | M | T3 | 13.119 | 11.327 | 0.049 | 0.967 | 0.047 | 0.969 | 0.969 | 0.047 |
| 842 | P-79-AZM | 27 | M | L1 | 9.170 | 9.927 | 0.027 | 0.965 | 0.027 | 0.968 | 0.968 | 0.027 |
| 843 | P-79-AZM | 27 | M | L2 | 7.514 | 9.800 | 0.036 | 0.944 | 0.035 | 0.964 | 0.964 | 0.035 |
| 844 | P-79-AZM | 27 | M | L3 | 19.613 | 9.291 | 0.045 | 0.966 | 0.046 | 0.964 | 0.966 | 0.045 |
| 845 | P-80-SIN | 78 | M | U1 | 14.392 | 12.855 | 0.064 | 0.952 | 0.071 | 0.949 | 0.952 | 0.064 |
| 846 | P-80-SIN | 78 | M | T1 | 22.670 | 22.018 | 0.057 | 0.974 | 0.059 | 0.972 | 0.974 | 0.057 |
| 847 | P-80-SIN | 78 | M | T4 | 24.580 | 22.908 | 0.054 | 0.970 | 0.062 | 0.957 | 0.970 | 0.054 |
| 848 | P-81-SEL | 57 | M | H1 | 8.151 | 7.127 | 0.032 | 0.941 | 0.031 | 0.944 | 0.944 | 0.031 |
| 849 | P-81-SEL | 57 | M | U1 | 10.953 | 13.872 | 0.048 | 0.954 | 0.048 | 0.948 | 0.954 | 0.048 |
| 850 | P-81-SEL | 57 | M | U3 | 5.477 | 5.855 | 0.028 | 0.934 | 0.028 | 0.933 | 0.934 | 0.028 |
| 851 | P-81-SEL | 57 | M | T1 | 10.826 | 7.891 | 0.046 | 0.950 | 0.044 | 0.955 | 0.955 | 0.044 |
| 852 | P-81-SEL | 57 | M | T2 | 26.745 | 8.909 | 0.051 | 0.988 | 0.050 | 0.988 | 0.988 | 0.050 |
| 853 | P-81-SEL | 57 | M | T3 | 13.755 | 14.000 | 0.040 | 0.964 | 0.041 | 0.951 | 0.964 | 0.040 |
| 854 | P-81-SEL | 57 | M | L1 | 22.161 | 17.182 | 0.057 | 0.979 | 0.055 | 0.980 | 0.980 | 0.055 |
| 855 | P-81-SEL | 57 | M | L2 | 28.783 | 16.927 | 0.066 | 0.961 | 0.072 | 0.953 | 0.961 | 0.066 |
| 856 | P-81-SEL | 57 | M | L3 | 20.377 | 28.508 | 0.059 | 0.987 | 0.063 | 0.986 | 0.987 | 0.059 |
| 857 | P-82-MAH | 51 | M | U1 | 6.495 | 8.145 | 0.036 | 0.940 | 0.036 | 0.940 | 0.940 | 0.036 |
| 858 | P-82-MAH | 51 | M | U2 | 8.533 | 13.872 | 0.038 | 0.966 | 0.038 | 0.967 | 0.967 | 0.038 |
| 859 | P-82-MAH | 51 | M | U3 | 6.750 | 3.054 | 0.023 | 0.947 | 0.023 | 0.947 | 0.947 | 0.023 |
| 860 | P-82-MAH | 51 | M | T1 | 12.609 | 12.091 | 0.050 | 0.961 | 0.050 | 0.960 | 0.961 | 0.050 |
| 861 | P-82-MAH | 51 | M | T2 | 6.878 | 8.527 | 0.036 | 0.956 | 0.036 | 0.956 | 0.956 | 0.036 |
| 862 | P-82-MAH | 51 | M | T3 | 16.939 | 13.363 | 0.032 | 0.975 | 0.039 | 0.966 | 0.975 | 0.032 |
| 863 | P-82-MAH | 51 | M | L1 | 12.226 | 11.582 | 0.025 | 0.995 | 0.027 | 0.993 | 0.995 | 0.025 |
| 864 | P-82-MAH | 51 | M | L2 | 4.967 | 5.345 | 0.021 | 0.911 | 0.021 | 0.911 | 0.911 | 0.021 |
| 865 | P-82-MAH | 51 | M | L3 | 5.732 | 7.000 | 0.040 | 0.937 | 0.039 | 0.943 | 0.943 | 0.039 |
| 866 | P-83-SHA | 34 | M | H1 | 4.967 | 4.073 | 0.031 | 0.966 | 0.031 | 0.966 | 0.966 | 0.031 |
| 867 | P-83-SHA | 34 | M | U1 | 17.067 | 7.891 | 0.039 | 0.971 | 0.043 | 0.966 | 0.971 | 0.039 |
| 868 | P-83-SHA | 34 | M | U2 | 11.335 | 22.908 | 0.052 | 0.965 | 0.049 | 0.957 | 0.965 | 0.052 |
| 869 | P-83-SHA | 34 | M | U3 | 7.896 | 8.654 | 0.057 | 0.912 | 0.065 | 0.905 | 0.912 | 0.057 |
| 870 | P-83-SHA | 34 | M | T1 | 14.647 | 26.090 | 0.051 | 0.978 | 0.056 | 0.973 | 0.978 | 0.051 |
| 871 | P-83-SHA | 34 | M | T2 | 33.114 | 16.799 | 0.073 | 0.978 | 0.072 | 0.979 | 0.979 | 0.072 |
| 872 | P-83-SHA | 34 | M | T3 | 29.038 | 13.745 | 0.070 | 0.956 | 0.069 | 0.958 | 0.958 | 0.069 |
| 873 | P-83-SHA | 34 | M | L1 | 14.902 | 25.072 | 0.059 | 0.966 | 0.061 | 0.964 | 0.966 | 0.059 |
| 874 | P-83-SHA | 34 | M | L2 | 30.948 | 17.181 | 0.064 | 0.990 | 0.066 | 0.988 | 0.990 | 0.064 |
| 875 | P-83-SHA | 34 | M | L3 | 16.684 | 24.690 | 0.085 | 0.938 | 0.085 | 0.941 | 0.941 | 0.085 |
| 876 | P-84-NOO | 49 | F | H1 | 18.977 | 15.527 | 0.040 | 0.962 | 0.042 | 0.958 | 0.962 | 0.040 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 877 | P-84-NOO | 49 | F | U1 | 8.915 | 20.745 | 0.056 | 0.942 | 0.055 | 0.941 | 0.942 | 0.056 |
| 878 | P-84-NOO | 49 | F | U2 | 15.283 | 20.491 | 0.088 | 0.982 | 0.087 | 0.982 | 0.982 | 0.087 |
| 879 | P-84-NOO | 49 | F | T1 | 10.316 | 18.454 | 0.061 | 0.905 | 0.058 | 0.913 | 0.913 | 0.058 |
| 880 | P-84-NOO | 49 | F | T3 | 24.071 | 14.126 | 0.072 | 0.921 | 0.075 | 0.913 | 0.921 | 0.072 |
| 881 | P-84-NOO | 49 | F | L1 | 14.519 | 19.981 | 0.032 | 0.988 | 0.037 | 0.986 | 0.988 | 0.032 |
| 882 | P-84-NOO | 49 | F | L2 | 17.449 | 17.817 | 0.061 | 0.913 | 0.059 | 0.916 | 0.916 | 0.059 |
| 883 | P-84-NOO | 49 | F | L3 | 23.689 | 16.672 | 0.058 | 0.979 | 0.072 | 0.972 | 0.979 | 0.058 |
| 884 | P-85-CHE | 61 | M | U1 | 11.335 | 7.127 | 0.026 | 0.970 | 0.026 | 0.968 | 0.970 | 0.026 |
| 885 | P-85-CHE | 61 | M | U2 | 7.514 | 12.218 | 0.048 | 0.926 | 0.049 | 0.922 | 0.926 | 0.048 |
| 886 | P-85-CHE | 61 | M | U3 | 13.245 | 17.054 | 0.071 | 0.945 | 0.067 | 0.949 | 0.949 | 0.067 |
| 887 | P-85-CHE | 61 | M | T1 | 10.699 | 13.236 | 0.043 | 0.960 | 0.043 | 0.960 | 0.960 | 0.043 |
| 888 | P-85-CHE | 61 | M | T2 | 14.774 | 12.981 | 0.040 | 0.924 | 0.041 | 0.923 | 0.924 | 0.040 |
| 889 | P-85-CHE | 61 | M | T3 | 16.684 | 15.527 | 0.068 | 0.933 | 0.068 | 0.958 | 0.958 | 0.068 |
| 890 | P-85-CHE | 61 | M | T4 | 30.439 | 27.999 | 0.093 | 0.931 | 0.091 | 0.933 | 0.933 | 0.091 |
| 891 | P-85-CHE | 61 | M | L1 | 16.684 | 13.618 | 0.038 | 0.957 | 0.038 | 0.958 | 0.958 | 0.038 |
| 892 | P-85-CHE | 61 | M | L2 | 9.552 | 23.672 | 0.060 | 0.979 | 0.060 | 0.979 | 0.979 | 0.060 |
| 893 | P-86-AKI | 47 | F | U2 | 2.674 | 3.818 | 0.022 | 0.972 | 0.022 | 0.972 | 0.972 | 0.022 |
| 894 | P-86-AKI | 47 | F | U3 | 4.585 | 7.000 | 0.023 | 0.925 | 0.023 | 0.924 | 0.925 | 0.023 |
| 895 | P-86-AKI | 47 | F | T1 | 5.349 | 7.636 | 0.037 | 0.934 | 0.036 | 0.967 | 0.967 | 0.036 |
| 896 | P-86-AKI | 47 | F | T2 | 5.604 | 5.982 | 0.020 | 0.957 | 0.020 | 0.958 | 0.958 | 0.020 |
| 897 | P-86-AKI | 47 | F | T3 | 4.840 | 5.091 | 0.043 | 0.957 | 0.042 | 0.958 | 0.958 | 0.042 |
| 898 | P-86-AKI | 47 | F | L1 | 5.477 | 4.454 | 0.026 | 0.964 | 0.027 | 0.963 | 0.964 | 0.026 |
| 899 | P-86-AKI | 47 | F | L2 | 5.604 | 5.982 | 0.046 | 0.931 | 0.046 | 0.933 | 0.933 | 0.046 |
| 900 | P-86-AKI | 47 | F | L3 | 5.986 | 7.763 | 0.026 | 0.924 | 0.026 | 0.920 | 0.924 | 0.026 |
| 901 | P-87-NOR | 33 | F | H1 | 8.024 | 6.363 | 0.029 | 0.932 | 0.030 | 0.920 | 0.932 | 0.029 |
| 902 | P-87-NOR | 33 | F | U1 | 18.340 | 14.508 | 0.050 | 0.953 | 0.049 | 0.954 | 0.954 | 0.049 |
| 903 | P-87-NOR | 33 | F | U2 | 11.972 | 12.091 | 0.054 | 0.995 | 0.054 | 0.996 | 0.996 | 0.054 |
| 904 | P-87-NOR | 33 | F | T1 | 19.359 | 16.418 | 0.054 | 0.933 | 0.053 | 0.935 | 0.935 | 0.053 |
| 905 | P-87-NOR | 33 | F | L1 | 16.303 | 14.636 | 0.036 | 0.972 | 0.035 | 0.973 | 0.973 | 0.035 |
| 906 | P-87-NOR | 33 | F | L3 | 11.972 | 9.163 | 0.045 | 0.948 | 0.034 | 0.937 | 0.948 | 0.045 |
| 907 | P-88-SAW | 57 | M | H1 | 12.227 | 13.364 | 0.040 | 0.962 | 0.036 | 0.958 | 0.962 | 0.040 |
| 908 | P-88-SAW | 57 | M | U1 | 9.552 | 5.727 | 0.024 | 0.971 | 0.023 | 0.973 | 0.973 | 0.023 |
| 909 | P-88-SAW | 57 | M | U2 | 6.622 | 5.091 | 0.028 | 0.976 | 0.028 | 0.976 | 0.976 | 0.028 |
| 910 | P-88-SAW | 57 | M | T1 | 9.042 | 13.872 | 0.041 | 0.923 | 0.041 | 0.925 | 0.925 | 0.041 |
| 911 | P-88-SAW | 57 | M | T2 | 11.972 | 5.346 | 0.042 | 0.929 | 0.040 | 0.934 | 0.934 | 0.040 |
| 912 | P-88-SAW | 57 | M | T3 | 10.316 | 7.891 | 0.046 | 0.935 | 0.043 | 0.943 | 0.943 | 0.043 |
| 913 | P-88-SAW | 57 | M | T4 | 11.717 | 12.600 | 0.054 | 0.936 | 0.057 | 0.924 | 0.936 | 0.054 |
| 914 | P-88-SAW | 57 | M | L1 | 10.062 | 10.309 | 0.035 | 0.960 | 0.034 | 0.962 | 0.962 | 0.034 |
| 915 | P-88-SAW | 57 | M | L2 | 7.387 | 8.018 | 0.042 | 0.945 | 0.039 | 0.950 | 0.950 | 0.039 |
| 916 | P-89-BAK | 48 | M | H1 | 3.439 | 4.072 | 0.020 | 0.966 | 0.020 | 0.966 | 0.966 | 0.020 |
| 917 | P-89-BAK | 48 | M | U1 | 13.500 | 10.945 | 0.070 | 0.950 | 0.066 | 0.956 | 0.956 | 0.066 |
| 918 | P-89-BAK | 48 | M | U2 | 10.953 | 12.981 | 0.060 | 0.957 | 0.060 | 0.957 | 0.957 | 0.060 |
| 919 | P-89-BAK | 48 | M | U3 | 26.491 | 14.764 | 0.086 | 0.944 | 0.075 | 0.954 | 0.954 | 0.075 |
| 920 | P-89-BAK | 48 | M | T2 | 21.524 | 11.455 | 0.080 | 0.964 | 0.084 | 0.960 | 0.964 | 0.080 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 921 | P-89-BAK | 48 | M | T3 | 9.424 | 20.108 | 0.048 | 0.973 | 0.048 | 0.974 | 0.974 | 0.048 |
| 922 | P-89-BAK | 48 | M | T4 | 6.877 | 9.545 | 0.046 | 0.946 | 0.041 | 0.956 | 0.956 | 0.041 |
| 923 | P-89-BAK | 48 | M | T6 | 31.840 | 26.090 | 0.070 | 0.948 | 0.073 | 0.942 | 0.948 | 0.070 |
| 924 | P-89-BAK | 48 | M | L1 | 16.175 | 16.036 | 0.042 | 0.981 | 0.037 | 0.985 | 0.985 | 0.037 |
| 925 | P-89-BAK | 48 | M | L2 | 21.906 | 17.436 | 0.063 | 0.964 | 0.062 | 0.965 | 0.965 | 0.062 |
| 926 | P-89-BAK | 48 | M | L3 | 9.934 | 9.036 | 0.037 | 0.946 | 0.035 | 0.950 | 0.950 | 0.035 |
| 927 | P-90-MOH | 51 | M | U1 | 8.405 | 5.473 | 0.026 | 0.920 | 0.026 | 0.920 | 0.920 | 0.026 |
| 928 | P-90-MOH | 51 | M | U3 | 4.712 | 4.709 | 0.021 | 0.948 | 0.020 | 0.954 | 0.954 | 0.020 |
| 929 | P-90-MOH | 51 | M | T1 | 4.968 | 5.473 | 0.038 | 0.926 | 0.025 | 0.955 | 0.955 | 0.025 |
| 930 | P-90-MOH | 51 | M | T2 | 6.496 | 7.127 | 0.025 | 0.960 | 0.025 | 0.932 | 0.960 | 0.025 |
| 931 | P-90-MOH | 51 | M | T3 | 2.929 | 3.436 | 0.020 | 0.935 | 0.021 | 0.935 | 0.935 | 0.021 |
| 932 | P-90-MOH | 51 | M | T4 | 11.844 | 15.527 | 0.032 | 0.940 | 0.031 | 0.941 | 0.941 | 0.031 |
| 933 | P-90-MOH | 51 | M | L2 | 4.076 | 4.582 | 0.031 | 0.939 | 0.029 | 0.947 | 0.947 | 0.029 |
| 934 | P-91-MOH | 28 | M | H1 | 8.915 | 10.309 | 0.024 | 0.951 | 0.025 | 0.946 | 0.951 | 0.024 |
| 935 | P-91-MOH | 28 | M | U1 | 12.481 | 13.363 | 0.042 | 0.966 | 0.041 | 0.966 | 0.966 | 0.041 |
| 936 | P-91-MOH | 28 | M | U2 | 23.944 | 18.073 | 0.068 | 0.962 | 0.067 | 0.962 | 0.962 | 0.067 |
| 937 | P-91-MOH | 28 | M | U3 | 9.807 | 9.927 | 0.042 | 0.956 | 0.041 | 0.958 | 0.958 | 0.041 |
| 938 | P-91-MOH | 28 | M | T1 | 26.618 | 10.945 | 0.072 | 0.970 | 0.068 | 0.973 | 0.973 | 0.068 |
| 939 | P-91-MOH | 28 | M | T2 | 23.052 | 13.108 | 0.071 | 0.930 | 0.071 | 0.932 | 0.932 | 0.071 |
| 940 | P-91-MOH | 28 | M | T3 | 17.066 | 12.854 | 0.066 | 0.920 | 0.065 | 0.921 | 0.921 | 0.065 |
| 941 | P-91-MOH | 28 | M | T4 | 9.170 | 7.764 | 0.028 | 0.939 | 0.028 | 0.939 | 0.939 | 0.028 |
| 942 | P-91-MOH | 28 | M | T5 | 28.656 | 26.090 | 0.116 | 0.909 | 0.132 | 0.789 | 0.909 | 0.116 |
| 943 | P-91-MOH | 28 | M | T6 | 25.727 | 25.326 | 0.085 | 0.933 | 0.076 | 0.931 | 0.933 | 0.085 |
| 944 | P-91-MOH | 28 | M | L1 | 18.468 | 18.454 | 0.071 | 0.978 | 0.069 | 0.980 | 0.980 | 0.069 |
| 945 | P-91-MOH | 28 | M | L2 | 17.321 | 18.072 | 0.059 | 0.956 | 0.059 | 0.957 | 0.957 | 0.059 |
| 946 | P-91-MOH | 28 | M | L3 | 23.307 | 13.363 | 0.056 | 0.951 | 0.054 | 0.956 | 0.956 | 0.054 |
| 947 | P-92-ASO | 55 | M | H1 | 9.043 | 12.854 | 0.029 | 0.927 | 0.030 | 0.926 | 0.927 | 0.029 |
| 948 | P-92-ASO | 55 | M | U1 | 17.703 | 11.199 | 0.049 | 0.983 | 0.049 | 0.982 | 0.983 | 0.049 |
| 949 | P-92-ASO | 55 | M | U2 | 11.335 | 7.382 | 0.026 | 0.953 | 0.025 | 0.955 | 0.955 | 0.025 |
| 950 | P-92-ASO | 55 | M | U3 | 7.133 | 8.655 | 0.028 | 0.959 | 0.029 | 0.956 | 0.959 | 0.028 |
| 951 | P-92-ASO | 55 | M | T1 | 6.750 | 13.873 | 0.046 | 0.962 | 0.041 | 0.967 | 0.967 | 0.041 |
| 952 | P-92-ASO | 55 | M | T2 | 4.585 | 5.473 | 0.024 | 0.929 | 0.024 | 0.930 | 0.930 | 0.024 |
| 953 | P-92-ASO | 55 | M | L1 | 12.609 | 12.855 | 0.050 | 0.977 | 0.048 | 0.977 | 0.977 | 0.048 |
| 954 | P-92-ASO | 55 | M | L2 | 9.934 | 13.872 | 0.032 | 0.982 | 0.030 | 0.983 | 0.983 | 0.030 |
| 955 | P-92-ASO | 55 | M | L3 | 19.995 | 14.127 | 0.057 | 0.959 | 0.055 | 0.962 | 0.962 | 0.055 |
| 956 | P-93-CHE | 40 | M | H1 | 17.321 | 18.708 | 0.080 | 0.939 | 0.085 | 0.947 | 0.947 | 0.085 |
| 957 | P-93-CHE | 40 | M | U1 | 10.062 | 11.327 | 0.036 | 0.949 | 0.035 | 0.938 | 0.949 | 0.036 |
| 958 | P-93-CHE | 40 | M | U2 | 8.916 | 8.781 | 0.037 | 0.981 | 0.033 | 0.986 | 0.986 | 0.033 |
| 959 | P-93-CHE | 40 | M | U3 | 12.991 | 11.200 | 0.028 | 0.967 | 0.029 | 0.966 | 0.967 | 0.028 |
| 960 | P-93-CHE | 40 | M | T1 | 23.052 | 14.000 | 0.063 | 0.969 | 0.063 | 0.970 | 0.970 | 0.063 |
| 961 | P-93-CHE | 40 | M | T2 | 17.958 | 9.673 | 0.040 | 0.928 | 0.037 | 0.925 | 0.928 | 0.040 |
| 962 | P-93-CHE | 40 | M | T3 | 20.632 | 9.927 | 0.060 | 0.917 | 0.059 | 0.919 | 0.919 | 0.059 |
| 963 | P-93-CHE | 40 | M | T4 | 21.524 | 18.581 | 0.055 | 0.967 | 0.054 | 0.969 | 0.969 | 0.054 |
| 964 | P-93-CHE | 40 | M | T5 | 32.222 | 25.199 | 0.051 | 0.969 | 0.058 | 0.970 | 0.970 | 0.058 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{\overline{S_{\text {final }}}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 965 | P-93-CHE | 40 | M | L1 | 15.028 | 17.308 | 0.033 | 0.986 | 0.032 | 0.986 | 0.986 | 0.032 |
| 966 | P-93-CHE | 40 | M | L3 | 16.812 | 11.964 | 0.062 | 0.949 | 0.060 | 0.951 | 0.951 | 0.060 |
| 967 | P-94-UVA | 22 | M | U1 | 5.477 | 5.345 | 0.013 | 0.965 | 0.013 | 0.965 | 0.965 | 0.013 |
| 968 | P-94-UVA | 22 | M | U2 | 5.349 | 6.873 | 0.024 | 0.928 | 0.023 | 0.930 | 0.930 | 0.023 |
| 969 | P-94-UVA | 22 | M | U3 | 8.279 | 5.600 | 0.019 | 0.971 | 0.019 | 0.973 | 0.973 | 0.019 |
| 970 | P-94-UVA | 22 | M | T1 | 6.750 | 5.218 | 0.025 | 0.907 | 0.021 | 0.930 | 0.930 | 0.021 |
| 971 | P-94-UVA | 22 | M | T2 | 3.184 | 2.927 | 0.027 | 0.919 | 0.027 | 0.919 | 0.919 | 0.027 |
| 972 | P-94-UVA | 22 | M | T3 | 6.113 | 4.709 | 0.021 | 0.945 | 0.021 | 0.946 | 0.946 | 0.021 |
| 973 | P-94-UVA | 22 | M | T4 | 7.005 | 4.327 | 0.015 | 0.968 | 0.016 | 0.966 | 0.968 | 0.015 |
| 974 | P-94-UVA | 22 | M | L2 | 4.330 | 3.946 | 0.017 | 0.948 | 0.017 | 0.948 | 0.948 | 0.017 |
| 975 | P-94-UVA | 22 | M | L3 | 4.330 | 4.454 | 0.021 | 0.985 | 0.022 | 0.984 | 0.985 | 0.021 |
| 976 | P-129-JEL | 43 | F | U11 | 12.482 | 12.345 | 0.031 | 0.969 | 0.029 | 0.973 | 0.973 | 0.029 |
| 977 | P-129-JEL | 43 | F | U12 | 12.736 | 12.218 | 0.026 | 0.952 | 0.027 | 0.951 | 0.952 | 0.026 |
| 978 | P-129-JEL | 43 | F | U3 | 13.373 | 12.855 | 0.061 | 0.926 | 0.062 | 0.923 | 0.926 | 0.061 |
| 979 | P-129-JEL | 43 | F | L1 | 11.844 | 11.073 | 0.035 | 0.973 | 0.040 | 0.938 | 0.973 | 0.035 |
| 980 | P-129-JEL | 43 | F | L2 | 11.590 | 16.163 | 0.048 | 0.929 | 0.049 | 0.915 | 0.929 | 0.048 |
| 981 | P-129-JEL | 43 | F | L3 | 9.425 | 11.327 | 0.041 | 0.879 | 0.038 | 0.905 | 0.905 | 0.038 |
| 982 | P-129-JEL | 43 | F | L32 | 9.934 | 10.308 | 0.041 | 0.964 | 0.041 | 0.964 | 0.964 | 0.041 |
| 983 | P-129-JEL | 43 | F | L33 | 9.297 | 11.581 | 0.051 | 0.964 | 0.051 | 0.963 | 0.964 | 0.051 |
| 984 | P-130-VIV | 67 | F | U1 | 3.311 | 3.818 | 0.051 | 0.920 | 0.051 | 0.919 | 0.920 | 0.051 |
| 985 | P-130-VIV | 67 | F | U2 | 1.656 | 2.545 | 0.020 | 0.935 | 0.019 | 0.936 | 0.936 | 0.019 |
| 986 | P-130-VIV | 67 | F | U3 | 2.802 | 2.546 | 0.019 | 0.932 | 0.020 | 0.934 | 0.934 | 0.020 |
| 987 | P-130-VIV | 67 | F | T2 | 12.736 | 10.945 | 0.054 | 0.766 | 0.042 | 0.900 | 0.900 | 0.042 |
| 988 | P-130-VIV | 67 | F | T3 | 21.651 | 12.982 | 0.070 | 0.931 | 0.068 | 0.936 | 0.936 | 0.068 |
| 989 | P-130-VIV | 67 | F | L1A | 5.476 | 4.709 | 0.029 | 0.922 | 0.029 | 0.922 | 0.922 | 0.029 |
| 990 | P-130-VIV | 67 | F | L2 | 3.821 | 4.073 | 0.017 | 0.916 | 0.017 | 0.915 | 0.916 | 0.017 |
| 991 | P-130-VIV | 67 | F | L3 | 3.948 | 5.218 | 0.028 | 0.964 | 0.027 | 0.965 | 0.965 | 0.027 |
| 992 | P-131-SIT | 58 | F | H | 8.278 | 12.218 | 0.042 | 0.950 | 0.041 | 0.952 | 0.952 | 0.041 |
| 993 | P-131-SIT | 58 | F | U1 | 8.024 | 12.854 | 0.043 | 0.962 | 0.043 | 0.963 | 0.963 | 0.043 |
| 994 | P-131-SIT | 58 | F | U2 | 15.920 | 23.544 | 0.041 | 0.966 | 0.040 | 0.969 | 0.969 | 0.040 |
| 995 | P-131-SIT | 58 | F | U3 | 8.406 | 10.436 | 0.027 | 0.926 | 0.028 | 0.924 | 0.926 | 0.027 |
| 996 | P-131-SIT | 58 | F | T1 | 11.717 | 10.054 | 0.041 | 0.939 | 0.039 | 0.945 | 0.945 | 0.039 |
| 997 | P-131-SIT | 58 | F | T2 | 12.481 | 13.745 | 0.036 | 0.944 | 0.034 | 0.948 | 0.948 | 0.034 |
| 998 | P-131-SIT | 58 | F | T3 | 10.188 | 12.218 | 0.037 | 0.922 | 0.049 | 0.913 | 0.922 | 0.037 |
| 999 | P-131-SIT | 58 | F | L1 | 14.646 | 16.036 | 0.041 | 0.921 | 0.041 | 0.921 | 0.921 | 0.041 |
| 1000 | P-131-SIT | 58 | F | L2 | 8.915 | 12.345 | 0.046 | 0.910 | 0.046 | 0.911 | 0.911 | 0.046 |
| 1001 | P-131-SIT | 58 | F | L3 | 9.679 | 10.436 | 0.042 | 0.960 | 0.042 | 0.960 | 0.960 | 0.042 |
| 1002 | P-132-ONG | 50 | M | U2 | 4.458 | 7.254 | 0.025 | 0.928 | 0.025 | 0.930 | 0.930 | 0.025 |
| 1003 | P-132-ONG | 50 | M | U3 | 5.221 | 5.472 | 0.028 | 0.947 | 0.028 | 0.947 | 0.947 | 0.028 |
| 1004 | P-132-ONG | 50 | M | T1 | 6.495 | 9.036 | 0.036 | 0.954 | 0.035 | 0.955 | 0.955 | 0.035 |
| 1005 | P-132-ONG | 50 | M | T2 | 6.114 | 5.727 | 0.032 | 0.934 | 0.032 | 0.947 | 0.947 | 0.032 |
| 1006 | P-132-ONG | 50 | M | T3 | 6.241 | 5.346 | 0.030 | 0.937 | 0.029 | 0.938 | 0.938 | 0.029 |
| 1007 | P-132-ONG | 50 | M | L2 | 5.222 | 4.963 | 0.023 | 0.926 | 0.023 | 0.927 | 0.927 | 0.023 |
| 1008 | P-132-ONG | 50 | M | L3 | 5.731 | 6.237 | 0.019 | 0.958 | 0.019 | 0.959 | 0.959 | 0.019 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{\overline{S_{\text {final }}}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1009 | P-133-SUB | 49 | M | U1 | 14.391 | 21.763 | 0.043 | 0.990 | 0.049 | 0.988 | 0.990 | 0.043 |
| 1010 | P-133-SUB | 49 | M | U2 | 21.906 | 9.164 | 0.080 | 0.994 | 0.085 | 0.993 | 0.994 | 0.080 |
| 1011 | P-133-SUB | 49 | M | U3 | 16.047 | 15.017 | 0.073 | 0.959 | 0.075 | 0.958 | 0.959 | 0.073 |
| 1012 | P-133-SUB | 49 | M | U4 | 34.515 | 27.745 | 0.167 | 0.955 | 0.160 | 0.958 | 0.958 | 0.160 |
| 1013 | P-133-SUB | 49 | M | T1 | 20.251 | 25.199 | 0.079 | 0.949 | 0.083 | 0.940 | 0.949 | 0.079 |
| 1014 | P-133-SUB | 49 | M | T2 | 27.510 | 11.327 | 0.053 | 0.990 | 0.055 | 0.989 | 0.990 | 0.053 |
| 1015 | P-133-SUB | 49 | M | T3 | 15.283 | 11.200 | 0.034 | 0.950 | 0.034 | 0.946 | 0.950 | 0.034 |
| 1016 | P-133-SUB | 49 | M | L1 | 30.949 | 15.018 | 0.128 | 0.943 | 0.111 | 0.956 | 0.956 | 0.111 |
| 1017 | P-133-SUB | 49 | M | L2 | 21.523 | 14.382 | 0.045 | 0.921 | 0.046 | 0.917 | 0.921 | 0.045 |
| 1018 | P-133-SUB | 49 | M | L3 | 19.358 | 13.363 | 0.050 | 0.911 | 0.064 | 0.855 | 0.911 | 0.050 |
| 1019 | P-134-JOH | 62 | M | U1 | 11.717 | 22.526 | 0.052 | 0.986 | 0.045 | 0.989 | 0.989 | 0.045 |
| 1020 | P-134-JOH | 62 | M | U2 | 13.118 | 16.036 | 0.044 | 0.968 | 0.042 | 0.970 | 0.970 | 0.042 |
| 1021 | P-134-JOH | 62 | M | U3 | 13.755 | 16.799 | 0.048 | 0.976 | 0.046 | 0.977 | 0.977 | 0.046 |
| 1022 | P-134-JOH | 62 | M | T1 | 12.100 | 14.381 | 0.036 | 0.925 | 0.036 | 0.922 | 0.925 | 0.036 |
| 1023 | P-134-JOH | 62 | M | T2 | 5.858 | 5.855 | 0.025 | 0.910 | 0.025 | 0.911 | 0.911 | 0.025 |
| 1024 | P-134-JOH | 62 | M | T3 | 4.839 | 6.109 | 0.029 | 0.928 | 0.029 | 0.929 | 0.929 | 0.029 |
| 1025 | P-134-JOH | 62 | M | L1 | 19.868 | 18.073 | 0.122 | 0.952 | 0.114 | 0.958 | 0.958 | 0.114 |
| 1026 | P-134-JOH | 62 | M | L2 | 11.335 | 14.127 | 0.029 | 0.988 | 0.031 | 0.986 | 0.988 | 0.029 |
| 1027 | P-134-JOH | 62 | M | L3 | 20.505 | 16.672 | 0.032 | 0.994 | 0.031 | 0.994 | 0.994 | 0.031 |
| 1028 | P-135-RUS | 44 | M | H | 11.081 | 7.000 | 0.024 | 0.973 | 0.022 | 0.976 | 0.976 | 0.022 |
| 1029 | P-135-RUS | 44 | M | U2 | 5.604 | 5.982 | 0.022 | 0.923 | 0.022 | 0.924 | 0.924 | 0.022 |
| 1030 | P-135-RUS | 44 | M | U3 | 6.877 | 9.291 | 0.033 | 0.967 | 0.033 | 0.967 | 0.967 | 0.033 |
| 1031 | P-135-RUS | 44 | M | T2 | 9.042 | 10.563 | 0.038 | 0.945 | 0.038 | 0.947 | 0.947 | 0.038 |
| 1032 | P-135-RUS | 44 | M | T3 | 12.226 | 12.982 | 0.044 | 0.925 | 0.040 | 0.933 | 0.933 | 0.040 |
| 1033 | P-135-RUS | 44 | M | L3 | 7.132 | 5.855 | 0.055 | 0.920 | 0.055 | 0.920 | 0.920 | 0.055 |
| 1034 | P-136-SAI | 51 | M | U1 | 11.335 | 13.109 | 0.063 | 0.708 | 0.044 | 0.911 | 0.911 | 0.044 |
| 1035 | P-136-SAI | 51 | M | U3 | 13.500 | 18.582 | 0.073 | 0.934 | 0.071 | 0.940 | 0.940 | 0.071 |
| 1036 | P-136-SAI | 51 | M | T1 | 21.015 | 6.109 | 0.154 | 0.916 | 0.134 | 0.932 | 0.932 | 0.134 |
| 1037 | P-136-SAI | 51 | M | T12 | 25.090 | 10.818 | 0.146 | 0.939 | 0.145 | 0.939 | 0.939 | 0.145 |
| 1038 | P-136-SAI | 51 | M | T2 | 10.953 | 16.036 | 0.036 | 0.972 | 0.032 | 0.978 | 0.978 | 0.032 |
| 1039 | P-136-SAI | 51 | M | T3 | 29.038 | 17.182 | 0.078 | 0.981 | 0.074 | 0.983 | 0.983 | 0.074 |
| 1040 | P-136-SAI | 51 | M | L2 | 16.939 | 27.872 | 0.168 | 0.955 | 0.161 | 0.957 | 0.957 | 0.161 |
| 1041 | P-136-SAI | 51 | M | L3 | 12.099 | 15.400 | 0.112 | 0.904 | 0.094 | 0.930 | 0.930 | 0.094 |
| 1042 | P-137-KRI | 57 | M | U1 | 8.915 | 10.436 | 0.036 | 0.948 | 0.036 | 0.949 | 0.949 | 0.036 |
| 1043 | P-137-KRI | 57 | M | U2 | 17.321 | 9.672 | 0.057 | 0.928 | 0.061 | 0.923 | 0.928 | 0.057 |
| 1044 | P-137-KRI | 57 | M | U3 | 19.996 | 18.327 | 0.058 | 0.984 | 0.064 | 0.979 | 0.984 | 0.058 |
| 1045 | P-137-KRI | 57 | M | T1 | 9.425 | 23.800 | 0.042 | 0.977 | 0.040 | 0.979 | 0.979 | 0.040 |
| 1046 | P-137-KRI | 57 | M | T3 | 15.410 | 11.963 | 0.076 | 0.932 | 0.071 | 0.928 | 0.932 | 0.076 |
| 1047 | P-137-KRI | 57 | M | L1 | 26.236 | 14.635 | 0.036 | 0.993 | 0.043 | 0.991 | 0.993 | 0.036 |
| 1048 | P-137-KRI | 57 | M | L2 | 25.472 | 9.418 | 0.027 | 0.994 | 0.033 | 0.992 | 0.994 | 0.027 |
| 1049 | P-137-KRI | 57 | M | L3 | 17.958 | 18.454 | 0.046 | 0.992 | 0.049 | 0.990 | 0.992 | 0.046 |
| 1050 | P-138-BAH | 44 | M | U1 | 18.722 | 14.891 | 0.060 | 0.957 | 0.063 | 0.955 | 0.957 | 0.060 |
| 1051 | P-138-ВАН | 44 | M | U2 | 20.377 | 15.654 | 0.072 | 0.980 | 0.076 | 0.978 | 0.980 | 0.072 |
| 1052 | P-138-BAH | 44 | M | U3 | 15.920 | 13.872 | 0.072 | 0.959 | 0.065 | 0.969 | 0.969 | 0.065 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1053 | P-138-BAH | 44 | M | T1 | 37.189 | 12.727 | 0.078 | 0.960 | 0.075 | 0.963 | 0.963 | 0.075 |
| 1054 | P-138-BAH | 44 | M | T2 | 10.444 | 13.491 | 0.056 | 0.924 | 0.053 | 0.933 | 0.933 | 0.053 |
| 1055 | P-138-BAH | 44 | M | T3 | 15.793 | 28.126 | 0.066 | 0.973 | 0.066 | 0.972 | 0.973 | 0.066 |
| 1056 | P-138-BAH | 44 | M | L1 | 14.391 | 29.272 | 0.070 | 0.989 | 0.078 | 0.986 | 0.989 | 0.070 |
| 1057 | P-138-BAH | 44 | M | L2 | 14.901 | 14.127 | 0.086 | 0.934 | 0.085 | 0.937 | 0.937 | 0.085 |
| 1058 | P-138-BAH | 44 | M | L3 | 14.010 | 16.800 | 0.071 | 0.976 | 0.069 | 0.978 | 0.978 | 0.069 |
| 1059 | P-139-SAR | 27 | F | U1 | 14.264 | 11.454 | 0.031 | 0.987 | 0.031 | 0.987 | 0.987 | 0.031 |
| 1060 | P-139-SAR | 27 | F | U2 | 7.133 | 8.528 | 0.025 | 0.987 | 0.025 | 0.986 | 0.987 | 0.025 |
| 1061 | P-139-SAR | 27 | F | U3 | 21.397 | 12.345 | 0.051 | 0.994 | 0.051 | 0.995 | 0.995 | 0.051 |
| 1062 | P-139-SAR | 27 | F | U32 | 12.863 | 10.182 | 0.049 | 0.989 | 0.049 | 0.988 | 0.989 | 0.049 |
| 1063 | P-139-SAR | 27 | F | T1 | 11.972 | 11.708 | 0.024 | 0.947 | 0.023 | 0.953 | 0.953 | 0.023 |
| 1064 | P-139-SAR | 27 | F | T3 | 5.477 | 5.345 | 0.017 | 0.981 | 0.017 | 0.981 | 0.981 | 0.017 |
| 1065 | P-139-SAR | 27 | F | L1 | 14.264 | 10.436 | 0.022 | 0.958 | 0.021 | 0.959 | 0.959 | 0.021 |
| 1066 | P-139-SAR | 27 | F | L2 | 18.722 | 16.673 | 0.052 | 0.962 | 0.050 | 0.963 | 0.963 | 0.050 |
| 1067 | P-139-SAR | 27 | F | L3 | 5.476 | 7.254 | 0.022 | 0.926 | 0.022 | 0.929 | 0.929 | 0.022 |
| 1068 | P-140-MUR | 25 | M | U1 | 14.137 | 13.236 | 0.039 | 0.951 | 0.037 | 0.955 | 0.955 | 0.037 |
| 1069 | P-140-MUR | 25 | M | U2 | 20.887 | 15.145 | 0.051 | 0.959 | 0.048 | 0.963 | 0.963 | 0.048 |
| 1070 | P-140-MUR | 25 | M | U3 | 10.189 | 23.800 | 0.068 | 0.974 | 0.065 | 0.977 | 0.977 | 0.065 |
| 1071 | P-140-MUR | 25 | M | T1 | 10.443 | 10.818 | 0.019 | 0.944 | 0.021 | 0.931 | 0.944 | 0.019 |
| 1072 | P-140-MUR | 25 | M | T2 | 9.170 | 11.072 | 0.045 | 0.970 | 0.042 | 0.973 | 0.973 | 0.042 |
| 1073 | P-140-MUR | 25 | M | T22 | 7.768 | 10.436 | 0.044 | 0.917 | 0.043 | 0.921 | 0.921 | 0.043 |
| 1074 | P-140-MUR | 25 | M | T3 | 6.622 | 6.746 | 0.021 | 0.967 | 0.022 | 0.966 | 0.967 | 0.021 |
| 1075 | P-140-MUR | 25 | M | L1 | 11.845 | 7.636 | 0.067 | 0.967 | 0.067 | 0.967 | 0.967 | 0.067 |
| 1076 | P-140-MUR | 25 | M | L2 | 35.916 | 10.054 | 0.062 | 0.933 | 0.051 | 0.955 | 0.955 | 0.051 |
| 1077 | P-140-MUR | 25 | M | L3 | 13.882 | 12.472 | 0.055 | 0.924 | 0.049 | 0.930 | 0.930 | 0.049 |
| 1078 | P-141-AHA | 46 | M | H | 13.373 | 7.000 | 0.026 | 0.977 | 0.025 | 0.978 | 0.978 | 0.025 |
| 1079 | P-141-AHA | 46 | M | H2 | 6.368 | 5.473 | 0.020 | 0.971 | 0.020 | 0.971 | 0.971 | 0.020 |
| 1080 | P-141-AHA | 46 | M | U1 | 7.387 | 18.326 | 0.079 | 0.931 | 0.073 | 0.938 | 0.938 | 0.073 |
| 1081 | P-141-AHA | 46 | M | U2 | 10.316 | 11.072 | 0.048 | 0.960 | 0.047 | 0.961 | 0.961 | 0.047 |
| 1082 | P-141-AHA | 46 | M | U3 | 5.986 | 8.400 | 0.043 | 0.917 | 0.044 | 0.912 | 0.917 | 0.043 |
| 1083 | P-141-AHA | 46 | M | T1 | 17.066 | 24.818 | 0.046 | 0.962 | 0.059 | 0.965 | 0.965 | 0.059 |
| 1084 | P-141-AHA | 46 | M | T2 | 21.524 | 16.418 | 0.051 | 0.954 | 0.051 | 0.953 | 0.954 | 0.051 |
| 1085 | P-141-AHA | 46 | M | T3 | 13.373 | 20.363 | 0.043 | 0.946 | 0.043 | 0.946 | 0.946 | 0.043 |
| 1086 | P-141-AHA | 46 | M | T4 | 36.679 | 28.381 | 0.058 | 0.989 | 0.063 | 0.986 | 0.989 | 0.058 |
| 1087 | P-141-AHA | 46 | M | L1 | 11.081 | 24.054 | 0.047 | 0.973 | 0.047 | 0.977 | 0.977 | 0.047 |
| 1088 | P-141-AHA | 46 | M | L2 | 9.934 | 11.836 | 0.038 | 0.954 | 0.038 | 0.954 | 0.954 | 0.038 |
| 1089 | P-141-AHA | 46 | M | L3 | 23.434 | 12.600 | 0.063 | 0.969 | 0.063 | 0.969 | 0.969 | 0.063 |
| 1090 | P-142-MAR | 55 | F | U2 | 4.967 | 8.399 | 0.052 | 0.904 | 0.085 | 0.745 | 0.904 | 0.052 |
| 1091 | P-142-MAR | 55 | F | U3 | 5.222 | 4.327 | 0.035 | 0.976 | 0.036 | 0.975 | 0.976 | 0.035 |
| 1092 | P-142-MAR | 55 | F | L1 | 17.448 | 13.363 | 0.030 | 0.962 | 0.031 | 0.963 | 0.963 | 0.031 |
| 1093 | P-142-MAR | 55 | F | L2 | 6.114 | 6.236 | 0.020 | 0.988 | 0.020 | 0.988 | 0.988 | 0.020 |
| 1094 | P-142-MAR | 55 | F | L3 | 7.642 | 16.545 | 0.082 | 0.950 | 0.071 | 0.958 | 0.958 | 0.071 |
| 1095 | P-143-RAJ | 46 | M | U1 | 9.806 | 9.418 | 0.024 | 0.962 | 0.025 | 0.960 | 0.962 | 0.024 |
| 1096 | P-143-RAJ | 46 | M | U2 | 5.986 | 8.655 | 0.031 | 0.937 | 0.031 | 0.938 | 0.938 | 0.031 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1097 | P-143-RAJ | 46 | M | U3 | 5.476 | 6.364 | 0.040 | 0.967 | 0.040 | 0.966 | 0.967 | 0.040 |
| 1098 | P-143-RAJ | 46 | M | T1 | 14.901 | 12.982 | 0.028 | 0.977 | 0.029 | 0.975 | 0.977 | 0.028 |
| 1099 | P-143-RAJ | 46 | M | T2 | 9.552 | 11.200 | 0.031 | 0.962 | 0.030 | 0.964 | 0.964 | 0.030 |
| 1100 | P-143-RAJ | 46 | M | T3 | 34.769 | 14.763 | 0.046 | 0.970 | 0.046 | 0.970 | 0.970 | 0.046 |
| 1101 | P-143-RAJ | 46 | M | L1 | 6.368 | 9.037 | 0.034 | 0.913 | 0.034 | 0.912 | 0.913 | 0.034 |
| 1102 | P-143-RAJ | 46 | M | L3 | 4.712 | 5.218 | 0.021 | 0.909 | 0.021 | 0.910 | 0.910 | 0.021 |
| 1103 P | P-144-ESM | 63 | F | U1 | 10.571 | 15.909 | 0.059 | 0.972 | 0.057 | 0.949 | 0.972 | 0.059 |
| 1104 P | P-144-ESM | 63 | F | U2 | 7.387 | 17.182 | 0.040 | 0.984 | 0.039 | 0.985 | 0.985 | 0.039 |
| 1105 | P-144-ESM | 63 | F | U3 | 9.807 | 7.255 | 0.084 | 0.954 | 0.078 | 0.960 | 0.960 | 0.078 |
| 1106 | P-144-ESM | 63 | F | L1 | 17.576 | 13.745 | 0.047 | 0.914 | 0.047 | 0.918 | 0.918 | 0.047 |
| 1107 P | P-144-ESM | 63 | F | L2 | 32.094 | 12.345 | 0.034 | 0.950 | 0.033 | 0.954 | 0.954 | 0.033 |
| 1108 | P-144-ESM | 63 | F | L4 | 33.369 | 27.490 | 0.101 | 0.962 | 0.103 | 0.955 | 0.962 | 0.101 |
| 1109 P | P-145-MUH | 26 | M | H | 7.387 | 7.891 | 0.030 | 0.928 | 0.029 | 0.930 | 0.930 | 0.029 |
| 1110 | P-145-MUH | 26 | M | U1 | 4.331 | 4.327 | 0.030 | 0.931 | 0.030 | 0.931 | 0.931 | 0.030 |
| 1111 P | P-145-MUH | 26 | M | U2 | 4.967 | 4.327 | 0.036 | 0.947 | 0.037 | 0.946 | 0.947 | 0.036 |
| 1112 | P-145-MUH | 26 | M | U3 | 5.222 | 5.218 | 0.039 | 0.941 | 0.039 | 0.942 | 0.942 | 0.039 |
| 1113 P | P-145-MUH | 26 | M | T1 | 10.062 | 8.400 | 0.057 | 0.941 | 0.054 | 0.950 | 0.950 | 0.054 |
| 1114 | P-145-MUH | 26 | M | T2 | 8.915 | 7.890 | 0.053 | 0.958 | 0.052 | 0.959 | 0.959 | 0.052 |
| 1115 | -145-MUH | 26 | M | T3 | 12.100 | 8.782 | 0.069 | 0.954 | 0.067 | 0.956 | 0.956 | 0.067 |
| 1116 | P-145-MUH | 26 | M | L1 | 5.094 | 5.472 | 0.042 | 0.937 | 0.040 | 0.944 | 0.944 | 0.040 |
| 1117 | P-145-MUH | 26 | M | L3 | 5.221 | 3.182 | 0.034 | 0.973 | 0.033 | 0.974 | 0.974 | 0.033 |
| 1118 | P-146-SAK | 48 | M | U1 | 12.991 | 12.854 | 0.038 | 0.956 | 0.037 | 0.956 | 0.956 | 0.037 |
| 1119 | P-146-SAK | 48 | M | U2 | 20.504 | 21.254 | 0.064 | 0.980 | 0.068 | 0.976 | 0.980 | 0.064 |
| 1120 | P-146-SAK | 48 | M | U3 | 20.505 | 17.944 | 0.090 | 0.947 | 0.091 | 0.945 | 0.947 | 0.090 |
| 1121 P | P-146-SAK | 48 | M | T1 | 7.515 | 6.618 | 0.033 | 0.929 | 0.033 | 0.929 | 0.929 | 0.033 |
| 1122 | P-146-SAK | 48 | M | T2 | 19.486 | 25.708 | 0.054 | 0.960 | 0.053 | 0.963 | 0.963 | 0.053 |
| 1123 P | P-146-SAK | 48 | M | T3 | 22.925 | 10.436 | 0.039 | 0.956 | 0.038 | 0.959 | 0.959 | 0.038 |
| 1124 | P-146-SAK | 48 | M | L1 | 16.939 | 21.890 | 0.051 | 0.971 | 0.049 | 0.973 | 0.973 | 0.049 |
| 1125 | P-147-THE | 50 | M | U2 | 15.538 | 11.709 | 0.052 | 0.955 | 0.053 | 0.954 | 0.955 | 0.052 |
| 1126 | P-147-THE | 50 | M | T1 | 13.118 | 8.018 | 0.026 | 0.991 | 0.026 | 0.991 | 0.991 | 0.026 |
| 1127 | P-147-THE | 50 | M | T2 | 14.519 | 8.399 | 0.029 | 0.991 | 0.028 | 0.992 | 0.992 | 0.028 |
| 1128 | P-147-THE | 50 | M | T3 | 17.194 | 10.436 | 0.055 | 0.985 | 0.056 | 0.984 | 0.985 | 0.055 |
| 1129 | P-147-THE | 50 | M | T32 | 18.849 | 10.309 | 0.058 | 0.971 | 0.057 | 0.973 | 0.973 | 0.057 |
| 1130 | P-147-THE | 50 | M | L1 | 14.647 | 17.945 | 0.049 | 0.984 | 0.044 | 0.986 | 0.986 | 0.044 |
| 1131 | P-147-THE | 50 | M | L2 | 11.335 | 18.964 | 0.056 | 0.963 | 0.057 | 0.960 | 0.963 | 0.056 |
| 1132 | P-147-THE | 50 | M | L3 | 18.467 | 17.436 | 0.053 | 0.954 | 0.053 | 0.955 | 0.955 | 0.053 |
| 1133 P | P-148-AHM | 33 | M | H | 8.279 | 11.836 | 0.028 | 0.937 | 0.027 | 0.940 | 0.940 | 0.027 |
| 1134 P | P-148-AHM | 33 | M | U1 | 14.010 | 25.708 | 0.055 | 0.976 | 0.056 | 0.975 | 0.976 | 0.055 |
| 1135 P | P-148-AHM | 33 | M | U2 | 17.193 | 10.690 | 0.057 | 0.944 | 0.057 | 0.943 | 0.944 | 0.057 |
| 1136 P | P-148-AHM | 33 | M | U3 | 25.090 | 15.273 | 0.075 | 0.991 | 0.074 | 0.991 | 0.991 | 0.074 |
| 1137 P | P-148-AHM | 33 | M | T1 | 36.425 | 10.691 | 0.049 | 0.985 | 0.051 | 0.984 | 0.985 | 0.049 |
| 1138 P | P-148-AHM | 33 | M | T2 | 22.415 | 16.672 | 0.078 | 0.954 | 0.079 | 0.953 | 0.954 | 0.078 |
| 1139 P | P-148-AHM | 33 | M | T3 | 39.099 | 11.582 | 0.071 | 0.921 | 0.072 | 0.919 | 0.921 | 0.071 |
| 1140 P | P-148-AHM | 33 | M | L1 | 15.028 | 27.490 | 0.097 | 0.961 | 0.090 | 0.966 | 0.966 | 0.090 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1141 | P-148-AHM | 33 | M | L2 | 29.929 | 21.636 | 0.095 | 0.989 | 0.097 | 0.988 | 0.989 | 0.095 |
| 1142 | P-148-AHM | 33 | M | L3 | 13.501 | 27.490 | 0.091 | 0.951 | 0.089 | 0.954 | 0.954 | 0.089 |
| 1143 | P-149-HAS | 28 | M | H | 15.028 | 10.945 | 0.028 | 0.973 | 0.027 | 0.973 | 0.973 | 0.027 |
| 1144 | P-149-HAS | 28 | M | U2 | 3.948 | 5.600 | 0.024 | 0.928 | 0.025 | 0.924 | 0.928 | 0.024 |
| 1145 | P-149-HAS | 28 | M | U3 | 3.439 | 3.691 | 0.019 | 0.962 | 0.018 | 0.962 | 0.962 | 0.018 |
| 1146 | P-149-HAS | 28 | M | T1 | 6.496 | 5.091 | 0.039 | 0.929 | 0.038 | 0.930 | 0.930 | 0.038 |
| 1147 | P-149-HAS | 28 | M | T2 | 4.457 | 4.836 | 0.031 | 0.926 | 0.031 | 0.927 | 0.927 | 0.031 |
| 1148 | P-149-HAS | 28 | M | L1 | 9.170 | 9.799 | 0.031 | 0.926 | 0.032 | 0.927 | 0.927 | 0.032 |
| 1149 | P-149-HAS | 28 | M | L2 | 3.821 | 3.436 | 0.021 | 0.940 | 0.021 | 0.941 | 0.941 | 0.021 |
| 1150 | P-149-HAS | 28 | M | L3 | 4.203 | 5.345 | 0.026 | 0.973 | 0.027 | 0.972 | 0.973 | 0.026 |
| 1151 | $\mathrm{P}-150-\mathrm{MOH}$ | 51 | M | U1 | 4.840 | 5.981 | 0.019 | 0.937 | 0.019 | 0.936 | 0.937 | 0.019 |
| 1152 P | P-150-MOH | 51 | M | U2 | 6.750 | 6.236 | 0.015 | 0.907 | 0.015 | 0.909 | 0.909 | 0.015 |
| 1153 | P-150-MOH | 51 | M | U3 | 2.802 | 2.673 | 0.019 | 0.950 | 0.019 | 0.949 | 0.950 | 0.019 |
| 1154 | P-150-MOH | 51 | M | T1 | 24.071 | 14.508 | 0.046 | 0.973 | 0.046 | 0.973 | 0.973 | 0.046 |
| 1155 | P-150-MOH | 51 | M | T2 | 11.335 | 17.691 | 0.038 | 0.956 | 0.038 | 0.957 | 0.957 | 0.038 |
| 1156 | P-150-MOH | 51 | M | T3 | 11.717 | 19.218 | 0.046 | 0.954 | 0.047 | 0.952 | 0.954 | 0.046 |
| 1157 | $\mathrm{P}-150-\mathrm{MOH}$ | 51 | M | L1 | 24.963 | 10.945 | 0.041 | 0.976 | 0.039 | 0.978 | 0.978 | 0.039 |
| 1158 | $\mathrm{P}-150-\mathrm{MOH}$ | 51 | M | L2 | 16.302 | 16.418 | 0.023 | 0.993 | 0.022 | 0.993 | 0.993 | 0.022 |
| 1159 | P-150-MOH | 51 | M | L3 | 20.887 | 8.145 | 0.024 | 0.993 | 0.023 | 0.994 | 0.994 | 0.023 |
| 1160 | P-151-ROS | 51 | F | H | 4.457 | 3.818 | 0.022 | 0.947 | 0.022 | 0.947 | 0.947 | 0.022 |
| 1161 | P-151-ROS | 51 | F | H2 | 4.203 | 3.945 | 0.029 | 0.857 | 0.021 | 0.901 | 0.901 | 0.021 |
| 1162 | P-151-ROS | 51 | F | U1 | 12.481 | 13.109 | 0.047 | 0.959 | 0.048 | 0.959 | 0.959 | 0.048 |
| 1163 | P-151-ROS | 51 | F | U3 | 16.302 | 13.618 | 0.056 | 0.930 | 0.055 | 0.933 | 0.933 | 0.055 |
| 1164 | P-151-ROS | 51 | F | U4 | 15.538 | 12.090 | 0.047 | 0.965 | 0.045 | 0.968 | 0.968 | 0.045 |
| 1165 | P-151-ROS | 51 | F | U5 | 11.589 | 13.109 | 0.031 | 0.963 | 0.032 | 0.958 | 0.963 | 0.031 |
| 1166 | P-151-ROS | 51 | F | U6 | 15.156 | 16.163 | 0.028 | 0.987 | 0.028 | 0.986 | 0.987 | 0.028 |
| 1167 | P-151-ROS | 51 | F | T1 | 12.736 | 9.545 | 0.033 | 0.987 | 0.032 | 0.988 | 0.988 | 0.032 |
| 1168 | P-151-ROS | 51 | F | T2 | 6.623 | 11.327 | 0.019 | 0.981 | 0.019 | 0.981 | 0.981 | 0.019 |
| 1169 | P-151-ROS | 51 | F | T3 | 14.519 | 15.527 | 0.066 | 0.911 | 0.064 | 0.916 | 0.916 | 0.064 |
| 1170 | P-151-ROS | 51 | F | T4 | 11.462 | 17.563 | 0.041 | 0.983 | 0.042 | 0.982 | 0.983 | 0.041 |
| 1171 | P-151-ROS | 51 | F | T5 | 9.425 | 11.964 | 0.038 | 0.950 | 0.035 | 0.957 | 0.957 | 0.035 |
| 1172 | P-151-ROS | 51 | F | T6 | 21.906 | 12.727 | 0.049 | 0.959 | 0.049 | 0.945 | 0.959 | 0.049 |
| 1173 | P-151-ROS | 51 | F | L1 | 13.882 | 21.000 | 0.056 | 0.976 | 0.057 | 0.959 | 0.976 | 0.056 |
| 1174 | P-151-ROS | 51 | F | L2 | 9.680 | 12.472 | 0.060 | 0.940 | 0.056 | 0.946 | 0.946 | 0.056 |
| 1175 | P-151-ROS | 51 | F | L3 | 11.717 | 14.382 | 0.037 | 0.984 | 0.040 | 0.981 | 0.984 | 0.037 |
| 1176 | P-152-SIT | 59 | F | H | 9.680 | 5.982 | 0.037 | 0.956 | 0.037 | 0.956 | 0.956 | 0.037 |
| 1177 | P-152-SIT | 59 | F | U1 | 14.901 | 16.163 | 0.037 | 0.967 | 0.036 | 0.969 | 0.969 | 0.036 |
| 1178 | P-152-SIT | 59 | F | U2 | 12.227 | 7.763 | 0.049 | 0.942 | 0.049 | 0.942 | 0.942 | 0.049 |
| 1179 | P-152-SIT | 59 | F | U3 | 23.434 | 12.472 | 0.072 | 0.922 | 0.070 | 0.923 | 0.923 | 0.070 |
| 1180 | P-152-SIT | 59 | F | T1 | 6.113 | 10.436 | 0.043 | 0.947 | 0.040 | 0.956 | 0.956 | 0.040 |
| 1181 | P-152-SIT | 59 | F | T2 | 33.623 | 10.564 | 0.050 | 0.932 | 0.056 | 0.914 | 0.932 | 0.050 |
| 1182 | P-152-SIT | 59 | F | T3 | 19.613 | 28.127 | 0.046 | 0.951 | 0.046 | 0.953 | 0.953 | 0.046 |
| 1183 | P-152-SIT | 59 | F | L1 | 11.081 | 16.290 | 0.034 | 0.960 | 0.033 | 0.961 | 0.961 | 0.033 |
| 1184 | P-152-SIT | 59 | F | L12 | 12.609 | 13.490 | 0.036 | 0.973 | 0.040 | 0.964 | 0.973 | 0.036 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{\overline{S_{\text {final }}}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1185 | P-152-SIT | 59 | F | L2 | 9.043 | 8.400 | 0.042 | 0.956 | 0.042 | 0.956 | 0.956 | 0.042 |
| 1186 | P-152-SIT | 59 | F | L3 | 11.972 | 10.690 | 0.062 | 0.959 | 0.062 | 0.959 | 0.959 | 0.062 |
| 1187 | P-153-ZUR | 31 | F | U1 | 4.203 | 4.454 | 0.029 | 0.929 | 0.030 | 0.926 | 0.929 | 0.029 |
| 1188 | P-153-ZUR | 31 | F | U2 | 4.967 | 8.909 | 0.026 | 0.951 | 0.025 | 0.953 | 0.953 | 0.025 |
| 1189 | P-153-ZUR | 31 | F | L1 | 7.005 | 10.563 | 0.017 | 0.995 | 0.017 | 0.995 | 0.995 | 0.017 |
| 1190 | P-153-ZUR | 31 | F | L2 | 6.241 | 5.345 | 0.023 | 0.970 | 0.023 | 0.969 | 0.970 | 0.023 |
| 1191 | P-153-ZUR | 31 | F | L3 | 4.458 | 5.090 | 0.022 | 0.958 | 0.022 | 0.957 | 0.958 | 0.022 |
| 1192 | P-154-GUN | 34 | M | H1 | 6.878 | 7.508 | 0.016 | 0.909 | 0.016 | 0.915 | 0.915 | 0.016 |
| 1193 | P-154-GUN | 34 | M | U1 | 7.514 | 9.290 | 0.022 | 0.961 | 0.020 | 0.964 | 0.964 | 0.020 |
| 1194 | P-154-GUN | 34 | M | U2 | 6.750 | 10.054 | 0.051 | 0.970 | 0.051 | 0.969 | 0.970 | 0.051 |
| 1195 | P-154-GUN | 34 | M | U3 | 5.603 | 4.836 | 0.036 | 0.912 | 0.036 | 0.911 | 0.912 | 0.036 |
| 1196 | P-154-GUN | 34 | M | T1 | 23.944 | 14.000 | 0.066 | 0.944 | 0.067 | 0.942 | 0.944 | 0.066 |
| 1197 | P-154-GUN | 34 | M | T2 | 21.014 | 12.472 | 0.055 | 0.933 | 0.055 | 0.933 | 0.933 | 0.055 |
| 1198 | P-154-GUN | 34 | M | T2C | 30.439 | 12.346 | 0.052 | 0.993 | 0.053 | 0.992 | 0.993 | 0.052 |
| 1199 | P-154-GUN | 34 | M | T2D | 18.849 | 19.599 | 0.045 | 0.939 | 0.047 | 0.935 | 0.939 | 0.045 |
| 1200 | P-154-GUN | 34 | M | T2E | 17.321 | 21.127 | 0.040 | 0.953 | 0.042 | 0.946 | 0.953 | 0.040 |
| 1201 | P-154-GUN | 34 | M | T3 | 5.604 | 6.490 | 0.024 | 0.960 | 0.024 | 0.960 | 0.960 | 0.024 |
| 1202 | P-154-GUN | 34 | M | L1 | 22.543 | 8.781 | 0.030 | 0.984 | 0.030 | 0.985 | 0.985 | 0.030 |
| 1203 | P-154-GUN | 34 | M | L1B | 21.778 | 14.636 | 0.029 | 0.980 | 0.031 | 0.978 | 0.980 | 0.029 |
| 1204 | P-154-GUN | 34 | M | L3 | 11.972 | 19.218 | 0.046 | 0.950 | 0.043 | 0.960 | 0.960 | 0.043 |
| 1205 | P-155-NOR | 41 | M | U1 | 9.425 | 14.508 | 0.023 | 0.986 | 0.023 | 0.987 | 0.987 | 0.023 |
| 1206 | P-155-NOR | 41 | M | U2 | 11.208 | 9.672 | 0.051 | 0.939 | 0.052 | 0.938 | 0.939 | 0.051 |
| 1207 | P-155-NOR | 41 | M | U3 | 8.660 | 7.255 | 0.029 | 0.954 | 0.028 | 0.945 | 0.954 | 0.029 |
| 1208 | P-155-NOR | 41 | M | U4 | 15.410 | 12.218 | 0.037 | 0.970 | 0.037 | 0.969 | 0.970 | 0.037 |
| 1209 | P-155-NOR | 41 | M | T1 | 14.265 | 9.164 | 0.050 | 0.933 | 0.049 | 0.935 | 0.935 | 0.049 |
| 1210 | P-155-NOR | 41 | M | T3 | 11.081 | 5.981 | 0.042 | 0.948 | 0.044 | 0.933 | 0.948 | 0.042 |
| 1211 | P-155-NOR | 41 | M | L4 | 31.076 | 27.363 | 0.061 | 0.964 | 0.085 | 0.969 | 0.969 | 0.085 |
| 1212 | P-155-NOR | 41 | M | L5 | 35.024 | 26.981 | 0.035 | 0.995 | 0.074 | 0.976 | 0.995 | 0.035 |
| 1213 | P-156-HAB | 38 | F | U1 | 6.241 | 6.237 | 0.019 | 0.987 | 0.019 | 0.987 | 0.987 | 0.019 |
| 1214 | P-156-HAB | 38 | F | U2 | 5.732 | 8.527 | 0.025 | 0.946 | 0.025 | 0.947 | 0.947 | 0.025 |
| 1215 | P-156-HAB | 38 | F | U3 | 11.971 | 17.182 | 0.017 | 0.995 | 0.017 | 0.994 | 0.995 | 0.017 |
| 1216 | P-156-HAB | 38 | F | L1 | 7.768 | 10.691 | 0.034 | 0.953 | 0.033 | 0.955 | 0.955 | 0.033 |
| 1217 | P-156-HAB | 38 | F | L2 | 9.424 | 11.200 | 0.022 | 0.980 | 0.021 | 0.982 | 0.982 | 0.021 |
| 1218 | P-156-HAB | 38 | F | L3 | 5.222 | 6.745 | 0.014 | 0.913 | 0.011 | 0.908 | 0.913 | 0.014 |
| 1219 | P-157-SET | 62 | M | U1 | 17.194 | 16.291 | 0.038 | 0.959 | 0.039 | 0.955 | 0.959 | 0.038 |
| 1220 | P-157-SET | 62 | M | U2 | 6.240 | 8.146 | 0.031 | 0.982 | 0.029 | 0.983 | 0.983 | 0.029 |
| 1221 | P-157-SET | 62 | M | U3 | 4.203 | 6.618 | 0.035 | 0.962 | 0.035 | 0.961 | 0.962 | 0.035 |
| 1222 | P-157-SET | 62 | M | T1 | 7.005 | 9.418 | 0.026 | 0.904 | 0.026 | 0.905 | 0.905 | 0.026 |
| 1223 | P-157-SET | 62 | M | T2 | 6.878 | 7.891 | 0.023 | 0.945 | 0.023 | 0.948 | 0.948 | 0.023 |
| 1224 | P-157-SET | 62 | M | L1 | 8.915 | 10.436 | 0.026 | 0.957 | 0.025 | 0.959 | 0.959 | 0.025 |
| 1225 | P-157-SET | 62 | M | L2 | 13.628 | 16.927 | 0.046 | 0.955 | 0.046 | 0.957 | 0.957 | 0.046 |
| 1226 | P-157-SET | 62 | M | L3 | 16.048 | 25.326 | 0.055 | 0.977 | 0.053 | 0.980 | 0.980 | 0.053 |
| 1227 | P-157-SET | 62 | M | L3B | 18.849 | 20.744 | 0.062 | 0.982 | 0.058 | 0.985 | 0.985 | 0.058 |
| 1228 | P-158-SEL | 46 | M | H | 15.920 | 10.818 | 0.041 | 0.929 | 0.045 | 0.917 | 0.929 | 0.041 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1229 | P-158-SEL | 46 | M | U1 | 17.448 | 15.654 | 0.041 | 0.963 | 0.038 | 0.965 | 0.965 | 0.038 |
| 1230 | P-158-SEL | 46 | M | U2 | 16.684 | 14.636 | 0.029 | 0.981 | 0.029 | 0.981 | 0.981 | 0.029 |
| 1231 | P-158-SEL | 46 | M | U3 | 24.580 | 14.509 | 0.040 | 0.967 | 0.039 | 0.969 | 0.969 | 0.039 |
| 1232 | P-158-SEL | 46 | M | T2 | 11.335 | 8.145 | 0.034 | 0.962 | 0.032 | 0.969 | 0.969 | 0.032 |
| 1233 | P-158-SEL | 46 | M | L1 | 9.934 | 8.909 | 0.030 | 0.963 | 0.028 | 0.980 | 0.980 | 0.028 |
| 1234 | P-158-SEL | 46 | M | L3 | 7.387 | 8.782 | 0.023 | 0.974 | 0.022 | 0.975 | 0.975 | 0.022 |
| 1235 | P-159-RAJ | 51 | M | H | 7.387 | 8.273 | 0.016 | 0.984 | 0.017 | 0.984 | 0.984 | 0.017 |
| 1236 | P-159-RAJ | 51 | M | U1 | 21.269 | 6.873 | 0.044 | 0.968 | 0.038 | 0.974 | 0.974 | 0.038 |
| 1237 | P-159-RAJ | 51 | M | U2 | 10.316 | 18.709 | 0.039 | 0.965 | 0.036 | 0.967 | 0.967 | 0.036 |
| 1238 | P-159-RAJ | 51 | M | U3 | 9.552 | 13.236 | 0.025 | 0.994 | 0.028 | 0.992 | 0.994 | 0.025 |
| 1239 | P-159-RAJ | 51 | M | L1 | 16.811 | 15.527 | 0.046 | 0.974 | 0.039 | 0.983 | 0.983 | 0.039 |
| 1240 | P-159-RAJ | 51 | M | L2 | 26.618 | 11.964 | 0.075 | 0.969 | 0.070 | 0.971 | 0.971 | 0.070 |
| 1241 | P-159-RAJ | 51 | M | L3 | 13.755 | 16.290 | 0.053 | 0.969 | 0.052 | 0.968 | 0.969 | 0.053 |
| 1242 | P-160-SAN | 35 | F | H | 11.717 | 9.163 | 0.045 | 0.955 | 0.044 | 0.957 | 0.957 | 0.044 |
| 1243 | P-160-SAN | 35 | F | U1 | 13.627 | 23.545 | 0.033 | 0.984 | 0.031 | 0.986 | 0.986 | 0.031 |
| 1244 | P-160-SAN | 35 | F | U2 | 10.443 | 10.182 | 0.049 | 0.977 | 0.049 | 0.977 | 0.977 | 0.049 |
| 1245 | P-160-SAN | 35 | F | U3 | 21.014 | 7.255 | 0.037 | 0.972 | 0.035 | 0.975 | 0.975 | 0.035 |
| 1246 | P-160-SAN | 35 | F | U4 | 24.198 | 26.726 | 0.050 | 0.979 | 0.052 | 0.980 | 0.980 | 0.052 |
| 1247 | P-160-SAN | 35 | F | T1 | 16.047 | 25.962 | 0.051 | 0.946 | 0.052 | 0.945 | 0.946 | 0.051 |
| 1248 | P-160-SAN | 35 | F | T3 | 4.839 | 3.946 | 0.015 | 0.957 | 0.020 | 0.940 | 0.957 | 0.015 |
| 1249 | P-160-SAN | 35 | F | L1 | 12.482 | 19.090 | 0.035 | 0.980 | 0.034 | 0.980 | 0.980 | 0.034 |
| 1250 | P-160-SAN | 35 | F | L2 | 21.906 | 18.708 | 0.042 | 0.993 | 0.042 | 0.993 | 0.993 | 0.042 |
| 1251 | P-160-SAN | 35 | F | L2B | 18.468 | 18.199 | 0.037 | 0.989 | 0.040 | 0.987 | 0.989 | 0.037 |
| 1252 | P-160-SAN | 35 | F | L3 | 6.368 | 15.272 | 0.041 | 0.929 | 0.040 | 0.931 | 0.931 | 0.040 |
| 1253 | P-160-SAN | 35 | F | L4 | 26.363 | 18.582 | 0.051 | 0.978 | 0.052 | 0.977 | 0.978 | 0.051 |
| 1254 | P-161-RAM | 56 | M | H | 6.113 | 8.655 | 0.026 | 0.965 | 0.026 | 0.964 | 0.965 | 0.026 |
| 1255 | P-161-RAM | 56 | M | U1 | 11.717 | 12.345 | 0.039 | 0.985 | 0.038 | 0.986 | 0.986 | 0.038 |
| 1256 | P-161-RAM | 56 | M | U2 | 5.349 | 8.527 | 0.023 | 0.927 | 0.022 | 0.928 | 0.928 | 0.022 |
| 1257 | P-161-RAM | 56 | M | U3 | 5.222 | 5.981 | 0.030 | 0.940 | 0.031 | 0.940 | 0.940 | 0.031 |
| 1258 | P-161-RAM | 56 | M | T1 | 8.024 | 8.018 | 0.034 | 0.922 | 0.035 | 0.919 | 0.922 | 0.034 |
| 1259 | P-161-RAM | 56 | M | L1 | 7.005 | 7.254 | 0.017 | 0.971 | 0.018 | 0.971 | 0.971 | 0.018 |
| 1260 | P-161-RAM | 56 | M | L2 | 25.599 | 11.582 | 0.064 | 0.975 | 0.059 | 0.979 | 0.979 | 0.059 |
| 1261 P | P-161-RAM | 56 | M | L4 | 31.330 | 27.618 | 0.058 | 0.947 | 0.068 | 0.948 | 0.948 | 0.068 |
| 1262 | P-162-BAL | 41 | M | L22 | 12.226 | 14.763 | 0.075 | 0.979 | 0.061 | 0.985 | 0.985 | 0.061 |
| 1263 P | P-163-NOR | 29 | F | H3 | 10.316 | 10.563 | 0.049 | 0.968 | 0.049 | 0.966 | 0.968 | 0.049 |
| 1264 | P-163-NOR | 29 | F | U22 | 4.840 | 6.490 | 0.028 | 0.961 | 0.028 | 0.961 | 0.961 | 0.028 |
| 1265 | P-163-NOR | 29 | F | U51 | 7.387 | 8.654 | 0.037 | 0.939 | 0.032 | 0.924 | 0.939 | 0.037 |
| 1266 | P-164-LOW | 68 | M | T21 | 11.972 | 16.036 | 0.030 | 0.920 | 0.031 | 0.918 | 0.920 | 0.030 |
| 1267 | P-165-BAL | 48 | M | H2 | 9.043 | 4.964 | 0.050 | 0.699 | 0.035 | 0.911 | 0.911 | 0.035 |
| 1268 | P-165-BAL | 48 | M | T32 | 16.175 | 16.036 | 0.049 | 0.948 | 0.051 | 0.943 | 0.948 | 0.049 |
| 1269 | P-165-BAL | 48 | M | L11 | 19.613 | 20.491 | 0.050 | 0.963 | 0.053 | 0.959 | 0.963 | 0.050 |
| 1270 | P-166-TEO | 58 | M | U31 | 3.439 | 3.054 | 0.016 | 0.940 | 0.016 | 0.939 | 0.940 | 0.016 |
| 1271 | P-167-SAN | 33 | M | H1 | 5.477 | 5.345 | 0.028 | 0.937 | 0.027 | 0.937 | 0.937 | 0.027 |
| 1272 P | P-168-NAG | 43 | M | T42 | 8.279 | 9.418 | 0.026 | 0.940 | 0.027 | 0.936 | 0.940 | 0.026 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{S_{\text {final }}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1273 P | P-168-NAG | 43 | M | T52 | 14.773 | 11.709 | 0.028 | 0.948 | 0.029 | 0.945 | 0.948 | 0.028 |
| 1274 | P-170-SIE | 56 | M | H1 | 6.496 | 6.745 | 0.040 | 0.937 | 0.038 | 0.942 | 0.942 | 0.038 |
| 1275 | P-171-NGS | 64 | M | T11 | 3.439 | 4.073 | 0.022 | 0.912 | 0.023 | 0.912 | 0.912 | 0.023 |
| 1276 | P-172-SUK | 35 | M | L52 | 38.080 | 24.181 | 0.087 | 0.911 | 0.118 | 0.508 | 0.911 | 0.087 |
| 1277 P | P-173-WON | 82 | M | U12 | 6.368 | 4.582 | 0.030 | 0.927 | 0.031 | 0.918 | 0.927 | 0.030 |
| 1278 | P-173-WON | 82 | M | T22 | 6.878 | 13.872 | 0.045 | 0.926 | 0.040 | 0.943 | 0.943 | 0.040 |
| 1279 P | P-173-WON | 82 | M | L22 | 5.222 | 5.473 | 0.036 | 0.973 | 0.036 | 0.973 | 0.973 | 0.036 |
| 1280 P | P-173-WON | 82 | M | L31 | 5.477 | 6.491 | 0.034 | 0.925 | 0.034 | 0.924 | 0.925 | 0.034 |
| 1281 P | P-176-ADH | 34 | M | T11 | 5.604 | 7.636 | 0.042 | 0.811 | 0.032 | 0.911 | 0.911 | 0.032 |
| 1282 P | P-176-ADH | 34 | M | T21 | 5.349 | 7.127 | 0.033 | 0.928 | 0.033 | 0.921 | 0.928 | 0.033 |
| 1283 P | P-176-ADH | 34 | M | L12 | 7.769 | 7.381 | 0.033 | 0.957 | 0.031 | 0.961 | 0.961 | 0.031 |
| 1284 P | P-177-MAH | 45 | M | L31 | 19.996 | 9.418 | 0.092 | 0.921 | 0.093 | 0.921 | 0.921 | 0.093 |
| 1285 | P-179-AAL | 31 | F | U12 | 17.448 | 11.836 | 0.090 | 0.939 | 0.090 | 0.938 | 0.939 | 0.090 |
| 1286 | P-181-NAK | 55 | M | L12 | 11.463 | 15.654 | 0.034 | 0.964 | 0.034 | 0.964 | 0.964 | 0.034 |
| 1287 P | P-182-WON | 68 | M | U32 | 3.820 | 3.818 | 0.022 | 0.918 | 0.022 | 0.918 | 0.918 | 0.022 |
| 1288 | P-183-NER | 64 | M | T21 | 13.882 | 14.254 | 0.052 | 0.938 | 0.052 | 0.939 | 0.939 | 0.052 |
| 1289 P | P-185-LOO | 71 | M | U32 | 7.005 | 6.618 | 0.045 | 0.936 | 0.042 | 0.945 | 0.945 | 0.042 |
| 1290 | P-184-RAT | 28 | F | U22 | 14.774 | 11.327 | 0.076 | 0.902 | 0.077 | 0.903 | 0.903 | 0.077 |
| 1291 P | P-184-RAT | 28 | F | T32 | 14.519 | 15.145 | 0.058 | 0.904 | 0.058 | 0.905 | 0.905 | 0.058 |
| 1292 P | P-187-TAN | 31 | F | T11 | 11.845 | 8.909 | 0.077 | 0.918 | 0.074 | 0.921 | 0.921 | 0.074 |
| 1293 P | P-187-TAN | 31 | F | T32 | 16.047 | 11.709 | 0.045 | 0.942 | 0.043 | 0.945 | 0.945 | 0.043 |
| 1294 P | P-187-TAN | 31 | F | T51 | 7.896 | 8.400 | 0.056 | 0.905 | 0.054 | 0.905 | 0.905 | 0.054 |
| 1295 | P-187-TAN | 31 | F | L22 | 11.589 | 10.054 | 0.039 | 0.930 | 0.038 | 0.933 | 0.933 | 0.038 |
| 1296 | P-187-TAN | 31 | F | L51 | 5.731 | 6.237 | 0.039 | 0.757 | 0.033 | 0.903 | 0.903 | 0.033 |
| 1297 P | P-188-MOH | 28 | M | U11 | 8.533 | 9.927 | 0.039 | 0.945 | 0.038 | 0.949 | 0.949 | 0.038 |
| 1298 P | P-188-MOH | 28 | M | U21 | 2.675 | 3.436 | 0.017 | 0.957 | 0.017 | 0.955 | 0.957 | 0.017 |
| 1299 P | P-188-MOH | 28 | M | L11 | 8.915 | 7.509 | 0.047 | 0.963 | 0.049 | 0.960 | 0.963 | 0.047 |
| 1300 P | P-188-MOH | 28 | M | L31 | 8.661 | 10.181 | 0.034 | 0.908 | 0.041 | 0.904 | 0.908 | 0.034 |
| 1301 P | P-189-MOH | 74 | M | T62 | 10.316 | 4.836 | 0.022 | 0.940 | 0.022 | 0.940 | 0.940 | 0.022 |
| 1302 P | P-189-MOH | 74 | M | T72 | 7.260 | 7.000 | 0.024 | 0.928 | 0.024 | 0.928 | 0.928 | 0.024 |
| 1303 P | P-190-OMA | 28 | M | U21 | 11.335 | 14.127 | 0.039 | 0.945 | 0.037 | 0.951 | 0.951 | 0.037 |
| 1304 P | P-190-OMA | 28 | M | U43 | 40.119 | 29.145 | 0.141 | 0.975 | 0.146 | 0.956 | 0.975 | 0.141 |
| 1305 P | P-190-OMA | 28 | M | T161 | 39.609 | 29.527 | 0.122 | 0.821 | 0.126 | 0.905 | 0.905 | 0.126 |
| 1306 P | P-190-OMA | 28 | M | T21 | 18.467 | 14.254 | 0.082 | 0.922 | 0.083 | 0.921 | 0.922 | 0.082 |
| 1307 P | P-190-OMA | 28 | M | L121 | 36.043 | 28.254 | 0.156 | 0.918 | 0.144 | 0.931 | 0.931 | 0.144 |
| 1308 P | P-190-OMA | 28 | M | L162 | 19.231 | 10.182 | 0.070 | 0.935 | 0.069 | 0.933 | 0.935 | 0.070 |
| 1309 P | P-190-OMA | 28 | M | L172 | 17.703 | 12.727 | 0.079 | 0.774 | 0.094 | 0.928 | 0.928 | 0.094 |
| 1310 | P-191-ALI | 56 | M | U32 | 10.061 | 3.564 | 0.041 | 0.906 | 0.041 | 0.912 | 0.912 | 0.041 |
| 1311 | P-191-ALI | 56 | M | T41 | 10.061 | 10.181 | 0.040 | 0.924 | 0.040 | 0.925 | 0.925 | 0.040 |
| 1312 | P-193-UMI | 50 | F | U22 | 5.349 | 6.491 | 0.025 | 0.972 | 0.026 | 0.971 | 0.972 | 0.025 |
| 1313 | P-193-UMI | 50 | F | T41 | 8.660 | 8.399 | 0.049 | 0.905 | 0.040 | 0.907 | 0.907 | 0.040 |
| 1314 | P-195-DEV | 42 | F | T81 | 23.943 | 28.381 | 0.054 | 0.924 | 0.055 | 0.917 | 0.924 | 0.054 |
| 1315 | P-162-BAL | 41 | M | U31 | 7.769 | 7.128 | 0.035 | 0.948 | 0.033 | 0.955 | 0.955 | 0.033 |
| 1316 | P-167-SAN | 33 | M | U11 | 4.203 | 4.836 | 0.041 | 0.913 | 0.040 | 0.917 | 0.917 | 0.040 |


| No | Patient | Age | Gender | Region | Width (mm) | Height (mm) | $\begin{gathered} S_{a, 2} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{2}^{2}$ | $\begin{gathered} S_{a, 3} \\ (\mathrm{~mm}) \end{gathered}$ | $R_{3}^{2}$ | $R_{\text {final }}^{2}$ | $\begin{aligned} & \overline{\overline{S_{\text {final }}}} \\ & (\mathrm{mm}) \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1317 P | P-168-NAG | 43 | M | T21 | 11.208 | 5.981 | 0.033 | 0.964 | 0.032 | 0.966 | 0.966 | 0.032 |
| 1318 P | P-168-NAG | 43 | M | L11 | 9.679 | 7.763 | 0.024 | 0.973 | 0.023 | 0.975 | 0.975 | 0.023 |
| 1319 | P-169-DUW | 51 | M | U21 | 5.604 | 11.709 | 0.037 | 0.961 | 0.036 | 0.963 | 0.963 | 0.036 |
| 1320 | P-169-DUW | 51 | M | T11 | 16.939 | 13.490 | 0.024 | 0.970 | 0.025 | 0.970 | 0.970 | 0.025 |
| 1321 | P-169-DUW | 51 | M | T21 | 17.321 | 14.636 | 0.021 | 0.926 | 0.025 | 0.901 | 0.926 | 0.021 |
| 1322 P | P-169-DUW | 51 | M | L31 | 16.811 | 16.800 | 0.026 | 0.991 | 0.024 | 0.992 | 0.992 | 0.024 |
| 1323 | P-170-SIE | 56 | M | U11 | 4.840 | 6.873 | 0.039 | 0.969 | 0.037 | 0.971 | 0.971 | 0.037 |
| 1324 | P-170-SIE | 56 | M | U21 | 6.368 | 8.527 | 0.027 | 0.967 | 0.029 | 0.963 | 0.967 | 0.027 |
| 1325 | P-170-SIE | 56 | M | U31 | 5.095 | 4.709 | 0.026 | 0.973 | 0.026 | 0.974 | 0.974 | 0.026 |
| 1326 | P-170-SIE | 56 | M | L21 | 8.024 | 9.672 | 0.033 | 0.923 | 0.032 | 0.926 | 0.926 | 0.032 |
| 1327 | P-170-SIE | 56 | M | L31 | 11.717 | 18.199 | 0.031 | 0.954 | 0.030 | 0.947 | 0.954 | 0.031 |
| 1328 | P-171-NG | 64 | M | U21 | 2.675 | 3.182 | 0.037 | 0.958 | 0.036 | 0.959 | 0.959 | 0.036 |
| 1329 P | P-172-SUK | 35 | M | U21 | 8.151 | 7.254 | 0.031 | 0.951 | 0.032 | 0.940 | 0.951 | 0.031 |
| 1330 | P-173-WON | 82 | M | U31 | 4.713 | 6.364 | 0.029 | 0.936 | 0.027 | 0.942 | 0.942 | 0.027 |
| 1331 | P-173-WON | 82 | M | T11 | 6.368 | 10.690 | 0.040 | 0.930 | 0.038 | 0.939 | 0.939 | 0.038 |
| 1332 | P-173-WON | 82 | M | T31 | 6.495 | 6.873 | 0.051 | 0.927 | 0.050 | 0.928 | 0.928 | 0.050 |
| 1333 | P-173-WON | 82 | M | L11 | 6.368 | 7.127 | 0.032 | 0.980 | 0.033 | 0.979 | 0.980 | 0.032 |
| 1334 | P-174-MAG | 33 | F | L21 | 15.283 | 14.636 | 0.040 | 0.995 | 0.043 | 0.994 | 0.995 | 0.040 |
| 1335 | P-175-DEN | 62 | M | U21 | 7.387 | 8.654 | 0.035 | 0.927 | 0.036 | 0.924 | 0.927 | 0.035 |
| 1336 | P-176-ADH | 34 | M | H1 | 7.132 | 5.218 | 0.038 | 0.943 | 0.039 | 0.942 | 0.943 | 0.038 |
| 1337 | P-176-ADH | 34 | M | U31 | 7.387 | 11.709 | 0.035 | 0.962 | 0.033 | 0.968 | 0.968 | 0.033 |
| 1338 | P-177-MAH | 45 | M | H1 | 9.807 | 10.563 | 0.030 | 0.926 | 0.030 | 0.929 | 0.929 | 0.030 |
| 1339 | P-178-YUR | 29 | F | L61 | 9.934 | 11.581 | 0.035 | 0.933 | 0.034 | 0.937 | 0.937 | 0.034 |
| 1340 | P-180-SEL | 65 | M | U11 | 22.161 | 16.799 | 0.045 | 0.994 | 0.046 | 0.994 | 0.994 | 0.046 |
| 1341 | P-181-NAK | 55 | M | L31 | 8.151 | 12.981 | 0.026 | 0.973 | 0.025 | 0.976 | 0.976 | 0.025 |
| 1342 | P-181-NAK | 55 | M | L41 | 8.788 | 15.399 | 0.035 | 0.956 | 0.031 | 0.957 | 0.957 | 0.031 |
| 1343 | P-182-WON | 68 | M | U21 | 4.967 | 5.727 | 0.029 | 0.971 | 0.029 | 0.970 | 0.971 | 0.029 |
| 1344 | P-185-LOO | 71 | M | T31 | 27.510 | 17.563 | 0.033 | 0.987 | 0.038 | 0.978 | 0.987 | 0.033 |
| 1345 | P-185-LOO | 71 | M | L21 | 9.170 | 15.909 | 0.035 | 0.942 | 0.033 | 0.955 | 0.955 | 0.033 |
| 1346 | P-188-MOH | 28 | M | U31 | 9.552 | 15.782 | 0.034 | 0.980 | 0.032 | 0.982 | 0.982 | 0.032 |
| 1347 | P-192-MUH | 52 | M | T61 | 10.953 | 5.600 | 0.037 | 0.922 | 0.027 | 0.929 | 0.929 | 0.027 |
| 1348 | P-192-MUH | 52 | M | L41 | 8.660 | 9.163 | 0.044 | 0.939 | 0.029 | 0.982 | 0.982 | 0.029 |
| 1349 | P-192-MUH | 52 | M | L51 | 6.240 | 8.145 | 0.029 | 0.985 | 0.029 | 0.985 | 0.985 | 0.029 |
| 1350 | P-193-UMI | 50 | F | L51 | 4.712 | 5.854 | 0.021 | 0.968 | 0.021 | 0.968 | 0.968 | 0.021 |
| 1351 | P-195-DEV | 42 | F | H1 | 6.750 | 12.218 | 0.022 | 0.934 | 0.020 | 0.947 | 0.947 | 0.020 |

## APPENDIX E: SCORING RESULTS BY APPLYING CLUSTERING

## ALGORITHM ON TESTING DATASET

| No | Patient | Age | Gender | Region | $S_{a}(\mathrm{~mm})$ <br> 1st Asm | $S_{a}(\mathrm{~mm})$ <br> nd Asm | $k$-Means <br> 1st Asm | $k$-Means <br> 2nd Asm | FCM <br> 1st Asm | FCM <br> 2nd Asm |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | P-162-BLM | 40 | M | U1 | 0.039 | 0.044 | 2 | 2 | 2 | 2 |
| 2 | P-162-BLM | 40 | M | U2 | 0.017 | 0.016 | 1 | 1 | 1 | 1 |
| 3 | P-162-BLM | 40 | M | U3 | 0.033 | 0.030 | 2 | 1 | 1 | 1 |
| 4 | P-162-BLM | 40 | M | T1 | 0.022 | 0.024 | 1 | 1 | 1 | 1 |
| 5 | P-162-BLM | 40 | M | T2 | 0.026 | 0.020 | 1 | 1 | 1 | 1 |
| 6 | P-162-BLM | 40 | M | T3 | 0.035 | 0.040 | 2 | 2 | 2 | 2 |
| 7 | P-162-BLM | 40 | M | L1 | 0.043 | 0.048 | 2 | 2 | 2 | 2 |
| 8 | P-162-BLM | 40 | M | L2 | 0.063 | 0.064 | 3 | 3 | 3 | 3 |
| 9 | P-163-NAS | 28 | F | H1 | 0.045 | 0.045 | 2 | 2 | 2 | 2 |
| 10 | P-163-NAS | 28 | F | U1 | 0.034 | 0.041 | 2 | 2 | 1 | 2 |
| 11 | P-163-NAS | 28 | F | U4 | 0.041 | 0.038 | 2 | 2 | 2 | 2 |
| 12 | P-163-NAS | 28 | F | T1 | 0.041 | 0.043 | 2 | 2 | 2 | 2 |
| 13 | P-163-NAS | 28 | F | T2 | 0.021 | 0.022 | 1 | 1 | 1 | 1 |
| 14 | P-163-NAS | 28 | F | T3 | 0.036 | 0.033 | 2 | 2 | 2 | 1 |
| 15 | P-163-NAS | 28 | F | L1 | 0.023 | 0.022 | 1 | 1 | 1 | 1 |
| 16 | P-163-NAS | 28 | F | L2 | 0.019 | 0.017 | 1 | 1 | 1 | 1 |
| 17 | P-163-NAS | 28 | F | L3 | 0.034 | 0.037 | 2 | 2 | 1 | 2 |
| 18 | P-163-NAS | 28 | F | L4 | 0.017 | 0.018 | 1 | 1 | 1 | 1 |
| 19 | P-163-NAS | 28 | F | L5 | 0.024 | 0.022 | 1 | 1 | 1 | 1 |
| 20 | P-163-NAS | 28 | F | L6 | 0.026 | 0.020 | 1 | 1 | 1 | 1 |
| 21 | P-163-NAS | 28 | F | L7 | 0.028 | 0.026 | 1 | 1 | 1 | 1 |
| 22 | P-164-LAT | 67 | M | H1 | 0.029 | 0.026 | 1 | 1 | 1 | 1 |
| 23 | P-164-LAT | 67 | M | U2 | 0.014 | 0.014 | 1 | 1 | 1 | 1 |
| 24 | P-164-LAT | 67 | M | T3 | 0.027 | 0.031 | 1 | 1 | 1 | 1 |
| 25 | P-164-LAT | 67 | M | T4 | 0.017 | 0.015 | 1 | 1 | 1 | 1 |
| 26 | P-164-LAT | 67 | M | T6 | 0.018 | 0.019 | 1 | 1 | 1 | 1 |
| 27 | P-164-LAT | 67 | M | L1 | 0.020 | 0.020 | 1 | 1 | 1 | 1 |
| 28 | P-164-LAT | 67 | M | L2 | 0.029 | 0.029 | 1 | 1 | 1 | 1 |
| 29 | P-164-LAT | 67 | M | L3 | 0.010 | 0.010 | 1 | 1 | 1 | 1 |
| 30 | P-165-BLK | 47 | M | U1 | 0.025 | 0.026 | 1 | 1 | 1 | 1 |
| 31 | P-165-BLK | 47 | M | U3 | 0.044 | 0.044 | 2 | 2 | 2 | 2 |
| 32 | P-165-BLK | 47 | M | T1 | 0.037 | 0.035 | 2 | 2 | 2 | 2 |
| 35 | P-165-BLK | 47 | M | T4 | 0.061 | 0.060 | 3 | 3 | 3 | 3 |
|  | P-165-BLK | 47 | M | L1 | 0.043 | 0.042 | 2 | 2 | 2 | 2 |
| 47 | M | L2 | 0.034 | 0.035 | 2 | 2 | 1 | 2 |  |  |


| No | Patient | Age | Gender | Region | $\begin{array}{\|l} \hline S_{a}(\mathrm{~mm}) \\ 1 \mathrm{st} \mathrm{Asm} \\ \hline \end{array}$ | $\begin{array}{\|c\|} \hline S_{a}(\mathrm{~mm}) \\ \text { 2nd Asm } \\ \hline \end{array}$ | $k$-Means 1st Asm | $k$-Means 2nd Asm | $\begin{gathered} \hline \text { FCM } \\ \text { 1st Asm } \\ \hline \end{gathered}$ | FCM <br> 2nd Asm |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 36 | P-165-BLK | 47 | M | L3 | 0.031 | 0.037 | 1 | 2 | 1 | 2 |
| 37 | P-165-BLK | 47 | M | L4 | 0.026 | 0.024 | 1 | 1 | 1 | 1 |
| 38 | P-166-TTL | 57 | M | U2 | 0.033 | 0.029 | 2 | 1 | 1 | 1 |
| 39 | P-166-TTL | 57 | M | T1 | 0.049 | 0.051 | 2 | 3 | 2 | 2 |
| 40 | P-166-TTL | 57 | M | T2 | 0.050 | 0.051 | 2 | 3 | 2 | 2 |
| 41 | P-166-TTL | 57 | M | T3 | 0.042 | 0.041 | 2 | 2 | 2 | 2 |
| 42 | P-166-TTL | 57 | M | T4 | 0.052 | 0.055 | 3 | 3 | 2 | 2 |
| 43 | P-166-TTL | 57 | M | L1 | 0.031 | 0.028 | 1 | 1 | 1 | 1 |
| 44 | P-166-TTL | 57 | M | L2 | 0.021 | 0.020 | 1 | 1 | 1 | 1 |
| 45 | P-166-TTL | 57 | M | L3 | 0.026 | 0.026 | 1 | 1 | 1 | 1 |
| 46 | P-166-TTL | 57 | M | L4 | 0.032 | 0.035 | 2 | 2 | 1 | 2 |
| 47 | P-167-SNR | 32 | M | U2 | 0.035 | 0.038 | 2 | 2 | 2 | 2 |
| 48 | P-167-SNR | 32 | M | L1 | 0.027 | 0.023 | 1 | 1 | 1 | 1 |
| 49 | P-167-SNR | 32 | M | L2 | 0.038 | 0.038 | 2 | 2 | 2 | 2 |
| 50 | P-167-SNR | 32 | M | L3 | 0.033 | 0.026 | 2 | 1 | 1 | 1 |
| 51 | P-167-SNR | 32 | M | L4 | 0.035 | 0.042 | 2 | 2 | 2 | 2 |
| 52 | P-168-NGJ | 42 | M | U1 | 0.019 | 0.021 | 1 | 1 | 1 | 1 |
| 53 | P-168-NGJ | 42 | M | U2 | 0.022 | 0.019 | 1 | 1 | 1 | 1 |
| 54 | P-168-NGJ | 42 | M | U3 | 0.017 | 0.019 | 1 | 1 | 1 | 1 |
| 55 | P-168-NGJ | 42 | M | U4 | 0.076 | 0.078 | 3 | 3 | 3 | 3 |
| 56 | P-168-NGJ | 42 | M | T1 | 0.024 | 0.025 | 1 | 1 | 1 | 1 |
| 57 | P-168-NGJ | 42 | M | T3 | 0.020 | 0.022 | 1 | 1 | 1 | 1 |
| 58 | P-169-DWA | 50 | M | H1 | 0.031 | 0.029 | 1 | 1 | 1 | 1 |
| 59 | P-169-DWA | 50 | M | U1 | 0.051 | 0.055 | 3 | 3 | 2 | 2 |
| 60 | P-169-DWA | 50 | M | T3 | 0.060 | 0.059 | 3 | 3 | 3 | 3 |
| 61 | P-169-DWA | 50 | M | T5 | 0.038 | 0.042 | 2 | 2 | 2 | 2 |
| 62 | P-169-DWA | 50 | M | T6 | 0.043 | 0.038 | 2 | 2 | 2 | 2 |
| 63 | P-169-DWA | 50 | M | L5 | 0.036 | 0.035 | 2 | 2 | 2 | 2 |
| 64 | P-170-SSG | 55 | M | T1 | 0.022 | 0.022 | 1 | 1 | 1 | 1 |
| 65 | P-170-SSG | 55 | M | T1 | 0.016 | 0.015 | 1 | 1 | 1 | 1 |
| 66 | P-170-SSG | 55 | M | T2 | 0.022 | 0.024 | 1 | 1 | 1 | 1 |
| 67 | P-170-SSG | 55 | M | T3 | 0.040 | 0.042 | 2 | 2 | 2 | 2 |
| 68 | P-170-SSG | 55 | M | T4 | 0.041 | 0.045 | 2 | 2 | 2 | 2 |
| 69 | P-170-SSG | 55 | M | T5 | 0.048 | 0.044 | 2 | 2 | 2 | 2 |
| 70 | P-170-SSG | 55 | M | T7 | 0.096 | 0.096 | 4 | 4 | 3 | 3 |
| 71 | P-170-SSG | 55 | M | L1 | 0.045 | 0.040 | 2 | 2 | 2 | 2 |
| 72 | P-170-SSG | 55 | M | L1 | 0.029 | 0.030 | 1 | 1 | 1 | 1 |
| 73 | P-170-SSG | 55 | M | L1 | 0.065 | 0.068 | 3 | 3 | 3 | 3 |
| 74 | P-170-SSG | 55 | M | L2 | 0.032 | 0.028 | 2 | 1 | 1 | 1 |
| 75 | P-170-SSG | 55 | M | L6 | 0.052 | 0.051 | 3 | 3 | 2 | 2 |


| No | Patient | Age | Gender | Region | $\begin{aligned} & \hline S_{a}(\mathrm{~mm}) \\ & 1 \mathrm{st} \text { Asm } \end{aligned}$ | $\left\lvert\, \begin{gathered} S_{a}(\mathrm{~mm}) \\ \text { 2nd Asm } \end{gathered}\right.$ | $k$-Means 1st Asm | $k$-Means 2nd Asm | FCM <br> 1st Asm | FCM <br> 2nd Asm |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 76 | P-170-SSG | 55 | M | L7 | 0.067 | 0.068 | 3 | 3 | 3 | 3 |
| 77 | P-170-SSG | 55 | M | L8 | 0.058 | 0.055 | 3 | 3 | 3 | 2 |
| 78 | P-171-NSH | 63 | M | U1 | 0.024 | 0.026 | 1 | 1 | 1 | 1 |
| 79 | P-171-NSH | 63 | M | U3 | 0.018 | 0.016 | 1 | 1 | 1 | 1 |
| 80 | P-171-NSH | 63 | M | L1 | 0.032 | 0.033 | 2 | 2 | 1 | 1 |
| 81 | P-171-NSH | 63 | M | L3 | 0.026 | 0.023 | 1 | 1 | 1 | 1 |
| 82 | P-171-NSH | 63 | M | L4 | 0.021 | 0.024 | 1 | 1 | 1 | 1 |
| 83 | P-171-NSH | 63 | M | L5 | 0.038 | 0.040 | 2 | 2 | 2 | 2 |
| 84 | P-172-SKM | 34 | M | H1 | 0.049 | 0.048 | 2 | 2 | 2 | 2 |
| 85 | P-172-SKM | 34 | M | U1 | 0.037 | 0.044 | 2 | 2 | 2 | 2 |
| 86 | P-172-SKM | 34 | M | U2 | 0.031 | 0.034 | 1 | 2 | 1 | 1 |
| 87 | P-172-SKM | 34 | M | U3 | 0.036 | 0.035 | 2 | 2 | 2 | 2 |
| 88 | P-172-SKM | 34 | M | U4 | 0.079 | 0.079 | 3 | 3 | 3 | 3 |
| 89 | P-172-SKM | 34 | M | U5 | 0.061 | 0.055 | 3 | 3 | 3 | 2 |
| 90 | P-172-SKM | 34 | M | U6 | 0.073 | 0.074 | 3 | 3 | 3 | 3 |
| 91 | P-172-SKM | 34 | M | U8 | 0.077 | 0.073 | 3 | 3 | 3 | 3 |
| 92 | P-172-SKM | 34 | M | T1 | 0.033 | 0.033 | 2 | 2 | 1 | 1 |
| 93 | P-172-SKM | 34 | M | T2 | 0.053 | 0.060 | 3 | 3 | 2 | 3 |
| 94 | P-172-SKM | 34 | M | T3 | 0.035 | 0.041 | 2 | 2 | 2 | 2 |
| 95 | P-172-SKM | 34 | M | T4 | 0.102 | 0.097 | 4 | 4 | 4 | 3 |
| 96 | P-172-SKM | 34 | M | T5 | 0.078 | 0.075 | 3 | 3 | 3 | 3 |
| 97 | P-172-SKM | 34 | M | L1 | 0.075 | 0.077 | 3 | 3 | 3 | 3 |
| 98 | P-172-SKM | 34 | M | L2 | 0.039 | 0.035 | 2 | 2 | 2 | 2 |
| 99 | P-172-SKM | 34 | M | L3 | 0.048 | 0.048 | 2 | 2 | 2 | 2 |
| 100 | P-172-SKM | 34 | M | L6 | 0.054 | 0.054 | 3 | 3 | 2 | 2 |
| 101 | P-172-SKM | 34 | M | L7 | 0.041 | 0.040 | 2 | 2 | 2 | 2 |
| 102 | P-173-WKW | 81 | M | U3 | 0.027 | 0.033 | 1 | 2 | 1 | 1 |
| 103 | P-173-WKW | 81 | M | L1 | 0.032 | 0.030 | 2 | 1 | 1 | 1 |
| 104 | P-174-MGH | 32 | F | U1 | 0.042 | 0.041 | 2 | 2 | 2 | 2 |
| 105 | P-174-MGH | 32 | F | U2 | 0.051 | 0.051 | 3 | 3 | 2 | 2 |
| 106 | P-174-MGH | 32 | F | U3 | 0.025 | 0.027 | 1 | 1 | 1 | 1 |
| 107 | P-174-MGH | 32 | F | T2 | 0.027 | 0.031 | 1 | 1 | 1 | 1 |
| 108 | P-174-MGH | 32 | F | L1 | 0.059 | 0.059 | 3 | 3 | 3 | 3 |
| 109 | P-174-MGH | 32 | F | L3 | 0.062 | 0.060 | 3 | 3 | 3 | 3 |
| 110 | P-175-DNH | 61 | M | H1 | 0.031 | 0.026 | 1 | 1 | 1 | 1 |
| 111 | P-175-DNH | 61 | M | U1 | 0.030 | 0.025 | 1 | 1 | 1 | 1 |
| 112 | P-175-DNH | 61 | M | U3 | 0.032 | 0.036 | 2 | 2 | 1 | 2 |
| 113 | P-175-DNH | 61 | M | T2 | 0.054 | 0.053 | 3 | 3 | 2 | 2 |
| 114 | P-175-DNH | 61 | M | L1 | 0.034 | 0.034 | 2 | 2 | 1 | 1 |
| 115 | P-175-DNH | 61 | M | L2 | 0.050 | 0.046 | 2 | 2 | 2 | 2 |


| No | Patient | Age | Gender | Region | $\begin{array}{\|l\|} \hline S_{a}(\mathrm{~mm}) \\ 1 \mathrm{st} \mathrm{Asm} \\ \hline \end{array}$ | $\begin{array}{\|c\|} \hline S_{a}(\mathrm{~mm}) \\ \text { 2nd Asm } \\ \hline \end{array}$ | $k$-Means 1st Asm | $k$-Means 2nd Asm | $\begin{gathered} \hline \text { FCM } \\ \text { 1st Asm } \\ \hline \end{gathered}$ | FCM <br> 2nd Asm |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 116 | P-176-MDA | 33 | M | U3 | 0.033 | 0.029 | 2 | 1 | 1 | 1 |
| 117 | P-176-MDA | 33 | M | U4 | 0.033 | 0.029 | 2 | 1 | 1 | 1 |
| 118 | P-176-MDA | 33 | M | L2 | 0.022 | 0.016 | 1 | 1 | 1 | 1 |
| 119 | P-176-MDA | 33 | M | L3 | 0.031 | 0.030 | 1 | 1 | 1 | 1 |
| 120 | P-177-MHD | 44 | M | H1 | 0.030 | 0.036 | 1 | 2 | 1 | 2 |
| 121 | P-177-MHD | 44 | M | U2 | 0.061 | 0.063 | 3 | 3 | 3 | 3 |
| 122 | P-177-MHD | 44 | M | U3 | 0.067 | 0.062 | 3 | 3 | 3 | 3 |
| 123 | P-177-MHD | 44 | M | T5 | 0.062 | 0.065 | 3 | 3 | 3 | 3 |
| 124 | P-177-MHD | 44 | M | L1 | 0.056 | 0.061 | 3 | 3 | 2 | 3 |
| 125 | P-177-MHD | 44 | M | L2 | 0.053 | 0.048 | 3 | 2 | 2 | 2 |
| 126 | P-177-MHD | 44 | M | L4 | 0.041 | 0.040 | 2 | 2 | 2 | 2 |
| 127 | P-177-MHD | 44 | M | L6 | 0.086 | 0.080 | 4 | 3 | 3 | 3 |
| 128 | P-177-MHD | 44 | M | L7 | 0.066 | 0.070 | 3 | 3 | 3 | 3 |
| 129 | P-178-YRL | 28 | F | H1 | 0.032 | 0.036 | 2 | 2 | 1 | 2 |
| 130 | P-178-YRL | 28 | F | U1 | 0.034 | 0.039 | 2 | 2 | 1 | 2 |
| 131 | P-178-YRL | 28 | F | U2 | 0.049 | 0.044 | 2 | 2 | 2 | 2 |
| 132 | P-178-YRL | 28 | F | U3 | 0.036 | 0.041 | 2 | 2 | 2 | 2 |
| 133 | P-178-YRL | 28 | F | T1 | 0.112 | 0.114 | 4 | 4 | 4 | 4 |
| 134 | P-178-YRL | 28 | F | T3 | 0.082 | 0.081 | 3 | 3 | 3 | 3 |
| 135 | P-178-YRL | 28 | F | L1 | 0.039 | 0.042 | 2 | 2 | 2 | 2 |
| 136 | P-178-YRL | 28 | F | L2 | 0.035 | 0.037 | 2 | 2 | 2 | 2 |
| 137 | P-178-YRL | 28 | F | L3 | 0.037 | 0.039 | 2 | 2 | 2 | 2 |
| 138 | P-178-YRL | 28 | F | L6 | 0.034 | 0.029 | 2 | 1 | 1 | 1 |
| 139 | P-179-AMI | 30 | F | H1 | 0.024 | 0.031 | 1 | 1 | 1 | 1 |
| 140 | P-179-AMI | 30 | F | U2 | 0.028 | 0.025 | 1 | 1 | 1 | 1 |
| 141 | P-179-AMI | 30 | F | U3 | 0.056 | 0.052 | 3 | 3 | 2 | 2 |
| 142 | P-179-AMI | 30 | F | T2 | 0.041 | 0.036 | 2 | 2 | 2 | 2 |
| 143 | P-179-AMI | 30 | F | L1 | 0.039 | 0.036 | 2 | 2 | 2 | 2 |
| 144 | P-179-AMI | 30 | F | L2 | 0.044 | 0.046 | 2 | 2 | 2 | 2 |
| 145 | P-180-SLV | 64 | M | U2 | 0.077 | 0.073 | 3 | 3 | 3 | 3 |
| 146 | P-180-SLV | 64 | M | U4 | 0.094 | 0.090 | 4 | 4 | 3 | 3 |
| 147 | P-180-SLV | 64 | M | U5 | 0.047 | 0.048 | 2 | 2 | 2 | 2 |
| 148 | P-180-SLV | 64 | M | T1 | 0.050 | 0.051 | 2 | 3 | 2 | 2 |
| 149 | P-180-SLV | 64 | M | T1 | 0.071 | 0.065 | 3 | 3 | 3 | 3 |
| 150 | P-180-SLV | 64 | M | T1 | 0.080 | 0.085 | 3 | 4 | 3 | 3 |
| 151 | P-180-SLV | 64 | M | T2 | 0.061 | 0.066 | 3 | 3 | 3 | 3 |
| 152 | P-180-SLV | 64 | M | T3 | 0.071 | 0.075 | 3 | 3 | 3 | 3 |
| 153 | P-180-SLV | 64 | M | T4 | 0.074 | 0.075 | 3 | 3 | 3 | 3 |
| 154 | P-180-SLV | 64 | M | T5 | 0.112 | 0.106 | 4 | 4 | 4 | 4 |
| 155 | P-180-SLV | 64 | M | T6 | 0.069 | 0.076 | 3 | 3 | 3 | 3 |


| No | Patient | Age | Gender | Region | $\begin{array}{\|l\|} \hline S_{a}(\mathrm{~mm}) \\ 1 \text { st Asm } \end{array}$ | $\begin{array}{\|c\|} \hline \begin{array}{c} S_{a}(\mathrm{~mm}) \\ \text { 2nd Asm } \end{array} \\ \hline \end{array}$ | $k$-Means 1st Asm | $k$-Means 2nd Asm | $\begin{gathered} \text { FCM } \\ \text { 1st Asm } \end{gathered}$ | FCM 2nd Asm |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 156 | P-180-SLV | 64 | M | T7 | 0.050 | 0.047 | 2 | 2 | 2 | 2 |
| 157 | P-180-SLV | 64 | M | T8 | 0.056 | 0.051 | 3 | 3 | 2 | 2 |
| 158 | P-180-SLV | 64 | M | L1 | 0.043 | 0.042 | 2 | 2 | 2 | 2 |
| 159 | P-180-SLV | 64 | M | L1 | 0.038 | 0.041 | 2 | 2 | 2 | 2 |
| 160 | P-180-SLV | 64 | M | L1 | 0.036 | 0.040 | 2 | 2 | 2 | 2 |
| 161 | P-180-SLV | 64 | M | L2 | 0.039 | 0.032 | 2 | 2 | 2 | 1 |
| 162 | P-180-SLV | 64 | M | L3 | 0.030 | 0.030 | 1 | 1 | 1 | 1 |
| 163 | P-180-SLV | 64 | M | L5 | 0.076 | 0.079 | 3 | 3 | 3 | 3 |
| 164 | P-180-SLV | 64 | M | L6 | 0.050 | 0.054 | 2 | 3 | 2 | 2 |
| 165 | P-181-NKR | 55 | M | H1 | 0.020 | 0.023 | 1 | 1 | 1 | 1 |
| 166 | P-181-NKR | 55 | M | U2 | 0.018 | 0.020 | 1 | 1 | 1 | 1 |
| 167 | P-181-NKR | 55 | M | T1 | 0.033 | 0.034 | 2 | 2 | 1 | 1 |
| 168 | P-181-NKR | 55 | M | T2 | 0.048 | 0.051 | 2 | 3 | 2 | 2 |
| 169 | P-181-NKR | 55 | M | T3 | 0.019 | 0.022 | 1 | 1 | 1 | 1 |
| 170 | P-181-NKR | 55 | M | T4 | 0.031 | 0.031 | 1 | 1 | 1 | 1 |
| 171 | P-181-NKR | 55 | M | T5 | 0.018 | 0.017 | 1 | 1 | 1 | 1 |
| 172 | P-181-NKR | 55 | M | T6 | 0.032 | 0.037 | 2 | 2 | 1 | 2 |
| 173 | P-181-NKR | 55 | M | T7 | 0.029 | 0.029 | 1 | 1 | 1 | 1 |
| 174 | P-181-NKR | 55 | M | L1 | 0.026 | 0.025 | 1 | 1 | 1 | 1 |
| 175 | P-181-NKR | 55 | M | L2 | 0.022 | 0.015 | 1 | 1 | 1 | 1 |
| 176 | P-181-NKR | 55 | M | L3 | 0.025 | 0.032 | 1 | 2 | 1 | 1 |
| 177 | P-181-NKR | 55 | M | L4 | 0.031 | 0.035 | 1 | 2 | 1 | 2 |
| 178 | P-182-WSS | 68 | M | U1 | 0.025 | 0.025 | 1 | 1 | 1 | 1 |
| 179 | P-182-WSS | 68 | M | U2 | 0.029 | 0.033 | 1 | 2 | 1 | 1 |
| 180 | P-182-WSS | 68 | M | T1 | 0.050 | 0.045 | 2 | 2 | 2 | 2 |
| 181 | P-182-WSS | 68 | M | T3 | 0.017 | 0.021 | 1 | 1 | 1 | 1 |
| 182 | P-182-WSS | 68 | M | T4 | 0.022 | 0.025 | 1 | 1 | 1 | 1 |
| 183 | P-182-WSS | 68 | M | T5 | 0.060 | 0.060 | 3 | 3 | 3 | 3 |
| 184 | P-182-WSS | 68 | M | T6 | 0.041 | 0.040 | 2 | 2 | 2 | 2 |
| 185 | P-182-WSS | 68 | M | L1 | 0.029 | 0.030 | 1 | 1 | 1 | 1 |
| 186 | P-182-WSS | 68 | M | L2 | 0.028 | 0.027 | 1 | 1 | 1 | 1 |
| 187 | P-183-NRM | 64 | M | T1 | 0.040 | 0.036 | 2 | 2 | 2 | 2 |
| 188 | P-183-NRM | 64 | M | T4 | 0.052 | 0.050 | 3 | 2 | 2 | 2 |
| 189 | P-183-NRM | 64 | M | L1 | 0.060 | 0.059 | 3 | 3 | 3 | 3 |
| 190 | P-183-NRM | 64 | M | L2 | 0.038 | 0.036 | 2 | 2 | 2 | 2 |
| 191 | P-183-NRM | 64 | M | L3 | 0.028 | 0.029 | 1 | 1 | 1 | 1 |
| 192 | P-185-LHM | 71 | M | H1 | 0.025 | 0.027 | 1 | 1 | 1 | 1 |
| 193 | P-185-LHM | 71 | M | U2 | 0.046 | 0.044 | 2 | 2 | 2 | 2 |
| 194 | P-185-LHM | 71 | M | T2 | 0.036 | 0.035 | 2 | 2 | 2 | 2 |
| 195 | P-185-LHM | 71 | M | T3 | 0.033 | 0.028 | 2 | 1 | 1 | 1 |


| No | Patient | Age | Gender | Region | $\begin{aligned} & \hline S_{a}(\mathrm{~mm}) \\ & 1 \text { st Asm } \\ & \hline \end{aligned}$ | $\begin{array}{\|c\|} \hline S_{a}(\mathrm{~mm}) \\ \text { 2nd Asm } \end{array}$ | $k$-Means 1st Asm | $k$-Means 2nd Asm | FCM <br> 1st Asm | FCM <br> 2nd Asm |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 196 | P-185-LHM | 71 | M | L1 | 0.049 | 0.046 | 2 | 2 | 2 | 2 |
| 197 | P-185-LHM | 71 | M | L2 | 0.033 | 0.031 | 2 | 1 | 1 | 1 |
| 198 | P-185-LHM | 71 | M | L3 | 0.032 | 0.034 | 2 | 2 | 1 | 1 |
| 199 | P-184-RTV | 28 | F | H1 | 0.087 | 0.084 | 4 | 4 | 3 | 3 |
| 200 | P-184-RTV | 28 | F | U3 | 0.065 | 0.071 | 3 | 3 | 3 | 3 |
| 201 | P-184-RTV | 28 | F | T1 | 0.124 | 0.129 | 4 | 4 | 4 | 4 |
| 202 | P-184-RTV | 28 | F | T4 | 0.111 | 0.114 | 4 | 4 | 4 | 4 |
| 203 | P-184-RTV | 28 | F | T5 | 0.158 | 0.160 | 4 | 4 | 4 | 4 |
| 204 | P-184-RTV | 28 | F | T8 | 0.068 | 0.074 | 3 | 3 | 3 | 3 |
| 205 | P-184-RTV | 28 | F | T9 | 0.182 | 0.184 | 4 | 4 | 4 | 4 |
| 206 | P-184-RTV | 28 | F | L2 | 0.037 | 0.038 | 2 | 2 | 2 | 2 |
| 207 | P-184-RTV | 28 | F | L4 | 0.072 | 0.069 | 3 | 3 | 3 | 3 |
| 208 | P-184-RTV | 28 | F | L5 | 0.035 | 0.034 | 2 | 2 | 2 | 1 |
| 209 | P-184-RTV | 28 | F | L6 | 0.057 | 0.056 | 3 | 3 | 3 | 2 |
| 210 | P-184-RTV | 28 | F | L7 | 0.033 | 0.033 | 2 | 2 | 1 | 1 |
| 211 | P-184-RTV | 28 | F | L9 | 0.056 | 0.054 | 3 | 3 | 2 | 2 |
| 212 | P-186-HSM | 52 | M | H1 | 0.027 | 0.021 | 1 | 1 | 1 | 1 |
| 213 | P-186-HSM | 52 | M | U1 | 0.044 | 0.044 | 2 | 2 | 2 | 2 |
| 214 | P-186-HSM | 52 | M | U3 | 0.035 | 0.035 | 2 | 2 | 2 | 2 |
| 215 | P-186-HSM | 52 | M | T1 | 0.039 | 0.036 | 2 | 2 | 2 | 2 |
| 216 | P-186-HSM | 52 | M | T2 | 0.037 | 0.035 | 2 | 2 | 2 | 2 |
| 217 | P-186-HSM | 52 | M | T3 | 0.035 | 0.034 | 2 | 2 | 2 | 1 |
| 218 | P-186-HSM | 52 | M | L1 | 0.056 | 0.053 | 3 | 3 | 2 | 2 |
| 219 | P-186-HSM | 52 | M | L2 | 0.036 | 0.039 | 2 | 2 | 2 | 2 |
| 220 | P-186-HSM | 52 | M | L3 | 0.043 | 0.045 | 2 | 2 | 2 | 2 |
| 221 | P-186-HSM | 52 | M | U2 | 0.041 | 0.036 | 2 | 2 | 2 | 2 |
| 222 | P-186-HSM | 52 | M | U3 | 0.037 | 0.035 | 2 | 2 | 2 | 2 |
| 223 | P-187-TLC | 31 | F | U4 | 0.052 | 0.046 | 3 | 2 | 2 | 2 |
| 224 | P-187-TLC | 31 | F | T4 | 0.048 | 0.048 | 2 | 2 | 2 | 2 |
| 225 | P-187-TLC | 31 | F | L1 | 0.040 | 0.039 | 2 | 2 | 2 | 2 |
| 226 | P-187-TLC | 31 | F | L3 | 0.047 | 0.042 | 2 | 2 | 2 | 2 |
| 227 | P-187-TLC | 31 | F | L4 | 0.035 | 0.040 | 2 | 2 | 2 | 2 |
| 228 | P-187-TLC | 31 | F | L6 | 0.041 | 0.043 | 2 | 2 | 2 | 2 |
| 229 | P-187-TLC | 31 | F | L7 | 0.079 | 0.073 | 3 | 3 | 3 | 3 |
| 230 | P-188-MNJ | 28 | M | U3 | 0.032 | 0.031 | 2 | 1 | 1 | 1 |
| 231 | P-188-MNJ | 28 | M | T1 | 0.040 | 0.041 | 2 | 2 | 2 | 2 |
| 232 | P-188-MNJ | 28 | M | T3 | 0.039 | 0.043 | 2 | 2 | 2 | 2 |
| 233 | P-188-MNJ | 28 | M | T4 | 0.026 | 0.031 | 1 | 1 | 1 | 1 |
| 234 | P-188-MNJ | 28 | M | T5 | 0.039 | 0.038 | 2 | 2 | 2 | 2 |
| 235 | P-188-MNJ | 28 | M | L2 | 0.021 | 0.023 | 1 | 1 | 1 | 1 |


| No | Patient | Age | Gender | Region | $\begin{array}{\|l} \hline S_{a}(\mathrm{~mm}) \\ 1 \text { st Asm } \end{array}$ | $\begin{array}{\|c\|} \hline S_{a}(\mathrm{~mm}) \\ \text { 2nd Asm } \end{array}$ | $k$-Means 1st Asm | $k$-Means <br> 2nd Asm | $\begin{gathered} \text { FCM } \\ \text { 1st Asm } \end{gathered}$ | $\begin{gathered} \text { FCM } \\ \text { 2nd Asm } \end{gathered}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 236 | P-189-MYM | 74 | M | T1 | 0.020 | 0.021 | 1 | 1 | 1 | 1 |
| 237 | P-189-MYM | 74 | M | T3 | 0.073 | 0.075 | 3 | 3 | 3 | 3 |
| 238 | P-189-MYM | 74 | M | T4 | 0.021 | 0.021 | 1 | 1 | 1 | 1 |
| 239 | P-189-MYM | 74 | M | T5 | 0.017 | 0.017 | 1 | 1 | 1 | 1 |
| 240 | P-189-MYM | 74 | M | T8 | 0.019 | 0.017 | 1 | 1 | 1 | 1 |
| 241 | P-189-MYM | 74 | M | T9 | 0.022 | 0.025 | 1 | 1 | 1 | 1 |
| 242 | P-190-OMR | 28 | M | H1 | 0.038 | 0.033 | 2 | 2 | 2 | 1 |
| 243 | P-190-OMR | 28 | M | U1 | 0.130 | 0.129 | 4 | 4 | 4 | 4 |
| 244 | P-190-OMR | 28 | M | U1 | 0.101 | 0.096 | 4 | 4 | 4 | 3 |
| 245 | P-190-OMR | 28 | M | U4 | 0.141 | 0.138 | 4 | 4 | 4 | 4 |
| 246 | P-190-OMR | 28 | M | U5 | 0.054 | 0.060 | 3 | 3 | 2 | 3 |
| 247 | P-190-OMR | 28 | M | T1 | 0.113 | 0.115 | 4 | 4 | 4 | 4 |
| 248 | P-190-OMR | 28 | M | T1 | 0.119 | 0.120 | 4 | 4 | 4 | 4 |
| 249 | P-190-OMR | 28 | M | T2 | 0.100 | 0.101 | 4 | 4 | 4 | 4 |
| 250 | P-190-OMR | 28 | M | T2 | 0.182 | 0.187 | 4 | 4 | 4 | 4 |
| 251 | P-190-OMR | 28 | M | T2 | 0.139 | 0.142 | 4 | 4 | 4 | 4 |
| 252 | P-190-OMR | 28 | M | T2 | 0.137 | 0.131 | 4 | 4 | 4 | 4 |
| 253 | P-190-OMR | 28 | M | T3 | 0.055 | 0.055 | 3 | 3 | 2 | 2 |
| 254 | P-190-OMR | 28 | M | T6 | 0.097 | 0.101 | 4 | 4 | 3 | 4 |
| 255 | P-190-OMR | 28 | M | T7 | 0.116 | 0.121 | 4 | 4 | 4 | 4 |
| 256 | P-190-OMR | 28 | M | T8 | 0.081 | 0.081 | 3 | 3 | 3 | 3 |
| 257 | P-190-OMR | 28 | M | L1 | 0.100 | 0.106 | 4 | 4 | 4 | 4 |
| 258 | P-190-OMR | 28 | M | L1 | 0.079 | 0.086 | 3 | 4 | 3 | 3 |
| 259 | P-190-OMR | 28 | M | L1 | 0.112 | 0.116 | 4 | 4 | 4 | 4 |
| 260 | P-190-OMR | 28 | M | L1 | 0.065 | 0.058 | 3 | 3 | 3 | 3 |
| 261 | P-190-OMR | 28 | M | L1 | 0.101 | 0.099 | 4 | 4 | 4 | 4 |
| 262 | P-190-OMR | 28 | M | L2 | 0.056 | 0.056 | 3 | 3 | 2 | 2 |
| 263 | P-190-OMR | 28 | M | L2 | 0.100 | 0.106 | 4 | 4 | 4 | 4 |
| 264 | P-190-OMR | 28 | M | L3 | 0.056 | 0.063 | 3 | 3 | 2 | 3 |
| 265 | P-190-OMR | 28 | M | L5 | 0.069 | 0.073 | 3 | 3 | 3 | 3 |
| 266 | P-190-OMR | 28 | M | L6 | 0.072 | 0.078 | 3 | 3 | 3 | 3 |
| 267 | P-190-OMR | 28 | M | L9 | 0.096 | 0.099 | 4 | 4 | 3 | 4 |
| 268 | P-191-ALD | 56 | M | H1 | 0.035 | 0.039 | 2 | 2 | 2 | 2 |
| 269 | P-191-ALD | 56 | M | T1 | 0.039 | 0.037 | 2 | 2 | 2 | 2 |
| 270 | P-191-ALD | 56 | M | T2 | 0.031 | 0.033 | 1 | 2 | 1 | 1 |
| 271 | P-191-ALD | 56 | M | T3 | 0.033 | 0.036 | 2 | 2 | 1 | 2 |
| 272 | P-191-ALD | 56 | M | L1 | 0.045 | 0.050 | 2 | 2 | 2 | 2 |
| 273 | P-191-ALD | 56 | M | L4 | 0.055 | 0.058 | 3 | 3 | 2 | 3 |
| 274 | P-192-MTA | 52 | M | H1 | 0.039 | 0.043 | 2 | 2 | 2 | 2 |
| 275 | P-192-MTA | 52 | M | U1 | 0.045 | 0.041 | 2 | 2 | 2 | 2 |


| No | Patient | Age | Gender | Region | $\begin{array}{\|l} \hline S_{a}(\mathrm{~mm}) \\ 1 \text { st Asm } \end{array}$ | $\left\lvert\, \begin{gathered} S_{a}(\mathrm{~mm}) \\ \text { 2nd Asm } \end{gathered}\right.$ | $k$-Means 1st Asm | $k$-Means 2nd Asm | FCM <br> 1st Asm | FCM <br> 2nd Asm |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 276 | P-192-MTA | 52 | M | U3 | 0.050 | 0.044 | 2 | 2 | 2 | 2 |
| 277 | P-192-MTA | 52 | M | T2 | 0.026 | 0.025 | 1 | 1 | 1 | 1 |
| 278 | P-192-MTA | 52 | M | T3 | 0.047 | 0.050 | 2 | 2 | 2 | 2 |
| 279 | P-192-MTA | 52 | M | T4 | 0.044 | 0.042 | 2 | 2 | 2 | 2 |
| 280 | P-192-MTA | 52 | M | T5 | 0.045 | 0.052 | 2 | 3 | 2 | 2 |
| 281 | P-192-MTA | 52 | M | L1 | 0.065 | 0.066 | 3 | 3 | 3 | 3 |
| 282 | P-192-MTA | 52 | M | L2 | 0.028 | 0.029 | 1 | 1 | 1 | 1 |
| 283 | P-192-MTA | 52 | M | L4 | 0.029 | 0.033 | 1 | 2 | 1 | 1 |
| 284 | P-192-MTA | 52 | M | L5 | 0.029 | 0.033 | 1 | 2 | 1 | 1 |
| 285 | P-193-UKY | 50 | F | H1 | 0.028 | 0.025 | 1 | 1 | 1 | 1 |
| 286 | P-193-UKY | 50 | F | U2 | 0.042 | 0.037 | 2 | 2 | 2 | 2 |
| 287 | P-193-UKY | 50 | F | T1 | 0.031 | 0.028 | 1 | 1 | 1 | 1 |
| 288 | P-193-UKY | 50 | F | T2 | 0.042 | 0.037 | 2 | 2 | 2 | 2 |
| 289 | P-193-UKY | 50 | F | L2 | 0.027 | 0.031 | 1 | 1 | 1 | 1 |
| 290 | P-193-UKY | 50 | F | L3 | 0.047 | 0.054 | 2 | 3 | 2 | 2 |
| 291 | P-193-UKY | 50 | F | L6 | 0.029 | 0.024 | 1 | 1 | 1 | 1 |
| 292 | P-194-ZHN | 34 | M | U1 | 0.114 | 0.109 | 4 | 4 | 4 | 4 |
| 293 | P-194-ZHN | 34 | M | U1 | 0.063 | 0.065 | 3 | 3 | 3 | 3 |
| 294 | P-194-ZHN | 34 | M | U3 | 0.086 | 0.083 | 4 | 3 | 3 | 3 |
| 295 | P-194-ZHN | 34 | M | U7 | 0.128 | 0.125 | 4 | 4 | 4 | 4 |
| 296 | P-194-ZHN | 34 | M | U9 | 0.078 | 0.074 | 3 | 3 | 3 | 3 |
| 297 | P-194-ZHN | 34 | M | T1 | 0.064 | 0.066 | 3 | 3 | 3 | 3 |
| 298 | P-194-ZHN | 34 | M | T3 | 0.062 | 0.061 | 3 | 3 | 3 | 3 |
| 299 | P-194-ZHN | 34 | M | T4 | 0.089 | 0.089 | 4 | 4 | 3 | 3 |
| 300 | P-194-ZHN | 34 | M | T5 | 0.068 | 0.073 | 3 | 3 | 3 | 3 |
| 301 | P-194-ZHN | 34 | M | T6 | 0.067 | 0.066 | 3 | 3 | 3 | 3 |
| 302 | P-194-ZHN | 34 | M | T7 | 0.101 | 0.102 | 4 | 4 | 4 | 4 |
| 303 | P-194-ZHN | 34 | M | L1 | 0.052 | 0.055 | 3 | 3 | 2 | 2 |
| 304 | P-194-ZHN | 34 | M | L5 | 0.065 | 0.066 | 3 | 3 | 3 | 3 |
| 305 | P-195-DVM | 42 | F | U1 | 0.044 | 0.042 | 2 | 2 | 2 | 2 |
| 306 | P-195-DVM | 42 | F | U2 | 0.039 | 0.038 | 2 | 2 | 2 | 2 |
| 307 | P-195-DVM | 42 | F | U4 | 0.035 | 0.036 | 2 | 2 | 2 | 2 |
| 308 | P-195-DVM | 42 | F | U5 | 0.043 | 0.039 | 2 | 2 | 2 | 2 |
| 309 | P-195-DVM | 42 | F | U6 | 0.043 | 0.036 | 2 | 2 | 2 | 2 |
| 310 | P-195-DVM | 42 | F | U7 | 0.038 | 0.037 | 2 | 2 | 2 | 2 |
| 311 | P-195-DVM | 42 | F | T1 | 0.067 | 0.065 | 3 | 3 | 3 | 3 |
| 312 | P-195-DVM | 42 | F | T2 | 0.060 | 0.060 | 3 | 3 | 3 | 3 |
| 313 | P-195-DVM | 42 | F | T3 | 0.044 | 0.047 | 2 | 2 | 2 | 2 |
| 314 | P-195-DVM | 42 | F | T5 | 0.032 | 0.033 | 2 | 2 | 1 | 1 |
| 315 | P-195-DVM | 42 | F | T6 | 0.039 | 0.039 | 2 | 2 | 2 | 2 |


| No | Patient | Age | Gender | Region | $S_{a}(\mathrm{~mm})$ <br> 1st Asm | $S_{a}(\mathrm{~mm})$ <br> 2nd Asm | $k$-Means <br> 1st Asm | $k$-Means <br> 2nd Asm | FCM <br> 1st Asm | FCM <br> 2nd Asm |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 316 | P-195-DVM | 42 | F | T9 | 0.029 | 0.030 | 1 | 1 | 1 | 1 |
| 317 | P-195-DVM | 42 | F | L1 | 0.047 | 0.043 | 2 | 2 | 2 | 2 |
| 318 | P-195-DVM | 42 | F | L1 | 0.029 | 0.025 | 1 | 1 | 1 | 1 |
| 319 | P-195-DVM | 42 | F | L1 | 0.053 | 0.056 | 3 | 3 | 2 | 2 |
| 320 | P-195-DVM | 42 | F | L2 | 0.035 | 0.042 | 2 | 2 | 2 | 2 |
| 321 | P-195-DVM | 42 | F | L5 | 0.023 | 0.017 | 1 | 1 | 1 | 1 |
| 322 | P-195-DVM | 42 | F | L6 | 0.046 | 0.049 | 2 | 2 | 2 | 2 |
| 323 | P-195-DVM | 42 | F | L7 | 0.072 | 0.072 | 3 | 3 | 3 | 3 |
| 324 | P-195-DVM | 42 | F | L9 | 0.052 | 0.052 | 3 | 3 | 2 | 2 |

# APPENDIX F: APPROVAL OF OBSERVATIONAL CLINICAL 

STUDY (NMRR-09-1098-4863)

## NATIONAL INSTITUTES OF HEALTH APPROVAL FOR CONDUCTING RESEARCH IN THE MINISTRY OF HEALTH MALAYSIA PENGESAHAN INSTITUSI PENYELIDIKAN NEGARA UNTUK MENJALANKAN PENYELIDIKAN DI KEMENTERIAN KESIHATAN

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| Unique NMRR <br> Registration ID : <br> [Nombor Pendaftaran] | NMRR-09-1098-4863 |
| :--- | :--- |
| Research Title: : <br> [Tajuk] | CBJECTIVE EVALUATION OF PSORIASIS SEVERITY USINGANEW <br> COMPUTERISED PASI SCORING SYSTEM (Alfa-PASI) |
| Protocol Number if <br> available: <br> [Nombor Protokol jika <br> ada] |  |


| $\#$ | Investigator Name <br> [Name Penyelidik] | Institution Name <br> [Nama Institusi] |
| :--- | :--- | :--- |
| 1 | Ahmad Fadzil M Hani | Universit Teknologi PETRONAS (UTP) |
| 2 | Azura Mohd Affandi | Kuala Lumpur Hospital |
| 3 | Chong Yew Thong | Kuala Lumpur Hospital |
| 4 | Suraiya Hani Hussein | Damansara Specialist Hospital |

I have reviewed the above titled research, and approve of its design and conduct.
Saya telah menyemak kajlian yang bertajuk seperti di atas dan meluluskan rekabentuk dan perlaksanaannya.

| Name of Director: <br> (Nama Pengarah] | Dr. Lim Teck Onn |
| :--- | :--- |
| NIH Institute (IMR, CRC, IPH, <br> IHM, IHSR and IHBR) <br> [Nama institus' di bawah NIH] | Clinical Research Centre (CRC) |
| Signature \& Official stamp: <br> [Tandatangan dan Cop <br> Rasmi] | This is computer generated document, <br> therefore no signature is required. |
| Date : <br> [Tarikh] | [T-04-2010 |

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## APPENDIX G: SOURCE CODE OF SURFACE ROUGHNESS DETERMINATION (MATLAB CODE)

```
% =================================================================================
% =========== FILTER AND OPEN A 3D SURFACE FILE
% ==============================================================================
close all; clear all; clc;
folderToBeOpened = 'C:\...\3D Lesion for scaliness\10';
% % read a surface file
cd(folderToBeOpened);
[txtFileNm, folderToBeOpened] = uigetfile('*.txt','Select the M-file');
% read 2D jpg file
N = length(txtFileNm);
imgFileNm = txtFileNm;
imgFileNm(N-2)='j';
imgFileNm(N-1) ='p';
imgFileNm(N)='g';
% ===============================================================================
% ========== End of FILTER AND OPEN A 3D SURFACE FILE
% ===============================================================================
% ==============================================================================
% =========== READ 2D IMAGE AND SEGMENT THE LESION AREA
% ================================================================================
I2d = imread([folderToBeOpened,imgFileNm]);
[m2D,n2D] = size(I2d);
n2D = n2D/3;
% % % % Normalise I 2D
for iii=1:m2D
    for jjj=1:n2D
        id = m2D - iii + 1;
        I2dT(id,jjj,:) = I2d(iii,jjj,:);
    end;
end;
I2d = I2dT;
clear I2dT;
figNameStr = ['File Name ',imgFileNm];
figure('Name',figNameStr);
imshow(I2d);
[BW,x,y] = roipoly;
close(figNameStr);
minX = round(min(min(x)));
maxX = round(max(max(x)));
minY = round(min(min(y)));
maxY = round(max(max(y)));
figure,imshow(I2d(minY:maxY,minX:maxX,:));
% ==========================================================================
% =========== End of READ 2D IMAGE AND SEGMENT THE LESION AREA
% ==========================================================================
```



```
=========== READ TXT FILE AND CONVERT IT INTO A SURFACE MATRIX
```



```
Function hdrload() is created by Jeff Daniels (Dec 20, 2002).
The file is obtained from
%http://www.mathworks.com/matlabcentral/fileexchange/2973-mhdrload-m
[hdr, A] = hdrload(txtFileNm);
initX = A(1,1);
newX = 1e10;
N = 1;
while initX ~= newX
    N = N + 1;
    newX = A(N,1);
end;
N = N - 1;
lengthA = length(A);
M = lengthA/N;
% % % Create X matrix
xArr = A(1:N,1);
for i=1:M
    I.X(i,:) = xArr;
end;
% % % Create Y matrix
for i=1:M
    idx = 1 + (i-1)*N;
    yArr(i) = A(idx,2);
end;
for j=1:N
    I.Y(:,j) = yArr;
end;
% % % Create Z matrix
idxA = 0;
for i=1:M
    for j=1:N
            idxA = idxA + 1;
            I.Z(i,j) = A(idxA,3);
        end;
end
% ==============================================================================
% =========== End Of READ TXT FILE AND CONVERT IT INTO A SURFACE MATRIX ====
% =============================================================================
```

```
% ==============================================================================
```

% ==============================================================================
% ========== SAMPLE THE LESION AREA FROM 3D SURFACE DATA ====================
% ========== SAMPLE THE LESION AREA FROM 3D SURFACE DATA ====================
% =============================================================================
% =============================================================================
k = 1; % for the highest resolution, image size 480 x 640 pixels
k = 1; % for the highest resolution, image size 480 x 640 pixels
minXSc = round(k*minX);
minXSc = round(k*minX);
maxXSc = round(k*maxX);
maxXSc = round(k*maxX);
minYSc = round(k*minY);
minYSc = round(k*minY);
maxYSc = round(k*maxY);
maxYSc = round(k*maxY);
Xcr=I.X(minYSc:maxYSc, minXSc:maxXSc);
Xcr=I.X(minYSc:maxYSc, minXSc:maxXSc);
Ycr=I.Y(minYSc:maxYSc, minXSc:maxXSc);
Ycr=I.Y(minYSc:maxYSc, minXSc:maxXSc);
Zcr=I.Z(minYSc:maxYSc, minXSc:maxXSc);
Zcr=I.Z(minYSc:maxYSc, minXSc:maxXSc);
I2.X = Xcr;
I2.X = Xcr;
I2.Y = Ycr;

```
I2.Y = Ycr;
```

```
I2.Z = Zcr;
figure
subplot(1,2,1), imshow(I2d);
subplot(1,2,2), imshow(I.Z,[]);
figure,
subplot(1,2,1), imshow(I2d(minY:maxY,minX:maxX,:));
subplot(1,2,2), imshow(I2.Z,[]);
% ================================================================================
% ========== End of SAMPLE THE LESION AREA FROM 3D SURFACE DATA ============
% ==============================================================================
% ============================================================================
% =========== DIVIDE THE SAMPLE INTO 4 SUB-DIVIDED SURFACES ==================
```



```
noDiv = 2;
[M,N] = size(I2.Z);
mD = floor(M/noDiv);
nD = floor(N/noDiv);
for i=1:(noDiv)
    cI(i) = 1 + (i-1)*mD;
    CJ(i) = 1 + (i-1)*nD;
end;
idxSd = 0;
for i=1:(noDiv)
    for j=1:(noDiv)
        idxSd = idxSd + 1;
        I2sDiv(idxSd).X = I2.X(cI(i):(cI(i) +mD-1), cJ(j):(cJ(j)+nD-1));
        I2sDiv(idxSd).Y = I2.Y(cI(i):(cI(i)+mD-1), cJ(j):(cJ(j)+nD-1));
        I2sDiv(idxSd).Z = I2.Z(cI(i):(cI(i)+mD-1), cJ(j):(cJ(j)+nD-1));
    end;
end;
figure,
subplot(2,2*noDiv,1), imshow(I2sDiv(1).Z,[]);
subplot(2,2*noDiv,2), imshow(I2sDiv(2).Z,[]);
subplot(2,2*noDiv,5), imshow(I2sDiv(3).Z,[]);
subplot(2,2*noDiv,6), imshow(I2sDiv(4).Z,[]);
subplot(2,2*noDiv,3), surf(I2sDiv(1).X,I2sDiv(1).Y,I2sDiv(1).Z);
idx=1;
miX = min(min(I2sDiv(idx).X));
mxX = max(max(I2sDiv(idx).X));
miY = min(min(I2sDiv(idx).Y));
mxY = max(max(I2sDiv(idx).Y));
miZ = min(min(I2sDiv(idx).Z));
mxZ = max(max(I2sDiv(idx).Z));
axis([miX mxX miY mxY miZ mxZ])
subplot(2,2*noDiv,4), surf(I2sDiv(2).X,I2sDiv(2).Y,I2sDiv(2).Z);
idx=2;
miX = min(min(I2sDiv(idx).X));
mxX = max(max(I2sDiv(idx).X));
miY = min(min(I2sDiv(idx).Y));
mxY = max(max(I2sDiv(idx).Y));
miZ = min(min(I2sDiv(idx).Z));
mxZ = max(max(I2sDiv(idx).Z));
axis([miX mxX miY mxY miZ mxZ])
subplot(2,2*noDiv,7), surf(I2sDiv(3).X,I2sDiv(3).Y,I2sDiv(3).Z);
idx=3;
miX = min(min(I2sDiv(idx).X));
```

```
mxX = max(max(I2sDiv(idx).X));
miY = min(min(I2sDiv(idx).Y));
mxY = max(max(I2sDiv(idx).Y));
miZ = min(min(I2sDiv(idx).Z));
mxZ = max(max(I2sDiv(idx).Z));
axis([miX mxX miY mxY miZ mxZ])
subplot(2,2*noDiv,8), surf(I2sDiv(4).X,I2sDiv(4).Y,I2sDiv(4).Z);
idx=4;
miX = min(min(I2sDiv(idx).X));
mxX = max(max(I2sDiv(idx).X));
miY = min(min(I2sDiv(idx).Y));
mxY = max(max(I2sDiv(idx).Y));
miZ = min(min(I2sDiv(idx).Z));
mxZ = max(max(I2sDiv(idx).Z));
axis([miX mxX miY mxY miZ mxZ])
% ==============================================================================
% =========== End Of DIVIDE THE SAMPLE INTO 4 SUB-DIVIDED SURFACES
% =============================================================================
```

```
% ================================================================================
```

% ================================================================================
% =========== APPLY 2nd ORDER SURFACE FITTING TO EXTRACT ROUGHNESS ===========
% =========== APPLY 2nd ORDER SURFACE FITTING TO EXTRACT ROUGHNESS ===========
=================-====-=-=-============================
=================-====-=-=-============================
noOfAcceptedFitting = 0;
noOfAcceptedFitting = 0;
RaAcc = 0;
RaAcc = 0;
RsqAcc = 0;
RsqAcc = 0;
for iSd=1:(noDiv^2)
for iSd=1:(noDiv^2)
[M,N] = size(I2sDiv(iSd).Z);
[M,N] = size(I2sDiv(iSd).Z);
X = I2sDiv(iSd).X(1:M,1:N);
X = I2sDiv(iSd).X(1:M,1:N);
Y = I2sDiv(iSd).Y(1:M,1:N);
Y = I2sDiv(iSd).Y(1:M,1:N);
Z = I2sDiv(iSd).Z(1:M,1:N);
Z = I2sDiv(iSd).Z(1:M,1:N);
%%%%%%%%%%%%%% Give A Real Surface %%%%%%%%%%%%%%%%%%%
%%%%%%%%%%%%%% Give A Real Surface %%%%%%%%%%%%%%%%%%%
rowIdx = 0;
rowIdx = 0;
for i=1:M %noDivSp
for i=1:M %noDivSp
for j=1:N %noDivSp
for j=1:N %noDivSp
rowIdx = rowIdx + 1;
rowIdx = rowIdx + 1;
Xv = X(i,j);
Xv = X(i,j);
Yv = Y(i,j);
Yv = Y(i,j);
Zv = Z(i,j);
Zv = Z(i,j);
nYv = 2;
nYv = 2;
V(rowIdx,1) = (Xv^2)* (Yv^nYv);
V(rowIdx,1) = (Xv^2)* (Yv^nYv);
V(rowIdx,2) = (Xv^1)* (Yv^nYv);
V(rowIdx,2) = (Xv^1)* (Yv^nYv);
V(rowIdx,3) = (Xv^0)* (Yv^nYv);
V(rowIdx,3) = (Xv^0)* (Yv^nYv);
nYv = 1;
nYv = 1;
V(rowIdx,4) = (Xv^2)* (Yv^nYv);
V(rowIdx,4) = (Xv^2)* (Yv^nYv);
V(rowIdx,5) = (Xv^1)* (Yv^nYv);
V(rowIdx,5) = (Xv^1)* (Yv^nYv);
V(rowIdx,6) = (Xv^0)* (Yv^nYv);
V(rowIdx,6) = (Xv^0)* (Yv^nYv);
nYv = 0;
nYv = 0;
V(rowIdx,7) = (Xv^2)* (Yv^nYv);
V(rowIdx,7) = (Xv^2)* (Yv^nYv);
V(rowIdx,8) = (Xv^1)* (Yv^nYv);
V(rowIdx,8) = (Xv^1)* (Yv^nYv);
V(rowIdx,9) = (Xv^0)* (Yv^nYv);
V(rowIdx,9) = (Xv^0)* (Yv^nYv);
Q(rowIdx,1) = Zv;
Q(rowIdx,1) = Zv;
end;
end;
end;
end;
VtV= V'*V;
VtV= V'*V;
cf = pinv(VtV)*V'*Q;
cf = pinv(VtV)*V'*Q;
% % % % % Reconstruct

```
    % % % % % Reconstruct
```

```
    Zcc = 0;
    nYv = 2;
    Zcc = Zcc + cf(1)*(X.^2).*(Y.^nYv);
    Zcc = Zcc + cf(2)*(X.^1).*(Y.^nYv);
    Zcc = Zcc + cf(3)*(X.^0).*(Y.^nYv);
    nYv = 1;
    Zcc = Zcc + cf(4)*(X.^2).*(Y.^nYv);
    Zcc = Zcc + cf(5)*(X.^1).*(Y.^nYV);
    Zcc = Zcc + cf(6)*(X.^0).*(Y.^nYv);
    nYv = 0;
    Zcc = Zcc + cf(7)*(X.^2).*(Y.^nYV);
    Zcc = Zcc + cf(8)*(X.^1).*(Y.^nYv);
    Zcc = Zcc + cf(9)*(X.^0).*(Y.^nYv);
    Zr = Zcc;
    err = abs(Z-Zr);
    Ra = mean(mean(err));
    meanZ = mean(mean(Z));
    SStot = sum(sum((Z-meanZ).^2));
    SSerr = sum(sum((Z-Zr).^2));
    Rsq = 1 - (SSerr/SStot);
    Ra 2nd(iSd) = Ra;
    Rsq_2nd(iSd) = Rsq;
    if and(Rsq_2nd(iSd)>=0.90,Rsq_2nd(iSd)<=1.0)
        noOfAcceptedFitting = noOfAcceptedFitting + 1;
        RaAcc = RaAcc + Ra 2nd(iSd);
        RsqAcc = RsqAcc + Rsq_2nd(iSd);
    end;
end;
Ra_2nd_Final = NaN;
Rsq_2nd_Final = NaN;
    if noOfAcceptedFitting>0
    Ra_2nd_Final = RaAcc/noOfAcceptedFitting;
    Rsq_2nd_Final = RsqAcc/noOfAcceptedFitting;
end;
ResMat(1,1) = Ra_2nd_Final;
ResMat (1,2) = Rsq_2nd_Final;
% =============================================================================
% =========== End Of APPLY 2nd ORDER SURFACE FITTING TO EXTRACT ROUGHNESS ==
% =============================================================================
% =============================================================================
% =========== APPLY 3rd ORDER SURFACE FITTING TO EXTRACT ROUGHNESS
noOfAcceptedFitting = 0;
RaAcc = 0;
RsqAcc = 0;
for iSd=1:(noDiv^2)
    [M,N] = size(I2sDiv(iSd).Z);
    X = I2sDiv(iSd).X(1:M,1:N);
    Y = I2sDiv(iSd).Y(1:M,1:N);
    Z = I2sDiv(iSd).Z(1:M,1:N);
    %%%%%%%%%%%%% Give A Real Surface %%%%%%%%%%%%%%%%%%%
    rowIdx = 0;
    for i=1:M %noDivSp
```

```
    for j=1:N %noDivSp
        rowIdx = rowIdx + 1;
        Xv = X(i,j);
        Yv = Y(i,j);
        Zv = Z(i,j);
        nYv = 3;
        V(rowIdx,1) = (Xv^3)*(Yv^nYv);
        V(rowIdx,2) = (Xv^2)*(Yv^nYv);
        V(rowIdx,3) = (Xv^1)*(Yv^nYv);
        V(rowIdx,4) = (Xv^0)* (Yv^nYv);
        nYv = 2;
        V(rowIdx,5) = (Xv^3)*(Yv^nYv);
        V(rowIdx,6) = (Xv^2)*(Yv^nYv);
        V(rowIdx,7) = (Xv^1)*(Yv^nYv);
        V(rowIdx,8) = (Xv^0)*(Yv^nYv);
        nYv = 1;
        V(rowIdx,9) = (Xv^3)*(Yv^nYv);
        V(rowIdx,10) = (Xv^2)*(Yv^nYv);
        V(rowIdx,11) = (Xv^1)*(Yv^nYv);
        V(rowIdx,12) = (Xv^0)*(Yv^nYv);
        nYv = 0;
        V(rowIdx,13) = (Xv^3)*(Yv^nYv);
        V(rowIdx,14) = (Xv^2)*(Yv^nYv);
        V(rowIdx,15) = (Xv^1)*(Yv^nYv);
        V(rowIdx,16) = (Xv^0)*(Yv^nYv);
    Q(rowIdx,1) = Zv;
    end;
end;
vtV= v'*V;
cf = pinv(VtV)*V'*Q;
% % % % % % % Reconstruct
Zcc = 0;
nYv = 3;
Zcc = Zcc + cf(1)*(X.^3).*(Y.^nYv);
Zcc = Zcc + cf(2)*(X.^2).*(Y.^nYv);
Zcc = Zcc + cf(3)*(X.^1).*(Y.^nYv);
Zcc = Zcc + cf(4)*(X.^0).*(Y.^nYv);
nYv = 2;
Zcc = Zcc + cf(5)*(X.^3).*(Y.^nYv);
Zcc = Zcc + cf(6)*(X.^2).*(Y.^nYv);
Zcc = Zcc + cf(7)*(X.^1).*(Y.^nYv);
Zcc = Zcc + cf(8)*(X.^0).*(Y.^nYv);
nYv = 1;
Zcc = Zcc + cf(9)*(X.^3).*(Y.^nYv);
Zcc = Zcc + cf(10)*(X.^2).*(Y.^nYv);
Zcc = Zcc + cf(11)*(X.^1).*(Y.^nYv);
Zcc = Zcc + cf(12)*(X.^0).*(Y.^nYv);
nYv = 0;
Zcc = Zcc + cf(13)*(X.^3).*(Y.^nYv);
Zcc = Zcc + cf(14)*(X.^2).*(Y.^nYv);
Zcc = Zcc + cf(15)*(X.^1).*(Y.^nYv);
Zcc = Zcc + cf(16)*(X.^0).*(Y.^nYv);
Zr = Zcc;
err = abs(Z-Zr);
```

```
    Ra = mean(mean(err));
    meanZ = mean(mean(Z));
    SStot = sum(sum((Z-meanZ).^2));
    SSerr = sum(sum((Z-Zr).^2));
    Rsq = 1 - (SSerr/SStot);
    Ra 3rd(iSd) = Ra;
    Rsq_}3rd(iSd) = Rsq
    if and(Rsq_3rd(iSd)>=0.90,Rsq_3rd(iSd)<=1.0)
        noOfAcceptedFitting = noOfAcceptedFitting + 1;
        RaAcc = RaAcc + Ra 3rd(iSd);
        RsqAcc = RsqAcc + Rsq_3rd(iSd);
    end;
end;
Ra_3rd_Final = NaN;
Rsq_3rd_Final = NaN;
if noOfAcceptedFitting>0
    Ra 3rd_Final = RaAcc/noOfAcceptedFitting;
    Rsq_3rd_Final = RsqAcc/noOfAcceptedFitting;
end;
ResMat(2,1) = Ra 3rd Final;
ResMat (2,2) = Rsq__3rd__Final;
% ==============================================================================
% =========== End Of APPLY 3rd ORDER SURFACE FITTING TO EXTRACT ROUGHNESS ==
% ==============================================================================
% =============================================================================
% =========== SELECT THE BEST Ra BASED ON ITS Rsq ============================
% =============================================================================
RsqMax = max(ResMat(:,2));
idxRa = find(ResMat(:,2)==RsqMax);
RaFinal = ResMat(idxRa(1),1);
disp(['Average roughness is ', num2str(RaFinal),' mm']);
```


## APPENDIX H: SOURCE CODE OF CLUSTERING (FCM) IMPLEMENTATION FOR SCORING SCALINESS (MATLAB CODE)

```
close all;
clear all;
clc;
% =============================================================================
% Load training dataset to build clustering system based on FCM algorithm.
% FCM is applied to cluster this combined dataset.
% =================================================================================
load('C:\...\xData.mat');
x=sort(x);
```



```
% ========= Membership degrees initialisation based on training dataset ====
% =================================================================================
L = x;
P = linspace(min(L),max(L),9);
sL = sort(L);
cPMi = P(1);
CPMx = P(9);
for i = 1:length(sL);
    Lv = sL(i);
    % score 1
    cPt = P(2);
    if Lv<cPt
        mD1(i,1) = (1/(cPt-cPMi))*(Lv-cPMi);
    end;
    if Lv==cPt
        mD1(i,1) = 1;
    end;
    if Lv>cPt
        mD1(i,1) = (-1/(cPMx-cPt))*(Lv-cPt) + 1;
    end;
```

    \% score 2
    cPt \(=P(4) ;\)
    if \(L v<c P t\)
        \(m D 2(i, 1)=(1 /(c P t-c P M i)) *(L v-c P M i) ;\)
    end;
    if \(L v==c P t\)
        mD2 (i,1) = 1;
    end;
    if Lv>cPt
        \(\operatorname{mD2}(i, 1)=(-1 /(c P M x-c P t)) *(L v-c P t)+1 ;\)
    end;
    \% score 3
    ```
    cPt = P(6);
    if Lv<cPt
        mD3(i,1) = (1/(cPt-cPMi))*(Lv-cPMi);
    end;
    if Lv==cPt
        mD3(i,1) = 1;
    end;
    if Lv>cPt
        mD3(i,1) = (-1/(cPMx-cPt))*(Lv-cPt) + 1;
    end;
    % score 4
    cPt = P(8);
if Lv<cPt
    mD4(i,1) = (1/(cPt-cPMi))*(Lv-cPMi);
end;
if Lv==cPt
    mD4(i,1) = 1;
end;
if Lv>cPt
    mD4(i,1) = (-1/(cPMx-cPt))*(Lv-cPt) + 1;
end;
end;
figure,
plot(sL,mD1, sL,mD2,sL,mD3, sL,mD4);
u=[mD1 mD2 mD3 mD4];
```



```
% == END of Membership degrees initialisation based on training dataset ====
% ===============================================================================
% ===============================================================================
% ================== SET THE NUMBER OF CLUSTER ==================================
% =============================================================================
k = 4;% number of cluster
N = length(x);
% Fuzziness coefficient is assigned. The best values are between 1.4 and
% 2.8 as mentioned in http://fuzziness.org/fcm
m = 2;
% Initialisation on objective function (OldJm), objective function changes
% during the iteration (deltaJm), and number of interation (noI)
OldJm = 1e9;
deltaJm = 1e9;
noI= 0;
% ==============================================================================
% ============= DETERMINE MEMBERSHIP DEGREES OF CLUSTERED DATASET ============
% ================================================================================
while deltaJm > 1e-5
    noI = noI + 1;
    % Centroids determination
    for j=1:k
        A = 0;
        B = 0;
        for i=1:N
            A = A + (u(i,j)^m*x(i));
            B = B + u(i,j)^m;
```

```
        end;
    C(j) = A/B;
    end;
    % Membership degrees determination
    pw = 2/(m-1);
    for j=1:k
        for i=1:N
            A= 0;
            for l=1:k
                A0 = abs(x(i)-C(j));
                BO = abs(x(i) - C(l));
                A = A + (A0/B0)^pw;
            end;
            u(i,j) = 1/A;
    end;
    end;
    % Objective function determination
    Jm = 0;
    for i=1:N
    for j=1:k
        Jm = Jm + (u(i,j)^m)*(x(i)-C(j))^2;
    end;
    end;
    % update on objective function
    deltaJm = abs(Jm - OldJm);
    OldJm = Jm;
% % showing the outputs in the command window
    JmA(noI) = Jm;
    disp([num2str(noI),' : ',num2str(Jm)])
% % to prepare some coloums containing cluster centroids
    for i=1:k
        c(i).data = ones(N,1)*C(i);
    end;
% finalMat is used to show the dataset, membership degrees, centroids,
% and objective function
    JmFinalMat = ones (N,1)* Jm;
    finalMat =[];
    finalMat = [x u];
    for i=1:k
        finalMat = [finalMat c(i).data];
    end;
    finalMat = [finalMat JmFinalMat];
    finalMatStruct(noI).data = finalMat;
    disp('----------------------------------------------
    disp(noI);
    disp(finalMat);
    disp(' ')
end;
% =================================================================================
% ========== End Of MEMBERSHIP DEGREES DETERMINATION ==========================
% ==============================================================================
============================================================================
%============ CONSTRUCT MEMBERSHIP FUNCTIONS OF CLUSTERED DATASET ==========
% =============================================================================
% arrange the columns of membership degrees according to the centroid
% ascending order
noMaxI = length(finalMatStruct);
```

```
finalMatData = finalMatStruct(noMaxI).data;
finalMatData = sortrows(finalMatData,1);
xData = finalMatData(:,1);
for i=1:k
    idxMat(i,1) = i+1;
    idxMat(i,2) = idxMat(i,1)+k;
    idxMat(i,3) = finalMatData(1, (k+1+i));
end;
idxMat2 = sortrows(idxMat,3);
for i=1:k
    idxMatInt = floor(idxMat2(i));
    mFunc(i).d = finalMatData(:,idxMatInt);
end;
centroids = sort(C);
x = xData;
miX = min(xData);
mxX = max(xData);
% Fit the membership degrees of clustered dataset into a gaussian function
% Matlab provides following Gaussian functions
% --------------------------------------------------------------------
% MODELNAME EQUATION
% gauss1 Y = a1*exp(-((x-b1)/c1)^2)
% gauss2 Y = a1*exp (-((x-b1)/c1)^2) +a2*exp (-((x-b2)/c2)^2)
% gauss3 Y = a1*exp(-((x-b1)/c1)^2)+...+a3*exp(-((x-
b3)/c3)^2)
% ...
% gauss8 Y = al*exp (-((x-b1)/c1)^2)+\ldots...a8*exp (-((x-
b8)/c8)^2)
% -----------------------------------------------------------------------
% Fit the membership degrees of clustered dataset to obtain k (number of
% cluster) gaussian
% functions. The function coefficients and the fitting performance
% parameters are stored
% in cfunCls(i).data and rsquareCls(i).data respectively.
for i=1:k
    y = mFunc(i).d;
    [cfun, rsquare] = fit(x,y,'gauss2');
    cfunCls(i).data = cfun;
    rsquareCls(i).data = rsquare;
end;
% ==============================================================================
% ============== End Of CONSTRUCTION ON MEMBERSHIP FUNCTIONS ===================
% ==============================================================================
% ===============================================================================
% ==== SHOW THE CLUSTERED DATASET AND ITS MEMBERSHIP FUNCTIONS IN THE PLOT =
% ==============================================================================
figure,
subplot(211);
for i=1:k
    plot(xData, mFunc(i).d)
    hold all
    end;
grid on
axis([miX mxX 0 1.2]);
labelFontSize = 12;
xlabel('Scaliness (mm)','FontSize',labelFontSize,'fontweight','b')
ylabel('Membership degree','FontSize',labelFontSize,'fontweight','b')
```

```
title('Membership degree of clustered
dataset','FontSize',labelFontSize,'fontweight','b')
su.bplot(212);
xC = linspace(miX,mxX,100);
for i=1:k
    fitR_sq = rsquareCls(i).data.rsquare;
    if fitR sq >= 0.9
        a1 = cfunCls(i).data.a1;
        b1 = cfunCls(i).data.b1;
        c1 = cfuncls(i).data.c1;
        a2 = cfunCls(i).data.a2;
        b2 = cfunCls(i).data.b2;
        c2 = cfunCls(i).data.c2;
    fittedMemFunc = a1*exp(-((xC-b1)/c1).^2) + a2*exp(-((xC-b2)/c2).^2);
    end;
    plot(xC,fittedMemFunc), hold all
end;
% % preparing title on the plot
plotTitle = '';
for i=1:k
    tVal = ['C(',num2str(i),')= ',num2str(centroids(i)),' '];
    plotTitle = [plotTitle tVal];
end;
title(plotTitle);
xlabel('Scaliness (mm)','FontSize',labelFontSize,'fontweight','b')
ylabel('Membership degree','FontSize',labelFontSize,'fontweight','b')
title('Membership function of clustered
dataset','FontSize',labelFontSize,'fontweight','b')
grid on
axis([miX mxX 0 1.2]);
% ================================================================================
% ==== End of SHOWING THE CLUSTERED DATASET & MEMBERSHIP FUNCTIONS ==========
% ===============================================================================
```

```
% ================================================================================
% ==== CLASSIFY AND INPUT INTO A CLUSTER
% ================================================================================
clc;
xVal = input('Input Roughness value ');
nLn = 5;
xLn = xVal*ones(nLn);
yLn = linspace(0,1,nLn);
line(xLn,yLn,'Color','k','LineWidth',3);
```

for $i=1: k$
fitR_sq = rsquareCls(i).data.rsquare;
if fitR sq >= 0.9
a1 = cfunCls(i).data.a1;
b1 = cfuncls(i).data.b1;
c1 $=$ cfuncls(i).data.c1;
a2 $=$ cfuncls(i).data.a2;
b2 = cfuncls(i).data.b2;
c2 $=$ cfuncls(i).data.c2;
probX(i) $=a 1^{*} \exp \left(-((x V a l-b 1) / c 1) .^{\wedge} 2\right)+a 2 * \exp (-((x V a l-b 2) / c 2) . \wedge 2)$;
end;
end;

\% =============== SUMMARY OF DATA CLASSIFICATION
\% =============================================================================12
clc;
maxProb $=\max ($ probx $)$;
clsOfX = find(probX == maxProb);

']);
disp([' Scaliness | Centroid |Roughness| Memb.Deg | Dec.Skin Tone |
']);
disp(['------------------+----------+---------+---------------------------------1
']);
for $i=1: k$
if $i==1$
sCls $=$ ' Score 1 ';
end;
if $i==2$ sCls = ' Score 2 ';
end;
if i==3 sCls = ' Score 3 '
end;
if $i==4$ sCls $=1$ Score 4 ';
end;
sCentroids = sprintf('\%3.4f',centroids(i));
if centroids(i)<10 sCentroids = [' ',sCentroids];
end;
sXVal $=$ sprintf('\%.2f',xVal);
sProbX = sprintf('\%.4f',probX(i));
if clsOfX == i sClsOfx = ' * ';
end;
if clsOfX ~= i sClsOfX = ' ';
end;
disp([' ',sCls,' | ',sCentroids,' | ',sXVal,' | ',sProbX,' |
',sClsOfx,' |']);
end;
disp (['------------------+-----------+---------+---------------------------------|
']);
disp(['Conclusion: ']);
disp(['xInput = ', sprintf('\%.2f',xVal),' is belong to cluster
',num2str(clsOfX),' with membership degree ',sprintf('\%.4f',maxProb)]);

## APPENDIX I: LIST OF PUBLICATIONS

## Journals

1. Ahmad Fadzil M.H., Esa Prakasa, Vijanth S.A., H. Nugroho, Suraiya H.H., Azura M.A., "3D Surface Roughness Measurement for Scaliness Scoring of Psoriasis Lesions", Computers in Biology and Medicine, (Accepted to be published: 16 August 2013), Elsevier, (Impact Factor in 2012: 1.162).
2. Ahmad Fadzil M.H., Esa Prakasa, H. Fitriyah, H. Nugroho, Suraiya H.H., Azura M.A., "Validation on 3D Surface Roughness Algorithm for Measuring Roughness of Psoriasis Lesion", International Journal of Biological and Life Sciences 8:4 2012, WASET Journal, (Indexed by Scopus).
3. Ahmad Fadzil M.H., Esa Prakasa, Vijanth S.A., H. Nugroho, C.H. Chong, Azura M.A., Suraiya H.H., "Development of Body Surface Area Measurement using Multi-View Imaging for Psoriasis Area Assessment", Australasian Physical and Engineering Sciences in Medicine Journal, Springer Verlag, Volume 34, Number 4, 559-637, 2011 (Impact Factor in 2012: 0.885).
4. Ahmad Fadzil M.H., H. Nugroho, Esa Prakasa, Vijanth S.A, Azura M.A, Suraiya H.H, "Soft Clustering of Lesion Erythema for Psoriasis Assessment", Journal of Investigative Dermatology 2012, Volume 132, Nature Publishing Group, (Impact Factor in 2012: 6.193).

## Book Chapter

1. Ahmad Fadzil M.H., Esa Prakasa, H. Fitriyah, H. Nugroho, Azura M.A., Suraiya H.H., "High Order Polynomial Surface Fitting for Measuring Roughness of Psoriasis Lesion". The paper is published as Book Chapter in Lecture Notes in Computer Science Volume 7066, Springer Verlag.
2. Ahmad Fadzil MH, Esa Prakasa, "Skin Surface Roughness Measurement for Assessing Scaliness of Psoriasis Lesion". The paper is published as Book Chapter in Surface Imaging for Biomedical Applications, Published: April 15, 2014 by CRC Press.
3. Ahmad Fadzil MH, Esa Prakasa, "Determination of Lesion Colour for Clustering Psoriasis Erythema". The paper is published as Book Chapter in Surface Imaging for Biomedical Applications, Published: April 15, 2014 by CRC Press.
4. Ahmad Fadzil MH, Esa Prakasa, "Body Surface Area Measurement for Lesion Area Assessment". The paper is published as Book Chapter in Surface Imaging for Biomedical Applications, Published: April 15, 2014 by CRC Press.

## Conferences

1. Ahmad Fadzil M.H., Esa Prakasa,"Digital Image Analysis-Assessment of Scaliness and Thickness for PASI Scoring", presented at The National Postgraduate Conference on Engineering, Science, and Technology (NPC) 2009, Universiti Teknologi PETRONAS, Seri Iskandar, Malaysia.
2. Ahmad Fadzil M.H., Esa Prakasa, H. Nugroho, H. Fitriyah, Suraiya H.H., Azura M.A., "Parallel Lines Extraction Approach to 3D Image Skin Surface for Roughness Characterization", presented at The WACBE World Congress on Bioengineering 2009, Hong Kong, China.
3. Ahmad Fadzil M.H., Esa Prakasa, H. Fitriyah, H. Nugroho, Suraiya H.H., Azura M.A., "Validation on 3D Surface Roughness Algorithm for Measuring Roughness of Psoriasis Lesion", presented at the International Conference on Medical Informatics and Biomedical Engineering (ICMIBE) 2010, Rio de Janeiro, Brazil.
4. Ahmad Fadzil M.H., Esa Prakasa, Vijanth S.A, H. Nugroho, Chi H.C, Azura M.A, Suraiya H.H, "Development of Body Surface Area Measurement using Multi-View Imaging for Psoriasis Area Assessment", presented at the Engineering and Physical Sciences in Medicine and the Australian Biomedical Engineering Conference (EPSM-ABEC) 2011, Darwin, Australia.
5. Ahmad Fadzil M.H., Esa Prakasa, H. Fitriyah, H. Nugroho, Azura M.A, S.H Hussein, "High Order Polynomial Surface Fitting for Measuring Roughness of Psoriasis Lesion", presented at The 2nd International Visual Informatics Conference (IVIC) 2011, Kuala Lumpur, Malaysia.
6. Ahmad Fadzil M.H., Esa Prakasa, H. Nugroho, Azura M.A, Suraiya H.H, " Sample Area for Surface Roughness Determination of Skin Surfaces", presented at The 4th International Conference on Intelligent and Advanced System (ICIAS) 2012, Kuala Lumpur, Malaysia.
7. Ahmad Fadzil M.H., Esa Prakasa, H. Nugroho, Azura M.A, Suraiya H.H, " Body Surface Area Measurement and Soft Clustering for PASI Area Assessment", presented at the 34th Annual International Conference of the IEEE Engineering in Medicine and Biology Society (EMBC 2012), San Diego, USA.
8. Ahmad Fadzil M.H., H. Nugroho, Esa Prakasa, Vijanth S.A., Azura M.A, Suraiya H.H, "Soft Clustering of Lesion Erythema for Psoriasis Assessment", presented at The 42nd Annual Meeting of the European Society for Dermatological Research (ESDR) 2012, Venice, Italy.
9. Ahmad Fadzil M.H., Esa Prakasa, H. Nugroho, Vijanth S.A., "Implementation of Fuzzy C-Means Clustering for Psoriasis Assessment on Lesion Erythema", presented at the The 2012 IEEE Symposium on Industrial Electronics and Applications (ISIEA 2012), Bandung, Indonesia.
10. Azura M.A, Suraiya H.H, R. Baba, H. Nugroho, Esa Prakasa, Ahmad Fadzil MH, "Objective Evaluation of Psoriasis Severity using a Computerized PASI Scoring System", presented at The 4th Congress of The Psoriasis International Network (PSORIASIS) 2013, Paris, France.

## Conferences, contributing as co-author

1. Ahmad Fadzil M.H., H. Nugroho, Esa Prakasa, Azura M.A., Suraiya H.H., "Skin Roughness Measurement for Lesion Classification", presented at The WACBE World Congress on Bioengineering 2009, Hong Kong, China.
2. Ahmad Fadzil M.H., H. Fitriyah, Esa Prakasa, H. Nugroho, Suraiya H.H., Azura M.A., "Reference Line Construction Approach for Thickness Characterization in 3D Skin Surface Images", presented at The WACBE World Congress on Bioengineering 2009, Hong Kong, China.
3. Ahmad Fadzil M.H., H. Fitriyah, Esa Prakasa, H. Nugroho, Suraiya H.H., Azura M.A., "Thickness Characterization of 3D Skin Surface Images Using Reference Line Construction Approach", presented at The 1st International Visual Informatics Conference (IVIC) 2009, Kuala Lumpur, Malaysia. The paper is published as Book Chapter published in Lecture Notes in Computer Science Volume 5857, Springer Verlag.
4. Ahmad Fadzil M.H., H. Fitriyah, Esa Prakasa, H. Nugroho, Suraiya H.H., Azura M.A., "Objective Assessment of Psoriasis Lesion Thickness for PASI Scoring using 3D Digital Imaging", presented at The International Conference on Medical Informatics and Biomedical Engineering (ICMIBE) 2010, Rio de Janeiro, Brazil.
5. Ahmad Fadzil M.H., H. Nugroho, Esa Prakasa, H. Fitriyah, Azura M.A., Suraiya H.H., "3D-based Skin Roughness Measurement for Lesion Classification", IEEE Symposium on Industrial Electronics \& Applications (ISIEA) 2009, Kuala Lumpur, Malaysia.
6. Ahmad Fadzil M.H., H. Fitriyah, Esa Prakasa, Vijanth S.A., Suraiya H.H., Azura M.A., "In vivo 3D Thickness Measurement of Skin Lesion", IEEE EMBS Conference on Biomedical Engineering and Sciences (IECBES) 2010, Kuala Lumpur, Malaysia.

## Technical Documents

1. Prof. Ir. Dr. Ahmad Fadzil M.H., Dr. Azura M.A., Dr. Chong Yew Thong, Puan Sri Dr. Suraiya H.H., Dr. Roshidah Baba, Esa Prakasa, Hurriyatul Fitriyah, Hermawan Nugroho, "Protocol for the Clinical Study: Objective Evaluation of Psoriasis Severity using A New Computerised Psoriasis Area and Severity Index ( $\alpha$-PASI)". The protocol was approved by Malaysian Research and Ethics Committee (MREC), Ministry of Health on 4 April 2010 (NMRR-09-10984863).
2. Dr. Azura M.A., Dr. Chong Yew Thong, Puan Sri Dr. Suraiya H.H., Prof. Ir. Dr. Ahmad Fadzil M.H., Esa Prakasa, H. Nugroho, H. Fitriyah, "Technical Report: Observational Clinical Study on Computerised System for Monitoring and

Scoring of Psoriasis Area and Severity Index", Centre for Intelligent Signal \& Imaging Research and Department of Dermatology, Hospital Kuala Lumpur, October 2011. The clinical study was conducted from March to October 2010.
3. Prof. Ir. Dr. Ahmad Fadzil M.H., F.A.Sc, FIEM, Esa Prakasa, H. Nugroho, "TechnoFund Completion Report: Computerised System for Monitoring and Scoring of Psoriasis Area and Severity Index (TF 0308 C041)", 26 October 2011.

## APPENDIX J: PATENT DOCUMENT OF SCALINESS ALGORITHM




APPENDIX K: CERTIFICATE OF AWARD (ITEX 2009)


APPENDIX L: CERTIFICATE OF AWARD (INNOVA 2010)


## APPENDIX M: CERTIFICATE OF AWARD (MTE 2012)



