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DOCTOR OF PHILOSOPHY

Developing nomograms and predictive models based on Multiparametric MRI characterised localised prostate cancer

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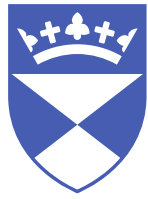
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**University
of Dundee**

**Developing nomograms and predictive
models based on Multiparametric MRI
characterised localised prostate cancer.**

by

Saeed Alqahtani

A Thesis Submitted in Fulfilment of the Requirements of the University of Dundee
for the Degree of Doctor of Philosophy (Ph.D.)

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Declarations:

I, Saeed Alqahtani, hereby certify that this thesis is written by me, is my own work and has not been submitted in any other degree qualification except as specified.

Signature of candidate

I confirm that Saeed Alqahtani has completed the minimum duration period of registration of full-time study at the University of Dundee and has fulfilled the conditions of the University of Dundee, thereby qualifying him to submit this thesis in application for the degree of Doctor of Philosophy.

Prof. Zhihong Huang

Prof. Ghulam Nabi

Contents:

Chapter 1 Introduction	3
1.1 Motivations and research objectives	3
1.2 Content of the thesis	7
1.3 Publication from the thesis	8
1.4 Publication related to the thesis	8
Chapter 2 Literature review	9
2.1 Anatomy and histology of the prostate gland.....	9
2.2 Prostate cancer (PCa)	11
2.2.1 Diagnosis	11
2.2.2 Grading and staging	12
2.2.3 Treatment	14
2.3 Multiparametric MRI of the prostate gland.....	16
2.3.1 Anatomical MR imaging for prostate gland	17
2.3.2 Functional MR imaging for prostate gland.....	24
2.4 Prostate Imaging-Reporting and Data system (PIRADS)	31
2.5 Histological grades between biopsy and radical prostatectomy.....	35
2.6 MRI-targeted biopsy.....	37
2.7 Nomogram construction.....	39
2.8 Decision-analysis curve	43
2.9 Summary	45
Chapter 3 Prediction of prostate cancer Gleason score upgrading	46
3.1 Materials and methods.....	46
3.1.1 Study population.....	46
3.1.2 MRI protocol and PIRADS score	50
3.1.3 Histopathology data and analysis	54
3.1.4 Mould design: a better standard reference	54
3.1.5 Statistical analysis.....	60
3.2 Result.....	63
3.2.1 Upgrading cohort characteristics	63
3.2.2 Predicting of GS upgrading	66
3.2.3 Impact of PIRADS score on prediction of GS upgrading in relation to other factors.....	69
3.2.4 Decision curve analysis	72
3.3 Discussion	74

Chapter 4 Predicting the performance of concurrent SB during TB sampling of MP-MRI	80
4.1 Materials and methods.....	80
4.1.1 Study population and power calculation.....	80
4.1.2 MRI techniques.....	81
4.1.3 Biopsy procedures and histopathological analysis	82
4.1.4 Outcomes	87
4.1.5 Statistical analysis.....	88
4.2 Result.....	90
4.2.1 Patient's characteristics.....	90
4.2.2 Comparison of the detection rate of significant prostate cancer between SB, TB and combined approaches.....	92
4.2.3 Univariate and multivariate logistic regression analysis and developed nomogram	97
4.2.4 Decision curve analysis	102
4.3 Discussion	104
Chapter 5 Conclusion and future work	109
References	111
Appendix	133
Programming for nomogram, calibration and decision curve analysis using RStudio ..	133
Raw data 1	137
Raw data 2	143

List of Figures:

Figure 2-1 The anatomical features of the prostate gland with other surrounding pelvic structures (24).....	9
Figure 2-2 Zonal anatomy of the prostate and their relation to each other (26).	10
Figure 2-3 Axial T1 showing diffuse hyperintense signal in the right PZ (arrowheads) due to post-biopsy haemorrhage with area free of T1 hyperintense blood (asterisks) matched the PCa lesions (T1 haemorrhage exclusion sign).	18
Figure 2-4 Normal prostate anatomy. (A) T2W images show the PZ and TZ in the axial plane. (B) Axial T2W image at the prostatic base shows the CZ (white arrow) as a hypointense area surrounding the ejaculatory ducts (55).	20
Figure 2-5 On (A) axial, (B) sagittal, and (C) coronal views, T2W imaging of a PCa lesion in the left PZ (white arrow) (55).	22
Figure 2-6 Axial T2W image of a PCa lesion in the prostate's right anterior part (white arrow) (55).	23
Figure 2-7 (A) DWI of PCa in the left PZ (white arrow) b1400 (B) ADCmap (55).	26
Figure 2-8 (A) DCE-MRI colour map shows early focal contrast enhancement in the TZ tumour (Red dotted circle). (B) DCE-curve which is a typical malignant curve with a high peak, rapid early enhancement (high wash-in rate) and early wash-out.	28
Figure 2-9 (A) The MRSI was acquired from the whole prostate showing a large low signal intensity lesion (arrows) in the left PZ. (B) The right voxel from the PZ (blue circle) shows the high signal intensity with the normal metabolic profile of the prostatic gland, high level of citrate and lower level of choline. (C) The corresponding spectrum from the left voxel shows increased choline and reduced citrate, indicative for a metabolic profile of prostatic cancer (orange circle) combined with the low signal intensity on MRI (87).	30
Figure 2-10 The nomogram depicts the impact of various predictive variables on various horizontal lines. The lengths of the different lines vary depending on the influence of each predictor. The greater the influence of a horizontal line, the longer it is. A number of points on the respective horizontal line represent the influence of each predictor. The anticipated magnitude of response can be read on the response horizontal line at the bottom of the nomogram by adding the points associated with each predictor (112).	41

Figure 2-11 At a threshold probability of < 5%, > 50%, the nomogram is irrelevant (115).	44
Figure 3-1 Flowchart of the study.....	47
Figure 3-2 lesions localised and graded in MRI	51
Figure 3-3 a: A 73-year-old man with GS 6 disease on prostate cancer on TRUS-Guided biopsies. b: The grade was upgrading to GS 7 on whole mount RP specimen c: Axial T2- weighted image shows ill-defined homogeneous low-signal-intensity on the CZ (d), ADC shows restricted diffusion in low-signal mass, and (e) DCE shows fast and strong enhancement and early contrast agent washout (type 3 curve) (e,f). The lesion was scored as PIRADS v2.0 5 (>1.5cm) and based on the parameters described here.	53
Figure 3-4 Steps of Method: (1) segmentation of MRI data in biomedical software MIMICS, (2) mould making in CAD software SolidWorks, (3) 3D printout from rapid prototyping machine Uprint, (4) post-radical prostatectomy specimen before dyeing and mould placement, (5) slicing of the prostate specimen with a single-blade, (6) final slices shown in the mould, and (7) the tissue slices are arranged from the apex to the base (120).	55
Figure 3-5 MRI's slices for a specific patient	57
Figure 3-6 Segmentation of the prostate (green area) of MR imaging	57
Figure 3-7 a view of all printed moulds	59
Figure 3-8 Cutting procedures and specimens after cutting.....	60
Figure 3-9 RP GS stratified according to PIRADS score.	65
Figure 3-10 ROC curve of the clinical variables with and without PIRADS score.....	68
Figure 3-11 The nomograms of GS upgrading prediction with (a1) and without PIRADS score (a2). Calibration plots of observed and predicted probability of GS upgrading with (b1) and without PIRADS score (b2).....	71
Figure 3-12 Decision analysis demonstrated a high net benefit of PIRADS score model across a wide range of threshold probabilities. Prediction model without PIRADS score (red line); prediction model with PIRADS score (blue line).	73
Figure 4-1 An MRI of the prostate taken before the biopsy is used as a map.	82

Figure 4-2 EMS with a trans-rectal ultrasound transducer	83
Figure 4-3 (A) The left window displays an MRI of the prostate, and (B) the right side window (US window) and the green circled region shows the index lesion where the biopsy can be taken from.	84
Figure 4-4 TRUS guided prostate biopsy.....	85
Figure 4-5 A: A 76 years old patient with a PIRADS 5 lesion detected from 3T MRI in anterior zone with a high PSA and abnormal DRE. B: Patient-specific 3D Mould based grossing of radical prostatectomy slice shows a 3+4 GS cancer located in the anterior zone	87
Figure 4-6 The detection rate of significant prostate cancer between SB, TB based on patients' level.	92
Figure 4-7 The detection rate of significant PCa via SB, TB and combined SB+TB on RP lesions.....	95
Figure 4-8 Nomogram with significant clinical variables to predict patients who will benefit from performing SB in addition to fusion TB.....	99
Figure 4-9 ROC curve and AUC for model discriminative ability.....	100
Figure 4-10 Model calibration plot for observed and predicted probability.....	101
Figure 4-11 Decision analysis demonstrated a high net benefit of the model across a wide range of threshold probabilities.....	103

List of Tables:

Table 2.1 T2W, DWI, and DCE imaging are classified using the PIRADS system. On T2W imaging, the PZ and TZ have significantly different anatomical appearances, so different PIRADS criteria are used for the two zones (43,93).....	32
Table 2.2 EPE risk scoring of extra prostatic extension (43).....	33
Table 2.3 Important Steps in constructing a Nomogram.	39
Table 3.1 Patient characteristics.....	49
Table 3.2 MRI acquisition parameters.	50
Table 3.3 Comparison between biopsy and RP GS sum.....	64
Table 3.4 Univariate and multivariate logistic regression analysis.....	67
Table 3.5 Comparison between previous and current studies.....	76
Table 4.1 SB report for all the 12 cores	86
Table 4.2 Descriptive characteristics of 198 patients who underwent MP-MRI of the prostate and subsequent systematic random and TB at our centre.....	91
Table 4.3 Biopsy findings per patient of SB compared with TB for the total cohort of 198 men. Orange shading indicates men in whom SB upgraded the prostate cancer risk category in relation to TB. Blue shading indicates men in whom TB upgraded prostate cancer risk.93	
Table 4.4 Upgrading lesions between SB and TB for the targeted lesions (112)	96
Table 4.5 Upgrading lesions between combined biopsy and TB for the targeted lesions (112)	96
Table 4.6 Upgrading lesions between combined biopsy and SB for the targeted lesions (112)	97
Table 4.7 Univariate and multivariate logistic regression analysis.....	98

Abbreviations

AS	Active surveillance
ADC	Apparent diffusion coefficient
AUC	Area under the curve
BIRADS	Breast imaging reporting and data system
BPH	Benign prostate hypertrophy
c-statistic	Concordance statistic
csPCa	Clinically significant cancer
cT	Clinical tumour stage
CZ	Central zone
DCE	Dynamic contrast enhanced
DRE	Digital rectal examination
DWI	Diffusion-weighted imaging
ECE	Extra capsular extension
EPE	Extra prostatic tumour extension
ERC	Endo-rectal coil
ESUR	European Society of Urogenital Radiology
GS	Gleason score
AJCC	American Joint Committee on Cancer
UICC	International Union Against Cancer
HIFU	High-intensity focused ultrasound
IQR	Interquartile range
ISUP	International Society of Urological Pathology
K_{ep}	Reverse reflux rate constant
K_{trans}	Transfer constant
m	Mean
M	Median
MP-MRI	Multiparametric MRI
MRI	Magnetic resonance imaging
MRSI	Magnetic resonance spectroscopic imaging
NVB	Neurovascular bundle
OR	Odds ratio

PIRADS	Prostate imaging reporting and data system
PCa	Prostate cancer
PPA-coil	Pelvic-phased-array coil
PSA	Prostate specific antigen
PSAD	Prostatic specific antigen density
PZ	Peripheral zone
RP	Radical prostatectomy
SB	Systematic random biopsy
SD	Standard deviation
SVI	Seminal vesicle invasion
T1W	T1-weighted
T2W	T2-weighted
TB	MR/US fusion targeted biopsy
TRUS	Transrectal ultrasound
TZ	Transitional zone
US	Ultrasound
TR	Transrectal
TP	Transperineal

Abstract

Multiparametric magnetic resonance imaging (MP-MRI) and MRI targeted biopsies (TB) are a new standard in prostate cancer (PCa) screening and diagnosis. Guidelines already include this approach for patients at risk. First, this thesis aimed to assess whether pre-biopsy MRI can narrow the discrepancy of histopathological grades between biopsy and radical prostatectomy (RP) using the Prostate Imaging Reporting and Data System (PIRADS). Second, this thesis aimed to develop a prediction model to identify patients who will benefit from performing systematic random biopsy (SB) at the time of TB.

330 men treated consecutively by RP with localised PCa were included in this study. The MRI and histopathology of the biopsies and RP specimens were assessed respectively. A multivariate model was constructed using logistic regression analysis to assess the ability of MRI to predict upgrading in biopsy Gleason score (GS) in a nomogram. A decision-analysis curve was constructed assessing the impact of the nomogram using different thresholds for probabilities of upgrading. In the SB and TB study, 198 patients with positive MRI findings who underwent both TB and SB were prospectively recruited in this study. The first outcome was to compare the detection rate of clinically significant prostate cancer (csPCa) in SB and TB. For the second outcome, a multivariate analysis using a logistic regression model and nomogram construction were used to identify patients who will benefit from SB in addition to TB. A decision-analysis curve was constructed assessing the impact of the nomogram using different thresholds for probabilities of our model. Statistical analyses were performed using IBM SPSS (version 23.0) and RStudio (version 4.0.3)

Using multivariate analysis, the PIRADS v2.0 score significantly improved the predictive ability of MRI scans for upgrading of biopsy GS ($p=0.001$, 95% CI [0.06-

0.034]), which improved the C-index of predictive nomogram significantly (0.90 vs. 0.64, $p < 0.05$). Moreover, the detection rate of csPCa using SB and TB was 51.0% (101/198) and 56.1% (111/198), respectively, adopting a patients-based biopsy approach. The detection rate of csPCa was higher using a combined biopsy (64.6%; 128/198) compared to a TB (56.1%; 111/198) alone. This was statistically significant ($\chi^2 = 15.06$, df (degree of freedom) = 1, $P < 0.001$). In the multivariate analysis, age, prostate-specific antigen density (PSAD) and PIRADS score were found to predict the detection of significant PCa by SB in addition to TB. A nomogram based on the model showed good discriminative ability (C-index; 78%).

In conclusion, MP-MRI using PIRADS score was shown to be an independent predictor of postoperative GS upgrading, and that this should be taken into consideration while offering treatment options to men with localised PCa. There was a significant difference in the detection of csPCa using a combined biopsy approach. The developed nomogram could help identify those patients at risk of having PCa who will benefit from adding SB biopsy in addition to TB.

Chapter 1 Introduction

1.1 Motivations and research objectives

Histology from biopsies categorised into GS is the only confirmatory test for cancer diagnosis and is most commonly used for risk stratification of men with a recent diagnosis of PCa. Based on the above, men are counselled for various treatment options. With increasing therapeutic options available to men with a diagnosis of PCa, scrutiny of information from biopsy grade becomes increasingly important. There is around 35.5% (range: 14–51%) upgrading of biopsy GS on LRP (1). Many factors contribute to the discrepancy between needle biopsy and corresponding radical surgery GS. Under calling of Gleason cribriform Gleason pattern 4 as pattern 3 or the presence of borderline grades due to barely appreciable glandular differentiation under microscope and lack of sampling of tertiary grade disease on biopsies are known contributors. Factors such as age, size of prostate, extent of cancer on biopsy needle and number of biopsy samples (extended/ or mapping) are also known to impact on the incidence of upgrading (2).

In light of a number of studies reporting upgrading or undergrading of PCa on needle biopsies, there is the potential for under treatment or overtreatment (i.e. radiotherapy and hormone duration). Several publications (3,4) and consensus updates on the Gleason grading system have partially addressed this issue including recommending deriving GS by adding the most common and highest Gleason pattern on the biopsy rather than original method of adding the primary and second most common patterns (5). Moreover, upgrading, if suspected, has long-term outcome implications. Corcoran et al (6) showed that even after adjusting for known preoperative variables (including clinical stage, PSA, number of positive cores, and percentage of positive cores) upgrade to a higher GS remained a strong and

independent predictor of biochemical recurrence after attempted local curative therapy, this underscores the importance of gaining more information to predict upgrading of biopsy GS in men diagnosed with PCa as this may serve as a marker of biologically aggressive disease.

Pre-biopsy MRI has recently been shown to hold great promise in the detection and characterisation of PCa (7). A negative scan (no lesion seen on the MRI scan) showed a high negative predictive value for the presence of significant PCa (8). Song et al (9) reported a high predictive value of PIRADS v2 in predicting upgrading of GS from biopsy, however this study was retrospective and MRI was obtained at least 3 weeks following biopsies, an approach known to impact the interpretation of images. Post-biopsy haemorrhage is the most common false-positive finding for PCa (10). In this study, there was no attempt to align histopathological sectioning to MRI using recently reported 3D-mould technology. Therefore, the objective of the first part of the thesis was:

1. To evaluate whether pre-biopsy MRI had the potential to narrow the discrepancy of histopathological grades between transrectal ultrasound (TRUS) biopsy and radical prostatectomy (RP) using PIRADS v2.0

Recent trends and evidence support pre-biopsy MRI with selective targeting of suspected malignant lesions using MRI/ultrasound (US) and TB methods (11). The burgeoning interest in fusion imaging has arisen to address the main limitations of SB: over detection of clinically insignificant cancers and possibly under detection of csPCa. A number of recent reports support the utility of pre-biopsy multi-parametric magnetic resonance imaging (mpMRI) to address the limitations of SB, and the advantage of increased csPCa detection (12–14). Pre-biopsy MRI in MRI directed pathways have been reported to detect a higher number of csPCa. However,

benefits of image-guided targeting performed in combination with systematic sampling or alone remains poorly defined. Drost et al. in a recent systematic review, used a mixed population (with or without image fusion targeting of suspicious areas) to answer this question, however image fusion methods were not used in all the cases of included studies and hence the benefits of targeting suspicious areas with the image fusion approach, with or without addition of systematic biopsy sampling, remains unclear (15) .

Image fusion approach makes use of information from MRI to direct biopsy needles under real-time US guidance (12,16,17) . Studies have shown that mpMRI combined with TB technology is a promising tool in the diagnosis of PCa (12,18,19) . In light of a number of previous trials showing the significant benefits of image TB, research is now focused on whether random biopsies are required at all in the detection of prostate cancer (20–22) . This question is pertinent to settle an ongoing debate as studies have also highlighted that TB with the addition of systematic random biopsy is superior to systematic random biopsy alone either in terms of capturing csPCa or even in terms of post-procedural morbidity (11) . In a large retrospective study, from centres in Europe and the USA, Dell'Oglio et al. aimed at findings a group of men where systematic biopsies could be avoided all together in men with MRI-facilitated targeted biopsy approach. The authors failed to achieve their objectives and concluded that systematic sampling should be combined with the image guided fusion biopsies (23) . In a large multicentre prospective study, researchers concluded a higher detection rate for clinically significant prostate cancer for a combined approach (TB and SB) biopsy method, however different image fusion systems including cognitive guidance were used in targeted biopsy approach (21). Furthermore, the study did not use PIRADS v2 categorisation and no standardised

protocol was used for MRI imaging. This allowed biases and heterogeneity into the reported data. Our study is a protocol-driven prospective investigation with standardised US/MRI fusion protocol. We assessed clinical variables that could help in identifying patients who may benefit from systematic random biopsies in addition to fusion targeted approach. The comprehensive analysis and outcomes using methodology of this study has not been reported in the literature (24,25), in particularly net clinical benefit of the approach.

The aim of the second part of the thesis was to:

1. compare the diagnostic accuracy of MRI/US fusion targeted biopsies, systematic biopsies and combined approaches in the detection of csPCa and define predictive factors where a combined approach could be used.
2. Quantify additional benefits of adding systematic biopsies to the targeted biopsies approach by constructing a nomogram and assessing its net-clinical benefits.

1.2 Content of the thesis

This thesis includes five chapters. Chapter I introduction: the purpose and background of this project.

Chapter II reviews the literature and summarises the knowledge related to my project, including prostate anatomy, diagnosis and treatment of prostate cancer, the principles and applications of MP-MRI are revealed. Finally, how to construct a nomogram and decision curve analysis.

Chapter III (Prediction of prostate cancer Gleason score upgrading): contains methods and materials of the first study, analysis, results and discussion. (First paper)

Chapter IV (Predicting the performance of concurrent SB during TB sampling of MP-MRI) contains methods and materials of the second study, analysis, results and discussion. (Second paper)

Chapter V Conclusion and future work: Pre-biopsy MRI and PIRADS score significantly and independently predict GS upgrading, a nomogram using clinical variables can assist decision-making during the counselling of patients to have systematic sampling of the prostate in addition to an image fusion biopsy approach. The further work also is mentioned in this chapter.

1.3 Publication from the thesis

1/ **Alqahtani S**, Wei C, Zhang Y, Szewczyk-Bieda M, Wilson J, Huang Z, Nabi G. Prediction of prostate cancer Gleason score upgrading from biopsy to radical prostatectomy using pre-biopsy multiparametric MRI PIRADS scoring system. Scientific reports. 2020 May 7;10(1):1-9.

(Chapter 3)

2/ **Alqahtani, S.**; Zhang, X.; Wei, C.; Zhang, Y.; Szewczyk-Bieda, M.; Wilson, J.; Huang, Z.; Nabi, G. Predicting the Performance of Concurrent Systematic Random Biopsies during Image Fusion Targeted Sampling of MultiParametric MRI Detected Prostate Cancer. A Prospective Study (PRESET Study). Cancers 2022, 14, 1

(Chapter 4)

1.4 Publication related to the thesis

1/Wei C, Zhang Y, Malik H, Zhang X, **Alqahtani S**, Upreti D, Szewczyk-Bieda M, Lang S, Nabi G. Prediction of postprostatectomy biochemical recurrence using quantitative ultrasound Shear wave elastography imaging. Frontiers in oncology. 2019 Jul 9; 9:572.

Chapter 2 Literature review

2.1 Anatomy and histology of the prostate gland

The prostate is a male organ surrounding the urethra, lying between the urinary bladder superiorly and the urogenital diaphragm inferiorly. It is a fibro-muscular glandular structure measuring approximately 3 cm in length.

The prostate is conical in shape and has a base superiorly against the bladder neck, and an apex lies on the superior surface of the urogenital diaphragm (Figure 2-1).

The mid-prostate locates between the base superiorly and the apex inferiorly (26).

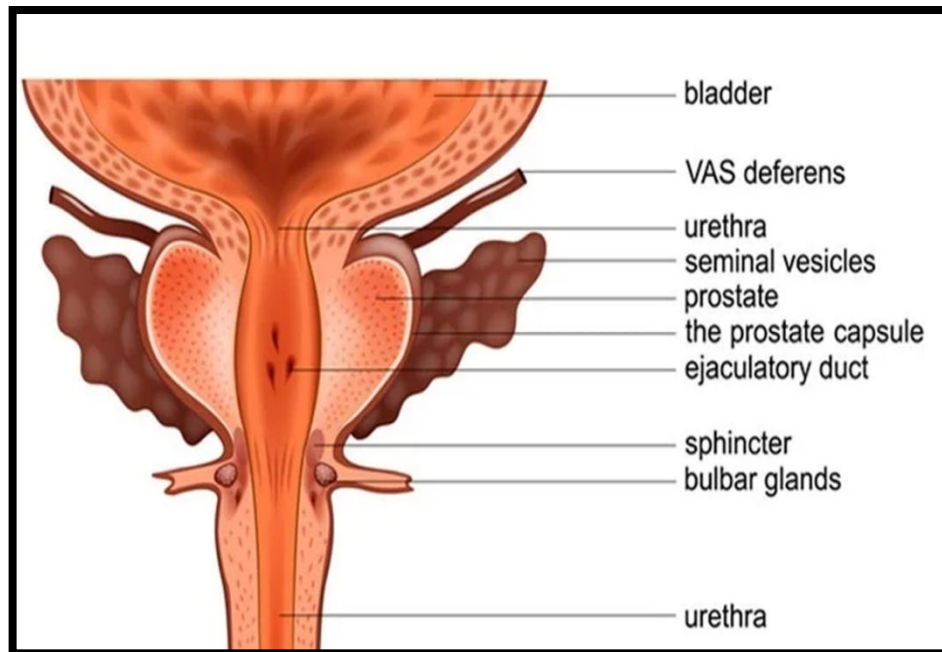


Figure 2-1 The anatomical features of the prostate gland with other surrounding pelvic structures (27).

McNeal (28) defined three separate zones of the prostate: the central zone (CZ); the peripheral zone (PZ); and the transition zone. All differed histologically and biologically. The CZ is located posterosuperiorly to the transitional zone (TZ), which is located centrally and surrounds the proximal part of the urethra. The PZ makes up the main body of the gland (approximately 65%) and is located in the posterior and

inferior parts of the prostate; 75% of malignancy is found in this zone (28) (Figure 2-2).

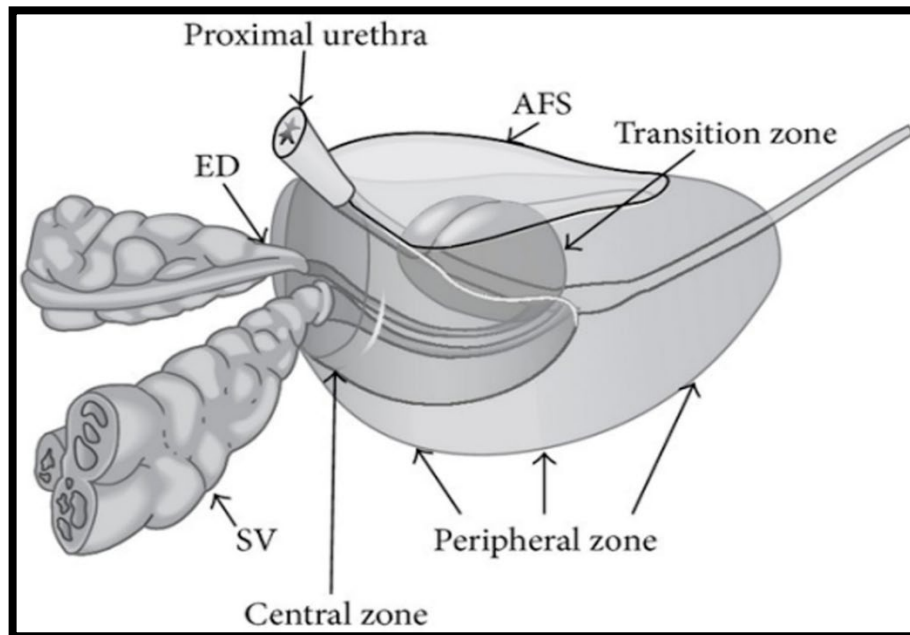


Figure 2-2 Zonal anatomy of the prostate and their relation to each other (29).

Histologically, the prostate gland is made up of epithelial and stromal components. The epithelial part is at most represented by the acinar glands that are lined with secretory stratified columnar epithelial cells. These glands are separated by the connective tissue stroma which consists of fibroblast, nerves, smooth muscle fibres, and rarely, adipocytes (30).

The function of the prostate gland is to produce a fluid that, together with sperm cells from the testicles and fluids from other glands, makes up semen. The muscles of the prostate also ensure that the seminal fluid is pressed into the urethra and then expelled outwards during ejaculation (28).

2.2 Prostate cancer (PCa)

The second most frequently diagnosed cancer in males worldwide, PCa continues to be a major health issue (31). Approximately 1.11 million men are diagnosed with PCa and 307,000 deaths by PCa occur per year (32). Higher than 95% of PCa are referred to as acinar or conventional types (30).

According to studies, the incidence of PCa relates to increases in age (33,34). From the results shown, almost 37% of PCa tumours were seen in males under the age of 65. Patients aged 65 to 74 years old with PCa were the largest group, accounting for 36% all cases. 27% of PCa was detected in males 75 years of age and older. Therefore, patients over the age of 65 accounted for 63% of all PCa.

GS is thought to be the gold standard for PCa aggressiveness and determination. It is based on the observation of acini under the microscope by a pathologist. Cancer epithelial cells have larger nuclei and more prominent nucleoli than normal epithelial cells, and there are no basal cells (35).

2.2.1 Diagnosis

High levels of PSA and an irregular digital rectal exam (DRE) are indicators of high PCa risk. A biopsy can be used to confirm a diagnosis of PCa. Low-grade indolent PCa may not progress to a clinically significant tumour. High-grade PCa can be aggressive, easily spread, and result in death. However, PSA detection, biopsies, and DRE have limited sensitivity and accuracy. This is because 70-80% of cases with high PSA have benign prostatic hyperplasia (BPH) or prostatitis, so the specificity of PSA to PCa is low. DRE can only be used to assess tumours from the posterior of the prostate. Moreover, it cannot detect the number of lesions, as tumour cells may be mixed with normal prostate tissue. A biopsy taken from a multifocal

PCa lesion can show Gleason underscoring (36). In addition, patients with low-grade PCa identified by biopsies may have aggressive lesions outside of the biopsied region, especially in bigger prostate. This might result in Gleason underscoring. Adding pre-biopsy MRI in the diagnosis and characterisation of PCa has recently been shown to be very promising (7).

A number of recent reports have supported the use of pre-biopsy MRI to address many of the limitations of SB, with the prominent advantage being the increased detection of csPCa (7,37–39).

2.2.2 Grading and staging

Used to grade PCa (40), the GS system was named after Gleason, a pathologist, who defined five degrees of increasing aggressiveness (grading) based on the extent to which tumour cells are structured into identifiably glandular structures (glandular differentiation) at low magnification:

Grade 1 – Small uniform glands with minimal nuclear changes.

Grade 2 – Medium sized acini still separated by stromal tissue but more closely arranged.

Grade 3 – These tumours have marked variation of glandular size and organization, as well as stromal tissue infiltration.

Grade 4 – Marked cytological atypia with extensive infiltration.

Grade 5 – Sheets of undifferentiated cancer cells.

Since PCa has more than two grades in heterogeneity, the GS system combines a primary score, which is the most prevalent, with a secondary grade, which is the second most prevalent, to assess PCa. GS is obtained by combining the primary and secondary grades. Consequently, the GS ranges from 2 (1+1) to 10 (5+5) (41).

The TNM classification, developed and maintained by the American Joint Committee on Cancer (AJCC) and the International Union Against Cancer (UICC) for the 7th edition in 2010 (42), is used to stage PCa.

Primary tumour assessment (T):

T1 - Clinically inapparent tumour neither palpable nor visible by imaging.

T1c - Tumour identified by needle biopsy (for example, because of elevated PSA).

T2 - Tumour confined within the prostate.

T3 - Tumour extends through the prostate capsule.

T3a - Extracapsular extension **T3b** = Tumour invades seminal vesicles

T4 - Tumour is fixed or invades adjacent structures other than seminal vesicles, such as external sphincter, rectum, bladder, levator muscles and/or pelvic wall.

Regional lymph node involvement (N):

NX - Regional lymph nodes were not assessed.

N0 - No regional lymph node metastasis.

N1 - Metastasis in regional.

Distant Metastasis (M):

M1 - Distant metastasis.

M1a - Non-regional lymph node(s).

M1b - Bone(s).

M1c - Other sites.

2.2.3 Treatment

PCa treatment requires the collaboration of a multidisciplinary team including urologists, histopathologists, radiologists, and radiation oncologists.

Age, original PSA, clinical staging (TNM), GS, urinary function, and comorbidities all play a role in treatment planning (43). A patient's life expectancy and the disease's biological characteristics, as well as the disease's predicted aggressiveness, are important factors to consider before starting any PCa treatment plan. A patient's preferences for various treatment choices should also be taken into account, by considering treatment risks before taking any decision.

Clinically localised PCa is defined by a tumour confined to the prostate gland or a Clinical stage of T2 or less (44). It is widely treated with the following:

1- Active surveillance (AS)

- Regularly monitoring of the disease is usually preferred for those with very-low-risk and low-risk PCa.

2. Radical prostatectomy (RP)

- Removing the whole prostate gland. Seminal vesicles and certain nearby lymph nodes may be removed based on tumour staging (44).

3. Radiotherapy

- Radiotherapy aims to destroy PCa cells without causing too much damage to healthy cells. External radiotherapy uses high-energy X-ray beams targeted at the prostate from outside the body. Brachytherapy is another type of radiotherapy where tiny radioactive seeds are internally put into the prostate.

4. Hormonal therapy

- Androgen suppression treatment is another name for hormone therapy. The goal is to lower or avoid male hormones, known as androgens, from fuelling PCa cells in the body. Hormonal therapy may also be used in conjunction with other treatments, such as radiotherapy, to improve the efficacy of the treatment (45).

5. Focal therapy

- High-intensity focused ultrasound (HIFU) uses high-frequency ultrasound energy to heat and destroy cancer cells in the prostate. Recently, patients with clinically significant nonmetastatic PCa can be given focal treatment using HIFU since it is effective in the medium term and has a low-risk of urinary and rectal side effects (46).

2.3 Multiparametric MRI of the prostate gland

MRI of the prostate is performed using a 1.5 or 3 Tesla MRI scanner and a pelvic phased-array coil (PPA-coil) placed over the pelvis, with or without an endorectal coil (ERC) depending on the clinical condition. The use of an ERC can increase image quality, as it is located in the rectum just posterior to the prostate gland as well as to fixate the prostate during the scan, potentially reducing motion artefacts. However, the ERC has some drawbacks, including longer scanning times, increased costs and lower patient compliance due to the coil's placement in the rectum. The additional image resolution of the ERC is valuable on 1.5 T MRI, whereas it is more questionable on 3 T. Most prostatic MRI examinations can be conducted with acceptable image quality without an ERC due to increased spatial resolution (the ability to distinguish two dense structures from each other) and increased signal-to-noise ratio on 3 T MRI (47). However, a recent study found that 3 T MRI with and without an ERC evidenced similar diagnosis of overall and index PCa (48). According to the European Society of Urogenital Radiology's (ESUR) MR prostate guidelines, the use of an ERC is optional for detection and preferable for staging at 3 T MRI (49).

The quality of an MRI image is also influenced by the patient's preparation. The administration of an oedema prior to the examination and injection of an antispasmodic agents may diminish rectal peristaltic motion and reduce intra-luminal air which can cause MRI artefacts (50).

The development of MP-MRI provides new possibilities in the detection, lesion characterisation and staging of PCa due to its high quality and soft-tissue contrast. Several published data (12,51–53) have illustrated the rapidly growing use of MP-MRI as the most sensitive and specific diagnostic imaging modality for PCa

management. MP-MRI can reveal details about morphological, metabolic and cellular changes in the prostate as well as to characterise tissue vascularity (54,55). The use of MP-MRI criteria to follow up malignancy recurrence and treatment decisions may decrease the population needing TRUS biopsies by allowing these biopsies to be targeted to high-risk groups. Pre-biopsy MP-MRI, used as a triage, could improve the detection of clinically significant cancer (12).

MP-MRI contains different MRI sequences: anatomical imaging; diffusion weighted imaging (DWI); dynamic contrast enhanced imaging (DCE); and spectroscopic imaging (MRSI). The information from these techniques can be combined to provide more accurate results.

2.3.1 Anatomical MR imaging for prostate gland

Anatomical MRIs include T2-weighted (T2W) and T1-weighted (T1W) images. These display the anatomy of the prostate and its adjacent structure with high resolution. As part of MP-MRI, anatomical MRIs are useful when combined with other functional MRI techniques such as DWI and DCE imaging to obtain the optimal accuracy of the result (49).

2.3.1.1 T1-weighted MRI

T1W imaging is used in conjunction with T2W imaging to detect post-biopsy haemorrhage and assess the prostate's contour and neurovascular bundles (NVB). Because of its low spatial resolution, T1W imaging cannot be used to assess intra-prostatic zonal anatomy (56). On T2W imaging, post-biopsy haemorrhage can mimic PCa because both cancerous lesions and haemorrhage can appear as dark (hypo-intense) areas. It has been reported that it affects between 28 and 95% of patients (10,57,58). However, only haemorrhage on T1W imaging will appear as a high signal

intensity area, which can be used to rule out false-positive findings on T2W imaging (10) (Figure 2-3).

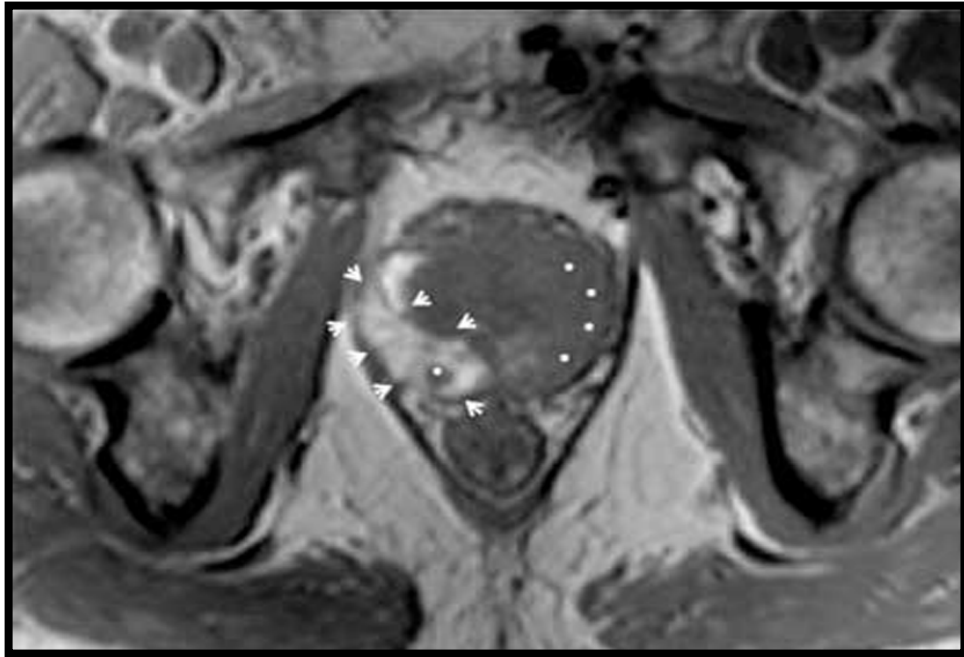


Figure 2-3 Axial T1 showing diffuse hyperintense signal in the right PZ (arrowheads) due to post-biopsy haemorrhage with area free of T1 hyperintense blood (asterisks) matched the PCa lesions (T1 haemorrhage exclusion sign).

The extent of haemorrhage in a PCa lesion is smaller than in adjacent benign tissue, and the presence of the excluded haemorrhage sign on T1W imaging combined with a homogeneous low signal intensity area on T2W imaging is highly accurate for PCa detection (58).

2.3.2.2 T2-weighted MRI

By providing a good picture of the prostatic zonal anatomy, T2W imaging with high spatial resolution can be used to detect, localise, and stage PCa (51). Often, the PZ has a higher signal intensity due to the high content of water in the glandular tissue as opposed to the transitional and CZ which often have a lower signal intensity (Figure 2.4A). The transitional and CZ is often referred in combination as the central

gland, as the two zones can be difficult to differentiate on MRI. However, awareness about the location and features of the CZ is important because its manifestation may mimic PCa, resulting in a false positive reading on MRI (Figure 2.4B). However, PCa in the CZ is uncommon, but when it is located there it is usually more aggressive (59).

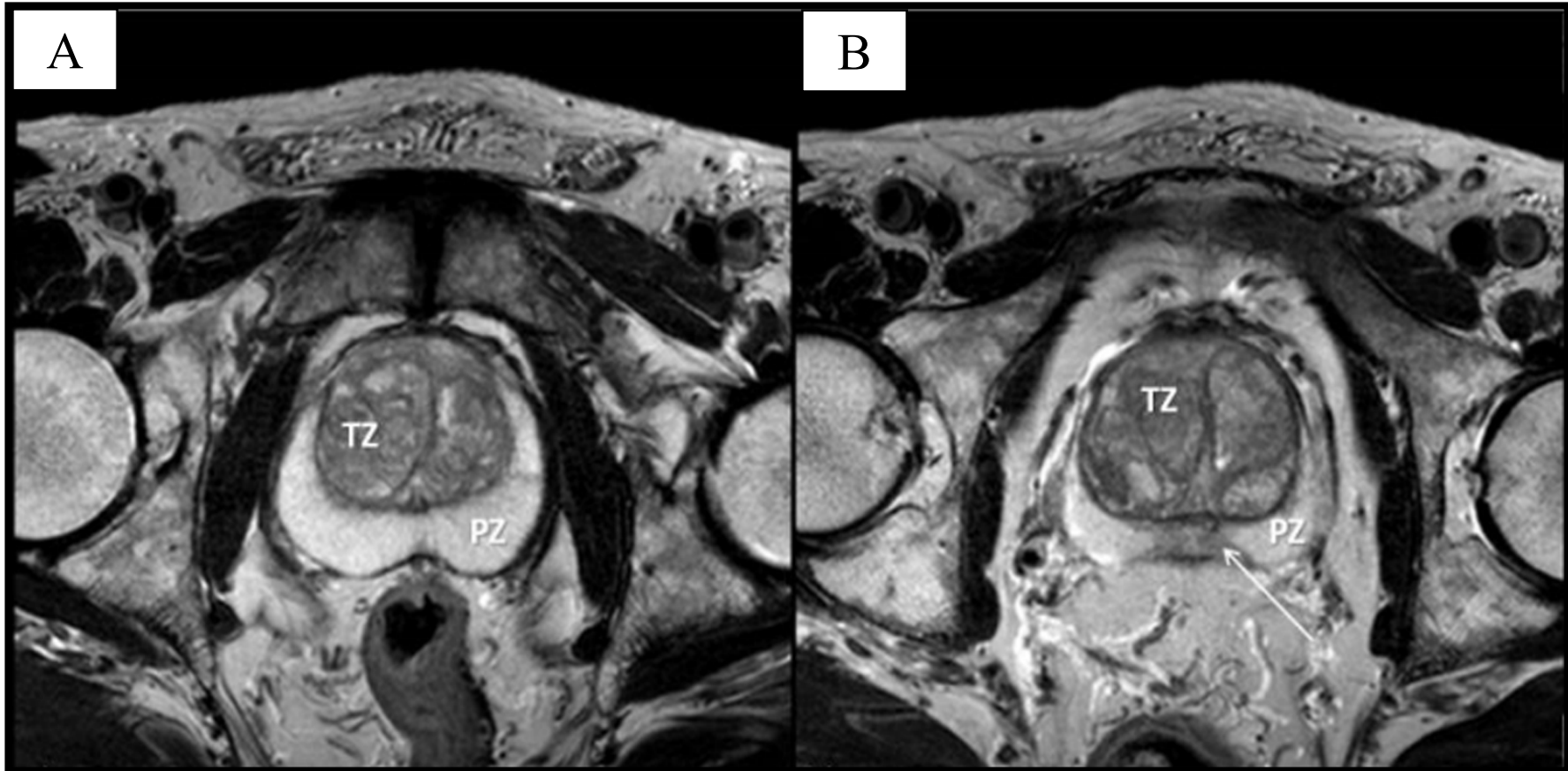


Figure 2-4 Normal prostate anatomy. (A) T2W images show the PZ and TZ in the axial plane. (B) Axial T2W image at the prostatic base shows the CZ (white arrow) as a hypointense area surrounding the ejaculatory ducts (60).

The prostatic capsule appears as a thin fibro-muscular fringe with lower signal intensity surrounding the prostate. In contrast to the higher signal intensity from the homogeneous benign PZ, PCa in the PZ typically appears as a round or oval area of low signal intensity (61,62) (Figure 2-5). On T2W imaging, however, some PCa lesions are iso-intense and cannot be seen. The use of T2W imaging alone therefore has limitations.

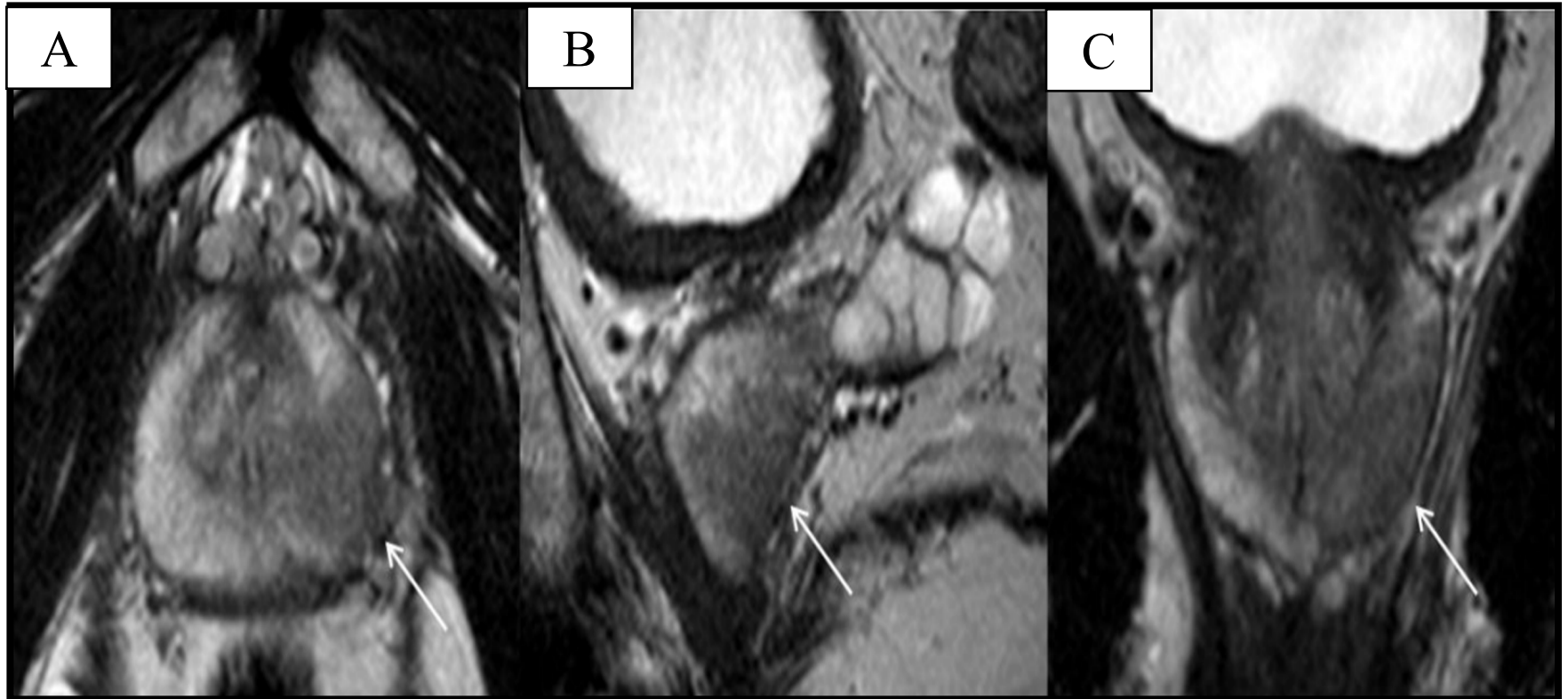


Figure 2-5 On (A) axial, (B) sagittal, and (C) coronal views, T2W imaging of a PCa lesion in the left PZ (white arrow) (60).

PCa occurring in the TZ is not as clearly defined because it often has lower and mixed signal intensities due to benign prostatic hyperplasia nodules that may interfere with diagnosis and mimic PCa. PCa in the TZ is characterised by a homogeneous low signal intensity area usually anteriorly located and has a lenticular shape (Figure 2-6) (63).

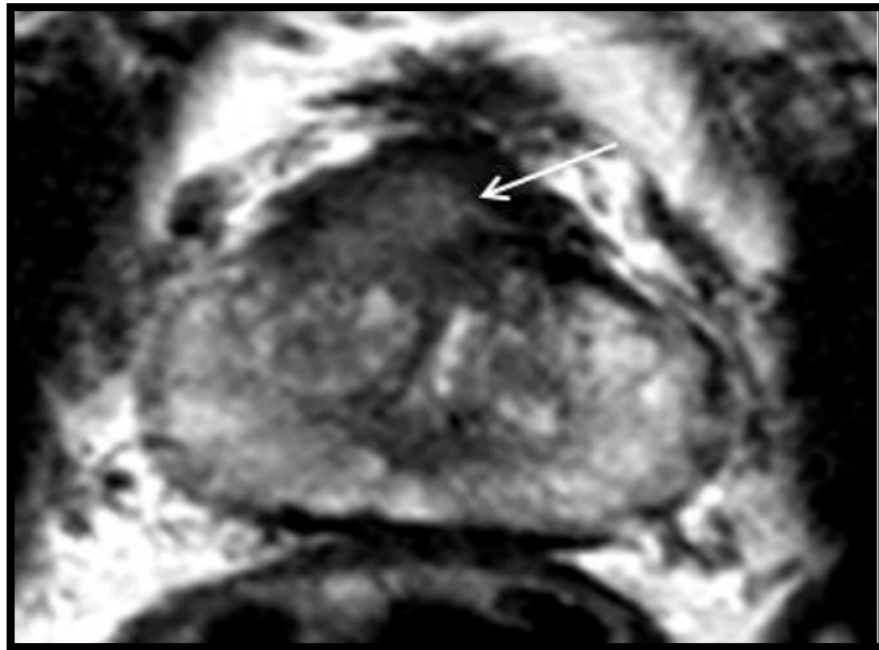


Figure 2-6 Axial T2W image of a PCa lesion in the prostate's right anterior part (white arrow) (60).

The degree of signal intensity on T2W imaging has been related to the GS as cancers with a Gleason grade 4 or 5 become more hypointense than cancers with a Gleason grade 3 (64). Further, the cancer's growth pattern may affect the appearance where sparse tumours with increased intermixed benign prostatic tissue appear more like normal PZ than more dense tumours (65). Moreover, several benign lesions in the prostate such as haemorrhage, atrophy, BPH, calcifications, and prostatitis may present as a low signal intensity region on T2W imaging, resulting in false-positive readings. According to a meta-analysis, T2W imaging alone had a sensitivity and specificity of 0.57-0.62 and 0.74-0.78, respectively (66). Due to this moderate sensitivity and specificity, T2W imaging can be used in

conjunction with other functional MRI techniques such as DWI and DCE imaging to improve diagnostic performance (67).

2.3.2 Functional MR imaging for prostate gland

2.3.2.1 Diffusion weighted imaging (DWI)

DWI is a non-invasive functional MRI technique that measures changes in water molecule diffusion as a result of microscopic structural changes. DWI generates different signal intensities that quantify the movement of free water molecules by applying different gradients (b-values) to the water protons in the tissue (68). The glandular structures in normal prostatic tissue, particularly in the PZ, allow water molecules to move freely without restriction. PCa often depletes glandular structures and causes restricted diffusion by containing more densely packed cells. Diffusion changes result in changes in the signal intensity on DWI: regions with restricted diffusion appear bright on DWI. DWI is usually performed with different b-values where low b-values (0-100) indicate a signal decay caused by perfusion in the tissue, whereas higher b-values represent water movement in the extracellular and intracellular compartment (69). DWI can be used to detect and characterise lesions by providing both qualitative and quantitative information about tissue cellularity and structure. Therefore, a qualitative assessment of a region with high signal intensity on high b-value DWI often represents a region with restricted diffusion caused by tightly packed cells.

The apparent diffusion coefficient (ADC) is calculated using the signal intensity changes of at least two b-values to quantitatively assess the degree of diffusion restriction. ADC is calculated using built-in software in the MRI scanner or workstation. An ADC_{map} is created based on the ADC value in each voxel of the

prostate. Restricted diffusion causes a reduction in the ADC value and appears dark on the ADC_{map} (69,70).

Compared to surrounding normal tissue, PCa has higher cellular density and restricted diffusion. As a result, PCa lesions appear bright on DWI with high b-values but dark on the ADC_{map} with lower ADC tumour values (70–72). As a result, DWI can help distinguish between malignant and benign tumours, and the use of DWI in the diagnosis of PCa has been shown to add sensitivity and especially specificity to T2W imaging carried out on its own (Figure 2-7) (73).

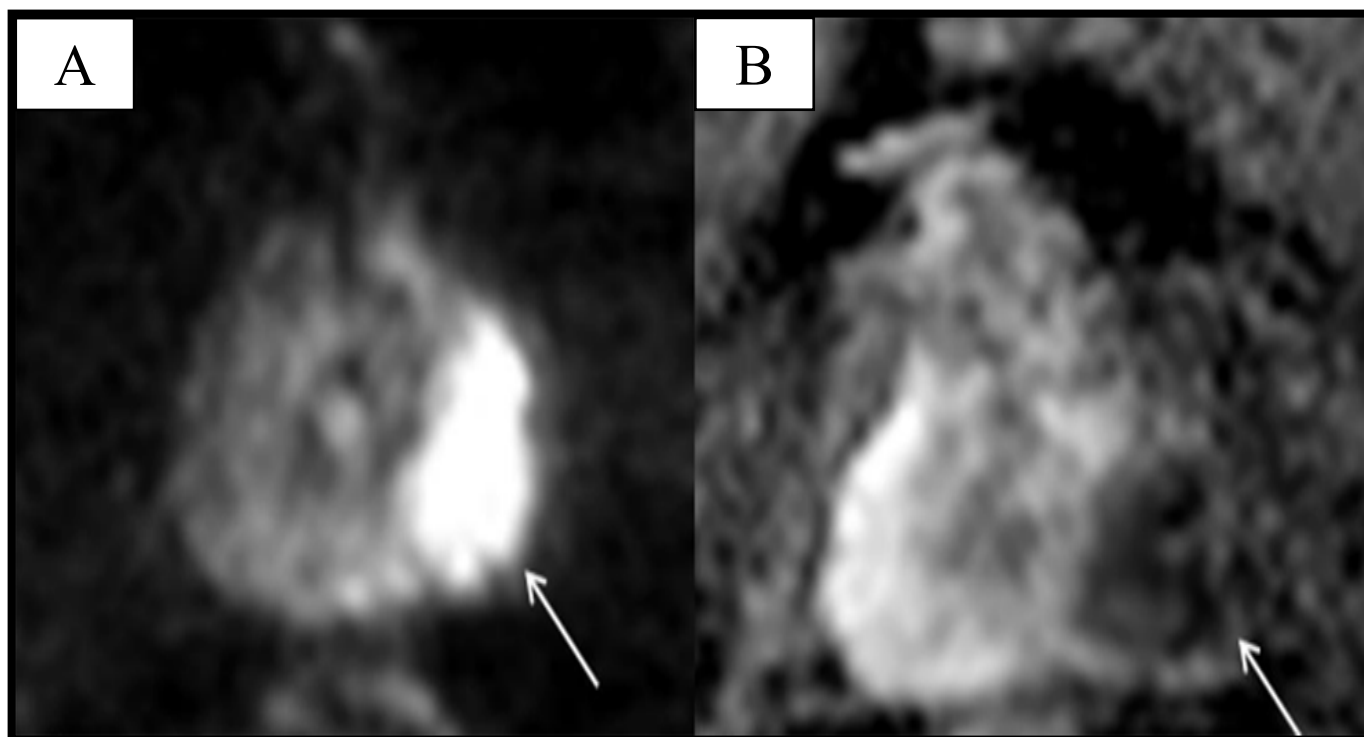


Figure 2-7 (A) DWI of PCa in the left PZ (white arrow) b1400 (B) ADC_{map} (60).

According to studies, the mean ADC tumour value measured from the cancerous lesion on the ADCmap and the GS have an inverse relationship (74–77), implying that ADC tumour values can be used as a non-invasive marker of tumour aggression. Attempts have been made to define precise cut-off values to distinguish malignant from healthy tissue and to further differentiate between GS groups. However, a wide range and inconsistency in mean ADC tumour values have been reported due to different study methodologies with different b-values, different MRI scan and field strengths, as well as patient variability between studies (74,76,78). Furthermore, remarkable overlap exists between ADC values from malignant and benign prostatic tissue, along with wide variability depending on the zonal origin. As a result, no agreement has been reached on absolute ADC tumour cut-off values for different GS (79–81).

2.3.2.2 Dynamic contrast-enhanced MRI

DCE-MRI exploits the fact that malignant and benign prostatic tissues frequently have distinct contrast enhancement profiles. The DCE-MRI method is based on changes in the pharmacokinetic features of the tissue mainly due to angiogenesis. DCE-MRI includes a series of fast high-temporal (the ability to make fast and accurate images in rapid succession) T1W images before, during and after a quick intravenous injection of a gadolinium-based contrast agent. Since prostatic tissue is usually heavily vascularised, a simple examination of pre- and post-contrast images is insufficient for PCa characterisation (82). Compared to normal prostatic tissue, PCa often causes angiogenesis and increased vascular permeability, resulting in a high and early contrast enhancement peak (increased enhancement) accompanied by fast washout of the contrast (Figure 2-8) (83,84).

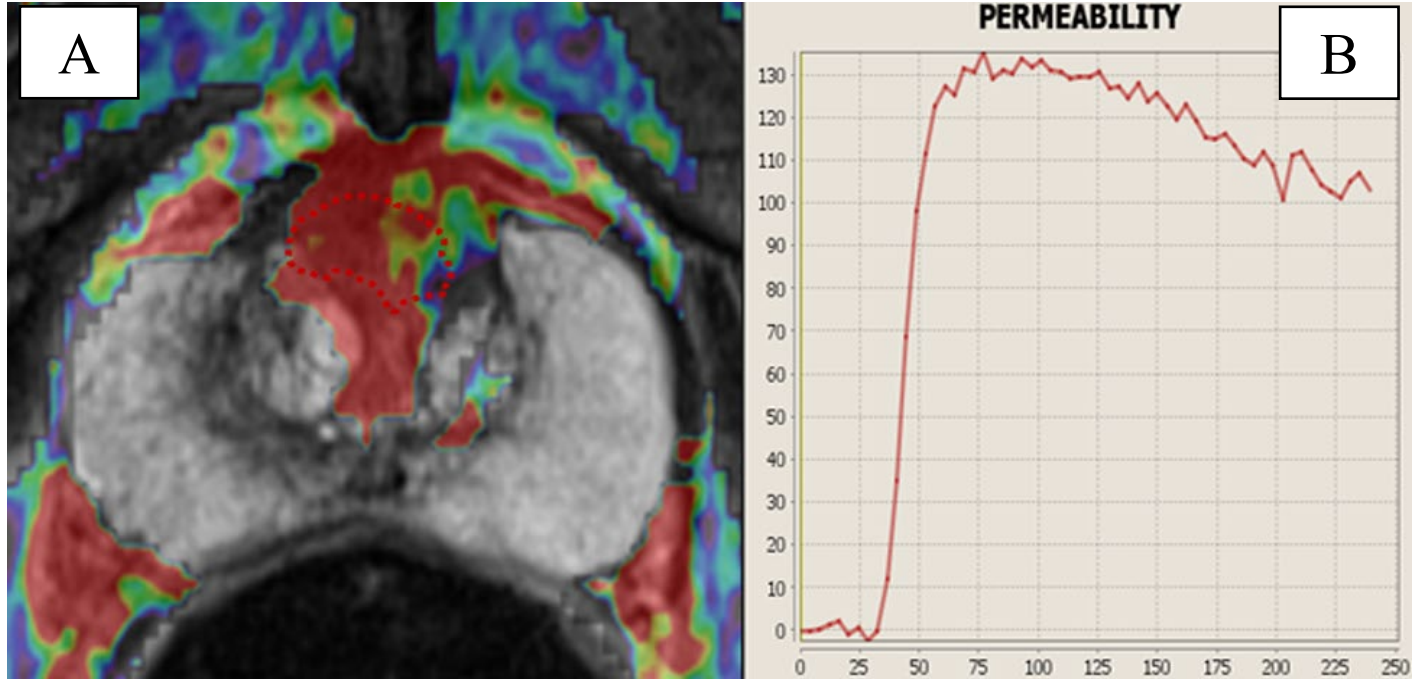


Figure 2-8 (A) DCE-MRI colour map shows early focal contrast enhancement in the TZ tumour (Red dotted circle). (B) DCE-curve which is a typical malignant curve with a high peak, rapid early enhancement (high wash-in rate) and early wash-out.

The pharmacokinetic features of the tissue can be defined using a number of methods. Qualitatively by visualising enhancement curves, and quantitatively by using detailed pharmacokinetic models to calculate the contrast exchange rate between different cellular compartments, or semi-quantitatively, by measuring various enhancement curve kinetic parameters such as wash-in/wash-out rate, time to peak. In addition, various post-processing software tools are used to analyse and describe the DCE-MRI, including overlaid colourised enhancement maps used to detect pathological changes and PCa. Previous research has shown that using DCE-MRI in combination with other MRI modalities can enhance PCa detection diagnostic accuracy (85,86) and improve extra capsular extension (ECE) detection (87). The utility of DCE-MRI primarily adds sensitivity to the MP-MRI performance and is essential for the detection of local recurrence. However, Baur et al (88) have stated that DCE-MRI did not add significant value to the detection of PCa. DCE-MRI has a low specificity as benign conditions such as hyper-vascularised BPH nodules and prostatitis can mimic pathological enhancement patterns. To reach the optimum sensitivity and accuracy for PCa evaluation, DCE-MRI should be used in combination with other MRI modalities such as T2W imaging and DWI (85,89).

2.3.2.3 Magnetic resonance spectroscopic imaging

MRSI displays the relative concentrations of chemicals within small volumes of interest, providing metabolic information about prostate tissue (voxels). Citrate is abundant in normal prostate tissue (higher in the PZ compared with the central or transition zone levels). Because the cells in PCa switch from a citrate producing to a citrate oxidising metabolism, citrate levels in PCa cells are decreased or undetectable. Choline levels are also elevated in proliferating malignant tissue due

to a high phospholipid cell membrane turnover (Figure 2-9). As a result, MRSI detects tumours based on an increased choline to citrate ratio (90).

However, because MRSI is technically challenging and requires a high level of expertise as well as a longer scan time, which is frequently combined with the use of an ERC, many centres do not include it in their standard protocol. MRSI is not required for prostate examination according to the ESUR MR prostate guidelines (91).

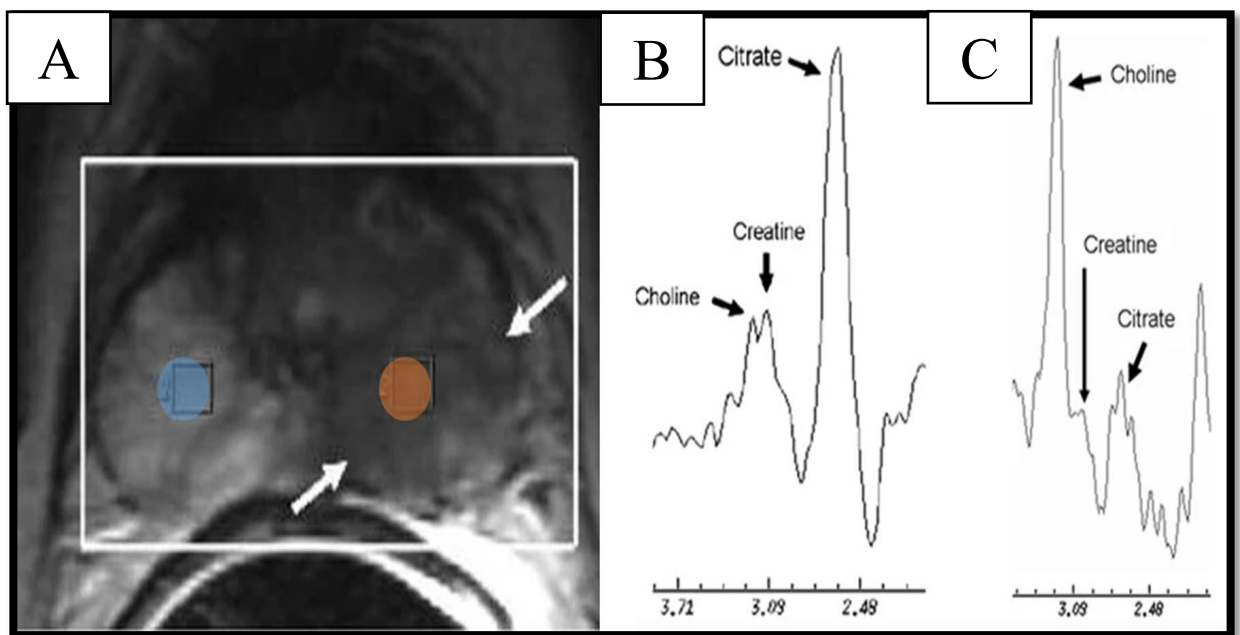


Figure 2-9 (A) The MRSI was acquired from the whole prostate showing a large low signal intensity lesion (arrows) in the left PZ. (B) The right voxel from the PZ (blue circle) shows the high signal intensity with the normal metabolic profile of the prostatic gland, high level of citrate and lower level of choline. (C) The corresponding spectrum from the left voxel shows increased choline and reduced citrate, indicative for a metabolic profile of prostatic cancer (orange circle) combined with the low signal intensity on MRI (92).

2.4 Prostate Imaging-Reporting and Data system (PIRADS)

The basic principle of a scoring system for MP-MRI readings is to detect abnormal regions and grade each region based on the appearance on the MP-MRI according to the degree of suspicion of PCa. However, interpreting prostate MP-MRI is difficult, requiring a steep learning curve; experienced readers are significantly more accurate than inexperienced readers (93,94). MP-MRI diagnostic accuracy varies between studies (95,96), owing to differences in study protocols' diagnostic criteria, MRI machines and expertise, which has sparked a debate about MP-MRI's readiness for routine use (97). Lack of standardisation and a consistent scoring system has been a source of debate for MP-MRI. As a result, clinical guidelines were recently published in order to promote high-quality MP-MRI acquisition and evaluation. The guidelines, including clinical indications for MP-MRI and a structured uniform PIRADS to standardise prostatic MP-MRI readings, are based on evidence from the literature and consensus from ESUR prostate MRI experts (49,98).

The PIRADS classification system, like the Breast imaging reporting and data system (BIRADS) for breast imaging, is a scoring system for prostate MP-MRI. It should include: 1) a graphic prostate scheme with 16-27 regions based on a five-point Likert scale; 2) a PIRADS score for each individual lesion; and 3) the largest lesion's maximum diameter. For each suspicious lesion within the prostate, all MRI modalities – such as T2W, DWI, and DCE imaging – are scored independently (1-5) on a five-point scale, and the sum of all individual scores (ranging from 3-15 for three modalities) constitutes the PIRADS summation score. Further, each lesion is assigned a final overall score (ranging from 1 to 5) based on the likelihood of clinically significant PCa being present (98) (Table 2-1).

Score	Criteria
T2WI imaging for PZ	
1	Signal intensity is uniformly hyperintense (normal).
2	Either Linear or wedge-shaped hypointensity or diffuse mild hypointensity with an indistinct margin.
3	Non-circumscribed, rounded, moderate hypointensity or heterogeneous signal intensity. Others that don't fit into the categories of 2, 4, or 5.
4	The focus/mass is circumscribed, homogenous, moderately hypointense, and confined to the prostate, with <1.5 cm in greatest dimension.
5	Identical to 4, but with ≥ 1.5 cm in greatest dimension or definite extraprostatic extension/invasive behaviour.
T2WI imaging for TZ	
1	A round, fully encapsulated nodule or a normal-appearing TZ (rare) (An example of a typical nodule).
2	A mostly encapsulated nodule or a circumscribed nodule that is homogeneous but not encapsulated. ("atypical nodule") or a homogeneous mildly hypointense area between nodules.
3	Non-circumscribed, rounded, moderate hypointensity or heterogeneous signal intensity. Others that don't fit into the categories of 2, 4, or 5.
4	Lenticular or non-circumscribed, homogeneous, moderately hypointense, and with <1.5 cm in greatest dimension
5	Identical to 4, but with ≥ 1.5 cm in greatest dimension or definite extraprostatic extension/invasive behaviour.
Diffusion weighted imaging	
1	No abnormality on ADC and high b-value DWI.
2	Hypointense linear/wedge shaped on ADC and/or hyperintense linear/wedge shaped on high b-value DWI.
3	On ADC, focal hypointense (distinct from background) and/or focal hyperintense on high b-value DWI; may be markedly hypointense on ADC or markedly hyperintense on high b-value DWI, but not both.
4	On ADC, the focal region is markedly hypointense, and on high b-value DWI, it is markedly hyperintense; <1.5cm in greatest dimension.
5	Identical to 4, but with ≥ 1.5 cm in greatest dimension or definite extraprostatic extension/invasive behaviour.
Dynamic contrast enhanced imaging	
-	No early or concurrent enhancement; or diffuse multifocal enhancement NOT corresponding to a focal finding on T2W and/or DWI; or focal enhancement corresponding to a lesion showing BPH characteristics on T2WI (including features of extruded BPH in the PZ).
+	Focal, it occurs before or at the same time as enhancement of adjacent normal prostatic tissues and it corresponds to a suspicious T2W and/or DWI finding.
Overall final score	
1	Clinically significant tumour is highly unlikely to be present.
2	Clinically significant tumour is unlikely to be present.
3	Clinically significant tumour is equivocal.
4	Clinically significant tumour is likely to be present.
5	Clinically significant tumour is highly likely to be present.

Table 2.1 T2W, DWI, and DCE imaging are classified using the PIRADS system. On T2W imaging, the PZ and TZ have significantly different anatomical appearances, so different PIRADS criteria are used for the two zones (49,98).

Each lesion should be evaluated for possible extra prostatic tumour extension (EPE) in addition to the PIRADS classification. The ESUR MR prostate guidelines include a table of MP-MRI findings along with a risk score stratified into different EPE criteria with concomitant tumour characteristics/findings (Table 2-2).

Criteria	Findings	Score
Extracapsular extension	Abutment	1
	Irregularity	3
	Thickening of the neurovascular bundle	4
	Bulge, capsule loss	4
	Extracapsular disease that can be measured	5
Seminal vesicles Expansion	Expansion of the seminal vesicles	1
	Low T2 signal	2
	Filling in of angle	3
	Enhancement and impeded diffusion	4
Distal sphincter	Adjacent tumour	3
	Effacement of the sphincter muscle with a low signal	3
	Sphincter enhancement that is abnormal	4
Neck of the bladder	Adjacent tumour	2
	Low T2 signal in bladder muscle	3
	Enhancement that extends into the bladder neck is abnormal	4

Table 2.2 EPE risk scoring of extra prostatic extension (49).

T4-disease includes invasion into the bladder neck, external distal sphincter, rectum, and/or side of the pelvic wall, although only the first two T4-findings are included in the ESUR EPE risk scoring. The most commonly used MRI modality for EPE assessment is anatomical T2W imaging. However, some of the categories (for

example, Seminal vesicle invasion (SVI)) also include functional imaging findings (enhancement and impeded diffusion risk score 4). Suspicion of EPE should be given an overall score ranging from 1 to 5 based on the likelihood of EPE being present. Therefore, the five-point scale is considered a continuum of risk with higher scores indicating a higher risk of EPE. However, not all categories include the total score range of 1-5, and functional imaging results, for example, are excluded from ECE evaluation. Functional imaging has been shown in previous studies to aid in the detection of ECE (99,100), particularly in less experienced readers. As a result, when applying functional imaging findings, personal opinion may influence the interpretation and overall impression of possible ECE.

2.5 Histological grades between biopsy and radical prostatectomy

Several studies have examined the relationship between the GS at needle biopsy and the RP (101). Following the consensus conferences of the International Society of Urological Pathology (ISUP), the Gleason grading system was updated in 2005 and again in 2014 (4,102). The ISUP recently endorsed the use of a validated Grade Group stratification system ranging from 1 (GS 6) to 5 (GS 9 or 10) in conjunction with the overall Gleason system to reduce the number of grading categories and facilitate more accurate disease stratification (102).

With an increase in men undergoing treatment other than RP, such as radiation therapy or AS, where the only tissue sampled is on the needle biopsy, biopsy grade has become more important in recent years. Not only is there a risk of undertreatment due to needle biopsy undergrading, but overtreatment (i.e. additional radiotherapy) is also a concern for men whose biopsies are overgraded. The issue of accounting for tertiary grade patterns is significant as they can be seen in nearly 20% of RP specimens (103). Despite recent advances, traditional diagnostic pathways still rely on TRUS guided prostate biopsy to obtain systematic needle biopsies of the prostate, which has recently been shown to have a sensitivity of only 48% for the diagnosis of 'clinically significant' cancer, defined as a GS of at least 4 + 3 or a maximum core length of at least 6 mm (7).

Epstein et al (2) found that the rate of upgrading from GS 6 on biopsy to GS 7 on RP was 36.3%. The incidence of upgrading can be influenced by several quantifiable factors including serum PSA levels, pathology weight, age, the extent of cancer on biopsy, and needle biopsy sampling. The difference in experience and skill in grading among pathologists is a more difficult variable to quantify, although it has been

shown that needle biopsy and RP GS are more closely matched when graded by urologic pathologists rather than general pathologists (101,104).

The incidence of downgrading was also reported by Epstein JI et al (2), who found that 12% of cases with a biopsy diagnosis of $3 + 4 = 7$ had GS 5–6 at RP. One explanation for these cases is that Gleason pattern 4 was overgraded due to tangentially sectioned small glands of pattern 3 on biopsy. Alternatively, the assigned biopsy GS may have been accurate in some cases, with such small foci of Gleason pattern 4 in the RP that it was either not recorded or unsampled deeper within the paraffin block. In cases where the GS 5–6 versus GS $3 + 4 = 7$ on biopsy is borderline, it is probably better to diagnose the case as GS 6 because undergrading due to sampling error is acceptable compared to overgrading due to grading error. Audenet et al (105) proposed a nomogram to predict the likelihood of upgrading patients on biopsy. This nomogram included several clinical variables such as age, PSA density, percentage of positive cores, and DRE in a first cohort of 896 patients. The accuracy of this model suggests that clinical factors are insufficient for assessing the disease's risk prior to treatment.

However, several studies have found that MP-MRI correlates with pathologic grade at surgery and can accurately distinguish between patients with low- and high-risk PCa (106,107). Ongoing efforts are being made to combine both imaging and conventional clinical variables in order to narrow the discrepancy between the GS of biopsy and RP Specimens and to improve the accuracy of predicting the upgrading.

2.6 MRI-targeted biopsy

In order to perform targeted prostate biopsy, three methods of MRI guidance are available: (1) cognitive fusion, in which the US operator simply points the biopsy needle at the prostate region reviewed prior to MRI that revealed a lesion; (2) direct MRI-guided biopsy, performed inside an MRI tube; and (3) software coregistration of stored MRI with real-time US, performed with a fusion unit. Each approach has its own set of benefits and drawbacks (108).

Cognitive fusion is simple, fast and requires no additional tools beyond the MRI and a conventional TRUS facility. The US operator does not need any additional training beyond that needed for traditional TRUS biopsy. In the comprehensive review of Moore et al (109), cognitive fusion was used in some 22 different studies. Although the evidence was minimal, cognitive fusion appears to be more accurate than conventional SB. The potential for human error in interpreting from MRI to TRUS without an actual overlay is a disadvantage of cognitive fusion.

Direct MRI-guided biopsy is conducted in-bore, that is within the MRI tube, by a radiologist who fuses a prior MRI demonstrating a lesion with a contemporaneous MRI to confirm the location of the biopsy needle. The patient is rescanned after each biopsy sample to confirm localisation. Only a few targeted cores are typically taken; systematic sampling is not done. The Barentsz group at Radboud University in Nijmegen, the Netherlands, published a major experience with in-bore biopsy (110). The small number of cores taken, the precise localisation of the biopsy, and the reduced detection of insignificant tumours are all advantages of this procedure. The drawbacks of this method include the time and expense required, along with the in-bore time and the two MRI sessions necessary to obtain the biopsy specimens. Furthermore, since only suspicious lesions are sampled, tissues with a 'normal' MRI

appearance are missed, which is problematic because any false-negative aspects of prostate MRI are unknown. The third approach for MRI guidance of prostate biopsy is MRI–TRUS fusion. In this method, the operator images the prostate using US, as performed for the past several decades. When viewing the prostate, the MRI of that prostate, which is performed beforehand and stored in the device, is fused with real-time ultrasound using a digital overlay, allowing the target(s) previously delineated by a radiologist to be brought into the aiming mechanism of the US. The fusion produces a three-dimensional reconstruction of the prostate, and the aiming and monitoring of biopsy sites takes place on the reconstructed model. This approach has the drawbacks of being indirect, requiring the use of an additional device, and requiring advanced operator training. The benefit is that it can be conducted within minutes in an outpatient clinic setting under local anaesthesia (108). The best MRI-TB method remains a matter of debate in the literature. However, several studies have concluded that MRI–TRUS fusion-guided biopsy is significantly more reliable and cost-effective than visual registration and in-bore biopsy (111,112).

MRI targeted fusion biopsy has been shown to outperform SB in the diagnosis of clinically significant cancer in many previous studies (12,17,113). Despite the improved detection of clinically significant cancers with MRI-targeted biopsies, debate still remains whether MRI-TB can be used instead of or in addition to SB. Ahdoot et al (20) found that combined biopsy improved the detection of all PCa in patients with MRI-visible lesions. However, MRI-TB alone underestimated the tumour's histologic grade. Following RP, upgrades to grade group 3 or higher on histopathological analysis were significantly lower after combined biopsy.

2.7 Nomogram construction

Iasonos et al (114) published the following steps (Table 2-3). The first step in developing a nomogram is to clearly define the population to which it will be applied in clinical practice. The population should be defined with detailed inclusion and exclusion criteria. When creating nomograms, it is preferable to use population-level data because it allows the model to be more generalisable. Clinicians' abilities to correctly apply a nomogram's predicted probabilities is entirely dependent on their understanding of the population on which the nomogram was based.

Initial steps in constructing nomogram
Define the patient population: Data source in Single institution, Multicentre
Identify the outcome of interest: What dose does the nomogram predict?
Make a list of all the predictors that you are considering.
Construct the nomogram: Model selection, Select variables
Validation: Data source (Internal data, External data)

Table 2.3 Important Steps in constructing a Nomogram.

The operational definition of the outcome of interest is the second step in creating a nomogram. When creating nomograms, it is crucial to precisely and accurately describe the outcome. The entire point of a nomogram is to be able to predict the probability of an outcome occurring. As a result, the result must be as precise and accurate as possible, as well as simple to understand for both clinicians and patients. The third step is to select several covariates (variables) with a pathophysiological link to the desired outcome. All potential variables associated with an outcome should be included in the model when creating a nomogram, as long as they fit into

the pathophysiology of developing the outcome. The covariates should be chosen in advance and based on clinical evidence rather than statistical significance.

The fourth step in creating a nomogram is to choose a prediction model, enter the covariates into the model, and make sure that the prediction model's statistical assumptions (regression) are met. The type of regression to use for a nomogram is determined by the outcome's scale of measurement. Logistic regression, multinomial logistic regression, and proportional odds regression are used to construct nomograms for categorical and ordinal outcomes. If the 'time-to-event' for developing an outcome is the basis for constructing a nomogram, then Cox regression analysis can be used. Multiple regression may be used to build nomograms for continuous outcomes. Poisson regression or negative binomial regression can be used to construct nomograms for count outcomes (naturally skewed distributions).

The prediction model is finalised using validation, discrimination, and calibration methods in the fifth step. Cross-validation (split-group validation and bootstrap validation) and external validation are two methods of validation (to see if the model holds up with a new population or dataset). The predictive accuracy of discrimination refers to how well a model distinguishes between patients who have and do not have the desired outcome. Discrimination is evaluated by the concordance index (or C-index) obtained from ROC analyses (c-statistics or area under the curve (AUC)). The calibration of regression models is calculated by plotting the predicted probabilities of the model against the actual probabilities.

The final step is to interpret the validated nomogram and evaluate it for concordance and model fit. Each covariate's relative effects are converted to a point scale ranging from 0 to 100. These 'scores' for each covariate are added together and correspond

to the regression model's predicted probability of a patient having the outcome of interest (Figure 2-10).

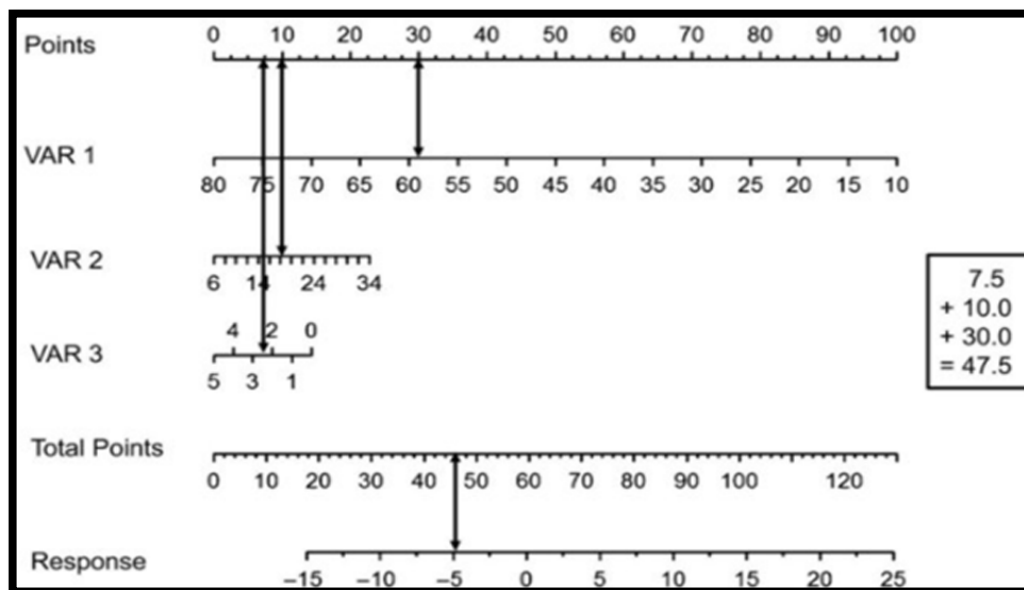


Figure 2-10 The nomogram depicts the impact of various predictive variables on various horizontal lines. The lengths of the different lines vary depending on the influence of each predictor. The greater the influence of a horizontal line, the longer it is. A number of points on the respective horizontal line represent the influence of each predictor. The anticipated magnitude of response can be read on the response horizontal line at the bottom of the nomogram by adding the points associated with each predictor (115).

The clinical utility of nomograms has been a source of debate. Increases in predictive accuracy related to the use of nomograms cannot only be statistically significant but also clinically meaningful. For example, preoperative nomograms estimating the risk of positive surgical margins, and lymph node metastases, may help clinicians in identifying patients who would derive greater benefit from more extensive surgery. Patients and physicians may benefit from postoperative nomograms that estimate recurrence, cancer specific survival, overall survival, the benefit of adjuvant therapies, and the impact of treatment on quality of life. However, their proper clinical application requires a thorough understanding of the nomogram-specific question,

study population, method of construction, and outcome, in order to accurately determine its applicability to a specific patient's clinical condition. Furthermore, the ability to analyse nomogram performance and assess limitations is needed in order to properly advise patients on the context, precision, and assumptions underlying nomogram risk estimations (116–118).

2.8 Decision-analysis curve

The definitive answer to determine if nomogram assisted decisions improve patient outcomes lies in prospective evaluation and randomising patients to nomogram or non-nomogram-based decisions and comparing the outcomes. However prospective validation of every nomogram prior to use is time consuming and impractical. Other method exists to evaluate the effects of prediction models on clinical decisions (118). Vickers and Elkin suggested decision analysis curves to estimate the clinical utility of prediction models using the threshold probability (probability that triggers a medical intervention by a physician or patient, equating to the probability at which the harm of a false-positive intervention exceeds the harm of a false-negative non-intervention) (119). The net benefit (defined as the fraction of true-positives subtracted by the fraction of false-positives weighted by the relative harm of a false-positive and false-negative result) is calculated using the threshold probability. A decision analysis curve is generated by plotting the net benefit against the threshold probability, which can then be used to assess the net benefit of nomogram at various threshold probabilities, compared to the net benefit of decisions made under the assumption that either all or none of the patients have the outcome of interest. For example, if a physician's threshold probability of dissecting the seminal vesicle during a RP is either $< 5\%$ or $> 50\%$ risk of SVI, decisions curve analysis at these threshold probabilities are irrelevant because the net benefit is the same if all or none of the patients have SVI (Figure 2-11).

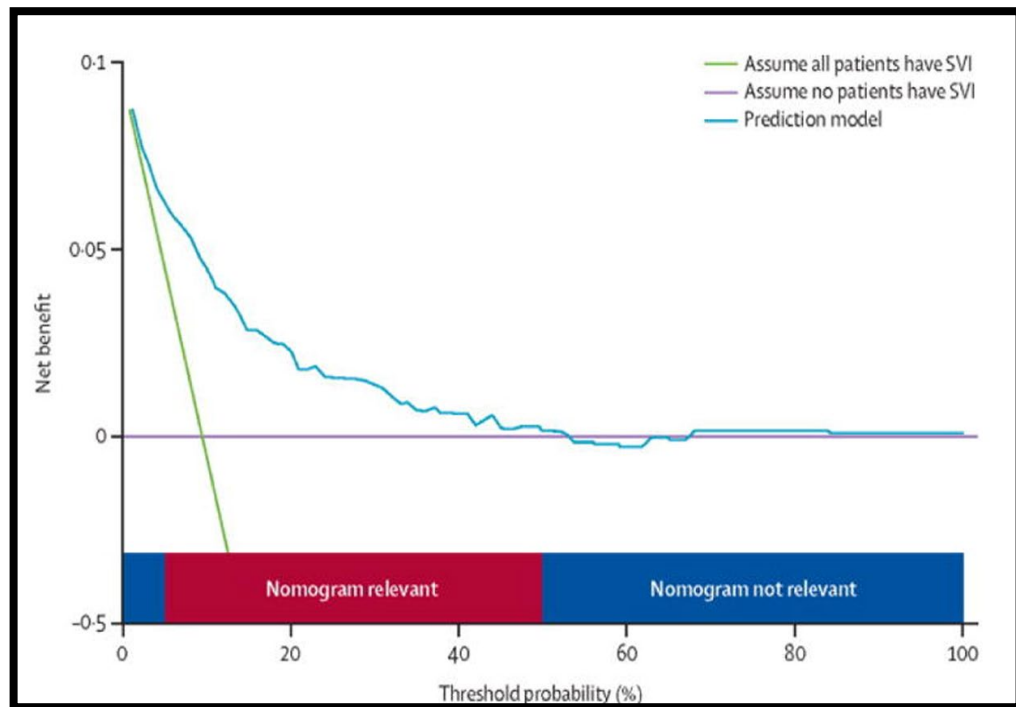


Figure 2-11 At a threshold probability of < 5%, > 50%, the nomogram is irrelevant (118).

2.9 Summary

The prostate gland is a conical fibromuscular pelvic organ containing three zones surrounded by extraperitoneal fat. PCa is an androgen-dependent malignancy which is the most common cancer in male patients and the second most common cause of cancer-related deaths. GS is believed to be the reference standard for PCa determination and aggressiveness. Several recent reports support the utility of pre-biopsy MP-MRI and image fusion of MRI with real-time US to address many of the limitations.

Firstly, an increase or 'upgrade' in GS in prostate cancer following biopsies remains a significant challenge to overcome. The second limitation addressed in this thesis was the development of a prediction model to identify patients who may benefit from performing SB at the time of TB in men suspected of having prostate cancer.

In the following chapters, the first part was aimed to evaluate whether pre-biopsy MRI has the potential to narrow the discrepancy of histopathological grades between biopsy and radical prostatectomy using PIRADS v2.0. Moreover, the second part was aimed to compare the diagnostic accuracy of TB, SB and combined approaches in the detection of csPCa and define predictive factors where combined approach could be used, and to quantify additional benefits of adding SB to TB methods by constructing a nomogram and assessing its net-clinical benefits.

Chapter 3 Prediction of prostate cancer Gleason score upgrading

3.1 Materials and methods

3.1.1 Study population

This is a study with prior Caldicott institutional approval (Caldicott/IGTCAL5626). All experiments including the study protocol followed approved institutional guidelines. The study had ethical approval (14/ES/1070) with each participant providing informed consent to the use of their imaging data. Between July 2014 and January 2019, 330 men treated by LRP who were diagnosed with localised PCa with raised PSA or/and abnormal DRE were retrospectively included in this study. They were offered MP-MRI and those with positive MRI results (PIRADS score 3 and above) subsequently had a TRUS-guided prostate biopsy (12 cores). Of these patients, eight were excluded because of contraindication to MRI such as a heart pacemaker and metallic foreign body including three claustrophobic patients. Further analysis occurred with the remaining 322 patients (Figure 3-1).

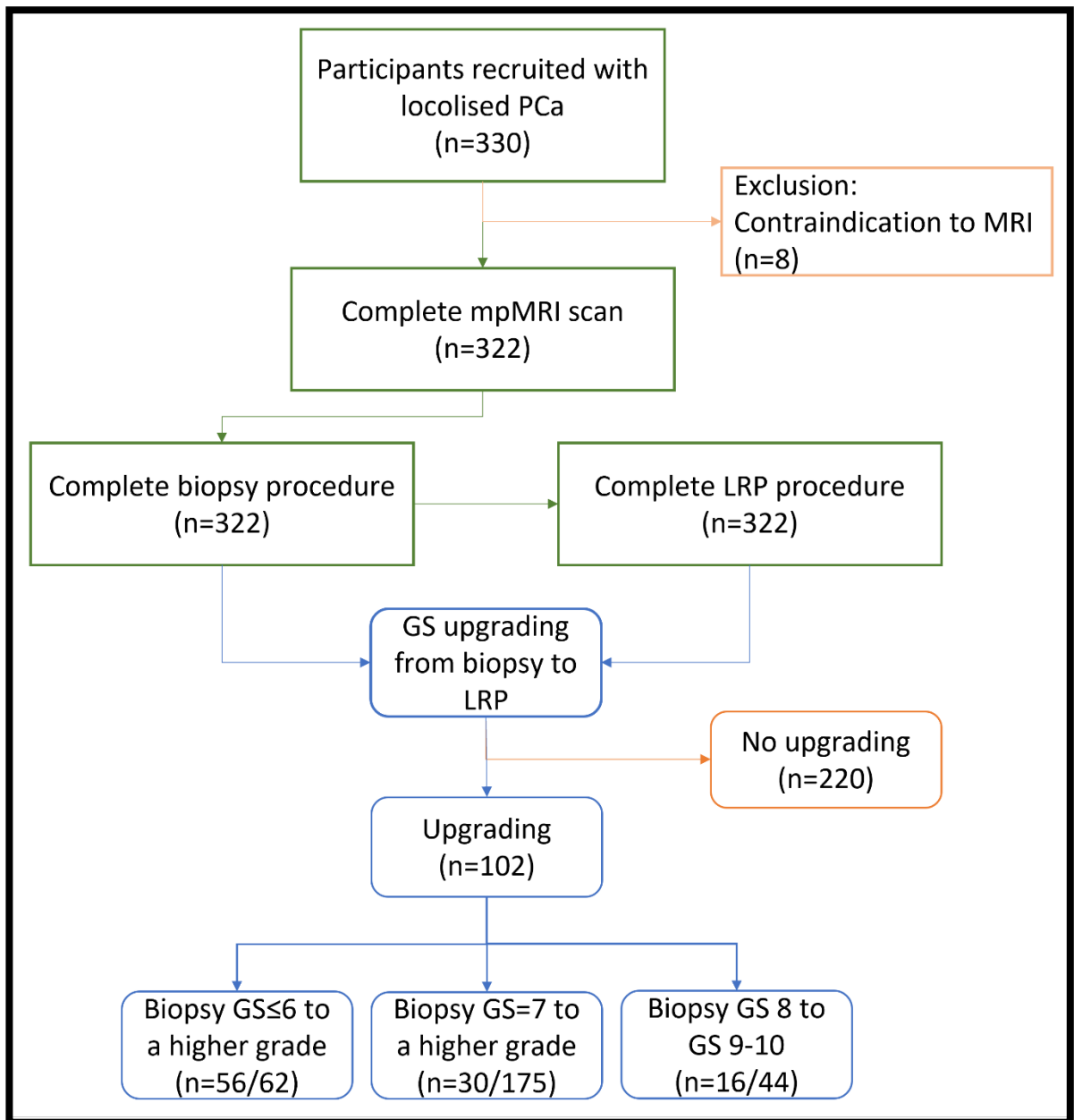


Figure 3-1 Flowchart of the study.

The clinical, pathological and imaging factors information of the patients, including age, weight, preoperative PSA, PSAD, number of positive cores, maximum percentage of cancer per core and PIRADS v2.0 score on MP-MRI were recorded. GS upgrading defined as a biopsy GS increasing from lower to higher grade on reported before (2). Table 3-1 summarises the baseline characteristics between upgraded and non-upgraded groups of the cohort.

We hypothesised that pre-biopsy MRI with PIRADS classification of a suspicious area in prostatic cancer would improve predictions of GS upgrading from biopsy to RP. Upgrading of GS on histology was defined as change of GS from lower to higher grade between biopsy and histology from RP.

	Total	Upgrading	No upgrading
Number of patients	322	102	220
Age (y), mean ± SD (range)	66.83±5.9(44-77)	66.82±6.12(49-77)	66.85±5.9(44-77)
Prostate Weight mean ± SD (range)	63.7±30.13(12-207)	65.3±26.2(20-155)	63.1±31.78(12-207)
PSA level (ng/ml), mean ± SD (range)	11.1±7.39(0.1-47.7)	12.6±9.98 (2-47.7)	10.39±5.7(0.1-41)
PSA Density (ng/ml²), mean ± SD (range)	0.261±0.234 (0.001-3.48)	0.212±0.183(0.035-1.11)	0.203±0.254(0.00198-3.48)
Number of positive cores	4.8±3.4(1-12)	4.1±3.07(1-12)	5.08±3.42(1-12)
Maximum percentage of cancer per core	50.2±30.4(5-100)	42.5±30.6(5-100)	53.3±29.52(5-100)
PIRADS from MP-MRI Benign (1,2)	17 (5%)	4 (4%)	13 (6%)
PIRADS 3	21 (7%)	6 (6%)	15(7%)
PIRADS 4	78 (24%)	26 (26%)	52 (23%)
PIRADS 5	206 (64%)	66 (65%)	140 (64%)

Table 3.1 Patient characteristics.

3.1.2 MRI protocol and PIRADS score

All patients' MP-MRI scans were performed on 3T scanner (TIM Trio, Siemens, Erlangen, Germany) 2 weeks before TRUS-guided biopsies. The MP-MRI protocol was derived from the European Society of Uro-radiology Guidelines 2012 for the detection of PCa and the subsequent publication of version 2 (120). Table 3-2 briefly summarises the MRI acquisition parameters. Localiser images were acquired in all three imaging planes, whereby the plane of the prostate was defined in relation to the rectal wall.

	T1WI	High resolution T2WI			DWI		DCE
	Axial	Sagittal	Axial	Coronal	DWI	DWI high b-value	Dyn Gd-MRI
Sequence	2DTSE	2DTS E	2D TSE	2DTS E	2DEPI	2DEPI	3D VIBE
TR (ms)	650	6000	400 0	5000	3300	3300	4.76
TE (ms)	11	102	100	100	95	95	2.45
Flip angle (°)	150	140	150	150	—	—	10
Slice thickness (mm)	3	3	3	3	3	3	3
Slice gap (mm)	0.6	0.6	0.6	0.6	0	0	0.6
Resolution (pixels)	320	320	320	320	192	192	192
FOV (mm)	200	200	200	200	280	280	280
b-values (s/mm²)	—	—	—	—	50,100,500, 1000	2000	—
Temporal resolution (s)	—	—	—	—	—	—	4

Table 3.2 MRI acquisition parameters.

The MP-MRI images were analysed and scored by experienced uro-radiologists using PIRADS v2.0; and the radiologists were blinded to all patients' pathology results (Figure 3-2).

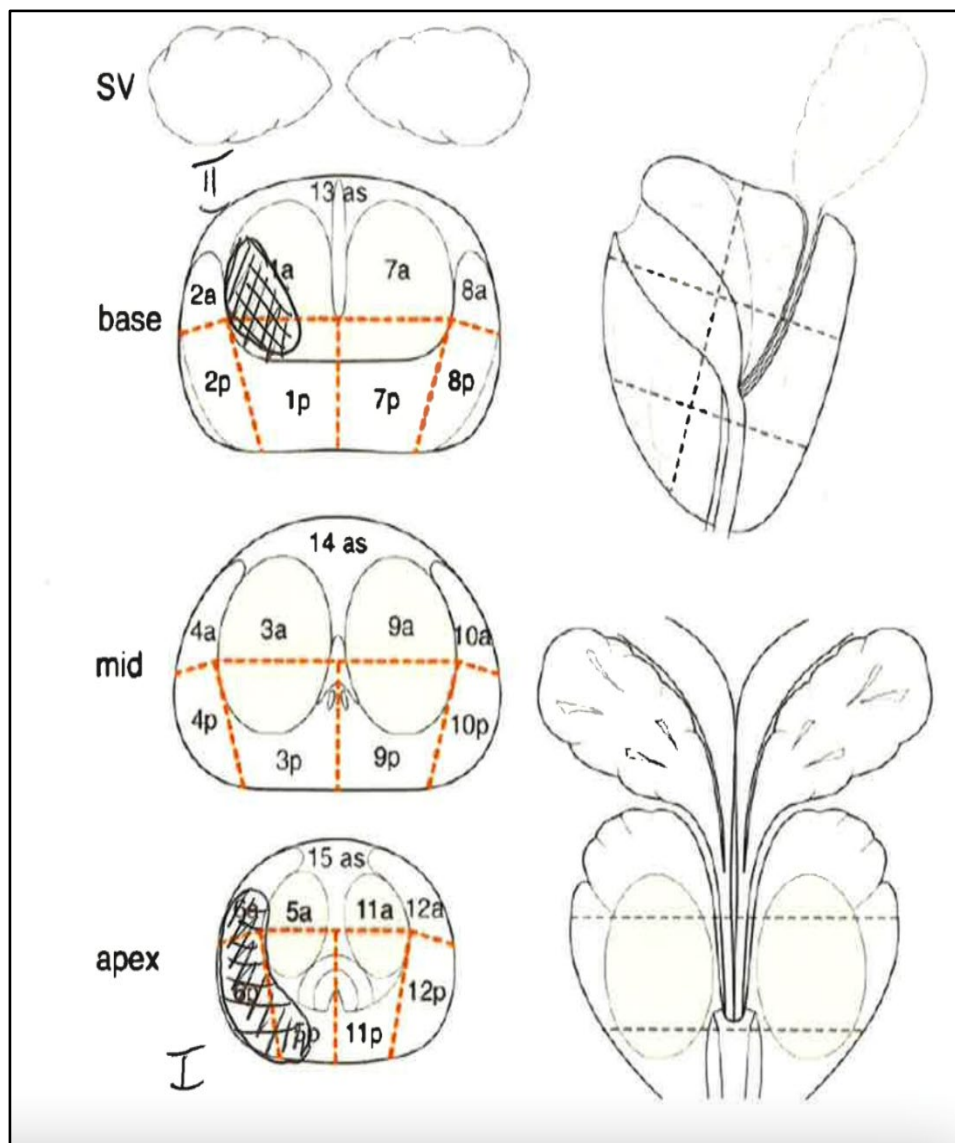


Figure 3-2 lesions localised and graded in MRI

PIRADS v2.0 assessment categories were described as follows: score 1, clinically significant cancer is highly unlikely to be present; score 2, clinically significant cancer is unlikely to be present; score 3, the presence of clinically significant cancer is

equivocal; score 4, clinically significant cancer is likely to be present; and score 5, clinically significant cancer is highly likely to be present (Figure 3-3).

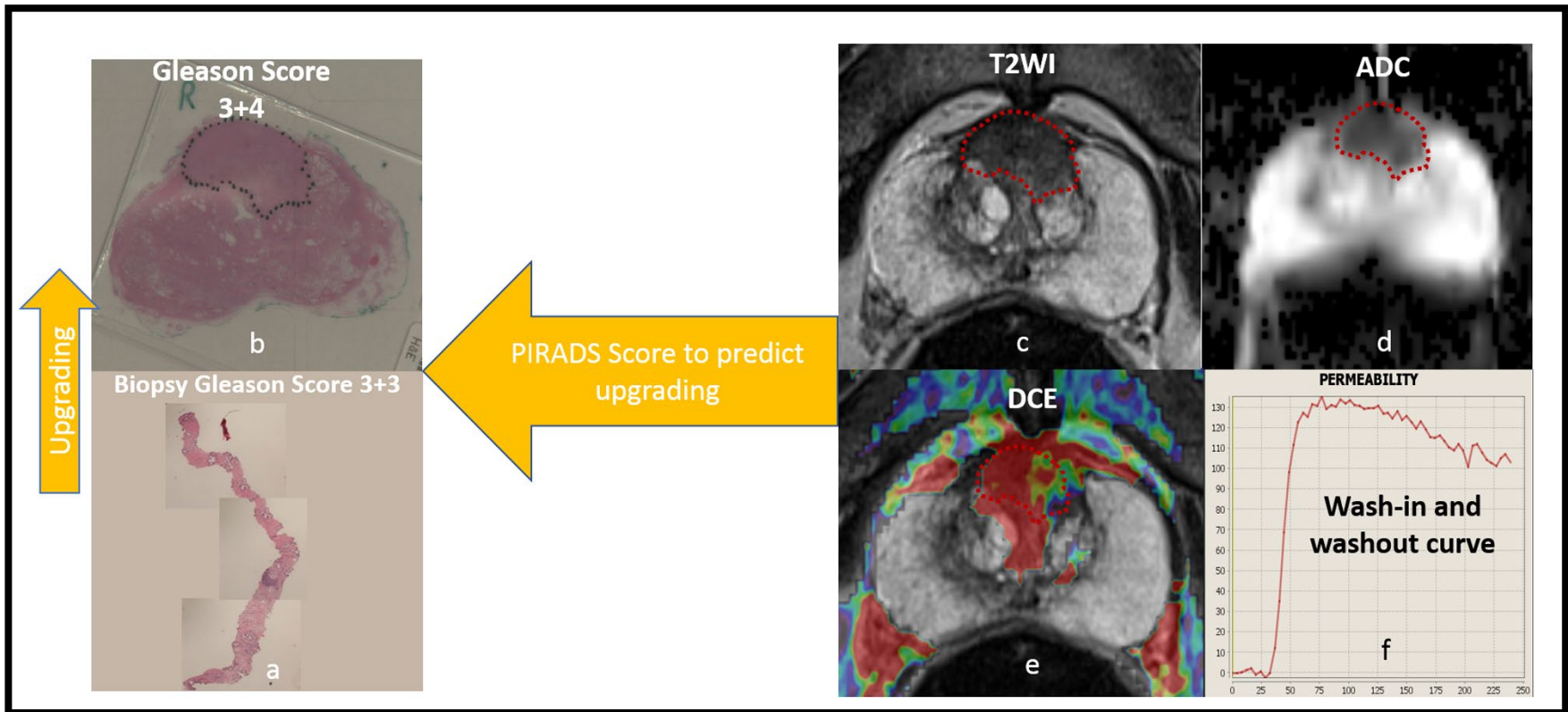


Figure 3-3 a: A 73-year-old man with GS 6 disease on prostate cancer on TRUS-Guided biopsies. b: The grade was upgrading to GS 7 on whole mount RP specimen c: Axial T2-weighted image shows ill-defined homogeneous low-signal-intensity on the CZ (d), ADC shows restricted diffusion in low-signal mass, and (e) DCE shows fast and strong enhancement and early contrast agent washout (type 3 curve) (e,f). The lesion was scored as PIRADS v2.0 5 (>1.5cm) and based on the parameters described here.

3.1.3 Histopathology data and analysis

The biopsy results were analysed by experienced pathologists, who were blinded to MRI findings. The GS for each patient was obtained. The radical prostate specimens for histology were sliced in patient-specific moulds to aid orientations between imaging and histology, which were fabricated using a 3D printer as described by our group and others previously (121,122). Specifically, patient specific 3D printed moulds were made prior to surgery based on the T2-weighted MRI prostate capsule the moulds were customised for each patient using MIMICS and Solidworks.

3.1.4 Mould design: a better standard reference

Figure 3-4 depicted the method for creating patient-specific moulds based on MR imaging and prostate sectioning during histopathology processing, but there is still a thorough discussion below.

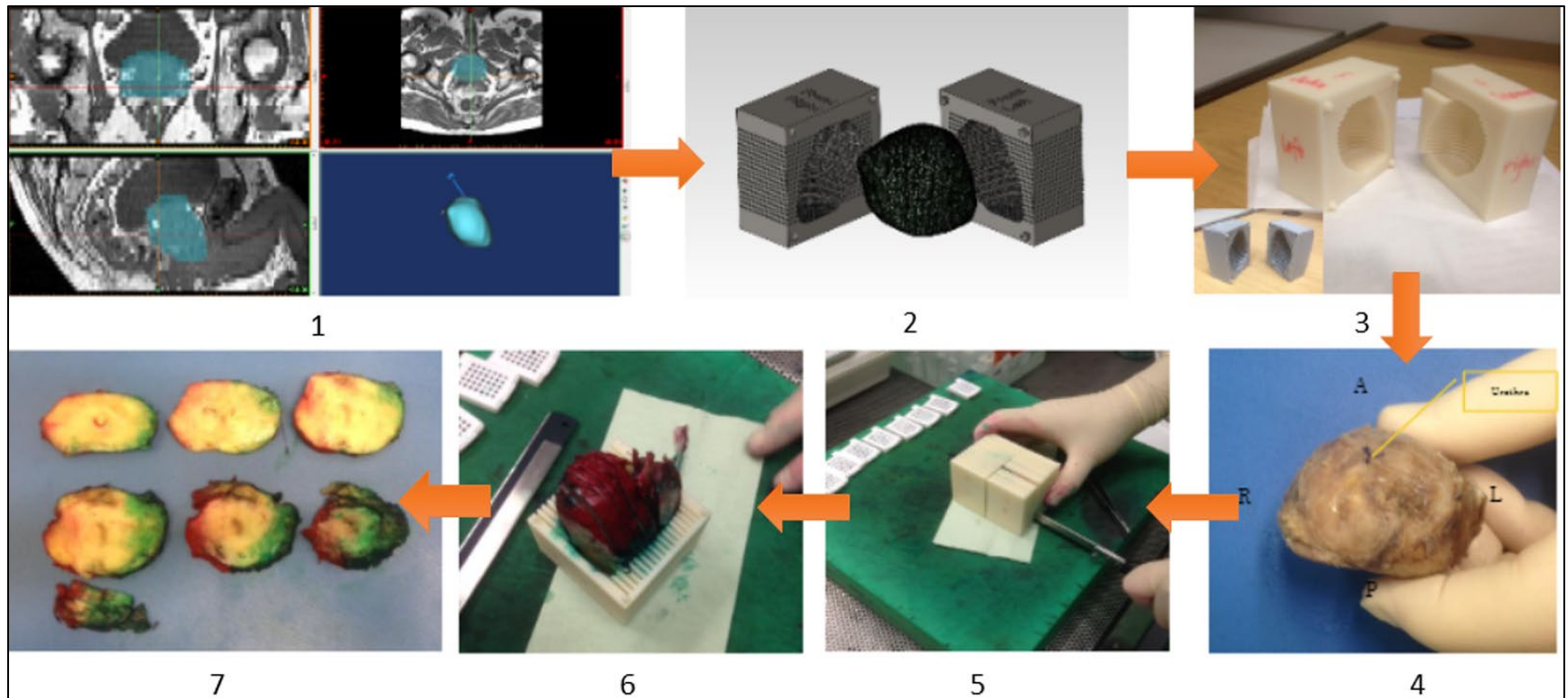


Figure 3-4 Steps of Method: (1) segmentation of MRI data in biomedical software MIMICS, (2) mould making in CAD software SolidWorks, (3) 3D printout from rapid prototyping machine Uprint, (4) post-radical prostatectomy specimen before dyeing and mould placement, (5) slicing of the prostate specimen with a single-blade, (6) final slices shown in the mould, and (7) the tissue slices are arranged from the apex to the base (123).

There are three factors to consider while creating such a mould and guaranteeing a higher pathological standard reference for prostate specimens:

1. Prostate gland segmentation in MR images: a critical step in customising each patient's prostate. The outer dimensions of the mould block had to be modified because the size of the gland differed across patients.
2. Measuring the real size of each prostate: following surgery, prostate specimens were somewhat altered, and the size measurement allowed the final design of the mould to be adjusted to match the prostate.
3. Aligning Cutting Slots: For each patient, the number of slots needs to be changed.

Participants received MRI scan. T2WI was used to produce three planar images of the prostate (axial, coronal, and sagittal). The scan resolution for the axial image was $0.63 \times 0.63 \text{ mm}^2$, and each slice was 3mm thick with a 0.6mm gap. After a thorough examination of these 2D pelvic pictures, skilled radiologists used the programme MIMICS (Medical Image Segmentation for Engineering on Anatomy) to determine the prostate capsule's borders in one direction, the prostate was segmented. There were 7-12 2D pictures in the axial direction (Figure 3-5 and Figure 3-6) that could be fused and verified to create a 3D model, saved as stereolithographic (STL) file.

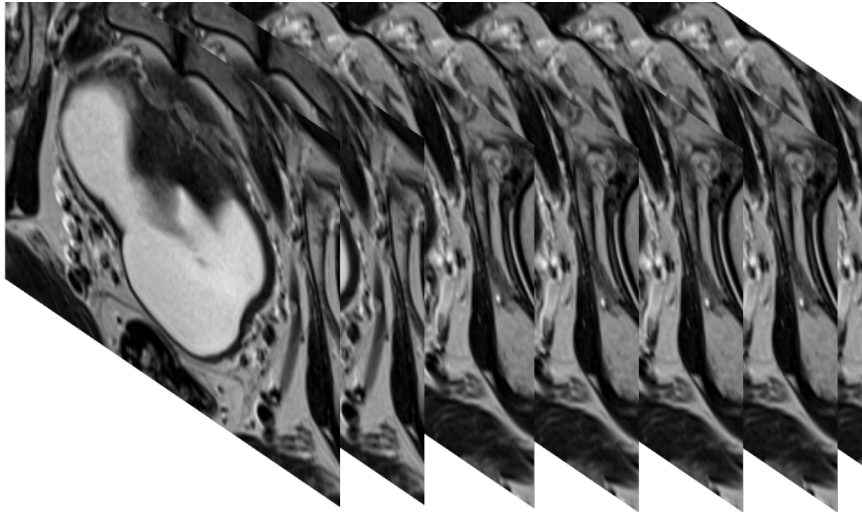


Figure 3-5 MRI's slices for a specific patient

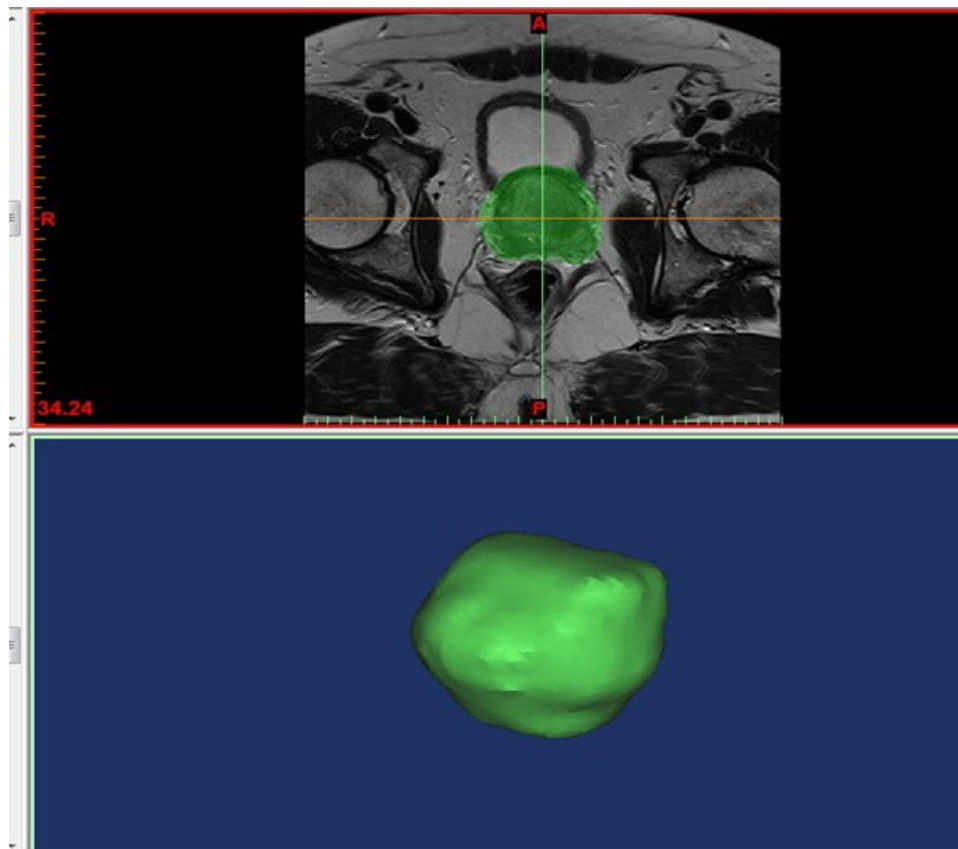


Figure 3-6 Segmentation of the prostate (green area) of MR imaging

One 3D model was merged and transformed into an object simulated in MIMICS software depicting the patient's prostate topography in the axial direction of segmentation and validation with the other two directions. To avoid any minor errors, the 3D prostate model was checked and then compared to the prior prostate measurement before any modifications were made. When the fused 3D model's capsule was verified, it was rough, thus smoothing was performed to the capsule's surface. The coarse model was loaded into the software of the CAD Meshmixer (2011 Autodesk, Inc.) to make the mould creation process easier by smoothing it out. To reduce file size, the smoothed model was also saved as a mesh model in STL format, which was then imported back into MIMICS for triangle reduction.

SolidWorks (3D CAD analysis software, design software, Dassault Systèmes SolidWorks Corp. USA) was used to import the triangle-reduced mesh model as a SolidWorks Part (SLDPRT) model. The mesh or the triangular surfaces of the SLDPRT model had to be regulated under a particular range for a successful conversion; otherwise, the importing process would be impacted and even collapse. After wrapping the solid simulated prostate model in a cubic or rectangular shaped block for the model and performing a "subtraction" (combine in feature) on the two models, the interior cavity that exactly housed the patient's prostate was generated in the newest model. Because patients' prostates had diverse forms and sizes, the outer dimensions of the moulds varied. After determining the prostates positioning in the mould, slots for specimens sectioning were put in the mould in an axial direction based on 2D images from axial MR imaging. The size is 1.2 mm thick of each slot, easy to cut off with a 0.245mm thick single trimming blade (Feather Safety Razor Co., LTD. Medical Division) with a 3-6 mm spacing that reflected the thickness of each axial imaging slice. The slots can either be sagittal or coronal in orientation,

depending on the patients' prostate models fusing direction. The mould was then split into two parts (left and right) utilising feature split for prostate specimen embedding. The two distinct sections that had been modified as SLDPRT files were re-saved as STL files.

Around 200 moulds were created with the assistance of the two 3D printers over the next three years, and Figure 3-7 shows some of the 3D printed models.

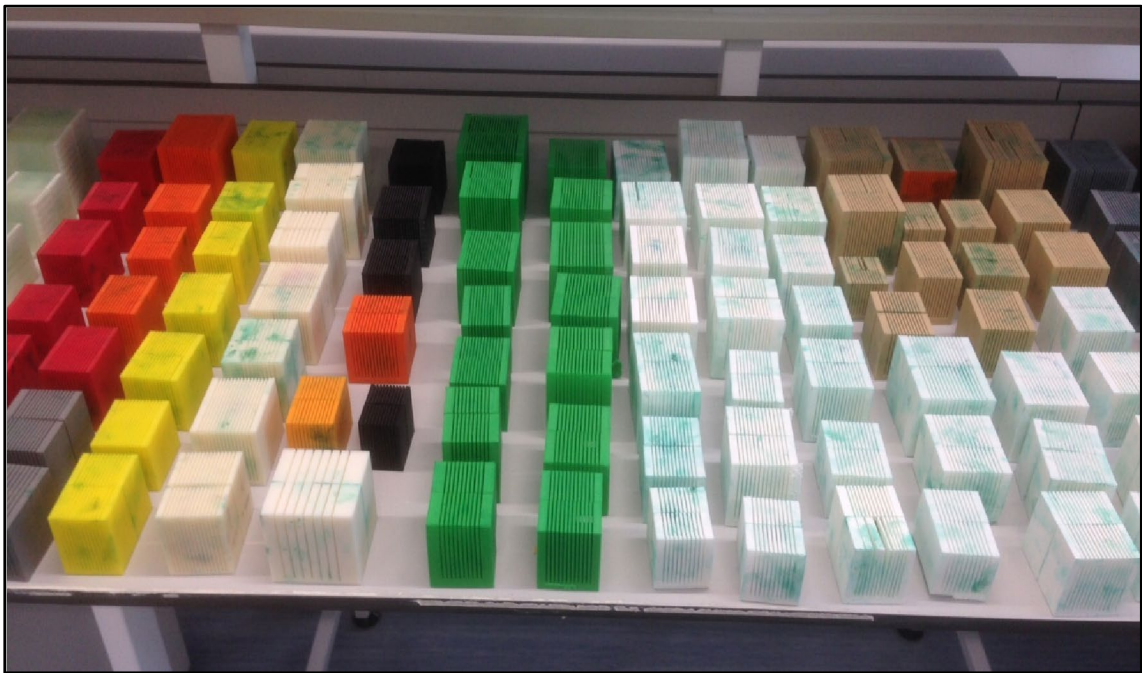


Figure 3-7 a view of all printed moulds

The average mould required 120 minutes to design, 4 to 7 hours to print and an expense for materials of less than \$7 (121).

The collection of the prostate specimen was made from patients in the operating room and sent to the pathology lab, where it was formalin fixed at room temperature for about 48-72 hours. The specimen was painted green on the left side and red on the right side before being cut in the mould, and the seminal vesicles were removed according to the mould from the apex to the prostate base. The proper orientation had to be confirmed while inserting the prostate specimen into the mould hence the

sliced histopathological specimens that would be showing their corresponding MRI scanning slices. Prostate is cut by using a single blade during sectioning, and by applying it gently and very slowly to prevent any specimen friction or movement (Figure 3-8). All histopathological sections were taken after slicing and kept separately in tissue blocks for future analysis.

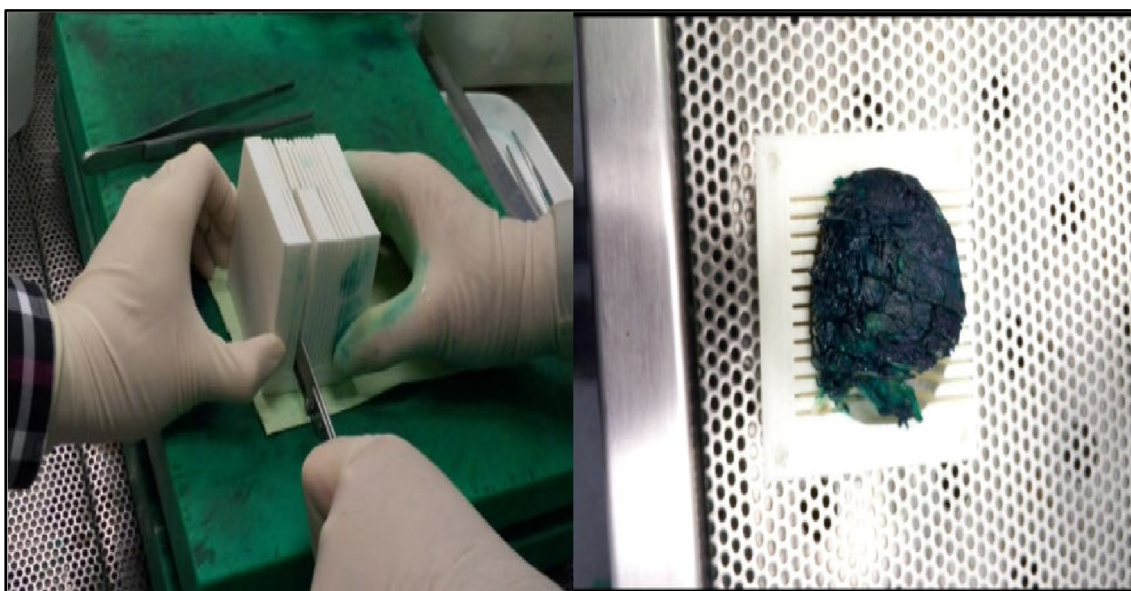


Figure 3-8 Cutting procedures and specimens after cutting

3.1.5 Statistical analysis

Baseline characteristics of patients and pathological outcomes were compared using a chi-square test for categorical data (PIRADS score) and a Student t-test or ANOVA for continuous data (age, weight, PSA level, PSAD, number of positive cores and maximum percentage of cancer per core).

Univariate logistic regression was applied to investigate the association of clinical variables with the upgrading of biopsy GS. Variables with $P < 0.05$ in the univariate analysis were further assessed using a multivariate logistic regression analysis to identify factors predictive of GS upgrading.

The parameter in the logistic regression model is measured. Specifically, In a logistic regression model, the probability of a binary response in an event is measured using a linear combination of independent variables. The dependent variable's binary value (0 and 1) represents two outcomes, such as death/survival, pass/fail, ill/healthy.

In order to estimate the area under the curve (AUC) for predicting GS upgrading to determine the diagnostic performance of clinical variables with or without PIRADS score, a receiver operating characteristic (ROC) curve analysis was conducted.

AUC reflects the discriminable ability between negative and positive probabilities based on independent variables. The accuracy of diagnosis could be improved by selecting single or combinable variables with optimal AUC as a test of diagnostic performance.

In addition, logistic regression model coefficients were used to perform a nomogram predicating the probability of GS upgrading. Non-informative or non-significant variables in univariate logistic regression for GS upgrading were removed.

The value of concordance indexes (c-index) were calculated and compared. The c-index should be between 0.5 and 1, with 0.5 indicating that the model is unable to predict the event. In contrast, when the C index is equal to 1, predicted results are completely consistent with actual results. Normally, a C index of 0.70 to 0.90 is considered to be of high accuracy. The bias-corrected calibrated values were generated from internal validation based on 200 bootstrap resamples, which ensures the excellent performance of a nomogram while it has been used in a new cohort.

A decision-analysis curve was constructed assessing impact of nomogram using different thresholds probabilities of upgrading (none of the GS upgrade to all GS upgrade).

Analyses were performed using SPSS 22 (IBM Corporation, New York, US) and R software. The alpha level was set at 0.05 to determine two-tailed significance.

3.2 Result

3.2.1 Upgrading cohort characteristics

In total, 322 men were included in our study. Table 3-3 shows the concordance between the biopsy and prostatectomy GS sums. Of these, (102/322; 31.6%) had GS upgrading from biopsy to LRP.

Almost half of this upgrading was from biopsy GS ≤ 6 disease (56/102; 55%). Majority of the low-risk patients GS ≤ 6 on biopsy were upgraded (56/62; 90%). More than half of whole cohort (175/322; 54%) had a GS 7 on prostate biopsy and (30/175; 17%) men had GS upgrading.

Finally, eighty-five of the cohort (85/322; 26%) had a GS ≥ 8 on prostate biopsy and (16/85; 18.8%) men had GS upgrading from GS 8 on prostate biopsy to GS > 8 at LRP.

Biopsy Gleason sum	Radical prostatectomy Gleason sum				
	6	7	8	9-10	Total
1-5	0	3	0	1	4
6	6	46	2	4	58
7	0	145	14	16	175
8	0	19	9	16	44
9-10	0	2	3	36	41
Total	6	215	28	73	322

Table 3.3 Comparison between biopsy and RP GS sum.

The correlation between PIRADS score and pathologic GS at LRP is demonstrated in Figure 3-9. Of the 322 patients, the distribution of PIRADS score was as follows: score 1 and 2 in 17 (17/322; 5%) patients, score 3 in 21 (21/322; 7%) patients, score 4 in 78 (78/322; 24%) patients, and score 5 in 206 (206/322; 64%) patients.

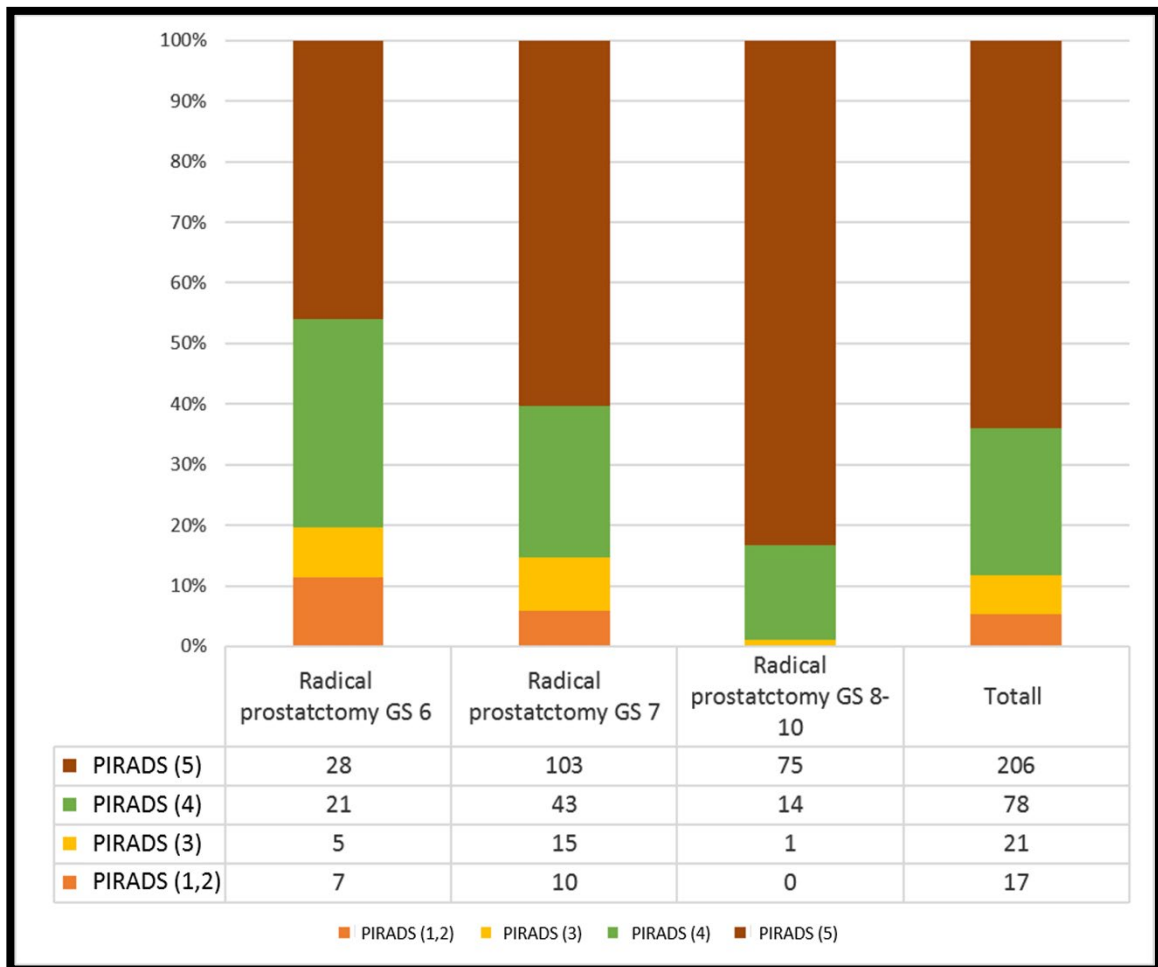


Figure 3-9 RP GS stratified according to PIRADS score.

3.2.2 Predicting of GS upgrading

Table 3-4 shows the outcomes of the logistic regression analysis and predictive variables of GS upgrading. On univariate analyses, increased preoperative prostate specific antigen levels, number of positive cores, maximum percentage of cancer per core and PIRADS ≥ 4 were all significantly associated with GS upgrading ($p < 0.05$). Age, weight of prostate and PSA density did not show any significance ($p > 0.05$) which were excluded from further analyses. In the multivariate analyses, PIRADS ≥ 4 and higher PSA level were both statistically significant and independently predictive of GS upgrading ($p = 0.001$ and 0.003 , respectively). In Figure 3-10, PIRADS v2.0 score with PSA value show a higher accuracy than PSA alone for predicting GS upgrading (AUC=0.90 and 0.64, respectively, $p < 0.001$).

	Univariate			Multivariate	
	OR (95% CI)	P-value		OR (95% CI)	P-value
Age	1.005 (0.962-1.041)	0.799			
Pathology weight	1.002(0.931-1.009)	0.540			
Number of positive cores	0.86(0.87-0.96)	0.005		0.970(0.98-1.01)	0.69
PSA level (ng/ml)	1.040 (1.009-1.073)	0.001		1.09(1.030-1.160)	0.003
PSA Density (ng/ml²)	1.15 (0.44-3.04)	0.76			
Maximum percentage of cancer per core	0.988 (0.980-0.96)	0.002		0.970 (0.84-1.12)	0.62
PIRADS ≤3 >3	1 (reference) 0.017(0.08-0.04)	- 0.001		1 (reference) 0.014 (0.06-0.034)	- 0.001

Table 3.4 Univariate and multivariate logistic regression analysis.

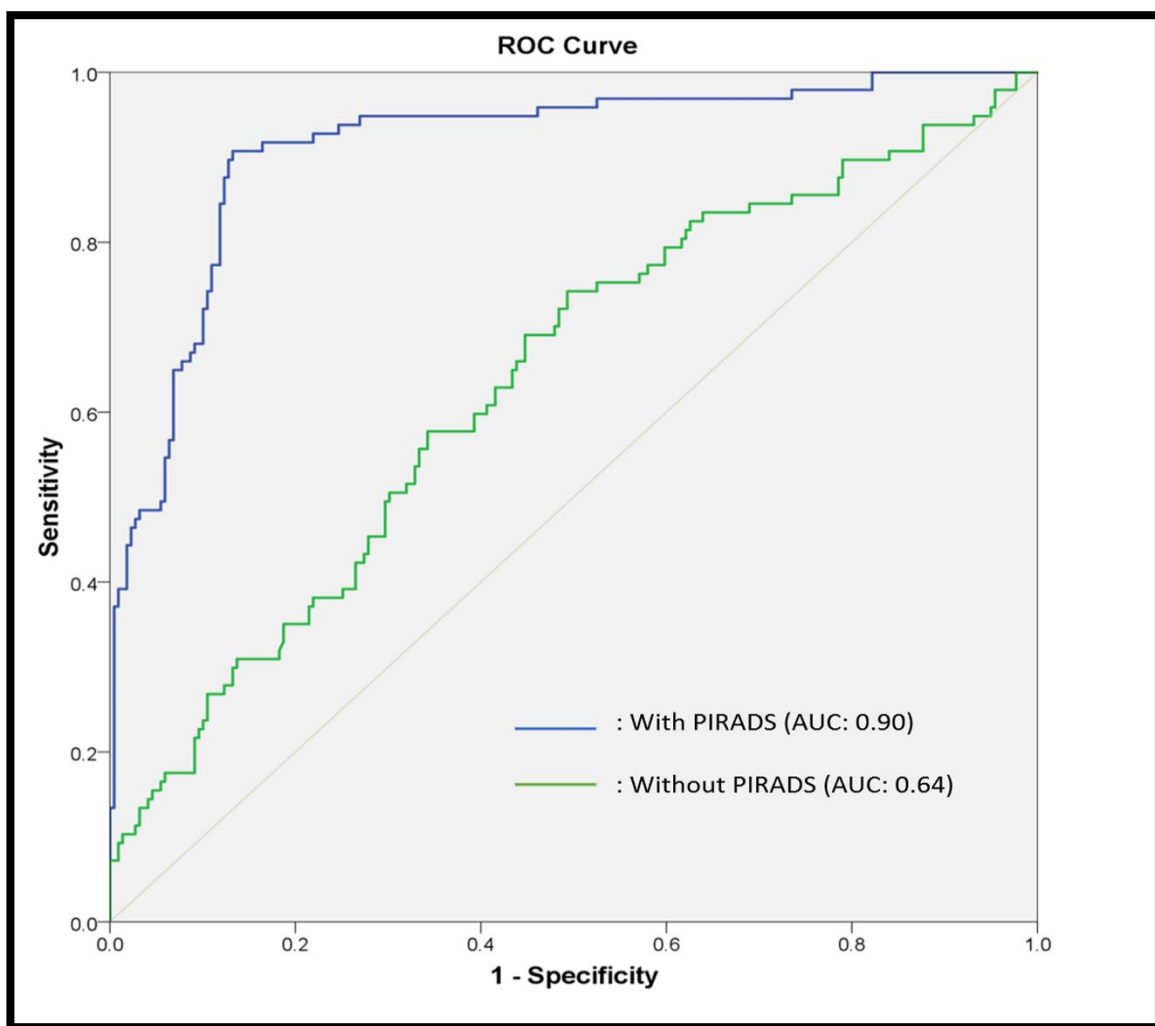


Figure 3-10 ROC curve of the clinical variables with and without PIRADS score.

3.2.3 Impact of PIRADS score on prediction of GS upgrading in relation to other factors

Figures 3-11 **a1** and **a2** show the nomograms constructed for upgradation of biopsy GS with and without PIRADS v2.0 score data. In the nomograms, a longer scale indicated higher percentage of impact and larger points were suggesting probability of upgrading. PIRADS score had the greatest impact followed by PSA level. The maximal point of each parameter was set at 10. The summation of the points of individual parameter could be found in the total points, and then the corresponding probability of GS upgrading was obtained. In the model including the PIRADS, the PIRADS occupied the longest bar in the nomogram, indicating PIRADS score had the greatest impact on the probability of GS upgrading, followed by PSA level, number of positive cores, maximum percentage of cancer per core. By contrast, the model without the PIRADS, PSA took up the largest points for the probability of GS upgrading. The second significant parameter was number of positive cores, with increase by about 3.5 point compared with the model including PIRADS.

C-index of the established nomogram which had PIRADS v2.0 score variable to predict the GS upgrading in the cohort was significantly higher than that of the nomogram without PIRADS score (0.90 [95% CI 0.87-0.89] vs. 0.64 [95% CI, 0.57-0.70], $p=0.001$). This result demonstrated that the model with PIRADS score had better performance to evaluate the predictive ability of GS upgrading.

The nomograms were then validated using 200 bootstrap samples, internal calibration curves are shown in Figures 3-11 **b1** and **b2**. The calibration plot for internal validation more intuitively reflected the predictive ability of two models. The internal calibration plot (b1) showed excellent agreement between the predictive

probabilities and actual probabilities. Compared with the model without PIRADS score (b2), left model covered additional fitting probability ranges involving below 20% and higher than 80%.

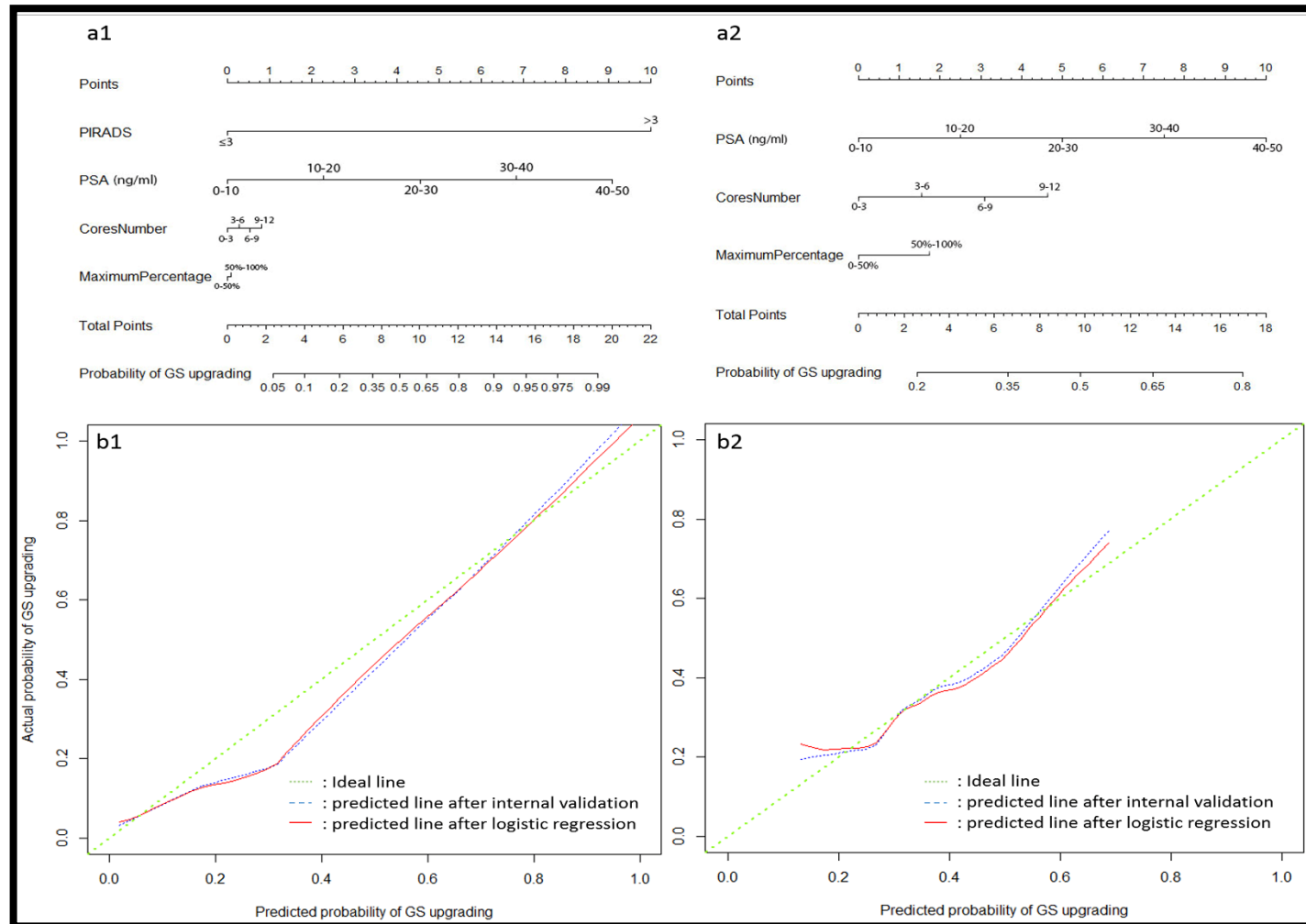


Figure 3-11 The nomograms of GS upgrading prediction with (a1) and without PIRADS score (a2). Calibration plots of observed and predicted probability of GS upgrading with (b1) and without PIRADS score (b2).

3.2.4 Decision curve analysis

The decision analysis curve is shown in Figure 3-12. The net benefit for the model using PIRADS score was significantly higher at all thresholds compared with the model without PIRADS score. As seen in Figure 3-12, the decision curve line (depicted by a red line) of the model without the PIRADS scores remained close to the line with threshold probabilities ranging from 0 to 0.25. In contrast, a higher positive net benefit was obtained in the range of threshold probabilities ranging from 0.05 to 1.0 in the model with PIRADS scores.

In the DCA curve, the line of the model without PIRADS score showed almost the same benefit to the line of 'intervention for all patient' from probability between 0%-20%, then dropped to about a low net benefit of around 0.01 at 40% probability of GS upgrading. However, the net benefit for the model with PIRADS was higher than 0.2 at 40% probability of GS upgrading. Even through at a high probability of > 80%, it still had net benefit.

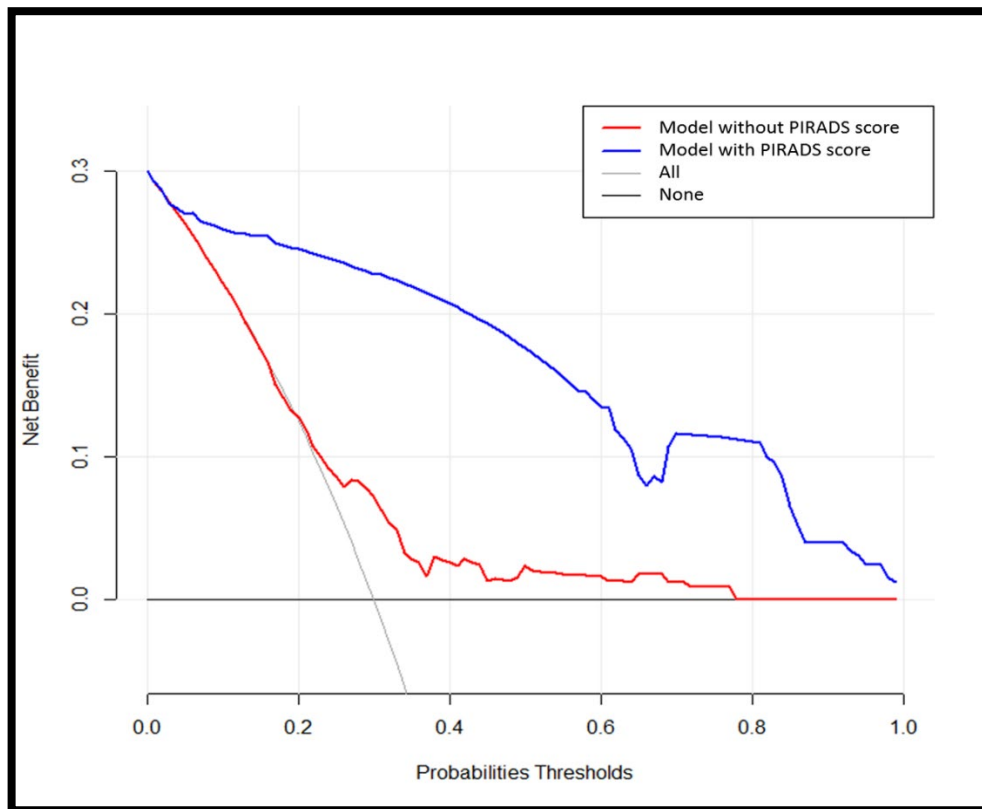


Figure 3-12 Decision analysis demonstrated a high net benefit of PIRADS score model across a wide range of threshold probabilities. Prediction model without PIRADS score (red line); prediction model with PIRADS score (blue line).

3.3 Discussion

This is the first study to bring together information of PIRADS scores in pre-biopsy MRI and an improved image oriented histopathological grossing of RP specimen by making the use of the mould, bridging the gap in the existing literature. As the T2 image is the best sequence for the anatomic information on MRI (123) we demonstrated that the mould can be used to accurately register histopathologic and MRI findings. Anatomical registration of preoperative MRI and prostate whole-mount obtained with 3D-printed, patient-specific, MRI-derived moulds versus conventional whole-mount sectioning was compared by Daniel et al (124). He concluded that anatomical registration of MRI and ex vivo prostate whole-mounts using MRI-derived moulds is significantly better than the conventional method (0.95/0.85 respectively) ($P < 0.0001$). Our results indicate a significant advantage (C-index 0.90 vs 0.64) of using the prediction model including PIRADS scores added to conventional clinic-pathological characteristics (PSA level, percentage of cancer on core-biopsies, gland size etc.) of men with PCa confirmed by systematic transrectal random biopsies relative to a model without PIRADS scores. Age has been considered an important predictor of GS upgrading in a previous study (125). However, in our study age was not associated with the upgrading of GS due to the limited sample size similar to the findings of Song (9). Prostate biopsy GS upgrading remains a challenge for physicians managing localised PCa, as better knowledge of contributing factors and how to narrow the gap is lacking (2). To inform any consensus, we need an improved understanding of the role imaging can play, in particular pre-biopsy MRI, in predicting GS change and adverse downstream oncological outcomes. Although, recent improvements have been made in refining biopsy strategies and in reducing

sampling errors, a significant and continued effort is still needed to identify men at risk of GS upgrading.

Wang et al (126) reported a nomogram with C-index of 0.795 using preoperative factors without imaging data in a non-screened population from China. This is similar to our study as the healthcare system for the cohort reported here did not have men screened for PCa. Table 3-5 shows the predictive ability of various reported nomograms in upgrading of biopsy GS of PCa in screened populations (1,126–129). The upgrading rate of biopsy GS seen in our cohort is similar to a larger cohort of 2982 patients reported previously (127). A higher percentage of men with a GS of 6 were upgraded in the present study. It is interesting that despite the higher number of cores obtained in the present study (12 in number) in comparison with 6-10 biopsy cores obtained in study by Chun FK et al (127); upgrading rates remain comparable. A number of previous studies have carried out multivariate analyses of factors responsible for upgrading of biopsy GS including construction of nomograms (Table 3-5).

Author	year	Number of patients	Performance (C-index)	Significant parameters on multivariate analysis
Chun, FK	2006	2982	0.804	PSA level, clinical stage and primary and secondary GS
Kulkarni, GS	2007	175	0.71	PSA level and the level of pathologist expertise
Budäus, L	2010	414	0.708	PSA level, clinical stage, prostate volume and percent of positive cores
Wang, JY	2014	220	0.789	PSA level, clinical stage, and primary and secondary GS
Biming, He	2016	411	0.753	Primary and secondary GS and obesity
This study	2019	322	0.90	PSA level and PIRADS score on MP-MRI

Table 3.5 Comparison between previous and current studies.

In predictive oncology, nomograms have huge potential to help clinicians determine the risk of disease progression and identify those who would experience a greater benefit from multimodality therapy. This approach may result in avoiding unnecessary treatment and improve quality of life by reducing side effects of therapy through better and more precise approach. However, a careful approach is needed to construct nomogram based on specific question, the study population, the method of construction, and its ability to apply to a particular clinical situation. We have followed guidelines described in previous publications (118,130,131) in constructing nomogram in this study including selection of variables and statistical methods. The nomogram in the present study has been internally validated (Cross-validation and bootstrapping).

External validation of nomogram was not carried out in the present study as this would require further prospective multi-centre recruitment of a cohort. Since D'Amico pioneered this approach (132), none of the reported predictive nomograms have included imaging features of the disease. Furthermore, the advantage of our nomogram is a higher overall accuracy (discriminant properties) and closer agreement between predicted and observed values (superior calibration). Estimating clinical utility of nomograms in prognosticating an outcome of intervention remains core value of translational research in precision medicine. Vickers and Elkin (119) have introduced decision analysis curves estimating probabilities of benefits and harms that a diagnostic test or intervention can trigger at various thresholds. Addition of PIRADS score to nomogram achieved a higher net benefit of decisions making in comparison to leaving out PIRADS score as shown in the decision analysis curve constructed in the present study. The thresholds ranged from no upgrading of disease to all men having upgrading of disease following LRP.

Predicting final histopathological GS of PCa remains a highly desirable information for physicians counselling men with localised PCa for various modalities of treatment and long-term disease recurrence. At present, various nomograms are used mainly taking into consideration clinical factors such as age, pre-operative PSA level and number of biopsy cores involved with the cancer. Notwithstanding this, there is still a large histopathological discrepancy between biopsy and final RP GS. The present study reports a nomogram based on pre-biopsy MP-MRI grade (PIRADS score) of cancer along with other known clinical parameters. The nomogram clearly showed an improved prediction of final GS and the findings have a large implication for clinicians and researchers in this area. The majority of low-risk patients were upgraded to a higher risk in this study. Low-risk patients with an elevated PSA or those for whom expert pathological review is not possible may benefit from a repeat biopsy, as evidence suggests that additional biopsy cores improve GS accuracy for patients opting for AS (128). We envisage that this and further research should take us close to precise prediction of final GS of histopathology in PCa and thereby an improved and informed decision making by stakeholders including patients in the management of localised PCa. This will have huge benefits for improved GS prediction for men opting for AS and focal therapy besides those opting for radical prostatectomy.

There are limitations to this study. The study was conducted retrospectively at a single centre with dedicated uro-radiologist and pathologist, and the rate of upgrading may be different in small centres. Moreover, overall accuracy of our model, although higher than previous was not perfect (90%). Additionally, performance of our model needs further validation in an external data set. PIRADS v2.0 is a step forward in simplifying the initial standardisation efforts made in PIRADS

v1, but it will undoubtedly require more changes as research progresses, experience grows, technology advances and this may impact the external validation of the present nomogram. Although, there may be inter-reader variability between the readers of MRI however, Greer et al (133) in a multi-reader study (showed high sensitivity and agreement (>90%) between prostate specialists for PIRADS v2.0. Finally, the accuracy of our model could still be improved by integrating additional predictor variables, such as the novel genomic and other biomarkers (134,135). The growing field of artificial intelligence and machine learning using radiomics approach may improve our ability to define tumour characteristics and classification. This undoubtedly may impact results of the study in the future. Finally, with emerging evidence supporting MRI facilitated biopsy targeting of suspicious areas using ultrasound, the rate of upgrading and the future implications for the practice may change (136). There is an emerging evidence that TB may improve our ability to narrow the upgrading gap between the biopsies and RP histology(137,138); however no predictive nomogram information was available from both the studies. Our ongoing work through randomised intervention in MR/US fusion should be able to provide more information (139)

Chapter 4 Predicting the performance of concurrent SB during TB sampling of MP-MRI

4.1 Materials and methods

4.1.1 Study population and power calculation

The study had ethical approval (14/ES/1070) and all participants provided informed consent for their imaging data to be used. The study also had Caldicott institutional approval (Caldicott/IGTCAL6358) to link data with electronic systems wherever needed for follow-up outcomes. The study period was between April 2015 and March 2020.

The inclusion criteria were age between 40 to 76; abnormal DRE; $PSA \leq 20$ ng/ml; and MRI $< T3$ disease. Exclusion criteria were repeat biopsies; prior radiotherapy to prostate; and diagnosis of acute prostatitis with 12 months or history of PCa. All participants had pre-biopsy MP-MRI and only MRI positive ($PIRADS \geq 3$) were recruited into the study ($n=198$). Of these patients, 78/198 (39%) underwent radical prostatectomy. Patients then underwent prostate biopsy by MR/US fusion technique (Hitachi HI-RVS; Europe Holding, Steinhausen, Switzerland) by operator 1. This was followed by a standard 12-core TRUS biopsy by a second operator (blinded to the MRI results).

In considering performing a McNemar matched test, a minimum sample size of 110 men undergoing both SB and TB prostate biopsy methods would be required to yield 90% power with a significance level of 0.05 ($\alpha = .05$) which would also allow 20% of dropout rate. csPCa detection rate via Sb and TB were found in previous study (140). Therefore, we recruited more patients than the minimum required number from the sample estimation to ensure achievement of study power and significance level.

4.1.2 MRI techniques

Two weeks before TRUS-guided biopsies, all patients had MP-MRI scans on a 3T scanner (TIM Trio, Siemens, Erlangen, Germany). The MP-MRI procedure was adapted from version 2 of the European Society of Uro-radiology Guidelines for the detection of prostate cancer, which were published in 2012. Phased-array coil was placed around the pelvis's patient while was lying on his back. A gadolinium contrast agent (Dotarem, Guerbet, France) was inserted to acquire DCE sequence with dose equal to 2 ml/kg of Dotarem. The scan took approximately an hour to include the following anatomical and functional sequences:

- High-resolution T2 spin echo (SE) pulse sequence, acquired in the sagittal, axial and coronal oblique planes.
- T1 SE pulse sequence of the pelvis acquired in the axial oblique plane.
- T1 SE axial sequence of abdomen and pelvis.
- T2 3D SPACE sequence for fusion biopsy planning.
- T2W fast spin echo sequence of abdomen and pelvis in coronal plane.
- DWI.
- DCE.

The MP-MRI images were analysed and scored by experienced uro-radiologists using PIRADS v2.0; and the radiologists were blinded to all patients' pathology results.

4.1.3 Biopsy procedures and histopathological analysis

Data from an MRI scan was obtained two weeks prior to the biopsy. MRI images were loaded into the HI-RVS software (Hitachi-Real-time Virtual Sonography) via a USB device. Prior to the MRI/US fusion biopsy, the prostate MRI was used as a map to target prostate lesions, register three points (the apex, base, and posterior of the prostate) and create a 3D-reconstruction of the prostate (Figure 4-1).

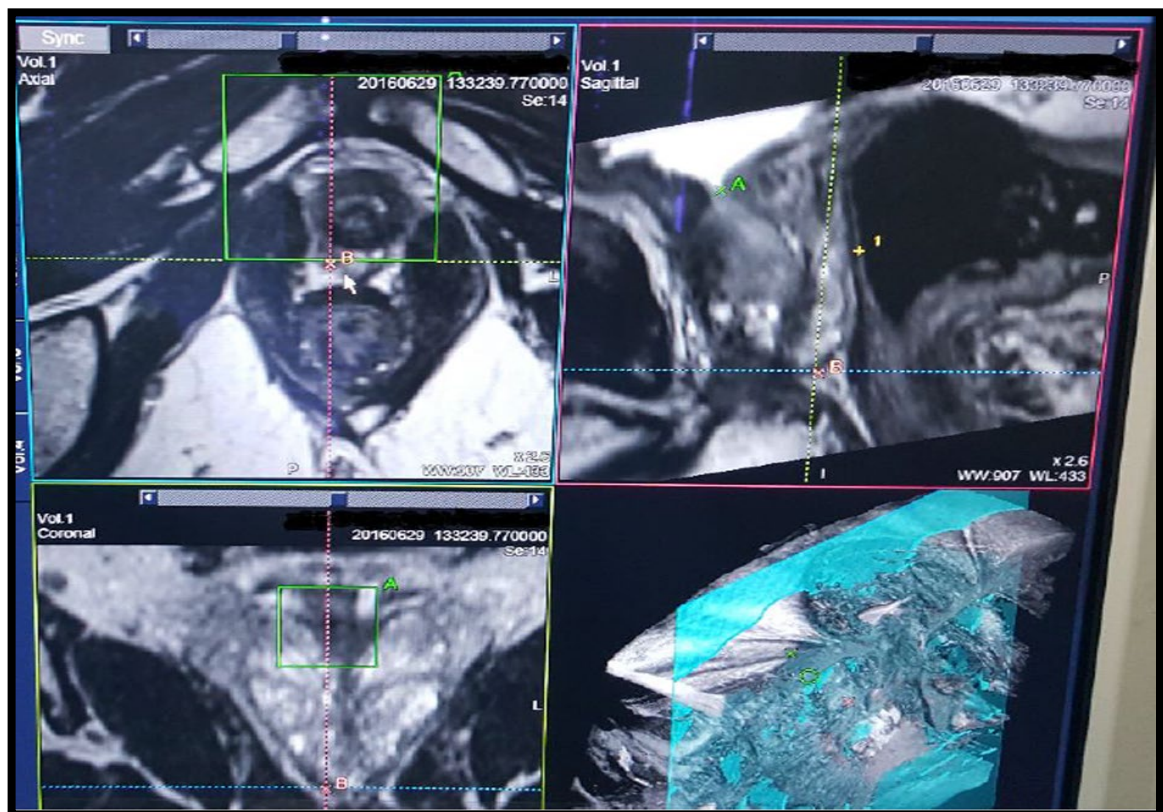


Figure 4-1 An MRI of the prostate taken before the biopsy is used as a map.

The patient was then placed on the table in left lateral decubitus so that an NHS-accredited, well-trained radiologist and/or urologist could perform an MRI/US fusion biopsy. While under local anaesthetic (Lidocaine, Hydrochloride injection 1%) diluted with normal saline, the operator inserted the endo-cavity (trans-rectal) biplane transducers (EUP-CC531) with a frequency between (8.0-4.0) MHz to perform the biopsy. The transducer was joined to a biopsy gun (BARD, 22mm penetration length,

18 gauge) to obtain the biopsy cores, and Electro-Magnetic (EM) sensors (EZU-RV2S) (Figure 4-2) were affixed to EM field generator stand. These were put close to the patient to transmit the signal to the sensor, allowing for more precise target location.



Figure 4-2 EMS with a trans-rectal ultrasound transducer

The fusion device's screen was divided into two screens: the left screen with a pre-biopsy MR image; and the right screen with real-time ultrasound to fuse and align the targeted lesion on the MRI. These would be aligned and fused with the US which would be used to guide the biopsy needle to the region of interest (ROI) in the MRI where the targeted lesion was located (Figure 4-3).

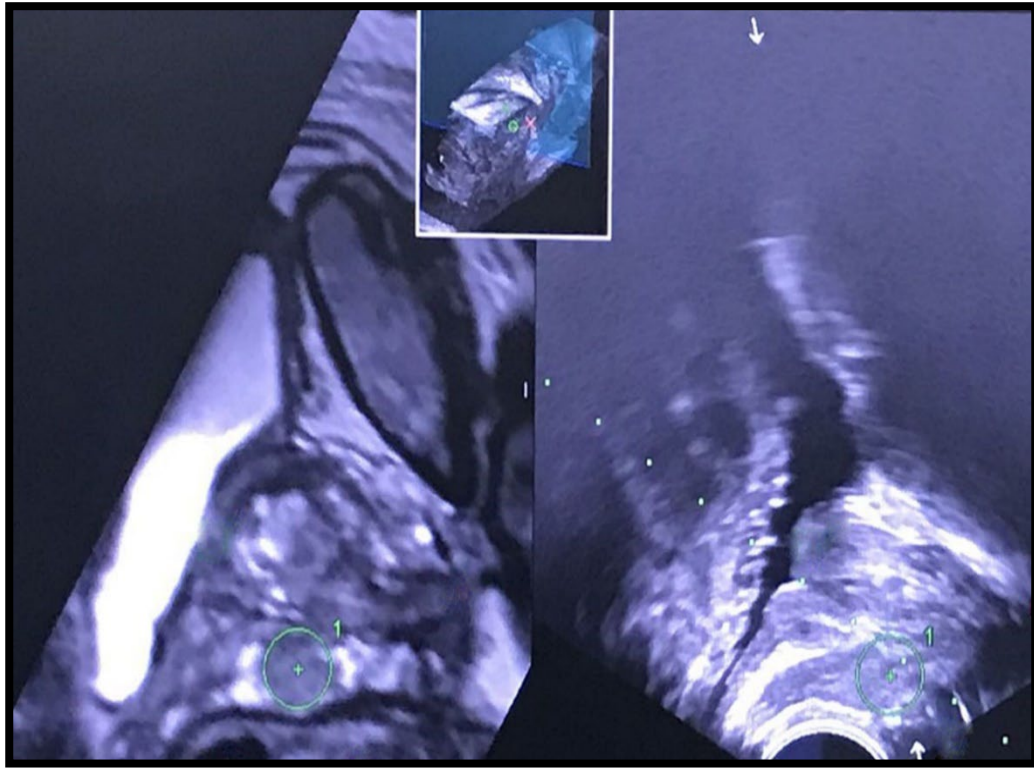


Figure 4-3 (A) The left window displays an MRI of the prostate, and (B) the right side window (US window) and the green circled region shows the index lesion where the biopsy can be taken from.

Three cores of tissue were obtained in the fusion TB approach from previously identified MP-MRI lesions using superimposed T2-weighted sequence on the real-time TRUS image.

On the exact same day as the MRI/US fusion biopsy, the systematic random 12-core biopsies (Figure 4-4) were performed by an experienced urologist or specialist nurse following targeting.

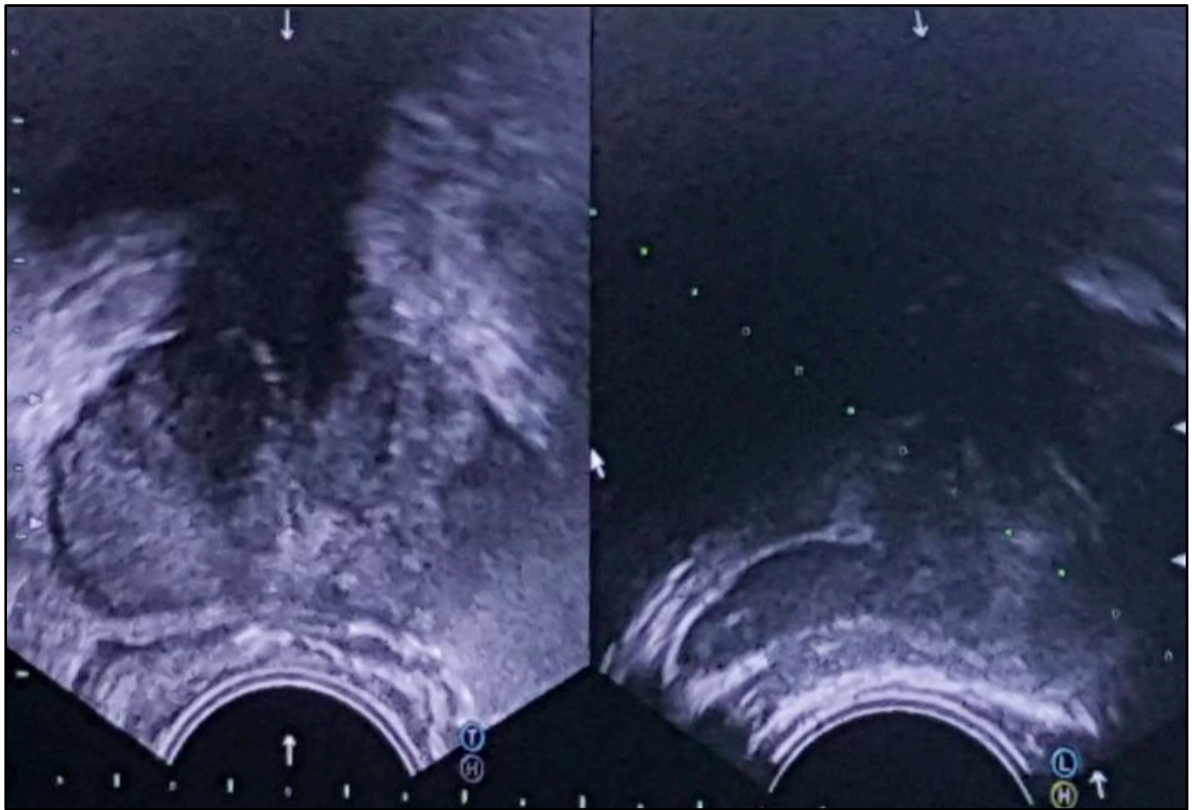


Figure 4-4 TRUS guided prostate biopsy

The SB was typically the 12 cores approach, collected in an extended sextant template of biopsies from the lateral and medial aspects of the base, mid, and apical prostate on the left and right sides

The twelve core locations are:

1. Right upper lateral
2. Right base
3. Right mid lateral
4. Right para-midline
5. Right lower lateral
6. Right apex
7. Left upper lateral
8. Left base

9. Left mid lateral
10. Left para-midline
11. Left lower lateral
12. Left apex

The pathologist identified all the cores and recorded their Gleason score (Table 4-1).

	Core Location	Score	% Involved
1	Right upper lateral	4+4	40
2	Right base	5+4	100
3	Right mid lateral	4+5	100
4	Right Para midline	4+5	100
5	Right lower lateral	4+5	100
6	Right apex	4+5	100
7	Left base	4+3	20
8	Left upper lateral	4+3	80
9	Left para midline	4+3	100
10	Left para midline	4+3	100
11	Left lower lateral	4+3	100
12	Left apex	4+3	50

Table 4.1 SB report for all the 12 cores

The biopsy results were analysed by experienced uro-pathologists (blinded to the MRI findings). The GS for each patient was obtained.

A subset of cohort (n=78) underwent RP and prostate glands were assessed by mould-based histopathological orientation as described in our previous study (121,122). A genitourinary pathologist with 10 years of experience independently reviewed step-section histologic slides of the prostatectomy specimen for each case. This pathologist was unaware of the MRI and biopsy findings. The pathologist identified all the lesions for each case and recorded their Gleason score (Figure 4-

5) summarises the study protocol in brief. This was used as a reference standard to assess the diagnostic accuracy of both SB and TB in detecting csPCa.

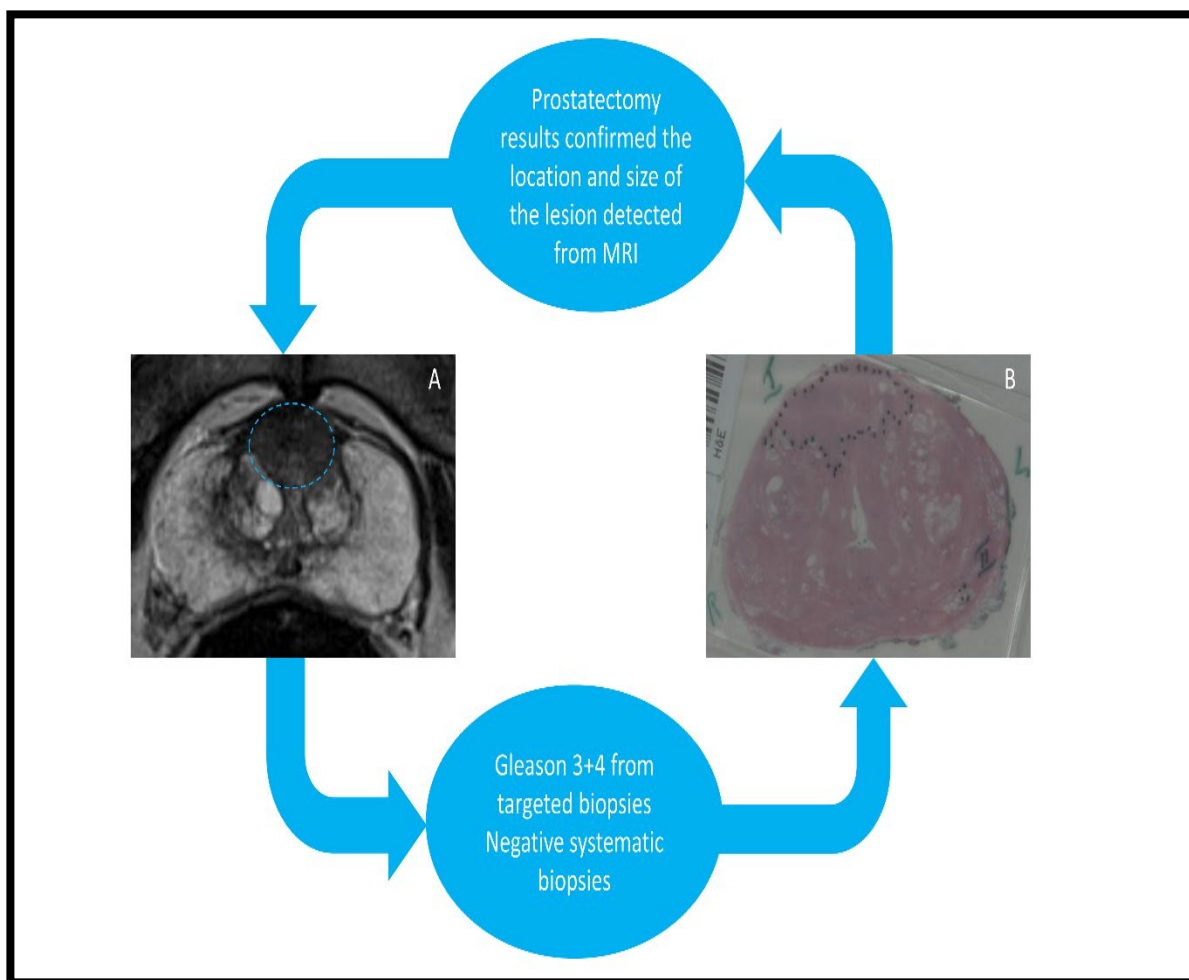


Figure 4-5 A: A 76 years old patient with a PIRADS 5 lesion detected from 3T MRI in anterior zone with a high PSA and abnormal DRE. B: Patient-specific 3D Mould based grossing of radical prostatectomy slice shows a 3+4 GS cancer located in the anterior zone

4.1.4 Outcomes

Primary outcome of the study was to compare the detection rate of clinically significant PCa using both SB and TB approaches alone and in combination. This was assessed both at biopsy and RP stages. Significant PCa was defined as the presence of PCa with GS $\geq 3+4$ (International Society for Urological Pathology [ISUP])

grade 2 or more). The secondary outcome was to assess net clinical benefit of the approaches using nomogram and decision-analysis methods.

4.1.5 Statistical analysis

Patient's age (in years), PSA, prostate volume (ml), and PSAD were collected. PSAD was calculated using PSA divided by the MRI-derived prostate volume (ellipsoid method). Number of lesions, index lesion size (mm), PIRADS category, and lesion location were measured by MP-MRI. Each lesion was counted only once. Index lesion size was the size of lesion with the highest PIRADS score. Continuous data were first tested to see if they were normally distributed by the Kolmogorov-Smirnov Test of Normality. The mean (m) and standard deviation (SD) were described if the variable followed a normal distribution. The median (M) and interquartile range (IQR) were presented if the variable was not normally distributed. Categorical variables were reported as frequencies and proportions. Cross tabulation was carried out in order to compare the proportions of significant PCa patients by SB, TB, and combined SB+TB.

The McNemar chi-square test was conducted in patients who were given both diagnostic tests. McNemar chi-square, degree of freedom (df) and P-value were calculated and presented. A two-step Logistic regression was performed to identify explanatory variables of significant PCa. First, the patient's age, PSAD, lesion size, PIRADS and number of lesions were individually put into a univariate logistic regression model where the outcome was defined as having significant PCa or not. Statistically significant variables were then put into the multivariate logistic regression model. Odds ratio (OR), 95% confidence interval (95% CI) of OR, and P values were recorded.

A nomogram was created based on the statistically significant variables in the final model. The discriminative ability of the predictive model was tested by ROC curve and the concordance statistic (c-statistic) was presented. The predicted probabilities of significant PCa were plotted against observed probabilities to test the calibration of the model. Decision curve analysis was applied to determine the benefit of the nomogram.

In the subgroup analysis, prostatectomy was performed in a group of 78 patients (112 lesions). The detection rate of true significant PCa lesions via SB, TB and combined SB+TB was compared using the McNemar chi-square test. Statistical analyses were conducted by SPSS V23.0 and R V4.0.3. The Bonferroni adjustment, which adjusted P value by times of tests, was used accounting for multiple testing. The alpha level (adjusted P- value) was set at 0.05/times of tests to determine two-tailed significance for the McNemar chi-square test.

4.2 Result

4.2.1 Patient's characteristics

The participating patients' demographic data are shown in Table 4-2. A total of 198 patients who underwent both SB and fusion TB in the same setting were recruited into the study.

Several clinical variables included baseline information (age, PSA level (ng/ml), PSAD (ng/ml²), and prostate volume (ml)) and MP-MRI features (number of lesions seen on MRI, index lesion size (mm), PIRADS score 3,4,5 and lesion location transition zone (TZ), PZ, and both zones (TZ+PZ)).

Variables		Overall (n=198)
Basic information	Age, median (IQR), in years	67 (71-61)
	Prostate specific antigen (PSA), median (IQR), ng/ml	8.2 (10.6-6.4)
	Prostate volume, median (IQR), ml	47 (63-33)
	PSA Density, median (IQR), ng/ml ²	0.18(0.27-0.11)
MP-MRI	Number of lesions, n (%)	
	1	102 (51.5%)
	2	75 (38%)
	3	14 (7%)
	4	6 (3%)
	5	1 (0.5%)
	Index lesion size, median (IQR), mm	16 (25-13)
	Prostate Imaging Reporting and Data System (PIRADS score), n (%)	
	PIRADS 3	22 (11%)
	PIRADS 4	55 (28%)
	PIRADS 5	121(61%)
	Lesion location, n (%)	
	Peripheral zone (PZ)	79 (40%)
	Transition zone (TZ)	44 (22%)
	Both zones (TZ-PZ)	75 (38%)
Targeted (TB)/Systematic random (SB) biopsy	Detection of prostate cancer (PCa) in TB, n (%)	129 (65%)
	Detection of prostate cancer (PCa) in SB, n (%)	127 (64%)

Table 4.2 Descriptive characteristics of 198 patients who underwent MP-MRI of the prostate and subsequent systematic random and TB at our centre.

4.2.2 Comparison of the detection rate of significant prostate cancer between SB, TB and combined approaches

The detection rate of csPCa using random biopsy was 51.0% (101/198) and using TB was 56.1% (111/198). This was not statistically significant (the McNemar chi-square test was $\chi^2=2.273$, $df=1$, $P=0.132$, $OR=0.63$ (95% CI, 0.34 to 1.16). There were 17 patients (17/198; 8.5%) where the TB approach alone missed clinically significant cancers (8 from the same site and 9 from normal looking prostate on MRI (Figure 4-6).

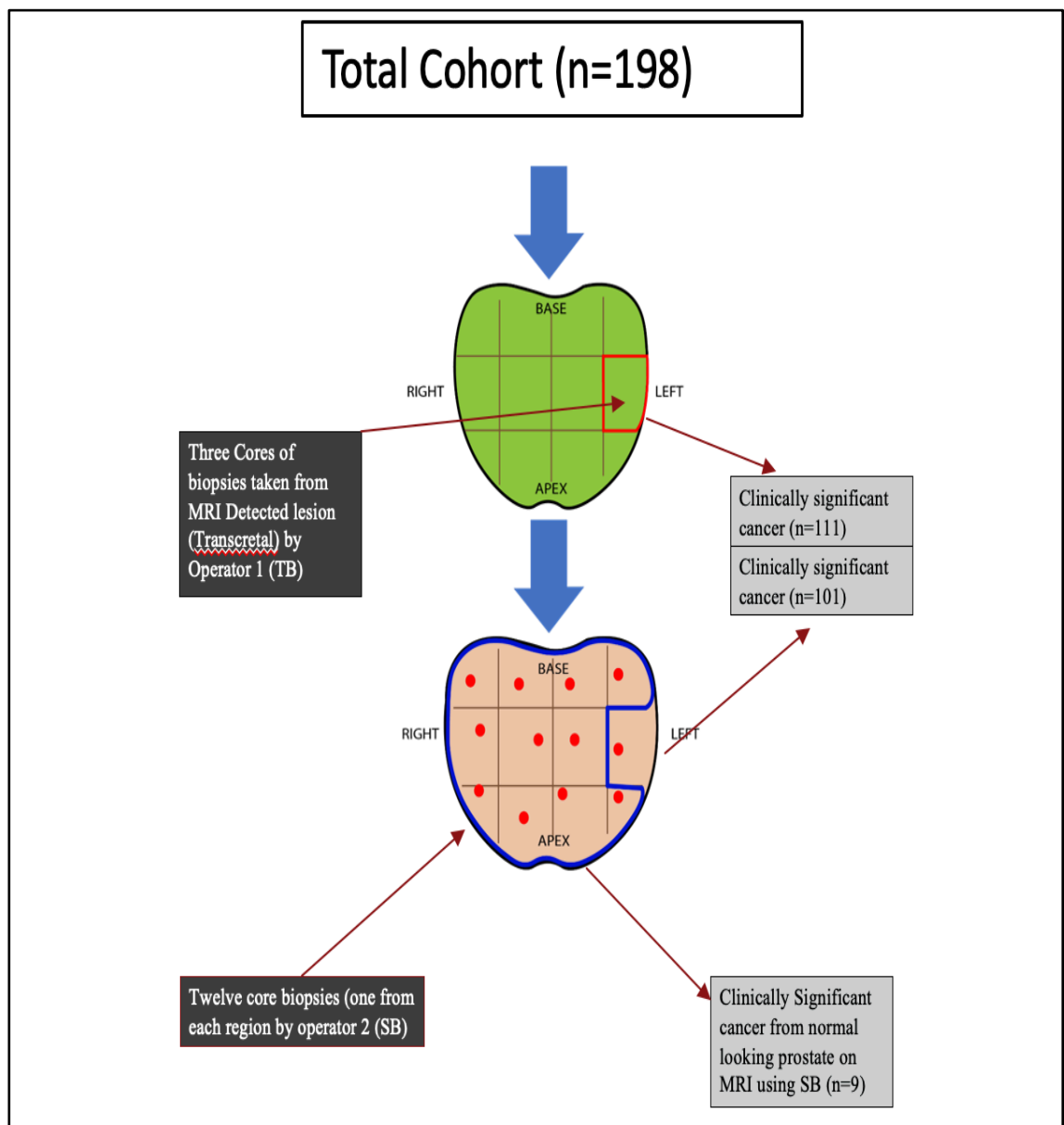


Figure 4-6 The detection rate of significant prostate cancer between SB, TB based on patients' level.

There were 84 patients (84/198;42.4%) where the positive cores on systematic sampling and TB detected csPCa (72 from the same sector of index lesion and 12 from different sectors away from index lesion). When the TB is negative (69/198; 34.8%), the SB detected clinically insignificant cancer in 12 patients (12/69; 17.3%) and detected csPCa in 8 patients (8/69;11.6%). Twenty-seven (27/198; 13.6%) men were upgraded to csPCa based on TB while 17 patients (17/198; 8.5%) were upgraded based on SB ($\chi^2=2.27$, $P=0.13$).

The detection rate of significant PCa was higher using combined biopsy (64.6%; 128/198) in comparison to TB (56.1%; 111/198). The McNemar chi-square test result with Yates correction was statistically significant ($\chi^2=15.06$, $df= 1$, $P<0.001$). There was an 8.5% increase in significant PCa detection rate at the patient level using combined biopsy methods compared to using TB alone. The results are shown in Table 4-3.

		Fusion Targeted biopsy (TB)				Total
		No Cancer	Low-Risk Cancer	Intermediate-Risk Cancer	High-Risk Cancer	
Systematic random Biopsy (SB)			Gleason 3+3	Gleason 3+4	Gleason > 3+4	
No Cancer		49	3	9	10	71
Low-Risk Cancer	Gleason 3+3	12	6	6	2	26
Intermediate-Risk Cancer	Gleason 3+4	4	8	15	6	33
High-Risk Cancer	Gleason > 3+4	4	1	18	45	68
Total		69	18	48	63	198

Table 4.3 Biopsy findings per patient of SB compared with TB for the total cohort of 198 men. Orange shading indicates men in whom SB upgraded the prostate cancer risk category in relation to TB. Blue shading indicates men in whom TB upgraded prostate cancer risk.

We further validated the findings using a subset of cohort where the histopathology of RP was used as a reference standard. There were 170 clinically significant cancers (170/190; 89.4%) seen on RP histopathology using mould-based approach and counting each focus of cancer. 112 were targeted using the MRI/US image fusion method. The TB approach detected 70 of these (70/112; 62.5%), whereas the SB approach detected 54 (54/112; 48.2%). The difference was statistically significant (the McNemar chi-square test result was $\chi^2=6.618$, $df=1$, $P=0.010$, $OR=0.36$ (95% CI, 0.17 to 0.77). The combined approach to 112 lesions detected more cancers than SB or TB alone (79/112; 70.5%). Compared to SB, the combined approach detected 22.3% more cancers (70.5% vs 48.2%). The McNemar chi-square test result with Yates correction was statistically significant ($\chi^2=23.04$, $df=1$, $P<0.001$). Similarly, the combined approach detected 8% more cancers in comparison to TB (70.5% vs. 62.5%). The McNemar chi-square test result with Yates correction was statistically significant ($\chi^2=7.111$, $df=1$, $P=0.008$).

Interestingly, there were 11 cancers (11/190; 5.8 %) which were labelled as clinically insignificant and all were upgraded to clinically significant using the TB approach. In comparison, there were 24 (24/190; 12.6%) cancers labelled as clinically insignificant and 20 (20/190; 10.5%) were upgraded using the SB approach. The McNemar chi-square test result with Yates correction was $\chi^2=0.450$, $df=1$, $P=0.502$, $OR=1.50$ (95% CI, 0.61 to 3.67. This was not statistically significant (Figure 4-7).

Radical prostatectomy cohort (n=78)

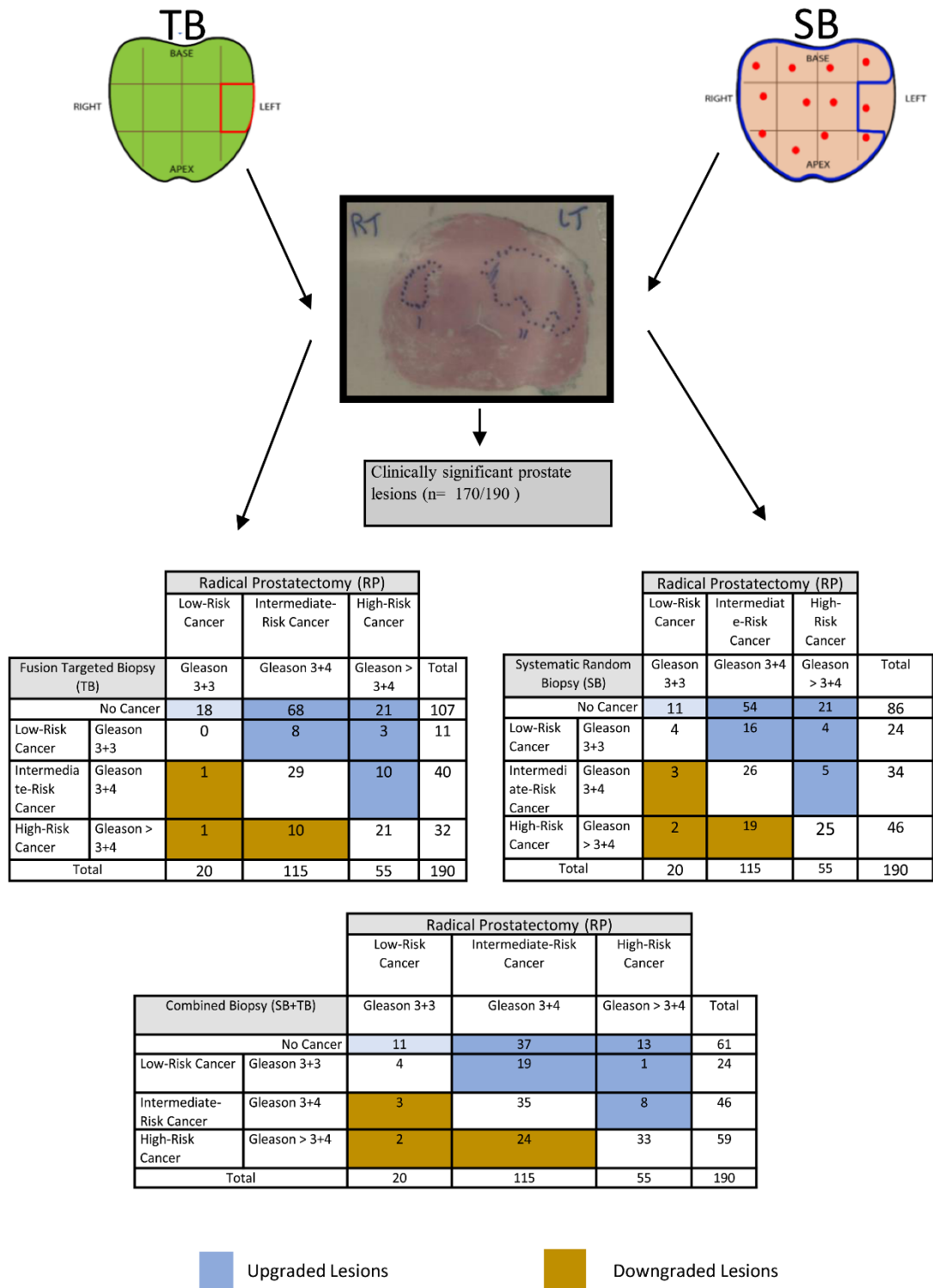


Figure 4-7 The detection rate of significant PCa via SB, TB and combined SB+TB on RP lesions.

Table 4-4 shows the difference in prostate cancer lesion upgrading between SB and TB (112)

SB	TB		Total lesions
	Upgrading	Not upgrading	
Upgrading	3	12	15
No upgrading	8	89	97
Total lesions	11	101	112

Table 4.4 Upgrading lesions between SB and TB (112)

The proportion of lesion upgrading between SB and TB are 9.8% (11/112) and 13.4% (15/112), respectively. This was not statistically significant (the McNemar chi-square test result with Yates correction is $\chi^2=0.450$, df= 1, P=0.502, Odds ratio OR=1.50 (95% CI, 0.61 to 3.67). Table 4-5 shows the difference in prostate cancer lesion upgrading between combined biopsies and TB

SB+TB	TB		Total lesions
	Upgrading	Not upgrading	
Upgrading	8	7	15
Not upgrading	3	94	97
Total lesions	11	101	112

Table 4.5 Upgrading lesions between combined biopsy (SB+TB) and TB (112)

The proportion of lesion upgrade between combined biopsy and TB are 13.4% (15/112) and 9.8% (11/112), respectively. This was not statistically significant (the McNemar chi-square test result with Yates correction is $\chi^2=0.900$, df= 1, P=0.343, Odds ratio OR=2.33 (95% CI, 0.60 to 9.02. Table 4-6 shows the

difference in prostate cancer lesion upgrading between combined biopsies and SB

SB+TB	SB		Total lesions
	Upgrade	Not upgrade	
Upgrading	10	5	15
No upgrading	5	92	97
Total lesions	15	97	112

Table 4.6 Upgrading lesions between combined biopsy (SB+TB) and SB (112)

The proportion of lesion upgrading between combined biopsy and SB are both 13.4% (15/112). The result was not statistically significant (the McNemar chi-square test result with Yates correction is $\chi^2=0.100$, df= 1, P=0.752, Odds ratio OR=1.00 (95% CI, 0.29 to 3.45).

4.2.3 Univariate and multivariate logistic regression analysis and developed nomogram

In univariate logistic regression, patient's age, PSAD, lesion size and PIRADS were all significant predictors of significant PCa detected by random biopsy (Table 4-7) and therefore were put into multivariate analysis. A 6% increase in odds of significant PCa by random biopsy was associated with each one-year increase in age (OR= 1.06, 95%CI 1.01-1.12). Per 1ng/ml² increase of PSAD was associated with an almost 26-fold increase in odds of significant PCa (OR= 25.63, 95%CI 1.93-341.27). Having PIRADS-5 was another significant predictor of significant PCa using random biopsy which was associated with a 6-fold increase in odds compared to those with PIRADS-3 (OR=5.94, 95%CI 1.77-19.93).

Covariate	N	Univariate Logistic Regression				Multivariate Logistic Regression			
		OR	95%CI		P value	OR	95%CI		P value
			Lower	Upper			Lower	Upper	
Age (year)	198	1.07	1.02	1.12	0.009	1.06	1.01	1.12	0.036
PSAD	198	92.79	7.61	1130.69	<0.001	25.63	1.93	341.27	0.014
Lesion size	198	1.06	1.03	1.10	<0.001	1.02	0.98	1.06	0.399
PIRADS	198				<0.001				0.001
3	22	Ref	-		-	Ref	-		-
4	55	1.69	0.49	5.80	0.406	1.51	0.42	5.43	0.525
5	121	9.46	3.00	29.84	<0.001	5.94	1.77	19.93	0.004
Number of Lesions					0.309				
1	102	Ref	-		-				
2	75	1.11	0.61	2.02	0.730				
3 and above	21	2.16	0.81	5.80	0.125				

Table 4.7 Univariate and multivariate logistic regression analysis.

The statistically significant independent variables from the multivariate logistic regression model (age, PSAD and PIRADS) were used to develop a nomogram to predict the probability of significant PCa using SB (Figure 4-8).

The model demonstrated good discriminative ability (C-statistic= 0.779, 95%CI 0.714-0.843) (Figure 4-9).

The calibration plot demonstrated good agreement between the model predictions and actual observations for detecting csPCa via SB (Figure 4-10).

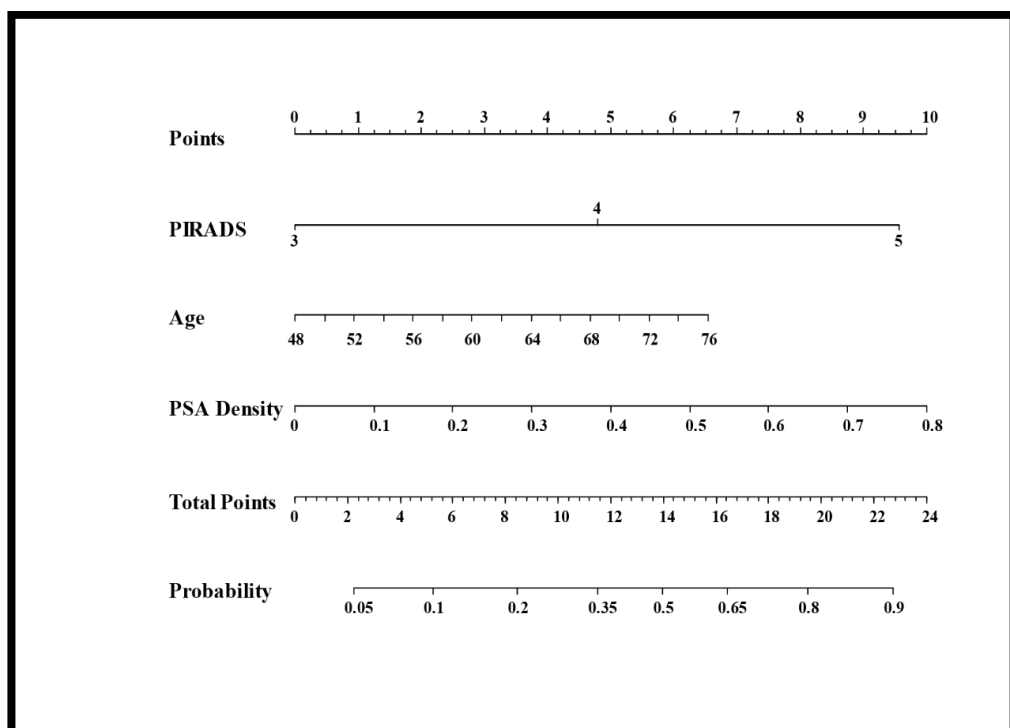


Figure 4-8 Nomogram with significant clinical variables to predict patients who will benefit from performing SB in addition to fusion TB.

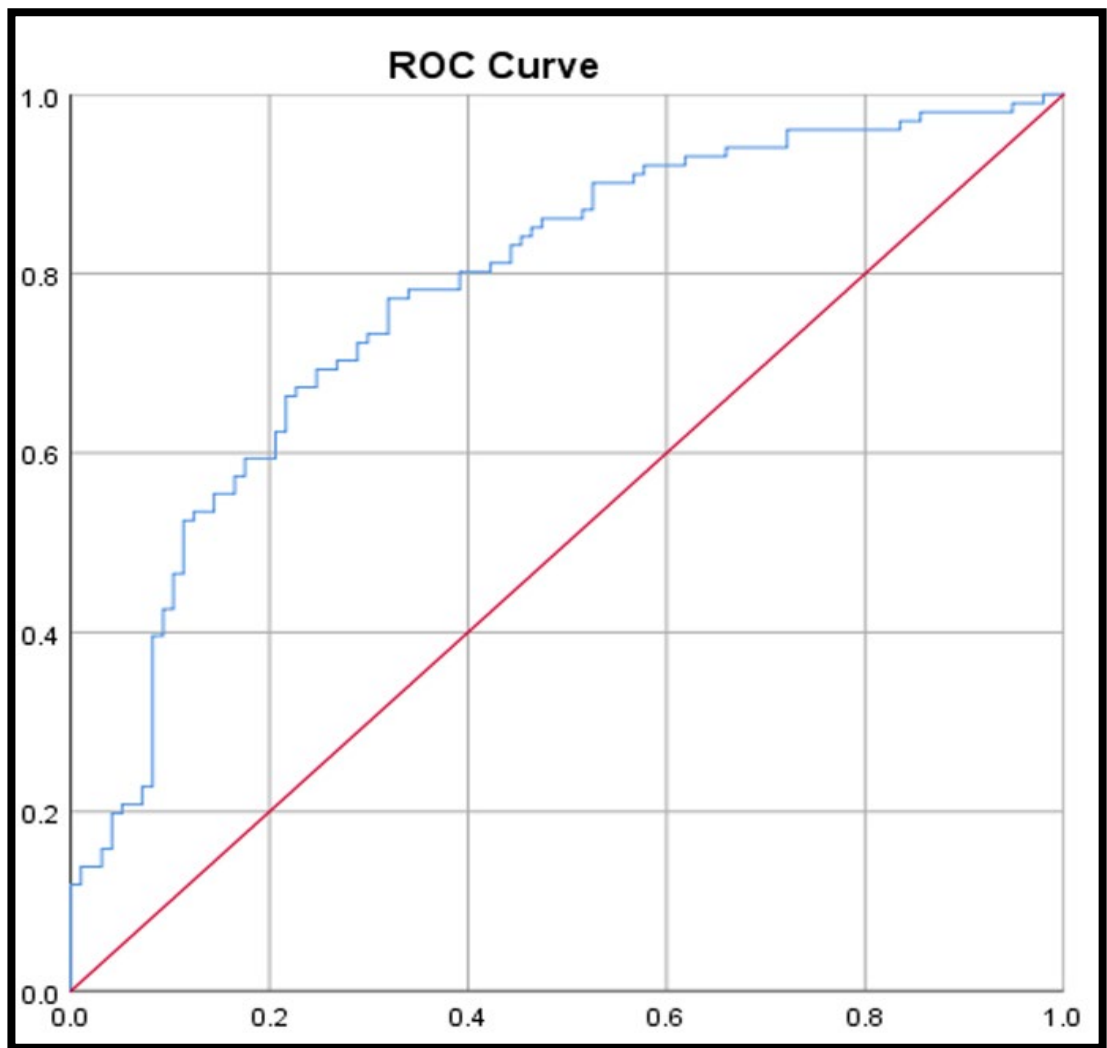


Figure 4-9 ROC curve and AUC for model discriminative ability.

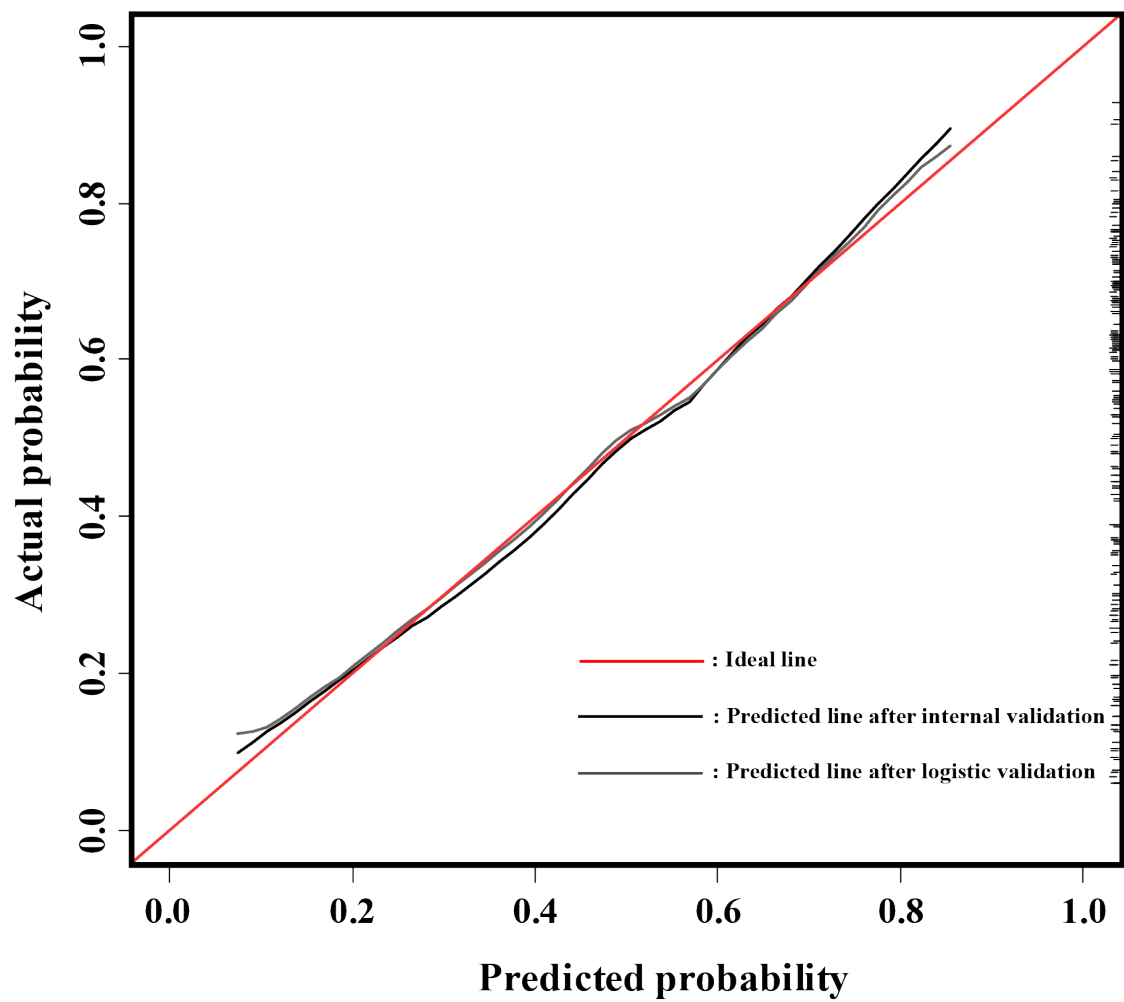


Figure 4-10 Model calibration plot for observed and predicted probability.

4.2.4 Decision curve analysis

The outcomes of the decision analysis curve are shown in Figure 4-11. The net benefit of performing SB in addition to TB on all cases is depicted by the grey line, whereas the black line represents the net benefit of not doing SB (only TB performed). The net benefit of using our prediction model was to identify patients who will benefit from SB in addition to TB at risk of having csPCa. Our nomogram increased the overall net clinical benefit when the threshold probability was <60% and improved the diagnosis of significant cancer compared to avoiding SB biopsy in all.

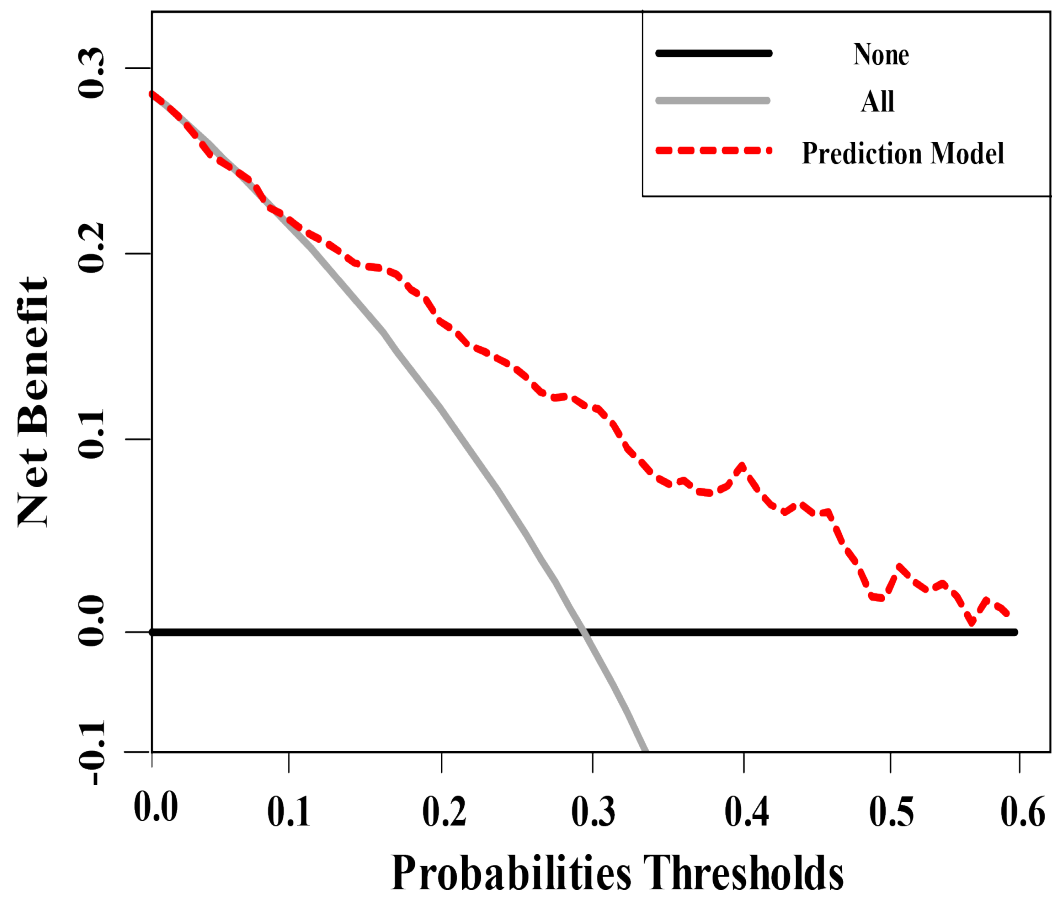


Figure 4-11 Decision analysis demonstrated a high net benefit of the model across a wide range of threshold probabilities.

4.3 Discussion

This study assessed the detection rate of clinically significant cancers using image fusion targeting, random systematic sampling and combination approaches. Patients-based analyses were further validated using lesion-based data from RP histology. There were statistically significant higher detection rates for the combined biopsy approach compared to SB or TB alone. The image-fusion TB approach alone would have missed 17 clinically significant PCa. Therefore, the combined approach detected more clinically significant cancers than either SB or TB alone. These results are similar to those reported by Filson et al (141) but differ to that reported in the PRECISION trial (11). Therefore, in our observations, omitting concurrent SB during image-fusion may run the risk of missing csPCa in around 8.5% of patients. Similar to Cash et al (142), we observed TB missing a small number clinically significant cancers in targeted areas. It is essential that we balance the advantages of concurrent sampling of prostate during targeting against the risks of side effects and increased detection of clinically insignificant cancer. Avoiding or adding SB at the time of fusion TB remains a challenge for physicians, as knowledge and evidence of decision-making contributing factors still remains unknown (143).

Our outcomes from the nomogram indicated the excellent advantage (C-index 78% vs. 70%) of using a multivariable prediction model adjusting for clinical and radiological features (age, PSAD and PIRADS). The nomogram could be used to assist in selecting a group of men where a combined biopsy approach would be more useful.

We also observed no significant advantage of improved characterisation of clinically significant PCa using the TB approach as all cancers labelled as clinically insignificant were upgraded on final histopathology of RP. The challenge of

upgrading or undergrading would continue with both biopsy approaches as seen in our previous study (144). There could be various reasons for this, including inadequate sampling due to cancer heterogeneity, poor visibility of cribriform architecture on MRI and in biopsies (2).

Several retrospective studies have assessed the outcomes of SB in addition to TB for the detection of clinically significant cancers. Sathianathan (145) et al reported a nomogram with C-index=70%. The nomogram was based on clinical variables (biopsy naivety, previous biopsy and AS patients), and imaging variables (number of MRI lesions and PIRADS score). The model provided a higher net clinical benefit at a threshold probability of <30%. The model was meant to predicts csPCa in systematic random cores only (when TB was negative). However, our finding focused on predicting those patients who will benefit from performing SB in addition to fusion TB (irrespective of the target biopsy being positive or negative). Also, unlike their study, our nomogram along with age, PSAD and PIRADS, found that adding these clinical variables to a model yielded a higher C-index (78% vs 70%). In contrast to the present study, Sathianathan et al (145) did not report on a validation cohort using RP as a reference standard. Further, and similar to our study, others have reported the possibility of missing significant cancers if the image fusion targeted approach was offered alone (38,146–148). Dell'Oglio et al (23) failed to identify patients who might benefit from fusion TB alone, therefore, they supported a combination of TB and SB as the preferred approach. In their study, there was no attempt to predict and assess the clinical variables that could help in identifying patients who will have a greater benefit from SB. Lastly, Falagario et al (149) highlighted that smaller lesions in a big prostate were more likely to be missed in TB biopsy, therefore, they developed a nomogram based on MRI volumetric parameters

and clinical information for deciding when SB should be performed in addition to TB. In their study, all patients underwent biparametric MRI; however, in our analysis we followed the standard method for prostate imaging (MP-MRI) using PIRADS (91). Moreover, the study was a multi-institutional retrospective analysis of the data from two previously published prospective trials with predominant fusion biopsies being cognitive rather than image-fusion using software. All men in the MULTI-IMPROD study (150,151) had transrectal systematic biopsies and these trials were not powered to answer the research question of the present study. In contrast to the present study, Falagario et al (149) provided a range of probabilities of missing clinically significant cancers in men if SB was to be avoided altogether. We reported a set of measurable imaging criteria which can predict the likely benefit of adding SB to TB.

To our knowledge, the present study is the first where lesion-based analyses were carried out using a mould-based approach for a comprehensive pathological analysis. This confirmed that most clinically significant cancers were detected using a combined biopsy approach. Clinically significant PCa were still missed by the biopsies and this may be due to smaller lesions or a cribriform pattern seen on histopathology (152).

Decision-making using critical analysis especially in situations of uncertainty, cost pressures and likely patient morbidity, is inevitably based on evidence or sets of observations. In this study, we have presented a decision-curve analysis estimating the net clinical benefits of offering a diagnostic test (combined approach to biopsy) in comparison to TB or SB approaches alone. The clinical and radiological observations have been used to construct a nomogram which provides the basis of the decision-making curve. The curve includes intervention for all and intervention

for none and provides background to facilitate discussions with patients. A balance has to be achieved between maximising the detection of clinically significant cancers and avoiding the detection of clinically insignificant cancers.

At present, various nomograms are used mainly for taking into consideration clinical factors such as age, pre-operative PSA level and PIRADS score of the suspicious cancers. The present study reports a nomogram based on clinical parameters (age, PSAD and PIRADS). The nomogram clearly showed an improved prediction which can be used to perform additional biopsy and the findings have substantial implications for clinicians and researchers in this area. We envisage that this and further research should take us close to precise decision-making. There will remain a group of men where a SB approach would bring value in addition to a fusion TB approach, and thereby improve informed decision making in the management of men suspected of PCa.

Even though many studies point to the accuracy of MRI/TRUS targeted biopsy in cancer diagnosis, there are few data about the standardisation of the procedure regarding the best approach (transrectal vs. transperineal). Recently there is a study investigating the detection rate of csPCa of TR versus TP targeted biopsy and found that the detection rate for csPCa was lower when the TR (66.7% of the cases) versus the TP (93.3% of the cases) approach was used. Moreover, the TP-targeted biopsies diagnosed more PCa (93.7%) located in the anterior zone of the prostate compared to the TR approach (25%) (153).

Our study had some limitations. This is a single centre study with dedicated uro-radiologists and pathologists. We wanted to explore the association between lesion location and csPCa via SB but due to low numbers of lesions in the TZ, this was not possible. It was not possible to use PIRADS v2.1 , since the enrollment to study

started before its publication, and this is a similar challenge to any other study published recently on this topic (21). We used fusion biopsy via transrectal approach however recent trends suggest that transperineal targeted biopsy could lead to detecting more clinically significant prostate cancer. The nomogram in the present study has been internally validated (Cross-validation and bootstrapping). External validation of the nomogram was not carried out in the present study as this would require further prospective multi-centre recruitment of a cohort.

Chapter 5 Conclusion and future work

In this thesis, the performance of pre-biopsy MRI for prostate cancer upgrading from biopsy to radical prostatectomy was tested. The results conclude that, a PIRADS score of 4 or 5 are associated with an increased risk of biopsy GS upgrading. In a multivariate analysis, the PIRADS score significantly improved prediction ability of pre-biopsy MRI scans for upgrading of Gleason Score ($p=0.001$, 95% CI [0.06-0.034]), which improved the C-index of predictive nomogram significantly (0.90 vs. 0.64, $p<0.05$). Currently, a variety of nomograms are used, with clinical parameters such as age, pre-operative PSA level and the number of biopsy cores involved in the cancer being taken into account. Despite this, there is still a significant histopathological difference between the biopsy and the final RP GS. This study presents a nomogram based on pre-biopsy MP-MRI grade (PIRADS score) and other known clinical parameters. We believe that this will bring us closer to a precise prediction of final GS of histopathology in prostate cancer, allowing stakeholders, including patients, to make better and more informed decisions in the treatment of localised prostate cancer.

In addition to prediction of upgrading GS, further findings also assessed the detection rate of csPCa using SB, TB and combined approaches. The image-fusion TB method would have missed 17 clinically significant PCa if it had been used alone. As a result, the combined approach detected more clinically significant cancers (64.6 %128/198) than either SB or TB alone. The results of the nomogram showed that using a multivariable prediction model that adjusts for clinical and radiological features (age, PSAD and PIRADS) has a significant benefit (C-index 78%). The nomogram might be used to help identify a group of men who would benefit from a

combined biopsy approach. The decision analysis curve confirmed a higher net clinical benefits of a combined approach (TB and SB) at an acceptable threshold. Future research should concentrate on incorporating additional predictor variables, such as novel genomic and other biomarkers to improve the accuracy of the model. The use of radiomics in conjunction with artificial intelligence and machine learning can improve our ability to identify tumour characteristics and classification. This would certainly have an effect on the study's findings in the future. Moreover, the nomogram in the study has been internally validated (Cross-validation and bootstrapping). External validation in a multi-institutional cohort is needed to predict the added values of SB to TB, which may help urologists avoid unnecessary biopsy sampling and reduce the detection of insignificant PCa while maintaining detection of clinically significant cancer.

In addition, understanding the limits of MP-MRI sensitivity for detecting tumours with Gleason pattern 4 is essential. MP-MRI is rapidly being adopted for PCa detection and surveillance, as well as a guide for focal therapy. GS pattern 4 is a heterogeneous group that can be further classified into architectural patterns such as poorly formed, cribriform, and fused glands. Improving risk assessment in this group is very crucial, as GS 7 prostate cancer on biopsy is an important clinical threshold for active treatment. Future study is warranted to evaluate the impact of Gleason pattern 4 architectural pattern (cribriform) on the sensitivity of MP-MRI for PCa detection and to compare the results with lesion-based data from RP histology.

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Appendix

Programming for nomogram, calibration and decision curve analysis using RStudio

```
library(foreign)
```

```
bc <- read.spss("E:/patients data.sav", use.value.labels=F, to.data.frame=T)
```

```
library(rms)
```

```
dd <- datadist(bc)
```

```
options(datadist="dd")
```

(First study)

1/With PI-RADS

```
f1 <- lrm(STATUS ~ PIRADS + PSA + CoresNumber + MaximumPercentage, data  
= bc)
```

```
lrm(formula = STATUS ~ PIRADS + PSA + CoresNumber + MaximumPercentage,  
data = bc)
```

```
nom <- nomogram(f1, fun= function(x)1/(1+exp(-x)), # or fun=plogis
```

```
lp=F, funlabel="Probability of GS upgrading",maxscale=10, fun.at=c(0.99, 0.975,  
0.95, 0.9, 0.8, 0.65, 0.5, 0.35, 0.2, 0.1, 0.05))
```

```
plot(nom)
```

```
f2 <- lrm(STATUS ~ PIRADS + PSA + CoresNumber + MaximumPercentage,  
x=T,y=T,data = bc)
```

```
cal2 <- calibrate(f2, cmethod="KM", method="boot", m=110, B=1000)
```

```
plot(cal2)
```

```
plot(cal2,xlab="Predicted probability of GS upgrading",ylab="Actual probability of  
GS upgrading")
```

2/without PI-RADS

```
f3 <- lrm(STATUS ~ PSA + CoresNumber + MaximumPercentage, data = bc)
```

```
lrm(formula = STATUS ~ PSA + CoresNumber + MaximumPercentage, data = bc)
```

```
nom <- nomogram(f3, fun= function(x)1/(1+exp(-x)), # or fun=plogis
```

```
lp=F, funlabel="Probability of GS upgrading",maxscale=10, fun.at=c(0.99, 0.975,  
0.95, 0.9, 0.8, 0.65, 0.5, 0.35, 0.2, 0.1, 0.05))
```

```
plot(nom)
```

```
f4 <- lrm(STATUS ~ PSA + CoresNumber + MaximumPercentage, x=T,y=T,data =  
bc)
```

```
cal4 <- calibrate(f4, cmethod="KM", method="boot", m=110, B=1000)
plot(cal4)
plot(cal4,xlab="Predicted probability of GS upgrading",ylab="Actual probability of
GS upgrading")
```

(Second study)

Nomogram:

```
bc <- read.spss("E:/filename.sav",
use.value.labels = F,
to.data.frame = T)
dd <- datadist(bc)
options(datadist = "dd")
f1 <-
lrm(Outcome ~ PIRADS + Age + PSAdensity, data = bc)
lrm(formula = Outcome ~ PIRADS + Age + PSAdensity,
data = bc)
# rt2$gender <- factor(rt2$gender,labels=c("F", "M"))
# rt2$stage <- factor(rt2$stage,labels=c("Stage1", "Stage2", "Stage3", "Stage4"))
# rt2$T <- factor(rt2$T,labels=c("T1", "T2", "T3", "T4"))4 rt2$M <-
factor(rt2$M,labels=c("M0", "M1"))
# rt2$N <- factor(rt2$N,labels=c("N0", "N1", "N2", "N3"))6 rt2$risk <-
factor(rt2$risk,labels=c("low", "high"))
nom <- nomogram(
f1,
fun = function(x)
1 / (1 + exp(-x)),
# or fun=plogis+lp = F,
lp=F,
funlabel = "Probability",
maxscale = 10,
fun.at = c(0.99, 0.975, 0.95, 0.9, 0.8, 0.65, 0.5, 0.35, 0.2, 0.1, 0.05))
plot(nom)
calibration plot:
f2 <- lrm(Outcome ~ PIRADS + Age + PSAdensity, x=T,y=T,data = bc)
```

```

cal2 <- calibrate(f2, cmethod="KM", method="boot", m=39, B=200)
p <- cal2[, "predy"]
p.app <- cal2[, "calibrated.orig"]
p.cal <- cal2[, "calibrated.corrected"]
xlim <- c(0, 1)
ylim <- c(0, 1)
at <- attributes(cal2)
predicted <- at$predicted
s <- !is.na(p + p.cal)
err <- predicted - approx(p[s], p.cal[s], xout=predicted, ties=mean)$y
cat('\nn=', n <- length(err), ' Mean absolute error=',
    round(mae <- mean(abs(err), na.rm=TRUE), 3), ' Mean squared error=',
    round(mean(err^2, na.rm=TRUE), 5),
    '\n0.9 Quantile of absolute error=',
    round(quantile(abs(err), .9, na.rm=TRUE), 3), '\n\n', sep="")
plot(p, p.app, xlim=xlim, ylim=ylim,
     xlab="Predicted probability",
     ylab="Actual probability",
     type="n")
lines(p, p.app, lty=1, col='blue') ## change color of apparent
lines(p, p.cal, lty=1, col='red') ## change color of bias-corrected
abline(0,
1,
lty = 1,
lwd = 2,
col = c(rgb(150, 250, 50, max = 255)))
do.call('scat1d', c(list(x=predicted), 'scat1d.opts'))
legend <- list(x=xlim[1] + .55*diff(xlim),
y=ylim[1] + .32*diff(ylim))
legend(legend, c("Apparent", "Bias-corrected", "Ideal"),
lty=c(3,1,2), bty="n")
title(sub=paste("B=", at$B, "repetitions,", at$method),
cex.sub=0.75, adj=0)

```

```

title(sub=paste('Mean absolute error=', round(mae,3),
' n=', n, sep="),
cex.sub=0.75, adj=1)
DCA analysis:
library(devtools)
Data <- read.spss("filename.sav",
use.value.labels = F,
to.data.frame = T)
simple <-
decision_curve(
Outcome ~ PIRADS + Age + PSAdensity,
data = Data,
family = binomial(link = 'logit'),
thresholds = seq(0, 0.6, by = 0.01),
confidence.intervals = 0.95,
study.design = 'case-control',
population.prevalence = 0.3)
List <- list(simple)
plot_decision_curve(
List,
curve.names = c('model'),
cost.benefit.axis = FALSE,
col = c('red'),
confidence.intervals = FALSE,
standardize = FALSE,
xlab="Probabilities Thresholds")
table=summary(simple,measure= 'NB')

```

Raw data 1

1	LRP	NO.	CHI	age	sitename	MRI	PIRAD	PSA	core	percenta	stage	TRUS	Pathology	Weigh	PSAD
2	16/07/2013	1	1309600031	53	CRC	3T	4	5.9	2	20	T0	3+3	3+4	34.5	0.17101
3	09/07/2013	2	2610500017	63	CRC	3T	4	8.5	3	10	T0	3+3	3+4	29	0.2931
4	15/08/2013	3	2107440077	69	CRC	3T*	4	11	3	20	T2b	3+4	4+5	58.5	0.18803
5	08/10/2013	4	1901410099	72	STRACATHRO	1.5T	5	11.1	7	50	T2	4+3	3+4	66	0.16818
6	08/10/2013	5	1604445254	69	STRACATHRO	1.5T	4	17	6	70	T2	4+4	4+3	40	0.425
7	12/11/2013	6	0311450059	68	NINEWELLS	1.5T	5	9	3	20	T0	3+3	3+4	58.5	0.15385
8	12/11/2013	7	0811430375	70	CRC	3T	4	6.6	4	20	T2	3+4	4+5	85	0.07765
9	19/11/2013	8	1708540253	59	STRACATHRO	1.5T	4	5.5	3	50	T2	3+4	3+4	26.5	0.20755
10	19/11/2013	9	0105480738	65	STRACATHRO	1.5T	5	4.9	10	50	T0	3+4	3+4	76.5	0.06405
11	10/12/2013	10	260290652	64	NINEWELLS	1.5T	5	23	6	70	T2	4+4	3+4	42	0.54762
12	16/12/2013	11	1505470951	66	STRACATHRO	1.5T	5	22.9	6	90	T2	4+5	4+5	48.5	0.47216
13	31/12/2013	12	0711430233	70	CRC	3T	5	6.5	3	20	T1c	3+4	3+4	39	0.16667
14	27/12/2013	13	2401430512	70	NINEWELLS	1.5T	5	11	10	90	T2	4+3	3+4	60.5	0.18182
15	25/02/2014	14	0205530052	61	NINEWELLS	1.5T	5	44	11	90	T2	4+4	4+5	61	0.72131
16	04/03/2014	15	2401570012	57	CRC	3T	3	7.8	2	100	T2	3+4	3+4	26	0.3
17	18/03/2014	16	2608495079	65	NINEWELLS	1.5T	4	21.6	1	10	T2	3+3	4+5	73	0.29589
18	18/03/2014	17	0810500574	64	NINEWELLS	1.5T	4	9	4	25	T0	3+3	3+3	114.5	0.0786
19	25/03/2014	18	0106410059	73	CRC	3T	5	7.4	4	10	T2	3+4	4+5	58	0.12759
20	15/04/2014	19	0903560038	58	CRC	3T	5	8.1	1	80	T2	4+5	4+5	53	0.15283
21	15/04/2014	20	2311540653	60	CRC	3T	5	34	7	90	T2	4+5	4+5	39.5	0.86076
22	29/04/2014	21	3006480171	66	NINEWELLS	1.5T	5	13.6	7	70	T1c	4+3	4+5	53	0.2566
23	29/04/2014	22	1505470293	67	NINEWELLS	1.5T	5	10.2	5	40	T1c	3+4	4+3	49.5	0.20606
24	07/05/2014	23	0407430318	71	STRACATHRO	1.5T	5	10	6	60	T2	3+4	3+4	37.5	0.26667
25	06/05/2014	24	1106410173	73	STRACATHRO	1.5T	5	10.6	10	100	T2b	4+5	4+5	94.5	0.11217
26	20/04/2014	25	2710450011	69	STRACATHRO	1.5T	5	33.2	1	70	T3	4+3	4+5	51.5	0.64466
27	27/05/2014	26	1511430079	71	CRC	3T	5	11.4	4	80	T2	4+4	4+5	61	0.18689
28	26/05/2014	27	1608509796	64	Queens Margaret	1.5T	5	6.9	5	90	T3c	4+4	4+5	52.5	0.13143
29	03/06/2014	28	0607536179	61	CRC	3T	5	18.4	8	80	T2B	3+5	3+5	72.5	0.25379
30	13/06/2014	29	2305460074	68	STRACATHRO	1.5T	5	6.5	7	100	T2	4+4	3+4	48.5	0.13402
31	18/06/2014	30	1508500312	64	CRC	3T	5	8	5	80	T2	4+3	4+3	52	0.15385
32	18/06/2014	31	2802550152	59	CRC	3T*	N	6.1	2	10	T1c	3+3	3+3	49.5	0.12323
33	01/07/2014	32	2704473595	67	CRC	3T	5	8.1	2	30	T1c	5+4	3+5	50.5	0.1604
34	08/07/2014	33	1604470275	67	STRACATHRO	1.5T	N	8	3	20	T1c	3+4	3+4	58	0.13793
35	09/07/2014	34	2605470113	67	CRC	3T*	3	16	2	20	T1c	3+4	3+4	160	0.1
36	29/07/2014	35	2705480390	66	CRC	3T	5	20.1	5	15	T1c	3+5	3+4	77.5	0.25935
37	29/07/2014	36	0712500154	64	STRACATHRO	1.5T	4	10	6	80	T2c	4+3	3+4	49.5	0.20202
38	12/08/2014	37	0106420313	72	NINEWELLS	1.5T	5	19.6	5	90	T2	3+3	3+4	52	0.37692
39	25/08/2014	38	0906472490	67	STRACATHRO	1.5T	5	23	9	70	T1c	3+4	3+4	118	0.19492
40	02/09/2014	39	0808500074	64	CRC	3T*	5	10.5	2	25	T1c	3+3	3+4	43	0.24419
41	02/09/2014	40	3008531350	61	STRACATHRO	1.5T	3	20.5	2	5	T1c	3+3	3+4	155	0.13226
42	09/09/2014	41	0909420211	72	STRACATHRO	1.5T	N	5.7	1	10	T1c	3+4	3+4	125.5	0.04542
43	15/09/2014	42	1902530330	61	STRACATHRO	1.5T	5	9.4	12	100	T1c	5+4	4+5	52.5	0.17905
44	23/09/2014	43	0603470017	67	CRC	3T	5	10.2	4	20	T1c	3+4	3+4	46	0.22174
45	23/09/2014	44	1804490350	65	VICTORIA HOS	1.5T	N	4.6	3	30	T1c	3+3	3+4	62.5	0.0736
46	30/09/2014	45	1408420058	72	STRACATHRO	1.5T	3	12.5	3	5	T1c	3+3	3+4	137	0.09124
47	30/09/2014	46	0604440197	70	Queens Margaret	1.5T	4	10	8	90	T2	4+4	4+3	48	0.20833

48	21/10/2014	47	2911430115	71	CRC	3T	5	42	3	50	T2	3+4	4+5	45.5	0.92308
49	21/10/2014	48	0509410472	73	CRC	3T*	5	25	7	100	T2	3+4	4+5	105.5	0.23697
50	14/10/2014	49	1205500235	64	CRC	3T	4	11	1	10	T1c	3+3	3+4	50.5	0.21782
51	03/11/2014	50	1811380077	76	STRACATHRO	1.5T	5	8.9	5	60	T1c	4+5	4+5	50.5	0.17624
52	01/11/2014	51	1802530355	61	CRC	3T	5	37	2	20	T2	3+3	3+4	63	0.5873
53	01/11/2014	52	1308530315	61	NINEWELLS	1.5T	5	26.5	11	100	T2	3+4	3+4	51.5	0.51456
54	28/11/2014	53	2903420130	72	NINEWELLS	1.5T	4	11.1	4	20	T2	5+3	3+5	48	0.23125
55	16/12/2014	54	0510490433	65	CRC	3T	3	8.5	1	20	T1c	4+3	4+3	127.5	0.06667
56	05/01/2015	55	1102530115	62	STRACATHRO	1.5T	5	6.9	2	20	T1c	3+4	3+4	34	0.20294
57	05/01/2015	56	1704493412	66	CRC	3T	4	9.8	2	20	T1c	4+4	4+3	67.5	0.14519
58	13/01/2015	57	0402475372	68	CRC	3T*	3	8.6	5	70	T1c	3+4	3+4	121	0.07107
59	13/01/2015	58	0512400059	75	CRC	3T*	5	8	4	70	T2	4+3	3+4	38	0.21053
60	09/01/2015	59	1503560074	59	CRC	3T*	N	6.8	1	10	T1c	3+3	3+3	74	0.09189
61	27/01/2015	60	1803410256	74	CRC	3T	3	11	12	90	T1c	3+3	3+4	34	0.32353
62	03/02/2015	61	2603425218	73	CRC	3T*	4	12.1	5	30	T0	4+4	4+3	53	0.2283
63	03/02/2015	62	1806485575	67	NINEWELLS	1.5T	5	15.2	4	30	T2	3+4	3+4	70	0.21714
64	17/02/2015	63	1011440091	71	NINEWELLS	1.5T	5	7.1	2	20	T2	3+3	3+4	59.5	0.11933
65	20/02/2015	64	0311461131	69	NINEWELLS	1.5T	5	47	3	40	T2	4+3	3+5	55	0.85455
66	24/02/2015	65	1305500474	65	CRC	3T*	5	28	12	80	T1c	3+5	3+5	54.5	0.51376
67	10/03/2015	66	2806490251	66	CRC	3T*	5	10	1	5	T1c	3+3	3+3	72	0.13889
68	17/03/2015	67	0803460198	69	CRC	3T*	4	0.9	3	60	T1c	4+4	4+4	36.5	0.02466
69	31/03/2015	68	1803535474	62	NINEWELLS	3T*	5	9	8	80	T1c	4+5	3+5	69	0.13043
70	31/03/2015	69	1211460193	69	STRACATHRO	1.5T	5	10	10	50	T2	4+5	4+5	44	0.22727
71	21/04/2015	70	0506530175	62	STRACATHRO	1.5T	5	15.5	8	80	T2	4+4	4+5	68	0.22794
72	28/04/2015	71	1809575451	58	NINEWELLS	1.5T	5	12.2	12	90	T1c	4+5	4+5	52	0.23462
73	28/04/2015	72	0302455310	70	NINEWELLS	1.5T	5	10	3	20	T2	4+4	4+4	55	0.18182
74	12/05/2015	73	0212412213	74	NINEWELLS	1.5T	5	19.4	3	70	T2	4+4	4+3	65	0.29846
75	19/05/2015	74	0110430018	72	CRC	3T	5	6.7	5	70	T2	4+3	4+3	70.5	0.09504
76	02/06/2015	75	1907530258	62	NINEWELLS	1.5T	4	9.4	9	50	T2	3+4	3+4	47	0.2
77	02/06/2015	76	0705410072	74	STRACATHRO	1.5T	5	7.5	6	50	T2	4+3	3+4	101	0.07426
78	09/06/2015	77	1110510012	64	STRACATHRO	1.5T	5	7.9	5	80	T0	4+5	4+5	59	0.1339
79	23/06/2015	78	2212430132	72	STRACATHRO	1.5T	5	13.3	2	10	T1c	4+3	3+5	108	0.12315
80	30/06/2015	79	2706440538	71	NINEWELLS	1.5T	4	4.5	2	30	T2	4+3	4+3	62	0.07258
81	14/07/2015	80	2512510514	64	CRC	3T	N	10	3	20	T1c	4+3	4+3	86	0.11628
82	21/07/2015	81	1211550095	60	CRC	3T	5	8.1	10	90	T2	3+4	3+4	68.5	0.11825
83	04/08/2015	82	2511520311	63	CRC	3T	3	4.2	3	20	T2	4+5	4+3	35.5	0.11831
84	11/08/2015	83	2111530131	62	CRC	3T	5	8.8	1	30	T1c	3+4	3+4	73.7	0.1194
85	11/08/2015	84	3001500131	65	NINEWELLS	1.5T	4	6.8	4	30	T1c	3+4	3+4	72.3	0.09405
86	01/09/2015	85	2405490072	66	STRACATHRO	1.5T	5	3.1	6	40	T0	4+4	4+3	57	0.05439
87	01/09/2015	86	1010410016	74	NINEWELLS	1.5T	4	7.6	1	20	T1c	3+4	3+4	120	0.06333
88	22/09/2015	87	1212420276	73	NINEWELLS	1.5T	5	2	6	90	T2	4+4	4+5	42	0.04762
89	07/10/2015	88	2711462196	69	STRACATHRO	1.5T	5	10.3	9	90	T1c	4+5	4+5	128.3	0.08028
90	05/10/2015	89	2312510138	64	NINEWELLS	1.5T	5	11.3	3	75	T2	N	4+5	71.5	0.15804
91	05/10/2015	90	2308450010	70	CRC	3T	5	11.7	1	30	T1c	3+4	4+3	45.6	0.25658
92	09/10/2015	91	1501560115	59	NINEWELLS	1.5T	4	2.1	6	75	T2	3+4	3+4	31.2	0.06731
93	09/10/2015	92	0807460079	69	STRACATHRO	1.5T	5	10.7	6	20	T1c	3+4	3+4	118	0.09068

94	09/10/2015	93	6605440034	71	CRC	3T	5	8.5	2	70	T1c	3+4	3+4	69.5	0.1223
95	26/10/2015	94	2003440275	71	NINEWELLS	1.5T	5	10.1	1	50	T1c	3+4	3+4	161.4	0.06258
96	26/10/2015	95	2709510677	64	CRC	3T	5	10.8	10	90	T2	4+3	4+3	51.5	0.20971
97	03/11/2015	96	2203460555	69	CRC	3T	5	10.9	11	90	T1c	4+3	3+4	40.5	0.26914
98	28/10/2015	97	604560037	59	CRC	3T	5	9.3	11	100	T2	4+3	4+5	59.5	0.1563
99	17/11/2015	98	309490171	66	CRC	3T	5	8.1	6	80	T2	5+4	4+3	43.2	0.1875
100	17/11/2015	99	0712480277	67	STRACATHRO	1.5T	5	9.2	2	30	T2	4+3	3+4	72.5	0.1269
101	24/11/2015	100	012420094	67	STRACATHRO	1.5T	5	11.4	3	10	T2	3+3	3+4	60.5	0.18843
102	08/12/2015	101	0207450235	70	STRACATHRO	1.5T	5	9	1	30	T2	3+3	3+5	69.7	0.12912
103	08/12/2015	102	0903395193	76	STRACATHRO	1.5T	5	22	1	10	T2	4+4	4+3	83	0.26506
104	15/12/2015	103	906570078	58	CRC	3T	3	0.9	1	60	T2	3+4	3+4	47	0.01915
105	05/01/2016	104	2405530236	62	STRACATHRO	1.5T	5	3.8	6	90	T2	4+4	4+3	32.8	0.11585
106	29/12/2015	105	0103420193	73	CRC	3T	4	5	2	80	T2	3+4	3+4	36	0.13889
107	29/12/2015	106	02011440157	71	NINEWELLS	1.5T	N	7.6	1	5	T2	3+3	3+4	99.25	0.07657
108	19/01/2016	107	2406530159	62	STRACATHRO	1.5T	4	6	1	20	T1c	3+4	3+4	75.9	0.07905
109	09/02/2016	108	2306330059	62	STRACATHRO	1.5T	4	11	2	10	T1c	3+3	3+3	115.4	0.09532
110	25/01/2016	109	2904450238	71	CRC	3T	5	11.2	14	90	T2	4+4	3+5	65	0.17231
111	25/01/2016	110	1605575577	59	STRACATHRO	1.5T	5	22	3	80	T3b	4+5	4+5	89	0.24719
112	26/01/2016	111	0402460111	70	not suit for MRI		no mri	7.1	7	70	T1c	N	3+4	71.8	0.09889
113	26/01/2016	112	2608540031	62	NINEWELLS	1.5T	5	9	1	30	T1c	3+4	3+4	39.8	0.22613
114	09/02/2016	113	270490055	67	STRACATHRO	1.5T	4	8.7	1	5	T1c	4+3	4+3	58.5	0.14872
115	09/02/2016	114	704510295	65	CRC	3T	3	9.7	6	80	T2	3+4	3+4	128	0.07578
116	16/02/2016	115	0906490170	67	CRC	3T*	5	8.3	2	25	T2b	3+4	3+4	62.5	0.1328
117	16/02/2016	116	1312430117	73	STRACATHRO	1.5T	5	9.4	8	80	T1c	3+4	3+4	58	0.16207
118	01/03/2016	117	1601390491	77	STRACATHRO	1.5T	5	47.7	1	10	T2	3+4	3+5	43	1.1093
119	01/03/2016	118	2507460156	70	STRACATHRO	1.5T	5	16	5	70	T3a	3+3	3+4	73.5	0.21769
120	08/03/2016	119	0212460013	70	CRC	3T*	5	4.6	1	50	T2c	3+4	3+4	76	0.06053
121	08/03/2016	120	1611520096	64	NINEWELLS	1.5T	5	15.8	10	60	T2	3+4	3+5	90	0.17556
122	22/03/2016	121	3103450036	71	CRC	3T	3	12.6	1	10	T2	3+3	3+4	90.8	0.13877
123	22/03/2016	122	1502450054	71	CRC	3T	4	11.2	12	80	T2	3+4	3+4	76	0.14737
124	29/03/2016	123	2702433359	73	CRC	3T	5	9.8	1	50	T3a	3+4	3+4	44.5	0.22022
125	29/03/2016	124	0309440017	72	NINEWELLS	1.5T	5	12	3	10	T2c	3+4	3+5	39	0.30769
126	12/04/2016	125	1205490213	67	Claustrophobic		cl	2.9	3	40	T2	4+3	3+5	81	0.0358
127	15/04/2016	126	1911520199	64	CRC	3T	5	9	2	70	T1c	3+4	3+4	54.5	0.16514
128	03/05/2016	127	2802445456	72	NINEWELLS	1.5T	5	17.4	5	100	T2c	3+4	4+3	89	0.19551
129	03/05/2016	128	1508400032	76	CRC	3T	5	19.1	7	60	T1c	4+3	4+3	41.5	0.46024
130	10/05/2016	129	1509440011	72	CRC	3T*	4	5.7	2	40	T2	3+4	4+4	77	0.07403
131	10/05/2016	130	0605460590	70	CRC	3T*	5	7.9	1	60	T2b	4+4	4+3	110	0.07182
132	19/05/2016	131	2801420077	74	NINEWELLS	1.5T	5	15.8	12	95	T3a	4+5	4+5	61.5	0.25691
133	31/05/2016	132	2905500115	66	NINEWELLS	1.5T	5	2.9	1	50	T1c	4+3	4+3	64	0.04531
134	31/05/2016	133	1003420257	74	NINEWELLS	1.5T	5	9	8	30	T1c	3+3	3+4	93.5	0.09626
135	14/06/2016	134	1304410196	75	CRC	3T	5	5.4	9	80	T0	3+3	4+3	31	0.17419
136	14/06/2016	135	1608390373	77	Claustrophobic		cl	5.6	10	70	T2b	4+5	4+5	20	0.28
137	21/06/2016	136	2309395277	77	PERTH	1.5T	5	7	4	80	T2c	4+3	3+4	61.5	0.11382
138	21/06/2016	137	1107390516	77	PERTH	1.5T	4	5.1	1	10	T3a	3+3	3+4	106.5	0.04789
139	05/07/2016	138	1508410097	75	NINEWELLS	1.5T	4	8	1	50	T2c	3+4	3+4	170	0.04706
140	05/07/2016	139	0911460136	70	STRACATHRO	1.5T	N	15	2	10	T2	3+3	4+5	66.5	0.22556
141	12/07/2016	140	010535552	63	STRACATHRO	1.5T	4	4.6	6	30	T2	4+5	4+5	39	0.11795
142	12/07/2016	141	2008490793	67	PERTH	1.5T	5	19	6	60	T1c	4+4	4+5	103	0.18447
143	27/07/2016	142	1705500056	66	CRC	3T	4	6.5	6	40	T2c	4+3	4+3	38.9	0.1671
144	27/07/2016	143	2111510033	65	STRACATHRO	1.5T	5	14.4	7	80	T2c	4+3	4+3	60	0.24
145	16/08/2016	144	2906532096	63	STRACATHRO	1.5T	5	10.8	3	10	T2	3+4	3+4	46.4	0.23276
146	16/08/2016	145	2110410094	75	CRC	3T	5	11.3	2	40	T2c	3+5	3+5	61.5	0.18374
147	19/08/2016	146	0406530831	63	STRACATHRO	1.5T	4	4	1	30	T1c	3+4	3+4	36	0.11111
148	23/08/2016	147	0710475594	69	CRC	3T	5	12	1	15	T1c	3+4	3+4	85.5	0.14035
149	23/08/2016	148	0311450113	71	CRC	1.5T	5	7.1	1	50	T2c	3+4	3+5	86	0.08256
150	06/09/2016	149	1103440055	72	NINEWELLS	1.5T	5	14.4	8	70	T1c	4+3	4+3	42	0.34286
151	13/09/2016	150	2209500079	66	CRC	3T	4	4.8	7	20	T1	3+3	3+4	35.5	0.13521
152	13/09/2016	151	1103430394	73	PERTH	1.5T	4	9	7	60	T1c	3+4	3+4	45.1	0.19956
153	27/09/2016	152	1709450339	71	NINEWELLS	1.5T	5	16	4	30	T1	4+4	4+5	54.2	0.2952
154	30/09/2016	153	0906580536	58	CRC	3T	5	7	8	40	T1c	4+3	3+4	54.3	0.12891
155	04/10/2016	154	0706510054	65	CRC	3T	5	14	3	60	T2c	3+4	4+5	87	0.16092

155	04/10/2016	154	0706510054	65	CRC	3T	5	14	3	60	T2c	3+4	4+5	87	0.16092
156	04/10/2016	155	2705430059	73	CRC	3T	5	13.9	1	30	T2c	3+4	3+4	207	0.06715
157	11/10/2016	156	0608555339	61	STRACATHRO	1.5T	5	9.3	5	40	T1c	3+4	3+4	68	0.13676
158	18/10/2016	157	0211442216	72	CRC	3T	3	30	1	10	T1c	4+3	4+5	100	0.3
159	18/10/2016	158	1609590058	57	CRC	3T	4	8.5	4	10	T2a	3+3	3+4	60	0.14167
160	25/10/2016	159	1712400398	76	CRC	3T	4	7.5	3	10	T2a	3+3	3+4	71.5	0.1049
161	25/10/2016	160	1704400236	76	STRACATHRO	1.5T	5	6	1	50	T2	4+4	4+5	118	0.05085
162	08/11/2016	161	2707500011	66	CRC	3T	5	3.8	2	5	T2c	3+3	3+4	46.3	0.08207
163	08/11/2016	162	2303420415	74	CRC	3T	5	9	10	80	T2c	4+4	4+5	88.5	0.10169
164	15/11/2016	163	2901470297	69	NINEWELLS	1.5T	4	7.1	8	40	T1c	3+4	3+4	61.5	0.11545
165	15/11/2016	164	1712509799	66	CRC	3T	3	8	2	20	T1c	3+4	3+4	78.8	0.10152
166	11/11/2016	165	1110560011	60	CRC	3T	5	11.8	12	80	T3b	4+4	4+5	36	0.32778
167	06/12/2016	166	0810640392	52	CRC	3T	3	8	6	50	T0	4+3	3+4	34.5	0.23188
168	06/12/2016	167	1908420030	74	CRC	3T	5	8	2	5	T2a	3+3	3+4	63	0.12698
169	29/11/2016	168	1903560934	60	CRC	3T	5	6.4	10	100	T2c	3+5	3+4	57	0.11228
170	20/12/2016	169	1207430153	73	CRC	3T	4	8	3	10	T2c	3+4	3+4	47.8	0.16736
171	23/12/2016	170	0104610158	55	CRC	3T	5	7.6	5	80	T2c	4+3	3+5	36.7	0.20708
172	10/01/2017	171	1104440016	72	STRACATHRO	1.5T	5	11.1	2	20	T2c	3+3	4+3	62	0.17903
173	10/01/2017	172	1507430019	73	STRACATHRO	1.5T	5	15	7	40	T2c	4+4	4+5	57.2	0.26224
174	23/01/2017	173	0903565412	61	CRC	3T	4	7	5	40	T2	3+3	3+4	120	0.05833
175	31/01/2017	174	2101500117	67	CRC	3T	5	8	2	30	T2c	4+4	3+4	47.5	0.16842
176	31/01/2017	175	1805532391	64	CRC	3T	5	10.6	2	80	T2c	4+3	4+3	66.5	0.1594
177	03/02/2017	176	1806461293	71	PERTH	1.5T	4	9.6	2	60	T3	3+4	3+4	52.7	0.18216
178	07/02/2017	177	1308551231	62	CRC	3T	5	14.5	10	90	T2c	4+3	4+3	51	0.28431
179	13/02/2017	178	0201420716	75	CRC	3T	4	14.2	6	90	T2c	4+3	3+4	82.8	0.1715
180	22/02/2017	179	0704430118	74	NINEWELLS	1.5T	5	10.1	4	20	T2	3+4	3+4	107	0.09439
181	22/02/2017	180	204520011	65	NINEWELLS	1.5T	5	5.6	3	20	T3a	3+3	3+4	68.5	0.08175
182	28/02/2017	181	1901490394	68	PERTH	1.5T	5	11.2	10	60	T3a	4+3	4+3	92	0.12174
183	02/03/2017	182	1503470296	70	NINEWELLS	1.5T	5	0.1	6	40	T2	3+4	3+4	52.3	0.00191
184	06/03/2017	183	2909440036	73	STRACATHRO	1.5T	5	6.5	1	10	T3a	3+3	4+5	54	0.12037
185	14/03/2017	184	0202450090	72	STRACATHRO	1.5T	5	17.7	12	95	T2c	3+4	4+3	46.5	0.38065
186	17/03/2017	185	1611540070	63	NINEWELLS	1.5T	5	19.2	1	40	T3a	3+3	3+4	90.5	0.21215
187	27/03/2017	186	0210410116	76	NINEWELLS	1.5T	5	8.2	9	90	T3a	4+3	3+4	33.7	0.24332
188	28/03/2017	187	2504420196	75	NINEWELLS	1.5T	5	25	6	80	T2c	4+3	4+3	56.3	0.44405
189	04/04/2017	188	0405550154	62	NINEWELLS	1.5T	5	6.5	8	50	T3a	4+4	4+5	45.5	0.14286
190	11/04/2017	189	0605523215	65	CRC	3T	5	10.8	3	40	T2c	4+3	4+5	75.5	0.14305
191	11/04/2017	190	1709470178	70	CRC	3T	5	9.3	4	30	T2b	3+4	3+4	50.5	0.18416
192	11/04/2017	191	1302560212	61	NINEWELLS	1.5T	4	3.5	7	90	T2	4+4	4+5	34.5	0.10145
193	25/04/2017	192	0501431152	74	NINEWELLS	1.5T	5	10.6	6	80	T3a	4+5	4+5	58	0.18276
194	28/04/2017	193	2205500295	67	CRC	3T	3	6.9	2	60	T2c	3+4	3+4	79.5	0.08679
195	02/05/2017	194	0704550636	62	CRC	3T	5	6.9	3	25	T1c	3+3	3+4	59	0.11695
196	02/05/2017	195	1610412133	77	STRACATHRO	1.5T	5	11.9	2	50	T2b	3+3	3+4	44	0.27045
197	16/05/2017	196	0404490190	68	CRC	3T	5	9.2	3	40	T2c	3+3	3+4	71	0.12958
198	16/05/2017	197	0910450293	72	STRACATHRO	1.5T	4	23	3	10	T2c	3+4	3+4	82.5	0.27879
199	17/05/2017	198	2703640617	53	CRC	3T	5	22	12	70	T3b	3+4	4+5	58.3	0.37736
200	23/05/2017	199	309510058	66	STRACATHRO	1.5T	3	10	3	40	T2a	3+4	3+4	147	0.06803
201	30/05/2017	200	2809610134	56	NINEWELLS	1.5T	5	13	3	75	T2c	3+4	3+5	71	0.1831
202	31/05/2017	201	1201640016	53	STRACATHRO	1.5T	5	7.3	11	90	T3a	4+4	3+5	45	0.16222
203	06/06/2017	202	1402470339	70	CRC	3T	5	11.7	9	70	T2c	4+5	3+5	68.9	0.16981
204	09/06/2017	203	0107730057	44			4	3.2	2	20	T2	4+3	3+4	28.8	0.11111
205	12/06/2017	204	1910550272	62		N	8.3	7	40	T2c	3+4	3+4	51.5	0.16117	
206	13/06/2017	205	0705550354	62	CRC	3T	4	15	6	85	T2c	3+4	4+5	76	0.19737
207	13/06/2017	206	1103430297	74			4	11.3	1	20	T2a	3+3	4+3	56.5	0.2
208	21/06/2017	207	1607520397	65	CRC	3T	4	7.2	6	75	T3a	4+3	4+5	91	0.07912
209	27/06/2017	208	0302480439	69			5	6.2	5	30	T2c	3+4	3+4	52	0.11923
210	30/06/2017	209	0007580390	59	CRC	3T	5	15.9	12	90	T3a	4+5	4+5	94	0.16915
211	04/07/2017	210	1802420592	75	CRC	3T	5	9.7	3	100	T2c	3+4	3+4		#DIV/0!
212	25/07/2017	211	2404470132	70	CRC	3T	4	5.9	1	80	T2a	3+4	4+3		#DIV/0!
213	25/07/2017	212	0108530493	64	CRC	3T	5	16.4	7	90	T2b	4+4	4+5		#DIV/0!
214	08/08/2017		1604490071	68			5	23.5	10	90	T2C	3+4	3+4	55.5	0.42342
215	15/08/2017		911580573	58			5	11.5	10	60	T2c	4+4	4+3	50.5	0.22772
216	15/08/2017		2405430290	76			5	8.5	20		T2c	3+4	4+3	32.5	0.26154

217	22/08/2017	1511610433	55			N	7.5	1	20	T2c	3+3	3+3	68	0.11029
218	22/08/2017	108470032	70			5	24.4	12	100	T3b	4+3	3+5	47.9	0.50939
219	29/09/2017	1608530310	64			5	9.8	10	80	T2c	3+4	3+4	29.2	0.33562
220	18/09/2017	2310500070	66			5	15.5	1	5		4+4	4+5	42	0.36905
221	25/09/2017	711540659	62			5	6.2				3+4	4+5		
222	03/10/2017	1602590095	58			5	6.4	5	30	T2c	3+4	3+4	57.8	0.11073
223	29/09/2017	2803490315	68			N	5.1	6	20	T2c	3+4	3+4	125.2	0.04073
224	10/10/2017	402470834	70			5	10	1	10	T2c	3+4	3+4	55	0.18182
225	06/10/2017	3112430158	73			5	9.2	4	20	T3b	4+4	4+5	54.2	0.16974
226	07/11/2017	1708550097	62			4	6.1	3	20	T2c	3+3	3+4	66.9	0.09118
227	04/11/2017	3006420217	75			4	12.7	2	10	T2c	3+4	3+4	182.5	0.06959
228	04/11/2017	2805490177	68			5	19.4	9	70	T2c	4+3	4+3	43.5	0.44598
229	22/11/2017	403540011	63			4	6.4	1	25	T2c	n	3+4	69.8	0.09169
230	14/10/2017	1809440297	73			5	34.9	3	40	T3a	4+4	4+5	70.2	0.49715
231	13/10/2017	509490611	68			4	8	5	80	T3a	3+4	3+4	54	0.14815
232	21/12/2017	2012500595	66			3	5.6	5	15	T2c	3+4	3+4	165.5	0.03384
233	21/12/2017	605520097	65			5	5.9	10	90	T3a	3+3	3+4	110.5	0.05339
234	19/12/2017	2104521270	65			5	6.2	5	40	T2c	4+3	4+3	52	0.11923
235	01/11/2017	1111410119	76			5	6.6	5	40	T3a	4+4	4+3	40.4	0.16337
236	24/01/2018	910630011	63			4	0.1	4	30	T2c	3+4	3+4	65	0.00154
237	18/01/2018	1402500416	68			4	7.5	3	30	T2c	4+3	4+3	83	0.09036
238	23/01/2018	911420096	76			4	10.6	6	40	T2c	3+4	3+4	112	0.09464
239	24/01/2018	2904450351	73			5	13	6	80	T3a	4+5	4+5	123	0.10569
240	26/01/2018	1710460210	72			4	6.8	4	40	T2c	4+5	4+5	36.9	0.18428
241	09/01/2018	1807510395	67			5	5.9	8	40	T2c	3+4	3+4	80.5	0.07329
242	11/01/2018	602450438	73			5	23.4	5	40	T2c	5+4	4+5	109	0.21468
243	16/01/2018	611500299	68			5	8.7	1	10	T3a	4+3	4+3	41	0.2122
244	22/01/2018	1209530112	65			5	12.7	12	90	T3a	4+5	4+5	50.5	0.25149
245	03/01/2018	907610293	57			5	7	3	10	T2c	3+3	3+4	28	0.25
246	06/02/2018	503530077	65			n	7.9	1	20	T2c	4+3	4+3	37.8	0.20899
247	15/02/2018	1706470630	70			5	8.5	1	5	T2c	4+4	4+3	82.4	0.10316
248	20/02/2018	212510037	67			4	9.6	2	10	T2c	3+3	3+4	76	0.12632
249	26/02/2018	803520018	66			5	8	7	80	T2c	3+4	3+4	36.2	0.22099
250	27/02/2018	1508560196	62			5	5.6	4	80	T2c	3+3	3+4	60	0.09333
251	06/03/2018	1001440153	74			4	6.6	3	60	T2c	4+4	4+3	48.2	0.13693
252	14/03/2018	1512500070	68			n	7.1	5	40	T2c	4+3	3+4	16	0.44375
253	15/03/2018	1305540239	64			4	11.1	3	40	T2c	4+3	4+5	41.7	0.26619
254	20/03/2018	1410530272	65			4	4.5	2	10	T2a	3+3	3+4	19.7	0.22843
255	26/03/2018	208520139	66			5	8.5			T2c	3+4	3+4	40.2	0.21144
256	27/03/2018	1401510434	67			n	8.4	3	10	T2c	4+3	3+4	142	0.05915
257	08/04/2018	1505690153	49			n	6.1	5	50	T2c	3+3	3+4	20	0.305
258	15/04/2018	810420015	76			n	7.1	6	30	T2c	4+3	4+3	35	0.20286
259	16/04/2018	1911506250	68			5	5.2	2	50	T2c	3+3	3+4	52	0.1
260	17/04/2018	3004560453	62			5	44	1	5	T2c	3+3	3+4	138.5	0.31769
261	24/04/2018	806505435	68			5	16.9	1	30	T2c	4+3	4+3	67.2	0.25149
262	10/05/2018	202590038	59			4	2.4	3	20	T2c	3+3	4+3	55	0.04364
263	25/05/2018	1001625277	56			4	15.2	5	60	T2c	3+4	3+4	48	0.31667
264	15/05/2018	503460079	72			5	6.9	4	30	T2c	3+4	3+4	38.5	0.17922
265	29/05/2018	1611600073	58			3	6	11	40	T3b	4+3	4+3	26.9	0.22305
266	30/05/2018	2202620230	56			4	11.2	10	90	T2c	3+3	3+4	80.5	0.13913
267	05/06/2018	1105615499	57			5	5.9	9	80	T2c	3+4	3+4	51	0.11569
268	21/06/2018	2512560015	62			5	18	7	90	T3a	4+5	4+5	65.6	0.27439
269	22/06/2018	708480314	70			5	6.7	9	80	T3a	4+5	4+5	49	0.13673
270	18/06/2018	1302480111	70			3	8.6	6	5	T2c	4+3	3+4	52.25	0.16459
271	19/06/2018	2007430096	75			5	14.6	5	40	T3a	4+5	4+5	43	0.33953
272	21/06/2018	2512560015	62			5	18	7	90	T3a	4+5	4+5	65.6	0.27439
273	22/06/2018	708480314	70			4	6.4	9	80	T3a	4+5	4+5	49	0.13061
274	02/07/2018	2805430573	75			5	9.4	2	20	T2c	3+4	3+4	44.5	0.21124
275	10/07/2018	1201550033	63			5	6.9	5	70	T2c	3+4	3+4	36.3	0.19008
276	12/07/2018	1601580118	60			5	6.9	6	90	T2c	3+4	3+4	44	0.15682
277	30/07/2018	209570172	61			5	18.7	11	80	T3b	4+5	4+5	45	0.41556

278	17/07/2018	2612450211	73			5	6.9	2	50	T2c	3+4	3+4	62.5	0.1104
279	20/07/2018	206580339	60			5	6.6	12	80	T3b	4+5	4+5	41	0.16098
280	14/08/2018	1012460533	72			5	10.7	8	90	T3b	5+5	5+4	66	0.16212
281	21/08/2018	2503430090	75			4	3.5	4	60	T2c	4+4	3+4	47	0.07447
282	22/08/2018	403500052	68			5	7	4	70	T2c	4+4	3+4	31	0.22581
283	30/08/2018	203420098	76			4	4.9	10	80	T3a	4+3	3+4	29	0.16897
284	11/09/2018	209460016	72			5	8.2	3	50	T2c	4+3	4+3	76	0.10789
285	10/09/2018	1109460198	72			4	8.5	3	15	T2c	3+3	4+3	43	0.19767
286	13/09/2018	1806450518	73			5	8.6	4	80	T2c	3+3	3+5	82	0.10488
287	18/10/2018	1201450195	70			4	13			T3a	3+3	4+5	39	0.33333
288	18/09/2018	704572273	61			5	41.7	5	100	T3b	4+5	4+5	12	3.475
289	20/09/2018	1611590019	59			no mri	16.4	10	90	T2c	3+4	3+4		
290	19/09/2018	1810440211	74			5	9.5			T3b	3+4	4+4	73	0.13014
291	02/10/2018	3010450079	75			5	27	6	15	T3a	4+4	4+4	85	0.31765
292	05/10/2018	2009440358	74			4	24.7	1	10	T2c	3+4	3+4	55.9	0.44186
293	19/10/2018	607590076	59			5	5.4	4	30	T3a	4+3	4+4	48	0.1125
294	23/10/2018	2002590958	59			5	5.4	3	20	T2a	4+5	4+5	35	0.15429
295	30/10/2018	1710490098	69			4	5.2	3	30	T3a	3+4	3+4	49	0.10612
296	01/11/2018	108420337	76			4	17.1	1	10	T2a	3+4	3+4	39	0.43846
297	06/11/2018	112645658	54			5	12	4	50	T3b	4+5	5+4	34	0.35294
298	13/11/2018	3003450433	73			5	8.8	2	60	T3a	4+3	4+4	68	0.12941
299	14/11/2018	2109420316	75			no mri	7.2	1	25	T3a	4+4	4+4		
300	19/11/2018	606550135	63			5	16	11	80	T3a	4+5	4+5	73	0.21918
301	20/11/2018	1012565556	62			4	5.9	3	10	T2c	3+3	3+4	30	0.19667
302	07/11/2018	2808605714	58			5	6	8	90	T2c	4+4	4+4	60	0.1
303	31/10/2018	604630492	55			n	7.1	1	10	T2a	3+4	3+4		
304	27/11/2018	2312550733	63			clu	6.7	5	90	T3a	4+3	4+3		
305	03/12/2018	2903570310	61			4	8.7	4	10	T2c	4+4	3+4	51	0.17059
306	04/12/2018	1609570014	61			4	5.4	6	100	T2c	3+4	3+4	25	0.216
307	05/12/2018	3103552335	63			5	8.2	2	20	T3a	4+3	4+3	53	0.15472
308	06/12/2018	1203515251	67			4	10	3	80	T3b	3+4	3+4	124	0.08065
309	11/12/2018	1507580118	60			5	18.5	10	90	T2c	3+4	3+4	31	0.59677
310	12/12/2018	806560630	62			5	11.2	3	10	T2c	4+3	3+4	57	0.19649
311	13/12/2018	1111490031	69			4	6.3	7	70	T2a	3+4	3+4	45	0.14
312	17/12/2018	805620435	56			3	6	0	0	T2c	n	3+4	31	0.19355
313	18/12/2018	2202540512	64			4	19.4	10	80	T2c	3+4	3+4	30	0.64667
314	19/12/2019	2106530137	65			5	9.6	4	20	T2c	4+4	4+3	35	0.27429
315	28/12/2018	708450016	73			5	8.2	7	70	T2c	3+3	4+3	34	0.24118
316	04/01/2019	2701456053	73			no mri	11	6	20	T3a	4+4	4+4		
317	08/01/2019	1812420412	76			5	13	4	90	T2c	4+3	3+4	49	0.26531
318	09/01/2019	2105512097	67			5	7.7	12	90	T3b	4+5	4+5	41	0.1878
319	14/01/2019	1106500016	68			4	18	12	90	T3b	4+5	4+5	29	0.62069
320	15/01/2019	803590032	59			5	19	12	90	T3a	4+5	4+5	31	0.6129
321	17/01/2019	2506630139	55			5	9.4	12	90	T3a	4+3	4+3	48	0.19583
322	24/01/2019	1402560117	62			5	9.4	10	70	T2c	4+3	3+4	70	0.13429
323	22/01/2019	1506470416	71			no mri	7.2	1	5	T2c	4+3	4+3		
324	21/01/2019	806559179	63			4	10.5	1	10	T2a	3+4	3+4	117	0.08974
325	29/01/2019	307440095	74			5	6.3	1	25	T2c	3+4	3+4	80	0.07875
326	05/02/2019	2904470158	71			5	5.4	6	60	T2c	3+4	4+3	38	0.14211
327	12/02/2019	206560095	62			5	8.3	12	100	T3b	4+3	4+3	36	0.23056
328	19/02/2019	406460094	72			5	7.9	12	100	T3a	4+5	5+4	45	0.17556
329	25/02/2019	412560038	62			5	7.4	5	50	T2c	3+4	3+4	37	0.2
330	06/03/2019	3107570370	61			5	8.1	1	60	T2c	4+5	4+5	64	0.12656
331	07/03/2019	1702620654	56			3	8.2	3	20	T2c	3+4	3+4	88	0.09318

Raw data 2

Basic information					198	MRI				198	TRUS Biopsy			Fusion biopsy	
No.	Study ID	age*	Date of MRI Scan	PSA (ng/ml)	MRI Result	lesion focus	Highest PIRADS	location	target lesion size (mm)	Biopsy Type	biopsy result	Highest G Grade	Location		Highest G Grade
1	T0001IC	64	17/02/2015	7.9	P	1	5	9p, 10p	14	F	P	3+4	1, 2, 4, 6, 9, 11	P	3+3
						2	3	1a	10						3+3
						3	3	9a	13						
						4	3	3a	20						
4	T0004AW	73	10/03/2015	7.3	P	1	5	1a	16	F	P	3+4	4, 6	P	3+3
						2	5	3p	25						NO
16	T0016RD	69	02/06/2015	11.7	P	1	5	(14,15)a,(10,12)a	25	F	N			P	3+4
17	T0017NM	58	02/06/2015	11.2	P	1	5	5p,6p,7,8,9,10,1	40	F	P	4+3	EXP 2	P	4+3
19	T0019NM	69	16/06/2015	9.7	P	1	5	(7-10)p	15	F	P	4+3	6	P	3+4
						2	5	1p, 2p, 4p, 9p	15						4+3
23	T0023DN	63	30/06/2015	10.8	P	1	5	1p, 2p, 7p	28	F	P	4+3	2-6, 10, 12	P	3+4
						2	5	5p, 11p	15						4+3
						3	5	13as	21						
24	T0024JN	65	30/06/2015	7.7	P	1	5	9p, 10p, 12p	16	F	P	4+4	7	P	5+4
						2	4	6a, 15as	12						NO
						3	4	1p, 7p	32						
27	T0027SF	60	15/07/2015	6.2	P	1	5	6a, 6p, 5p	16	F	N			N	NO
						2	3	1a	25						NO
28	T0028JS	73	15/07/2015	5	P	1	4	11p, 12p	11	F	N			P	3+4
						2	3	7a	19						3+4
34	T0034ED	68	13/08/2015	13.8	P	1	3	1a	20	F	N			N	NO
35	T0035RM	70	18/08/2015	12.4	P	1	5	all	44	F	P	4+4	10	P	4+3
38	T0038IM	70	01/11/2015	12.6	P	1	3	14as	22	F	P	3+3	1	N	NO
39	T0039TM	72	18/11/2015	9.9	P	1	5	10p, 10a	17	F	N			P	3+4
						2	5	13as	16						NO
40	T0040PH	66	18/11/2015	5.9	P	1	4	1a	10	F	P	3+3	7	P	3+3
						2	3	12p	9						3+3
42	T0042DM	70	02/12/2015	8.7	P	1	4	6p, 5p	12	F	P	3+4	2,3,6,7,8,9,10	P	3+4
						2	4	12p	10						NO
						3	3	9a	19						
46	T0046JS	62	22/12/2015	7.3	P	1	4	3a, 5a	13	F	N			P	3+3
						2	4	12p	7						3+4
						3	4	12a, 12p	6						
49	T0049RM	75	13/01/2016	5.7	P	1	5	2a, (4-6)a, 9p	36	F	P	5+4	4, 6	P	4+5
51	T0051LE	74	20/01/2016	6.8	P	1	5	(1-4)a, 2p, 4p	15	F	P	3+4	5,6	P	3+4
						2	4	10p	10						3+3
55	T0055AR	51	10/02/2016	14.5	P	1	4	12p	11	F	N			N	NO
58	T0058JT	59	02/03/2016	9.1	P	1	5	7-12)p, 3p, 5	36	F	P	4+3	7,8,11,12	P	4+5
						2	4	4a, 6a	11						3+3
59	T0059JM	62	09/03/2016	6.6	P	1	5	8a 8p 10a 10p	18	F	P	3+3	5,9,12	P	3+4
						2	5	1p 3p 4p	15						NO
63	T0063GR	69	23/03/2016	14.4	P	1	5	3,7,9)a, (13-15	33	F	N			N	NO
						2	5	9a,12a,12p	15						NO
64	T0064TD	57	30/03/2016	4.6	P	1	5	(7-10)p	18	F	P	4+3	1,3,4,6	P	NO
						2	4	10a	9						4+3
68	T0068IM	57	19/04/2016	4.5	P	1	4	2p, 1p	10	F	P	3+3	7, 9	N	NO
						2	3	10a	9						NO
70	T0070SS	74	04/05/2016	11.3	P	1	5	5a, 5a, 6p, 15a	20	F	N			P	3+3
						2	5	9a	15						3+5
						3	4	10a, 10p	10						
						4	4	12a	10						
71	T0071IC	52	04/05/2016	10.2	P	1	4	2A,2p, 4a,4p	11	F	N			N	NO

73	T0073AD	70	18/05/2016	7.1	P	1	5	a,p	33	F	N			P	3+4
						2	5	p	15						NO
75	T0075JS	74	25/05/2016	8.2	P	1	5	10p,12p	15	F	P	3+3	7, 8	N	NO
78	T0078DO	57	08/06/2016	4.8	P	1	4	4p	10	F	N			N	NO
						2	4	10p	9						NO
80	T0080SS	65	15/06/2016	14	P	1	5	(3-6)p	16	F	P	3+4	6	P	3+4
						2	4	12a	8						NO
83	T0083DF	63	05/10/2016	5.5	P	1	5	3p,9p	15	F	N			P	3+3
						2	4	6p,5p	8						3+3
84	T0084AH	62	29/06/2016	3	P	1	3	3a	10	F	N			N	NO
90	T0090WL	74	20/07/2016	14.4	P	1	4	5p,6p	12	F	N			N	NO
						2	4	6a	10						NO
93	T0093AD	66	09/08/2016	7	P	1	4	15as	12	F	N			N	NO
94	T0094mm	51	10/08/2016	9.1	P	1	3	1a,13as	15	F	P	4+3	9,11	P	NO
						2	3	15as	9						3+3
99	T0099PT	63	07/09/2016	10.2	P	1	5	10p,12a,12p	15	F	N			P	4+3
						2	4	1p,2p,3p,4p	9						4+3
						3	3	1a,7a	18						
101	T0101DM	59	14/09/2016	1.7	P	1	5	10p,8a,8p,11,3p,4a,4p,6p	31	F	N			N	NO
						2	5		19						NO
107	T0107GP	63	12/10/2016	11	P	1	5	13as,7a	30	F	N			P	3+3
						2	4	9p,10p	19						NO
112	T0112JR	57	09//11/2016	4.4	P	1	4	1A	10	F	N			N	NO
						2	4	11A	7						NO
						3	3	9A	12						
114	T0114DK	68	16/11/2016	2.2	P	1	5	12P	16	F	N			N	NO
116	T0116SB	61	23/11/2016	8.6	P	1	5	3P,4P	20	F	P	4+4	ALL	P	4+4
						2	4	10P,9P	18						NO
						3	4	3P	8						
						4	4	3P,9P	6						
118	T0118JW	70	02/12/2016	9.9	P	1	5	3P	13	F	P	4+4	3 7 8 11	P	4+4
120	T0120ER	71	07/12/2016	10.1	P	1	3	11p, 12p	17	F	N			N	NO
123	T0123BA	55	15/12/2016	6.1	P	1	5	13as, 14as	24	F	P	3+3	1, 9, 10	N	NO
						2	4	12p	8						NO
124	T0124RW	68	21/12/2016	7.6	P	1	3	7a	20	F	N			N	NO
129	T0129JC	69	11/01/2017	5.4	P	1	4	10a, 10p	10	F	N			N	NO
131	T0131FR	64	18/01/2017	12.1	P	1	5	a,3a,4a,13as,1	22	F	P	3+3	9 10	P	4+3
						2	4	10p	10						NO
133	T0133PM	69	25/01/2017	11.7	P	1	5	10p,9p,12p	16	F	P	4+5	10 11 12	P	4+4
						2	4	4a	13						
						3	4	10a	7						
135	T0135ID	71	01/02/2017	4.7	P	1	3	m4	7	F	N			N	NO
						1	4	m4	27						NO
138	T0138BF	66	15/02/2017	12.2	P	2	4	a3	5	F	N			N	NO
						3	4	a4	4						
140	T0140RM	70	22/02/2017	9.7	P	1	5	m2	17	F	P	3+4	9	P	NO
						2	4	a3	9						3+4
142	T0142IW	60	01/03/2017	7.7	P	1	3	m2	14	F	N			N	NO
145	T0145GC	61	08/03/2017	14.4	P	1	4	m3	11	F	P	4+3	3	P	NO
						2	3	a4	10						3+4
149	T0149RC	67	29/03/2017	8.5	P	1	5	b1,b2,m1	25	F	N			N	NO
150	T0150AM	63	29/03/2017	16.4	P	1	5	b4,m2,m4,a2	30	F	P	4+3	8 9 10 11 12	P	4+4
154	T0154NJ	64	12/04/2017	9.3	P	1	5	b1,m1,b3,m3	32	F	P	3+4	10	P	4+5
						2	5	m2	22						3+3
159	T0159NM	67	26/04/2017	9.1	P	1	4	m1,m3	21	F	N			N	NO

162	T0162RH	73	10/05/2017	9.3	P	1	4	m3	16	F	P	4+3	1 2 5	N	NO
165	T0165HH	68	17/05/2017	8.3	P	1	5	m3,m4	20	F	N			N	NO
167	T0167JS	73	24/05/2017	8.2	P	1	4	a3	11	F	P	3+4	2 3 4 6	P	NO
169	T169ND	58	01/06/2017	6.5	P	2	4	a4	10	F	P	3+4	3	P	3+4
						1	5	b2,m1,m2,a1	35						
						2	5	a3	22						
						3	4	a4	13						
170	T0170LB	62	01/06/2017	16.4	P	1	5	b4,m1,m2,m3	39	F	P	5+4	12	P	4+4
						2	5	m2	15						
173	T0173CS	64	14/06/2017	6.3	P	1	3	m2	11	F	N			P	3+3
174	T0174FC	67	14/06/2017	19.6	P	1	5	m4,a4	21	F	P	4+3	8,9,10,11	P	3+4
						2	5	b3,m3	16						
179	T0179DB	69	29/06/2017	6.1	P	1	3	m3,m4,a3,a4	37	F	N			N	NO
181	T0181SM	50	05/07/2017	10.8	P	1	5	b3,b4,m2,m3	30	F	P	5+4	5, 7...11	P	4+5
						2	4	a4	5						
184	T0184AG	66	02/08/2017	6.3	P	1	3	m1,m3	16	F	N			N	NO
187	T0187GE	69	16/08/2017	5.4	P	1	5	1,b2,m1,m3,a	22	F	P	4+5	7	P	4+5
						2	4	a4	9						
188	T0188JM	74	16/08/2017	9.3	P	1	5	b2,b4,m2,m4	22	F	N			N	NO
191	T0191AT	71	23/08/2017	18.3	P	1	5	b4,m3,m4,a3	36	F	P	4+5	3 5 11	P	4+5
193	T0193RH	73	30/08/2017	6.7	P	1	4	a4	14	F	P	4+5	7 9 10	N	NO
						2	4	a4	4						
196	T0196DH	67	13/09/2017	6.9	P	1	4	m4	13	F	P	4+3	11	P	3+4
						2	4	a3,a4	12						
204	T0204BA	55	04/10/2017	6.5	P	1	3	m4	15	F	N			N	NO
207	T0207SP	60	20/10/2017	7.2	P	1	4	m4	12	F	P	3+3	12	P	3+3
210	T0210AR	70	01/11/2017	8.5	P	1	5	b1,b2	18	F	N			P	4+4
						2	3	m4	13						
213	T0213JS	61	08/11/2017	12.5	P	1	3	m2	15	F	N			N	NO
215	T0215PN	64	15/11/2017	7	P	1	4	m3	13	F	N			P	4+4
						2	3	m1	10						
219	T0219IM	75	29/11/2017	3.3	P	1	5	b4,m2,m4,a2	28	F	P	4+5	11	P	3+4
223	T0223RR	71	13/12/2017	6.5	P	1	5	b4	15	F	P	3+4		P	3+3
226	T0226HA	57	03/01/2018	6	P	1	3	a4	11	F	P	4+3	8,12	P	3+4
228	T0228TM	67	10/01/2018	10.2	P	1	5	m2,m4	19	F	P	3+3	7,8	N	NO
						2	4	m4	13						
						3	4	b3	11						
231	T0231SD	65	24/01/2018	6.9	P	1	5	b1,b3	30	F	P	3+4	3,5,6	N	NO
234	T0234RP	67	31/01/2018	6.7	P	1	5	b4	30	F	P	4+5	7-9,11	P	4+5
237	T0237NS	71	14/02/2018	10.6	P	1	5	m3,a3	15	F	P	3+4	12	P	3+4
						2	4	a4	12						
						3	4	m1	8						
						4	4	b4	6						
						5	4	m2	5						
240	T0240JP	74	21/02/2018	10.5	P	1	5	b4,m4	16	F	P	3+4	3	P	3+3
						2	4	m2	6						
241	T0241TJ	71	28/02/2018	14.7	P	1	5	b2,b4,m2,m4	16	F	P	4+3	7,8,9,19	P	3+4
243	T0243WM	56	07/03/2018	11.4	P	1	4	b4,m4	12	F	P	3+3	ALL	N	NO
248	T0248GK	68	21/03/2018	9.3	P	1	3	b3	10	F	P	3+4		P	3+3
249	T0249RR	69	28/03/2018	6.7	P	1	5	b1,b3,m1,m3	18	F	P	4+5	3,6	P	4+3
						2	4	m2	8						
253	T0253JK	73	09/04/2018	13.4	P	1	4	m2	14	F	N			P	3+4
255	T0255TC	61	11/04/2018	18.9	P	1	5	4,m1,m2,m3	45	F	P	4+4		P	4+5
257	T0257SP	52	18/04/2018	6.7	P	1	4	m1,m2	11	F	P	3+3	1,3	N	NO
260	T0260PH	72	30/04/2018	10.3	P	1	5	m1,a1	18	F	N			N	NO



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Prediction of prostate cancer Gleason score upgrading from biopsy to radical prostatectomy using pre-biopsy multiparametric MRI PIRADS scoring system

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An increase or 'upgrade' in Gleason Score (GS) in prostate cancer following Transrectal Ultrasound (TRUS) guided biopsies remains a significant challenge to overcome. To evaluate whether MRI has the potential to narrow the discrepancy of histopathological grades between biopsy and radical prostatectomy, three hundred and thirty men treated consecutively by laparoscopic radical prostatectomy (LRP) between July 2014 and January 2019 with localized prostate cancer were included in this study. Independent radiologists and pathologists assessed the MRI and histopathology of the biopsies and prostatectomy specimens respectively. A multivariate model was constructed using logistic regression analysis to assess the ability of MRI to predict upgrading in biopsy GS in a nomogram. A decision-analysis curve was constructed assessing impact of nomogram using different thresholds for probabilities of upgrading. PIRADS scores were obtained from MRI scans in all the included cases. In a multivariate analysis, the PIRADS v2.0 score significantly improved prediction ability of MRI scans for upgrading of biopsy GS ($p = 0.001$, 95% CI [0.06–0.034]), which improved the C-index of predictive nomogram significantly (0.90 vs. 0.64, $p < 0.05$). PIRADS v2.0 score was an independent predictor of postoperative GS upgrading and this should be taken into consideration while offering treatment options to men with localized prostate cancer.

Histology from biopsies categorised into Gleason score is the only confirmatory test for cancer diagnosis and is most commonly used for risk stratification of men with a recent diagnosis of prostate cancer. Based on this men are counselled for various treatment options. MR imaging data is not considered in risk stratification at present. With increasing therapeutic options available to men with a diagnosis of prostate cancer, scrutiny of information from biopsy grade becomes increasingly important. There is around 35.5% (range: 14–51%) upgrading of biopsy GS on LRP¹. Many factors contribute to the discrepancy between needle biopsy and corresponding radical surgery GS. Under calling of Gleason cribriform Gleason pattern 4 as pattern 3 or the presence of borderline grades due to barely appreciable glandular differentiation under microscope and lack of sampling of tertiary grade disease on biopsies are known contributors. Factors such as age, size of prostate, extent of cancer on biopsy needle and number of biopsy samples (extended/ or mapping) have also been known to impact on the incidence of upgrading².

In light of a number of studies reporting upgrading or under-grading of prostate cancer on needle biopsies, there is the potential for under treatment or overtreatment (i.e. radiotherapy and hormone duration). Several publications^{3,4} and consensus updates on the Gleason grading system have partially addressed this issue including recommendation of deriving GS by adding the most common and highest Gleason pattern on biopsy rather than

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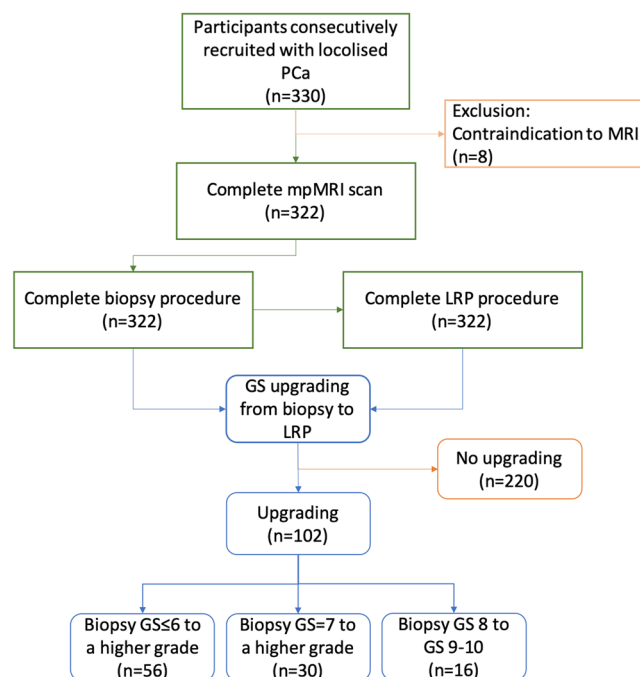


Figure 1. Flowchart of the study.

original method of adding the primary and second most common patterns⁵. Moreover, upgrading if suspected, has long-term outcome implications. Corcoran *et al.*⁶ have shown that even after adjusting for known preoperative variables (including clinical stage, prostate-specific antigen (PSA), number of positive cores and percentage of positive cores) upgrade to a higher Gleason Score (GS) remained a strong and independent predictor of biochemical recurrence after attempted local curative therapy, this underscores the importance of gaining more information to predict upgrading of biopsy GS in men diagnosed with prostate cancer as this may serve as a marker of biologically aggressive disease.

Pre-biopsy MRI has recently been shown to hold great promise in the detection and characterisation of prostate cancer⁷. A negative scan (no lesion seen on the MRI scan) showed a high negative predictive value for the presence of significant prostate cancer⁸. Song *et al.*⁹ reported a high predictive value of PIRADS v2 in predicting upgrading of GS from biopsy, however this study was retrospective and MR Imaging was obtained at least 3 weeks following biopsies - an approach known to impact interpretation of images. Post-biopsy haemorrhage is the most common false-positive finding for prostate cancer¹⁰. In this study, there was no attempt to align histopathological sectioning to MRI using recently reported 3D-mould technology. Therefore, this is the first report describing predictive accuracy of pre-biopsy MRI in upgrading biopsy GS following LRP using patient-specific 3D moulds to ensure permitted alignment of excised prostates with MRI scans.

Patients and methods

Study population. This is a study with prior Caldicott institutional approval (Caldicott/IGTCAL5626). All experiments including the study protocol study followed approved institutional guidelines. The study had ethical approval (14/ES/1070) with each participant informed consenting to the use of their imaging data. Between July 2014 and January 2019, 330 men consecutively treated by LRP who were diagnosed with localised prostate cancer with raised PSA or/and abnormal digital rectal examination were included in this study. They were offered mpMRI and those with positive MRI results (PIRADS score 3 and above) were performed transrectal ultrasound (TRUS)-guided prostate biopsy (12 cores) followed. Of these patients, eight were excluded because of contraindication to MRI such as a heart pacemaker and metallic foreign body including three claustrophobic patients. Further analysis included remaining 322 patients (Fig. 1). The clinical, pathological and imaging factors information of the patients, including age, weight, preoperative PSA, PSA density, number of positive cores, maximum percentage of cancer per core and PIRADS v2 score on multiparametric MRI (mpMRI) were recorded. GS upgrading defined as a biopsy GS increasing from lower to higher grade on reported before². Table 1 summarises the baseline characteristics between upgraded and non-upgraded groups of the cohort.

Hypotheses of the study. We hypothesised that pre-biopsy MRI with PIRADS classification of suspicious area in prostatic cancer improve prediction of GS upgrading from biopsy to radical prostatectomy. Upgrading of GS on histology was defined as change of GS from lower to higher grade between biopsy and histology from radical prostatectomy.

MRI protocol and PIRADS score. All patients' mpMRI scans were performed on 3 T scanner (TIM Trio, Siemens, Erlangen, Germany) 2 weeks before TRUS-guided biopsies. The mpMRI protocol was derived from the European Society of Uro-radiology Guidelines 2012 for the detection of prostate cancer and the subsequent

	Total	Upgrading	No upgrading
Number of patients	322	102	220
Age (y), mean \pm SD (range)	66.83 \pm 5.9(44–77)	66.82 \pm 6.12(49–77)	66.85 \pm 5.9(44–77)
Prostate Weight mean \pm SD (range)	63.7 \pm 30.13(12–207)	65.3 \pm 26.2(20–155)	63.1 \pm 31.78(12–207)
PSA level (ng/ml), mean \pm SD (range)	11.1 \pm 7.39(0.1–47.7)	12.6 \pm 9.98 (2–47.7)	10.39 \pm 5.7(0.1–41)
PSA Density (ng/ml ²), mean \pm SD (range)	0.261 \pm 0.234 (0.001–3.48)	0.212 \pm 0.183(0.035–1.11)	0.203 \pm 0.254(0.00198–3.48)
Number of positive cores	4.8 \pm 3.4(1–12)	4.1 \pm 3.07(1–12)	5.08 \pm 3.42(1–12)
Maximum percentage of cancer per core	50.2 \pm 30.4(5–100)	42.5 \pm 30.6(5–100)	53.3 \pm 29.52(5–100)
PIRADS from mpMRI Benign (1,2)	17 (5%)	4 (4%)	13 (6%)
PIRADS 3	21 (7%)	6 (6%)	15(7%)
PIRADS 4	78 (24%)	26 (26%)	52 (23%)
PIRADS 5	206 (64%)	66 (65%)	140 (64%)

Table 1. Patient characteristics.

	T1WI		High resolution T2WI			DWI	DCE
	Axial	Sagittal	Axial	Coronal	DWI	DWI high b-value	Dyn Gd-MRI
TR (ms)	650	6000	4000	5000	3300	3300	4.76
Sequence	2DTSE	2DTSE	2DTSE	2DTSE	2DEPI	2DEPI	3D VIBE
TE (ms)	11	102	100	100	95	95	2.45
Flip angle (°)	150	140	150	150	—	—	10
Slice thickness (mm)	3	3	3	3	3	3	3
Slice gap (mm)	0.6	0.6	0.6	0.6	0	0	0.6
Resolution (pixels)	320	320	320	320	192	192	192
FOV (mm)	200	200	200	200	280	280	280
b-values (s/mm ²)	—	—	—	—	50,100,500,1000	2000	—
Temporal resolution (s)	—	—	—	—	—	—	4

Table 2. MRI acquisition parameters.

publication of version 2¹¹. Table 2 briefly summarizes the MRI acquisition parameters. Localiser images were acquired in all three imaging planes, whereby the plane of the prostate was defined in relation to the rectal wall.

The mpMRI images were analysed and scored by experienced uro-radiologists using PIRADS v2.0; and the radiologists were blinded to all patients' pathology results. PIRADS v2.0 assessment categories were described as follows: score 1, clinically significant cancer is highly unlikely to be present; score 2, clinically significant cancer is unlikely to be present; score 3, the presence of clinically significant cancer is equivocal; score 4, clinically significant cancer is likely to be present; and score 5, clinically significant cancer is highly likely to be present (Fig. 2).

Histopathology data and analysis. The biopsy results were analysed by experienced pathologists; who were blinded to MRI findings. The GS for each patient was obtained. The radical prostate specimens for histology were sliced in patient-specific moulds to aid orientations between imaging and histology, which were fabricated using a 3D printer as described by our group and others previously^{12,13}. Specifically, patient specific 3D printed moulds were made prior to surgery based on the T2-weighted MRI prostate capsule the moulds were customised for each patient using MIMICS and Solidworks. Moulds were printed at 200 micro resolution using a consumer grade 3D printer (MakerBot Replicator 5th generation). The average mould required 120 minutes to design, 4 to 7 hours to print and an expense for materials of less than \$7¹².

Statistical analysis. Baseline characteristics of patients and pathological outcomes were compared using a chi-square test for categorical data (PIRADS score) and a Student t-test or ANOVA for continuous data (age, weight, PSA level, PSAD, number of positive cores and maximum percentage of cancer per core). Univariate logistic regression was applied to investigate the association of clinical variables with the upgrading of biopsy GS. Variables with $P < 0.05$ in the univariate analysis were further assessed using a multivariate logistic regression analysis to identify factors predictive of GS upgrading. In order to estimate the area under the curve (AUC) for predicting GS upgrading to determine the diagnostic performance of clinical variables with or without PIRADS score, a receiver operating characteristic (ROC) curve analysis was conducted.

In addition, logistic regression model coefficients were used to perform a nomogram predicating the probability of GS upgrading. Non-informative or non-significant variables in univariate logistic regression for GS upgrading were removed. The value of concordance indexes (c-index) were calculated and compared. The bias-corrected calibrated values were generated from internal validation based on 200 bootstrap resamples.

A decision-analysis curve was constructed assessing impact of nomogram using different thresholds probabilities of upgrading (none of the GS upgrade to all GS upgrade).

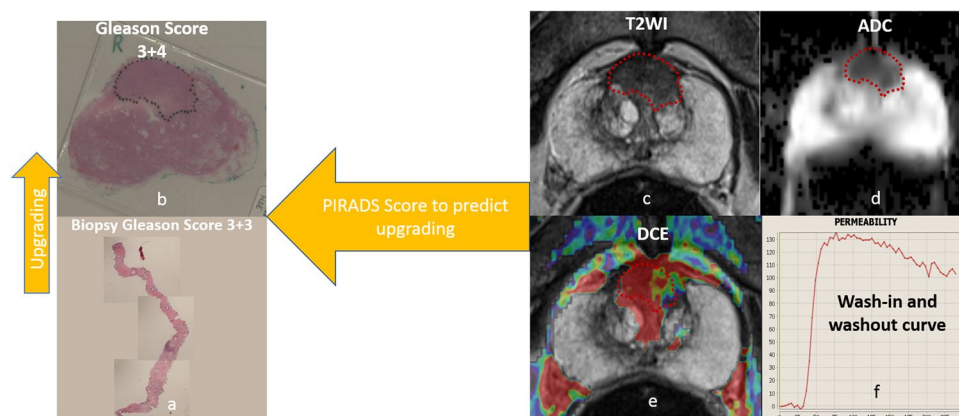


Figure 2. (a) A 73 year-old man with Gleason score 6 disease on prostate cancer on TRUS-Guided biopsies. (b) The grade was upgrading to GS 7 on whole mount radical prostatectomy specimen (c) Axial T2-wighted image shows ill-defined homogeneous low-signal-intensity on the central zone (d), Apparent diffusion coefficient (ADC) shows restricted diffusion in low-signal mass and (e) dynamic contrast enhanced (DCE) shows fast and strong enhancement and early contrast agent washout (type 3 curve), (e,f) The lesion was scored as PIRADSV2 5 (>1.5 cm) and based on parameters described here.

Biopsy Gleason sum	Radical prostatectomy Gleason sum				Total
	6	7	8	9–10	
1–5	0	3	0	1	4
6	6	46	2	4	58
7	0	145	14	16	175
8	0	19	9	16	44
9–10	0	2	3	36	41
Total	6	215	28	73	322

Table 3. Comparison between biopsy and radical prostatectomy Gleason score sum.

Analyses were performed using SPSS 22 (IBM Corporation, New York, US) and R software. The alpha level was set at 0.05 to determine two-tailed significance.

Results

Upgrading cohort characteristics. In total, 322 men were included in our study. Table 3 shows the concordance between the biopsy and prostatectomy Gleason score sums. Of these, (102/322; 31.6%) had GS upgrading from biopsy to LRP. Almost half of this upgrading was from biopsy GS ≤ 6 disease (56/102; 55%). More than half of whole cohort (175/322; 54%) had a GS 7 on prostate biopsy and (30/175; 17%) men had GS upgrading. Finally, eighty five of the cohort (85/322; 26%) had a GS ≥ 8 on prostate biopsy and (16/85; 18.8%) men had GS upgrading from GS 8 on prostate biopsy to GS > 8 at LRP.

The correlation between PIRADS score and pathologic GS at LRP is demonstrated in Fig. 3. Of the 322 patients, the distribution of PIRADS score was as follows: score 1 and 2 in 17 (17/322; 5%) patients, score 3 in 21 (21/322; 7%) patients, score 4 in 78 (78/322; 24%) patients, and score 5 in 206 (206/322; 64%) patients.

Predictions of GS upgrading. Table 4 shows the outcomes of the logistic regression analysis and predictive variables of GS upgrading. On univariate analyses, increased preoperative PSA levels, number of positive cores, maximum percentage of cancer per core and PIRADS ≥ 4 were all significantly associated with GS upgrading ($p < 0.05$). Age, weight of prostate and PSAD did not show any significance ($p > 0.05$) which were excluded from further analyses. In the multivariate analyses, PIRADS ≥ 4 and higher PSA level were both statistically significant and independently predictive of GS upgrading ($p = 0.001$ and 0.003 , respectively).

In Fig. 4, PIRADS v2 score with PSA value show a higher accuracy than PSA alone for predicting GS upgrading (AUC = 0.90 and 0.64, respectively, $p < 0.001$).

Impact of PIRADS score on prediction of GS upgrading in relation to other factors. Figure 5a₁,a₂ show the nomograms constructed for upgradation of biopsy GS with and without PIRADS v2 score data. Longer scales indicated higher percentage of impact and larger points were suggesting probability of upgrading. PIRADS score had the greatest impact followed by PSA level. C-index of the established nomogram which had PIRADS v2 score variable to predict the GS upgrading in the cohort was significantly higher than that of the nomogram

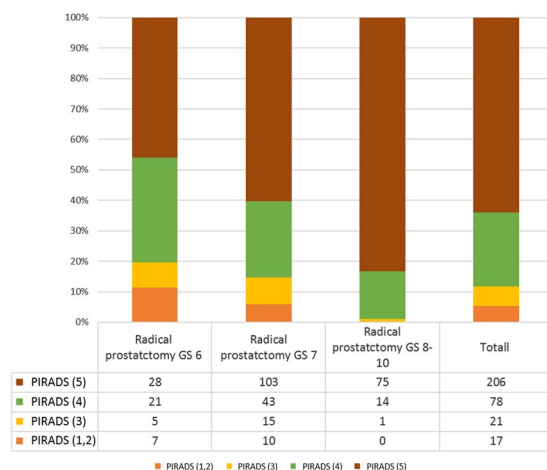


Figure 3. Radical prostatectomy Gleason score stratified according to PIRADS score.

	Univariate		Multivariate	
	OR (95% CI)	P-value	OR (95% CI)	P-value
Age	1.005 (0.962–1.041)	0.799		
Pathology weight	1.002(0.931–1.009)	0.540		
Number of positive cores	0.86(0.87–0.96)	0.005	0.970(0.98–1.01)	0.69
PSA level (ng/ml)	1.040 (1.009–1.073)	0.001	1.09(1.030–1.160)	0.003
PSA Density (ng/ml ²)	1.15 (0.44–3.04)	0.76		
Maximum percentage of cancer per core	0.988 (0.980–0.996)	0.002	0.970 (0.84–1.12)	0.62
PIRADS				
≤ 3	1 (reference)	—	1 (reference)	—
> 3	0.017(0.08–0.04)	0.001	0.014 (0.06–0.034)	0.001

Table 4. Univariate and multivariate logistic regression analysis.

without PIRADS score (0.90 [95% CI 0.87–0.89] vs. 0.64 [95% CI, 0.57–0.70], $p = 0.001$). The nomograms were then validated using 200 bootstrap samples, internal calibration curves are shown in Fig. 5b₁,b₂.

Decision curve analysis. The decision analysis curve is shown in Fig. 6. The net benefit for the model using PIRADS score was significantly higher at all thresholds compared with the model without PIRADS score. As seen in Fig. 6, the decision curve line (depicted by a red line) of the model without the PIRADS scores remained close to the line with threshold probabilities ranging from 0 to 0.25. In contrast, a higher positive net benefit was obtained in the range of threshold probabilities ranging from 0.05 to 1.0 in the model with PIRADS scores.

Discussion

Principal findings of the study. This is the first study to bring together information of PIRADS scores in pre-biopsy MRI and an improved image oriented histopathological grossing of RP specimen by making the use of the mould, bridging the gap in the existing literature. Our results indicate a significant advantage (C-index 0.90 vs 0.64) of using the prediction model including PIRADS scores added to conventional clinic-pathological characteristics (PSA level, percentage of cancer on core-biopsies, gland size etc.) of men with prostate cancer confirmed by systematic transrectal random biopsies relative to a model without PIRADS scores. Prostate biopsy GS upgrading remains a challenge for physicians managing localised prostate cancer, as better knowledge of contributing factors and how to narrow the gap is lacking². To inform any consensus, we need an improved understanding of the role imaging can play, in particular pre-biopsy MRI, in predicting GS change and adverse downstream oncological outcomes. Although, recent improvements have been made in refining biopsy strategies and in reducing sampling errors, a significant and continued effort is still needed to identify men at risk of GS upgrading.

Study findings in context of the reported literature. Wang JY *et al.*¹⁴ reported a nomogram with C-index of 0.795 using preoperative factors without imaging data in a non-screened population from China. This is similar to our study as the healthcare system for the cohort reported here did not have men screened for prostate cancer. Table 5 shows the predictive ability of various reported nomograms in upgrading of biopsy GS of prostate cancer in screened populations^{1,14–17}. The upgrading rate of biopsy GS seen in our cohort is similar to a larger cohort of 2982 patients reported previously¹⁵. A higher percentage of men with a GS of 6 were upgraded in the present study. It is interesting that despite the higher number of cores obtained in the present study (12 in

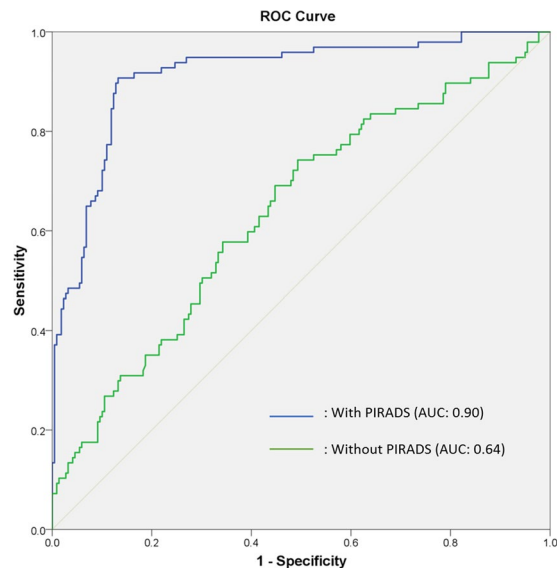


Figure 4. Receiver operating characteristic (ROC) curve of the clinical variables with and without PIRADS score.

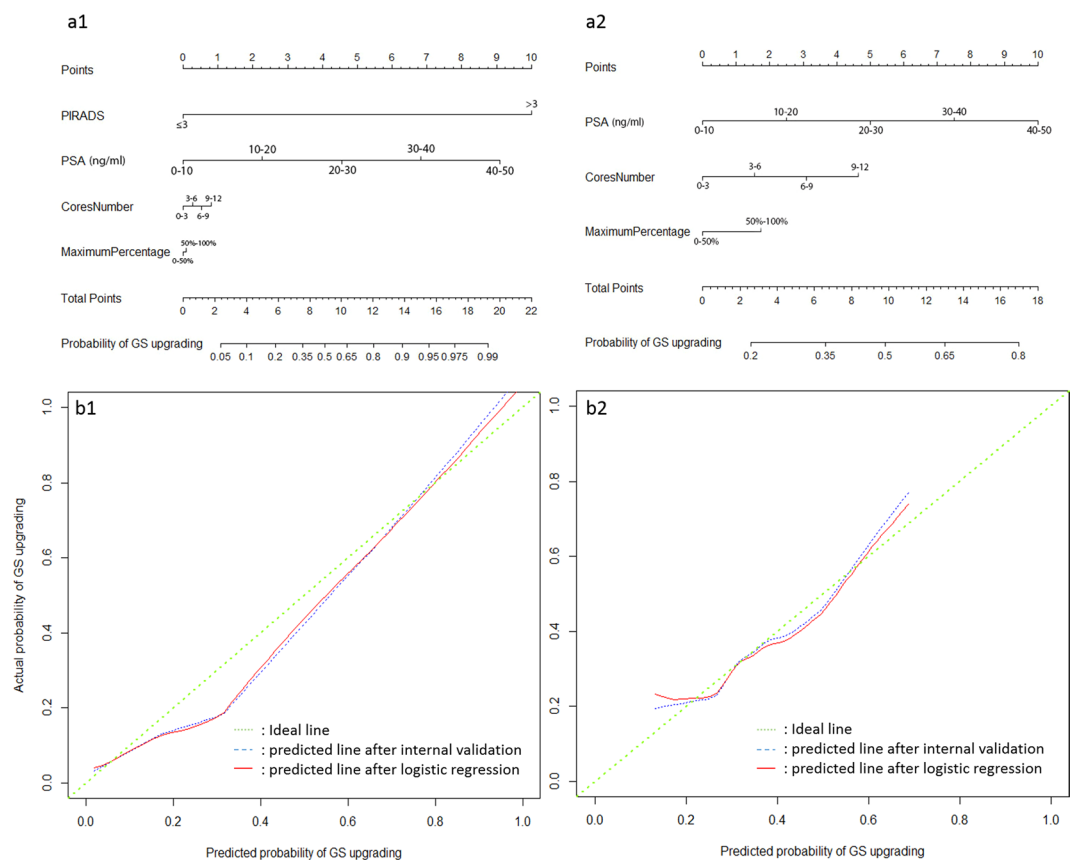


Figure 5. The nomograms of Gleason score upgrading prediction with (a1) and without PIRADS score (a2). Calibration plots of observed and predicted probability of GS upgrading with (b1) and without PIRADS score (b2).

number) in comparison with 6–10 biopsy cores obtained in study by Chun FK *et al.*¹⁵; upgrading rates remain comparable.

A number of previous studies have carried out multivariate analyses of factors responsible for upgrading of biopsy GS including construction of nomograms (Table 5). In predictive oncology, nomograms have huge

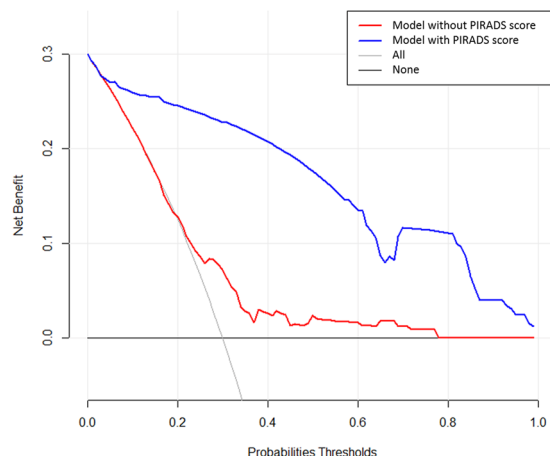


Figure 6. Decision analysis demonstrated a high net benefit of PIRADS score model across a wide range of threshold probabilities. Prediction model without PIRADS score (red line); prediction model with PIRADS score (blue line).

Author	year	Number of patients	Performance (C-index)	Significant parameters on multivariate analysis
Chun, FK	2006	2982	0.804	PSA level, clinical stage and primary and secondary GS
Kulkarni, GS	2007	175	0.71	PSA level and the level of pathologist expertise
Budäus, L	2010	414	0.708	PSA level, clinical stage, prostate volume and percent of positive cores
Wang, JY	2014	220	0.789	PSA level, clinical stage, and primary and secondary GS
Biming, He	2016	411	0.753	Primary and secondary GS and obesity
This study	2019	322	0.90	PSA level and PIRADS score on mp-MRI

Table 5. literature review and comparison between previous and current studies.

potential to help clinicians determine the risk of disease progression and identify those who would experience a greater benefit from multimodality therapy. This approach may result in avoiding unnecessary treatment and improve quality of life by reducing side effects of therapy through better and more precise approach. However, a careful approach is needed to construct nomogram based on specific question, the study population, the method of construction, and its ability to apply to a particular clinical situation. We have followed guidelines described in previous publications^{18–20} in constructing nomogram in this study including selection of variables and statistical methods. The nomogram in the present study has been internally validated (Cross-validation and bootstrapping). External validation of nomogram was not carried out in the present study as this would require further prospective multi-centre recruitment of a cohort. Since D'Amico pioneered this approach²¹, none of the reported predictive nomograms have included imaging features of the disease. Furthermore, the advantage of our nomogram is a higher overall accuracy (discriminant properties) and closer agreement between predicted and observed values (superior calibration). Estimating clinical utility of nomograms in prognosticating an outcome of intervention remains core value of translational research in precision medicine. Vickers and Elkin²² have introduced decision analysis curves estimating probabilities of benefits and harms that a diagnostic test or intervention can trigger at various thresholds. Addition of PIRADS score to nomogram achieved a higher net benefit of decisions making in comparison to leaving out PIRADS score as shown in the decision analysis curve constructed in the present study. The thresholds ranged from no upgrading of disease to all men having upgrading of disease following LRP.

Clinical implications of the study findings. Predicting final histopathological Gleason score of prostate cancer remains a highly desirable information for physicians counselling men with localised prostate cancer for various modalities of treatment and long-term disease recurrence. At present, various nomograms are used mainly taking into consideration clinical factors such as age, pre-operative PSA level and number of biopsy cores involved with the cancer. Notwithstanding this, there is still a large histopathological discrepancy between biopsy and final radical prostatectomy Gleason score. The present study reports a nomogram based on pre-biopsy multiparametric MRI grade (PIRADS score) of cancer alongwith other known clinical parameters. The nomogram clearly showed an improved prediction of final Gleason score and the findings have a large implications for clinicians and researchers in this area. We envisage that this and further research should take us close to precise prediction of final Gleason score of histopathology in prostate cancer and thereby an improved and informed decision making by stakeholders including patients in the management of localised prostate cancer. This will have huge benefits for improved GS prediction for men opting for active surveillance and focal therapy besides those opting for radical prostatectomy.

Limitation of the study. There are limitations to our study. This is a single centre study with dedicated uro-radiologist and pathologist, and the rate of upgrading may be different in small centres. Moreover, overall accuracy of our model, although higher than previous was not perfect (90%). Additionally, performance of our model needs further validation in an external data set. Finally, the accuracy of our model could still be improved by integrating additional predictor variables, such as the novel genomic and other biomarkers^{23,24}. The growing field of artificial intelligence and machine learning using radiomics approach may improve our ability to define tumour characteristics and classification. This, undoubtedly may impact results of the study in the future. Finally, with emerging evidence supporting MRI facilitated biopsy targeting of suspicious areas using ultrasound, the rate of upgrading and the future implications for the practice may change²⁵. There is an emerging evidence that targeted biopsy may improve our ability to narrow the upgrading gap between the biopsies and radical prostatectomy histology^{26,27}; however no predictive nomogram information was available from both the studies. Our ongoing work through randomised intervention in MR/US fusion should be able to provide more information²⁸.

In Conclusions, PIRADS version 2 score of 4 or 5 are associated with an increased risk of biopsy Gleason Score upgrading. Pre-biopsy MRI and PIRADS score significantly and independently predict GS upgrading. If proven by external validation, this information should help in decision making by offering treatment options to men with localised prostate cancer.

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Author contributions

All authors had input into the manuscript and have approved the manuscript for publication. Conceptualization, G.N. and C.W.; methodology, C.W.; validation, M.S.B., J.W. and Z.H.; formal analysis, Y.Z.; investigation, C.W.; resources, S.A., M.S.B., J.W.; writing—original draft preparation, S.A.; writing—review and editing, C.W. Z.H., G.N.; visualization, S.A.; supervision, G.N.

Competing interests

The authors declare no competing interests.

Additional information

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Article

Predicting the Performance of Concurrent Systematic Random Biopsies during Image Fusion Targeted Sampling of Multi-Parametric MRI Detected Prostate Cancer. A Prospective Study (PRESET Study)

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Simple Summary: The study provides a predictive model by using clinical factors in selecting men who may benefit from the addition of systematic biopsies with an image fusion targeted approach. The approach is likely to improve the detection of csPCa and avoid unnecessary detection of indolent prostate cancers.

Abstract: The study was aimed to develop a predictive model to identify patients who may benefit from performing systematic random biopsies (SB) in addition to targeted biopsies (TB) in men suspected of having prostate cancer. A total of 198 patients with positive pre-biopsy MRI findings and who had undergone both TB and SB were prospectively recruited into this study. The primary outcome was detection rates of clinically significant prostate cancer (csPCa) in SB and TB approaches. The secondary outcome was net clinical benefits of SB in addition to TB. A logistic regression model and nomogram construction were used to perform a multivariate analysis. The detection rate of csPCa using SB was 51.0% (101/198) compared to a rate of 56.1% (111/198) for TB, using a patient-based biopsy approach. The detection rate of csPCa was higher using a combined biopsy (64.6%; 128/198) in comparison to TB (56.1%; 111/198) alone. This was statistically significant ($p < 0.001$). Age, PSA density and PIRADS score significantly predicted the detection of csPCa by SB in addition to TB. A nomogram based on the model showed good discriminative ability (C-index; 78%). The decision analysis curve confirmed a higher net clinical benefit at an acceptable threshold.

Keywords: prostate cancer; magnetic resonance imaging; prostatectomy; systematic random biopsy; targeted biopsy

1. Introduction

Recent trends and evidence support pre-biopsy MRI with selective targeting of suspected malignant lesions using MRI/ultrasound (US) and TB methods [1]. The burgeoning interest in fusion imaging has arisen to address the main limitations of SB: over detection of clinically insignificant cancers and possibly under detection of csPCa. A number of recent

reports support the utility of pre-biopsy multi-parametric magnetic resonance imaging (mpMRI) to address the limitations of SB, and the advantage of increased csPCa detection [2–4]. Pre-biopsy MRI in MRI directed pathways have been reported to detect a higher number of csPCa. However, benefits of image-guided targeting performed in combination with systematic sampling or alone remains poorly defined. Drost et al., in a recent systematic review, used a mixed population (with or without image fusion targeting of suspicious areas) to answer this question; however, image fusion methods were not used in all the cases of included studies, and hence, the benefits of targeting suspicious areas with the image fusion approach, with or without addition of systematic biopsy sampling, remains unclear [5].

Image fusion approach makes use of information from MRI to direct biopsy needles under real-time US guidance [2,6,7]. Studies have shown that mpMRI combined with TB technology is a promising tool in the diagnosis of PCa [2,8,9]. In light of a number of previous trials showing the significant benefits of image TB, research is now focused on whether random biopsies are required at all in the detection of prostate cancer [10–12]. This question is pertinent to settle an ongoing debate as studies have also highlighted that TB with the addition of systematic random biopsy is superior to systematic random biopsy alone either in terms of capturing csPCa or even in terms of post-procedural morbidity [1]. In a large retrospective study, from centres in Europe and the USA, Dell'Oglio et al. aimed at findings a group of men where systematic biopsies could be avoided altogether in men with MRI-facilitated targeted biopsy approach. The authors failed to achieve their objectives and concluded that systematic sampling should be combined with the image-guided fusion biopsies [13]. In a large multicentre prospective study, researchers concluded a higher detection rate for clinically significant prostate cancer for a combined approach (TB and SB) biopsy method; however, different image fusion systems including cognitive guidance were used in the targeted biopsy approach [11]. Furthermore, the study did not use PIRADS v2 categorisation and no standardised protocol was used for MRI imaging. This allowed biases and heterogeneity into the reported data. Our study is a protocol-driven prospective investigation with a standardised US/MRI fusion protocol. We assessed clinical variables that could help in identifying patients who may benefit from systematic random biopsies in addition to fusion targeted approach. The comprehensive analysis and outcomes using methodology of this study has not been reported in the literature [14,15], in particularly the net clinical benefit of the approach.

The aim of this study was to:

- (1) Compare the diagnostic accuracy of MRI/US fusion targeted biopsies, systematic biopsies and combined approaches in the detection of csPCa and define predictive factors where a combined approach could be used.
- (2) Quantify additional benefits of adding systematic biopsies to the targeted biopsies approach by constructing a nomogram and assessing its net clinical benefits.

2. Materials and Methods

2.1. Study Population

The study had ethical approval (14/ES/1070) and all participants provided informed consent for their imaging data to be used. The study also had Caldicott institutional approval through the East of Scotland Ethical committee and Caldicott permission (Caldicott/IGTCAL6358) to link data with electronic system wherever follow-up outcomes were needed. The study period was between April 2015 and March 2020.

The inclusion criteria were age between 40 to 76, abnormal digital rectal examination (DRE), PSA ≤ 20 ng/mL and MRI $< T3$ disease. Exclusion criteria were repeat biopsies, prior radiotherapy to prostate and diagnosis of acute prostatitis with the last 12 months or a history of PCa. All participants had pre-biopsy mpMRI and only MRI positive (PIRADS ≥ 3) were recruited into the study ($n = 198$). Patients then underwent prostate biopsy by the MRI/US fusion technique (Hitachi HI-RVS; Europe Holding, Steinhausen, Switzerland) by

operator 1. This was followed by a standard 12-core TRUS biopsy by a second operator (blinded to the MRI results). In total, 78/198 (39%) underwent radical prostatectomy (RP).

2.2. Outcomes

The primary outcome was to compare detection rates of csPCa using both SB and TB approaches alone and in combination. This was assessed both at biopsy and RP stages. csPCa was defined as the presence of prostate cancer with Gleason Score $\geq 3 + 4$ (International Society for Urological Pathology [ISUP] grade 2 or more).

The secondary outcome was to assess the net clinical benefit of the approaches using nomogram and decision-analysis methods.

2.3. Sample Size Estimation

In considering performing a McNemar matched test, a minimum sample size of 110 men undergoing both SB and TB prostate biopsy approaches would be required to yield 90% power with a significance level of 0.05 ($\alpha = 0.05$), which would also allow 20% of the dropout rate. The csPCa detection rate via Sb and TB were found in a previous study [16]. Therefore, we recruited more patients than the minimum required number from the sample estimation to ensure achievement of study power and significance level.

2.4. Multi-Parametric MRI

All mpMRI scans were performed using 3T scanner (TIM Trio, Siemens, Erlangen, Germany) approximately 2 weeks before TRUS and MRI/US fusion biopsies. The mpMRI protocol was derived from the European Society of Uro-radiology Guidelines 2012 for the detection of prostate cancer and the subsequent publication of version 2 (Table S1—Supplementary file). Briefly, Table S1 summarises the MRI acquisition parameters. Prostate images were acquired in all three imaging planes, whereby the plane of the prostate was defined in relation to the rectal wall.

The mpMRI images were analysed and scored by experienced uro-radiologists (with more than 5 years post-certification experience using PIRADS v2.0). PIRADS v2.0 assessment categories were described as follows: score 1, clinically significant cancer is highly unlikely to be present; score 2, clinically significant cancer is unlikely to be present; score 3, the presence of clinically significant cancer is equivocal; score 4, clinically significant cancer is likely to be present; and score 5, clinically significant cancer is highly likely to be present.

2.5. Biopsy Procedures

All mpMRI scans were prepared by an experienced uro-radiologist (MSB) and fusion-targeted biopsies were performed by an experienced radiologist or in their presence using the Hitachi HI-RVS platform (Europe Holding, Steinhausen, Switzerland) using a pre-recorded lesion location. Three cores of tissue were obtained in TB approach from previously identified mpMRI lesions using a superimposed T2-weighted sequence on the real-time TRUS image. The systematic random 12-core biopsies were performed by an experienced urologist or specialist nurse following targeting.

The systematic random biopsy was typically a 12-core approach, collected in an extended sextant template of biopsies from the lateral and medial aspects of the base, mid, and apical prostate from the left and right sides. The biopsy results were analysed by experienced uro-pathologists who were blinded to the MRI findings. The Gleason Score for each patient was obtained.

A subset of the cohort ($n = 78$) underwent RP and their pathological stages were as follows: T2a = 2, T2c = 47, T3a = 24 and T3b = 5. Figure 1 summarises the study protocol in brief. The radical prostate specimens for histology were sliced in patient-specific moulds to aid orientation between imaging and histology per lesion, which were then fabricated using a 3D printer, as described previously by our group and others [17]. Specifically, patient-specific 3D printed moulds were made prior to surgery based on the T2-weighted MRI prostate capsule. The moulds were then customised for each patient using MIMICS

and Solidworks. This was used as a reference standard to assess the diagnostic accuracy of both SB and TB in detecting csPCa.

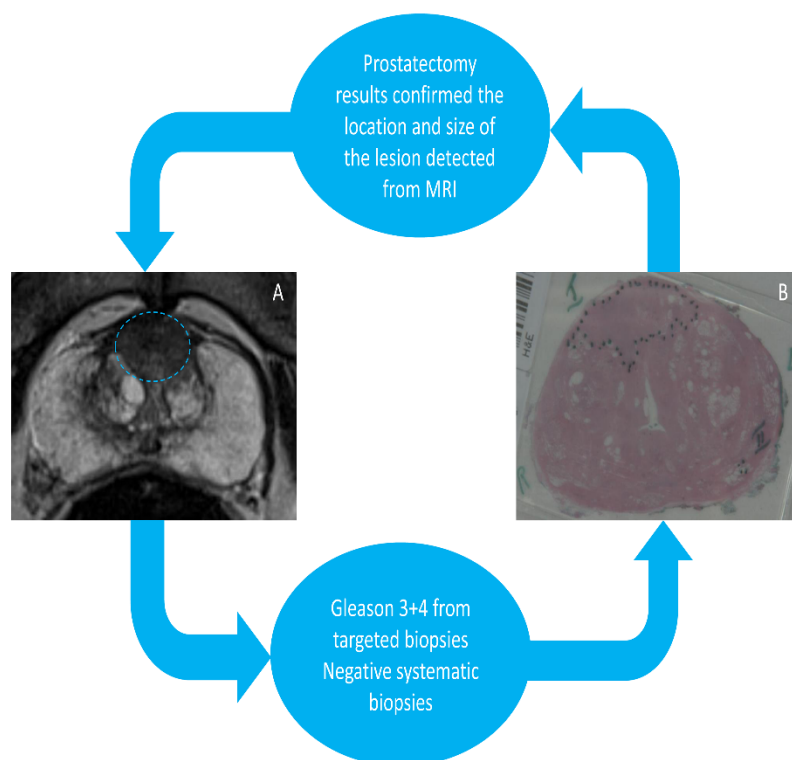


Figure 1. (A) A 76-year-old patient with a PIRADS 5 lesion detected from 3T MRI in anterior zone with a high PSA and abnormal DRE. (B) Patient-specific 3D mould-based grossing of a radical prostatectomy slice shows a 3 + 4 GS cancer located in the anterior zone.

2.6. Statistical Analysis

Patients' age, PSA, prostate volume (mL) and PSAD were collected. PSAD was calculated using PSA divided by the MRI-derived prostate volume (ellipsoid method). The number of MRI lesions, index lesion size (mm), PIRADS category and lesion location were measured by mpMRI. Each lesion was counted only once. The index lesion size was the size of the lesion with the highest PIRADS score. Continuous data were first tested to see if they were normally distributed by the Kolmogorov–Smirnov Test of Normality. The mean (m) and standard deviation (SD) were described if the variable followed a normal distribution. The median (M) and interquartile range (IQR) were presented if the variable was not normally distributed. Categorical variables are reported as frequencies and proportions. Cross tabulation was carried out in order to compare the proportions of csPCa patients by SB, TB, and combined SB + TB. The McNemar chi-square test was conducted in patients who were given both diagnostic tests. McNemar chi-square, degree of freedom (df) and *p*-value were calculated and presented. A two-step logistic regression was performed to identify explanatory variables of csPCa. First, patients' age, PSAD, lesion size, PIRADS and number of lesions were individually put into a univariate logistic regression model, where the outcome was defined as having csPCa or not. Statistically significant variables were then put into the multivariate logistic regression model. Odds ratio (OR), 95% confidence interval (95% CI) of odds ratio and *p* values were recorded. A nomogram was created based on the statistically significant variables in the final model. The discriminative ability of the predictive model was tested by receiver operating characteristics (ROC) curve and the concordance statistic (c-statistic) was presented. The predicted probabilities of csPCa were plotted against observed probabilities to test the calibration of the model. Decision curve analysis was applied to determine the benefit of the nomogram. In the subgroup analysis,

prostatectomies were performed in a group of 78 patients (112 lesions). The detection rate of true significant prostate cancer lesions via SB, TB and combined SB+TB was compared using McNemar chi-square test. Statistical analyses were conducted by SPSS V23.0 and R V4.0.3. The Bonferroni adjustment, which adjusted p value by times of tests, was used accounting for multiple testing. The alpha level (adjusted p -value) was set at 0.05/times of tests to determine two-tailed significance for McNemar chi-square test.

3. Results

3.1. Patient Characteristics

The participating patients' demographic data are shown in in Table 1. A total of 198 patients who underwent both systematic random and TB in the same setting were recruited into the study. Several clinical variables included baseline information (age, PSA level (ng/mL), PSA density (ng/mL²) and prostate volume (mL)), multi-parametric magnetic resonance imaging features (number of lesions seen on MRI, index lesion size (mm)) and PIRADS score.

Table 1. Characteristics of participating patients.

Variables		Overall ($n = 198$)
Basic information	Age, median (IQR), in years	67 (71–61)
	Prostate specific antigen (PSA), median (IQR), ng/mL	8.2 (10.6–6.4)
	Prostate volume, median (IQR), mL	47 (63–33)
	PSA Density, median (IQR), ng/mL ²	0.18 (0.27–0.11)
mp-MRI	Number of lesions, n (%)	
	1	102 (51.5%)
	2	75 (38%)
	3	14 (7%)
	4	6 (3%)
	5	1 (0.5%)
	Index lesion size, median (IQR), mm	16 (25–13)
	Prostate Imaging Reporting and Data System (PIRADS score), n (%)	
	PIRADS 3	22 (11%)
	PIRADS 4	55 (28%)
	PIRADS 5	121 (61%)
	Lesion location, n (%)	
	Peripheral zone (PZ)	79 (40%)
	Transition zone (TZ)	44 (22%)
	Both zones (TZ-PZ)	75 (38%)
Targeted (TB)/Systematic random (SB) biopsy	Detection of prostate cancer in TB, n (%)	129 (65%)
	Detection of prostate cancer in SB, n (%)	127 (64%)

3.2. Comparison of the Detection Rate of csPCa between SB, TB and Combined Approaches

The detection rate of csPCa using random biopsy was 51.0% (101/198) and using targeted biopsy was 56.1% (111/198). This was not statistically significant (McNemar chi-square test was $\chi^2 = 2.273$, $df = 1$, $p = 0.132$, Odds ratio (OR) = 0.63 (95% CI, 0.34 to 1.16). The results are shown in Figure 2. There were 17 patients (17/198; 8.5%) where the TB approach alone missed csPCa (eight from the same site and nine from normal-looking prostate on MRI). There were 84 patients (84/198; 42.4%) where the positive cores on systematic sampling and TB detected csPCa (72 from the same sector of index lesion and 12 from different sectors away from index lesion). When the TB is negative (69/198; 34.8%), the SB detected clinically insignificant cancer in 12 patients (12/69; 17.3%) and detected csPCa in eight patients (8/69; 11.6%). Twenty-seven (27/198; 13.6%) men were upgraded to csPCa based on TB, while 17 patients (17/198; 8.5%) were upgraded based on SB ($\chi^2 = 2.27$, $p = 0.13$).

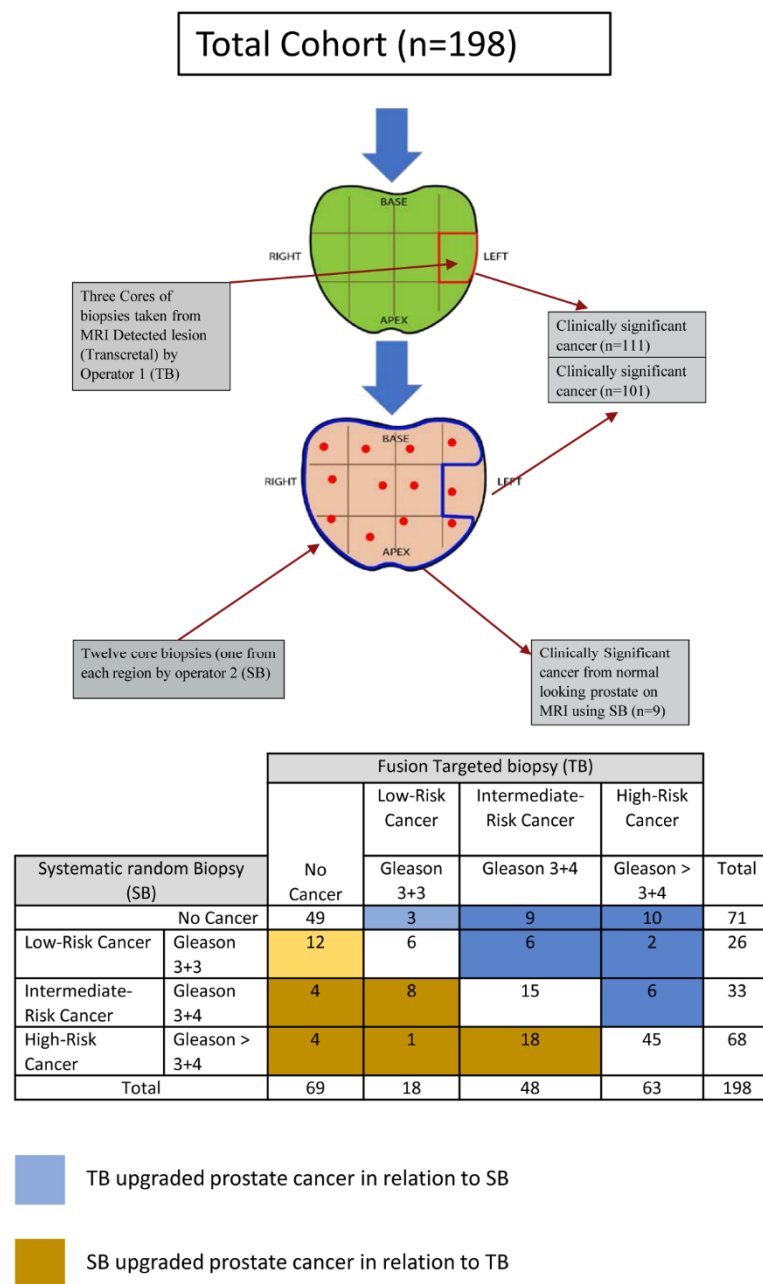


Figure 2. The detection rate of significant prostate cancer between SB and TB based on patients' level.

The detection rate of csPCa was higher using combined biopsy (64.6%; 128/198) in comparison to TB (56.1%; 111/198). The McNemar chi-square test result with the Yates correction was statistically significant ($\chi^2 = 15.06$, $df = 1$, $p < 0.001$). There was an 8.5% increase in significant prostate cancer detection rate at the patient level using combined biopsy methods compared to using TB alone.

We further validated findings using a subset of the cohort, where the histopathology of RP was used as a reference standard (Figure 3). There were 170 csPCa (170/190; 89.4%) seen on RP histopathology using mould-based approach and counting each focus of cancer. In total, 112 were targeted using MRI/US image fusion method. The TB approach detected 70 of these (70/112; 62.5%), whereas the SB approach detected 54 (54/112; 48.2%). The difference was statistically significant (the McNemar chi-square test result with the Yates correction was $\chi^2 = 6.618$, $df = 1$, $p = 0.010$, $OR = 0.36$ (95% CI, 0.17 to 0.77)). The combined approach to 112 lesions detected more cancers than SB or TB alone (79/112; 70.5%). Compared to SB, the combined approach detected 22.3% more cancers (70.5% vs. 48.2%).

The McNemar chi-square test result with the Yates correction was statistically significant ($\chi^2 = 23.04$, $df = 1$, $p < 0.001$). Similarly, the combined approach detected 8% more cancers in comparison to TB (70.5% vs. 62.5%). The McNemar chi-square test result with the Yates correction was statistically significant ($\chi^2 = 7.111$, $df = 1$, $p = 0.008$).

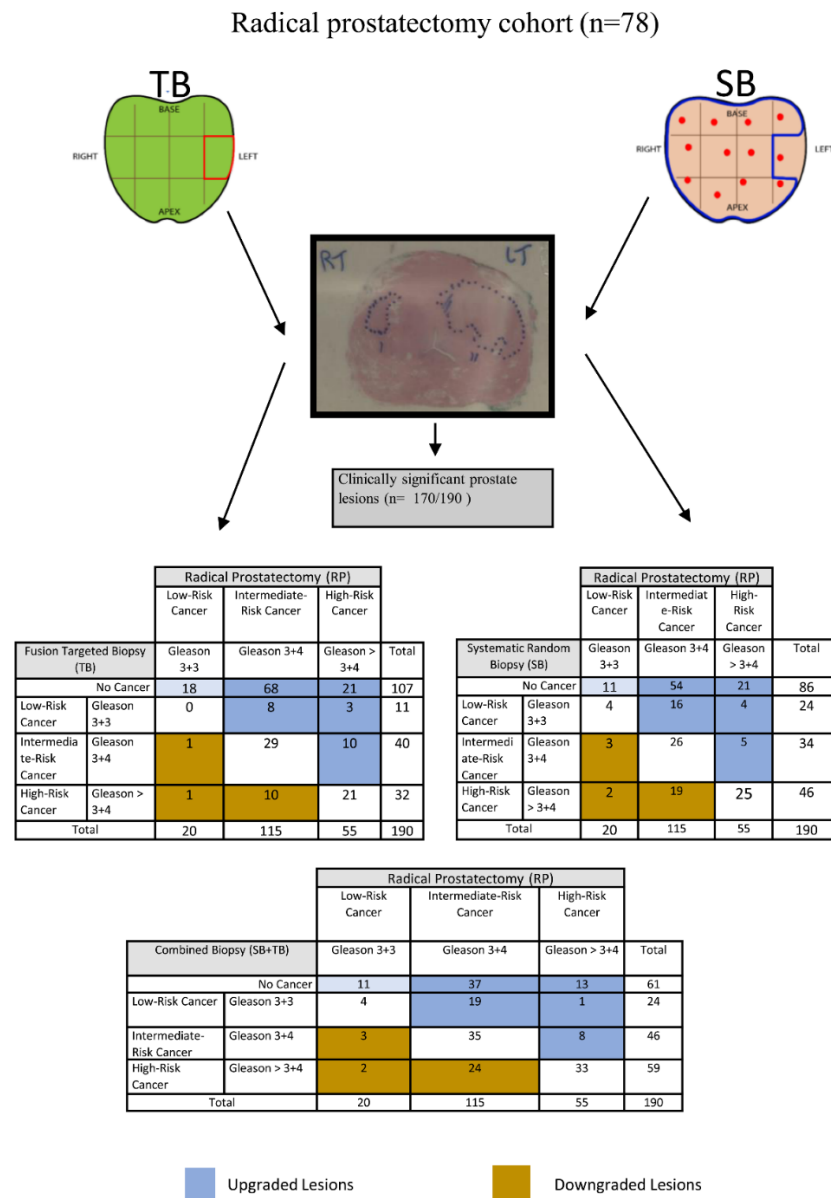


Figure 3. The detection rate of significant prostate cancer via SB, TB and combined SB+TB on RP lesions.

Interestingly, there were 11 cancers (11/190; 5.8%) which were labelled as clinically insignificant and all were upgraded to clinically significant using the TB approach. In comparison, there were 24 (24/190; 12.6%) cancers labelled as clinically insignificant and 20 (20/190; 10.5%) were upgraded using the SB approach. The McNemar chi-square test result with the Yates correction was $\chi^2 = 0.450$, $df = 1$, $p = 0.502$, OR = 1.50 (95% CI, 0.61 to 3.67). This was not statistically significant.

3.3. Univariate and Multivariate Logistic Regression Analysis and Developed Nomogram

In univariate logistic regression, patient's age, PSAD, Index lesion size and PIRADS were all significant predictors of csPCa detected by random biopsy (Table 2) and were,

therefore, inputted into the multivariate analysis. A 6% increase in odds of csPCa by random biopsy was associated with each one-year increase in age (OR = 1.06, 95% CI 1.01–1.12). A PSAD increase of 1 ng/mL was associated with an almost 26-fold increase in odds of csPCa (OR = 25.63, 95% CI 1.93–341.27). Having PIRADS-5 was another significant predictor of csPCa using random biopsy, which was associated with a six-fold increase in odds compared to those with PIRADS-3 (OR = 5.94, 95% CI 1.77–19.93).

Table 2. Univariate and multivariate logistic regression analysis.

Covariate	N	Univariate Logistic Regression				Multivariate Logistic Regression			
		OR	95% CI Lower	Upper	p Value	OR	95% CI Lower	Upper	p Value
Age (year)	198	1.07	1.02	1.12	0.009	1.06	1.01	1.12	0.036
PSAD	198	92.79	7.61	1130.69	<0.001	25.63	1.93	341.27	0.014
Index lesion size	198	1.06	1.03	1.10	<0.001	1.02	0.98	1.06	0.399
PIRADS	198				<0.001				0.001
3	22	Ref	-		-	Ref	-		-
4	55	1.69	0.49	5.80	0.406	1.51	0.42	5.43	0.525
5	121	9.46	3.00	29.84	<0.001	5.94	1.77	19.93	0.004
Number of Lesions					0.309				
1	102	Ref	-		-				
2	75	1.11	0.61	2.02	0.730				
3 and above	21	2.16	0.81	5.80	0.125				

The statistically significant independent variables from the multivariate logistic regression model (age, PSAD and PIRADS) were used to develop a nomogram to predict the probability of csPCa using SB (Figure 4). The model demonstrated good discriminative ability (C-statistic = 0.779, 95% CI 0.714–0.843 (Figure S1—Supplementary Material)).

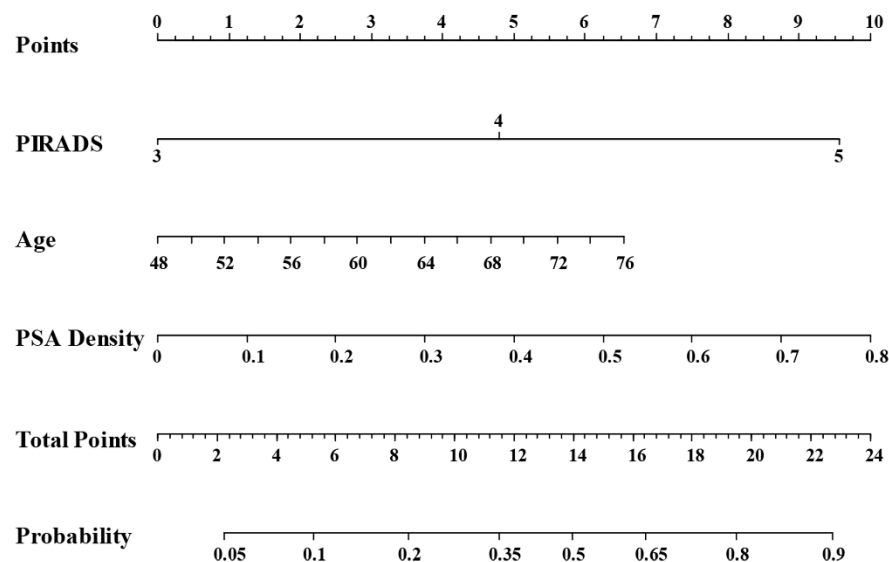


Figure 4. Nomogram with significant clinical variables to predict patients who will benefit from performing systematic random biopsy in addition to TB.

The calibration plot demonstrated a good agreement between the model predictions and actual observations for detecting csPCa via SB (Figure 5).

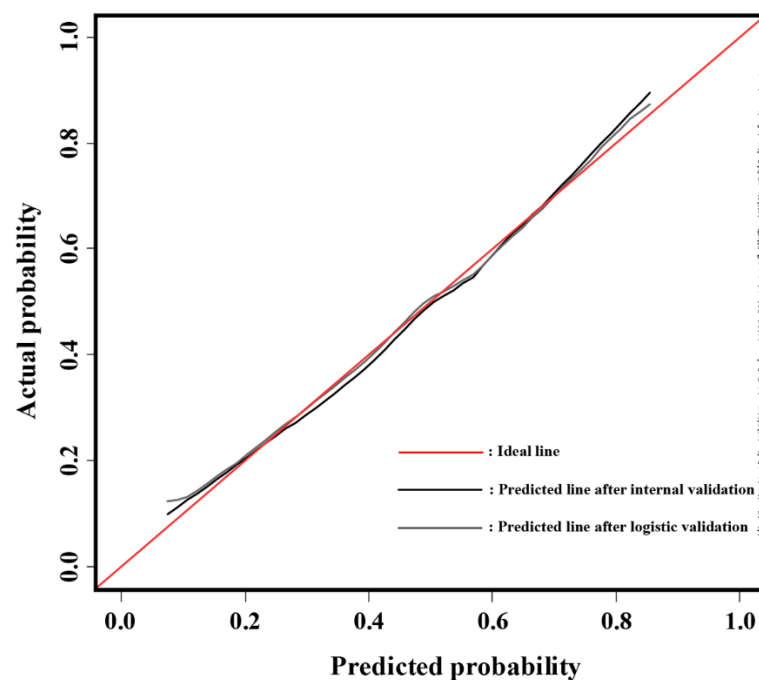


Figure 5. Model calibration plot for observed and predicted probability.

3.4. Decision Curve Analysis

The outcomes of the decision analysis curve are shown in Figure 6. The net benefit of performing SB in addition to TB on all cases is depicted by the grey line, whereas the black line represents the net benefit of not performing SB (only TB performed). To avoid the harm of unnecessarily intervening on the patients who are disease free and over intervening in the patients with disease, the net benefit of performing SB in addition to TB based on our prediction model with a reasonable range of threshold probabilities is shown as a red line in Figure 6. The net benefit of using our prediction model is to identify patients at risk of having csPCa who will benefit from SB in addition to TB. Our nomogram increased the overall net clinical benefit when the threshold probability was <60% and improved the diagnosis of csPCa compared to avoiding SB biopsy in all.

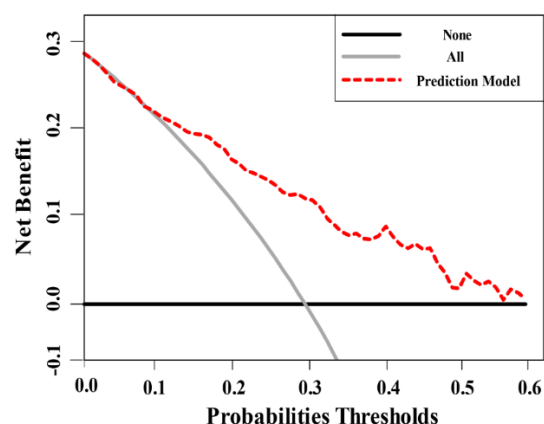


Figure 6. Decision analysis demonstrated a high net benefit of the model across a wide range of threshold probabilities.

4. Discussion

4.1. Principal Findings of the Study

This study assessed detection rate of csPCa using image fusion targeting, random systematic sampling and combination approaches. Patient-based analyses were further

validated using lesion-based data from RP histology. There were statistically significant higher detection rates for the combined biopsy approach in comparison to SB or TB alone. The TB approach alone would have missed 17 csPCas. Therefore, the combined approach detected more csPCa than either SB or TB alone. These results are similar to those reported by Filson et al. [18]; however, they were different to those reported in the PRECISION trial [1]. Therefore, in our observations, omitting concurrent SB during image-fusion may run the risk of missing csPCas in around 8.5% of patients. Similar to Cash et al. [19], we observed TB missing a small number csPCa in targeted areas. It is essential that we balance the advantages of concurrent sampling of the prostate during targeting against the risk of side effects and increased detection of clinically insignificant prostate cancers. Avoiding or adding systematic random biopsy at the time of TB remains a challenge for physicians, as knowledge and evidence of decision-making contributing factors still remains known [20]. Our outcomes from the nomogram indicated the excellent advantage (C-index 78% vs. 70%) of using a multivariable prediction model adjusting for clinical and radiological features (age, PSAD and PIRADS). The nomogram could be used to assist in selecting a group of men where a combined biopsy approach would be more useful.

We have also observed no significant advantage of improved characterisation of csPCa using the TB approach as all cancers labelled as clinically insignificant were upgraded on the final histopathology of RP. The challenge of upgrading or under grading would continue with both biopsy approaches as seen in our previous study [21]. There could be various reasons, such as inadequate sampling due to cancer heterogeneity and poor visibility of cribriform architecture on MRI and in biopsies [22].

4.2. Study Findings in Context of the Reported Literature

Several retrospective studies have assessed the outcomes of SB in addition to TB for the detection of csPCa. Sathianathan [23] et al. reported a nomogram with C-index = 70%. This nomogram was based on the clinical variables (biopsy naïve, previous biopsy and active surveillance patients) and imaging variables (number of MRI lesions and PIRADS score). The model provided a higher net clinical benefit at a threshold probability of <30%. The model was meant to predict csPCa in systematic random cores only (when TB was negative); however, our findings focused on predicting those patients who will benefit from performing systematic random biopsy in addition to TB (irrespective of target biopsy being positive or negative). Additionally, unlike their study, our nomogram, along with age, PSAD and PIRADS, found that adding these clinical variables to a model yielded a higher C-index (78% vs. 70%). In contrast to the present study, Sathianathan et al. [23] did not report on a validation cohort using RP as a reference standard. Furthermore, and similar to our study, others have reported the possibility of missing significant cancers if the image fusion targeted approach was offered alone [3,24–26]. Dell'Oglio et al. [13] failed to identify patients who might benefit from TB alone; therefore, they supported a combination of TB and SB as the preferred approach. In their study, there was no attempt to predict and assess the clinical variables that could help in identifying patients who might derive a greater benefit from systematic random biopsies. Lastly, Falagarío et al. [27], highlighted that smaller lesions in big prostates are more likely to be missed in TB biopsies; therefore, they developed a nomogram based on MRI volumetric parameters and clinical information for deciding when SB should be performed in addition to TB. In their study, all patients underwent biparametric MRI; however, in our analysis, we followed the standard mpMRI using PIRADS [28]. Moreover, the study was a multi-institutional retrospective data analysis of two previously published prospective trials with predominant fusion biopsies being cognitive rather than image-fusion using software. All men in the MULTI-IMPROD study [29,30] had transrectal systematic biopsies; therefore, these trials were not appropriate in answering the research question of the present study. In contrast to this study, however, Falagarío et al. [27] provided a range of probabilities of men missing clinically significant cancers, if SB was to be avoided altogether. We reported a set of measurable imaging criteria which could predict the likely benefit of adding SB to TB.

To our knowledge, this study is the first where lesion-based analyses were carried out using mould-based approach for a comprehensive pathological analysis. This confirmed that most csPCas were detected using a combined biopsy approach. csPCas were still missed by biopsies, which may be due to smaller lesions or the cribriform pattern seen on histopathology [31].

4.3. Clinical Implications of the Study Findings and Limitation of the Study

Decision-making using critical analysis, especially in situations of uncertainty, cost pressures and likely patient morbidity, is inevitably based on evidence or on a set of observations. In this study, we presented a decision-curve analysis estimating the net clinical benefits of offering a diagnostic test (combined approach to biopsy) in comparison to TB or SB approaches alone. The clinical and radiological observations were used to construct a nomogram, which is then the basis of a decision-making curve. The curve includes intervention for all and intervention for none and provides a background to facilitate discussions with patients. A balance has to be achieved between maximising detection of csPCas and avoiding detection of clinically insignificant cancers.

At present, various nomograms are used mainly for taking into consideration clinical factors, such as age, pre-operative PSA level and PIRADS score of the suspicious cancers. The present study reports a nomogram based on clinical parameters (age, PSAD and PIRADS). The nomogram clearly showed an improved prediction rate, which can be used to perform additional biopsies and the findings have substantial implications for clinicians and researchers in this area. We envisage that this and further research should bring us closer to precise decision-making. There will remain a group of men where systematic the random biopsy approach would bring value in addition to the TB approach, and thereby, improve informed decision-making in the management of men suspected of having prostate cancer.

However, there are some limitations to this study. This is a single-centre study with dedicated uro-radiologist and pathologists. We wanted to explore the association between lesion location in prostate and csPCa via SB but due to low numbers of lesions in TZ, which was not possible. It was not possible to use PIRADS v2.1, since the enrolment to study started before its publication, and this is a similar challenge to any other study published recently on this topic [11]. The nomogram in the present study has been internally validated (cross-validation and bootstrapping). External validation of the nomogram was not carried out in this study, as this would require further prospective multi-centre recruitment of a cohort to test external validity.

5. Conclusions

The study reports a nomogram using clinical variables which can assist decision-making during counselling. Patients could be directed towards having systematic sampling of the prostate in addition to an image fusion biopsy approach. The decision analysis curve confirmed a higher net clinical benefit of a combined biopsy approach compared to targeted or random sampling at an acceptable threshold.

Supplementary Materials: The following are available online at <https://www.mdpi.com/article/10.3390/cancers14010001/s1>, Figure S1: Receiver operating characteristics (ROC) curve and area under the curve (AUC) for a model's discriminative ability, Table S1: MRI acquisition parameters.

Author Contributions: Conceptualisation, G.N. and Z.H.; methodology, S.A., X.Z., M.S.-B., Y.Z., C.W., J.W., Z.H. and G.N.; investigation, S.A., X.Z., C.W., Y.Z., J.W., M.S.-B., Z.H. and G.N.; writing—original draft, S.A. and G.N.; reviewing and editing, S.A., X.Z. and G.N.; funding acquisition, G.N.; supervision, G.N. and Z.H. All authors have read and agreed to the published version of the manuscript.

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Data Availability Statement: The data are available for scrutiny from external requests.

Conflicts of Interest: The authors declare no conflict of interest.

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