STATISTICAL MODELLING OF CARDIOVASCULAR DISEASE PATIENTS USING BAYESIAN APPROACHES

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DEDICATION

This thesis is dedicated to my beloved parents and husband, who pray all day and night for my success, instilled in me the virtues of perseverance and commitment and relentlessly encouraged me to strive for excellence.

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ABSTRACT

This study focuses on statistical modelling on cardiovascular disease (CVD) patients in Malaysia. A secondary dataset from the National Cardiovascular Disease Database-Acute Coronary Syndrome (NCVD-ACS) registry for the years 2006 to 2013 is utilised. Studies have shown that CVD affects males and females differently. Thus, a gender-specific analysis with regard to the risk factors and mortality among ST-Elevation Myocardial Infarction (STEMI) patients is needed. Initially, this study performed the standard multivariate logistic analysis where the aims are to identify risk factors associated with mortality for each gender and to compare differences, if any, among STEMI patients. The results showed that gender differences existed among STEMI patients. Even though females share the same risk factors as males, there are risk factors that relate only to females which may have increased their tendency to develop and increase the risk of mortality of CVD patients. An important contribution of this analysis is that it gives an understanding of possible gender-based differences in baseline characteristics, risk factors, treatments and outcomes which will help cardiac care specialists in improving current management of patients with CVD. Next, Bayesian analysis is proposed to develop a prognostic model of the STEMI patients. Bayesian Markov Chain Monte Carlo (MCMC) simulation approach is applied. Beside that, comparisons of the parameter estimates from the proposed Bayesian and frequentist models are made. The results showed that the proposed Bayesian modelling can deal correctly with the probabilities and provides parameter estimates of the posterior distribution which have natural clinical interpretations. In doing so, several programming codes for the Bayesian model development and convergence diagnostics in the Just Another Gibbs Sampler (JAGS) software in R interface are developed. In the final part of this study, a graphical probabilistic model framework defined using a Bayesian Network (BN) is proposed to identify and interpret the dependence structure between the predictors and health outcomes of STEMI patients. In doing so, the two learning processes are involved in obtaining the BN model from the data namely the structural learning and parameter learning. From the structural learning, 25 and 20 arcs were considered significant for males' and females' BN respectively. A few variables namely, Killip class, renal disease and age group were classified as key predictors as they were the most influential variables directly associated with the outcome of patients' status. Moreover, conditional probabilities for each feature were obtained. The novelty of this study is that it provides an indication on the strength of each arc in the network by exploiting the bootstrap resampling method in the structural learning. A graphical model is developed where the relationships in a diagrammatical form is capable to be displayed and the causeeffect relationships can be illustrated. An important implication of this model is that it identifies dependencies based on the different features of variables. It can also include expert knowledge to improve predictability for data driven research when information or resources regarding the variables are limited.

ABSTRAK

Kajian ini memberi tumpuan kepada pemodelan statistik pesakit kardiovaskular (CVD) di Malaysia. Satu set data sekunder daripada Pangkalan Data Kebangsaan Penyakit Kardiovaskular-Sindrom Koronari Akut (NCVD-ACS) bagi tahun 2006 hingga 2013 telah digunakan. Kajian menunjukkan bahawa CVD memberi kesan kepada lelaki dan wanita secara berbeza. Dengan itu, analisis khusus-jantina berkaitan dengan faktor risiko dan kematian di kalangan pesakit Infarksi Miokardium ST-aras tinggi (STEMI) diperlukan. Pada mulanya, kajian ini menjalankan analisis logistik multivariat yang bertujuan untuk mengenal pasti faktor risiko yang berkaitan dengan kematian bagi setiap jantina dan membandingkan perbezaan jantina, jika ada, di kalangan pesakit STEMI. Hasilnya menunjukkan perbezaan jantina wujud di kalangan pesakit STEMI. Walaupun wanita mempunyai faktor risiko yang sama dengan lelaki, terdapat faktor risiko yang berkaitan hanya dengan wanita yang mungkin meningkatkan kecenderungan mereka untuk mendapat dan meningkatkan risiko kematian pesakit CVD. Sumbangan penting dalam analisis ini adalah ia memberikan kefahaman tentang kemungkinan perbezaan jantina dalam ciri asas, faktor risiko, rawatan dan hasil yang akan membantu pakar penjagaan jantung dalam meningkatkan pengurusan pesakit CVD masa kini. Seterusnya, analisis Bayesian dicadangkan untuk membangunkan model prognostik pesakit STEMI. Pendekatan simulasi Rantai Markov Monte Carlo (MCMC) Bayesian digunakan. Di samping itu, perbandingan anggaran parameter daripada model Bayesian yang dicadangkan dan model frekuentis telah dibuat. Hasil kajian menunjukkan bahawa pemodelan Bayesian yang dicadangkan boleh mengganggarkan kebarangkalian dengan betul dan menyediakan anggaran parameter bagi taburan posterior yang mempunyai tafsiran klinikal semulajadi. Oleh yang demikian, beberapa kod pengaturcaraan untuk pembangunan model Bayesian dan diagnostik penumpuan dalam perisian Just Another Gibbs Sampler (JAGS) dengan antaramuka R telah dibangunkan. Di bahagian akhir kajian ini, satu rangka kerja model kebarangkalian grafik yang ditakrifkan menggunakan Rangkaian Bayesian (BN) telah dicadangkan untuk mengenal pasti dan mentafsir struktur kebersandaran antara pesakit dan hasil kesihatan pesakit STEMI. Dengan berbuat demikian, dua proses pembelajaran telah terlibat dalam mendapatkan model BN dari data iaitu pembelajaran berstruktur dan pembelajaran parameter. Dari pembelajaran berstruktur, masing-masing 25 dan 20 lengkok dianggap penting bagi BN lelaki dan wanita. Beberapa pembolehubah iaitu kelas Killip, penyakit buah pinggang dan kumpulan umur diklasifikasikan sebagai peramal utama kerana pembolehubah-pembolehubah ini adalah yang paling berpengaruh secara langsung dengan hasil status pesakit. Selain itu, kebarangkalian bersyarat untuk setiap ciri telah diperolehi. Penemuan baru kajian ini adalah ia memberi petunjuk kepada kekuatan setiap lengkok dalam rangkaian dengan mengeksploitasi kaedah persampelan cangkuk but dalam pembelajaran berstruktur. Model grafik telah dibangunkan di mana hubungan dalam bentuk rajah mampu dipaparkan dan hubungan sebab-akibat boleh digambarkan. Implikasi penting dalam model ini ialah ia mengenalpasti kebersandaran berdasarkan ciri-ciri pembolehubah yang berbeza. Ia juga boleh memasukkan pengetahuan pakar untuk meningkatkan kebolehramalan untuk penyelidikan yang berasaskan data apabila maklumat atau sumber berkaitan pemboleubah adalah terhad.

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LIST OF ABBREVIATIONS

ACS	-	Acute Coronary Syndrome
AIC	-	Akaike Information Criterion
ANCOVA	-	Analysis of Covariance
ANN	-	Artificial Neural Network
aSAH	-	Acute Aneurysmal Subarachnoid Haemorrhage
AUC	-	Area Under Receiver Operating Characteristic Curve
BDe	-	Bayesian Dirichlet equivalent
BDeu	-	Bayesian Dirichlet equivalent uniform
BMI	-	Body-Mass Index
BN	-	Bayesian Network
CI	-	Confidence Interval
CVD	-	Cardiovascular Disease
DAG	-	Directed Acyclic Graph
DIC	-	Deviance Information Criterion
ETS	-	Environmental Tobacco Smoke
FDA	-	Food and Drug Administration
FPR	-	False Positive Rate
GS	-	Grow-Shrink
GVIF	-	Generalized Variance-Inflation Factors
HL	-	Hosmer and Lemeshow
ICU	-	Intensive Care Unit
IgAN	-	Immunoglobulin A Nephropathy
JAGS	-	Just Another Gibbs Sampling
MC	-	Monte Carlo
MCMC	-	Markov Chain Monte Carlo
MetS	-	Metabolic Syndrome
MI	-	Myocardial Infarction
MLE	-	Maximum Likelihood Estimation
MMHC	-	Max-Min Hill-Climbing
MMPC	-	Max-Min Parents and Children

MOH	_	Ministry of Health
		·
MREC	-	Medical Review & Ethics Committee
NCVD	-	National Cardiovascular Disease Database
NHMS	-	National Health and Morbidity Survey
NSAIDs	-	Non-Steroidal Anti-Inflammatory Drugs
NSTEMI	-	Non-ST Elevation Myocardial Infarction
OR	-	Odds Ratio
PCAD	-	Premature Coronary Artery Disease
PCI	-	Primary Percutaneous Coronary Intervention
pdf	-	Probability Density Function
PK-BUGS	-	Pharmacokinetic BUGS
PSRF	-	Potential Scale Reduction Factors
ROC	-	Receiver Operating Characteristic
SE	-	Standard Error
SHS	-	Second-Hand Smoke
STEMI	-	ST-Elevation Myocardial Infarction
TPR	-	True Positive Rate
UA	-	Unstable Angina
VIF	-	Variance Inflation Factor
WHO	-	World Health Organization
WHO/ISH	-	World Health Organization/ International Society of
		Hypertension

LIST OF SYMBOLS

χ^{2}	-	Chi-square
O_{i}	-	Number of observations of type <i>i</i>
п	-	Number of cells in the table for the Chi-square test
E_{i}	-	Expected frequency of type <i>i</i>
R_i^2	-	Coefficient of determination for a model where the
1		<i>i</i> th variable is fit against all other predictor
		variables in the model
df	-	Degree of freedom
$t \in \mathbb{R}$	-	Real input in logistic function
$\pi(t)$	-	Predicted probability of mortality
F(x)	-	Probability of the dependent variable equalling a
		case
$oldsymbol{eta}_0$	-	Intercept value
eta_i	-	Regression coefficient for the <i>i</i> th variable
$x' = (x_0, x_1, x_2,, x_p)$	-	CVD risk factors associated to mortality of patients
		in the NCVD-ACS data
l	-	Log-likelihood
${\cal Y}_i$	-	Vector of observed class
а	-	Number of exposed cases
b	-	Number of exposed non-cases
С	-	Number of unexposed cases
d	-	Number of unexposed non-cases
G	-	Likelihood ratio test which follows a Chi-squared
		distribution with k degrees of freedom
k	-	Difference in the number of parameters between the
		two models
j	-	Iteration count
$p(\theta)$	-	Prior distribution

$p(y \theta)$	-	Likelihood function
p(y)	-	Distribution of the data
μ	-	Parameter of Bernoulli distribution
x	-	Directly proportional to
	-	Conditional event or given
W	-	Within chain variance
В	-	Between chain variance
Ŕ	-	Potential scale reduction factors
m	-	Number of chains
n	-	Number of iterations
s_j^2	-	Variance of the <i>j</i> th chain
$ heta_{ij}$	-	Model parameter <i>i</i> in <i>j</i> th chain
$\overline{oldsymbol{ heta}}_j$	-	Sample posterior mean
$V\hat{a}r(\theta)$	-	Estimated variance
<i>S</i> (0)	-	Spectral density evaluated at frequency zero
T_n	-	Empirical distribution function constructed from
		the sample x_j
α	-	Significance level in the hypothesis tests
G	-	Directed acyclic graph
V	-	A set of nodes in the directed acyclic graph
Α	-	A set of directed arcs in the directed acyclic graph
X	-	A set of random variables in the directed acyclic
		graph
$\perp \perp_G$	-	d-separate
\rightarrow	-	Direction arrow (to)
v	-	Intermediate node
В	-	Bayesian network
D	-	Dataset used in the Bayesian network
Θ	-	Parameters of the global distribution
$P(B D) = P(G,\Theta D)$	-	Bayesian network learning
Pig(G Dig)	-	Structural learning

$Pig(\Theta ig G, Dig)$	-	Parameter learning
α	-	Imaginary sample size
r_i	-	Number of categories for the node X_i
q_{i}	-	Number of configurations of the categories of the
		parents of X_i
n _{ijk}	-	Number of samples which have the <i>j</i> th category for
		node X_i and the kth configuration for its parents
G^*	-	Modified network
М	-	Number of bootstrap resamples
$P_{\scriptscriptstyle B}$	-	Joint probability of a CVD event in relation with a
		set of risk factors using BN
$\theta_{_{X_i\mid\Pi_{X_i}}}$	-	Joint probability distribution over risk factor X_i
S	-	Score
$f_k(s)$	-	Probability density function of the scores for class k
$F_k(s)$	-	Cumulative density function of the scores for class
K		k
$F_0(t)$	-	Sensitivity
$F_0(t)$ $F_1(t)$	-	Sensitivity Specificity
	- -	
$F_1(t)$	- - -	Specificity
$F_1(t)$ $G_{_{HL}}$	- - - -	Specificity Hosmer and Lemeshow test statistic
$F_{1}(t)$ $G_{_{HL}}$ $\widehat{\pi}_{_{ij}}$	- - - -	Specificity Hosmer and Lemeshow test statistic Estimated probability for observation <i>j</i> in group <i>i</i>
$F_1(t)$ G_{HL} $\widehat{\pi}_{ij}$ D^*		Specificity Hosmer and Lemeshow test statistic Estimated probability for observation <i>j</i> in group <i>i</i> Deviance

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CHAPTER 1

INTRODUCTION

1.1 Background of the Study

Cardiovascular disease (CVD) is the number one cause of death in Malaysia (Mohammad *et al.*, 2018) and globally (WHO, 2017; Gutierrez *et al.*, 2018). CVD is defined as a group of disorders of the heart and blood vessels which include coronary heart disease, cerebrovascular disease, peripheral arterial disease, rheumatic heart disease and congenital heart disease (WHO, 2017). More people die annually from this non-communicable disease than from any other cause. Of these deaths, an estimated 7.4 million were due to coronary heart disease and 6.7 million were due to stroke (WHO, 2017). It was estimated that 23.6 million people will die by the year 2030 due to CVD (Benjamin *et al.*, 2017). Amongst the more developed countries, the highest death rates from CVD are in Ukraine and Russian Federation with 718 and 654 deaths per 100,000 population respectively, while the lowest are in South Korea and Japan with 36.5 and 47.0 deaths per 100,000 respectively (Mozaffarian *et al.*, 2015).

Although various research has been done on CVD worldwide, it is important to better understand the disease pattern in Malaysia specifically and analyse the impacts of the study on clinical practice locally. It is worthwhile to note that, CVD accounted for 98.9 deaths per 100,000 population in Malaysia in 2012, or 29,400 deaths which is 20.1% of all deaths (WHO, 2017). Even worse, CVD remain as a principal cause of death in Malaysia for the last ten years, from 2005 to 2014 (Department of Statistics Malaysia Official Portal, 2016). Ample information on the burden of disease has also been obtained from death certifications and hospital admission records from the Malaysia Ministry of Health (MOH) hospitals where circulatory disease accounted for 6.99% of the total hospital admissions and 23.34% of all hospital deaths in 2014 (Ang and Chan, 2016; Ministry of Health Malaysia, 2016). In Malaysia, the National Cardiovascular Disease Database (NCVD), a service supported by the MOH, plays an important role to collect information about CVD across Malaysia. NCVD enables us to obtain the incidence of CVD, and to evaluate its risk factors and treatment in the country. This information is useful in supporting the MOH, non-governmental organizations, private healthcare providers and industry in programme planning and evaluation, leading to CVD prevention and control. The NCVD is established to integrate various existing databases for individual hospitals either in MOH hospitals or private and other data sources to achieve a nationwide cardiovascular database.

The Acute Coronary Syndrome (ACS) registry was officially launched on 31st March 2006. The Malaysian National Cardiovascular Disease-Acute Coronary Syndrome (NCVD-ACS) registry was the first stage in realising the rationale of a nationwide cardiovascular database. Previously, the risk prediction of ACS is unclear and might be different from cardiovascular disease patients with chronic stable angina. These are the reasons why NCVD-ACS registry is needed, as it provides the real-life data that would represent the population of Malaysia (Wan Ahmad and Sim, 2006).

Prevention of CVD requires timely identification of patients at increased risk to target effective dietary, lifestyle, drug interventions or treatments (WHO, 2016). These can be done by studying the risk factors, analysing and interpreting them using various statistical methods. Therefore, the existence of high-quality data as well as suitable statistical methods to analyse data is of significant importance.

Understanding a disease often requires research that examines multiple variables and their relationships. These include clinical and laboratory data and other attributes such as risk factors, socioeconomic factors among others. In order to integrate these multiple variables, statistical model is often used where it can be expressed in mathematical expression that portrays the relationships among the variables. The model is able to provide prediction and explanation on the nature of the illness. One commonly used model is the regression model where we understand how the typical value of the dependent variable changes when any one of the independent

variables is varied, while the other independent variables are held fixed (Suárez *et al.*, 2017).

Over the past few years, numerous regression models have been developed for predicting risk factors such as models based on logistic regression model (Awad and Al-Nafisi, 2014; Umamahesh *et al.*, 2014; Jousilahti *et al.*, 2016; Keto *et al.*, 2016; Yuan *et al.*, 2017) and Cox regression model (McClelland *et al.*, 2015; Wulsin *et al.*, 2015). This study commences by considering the logistic regression model with independent variables that include the risk factors associated with mortality among Malaysian CVD patients. For initial analysis, univariate analysis is performed to identify significant variables. This is followed by the multivariate analysis using the purposeful selection method to obtain the best model.

Using various statistical methods, this study attempts to model the data by validating, making head-to-head comparisons, and thus providing a parsimonious model that describes the data. Bayesian model is also considered to describe the data where Bayesian Markov Chain Monte Carlo (MCMC) simulation approach is applied in the analysis. The dataset used for this study is the NCVD-ACS dataset from the year of its establishment, 2006 until 2013.

Model performance is assessed through convergence diagnostics, overall model fit, model calibration and discrimination. Additionally, comparisons of the parameter estimates are made between Bayesian model and frequentist model. To our knowledge, Bayesian model using the MCMC method has not been used extensively in the analysis of CVD data in Malaysia.

Also, another aim of the study is to identify the dependence structures between variables and the outcome of CVD patients, graphical model based on the Bayesian network (BN) approach has been considered. The BN approach incorporated a few learning techniques namely structural learning and parameter learning. In addition, this study applied a bootstrap resampling approach to the structure learning, in the interest of estimate the strength of each identified dependence. Validation and the performance

of the BN model are assessed using the area under receiver operating characteristic curve (AUC) and the accuracy test.

1.2 Problem Statement

A few studies have been carried out for the CVD datasets worldwide such as the treatment attitude and clinical outcome of elderly patients with ACS (Zuhdi et al., 2016), prediction on CVD mortality among working men and women (Jousilahti et al., 2016), determination of the population trends of CVD risk factors in the national capital region (Prabhakaran et al., 2017) and among others. However, successful implementation of statistical methods remains a challenge among researchers as they are many common mistakes applied especially in biomedical research such as improper research design, inadequate sample size, unsuitable statistical test and overfitting regression model (Nuzzo, 2014; Moyé, 2016). A review of the literature reveals general awareness of this issue (George, 1985; Ioannidis, 2005; Nuzzo, 2014a; Moyé, 2016). In medical research, the purpose of statistical model is often used to describe relationships between variables and identify risk factors associated with the outcome of some illness; for example, Shah et al. (2015), examined if the likelihood of having coronary artery disease is influenced by factors involving age and smoking. Measures of association provide an initial impression of the extent of statistical dependence between variables. Besides, it is important to eliminate and evaluate the best set of variables to be used for building predictive models and identify the risk factors that influence the outcome.

In Malaysia, there is a lack of comprehensive studies on the risk factors of CVD. Previous studies of CVD only focused on treatment attitude (Zuhdi *et al.*, 2016; Venkatason *et al.*, 2016) and the samples are based only on Kuala Lumpur's population (Amiri *et al.*, 2014) and limited to certain states such as Kelantan (Mohd Noor *et al.*, 2013) and Pahang (Mohammad *et al.*, 2018). Moreover, there were also studies which focused only on certain age group such as adolescents (Thangiah *et al.*, 2017) and elderly (Azahar *et al.*, 2016). Most of these studies used descriptive and regression analyses. Besides, the prevalence of CVD in males is higher than in females

and mortality rates are also different between the two genders (Department of Statistics Malaysia Official Portal, 2016). This suggest that risk factors associated with mortality between males and females are different. Hence, this study will demonstrate the development of logistic regressions using the NCVD-ACS dataset particularly on the ST-Elevation Myocardial Infarction (STEMI) patients in Malaysia. Based on the regression model, we would like to find out what are the risk factors and what is the prognosis of the patients. Is there any difference between male and female patients' prognosis?

An alternative to traditional frequentist statistical analysis is the Bayesian approach. The flexibility of the Bayesian approach is that we are able to update prior information on the underlying parameters with information from cumulative or past experience (Ntzoufras, 2009; Torman & Camey, 2015) and this has become the motivation in this study. A search through the literature revealed that implementation of Bayesian approach in CVD dataset are somewhat limited in Malaysia. Difficulties in performing Bayesian analyses for multivariate data and the availability of other easier methods are some of the reasons why the Bayesian approach is less popular compared to other traditional methods. Bayesian approach rests on the assumption that all model parameters are random quantities and hence can incorporate prior knowledge. Its use in predicting risk of mortality in CVD has been rather underutilized (Hannan et al., 2005) even though this method is widely used for predictive analysis in other medical applications such as modelling risk of death in an intensive care unit (Wong and Ismail, 2016), identifying risk genes for schizophrenia and neurodevelopmental disorders (Nguyen et al., 2017) and multiple treatment comparisons in female urinary incontinence (Carlin et al., 2013). In this study, we will incorporate prior knowledge and investigate how does this affect the risk factors, the prognosis of the patients and how does this improve the prediction of the model. To our knowledge, there is no published study that discussed the performance of Bayesian approach using Malaysian medical data, particularly CVD patients in the NCVD-ACS registry. Therefore, this study considers MCMC approach where development of Bayesian model using Malaysian STEMI patients from the NCVD-ACS dataset is shown.

Modelling the risk factors associated with mortality of CVD taking all the related variables into account is rather challenging from a theoretical point of view, as the risk factors-mortality nexus is expected to be characterised by a rather complex dependence structure. Obvious computational challenges can be seen, especially when it involves quite a large dataset in which it requires long processing times and can only be performed with an adequate computer infrastructure in place (Torman and Camey, 2015; Liu, 2018). When research is driven mostly by available data and resources or information regarding the variables are limited, a model that can essentially identify dependencies based on the different features of variables and can include expert knowledge to improve predictability is needed. In this setting, graphical model such as Bayesian network seem to be a perfect fit. This model was increasingly being used in computer science (Lacave et al., 2018; Scutari et al., 2018), business analytics problems (van Wagenberg et al., 2018) as well as medicine (Foroushani et al., 2016; Zador et al., 2017), and genetics (Scutari et al., 2014). Both categorical and numerical data were involved in these previous studies. However, all of these studies have been based on international samples such as Spain, United State of America, United Kingdom and Iran and this model is relatively less explored in Malaysian CVD dataset. Also, most of these Bayesian network studies do not contain indication on the degree of confidence or strength of each arc in the network. In order to address this limitation, a graphical model based on Bayesian network will be applied to discover the dependence structure between variables in Malaysian CVD dataset and the conditional independence of a variable or groups of variables from a given variable or variables. Bootstrap resampling method will be considered in this study in which the degree of confidence of each arc can be obtained. In short, we would like to know what are the relationship defined by the Bayesian network where the prognosis is different.

In machine learning, fitting a model to our training dataset is one thing (Witten *et al.*, 2016), but how do we know that it generalizes well to unobserved data? How to make sure that it doesn't simply memorize the data we feed it and able to make good predictions on future samples? In order to address this constraint, this study will assess the performance of both proposed Bayesian regression and Bayesian network models and carry out the validation studies to identify whether the predictions are sufficiently accurate across different settings and populations.

1.3 Research Questions

In this study, we aim to address the following research questions:

- (a) What are the CVD risk factors associated with mortality among the STEMI patients in Malaysia? Is there any difference for each gender?
- (b) What is the alternative technique in identifying prognostic factor of STEMI patients?
- (c) How well can the proposed statistical model fit the Malaysian STEMI data?
- (d) What is the suitable model to describe the dependence structure between the predictors and health outcomes of STEMI patients?
- (e) How accurate the proposed model in identifying the dependencies between variables?

1.4 Objective of the Study

The objectives of the study are as follow:

- (a) To identify the CVD risk factors associated with mortality for each gender and compare differences, if any, among STEMI patients in Malaysia using frequentist approach.
- (b) To propose prognostic model using Bayesian MCMC method for STEMI patients.
- (c) To assess Markov chains convergence, model performance and result validation of the proposed Bayesian model.
- (d) To establish a framework of dependence structure for STEMI patients based on graphical probabilistic model defined using a Bayesian network.
- (e) To assess the accuracy of the proposed Bayesian network model.

1.5 Significance of the Study

In general terms, the analyses will give an accurate information of the illness nature for patients with CVD in Malaysia. The findings from this study will be beneficial in the following ways:

- (a) This study gives an understanding of CVD-STEMI cases in Malaysia and provides accurate information to clinicians.
- (b) This study provides predictive models using frequentist approach and Bayesian MCMC where risk factors, outcome of illness and prognosis of the disease can be identified.
- (c) A graphical model is developed where it is capable of displaying relationships in a diagrammatical form and the cause-effect relationships can be illustrated to the clinicians.
- (d) It is hoped that the results of the statistical analyses will be of use to clinicians treating CVD-STEMI patients by providing robust decision support system.

1.6 Scope of Study

The scope of the study is on patients with acute coronary syndromes (ACS) who are registered with the National Cardiovascular Disease Database (NCVD) with the main focus on ST-Elevation Myocardial Infarction (STEMI). The analyses were done on the information obtained from 2006 to 2013.

1.7 Organization of the Thesis

This thesis is divided into seven chapters:

- (a) **Chapter 1** provides the background of the study, problem statement, objectives, significance of the study, scope and outline of chapters in the thesis.
- (b) **Chapter 2** presents the literature reviews related to the study.
- (c) **Chapter 3** elaborates the background of the source of data and the methodology for the construction of the proposed models in this study.
- (d) **Chapter 4** presents the results and findings of modelling the mortality of STEMI male and female patients using multivariate logistic regression.
- (e) Chapter 5 discusses the performance of the proposed Bayesian modelling of STEMI patients using MCMC approach. This is followed by convergence diagnostics for the Markov chains and comparison between Bayesian and frequentist estimates in the final model.
- (f) Chapter 6 focuses on the development of BN models and parameter estimation for the local distributions. The performances of the BN models are also discussed.
- (g) **Chapter 7** summarises and concludes the overall findings of this research. This chapter also includes the research contribution to knowledge and recommendations for future work.

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LIST OF PUBLICATIONS

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