# PATIENT DETERIORATION PREDICTIVE MODEL USING LONG SHORT-TERM MEMORY RECURRENT NEURAL NETWORK WITH GENETIC ALGORITHM OPTIMIZATION

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A thesis submitted in fulfilment of the requirements for the award of the degree of Doctor of Philosophy

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### DEDICATION

This thesis is dedicated to my father, who taught me that the best kind of knowledge to have is that which is learned for its own sake. It is also dedicated to my mother, who taught me that even the largest task can be accomplished if it is done one step at a time. Not to forget my beloved wife Dr. Mayyada and my lovely kids Doha, Ibrahim, Salma, and Shatha who stand with me from the first step with all their power. It is also dedicated to my great brothers Mohammad, Abdel Latif, Hamza, and Ghaith as well as my little sister who do not avoid (or delay) to provide any kind of help.

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iii

### ABSTRACT

The clinical investigation found that early recognition and intervention are crucial for preventing clinical deterioration in patients in Intensive Care units (ICUs) as well as in general wards. Deterioration of patients is predictable and can be avoided if early risk factors are recognized and developed in the clinical setting. Existing patient deterioration prediction methods generally have some disadvantages such as limited to specific patient groups or diseases that lead to lack of generalization, low prediction performance, and less optimized model parameter setting. This thesis proposes a patient deterioration predictive model based on Long Short-Term Memory Recurrent Neural Network (LSTM-RNN) with Genetic Algorithm (GA) optimization. The LSTM-RNN predictive model able to accept multiple input and data types in both static and dynamic parameters to predict patient deterioration, in terms of mortality and sudden transfer of patients from general wards to ICU with good accuracy. Another main strength of this predictive model is the input dataset is based on minuteby-minute time-series data obtained from open-source MIMIC-III research database for both model training and testing, hence also contribute to good prediction performance. To identify the baseline reference model with optimal performance, the setting of LSTM-RNN predictive model is explored using heuristically approach in terms of number of hidden layers, number of neurons in the first hidden layer, number of epochs, feature selection approach, as well as the impact of data cleaning in data pre-processing. On the other hand, the GA acts as an optimization model to further enhance the prediction performance of the baseline reference LSTM-RNN predictive model by exploration and identification of the optimum parameter settings, which include observation window size, prediction window size, and number of neurons in the first hidden layer. In this study, the proposed predictive model is benchmarked with other related work in terms of various prediction model, data sequence type, patient's age involved, number and types of features, dataset splitting ratios, prediction and observation window size and data source. For standard benchmarking result comparison, the selected performance metrics includes accuracy, area under receiver operating curve (AUROC), and test loss. The benchmarking results show that the proposed model outperforms other related models in general as it is capable to predict patient deterioration up to six hours before the onset with minimum prediction accuracy above 0.80 as recommended in the clinical setting. In specific, the best optimum LSTM-RNN predictive model after GA optimization able to achieve AUROC of 0.933, prediction accuracy of 0.921, test loss of 0.435, longer prediction window of 4.77 hours while reducing the observation window from 24 hours to 9.6 hours (60%) at the same time. The proposed patient deterioration prediction model based on LSTM-RNN, and GA will be very useful to clinical team as they have more sufficient time window to take prompt medical action before the onset of deterioration. As a result, this will help to reduce the mortality rate of patients or sudden transfer of patients from general wards to ICU.

### ABSTRAK

Penyelidikan klinikal mendapati bahawa pengecaman dan intervensi awal adalah sangat penting untuk mencegah kemerosotan klinikal pada pesakit di Unit Rawatan Rapi (ICU) dan juga di wad umum. Kemerosotan pesakit dapat diramal dan dielakkan sekiranya faktor risiko awal dikesan dan dikembangkan dalam penetapan klinikal. Kaedah ramalan kemerosotan pesakit yang sedia ada secara umumnya mempunyai beberapa kelemahan tertentu seperti terhad kepada kumpulan pesakit atau penyakit tertentu yang menyebabkan kekurangan generalisasi, prestasi ramalan yang rendah, dan tetapan parameter model yang kurang dioptimumkan. Tesis ini mencadangkan sebuah model ramalan kemerosotan pesakit berdasarkan Rangkaian Jangka-Pendek Neural Berulang Ingatan Panjang (LSTM-RNN) dengan pengoptimuman Algoritma Genetik (GA). Model ramalan LSTM-RNN ini dapat menerima pelbagai input dan jenis data dalam kedua-dua parameter statik dan dinamik untuk meramalkan kemerosotan pesakit, dari segi mortaliti dan pemindahan pesakit dari wad umum ke ICU secara tiba-tiba dengan ketepatan yang baik. Satu lagi kekuatan utama model ramalan ini adalah set data inputnya adalah berdasarkan data siri masa minit-demi-minit yang diperolehi dari pangkalan data penyelidikan sumber terbuka MIMIC-III untuk latihan dan ujian model, oleh itu juga menyumbang kepada prestasi ramalan yang baik. Untuk mengidentifikasikan model rujukan asas dengan prestasi yang optimum, penetapan model ramalan LSTM-RNN adalah dieksplorasi dengan menggunakan pendekatan heuristik dari segi jumlah lapisan tersembunyi, bilangan neuron pada lapisan tersembunyi pertama, bilangan zaman, pendekatan pemilihan ciri, serta impak pembersihan data dalam pra-pemprosesan data. Dari sudut lain, GA bertindak sebagai model pengoptimuman untuk meningkatkan lagi prestasi ramalan pada model ramalan LSTM-RNN rujukan asas dengan penerokaan dan pengenalpastian tetapan parameter yang optimum, di mana meliputi ukuran tetingkap pemerhatian, ukuran tetingkap ramalan, dan jumlah neuron pada lapisan tersembunyi pertama. Dalam kajian ini, model ramalan yang dicadangkan adalah dibandingkan dengan karya lain yang berkaitan dari segi model ramalan yang pelbagai jenia, jenis urutan data, usia pesakit yang terlibat, bilangan dan jenis ciri, nisbah pemisahan set data, ukuran tetingkap ramalan dan tetingkap pemerhatian serta sumber data. Untuk perbandingan keputusan yang standard, metrik prestasi yang dipilih merangkumi ketepatan, kawasan di bawah keluk operasi penerima (AUROC), dan kehilangan ujian. Hasil perbandingan menunjukkan bahawa model yang dicadangkan mengatasi modelmodel lain pada umumnya kerana ia mampu meramalkan kemerosotan pesakit sehingga enam jam sebelum permulaan dengan ketepatan ramalan minimum yang melebihi 0.80 seperti yang disarankan dalam tetapan klinikal. Secara khusus, model ramalan LSTM-RNN optimum yang terbaik setelah pengoptimuman GA dapat mencapai AUROC 0.933, ketepatan ramalan 0.921, kehilangan ujian 0.435, tetingkap ramalan yang lebih panjang selama 4.77 jam sambil mengurangkan tetingkap pemerhatian dari 24 jam kepada 9.6 jam (60%) pada masa yang sama. Model ramalan kemerosotan pesakit ini yang dicadangkan berdasarkan LSTM-RNN dan GA akan sangat berguna kepada pasukan klinikal kerana mereka mempunyai masa yang lebih mencukupi untuk mengambil tindakan perubatan yang cepat sebelum permulaan kemerosotan. Justeru itu, ini akan membantu mengurangkan kadar mortaliti pesakit atau pemindahan pesakit secara tiba-tiba dari wad umum ke ICU.

## TABLE OF CONTENTS

# TITLE

13

D	ECLARATION	i
D	EDICATION	ii
A	CKNOWLEDGEMENT	iii
A	BSTRACT	iv
A	BSTRAK	v
T	ABLE OF CONTENTS	vi
L	ST OF TABLES	xi
L	xiii	
L	XV	
L	XX	
L	ST OF APPENDICES	xxii
CHAPTER 1	INTRODUCTION	1
1.	Background of the Study	1
1.2	2 Problem Statement	4

1.3	Objective	8
1.4	Scope of Work	8
1.5	Significance of the Study	9
1.6	Thesis Organization	12

# CHAPTER 2 LITERATURE REVIEW

2.1	Introduction	13
2.2	Common Parameters for Patient Deterioration Detection and Prediction	13
	2.2.1 Vital Signs	14
	2.2.2 Laboratory Measurement Tests	19
	2.2.3 Level of Consciousness	24
	2.2.4 Static Parameters	25

2	.3 I	Early V	Varning Scoring Systems	27
2		Predicti Models	ion of Deterioration using Machine Learning	30
	2		Predictive Models based on Logistic Regression	31
	2		Predictive Models based on Support Vector Machine	33
	2		Predictive Models based on Artificial Neural Networks	35
	2		Predictive Models based on Other Machine Learning Approaches	37
2		Predicti Models	ion of Deterioration based on Deep Learning	41
	2	-	Predictive Models based on Markov Models and Convolutional Neural Networks	43
	2		Predictive Models based on Recurrent Neural Networks	47
2		-	zation Gap in Predictive Models for pration of Patients	57
2	.7 (	Chapter	r Summary	64
		-	5	
CHAPTER		TOP-L FOR	LEVEL PROPOSED PREDICTION FRAMEWORK	67
			LEVEL PROPOSED PREDICTION FRAMEWORK PATIENT DETERIORATION	
3	.1 I	FOR Introdu	LEVEL PROPOSED PREDICTION FRAMEWORK PATIENT DETERIORATION	67
3	.1 I .2 7	FOR Introdu Top-Le	LEVEL PROPOSED PREDICTION FRAMEWORK PATIENT DETERIORATION ction	<b>67</b> 67
3	.1 I .2 7 .3 I	FOR Introdu Top-Le Dataset	LEVEL PROPOSED PREDICTION FRAMEWORK PATIENT DETERIORATION action evel Prediction Framework Architecture	<b>67</b> 67 67
3	.1 I .2 7 .3 I	FOR Introdu Top-Le Dataset 3.3.1 3.3.2	LEVEL PROPOSED PREDICTION FRAMEWORK PATIENT DETERIORATION action evel Prediction Framework Architecture t Layer	<b>67</b> 67 67 69
3	.1 I .2 7 .3 I	FOR Introdu Top-Le Dataset 3.3.1 3.3.2 3.3.3	LEVEL PROPOSED PREDICTION FRAMEWORK PATIENT DETERIORATION ection evel Prediction Framework Architecture t Layer Static Parameters in MIMIC-III Database Dynamic Parameters in MIMIC-III Database:	67 67 67 69 75
3	.1 I .2 7 .3 I	FOR Introdu Top-Le Dataset 3.3.1 3.3.2 3.3.3 3.3.3	LEVEL PROPOSED PREDICTION FRAMEWORK PATIENT DETERIORATION action evel Prediction Framework Architecture t Layer Static Parameters in MIMIC-III Database Dynamic Parameters in MIMIC-III Database: Vital Signs Dynamic Parameters in MIMIC-III Database:	<ul> <li>67</li> <li>67</li> <li>67</li> <li>69</li> <li>75</li> <li>76</li> </ul>
3 3 3	.1 I .2 7 .3 I .3	FOR Introdu Top-Le Dataset 3.3.1 3.3.2 3.3.3 3.3.3	<b>LEVEL PROPOSED PREDICTION FRAMEWORK</b> PATIENT DETERIORATION extion evel Prediction Framework Architecture t Layer Static Parameters in MIMIC-III Database Dynamic Parameters in MIMIC-III Database: Vital Signs Dynamic Parameters in MIMIC-III Database: Lab Tests Dynamic Parameters in MIMIC-III Database:	<ul> <li>67</li> <li>67</li> <li>67</li> <li>69</li> <li>75</li> <li>76</li> <li>78</li> </ul>
3 3 3	.1 I .2 7 .3 I .3 .3	FOR Introdu Top-Le Dataset 3.3.1 3.3.2 3.3.3 3.3.3 Predicti	LEVEL PROPOSED PREDICTION FRAMEWORK PATIENT DETERIORATION evel Prediction Framework Architecture t Layer Static Parameters in MIMIC-III Database Dynamic Parameters in MIMIC-III Database: Vital Signs Dynamic Parameters in MIMIC-III Database: Lab Tests Dynamic Parameters in MIMIC-III Database: Level of Consciousness	<ul> <li>67</li> <li>67</li> <li>69</li> <li>75</li> <li>76</li> <li>78</li> <li>80</li> </ul>
3 3 3	.1 I .2 7 .3 I .3 .3 .4 I	FOR Introdu Top-Le Dataset 3.3.1 3.3.2 3.3.3 3.3.3 3.3.4 Predicti 3.4.1	LEVEL PROPOSED PREDICTION FRAMEWORK PATIENT DETERIORATION externation and the second structure of the second structure structure of the second structure structure of the second structure structure structure structure of the second structure structu	<ul> <li>67</li> <li>67</li> <li>69</li> <li>75</li> <li>76</li> <li>78</li> <li>80</li> <li>81</li> </ul>

	3.4.4	Data Merging	84
	3.4.5	Prediction Model based on LSTM-RNN	85
		Predictive Models based on Support Vector Machine and Logistic Regression	86
3.5	Optimiz	zation Layer based on Genetic Algorithm	88
3.6	Explora	tion Layer	92
3.7	Framew	vork Evaluation Layer	93
3.8	Modell	ing and Verification Tools	94
3.9	Framew Metrics	vork Verification Strategy and Performance	96
3.1	) Chapter	Summary	101
CHAPTER 4		OSED PREDICTIVE LAYER BASED ON LONG T-TERM MEMORY-RECURRENT NEURAL /ORK	103
4.1	Introdu	ction	103
4.2	Parame	ter Configuration from the Exploration Layer	103
4.3	Data Pr	e-processing	105
	4.3.1	Cohort Selection	105
	4.3.2	Data Extraction	107
	4.3.3	Data Cleaning	110
4.4	Feature	s Selection	112
4.5	Data M	erging	114
4.6	Dataset	Splitting for Framework Generalization	117
4.7	Predicti	on Model based on LSTM-RNN Architecture	118
	4.7.1	Batch Normalization	128
	4.7.2	Dropout	130
	4.7.3	Learning Rate	132
	4.7.4	Early Stopping	134
		Experimental Settings of the Proposed Framework	135
4.8	Chapter	Summary	136

CHAPTER 5	PROPOSED OPTIMIZATION MODEL BASED ON MULTI OBJECTIVE GENETIC ALGORITHM	139
5.1	Optimization Layer based on Multi Objective Genetic Algorithm	139
5.2	Search and Optimization Problem	141
5.3	Detail Operation of Optimization Model based on Genetic Algorithm	146
	5.3.1 Chromosome Representation	147
	5.3.2 Initialization	148
	5.3.3 Fitness Evaluation	148
	5.3.4 Selection	150
	5.3.5 Crossover	151
	5.3.6 Mutation	153
5.4	Genetic Algorithm Implementation using DEAP Library	156
5.5	Random Number Generator and Seed Number	157
5.6	Genetic Algorithm based on LSTM Optimization	157
5.7	Chapter Summary	159
CHAPTER 6	RESULT AND DISCUSSION	161
6.1	Optimal Architecture of the Proposed LSTM-RNN Model	161
	6.1.1 Optimal Number of Hidden Layers of the LSTM-RNN Model	161
	6.1.2 Optimal Number of Epochs of the LSTM-RNN Model	163
	6.1.3 Optimal Number of Neurons in the 1 <sup>st</sup> Hidden Layer of the LSTM-RNN Model	165
6.2	Prediction Performance Analysis of Patient Deterioration	166
	6.2.1 Prediction of Patient Deterioration using Auto Feature Engineering by LSTM-RNN, Support Vector Machine, and Logistic Regression Models	166
	6.2.2 Prediction Performance of Patient Deterioration using Vital Sign Features Only	168
	6.2.3 Prediction of Patient Deterioration using Laboratory Tests Only	171

		Patient Deteriorati ature Selection App		.73
	5.2.5 Prediction of Exploration or	f Patient Deterio Data Pre-Processin		.77
6.3	Dptimization Perform Dptimum Solution Framework		d Prediction	.78
6.4	Results Benchmarking	g with Previous Wo	rks 1	82
	5.4.1 Results Bench Learning Mod	hmarking with Pr els	_	82
	5.4.2 Results Ben Window of 24	ē		83
	5.4.3 Results Ben Windows $\geq 48$		Observation 1	87
	6.4.4 Results Ben Windows (≤ 1	U	Observation 1	.89
	1	Performance B rent Predictive Mo	enchmarking lels 1	89
6.5	Chapter Summary		1	94
CHAPTER 7	CONCLUSION		1	95
7.1	Concluding Remarks		1	95
7.2	Research Contribution	18	1	97
7.3	Future Works		1	98
REFERENCES			2	201
LIST OF PUBLICATIONS		2	241	

## LIST OF TABLES

TABLE NO.	TITLE	PAGE
Table 2.1	Different Blood Pressure Categories according to the American Health Association	16
Table 2.2	Normal Ranges employed in several EWS Systems	28
Table 2.3	Summary of Previous Works based on Statistical and Machine Learning Prediction Models	40
Table 2.4	Summary of Previous Works based on Markov Models and CNN Models	46
Table 2.5	Summary of Previous Works based on RNN Prediction Models	55
Table 2.6	Description of Predictive Models Adopted GA as the Optimization Algorithm	63
Table 3.1	Input Data Summary of Dataset Layer	71
Table 3.2	Description of Tables in MIMIC-III Database	74
Table 3.3	CHARTEVENTS Table in MIMIC-III Database	77
Table 3.4	Vital Signs extracted from CHARTEVENTS Table of MIMIC-III Database	78
Table 3.5	LABEVENTS Table in MIMIC-III Database	79
Table 3.6	Description of different Lab Tests in MIMIC-III Database	80
Table 3.7	Description of GCS in MIMIC-III Database	81
Table 3.8	Software Libraries and Hardware Tools used	96
Table 4.1	Baseline Characteristics of Patients	108
Table 4.2	Selected Features after Performing Selecting a Percentile Technique	114
Table 5.1	Parameter Setting for Genetic Algorithm used in this Research	154
Table 5.2	Optimizable Settings	159
Table 6.1	Prediction Results to assign the Number of Hidden Layers for the Proposed LSTM-RNN Model	162
Table 6.2	Prediction Results to assign the Number of Epochs for the Proposed LSTM-RNN Model	164

Table 6.3	Prediction Results to assign the Number of Neurons in the 1stHidden Layer for the Proposed LSTM-RNN Model1			
Table 6.4	Accuracy Performance of Logistic Regression, Support Vector Machine and LSTM-RNN Models			
Table 6.5	Framework Prediction based on Vital Sign Features only	170		
Table 6.6	Framework Prediction based on Laboratory Measurements only	171		
Table 6.7	Framework Prediction based on Chi-Square "Selecting a Percentile" Feature Selection Approach	174		
Table 6.8	Performance Comparison Summary of Performance Trade-off	176		
Table 6.9	Performance Comparison Summary of Different Models	177		
Table 6.10	Performance Comparison Summary	178		
Table 6.11	Comparison of the Test Loss, Accuracy, and AUROC of the Baseline Reference Model with Optimized Model based on MOGA 1			
Table 6.12	Summary of the Selected Prediction Framework Configuration for Performance Benchmarking with Related Work	182		
Table 6.13	Benchmarking Results against Previous Works that used Deep Learning Model	184		
Table 6.14	Benchmarking Results against Previous Works with Observation Window of 24 Hours	185		
Table 6.15	Result Benchmarking against Previous Works with Observation Window $\geq$ 48 Hours	188		
Table 6.16	Benchmarking Results against Previous Works with Observation Window $\leq 12$ Hours	190		
Table 6.17	Specifications of Virtual Machine provided by Google Colaboratory	191		
Table 6.18	Computation Timing Performance Comparison of LSTM and GRU at various Computation Platform	191		
Table 6.19	Computation Timing Performance Comparison with Related Work	193		

# LIST OF FIGURES

FIGURE NO	. TITLE	PAGE	
Figure 1.1	Scope of the proposed patient deterioration prediction framework for ICU patients	10	
Figure 2.1	Blood Pressure Vital Sign	15	
Figure 3.1	Top-Level Prediction Framework for Patient Deterioration	68	
Figure 3.2	Structure of the Dataset Layer	70	
Figure 3.3	Overview of the MIMIC-III Database	71	
Figure 3.4	Glasgow Coma Scale Scoring of Consciousness Responses of Each Component	81	
Figure 3.5	Prediction Layer	82	
Figure 3.6	Observation Window and Prediction Window with respect to Time	82	
Figure 3.7	Population, Chromosome and Gene	88	
Figure 3.8	Phases of Genetic Algorithm	89	
Figure 3.9	Processes of Crossover and Mutation in Genetic Algorithm	90	
Figure 3.10	Exploration Layer	92	
Figure 3.11	Framework Evaluation Layer	93	
Figure 3.12	An Overview of the Experiments performed in this Research	97	
Figure 3.13	A Confusion Matrix and Evaluation Metrics derived from the Matrix	99	
Figure 4.1	Data Flow and Parameter Configuration of Prediction Layer	104	
Figure 4.2	Patient Distribution according to Age at Admission from total of 399 Patients	109	
Figure 4.3	A Simple Case to Merge Different Variables	116	
Figure 4.4	Data Merging Output Concatenation between Hidden State and Dense Layer of LSTM at every Time Stamp	116	
Figure 4.5	Training, Validation, and Testing Process Pipelines		
Figure 4.6	Forward Architecture of an LSTM cell	120	
Figure 4.7	Backward Architecture of an LSTM cell	123	

Figure 4.8	Data Structure of the Input Data used with LSTM-RNN DeepLearning Model (Dim = dimension and t = time in minutes)126		
Figure 4.9	The Architecture of the Proposed LSTM-RNN Approach	127	
Figure 4.10	An Example of Dropout. The Left Network is Fully Connected, and the Right has had Neurons Dropped with Probability of 0.5 in the 1 <sup>st</sup> Hidden Layer	132	
Figure 4.11	An Example of Adaptive Imputation Approach	136	
Figure 5.1	Pseudo Code for Genetic Algorithm	142	
Figure 5.2	Modified Genetic Representation of a Solution	147	
Figure 5.3	Formulation of the Selection Scheme	151	
Figure 5.4	Crossover Operation	152	
Figure 5.5	Algorithmic Formulation of the Crossover Scheme	152	
Figure 5.6	Algorithmic Formulation of the Mutation Scheme	153	
Figure 5.7	Algorithmic Formulation of all Operations	155	
Figure 6.1	Number of Hidden Layers vs Test Accuracy & AUROC	162	
Figure 6.2	Number of Epochs vs Test Accuracy & AUROC	164	
Figure 6.3	Prediction Performance Trend Analysis based on Vital Sign Features Only	170	
Figure 6.4	Prediction Performance Trend Analysis based on Laboratory Measurements Only	172	
Figure 6.5	Prediction Performance Trend Analysis based on Chi-Square Feature Selection Approach	174	

# LIST OF ABBREVIATIONS

AD	-	Alzheimer's Disease
ADNI	-	Alzheimer's Disease Neuroimaging Initiative
AHA	-	American Health Association
AKI	-	Acute Kidney Injury
ANN	-	Artificial Neural Network
APACHE	-	Acute Physical and Chronic Health Evaluation
aPTT	-	Activated Partial Thromboplastin Time
AUC	-	Area Under the Curve
AUROC	-	Area Under the Receiver Operating Curve
AVPU	-	Alert, Voice, Pain, and Unresponsive
BiLSTM	-	Bidirectional Long Short-Term Memory
BN	-	Bayes Net
BP	-	Blood Pressure
BRNN	-	Bidirectional Recurrent Neural Network
BUN	-	Blood Urea Nitrogen
CAC	-	Coronary Artery Calcium
$Ca^+$	-	Calcium ion
CCHS	-	Christiana Care Health System
CCU	-	Coronary Care unit
CDSS	-	Clinical Decision Support System
CHMM	-	Coupled Hidden Markov Model
CITI	-	Collaborative Institutional Training Initiative
CNMF	-	Constrained Non-negative Matrix Factorisation
CNN	-	Convolutional Neural Network
CPT	-	Current Procedural Terminology
CRI	-	Cardiorespiratory Insufficiency
CSRU	-	Cardiac Surgery Recovery Unit
CT	-	Computed Tomography
DEAP	-	Distributed Evolutionary Algorithms in Python
DI		Decilitre

DiasBP	-	Diastolic Blood Pressure
Dim	-	Dimension
DOB	-	Date of Birth
DRG	-	Diagnoses Related Group
ED	-	Emergency Department
EHR	-	Electronic Health Record
ELM	-	Extreme Learning Machine
EM	-	Expectation Minimization
EMS	-	Emergency Medical service
EWS	-	Early Warning Score
FC	-	Fully Connected
FFNN	-	Feed-Forward Neural Network
FOC	-	Free of Charge
GA	-	Genetic Algorithm
GB	-	Gradient Boosting
GB	-	Giga Bytes
GCS	-	Glasgow Coma Scale
GHz	-	Giga Hertz
GMM	-	Gaussian Mixture Model
GPU	-	Graphical Processing unit
GRNN	-	General Regression Neural Network
GRU	-	Gated Recurrent Unit
GUI	-	Graphical User Interface
HADM	-	Hospital Admission
HDP	-	Hierarchical Dirichlet Processes
HE	-	Hypotensive Episodes
HER	-	Electronic Health Record
HIPAA	-	Health Insurance Portability and Accountability Act
HIS	-	Hospital Information System
HMM	-	Hidden Markov Model
HR	-	Heart Rate
HRV	-	Heart Rate Variability
ICD	-	International Classification of Diseases

ICU	-	Intensive Care Unit
ID	-	Identifier
IMCU	-	Intermediate Care Unit
INR	-	International Normalized Ratio
$K^+$	-	Potassium
KB	-	Kilo Byte
kNN	-	k-Nearest Neighbour
LDCT	-	Low Dose Computed Tomography
LOOCV	-	Leave-One-Out Cross Validation
LOS	-	Length of Stay
LSTM	-	Long Short-Term Memory
MAD	-	Mean Absolute Difference
MCI	-	Mild Cognitive Impairment
MDRP	-	Multi Modal Disease Risk Prediction
MeanBP	-	Mean Blood Pressure
MEWS	-	Modified Early Warning System
Mg	-	Milli gram
MICU	-	Medical Intensive Care Unit
MIMIC	-	Medica Information Mart for Intensive Care
MLP	-	Multilayer Perceptron
MMDL	-	Multi Modal Deep Learning
MOGA	-	Multi Objective Genetic Algorithm
MVCC	-	Multi-Version Concurrency Thromboplastin
$Na^+$	-	Sodium
NaN	-	Not a Number
NB	-	Naïve Bayes
NBP	-	Non-Invasive Blood Pressure
NC	-	Normal Control
NEWS	-	National Early Warning Scoring
NP	-	Non-deterministic Polynomial
OW	-	Observation Window
PCA	-	Principle Component Analysis
PDPA	-	Probability Distribution Patterns Analysis

PHI	-	Protected Health Information
PICU	-	Paediatric Intensive Care Unit
pMCI	-	Progressive state Mild Cognitive Impairment
PPV	-	Positive Predictive Value
PSM	-	Patient Similarity Metric
PT	-	Prothrombin Time
PTT	-	Partial Thromboplastin Time
PW	-	Predictive Window
RDS	-	Real-Time Data Sensing
RF	-	Random Forest
RLR	-	Regularized Logistic Regression
RNN	-	Recurrent Neural Network
ROC	-	Receiver Operating Curve
RR	-	Respiratory Rate
RRV	-	Respiratory Rate Variability
SANMF	-	Subgraph Augmented Non-Negative Matrix Factorization
SAPS	-	Simplified Acute Physiology Score
SAX	-	Symbolic Aggregate Approximation
sCr	-	Serum Creatinine
SICU	-	Surgical Intensive Care Unit
sMCI	-	Stable state Mild Cognitive Impairment
SPO <sub>2</sub>	-	Saturation of Oxygen in the Blood
SQL	-	Structured Query Language
SVM	-	Support Vector Machine
SysBP	-	Systolic Blood Pressure
Temp	-	Temperature
TN	-	True Negative
TP	-	True Positive
TSC	-	Tissue Sodium Concentration
TSICU	-	Trauma/Surgical Intensive Care Unit
T3	-	Triiodothyronine
T4	-	Thyroxine
T-data	-	Text Data

T&T	-	Track and Trigger
UDRP	-	Unimodal Disease Risk Prediction
ViSiBiD	-	Vital Sign Big Data
VT	-	Ventricular Tachycardia
WEKA	-	Waikato Environment for Knowledge Analysis
WFDB	-	Waveform Database

# LIST OF SYMBOLS

$C_t$	-	Cell State at time t
t	-	Time
$h_t$	-	Hidden Representation at time t
i	-	Input gate
f	-	Forget gate
0	-	Output gate
<i>pi</i>	-	Probability
b	-	Bias vector
tanh	-	hyperbolic tangent
Xt	-	new event
bf	-	forget bias
<i>ht</i> -1	-	output of the previous cell
<i>a</i> t	-	input activation
Ua	-	input activation update
Wa	-	input activation weight
ba	-	input activation bias
İt	-	Input
$U_i$	-	input update
$W_i$	-	W <sub>i</sub> input weight
bi	-	input bias
Ot	-	output gate
$W_{o}$	-	output weight
$U_o$	-	output update
$b_o$	-	output bias
$\Delta t$	-	output difference as computed by any subsequent layers
$\Delta h_t$	-	output difference as computed by the next time step
$\delta C_t$	-	change in the output
$\delta h_t$	-	change in the state
$\delta a_t$	-	change in the activation input

δit	-	change in the input gate
$\delta f_t$	-	change in the forget gate
Ct-1	-	previous state
$\Delta o_t$	-	change in the output gate
$\delta x_t$	-	change in the new event
$\Delta h_{t-1}$	-	change in the output difference as computed by the next time
		step LSTM
$\delta gates_t$	-	change in gate values at time t
$\delta W$	-	change in weight
$\delta U$	-	change in update
$\delta b$	-	change in bias
κ	-	selective pressure
R(i)	-	rank of individual <i>i</i>
т	-	number of individuals
Pm	-	probability that one gene has been modified for a binary
S	-	a given binary
S	-	Space
N	-	Length of gene
indpb	-	individual input probability
Zfi	-	forecast value
Zoi	-	observed value

## LIST OF APPENDICES

APPENDIX	TITLE	PAGE
Appendix A	Certificates of Completion Various Courses to Obtain Access from MIMIC-III Database	230
Appendix B	The Proposed Prediction Algorithm Pseudo Code	234
Appendix C	The Proposed Optimization Algorithm Pseudo Code	239

#### **CHAPTER 1**

### **INTRODUCTION**

This chapter discusses the study background to illustrate the research motivation, followed by the problem statement, the research objectives and its associated scopes, as well as research significance.

### **1.1 Background of the Study**

Identifying of patients who have a high deterioration risk is vital so that treatment decisions, quality assurance, and resource use management can be guided to reduce mortality rate. Patients who are admitted to ICUs and survive hospitalization have a high mortality rate in the six months after discharge (Wunsch et al., 2010). A lot of these post-discharge deaths are within patients transferred to other acute-care hospitals (Vasilevskis et al., 2009) or long-term acute care facilities (Hall et al., 2012). Unidentified deteriorations could delay the ICU transfer of patients, which would necessitate resuscitation in as much as 67% of cases or eventually result in deaths (Wellner et al., 2017). A report by the American Health Association (AHA) in 2015 showed that about 209,000 in-hospital cardiac arrests occur annually in the United States of America (USA) (Kolte et al., 2015). There are approximately 2,300 annual cases of cardiac arrests in Swedish hospitals as reported by the Swedish Resuscitation Council, which oversees 95% of Swedish hospitals (Spångfors et al., 2016). It was also found by the 2010 USA government investigation that 44% of adverse events could have been clearly or likely prevented (Levinson and General, 2010). Some researchers in New Zealand (Davis et al., 2003), the United Kingdom (UK) (Vincent et al., 2001), and Canada (Baker et al., 2004) used deterioration as defined by the result of health care management instead of the underlying disease process in the assessment of more than 25,000 patient records, from which 8% - 17% of admissions were related

to unfavourable events, preventable deteriorations made thought to be around 37% - 51%, and 7% - 19% ended in disability or death.

To this end, several studies have put forward different definitions of deterioration that are dependent on the various causes and the involved critical procedure. For instance, some studies (Churpek et al., 2013; Churpek et al., 2014a; Hu et al., 2016b; Smith et al., 2013) defined the deterioration as the patient being transferred to an ICU or experiencing a cardiac arrest, while there are other researchers related the term to patients who are admitted, transferred to another specialised hospital for emergency surgical treatment, or died after revisiting the emergency department (ED) (Mochizuki et al., 2017). There is also a research demonstrated that deterioration is primarily connected with organ dysfunctions like liver failure, kidney injury, respiratory failure, ICU admission, or death at a hospital (Quinten et al., 2018). Further, deterioration have also been defined by several studies to be a patient's sudden transfer from the general ward to an ICU with positive pressure ventilation, vasopressors, fluid resuscitation, or any immediate procedure that may be conducted between 2 hours pre or 12 hours post transfer (Bonafide et al., 2014; Wellner et al., 2017). (Henriksen et al., 2014) has also defined the deterioration as a patient deviating from the specified normal range in the 2 - 24 hours interval after hospital admission. Nevertheless, in the present, the physiological importance of deterioration is appreciated and the exact definition of it is still vague among the scientific community (Zheng and Shi, 2018).

Deterioration of patients in can be avoided by utilizing technologies that detect deterioration in a timely manner, by logging several data types in health informatics systems, and processing the data by utilizing software analysis models with accurate performance (Bonnici *et al.*, 2013; Findlay *et al.*, 2012; Stewart, 2009; Stewart 2011). There are many excellent data-driven learning models could be implemented in clinical decision support system by the implementation of electronic health records (EHRs), Markov models (Santamaria Ariza *et al.*, 2020) and dynamic Bayesian network (Abebe and Tesfamariam, 2020) to study disease development through modelling the temporal characteristics of EHRs. Moreover, preventing the occurrence of patients' deterioration in an adequate time window turns into a need in medicinal

services communities and biomedical research fields. It is also imperative that hospital care quality is enhanced significantly so that unwanted results are reduced. The notable hypothesis is recent technology can be used so that models that were developed using dynamic variables (e.g., vital signs and/or lab tests) and static variables (e.g., age, gender, and admission type) are utilized to build and strengthen an automated classification algorithm that can predict deterioration accurately.

In this study, the patient deterioration is defined as the patients either suddenly being transferred to ICUs from general wards (i.e., urgent admission type), or ICU patients suddenly dying (Churpek *et al.*, 2013; Churpek *et al.*, 2016; Edelson *et al.*, 2018; Smith *et al.*, 2013). Studies by a few researchers (Goldhill and Sumner, 1998; Lundberg *et al.*, 1998) showed a sudden ICU transfer is related with worse outcomes and increased mortality. The complex patterns in patients' longitudinal data affect the clinical interventions and ICU deaths (Catling and Wolff, 2020). As such, this study intends to forecast these events more reliably prior to their occurrence so suitable preemptive action can be taken by the hospital staff.

The Early Warning Score (EWS) systems are currently the common utilized models to improve the early detection of deteriorating patients (Hu *et al.*, 2016b; Kivipuro *et al.*, 2018; Panday *et al.*, 2017; Quinten *et al.*, 2018; Singer *et al.*, 2016). These systems provide early notification or warning to medical teams to take suitable and prompt medical action to save patients' lives. The design of these systems aims to solve sudden harmful events by combining various measures into an exact score that is quantifiable. The systems normally are integrated with the hospital equipment, such as patient monitor to track when patients reach certain thresholds. For example, "Track and Trigger" (T&T) systems track vital signs based on their periodic measurement and act (triggered) when patient vital sign reaches a specific threshold value. Fletcher et al. (Fletcher and Cuthbertson, 2010) showed that T&T systems are based on an erroneous foundation which derives from huge datasets that are regressed logistically, resulting in the prediction of death by using certain parameters. Some hospitals have asserted that the problem of deteriorating patients can be solved by continuous monitoring via measuring the impact of diseases on patients' daily lives (Edelson *et al.*, 2018;

Newman, 2017; Tilly *et al.*, 1995). However, the long hours continuous monitoring would consume great human resources of medical teams in hospitals.

Identifying deterioration prior to its onset is a huge and challenging issue in modern healthcare. Much research have proposed different predictive models to reliably predict such occurrences. Based on sufficient observation window and prediction window, the crucial techniques used to solve patient deterioration problems are machine learning and deep learning models (Bonnici et al., 2013; Choi et al., 2017; Goodfellow et al., 2016; Shotton and Findlay, 2012; Stewart, 2009; Stewart 2011; Ward et al., 2016b). Machine learning applies computational methods that depend on past experience to predict a task or outcome perfectly (Ward et al., 2016a). In contrast, deep learning constitutes an operation to refine information in multiple stages, where highly purified information is gained after being put through successive filters (Bengio et al., 2017). However, unlike deep learning models, machine learning-based models cannot frequently provide accurate performance and explicit interpretability; as a result, this research aims to propose a generic prediction framework based on deep learning models. Technologies used for deep learning produce approximately 2.5 quintillion bytes of data daily, the volume, velocity, and variety of information enable the "Big Data" analysis (Masud and Al Harahsheh, 2016; Nepal et al., 2015). The highest quality personalised healthcare is provided by big medical data and it is a vital factor in the success of a healthcare industry that has been revolutionised (Cheng et al., 2016; Madsen, 2014). Therefore, this study aims to utilize data from various types of patients so that patient deterioration can be detected in real-time, and the occurrence can be predicted. This study is vitally needed so that its results can be used to save the lives of more patients and in the provision of better healthcare services for people in general.

### **1.2 Problem Statement**

This research takes into consideration of three major research problems of prediction of deterioration for patients. The first problem is the framework of deterioration-based predictive models. The second problem is the shortcoming of current predictive models based on machine learning / deep learning. The third issue is the optimization problem for prediction model.

The first outstanding issue in the framework is the impractical workflow (i.e., sequence of processes) embedded in a predictive model. One example of an unrealistic approach is using laboratory tests as the only variables in the patient deterioration prediction (AlNuaimi *et al.*, 2015; Masud and Al Harahsheh, 2016). Each patient undergoes different medical tests, and a particular patient might need undergo the same tests more than once. Patients are initially placed into different groups according to their demographic profile, and every group is further clustered into groups of patients with similar test profile. After that, every group is utilized to implement a predictive model. However, the results from such models cannot be benchmarked (Masud and Al Harahsheh, 2016). Also, some previous works (Clifton *et al.*, 2011; Hu *et al.*, 2016b) performed studies on hospitals specialized in certain diseases, as well as patients with targeted diseases like cancer and suspected infection or sepsis (Masud and Al Harahsheh, 2016). As a result, a strong influence on the final model causes a high variance in behaviour and performance. This also negatively affects the generalization of the proposed prediction framework based on selected models.

Besides, certain proposed deterioration prediction framework is most focusing on in-hospital deterioration outcome evaluation based on specific factor. For example, (Jones *et al.*, 2013) evaluates an in-hospital deterioration in USA focus on medical neglect. This predictive model-based framework takes a long time, uses a design that is retrospective, sometimes considers adverse events caused by pre-hospital treatment, and ignores the fact that sometimes part of the natural dying process involves deterioration. Superior frameworks are required to categorise patient risks in a prospective and stratified approach while they are being admitted, and at the same time updating the approaches to educating and care models in preventing, identifying, and improving care for clinical deterioration.

Another research issue is overcoming the performance issue of existing deterioration predictive models. A few researches (Garla and Brandt, 2012; Seide *et al.*, 2011) have proposed patient deterioration prediction model based on machine

learning, but they are inadequate because they only consider the most crucial of feature engineering in machine learning workflow. A number of researches also utilized logistic regression statistical model to build patient deterioration prediction model in ICUs (Churpek *et al.*, 2016). These researches (Churpek *et al.*, 2016; Kate *et al.*, 2016; Mao *et al.*, 2012; Quinten *et al.*, 2018; Spångfors *et al.*, 2016), obtained a minimum Area Under Receiver Operating Curve (AUROC) results of 0.679, 0.68, 0.74, 0.76, and 0.77, respectively. In fact, Machado et al. (Machado and Cortez-Pinto, 2013) illustrated that models with the AUROC value of 0.5 are considered randomly predictive models, whereas models with the AUROC value higher than 0.8 represent good discriminatory models.

There are also previous works (Mao *et al.*, 2012; Ong *et al.*, 2012), proposed support vector machine (SVM) to predict cardiac arrest within the next 72 hours and achieved clinical deterioration with AUROC of 0.781 and 0.775, respectively. Alnuaimi *et al.* (2015) used a decision tree model to predict mortality and obtained an accuracy of 0.77. Ghosh et al. (Ghosh *et al.*, 2017) used coupled hidden Markov models (CHMMs) to predict septic shock and received a likelihood of 0.71. As a result, it can be observed that most of the obtained AUROC values are less than recommended 0.8 and hence conventional machine learning predictive models suffer from inaccurate performance.

Current models are also subject to robustness due to data uncertainty problems like missing data, null values, and irregular sampling clinical measurements. Quantitative research regards missing data as the norm, but the effects of missing data in quantitative studies are occasionally severe, resulting in biased parameter estimates, loss of information, and an inferior findings' generalisability (Dong and Peng, 2013; PANDA). To compensate the problem of missing data, a few researchers (Mochizuki *et al.*, 2017) limited their analyses to patients with full data (i.e., no missing data is associated with any selected variable), which results in an approach that is not applicable in the real world. A few current models (Wellner *et al.*, 2017) made no effort to account for missing values. Thus, the undertaking of this work is proposing a predictive patient deterioration model that adopt superior techniques for raw data pre-processing to evolve data uncertainty issues.

Moreover, it is necessary to determine an optimal interval of time-series windows involved in predictive models. Past researchers mainly took parameter values and settings that were defined by other studies to use in their work. However, the parameters of predictive models need to be adjusted for various applications and databases to enhance their discrimination capability (Rashedi et al., 2013). Previous works (Caballero Barajas and Akella, 2015; Celi et al., 2012; Ghassemi et al., 2014; Ghassemi et al., 2015; Hoogendoorn et al., 2016; Hug and Szolovits, 2009; Johnson et al., 2017a; Joshi and Szolovits, 2012; Knaus et al., 1981; Le Gall et al., 1993; Lee and Maslove, 2017; Lee et al., 2015; Lehman et al., 2012; Pirracchio, 2016; Potes et al., 2017; Ripoll et al., 2014; Vincent et al., 1996) involved a 24-hour observation window, whereas others (Che et al., 2018; Deng et al., 2009; Harutyunyan et al., 2017; Joshi et al., 2016) involved a 48-hour observation window to achieve acceptable prediction performance; hence increasing the volume of data. The long duration of observation window will have great demand in data storage and compute intensive operations. Thus, this study aims to solve the research problem of decreasing the window of observation from 24 hours to only 4 hours (i.e., an 83% reduction) while still maintaining an acceptable accuracy rate in its prediction.

Further, this issue also facing challenge due to inability of automatic techniques to tune several models at the same time without conducting a grid search or using the trial and error approach (Yuan *et al.*, 2018). In predictive models, it is well known that bias will be larger for smaller observation windows. This issue can be mitigated if the observation window is sufficiently long. It is important to carry out the optimization and monitoring of individual units as well as the whole process, which can largely improve the structure of predictive models of deterioration of patients. As a result, it is necessary to build an optimization algorithm that can automatically tune several important parameters and settings at the same time, such as observation window, prediction window, and the number of units in hidden layers (either separately or in combination) to maximize the prediction performance. Towards this end, the optimization problem will be solved using a modified genetic algorithm (GA). Furthermore, implementing the proposed models using advanced hardware to overcome challenges of gain (i.e., execution time), estimation time, and testing processing time is a necessity in proposing predictive models.

### 1.3 **Objective**

Based on the research problems, this study aims to develop a generic and robust patient deterioration prediction framework targeted for ICU patients. The detail objectives are as stated below:

- (a) To develop an accurate predictive model that can accept multiple input and data types using Long Short-Term Memory (LSTM).
- (b) To design an automated optimization approach using the Genetic Algorithm (GA) for identifying optimum parameters for accurate prediction of patient deterioration.

### 1.4 Scope of Work

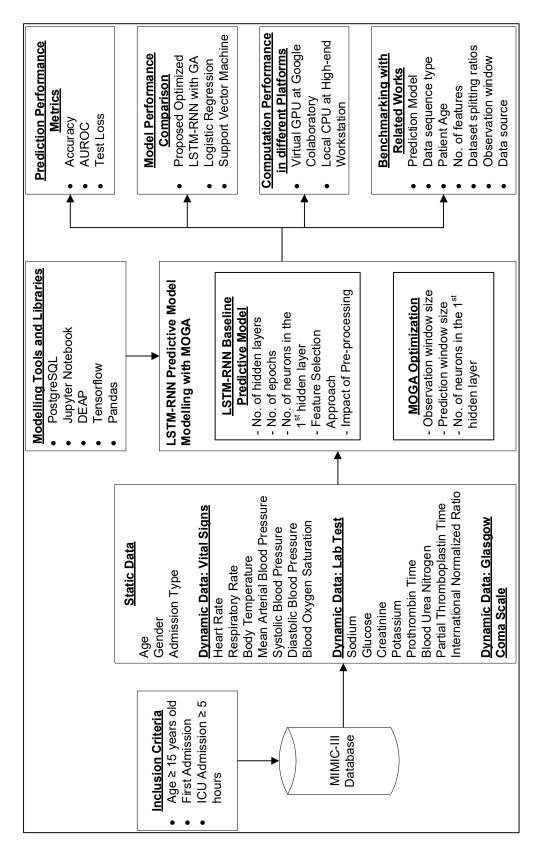
To fulfil the aforementioned research aims and objectives, this study has limited its research scope as shown in Figure 1.1. The prediction model is developed based on long-short term memory - recurrent neural network (LSTM-RNN) deep learning algorithm, to predict the patient deterioration in terms of patients either suddenly being transferred to ICUs from general wards, or ICU patients suddenly dying. The input of LSTM-RNN predictive model consists of two categories of data, which are static data and dynamic data obtained from Medical Information Mart for Intensive Care (MIMIC-III) version 1.4 "restricted access" database based on inclusion criteria of patient age more than 15 years old, first admission, and patients who have stayed more than 5 hours in ICU. Referring to figure 1.1, the static data consists of age, gender, and types of admission, whereas the dynamic data consists of seven vital signs, eight laboratory measurements, and Glasgow Come Scale (GCS) in the form of minute-by-minute time trends where each patient's selected parameter of interest is updated in every minute. The impact of data pre-processing technique and different feature selection approach to prediction performance is studied and compared. The baseline reference model is developed using heuristic approach to

identify optimal parameter setting in terms of number of hidden layers, number of epochs, and number of neurons in the 1<sup>st</sup> hidden layer.

On the other hand, the optimization model is designed based on Genetic Algorithm (GA) to further optimize the performance of the developed LSRM-RNN baseline reference model. It is conducted by auto exploration of different configuration setting in terms of observation window size, prediction window size and number of neurons in the first hidden layer through performance trade-off analysis. The performance metrics used for predictive performance evaluation and comparison includes accuracy, AUROC, and testing loss. For performance comparison, this work also develop other two different predictive models based on logistic regression (LR) and Support Vector Machine (SVM) using standard libraries provided in PostgreSQL and Jupyter Notebook. This research employs comprehensive benchmarking experiments with related previous works based on prediction task, sequence type, ages involved, number of features, splitting ratios, observation window, data source, performance metrics, and hardware features. All the modelling and performance analysis are executed in a virtual Graphical Processing Unit (GPU) provided by Google Collaboratory as well as a conventional Central Processing Unit (CPU).

## 1.5 Significance of the Study

The implementation of the proposed deep learning approach is expected to result in a new model that possesses reliable accuracy to predict patient deterioration. It is expected that the proposed model will assist in building a prediction model based on "Big Data" which has enhanced prediction accuracy. The clinical state of patients will be identified using this model via present and past data that comprise several parameters and measurements (i.e., periodic data). The interaction between various parameters is currently ignored by the existing prediction models. This study contributes by revealing previously unknown relationships between many variables (predictors) which could result in useful diagnostic or prognostic insights. The study also proposed the required clinical intervention to alleviate the effect of these events.





Moreover, this research uses definitions of deterioration, where its endpoint measure will be either mortality or sudden transfer to ICUs, which is used by researchers to obtain a better classification of patients.

In this study, the proposed predictive model is implemented using the state-ofthe-art GPU virtual machine provided by Google Colaboratory. Moreover, the study uses a minute-by-minute time-series approach. This approach enables the proposed model to obtain highly accurate results. The deep learning predictive model's ability to identify patterns in multivariate time-series of different clinical measurements is empirically evaluated by this research. To overcome the impractical workflow of predictive models that use one form of data, this study utilizes the individual and combined effectiveness of different types of variables (i.e., vital signs, laboratory measurements, GCS, and demographic data). Previous works face the problem of generalization due to data from hospitals specialized in certain diseases, or patients with certain diseases. In this study, data extracted from an open source that can be easily benchmarked and generalizing the results achieved.

Current predictive models suffer from weakness in performance due to using machine learning models that require feature engineering. However, this research proposes an LSTM-RNN deep learning model that does not require feature engineering. Existing predictive models use conventional hardware suffers from challenges in gain, estimation time and testing processing time. This work proposes an advanced hardware that overcome challenges in gain, estimation time via using a virtual GPU. The ad-hoc frameworks proposed by previous studies can be improved by the generic prediction framework proposed in this research, which will result in predictions of higher accuracy. The proposed predictive model could reduce the required observation window for the prediction task while improving the performance. In fact, the proposed significant small size of observation window could obtain higher results which outperformed all previous works that utilize different sizes of observation window (i.e., 48 hours and 24 hours). The proposed predictive model achieved accurate performances when using a prediction window with sizes longer than 1 hour.

The proposed optimization algorithm based on GA could improve the accuracy obtained by the predictive model. It also could increase the prediction window. It also reduced the observation window by 60% compared to the size of observation windows used by most of the studies used in the literature to predict the deterioration of patients (i.e., 24 hours). In addition, the proposed optimization algorithm could reduce test loss. The study identifies the most important medical lab tests without using any informed domain knowledge. Some current predictive models implement the structure of the models via trial and error, whereas this study propose an optimization model based on GA to determine the size of the observation window, prediction window, and number of units in the hidden layer.

### 1.6 Thesis Organization

The rest of the thesis is organised as described below. A comprehensive literature review of works related to this study is presented in Chapter 2. The methodology for proposing a generic prediction framework is discussed in Chapter 3. It also discusses the detail of different layers that form the generic prediction framework. Moreover, modelling software libraries and tools is demonstrated in this chapter. It also illustrates the performance metrics and framework verification strategy. Chapter 4 covers the details for the modelling and algorithmic development in the proposed predictive model based on LSTM-RNN performed in this thesis. It also includes the findings of performing the research methodology to obtain the dataset and the results of performing feature selection. A description of different sizes of observation window and prediction window is also included. Chapter 5 illustrates the test works that confirm the usefulness and dissect the presentation of the proposed modified optimization model based on multi objective GA. Chapter 6 presents the experimental works that verify the functionality and analyse the performance of the proposed predictive algorithm based on LSTM-RNN and the proposed optimized algorithm based on GA described in Chapters 4 and 5, respectively. It also includes the benchmarked results of performance against the related previous works. Chapter 7 shows the future works, contributions, and conclusions of this research.

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## LIST OF PUBLICATIONS

This list contains the papers that are generated based on this research. They are:

- Tariq Ibrahim Abdel Latif Al-Shawaheen, Mehrdad Moghbel, Yuan Wen Hau, Chia Yee Ooi, Use of learning approaches to predict clinical deterioration in patients based on various variables: a review of the literature, *Artificial Intelligence Review*, 13 March 2021 (Q1, IF = 5.747).
- Tariq Ibrahim Abdel Latif Al-Shawaheen, Yuan Wen Hau, Nizar Ass'Ad, Mahmoud M. Abualsamen, A Novel and Reliable Framework of Patient Deterioration Prediction in Intensive Care Unit Based on Long Short-Term Memory-Recurrent Neural Network, *IEEE Access*, Vol. 9, pg. 3894-3918, 24 December 2020 (Q1, IF = 3.745).
- Tariq Ibrahim Abdel Latif Al-Shawaheen and Yuan Wen Hau, A New Model for Tracking Deterioration of Vital Signs by Means of Artificial Neural Network, *Journal of Theoretical and Applied Information Technology* (*JATIT*), Vol. 97, No. 14, pg. 3809-3818, 31 July 2019. (Q1, IF = 0.628)