Optimal Combination of Centrality Measures for Node Importance Determination

Ho Khai Shen (U2103646)



1. Abstract and Objectives

Introducing centrality measures, a way to rank nodes, plus how and why to construct a convex combination of them. We mainly put our focus on convex combinations of two centrality measures. We introduce some well-known centrality measures and discuss what aspects they individually emphasise to rank nodes of a graph. We will see that different centrality measures provide a variety of valuable insights into node importance, and forming a combination of multiple centrality measures is necessary to capture a graph's whole structure. Objectives To determine a t_{μ} such that

- it maximise the correlation value between c_0 and $c_{t_{\mu}}$ and between c_1 and $c_{t_{\mu}}$
- it induces the most number of ranks (regular points)
- $c_{t_{\mu}}$ mimics both c_0 and c_1 better than c_0 and c_1 do to each other ($min\{\tau_b(c_0, c_{t_{\mu}}), \tau_b(c_1, c_{t_{\mu}})\} \ge \tau_b(c_0, c_1)$)

Definition 1: Centrality measures

A centrality measure of a graph G is a function $c: V \mapsto [0, \infty)$ with the following properties:

- If G is a star with centre u, then c(u) > c(v) for all nodes $v \in V \{u\}$.
- 2 If $G \cong H$ with bijection $\theta: V(G) \to V(H)$, then $c(v) = c(\theta(v))$ for all nodes $v \in V$. [1]

2. Well-known Centralities

Degree centrality

Harmonic centrality

- Closeness centrality
- Betweenness centrality Eigenvector centrality

Katz centrality PageRank centrality

Example: Different centrality measures give different rankings



Figure 1. A custom drawn graph

Table 1. Ranks of nodes of Figure 1 by degree, closeness, harmonic, betweenness, eigenvector, Katz and PageRank centrality.

Different centrality measures yield significantly different rankings of nodes, for this reason, a single centrality measure may not be enough to capture the whole network structure and the importance of nodes. Therefore, it is crucial to take into account the view of multiple centrality measures simultaneously, by taking the combination of them when ranking nodes. [2]

3. Convex Combinations of Centrality Measures

Given centrality measures c_0 and c_1 and a real number t, we define the functions tc_0 and $c_0 + c_1$ by setting

$$(tc_0)(v) = tc_0(v)$$
 and $(c_0 + c_1)(v) = c_0(v) + c_1(v)$

for all $v \in V$.

Definition 2: Convex combinations of centrality measures

Given centrality measures $c_0, c_1, ..., c_n$ and non-negative real numbers $t_0, t_1, ..., t_n$ such that $\sum_{i=0}^{n} t_i = 1$, we define the convex combination c_t as follows:

$$c_t = t_0 c_0 + t_1 c_1 + \dots + t_n c_n = \sum_{i=0}^n t_i c_i.$$
(1)

Theorem 3:

Let $c_0, c_1, ..., c_n$ be centrality measures. For any $(t_0, t_1, ..., t_n) \in \Delta^n$, the convex combination c_t is a centrality measure. [1]

Each score of node assigned by individual c_i contributes some weightage to c_t , which is determined by the value t_i . One way to determine the value of each t_i is by making sure c_t truly represents all individual c_i 's, that is, the correlation between c_t and each individual c_i 's are as large as possible.

4. Correlation Methods

5. Convex Combinations of Two Centrality Measures

A convex combination of two centrality measures, $c_t = (1 - t)c_0 + tc_1$, c_0, c_1 spanned by $\Delta^1 = \{(1 - t, t) \in | t \in [0, 1]\}$

Lemma 6:

If any pair of distinct nodes of G are concordant with respect to or tied on both c_0 and c_1 , then c_t will preserve this property for all $t \in [0, 1]$. [1]

Lemma 7:

Any pair of distinct nodes of G are either tied on c_{t^*} for at most one $t^* \in [0,1]$ or tied on c_t for all $t \in [0,1]$. [1]

Definition 8: Regular points

A point $t^* \in [0,1]$ is a critical point if there exist distinct nodes u and v of G that are ranked equally by c_{t^*} but u and v are either not tied on c_0 nor on c_1 . [1]

Definition 9: Critical points

A point $t^* \in [0,1]$ is a critical point if there exist distinct nodes u and v of G that are ranked equally by c_{t^*} but u and v are either not tied on c_0 nor on c_1 . [1]

Lemma 10:

For any pair of distinct nodes of G that is neither concordant wrt nor tied on both c_0 and c_1 , there exists a unique critical point $t^* \in [0, 1]$. Furthermore, for any $r, s \in [0, 1]$, the pair of nodes are concordant with respect to c_r and c_s if and only if either $r, s \in [0, t^*)$ or $r, s \in (t^*, 1]$. [1]

Lemma 11:

The ranking of nodes of a graph G induced by c_t , as t varies in [0, 1], changes only finitely many times and the number of times the ranking changes is

$$2|T^*| - |T^* \cap \{0, 1\}|,\tag{4}$$

where T^* is the set of critical points $t^* \in [0, 1]$ and $|T^*|$ is the cardinality of the set T^* . [1]

Lemma 12:

For a regular point t, the ranking of nodes of a graph G induced by c_t is insensitive to sufficiently small variation in t. Particularly, for any two consecutive critical points t_i^* and t_j^* , where $t_i^* < t_j^*$, the ranking induced by c_t stays the same as t varies in $t \in (t_i^*, t_j^*) \cap [0, 1]$. [1]

6. Optimizing Convex Combinations of Two Centrality Measures

Theorem 13:

Using Kendall's tau-b, there is always a $t_b \in [0,1]$ such that t_b is a regular point and c_{t_b} mimics both c_0 and c_1 better than c_0 and c_1 do to each other. [1]

Via minimizing the objective function $g(t; c_0, c_1) = (1 - \tau_b(c_0, c_t))^2 + (1 - \tau_b(c_1, c_t))^2$

- By Lemma 11, each $\tau_b(c_0, c_t)$ and $\tau_b(c_1, c_t)$ changes its values finitely many times as t varies in [0, 1]. By Lemma 6 and 10, changes in values occur only when it passes through a critical point.
- Therefore together with the Theorem 13, it is feasible to determine t_b and hence compute the optimum c_{t_b} by minimizing the objective function $g(t; c_0, c_1)$ with regular points.

Theorem 14:

Using Pearson's r, there is always a $t_r \in [0, 1]$ such that c_{t_r} mimics both c_0 and c_1 better than c_0 and c_1 do to each other.

Via minimizing the objective function $h(t; c_0, c_1) = (1 - r(c_0, c_t))^2 + (1 - r(c_1, c_t))^2$

With similar sayings as before, by Lemma 6, 10, 11 and Theorem 14, it is feasible to determine t_r and hence compute the optimum c_{t_r} by minimizing the objective function $h(t; c_0, c_1)$.

Example: c_{t_b} and c_{t_r} splits ranks of nodes better



Definition 4: Kendall's tau-b

Given normalized centrality measures c_0 and c_1 , the Kendall's tau-b of the two ranks induced by c_0 and c_1 on a graph G on n vertices is

$$\tau_b(c_0, c_1) = \frac{m_C - m_D}{\sqrt{(m - T_0)(m - T_1)}},$$
(2)

where m = #pairs of distinct nodes in $G = \frac{n(n-1)}{2}$, $m_C = \#$ concordant pairs in G with respect to c_0 and c_1 , $m_D = \#$ discordant pairs in G with respect to c_0 and c_1 , $T_0 = \#$ pairs of distinct nodes in G tied on c_0 , $T_1 = \#$ pairs of distinct nodes in G tied on c_1 .

Definition 5: Pearson's r Given normalized centrality measures c_0 and c_1 , the Pearson's r between the two ranks induced by c_0 and c_1 on a graph G on n vertices is

$$r(c_0, c_1) = \frac{n(\sum_{i=1}^n c_0(v_i)c_1(v_i)) - (\sum_{i=1}^n c_0(v_i))(\sum_{i=1}^n c_1(v_i))}{\sqrt{[n\sum_{i=1}^n (c_0(v_i))^2 - (\sum_{i=1}^n c_0(v_i))^2][n\sum_{i=1}^n (c_1(v_i))^2 - (\sum_{i=1}^n c_1(v_i))^2]}}.$$
(3)

7. Conclusion

In conclusion, we see different centralities emphasise different aspects. To construct a convex combination, we need it to represent all c_i 's, where we used correlation methods. Finally, we demonstrated how to use Kendall's tau-b and Pearson's r to find the optimal t for c_t .

0.0175 - 0.0150 - 0.0	0.2	0.4 0.6 t	0.8	1.0	0.001 -	0.2	0.4 t	
Figure 2 Figure 1	. g(t	$c; c_0, c_1)$ f	or		Figure 3 Figure 1	8. h($t; c_0, c_1$) for



Table 2. Ranking of nodes by $c_0, c_1, c_{t_b}, c_{t_r}$ on Figure 1

When $c_0, c_1 =$ closeness, betweenness centrality,

■ $g(t; c_0, c_1)$ minimized at $t_b \approx 0.1959$, $h(t; c_0, c_1)$ minimized at $t_r \approx 0.3013$ ■ $c_{t_b} = 0.8041c_0 + 0.1959c_1$, $c_{t_r} = 0.6987c_0 + 0.3013c_1$ ■ $\min\{\tau_b(c_0, c_{t_b}) \approx 0.9271$, $\tau_b(c_1, c_{t_b}) \approx 0.8997\} \ge \tau_b(c_0, c_1) \approx 0.8315$, $\min\{r(c_0, c_{t_b}) \approx 0.9795, r(c_1, c_{t_b}) \approx 0.9795\} \ge r(c_0, c_1) \approx 0.9188$

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UNSTEADY MAGNETOHYDRODYNAMIC (MHD) FLUID FLOW OVER A STRETCHING SHEET WITH VISCOUS DISSIPATION AND HEAT SOURCE EFFECTS



SIN3015 MATHEMATICAL SCIENCE PROJECT MOHAMAD ADAM DANIAL BIN SHAHRIMAN (U2004713) SUPERVISOR: DR KOHILAVANI NAGANTHRAN

Abstract

This study analyzes the unsteady magnetohydrodynamic (MHD) boundary layer flow and heat transfer of a fluid over a stretching sheet in the presence of viscous dissipation and heat source. By using the similarity transformation, the governing non-linear partial differential equations are transformed into a solvable form, which is ordinary differential equations, before being solved numerically by a collocation method using the bvp4c function in MATLAB software. The influence of physical parameters such as the local Nusselt number and skin friction coefficient on the developed model is presented and discussed. The impacts of various physical parameters on the dimensionless temperature profiles are graphically illustrated and analyzed. The graph should be asymptotic to the x-axis as the similarity variable increases. From the result, we could see that as the magnetic parameter M increases, the skin friction coefficient, increases while the Nusselt number decreases.

Research Objectives

To construct a mathematical model which describes the unsteady two-dimensional laminar boundary layer flow past a continuously stretching sheet immersed in an incompressible electrically conducting field.

- To solve the mathematical model by using the collocation method (bvp4c function) available in MATLAB software.
- To analyze the transport phenomena in the fluid regime when the governing parameter varies in terms of the physical quantities.

Literature Review

Boundary layer flow over a stretching sheet

- Sakiadis (1961) analyzed boundary layer flow on a continuously stretching surface at a constant speed.
- Chen and Char (1988) analyzed laminar boundary layer flow and heat transfer on a stretching sheet with various thermal conditions.

2. Magnetohydrodynamics (MHD)

- In 1940s, geophysicists theorized that Earth's magnetic field was generated by the dynamo action in its liquid-metal core, an idea first proposed in 1919 by Larmor for the Sun.
- In the 1950s, plasma physicists became interested in MHD for controlled thermonuclear fusion, focusing on the stability of magnetically confined plasmas.

3. Unsteady Flow

- Stokes has laid the foundation for the Navier-Stokes equations, which describe the motion of viscous fluids and are fundamental to the analysis of unsteady flows.
- Ludwig Prandtl developed the concept of the boundary layer and made significant advances in the understanding of unsteady flow phenomena, particularly in the context of aerodynamics.
- Unsteady flow is characterized by flow parameters at any point that vary with time.

4. Viscous Dissipation

- Morini (2013) stated that the irreversible process by means of which the work done by a fluid on adjacent layers due to the action of shear forces is transformed into heat is defined as viscous dissipation.
- Stokes (1851) studied the motion of spheres through a viscous fluid, leading to the formulation of Stokes' law, which describes





- An increment in *Ec* infers that the viscous dissipation in the boundary layer becomes more significant. Viscous dissipation refers to the conversion of mechanical energy into thermal energy due to the internal friction within the fluid. Thus, the fluid temperature increases as *Ec* increases from 0.1 to 3.
- From the table, we can observe that when parameters are fixed at M = 2, A = 7, R = 1, y = 1, as Ec increase, the local Nusselt number decreases as stated by Reddy et al. (2015).



- The heat source parameter represents the strength of the internal heat generation or absorption within the boundary layer. As the heat source
 parameter increases, the internal heat generation within the boundary layer also increases. Thus, the fluid temperature increases as
 γ increases
 from 0.1 to 0.4.
- It is observed that as the heat source parameter γ increases, the local Nusselt number decreases as proved by Reddy et al. (2015).
- The decreased temperature gradient between the stretching sheet and the fluid, caused by the increased fluid temperature, leads to a reduction
 in the convective heat transfer from the stretching sheet to the fluid.



Figure 3: Temperature profiles for different values of M.

- The magnetic parameter represents the strength of the applied magnetic field perpendicular to the flow direction. As the magnetic parameter increases, the Lorentz force acting on the electrically conductive fluid within the boundary layer also increases. The decreased fluid velocity within the boundary layer due to the increased Lorentz force leads to a reduction in the convective heat transfer from the stretching sheet to the fluid. Thus, when *M* increases the fluid temperature increase.
- We could see that as the magnetic parameter M increases, the skin friction coefficient increases while the local Nusselt number decreases.
- The Lorentz force introduced by the increased magnetic parameter acts to oppose the fluid motion within the boundary layer. This affect the
 wall shear stress to rise and results in an increase of -f^{or}(0).
- The decreased fluid velocity within the boundary layer due to the increased Lorentz force leads to a reduction in the convective heat transfer from the stretching sheet to the fluid. This affect the fluid temperature to rise and results in the decrease of $-\theta'(0)$.



The radiation parameter represents the contribution of thermal radiation to the overall heat transfer in the boundary layer. The fluid within the
 boundary layer absorbe the incoming radiation heat leading to an increase in the internal energy of the fluid. This additional heat absorbed he

the drag force experienced by a sphere moving through a viscous fluid.

5. Heat Source/Sink

- In the context of heat transfer, a heat source adds thermal energy to a system, increasing its temperature, while a heat sink removes thermal energy, decreasing the temperature.
- According to West (2014), Joseph Black was a pioneer in careful measurements of heat transfer in 1761 who noticed that the change of state can occur over a prolonged period when a substance is heated or cooled without a change in temperature.

Methodology

This section explains how viscous dissipation and heat sources affect unsteady MHD flow over a stretching sheet. It derives the governing equations which are continuity equation, momentum equation and energy equation and uses similarity transformation to turn them into ordinary differential equations. The chapter also discusses using the bvp4c function in MATLAB to compute numerical solutions, which helps understand how the governing parameters change. boundary layer absorbs the incoming radiative heat, leading to an increase in the internal energy of the fluid. This additional heat absorbed by the fluid through radiative heat transfer causes the fluid temperature to rise. Thus, the fluid temperature increases as R increases from 1 to 4. We could see that as the radiation parameter R increases, the local Nusselt number decreases, as stated by Reddy et al. (2015).

Conclusion

- Temperature profiles decrease with an increase in the magnetic parameter, M.
- · Temperature distribution increases as the radiation parameter, Eckert number, and the heat source parameter increase.
- The magnitude of the skin friction coefficient increases with an increase in magnetic parameter whereas the magnitude of the Nusselt number decreases.
- The heat transfer rate at the surface decreases with an increase in the radiation parameter, magnetic parameter, Eckert number and heat source parameter.
- · The contribution of the present work has rectified the mistake which has been made by Reddy et al. (2015).

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ASSESSMENTS OF MACHINE LEARNING REGRESSION MODELS FOR PREDICTING THE DEVELOPMENT OF THE MIDFACE IN SYNDROMIC CRANIOSYNOSTOSIS

HaoRan Xin S2106630



UNIVERSITY MALAYA

Introduction

Syndromic Craniosynostosis is a congenital condition in which the sutures of the skull fuse prematurely, resulting in abnormal head and face shape. In order to develop a targeted personalized treatment plan, the prediction of facial growth patterns is particularly important. By examining the accuracy of a specific regression model for the face measurement, it provides the basis for personalized and precise treatment programs. This study aimed to determine the effectiveness of three regression modelsleast square linear regression, support vector machine (SVM) regression, and random forest-in predicting the middle facial area of Syndromic Craniosynostosis. Identify the most reliable models for high-precision prediction in clinical applications.

Analysis Methodology

Least Squares Linear Regression

Least squares linear regression minimizes the sum of squared errors to fit a line through the data, simplifying the analysis when the variables have an almost linear relationship.

Support Vector Machine Regression

SVM regression uses kernel functions to map input features to higher dimensional Spaces, thus achieving linear separation and suitable for handling complex interactions

Random Forest

The random forest builds multiple decision trees and averages their predictions to reduce errors and improve confidence. Its efficient handling of nonlinear relationships and complex interactions makes it suitable for predicting high-precision midplane measurements.

Random Forest Results

Variable	Mean Squared Error (MSE)	R-squared (R ²)
ZMs-ZMsR	6.698	0.877
N-ANS	6.042	0.992
ANS-PNS	10.123	0.836
ZF-ZFR	18.989	0.904
ZMs-ZTi	13.462	0.934

The outcome of Random Forest Regression indicates the sufficiently good seemingly in each dependent variable, with the range of R-squared (R²) values fluctuation from 0.836 to 0.992. The N-ANS factor is depicted as bearing a value of 0.992 at R², which could mean that the model explains 0.992. And the N-ANS variable MSE also comes out low in correlation





Least Squares Linear Regression Results

Variable	Mean Squared Error (MSE)	R-squared (R ²)
ZMs-ZMsR	6.484	0.906
N-ANS	12.840	0.888
ANS-PNS	8.815	0.860
ZF-ZFR	20.022	0.870
ZMs-ZTi	13.115	0.835

ZMs-ZMsR and ANS-PNS are the most suitable predictors because they have high R² values and relatively low MSE values, indicating that the model has strong explanatory power for these variables and small prediction errors.

ZMs-ZMsR Function:

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ZMs-ZMsR=0.42(S-N)-1.62(N-SO)-0.46(SO-BA)-0.15(N-S-BA)+0.05(S-SO-BA)-21.38
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ANS-PNS Function:

ANS-PNS=0.12(S-N)+1.74(N-SO)+0.91(SO-BA)+1.15(N-S-BA)+1.24(S-SO-BA)+45.01

ZF-ZFR Function:

ZF-ZFR=0.15(S-N)-0.42(N-SO)-0.22(SO-BA)+0.19(N-S-BA)+0.24(S-SO-BA)-0.54

ZMs-ZTi Function:

ZMs-ZTi=-0.05(S-N)-0.13(N-SO)-0.04(SO-BA)-0.06(N-S-BA)+0.01(S-SO-BA)-48.03

These formulas are derived by linear regression model, they describe the linear influence of different independent variables on the target variable, and can be used to predict the corresponding target variable value.

Actual vs Predicted: ZMs_ZMsR 60 ZMsR 20 ZMs ed 45 44 46 48 Actual ZMs ZMsF LEAST SQUARES LINEAR REGRESSION ACTUAL VS PREDICTED VALUES FOR ZMS ZMSR At the point where the X-axis is close to 40, the predicted value is significantly lower than the actual value, and the positive residual is larger in the corresponding residual plot. For points where the X-axis is close to 60, the predicted value is significantly higher than the actual value, and the negative residual is larger in the corresponding residual plot. The residual error in the middle region is small, indicating that the model is more accurate in this interval.



FOR ZMS_ZMSR The residual distribution is relatively random,

In the least square linear regression model, ZMs-ZMsR is chosen as the best predictor. In order to compare the performance of different models on the same variable, we also use the ZMs-ZMsR variable for analysis in the random forest model.Although not as good as N-ANS, it still has higher explanatory power and lower prediction error.The random forest model shows high prediction accuracy when predicting the variable "ZMs_ZMsR".Most of the data points are close to the ideal 1:1 relationship line, indicating that the random forest model can fit the data well and provide accurate predictions. Most of the residials are between -0.5 and 1.5, showing that the random forest model's predictions of the "ZMs_ZMsR" variable are very accurate with very little error. While there are some positive and negative residuals near the predicted values of 40, 48, and 54, the values are very small, showing that the model's prediction error in these intervals is also small.The random forest model performs well in predicting the "ZMs_ZMsR" variable, with small residual values, relatively random error distribution, and no obvious systematic bias. and there is no obvious systematic bias, indicating that the error of the model is random in most of the predicted values. Large residuals appear at the extremes of the predicted values (near 45 and 60), suggesting that the model has a large error problem in these intervals.

Support Vector Machine (SVM)regression Result

Variable	Mean Squared Error (MSE)	R-squared (R ²)
ZMs-ZMsR	56.512	0.178
N-ANS	92.9762	0.191
ANS-PNS	45.671	0.274
ZF-ZFR	130.316	0.150
ZMs-ZTi	66.714	0.160

The mean square error (MSE) values of all variables are relatively high, which indicates that the SVM model has a large prediction error on these variables.

All variables had low r-squared (R²) values, indicating that the SVM model had limited ability to explain these variables.

These conditions indicate that the model cannot effectively predict these mid-face measurement parameters in the current

Future Research Directions

 For future research, the parameters of the random forest model can be further optimized and more relevant variables can be included to further improve the prediction accuracy.
 Removing features with high VIF values or dimensionality reduction through principal component analysis (PCA) can reduce the influence of multicollinearity on the model and improve the stability and prediction accuracy of the model.

3. Use regularization methods such as Ridge regression and Lasso regression to introduce penalty terms and reduce multicollinearity effects

4. Hyperparameter adjustment of SVM , such as adjusting regularization parameters, can significantly improve the model's prediction performance.

5. Data expansion: Expand and enrich the data set by collecting more patient data, especially patient data of different age groups and different disease severity.

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Summary and Conclusion

Least square linear	Support vector machine	Random forest regression
regression	regression (SVM)	
It performed well in predicting the	It performed poorly in predicting the	It performed well in predicting the "ZMs-ZMsR"
"ZMs-ZMsR" variable with an MSE of	"ZMs-ZMsR" variable, with MSE of 56.512	variable, with MSE of 6.698 and R ² of 0.877,
6.484 and an R ² of 0.906, showing	and R ² of only 0.178, showing large	showing good predictive performance and
trong explanatory power and a small	prediction error and weak explanatory	explanatory power.
prediction error.	power.	The prediction of extreme values is stable and
lowever, it does not perform well on	It performs poorly on all variables and	the residual is small.
extreme values, and the residual is	may require further tuning and	It also performs well on other variables and is
large.	optimization.	suitable for data with complex and non-linear
-		relationships.

The prediction of extreme value by Random forest is stable and the residual is small. It also performed well on other variables. And Random forest model hasn obvious advantages in dealing with data complexity and multicollinearity, and is the best predictive model among the three models. Its the best choice for predicting facial development in patients with Syndromic Craniosynostosis

Predicting Midface Development in Syndromic Craniosynostosis Using Penalized Regression Models in Machine Learning



WANG JIALE

Affiliations

Institute of Mathematical Science, University of Malaya, Kuala Lumpur, Malaysia

	age	ANS-PI	NS N-ANS	ZMs-ZM	[s]N-S-BA	N-SO	S-N	SO-BA	S-SO-BA	A ZF-ZFR	ZMs-ZTi	
	15y 10m	52.11	54.87	54.44	142.09	78.62	66.65	22.13	137.83	98.26	49.56	
Introduction	11.3m	33.81	29.79	42.13	147.15	58.75	49.65	14.21	150.00	79.56	34.69	The dataset for this study
	9y 6m	44.65	47.35	50.59	128.24	67.50	59.74	18.07	145.26	91.94	44.70	consisted of 21 nationts arch.
Syndromic craniosynostosis is a aroup of rare genetic disorders characterized	4y 11m	38.50	36.46	46.16	139.48	70.21	59.21	14.57	145.47	82.97	44.10	consisted of 24 putients, euch
	5y 10m	43.35	46.17	43.58	134./1	64.32	55.77	21.09	153.20	80.58	44.05	representing a patient diagnosed
by the premature fusion of one or more of the cranial sutures, the joints	4y 5m	37.49	28.77	50.25 40.79	137.81	04.12 57.01	22.77 10.23	10.05	140.08	87.14 78.54	36.50	
botwoon the bonce of the ekull. This prometure fusion can load to an	1 75v	35.37	31.97	40.79	141 19	60.98	50.06	14.34	163.36	77.17	37.48	with sydromic craniosynostosis.
between the pones of the skutt. This premuture rusion cuntedu to un	3v 3m	36.91	34.06	47.59	137.61	61.54	54.37	16.48	137.04	83.11	37.94	The response variables include
abnormal head shape and, in some cases, associated physical and coanitive	4.6m	30.92	28.93	35.66	149.76	51.24	41.93	13.10	156.28	73.66	32.54	
	3y 8m	38.83	39.69	49.22	144.05	68.46	57.99	17.64	146.36	87.47	45.47	'ZMs-ZMsR', 'N-ANS', 'ANS-PNS',
impairments. I reatment for syndromic craniosynostosis typically involves a	4y 10m	40.66	41.67	49.31	140.74	70.10	58.29	17.61	165.75	89.82	44.70	'ZE ZED' 'ZNA ZT:' likely derived
multidisciplinary approach with input from pourosurgeons craniofacial	13y 2m	44.46	48.02	55.52	141.97	71.99	60.19	20.24	134.89	101.31	52.05	ZF-ZFR, ZIVIS-ZTT, LIKELY derived
mattaisciptinary approach, with input normed osargeons, cramoraciat	9m	32.88	28.85	38.08	141.28	55.27	45.82	12.95	153.39	72.80	33.31	from anatomical or physiological
surgeons, geneticists, and other specialists. The primary goal is to allow for	10y 1m	45.85	50.36	50.30	135.22	71.60	62.41	22.54	139.08	89.95	49.45	nonnanaconnoac or priyolocogioac
	2y 5m	36.66	34.97	44.47	135.25	65.42 54.20	56.13	14.46	150.39	80.59	35.86	measurements, used for
normal brain growth and development by surgically correcting the abnormal	10.8m 17m 13d	35.00	29.57	37.20	141.10	57.02	44.97	12.00	152.75	75.13	32.29	annoning conditions or
skull shape Additional therapies may address other medical issues that	1/m 13u 1.2v	31.87	30.55	42.73	138.83	58 59	47.85	12.67	133.13	75.15	35.01	assessing conditions of
Skall shape. Additional therapies may address other medioat issues that	9v 4m	42.61	44.93	51.85	134.80	70.82	62.30	20.17	140.16	92.40	46.43	anatomical relationships.
arise.	10y 3m	43.33	46.14	55.92	145.64	74.26	59.73	17.67	157.32	100.53	52.79	
	8.3m	33.16	28.15	38.74	144.95	55.75	46.79	13.96	153.75	73.88	34.35	
	6y 4m	41.63	43.29	46.33	138.96	63.99	55.26	20.68	148.36	91.63	51.14	
	15y 4m	47.13	57.86	56.70	135.50	84.57	63.02	17.02	169.01	98.92	50.07	

Objective

The primary objective of this study is to compare three regression models-Least squares linear Regression, Ridge Regression, and Lasso regression – for predicting midface region measurements based on cranial base parameters.

Methodology

Data Description Multicollinearity Check (VIF Test) Normality Test (Shapiro-Wilk Test) Analysis Methodology (1) Least Squares Linear Regression (2) Ridge Regression (3) Lasso Regression

Multicollinearity is a common problem in regression analysis where the predictors are highly correlated. It was to detect a degree of multicollinearity which is the correlation between independent variables. We will perform the Shapiro-Wilk Test to check their normality. This determines whether the variable is suitable for linear regression analysis. The Shapiro-Wilk test is used to test the residuals of each model and to test the normality of the distribution, which is the basic assumption of linear regression analysis

Results

Variable	Mean Squared Error (MSE)	R-squared (R ²)	Variable	Mean Squared Error (MSE)	R-squared (R ²)	Variable	Mean Squared Error (MSE)	R-squared (R ²)
ZMs-ZMsR	6.48	0.91	ZMs-ZMsR	5.81	0.80	ZMs-ZMsR	1.95	0.93
N-ANS	12.84	0.89	N-ANS	5.90	0.93	N-ANS	2.76	0.97
ANS-PNS	8.82	0.86	ANS-PNS	6.23	0.88	ANS-PNS	9.08	0.83
ZF-ZFR	20.02	0.87	ZF-ZFR	6.11	0.94	ZF-ZFR	3.90	0.96
ZMs-ZTi	13.12	0.84	ZMs-ZTi	10.27	0.75	ZMs-ZTi	4.33	0.89

Least Squares Linear Regression

The Mean Squared Error (MSE) for the predictions is approximately 6.78.

MSE is a measure of the average squared difference between the estimated values and the actual value. The R-squared (R^2) value is about 0.856. This value represents the proportion of the variance for the dependent variable that's explained by the independent variables in the model. An R² value closer to 1 indicates a model that explains a large portion of the variance.

Ridge Regression

The mean square error (MSE) value represents the mean square error between the estimate and the actual value. The greater the MSE value, the greater the prediction error. In this analysis, the mean square error value ranges from 5.81 to 10.27, and there is a significant difference in prediction accuracy between different variables. The R Squared value represents the proportion of the difference in the dependent variable that can be predicted by the independent variable. The R Squared value is between 0.75 and 0.94.

Lasso Regression

The Lasso Regression results show that each dependent variable is superficially good enough, and the R Squared value fluctuates from 0.836 to 0.992. The N-ANS factor has a value of 0.97 at R Squared, which probably means that the model explains 97%. The correlation of the ZMs-ZMsR variable is also low (R²=0.93). Although this value is reduced from the original 100%, the model still describes the proposed phenomenon fairly well.



Discussion

We evaluated the Least Squares Linear Regression, Ridge Regression and Lasso Regression, revealed large differences in the predictive power of facial measurements in patients with Sydromic Craniosynostosis. The Lasso Regression, which has unique features for handling multicollinearity problems and variable selection, has an R² of 0.97 for N-ANS, suggesting that the model can explain 97% of the variance. Under the least squares model, VIF values can indicate problems of high multicollinearity. Processing multicollinearity with Ridge Regression and Lasso Regression can reduce the effect of insignificant predictors, thereby improving the reliability of the model.

Conclusion

The primary objective of this study was to compare the performance of three regression models–Least Squares Linear Regression, Ridge Regression, and Lasso Regression-in predicting midface region measurements in patients with syndromic craniosynostosis. Our analysis demonstrates that the choice of regression model significantly impacts the accuracy and reliability of predictions, with Lasso Regression showing the most promising results. The lowest MSE and highest R-squared values obtained with Lasso Regression indicate its superior capability in predicting midface development in patients with syndromic craniosynostosis. This makes it an invaluable tool for clinicians involved in treatment planning and intervention strategies.

Forecasting Ricketsiosis Case Numbers in Malaysia: Analysis of Count Time Series Following Generalized Linear Model

By: WANG CAN S2037359

What are Rickettsia and Rickettsiosis?

Rickettsia are small, obligately intracellular Gramnegative bacilli. They are distributed among various hematophagous arthropod vectors and cause Rickettsioses, an acute undifferentiated febrile illness, and are often accompanied by headache, myalgias, and malaise.

Cases of Rickettsiosis in Malaysia



As shown in Figure 1, we can see a clear trend of increasing Rickettsiosis cases over the years from May 2016 to December 2019.

Additionally, Figure 2 shows the distribution of cases by state. From this pie chart, we can see that a significant proportion of cases are concentrated in East Malaysia, particularly in the states of Sabah and Sarawak.



The Seasonal Autoregressive Integrated Moving Average (SARIMA) model is an extension of the ARIMA model that supports time series data with a seasonal component. It combines autoregressive (AR), differencing (I), and moving average (MA) components to capture both the seasonal and non-seasonal behavior of the time series.

Components of SARIMA:

The SARIMA model is represented as:

ARIMA (p, d, q) (P, D, Q) m Uppercase letters Lowercase letters





Residual Diagnostics (SARIMA)



The ACF plot indicates that most residuals fall within the confidence intervals, suggesting no autocorrelation.

The histogram reveals that the residuals are approximately normally distributed but exhibit a slight skew and a few extreme values, indicating the presence of outliers. The residual diagnostics suggest that the SARIMA(0,1,1)(0,1,1)[12] model provides a reasonable fit to the data.

Forecast Value Vs. Actual Value (SARIMA)



Why would we want to develop a GLM?

The monthly cases of Rickettsiosis form a count time series with low counts, ranging from 2 to 27. As noted by Stephan (2016), such series are not well-suited for methods designed for continuous distributions. Traditional SARIMA methods, which assume a continuous sample space, are therefore inappropriate for this type of data. Given the absence of a generalized approach, we decided to investigate a specific model for the data. Our analysis focuses on the Generalized Linear Model (GLM), which is particularly effective for this purpose, and we compared its performance against SARIMA models.

GLM

- · Generalized Linear Models (GLM): is the most common model for analyzing count data
- · Compared to the usual GLM, the innovative GLM we used from Tobias Liboschik
- considers the empirical autocorrelation of the count time series data.
- · It can be defined in the following form: $g_{(\lambda_t)} = eta_0 + \sum_{i} eta_k \widetilde{g}(Y_{t-ik}) + \sum_{i} lpha_l g(\lambda_{t-jl}) + \eta^T X_t$

Fit GLM Model

R > summary(fit)

Call:

tsglm(ts = train timeseries, model = list(past obs = c(1, 12)),xreg = train regressors, link = "log", distr = "nbinom")

Residual Diagnostics (GLM)

ACF of response residuals

Capture short-term serial dependencies through a first-order autoregressive term and annual seasonality twelfth-order through a autoregressive term, both specified in the model parameter list element named 'past_obs'. Due to the more severe infections in East Malaysia, we include the number of cases in East Malaysia as a covariate. For distribution, we chose the negative binomial. Due to our data being discrete and not satisfying the Poisson distribution assumption (variance equals mean).

- (p, d, q): These parameters represent the non-seasonal part of the model.
- p: Non-seasonal autoregressive order.
- It indicates the number of lag observations included in the model.
- d: Non-seasonal differencing order.
- It indicates the number of times the raw observations are differenced to make the time series stationary
- q: Non-seasonal moving average order.
- It indicates the size of the moving average window used to smooth the time series.

Years	Manth	Positive	Years	Month	Pestler
21210	May-30	4	N 197	Nac18	17
	Am 16	3		Apr 38	12
	34.16			Netti	2
	Aug 15	2		Jun 18	
	Sep-10	3		266-38	
	Oct-28-	2		Aug-18	
	Mov-TD	- 6		Sep.18	
	Dec/16	3		03.18	3
7037	Jan 17	4		Nor 15	. 3.
	Feb:17	8		Day:18	12
	Mar-17	3	2019	3ar-19	35
	Apr 17	2		1eb-19	25
	Hay-17	4		Har:19	27
	349.17	7		Apr.35	
	34.17	7		May 19	-128
	Aug:17	2		30+19	13
	Sep:17			34-19	25
	061-17	- 6		Aug-19	30
1	Mon-17	X.		5ep 10	13
	fbx:17	4		0.1.21	-11
anan.	Jan 36			Nov:10	310

Seasonal Check

The radar plot shows the monthly distribution of cases from 2016 to 2019. **Seasonality Detection:** The plot indicates a recurring peak in March across multiple years, suggesting the presence of seasonal patterns in the data. Given the observed seasonality, we could train a SARIMA model to capture these seasonal effects for more accurate forecasting.

Fit SARIMA Model

1.Split train and test set:

2. Model Selection

To choose the best model, we used three information criterias: AIC, AICc, and BIC. They are information criteria to measure the goodness of fit of a statistical model. Lower values mean that the model fits better.

3.SARIMA Model

(p, d, q) (P, D, Q) m

Model: SARIMA(0,1,1)(0,1,1)[12]

- No autoregressive terms, so p = 0 and P = 0.
- One differencing term (d = 1) and one seasonal differencing term (D = 1).
- One moving average term (q = 1) and one seasonal moving average term (Q = 1).

Model

SARIMA(0,1,

• A seasonal period of 12 months (m = 12).

- (P, D, Q): These parameters represent the seasonal part of the model. P: Seasonal autoregressive order.
- It captures the relationship between a value and its seasonal lags.
- D: Seasonal differencing order.
- It helps to remove seasonal trends in the data, making it easier to model. Q: Seasonal moving average order.
- It captures the relationship between a value and past forecast errors from seasonal lags



Monthly Rickettsiosis Case Numbers in Malaysia

- Study Area :
- **Includes 8 states in Malaysia**
- Time Period :
- May 2016 ~ December 2019
- (Total of 44 observations)
- Data Type: **Discrete Time Series**



Train Set: 2016 May ~ 2019 Jun Test Set: 2019 Jul ~ 2019 Dec

Table	l. Informat	tion Criteria	
	AIC	AICc	BIC
1)(0,1,1)[12]	159	160	162

SARIMA(0,1,1)(0,0,1)[12] 219 220 224 SARIMA(1,1,0)(0,0,1)[12] 219 220 224 As shown in Table 1, our best model is

SARIMA(0,1,1)(0,1,1)[12], which has the lowest AIC, AICc, and BIC scores.



The ACF plot helps identify any correlations between residuals over different time lags. All autocorrelations lie within the 95% confidence bands (blue dashed lines), suggesting that the residuals do not exhibit significant autocorrelation. It shows that does not exhibit any autocorrelation or seasonality which has not been taken into account by the model. This is a desirable property indicating a good model fit.

Forecast Value Vs. Actual Value (SARIMA)



The forecasts (red lines) have generally trended in line with the actuals and in most months the forecasts have almost accurately predicted the number of cases, e.g. September and October 2019. But there are still months that deviate significantly from the actual values, which is common because our prediction horizon, i.e. the test set, is too short. We can look forward to the predictive performance of this model over longer time periods.

omo	son	

Table 2. Forecast Accuracy	Comparison of SARIMA and
----------------------------	--------------------------

Model	RMSE	MAE	MPE	MAPE
SARIMA	4.945685	4.015738	-12.67336	26.86456
GLM	4.103619	3.543842	-0.1469728	23.89788

lusion: the table 2, The GLM rforms the SARIMA in all measures. mendation: recasting Rickettsiosis umbers in Malaysia, LM is preferred due to her accuracy and its ility for discrete data.

Lower values indicate that the model performs better.

· Accuracy: GLM model shows lower RMSE and MAE, indicating higher accuracy compared to the SARIMA model.

- · Prediction Bias: GLM's MPE is closer to zero, suggesting it has less prediction bias.
- nonstrates better performance in percentage error terms.





The public health department can use statistical models to develop more precise prevention and control plans for rickettsiosis.

03

	 Error Percentage: GLM's lower 	MAPE den
n	clusions	
	74 hills	
	01	

Comparative Analysis of ARIMA Model and Generalized Linear Model for Predicting MRSA Infection Cases in Malaysia



VANG CU S2100923

Introduction

Methicillin-resistant Staphylococcus aureus (MRSA) bacterium is a significant hospitalacquired pathogen. It resists many antibiotics that treat regular staph infections, posing a serious public health risk . *Staphylococcus* skin infection usually begin as swollen, painful red bumps that may resemble pimples or spider bites. Sometimes the bacteria stay on the skin, but they can also go deeper and cause serious infections in bones, joints, surgical wounds, the bloodstream, heart valves, and lungs.

Data Background



Motivation for the Study

The monthly infections number of MRSA is a typically count time series with low counts. According to Stephan (2016), This kind series can no longer be dealt with using the approximative methods that are appropriate for continuous distributions. Given this requirement, it is clear that traditional ARIMA methods are inappropriate as they assume a continuous sample space for the data. In the absence of a generalized methodology, we are interested in considering some specific classes of models and noting the suitability of particular models for the count data. As a result, the Generalized Linear Model (GLM), which is the most common model for analyzing count data, became the model we wanted to analyze and compare with ARIMA.



01. Fitting and forecasting with Auto Regression Integrated Moving

3. Model Selection

Table 1. Information Criteria					
Model Name	AIC	AICc	BIC		
ARIMA (1,2,1)	143.6286	145.2286	146.4619		
ARIMA(0,2,1)	144.6893	145.4393	146.5782		
ARIMA(1,2,0)	151.1609	151.9109	153.0498		
ARIMA(0,2,0)	159.0685	159.3038	160.0129		

ARIMA (p=1,d=2,q=1)



Parameters P=1,Q=1 in this model means that the model uses 1 lagged data and one prediction error to make predictions.

d=2: minimum number of differencing needed to make the series stationary.

Step2:Residual Diagnosis





Figure 3: ACF Plot of ARIMA(1,2,1) models residual Figure 4: Normal Q-Q Plot of ARIMA(1,2,1) models

residual

• The ACF plot shows that most of the residuals are within the confidence interval, indicating little autocorrelation. • The Q-Q plot of residual shows normally distributed since the points appears as roughly a straight line.

The residual diagnostics suggest that the ARIMA(1,2,1) model provides a reasonable fit to the data.

Objective 02

Model Selection

1.AIC & QIC method

• Select the model with the smallest AIC and QIC, the result (p=1,q=1 obtained is model with parameters **p=1 and q=1**

2.Compare the Scoring values

Model name	logarithmic	quadratic	spherical	rankprob	dawseb	normsq	sqerror
Poisson GLM	3.582194	-0.0240271	-0.1715512	4.227328	5.294492	2.20975	48.38627
Negative-Binomial GLM	3.367235	-0.0373987	-0.1933294	4.020282	4.891814	0.85000	48.38627

Table 2: Scoring values of Poisson and **Negative Binomial**

Similar to ARIMA, the parameters P=1,Q=1 in this model means that the model uses 1 lagged data and one prediction error to make predictionsThe selected distribution is Negative Binomial, it means that the model can be applied to overdispersion data.

Residual Diagnosis

Negative binomial with

Average (ARIMA) Model

02.

Fitting and forecasting with Generalised Linear Model (GLM) 03.

Compare the performance of the ARIMA model and Generalized Linear Models (GLM) in forecasting MRSA Infection Number.

Methodology

ARIMA Model

(Auto Regression Integrated Moving Average)

- Is a statistical tool used for analyzing and forecasting time series data.
- It combines differencing with autoregressive(AR) and moving average(MA) components.

The model is defined as:

$$X_t = c + \sum_{i=1}^p \phi_i X_{t-i} + arepsilon_t + \sum_{i=1}^q heta_j arepsilon_{t-i}$$

where

- p:order of the AR part
- q:order of the MA part
- $\phi i, \theta j$ are coefficients
- c is a constant
- ϵt is the error term

(Generalized Linear Model) The Generalised Linear Model (GLM), which is

GLM

the most common model for analyzing count data

The model is defined as:

$$g_{(\lambda_t)}=eta_0+\sum_{k=1}^peta_k\widetilde{g}(Y_{t-ik})+\sum_{l=1}^qlpha_lg(\lambda_{t-jl})+oldsymbol\eta^TX_t$$
 where

- { Yt : $t \in N$ } stands for a count time series • { Xt : $t \in N$ } is a time-varying r-dimensional covariate vector
- $\lambda t = E(Yt|Ft-1)$ is the conditional mean of the count time series.
- The distributions we are interested in are the Poisson distribution and the negative binomial distribution.

Similar to ARIMA

- p:order of the AR part
- q:order of the MA part

• α,β are coefficients

Resu Íscus

Step1:Fitting the ARIMA model

1.Split train and test set

Train Set: 2020 February ~ 2021 October

Test Set: 2021 November~ 2022 March

(The training set and test set account for 80% and 20% of the total data, respectively) 2. Stationary check

 ADF Test:(Null hypothesis-Nonstationary) Augmented Dickey-Fuller Test

data: train.ts_adj Dickey-Fuller = -2.5332, Lag order = 2, p-value = 0.3692 alternative hypothesis: stationary ADF Test Result of d=0

Augmented Dickey-Fuller Test

data: difference_train.ts_adj Dickey-Fuller = -4.0548, Lag order = 2, p-value = 0.02134 alternative hypothesis: stationary ADF Test Result of d=1 Augmented Dickey-Fuller Test

data: difference_train.ts_adj Dickey-Fuller = -4.0548, Lag order = 2, p-value = 0.02134 alternative hypothesis: stationary

ADF Test Result of d=2



The residual diagnostics suggest that the Negative Binomial generalized linear model provides a reasonable fit to the data.

Figure 6: ACF Plot of GLM models residual Figure 7: Histogram Plot of GLM models residue

Objective 03

Comparison

1.Compare Forecasting Plots



- ARIMA model and generalized linear model (GLM) both roughly capture the overall trend of the actual values
- Generalized Linear Model (GLM) seems better captures the upward trend of the actual

2.Compare Accuracy Values

Model name	RMSE	MAE	MAPE
ARIMA(1,2,1)	5.724014	4.716344	35.32365
Negative-Bin omial GLM	7.758253	6.855503	65.89999

Table 4 Forecast Accuracy Valus of ARIMA and GLM for MRSA Infection Cases

Both ARIMA and generalized linear model (GLM) capture the overall trend in MRSA infection numbers in two major hospitals in Klang Valley.

In comparison, generalized linear model (GLM) is more appropriate for 2 forecasting MRSA infection number.

3 Continuous data collection on MRSA infection cases in Malaysia is essential. Addressing the data's limitations is crucial for developing and analyzing more robust models in the future.

Objective 01

0

we chose the difference order d=2, the p-value is less than 0.05, means that the series is stationary.

ELECTRON SCATTERING FROM ZINC ATOM SIM3021 MATHEMATICAL SCIENCE PROJECT

SIM3021 MATHEMATICAL SCIENCE PROJECT RANJITHAA A/P VASU (U2103400)

Supervisor : Dr. Mohd Zahurin Mohamed Kamali Co-Supervisor : Dr.Shahizat Bin Amir Institute of Mathematical Sciences (ISM), Faculty Science

UNIVERSITI

MALAYA

Abstract

This study investigates electron scattering from zinc atoms, emphasizing the calculation of phase shifts and differential crosssections. The research employs the Schrödinger equation to model electron interactions, converting complex equations into a solvable format using a similarity transformation. Numerical solutions are obtained via the Runge-Kutta method solved using R studio software. Results indicate that higher electron energy increases both phase shifts and differential cross-section values, revealing significant changes in scattering amplitude and cross-section distribution. Additionally, the influence of potential energy affected by exchange and polarization effects becomes more prominent with increased electron energy, highlighting the physical parameters' critical role in scattering dynamics. These insights have implications for material analysis, radiation therapy, and semiconductor technology, providing a basis for optimizing electron scattering conditions.

Objective

a)To compute elastic scattering amplitude and differential cross-sections for electron-Zn atom interactions spanning an energy range from 25 eV to 80eV.
 b)To validate the computational approach by comparing the computed differential

cross-sections with theoretical models

c)To develop numerical algorithms for each formulated problems and programming for computations in generating numerical solutions via R Studio software.

Literature Review

1. Electron Scattering

- The study of electron scattering using cathode ray was initially founded by Sir J.J. Thomson in 1897, which is also the groundbreaking experiment in the late 19th century.
- Garcia and Wang (2024) developed a sophisticated theoretical model to mimic electron scattering in intricate molecular systems by combining quantum mechanics with machine learning algorithms.

2. Elastic and inelastic scattering

• The first investigations by Sir J.J. Thomson in 1897, which showed that cathode rays were made of negatively charged particles subsequently dubbed electrons, laid the groundwork for our understanding of electron scattering.

3. Cross-section Analysis

- In 1984, Trajmar et al. published a comprehensive compilation and critical assessment of available cross-sections for electron-molecule scattering.
- Lee and Park (2017) defined total cross-sections as the likelihood that an electron will
 experience any kind of scattering event, providing a fundamental understanding of
 these processes.
- Smith and Jones (2019) investigated differential cross-sections, shedding light on how scattering probabilities change with energy and direction

4. Electron-Zinc (Zn) scattering processes

- The optical excitation functions of Zn for many transitions have been measured in Shpenik et al. (1973) and Souter et al. (1974).
- Zn represents a quasi-two-electron atom, like helium, beryllium, and magnesium, making it valuable for testing theoretical models, especially where relativistic effects might be important (Marinković et al., 2019).



Results





Figure 5.3. Differential cross section for electrons incident at 80 eV on

bound zinc atom in the SPX approximation. The significant differences between the DCS results obtained in my study using the SPX approximation and Kelemen et al (2023)

is be largely attributed to the simplifications inherent in the SPX model. The significant differences observed can be attributed to several factors:

- The SPX model employs a straightforward combination of static, polarization, and exchange potentials. This contrasts
 with the detailed and parameterized potentials in the RSEPA and RSEP models, which include contributions from
 spin-orbit interactions and relativistic corrections.
- Kelemen et al (2023) include absorption potentials that account for inelastic scattering processes. The SPX model, however, is purely elastic and does not include such effects.
- The Kelemen et al (2023) model predicts detailed patterns in forward and backward scattering, including the impact of
 various partial waves and the inelastic contributions. The SPX model, with its simplified approach, may not fully
 replicate these detailed patterns, resulting in differences in the observed local maxima and minima, as well as
 variations in the intensity of scattering at extreme angles.
- Kelemen et al (2023) include detailed treatments of exchange and correlation effects through various potentials, such as the Furness-McCarthy (FM) potential and its modifications.
- The accuracy of the scattering computations is strongly dependent on the choice of angular momentum, especially at higher angles when the contribution of higher-order partial waves becomes greater.

Conclusion

Overall, this work serves an extension study of Kelemen et al (2023) who use complex model but in my study we used simple model. The significant differences between the DCS results obtained in my study using the SPX approximation and Kelemen et al (2023) can be largely attributed to the simplifications inherent in the SPX model. These simplifications, while beneficial for computational efficiency, result in the exclusion of complex interactions and inelastic effects that are critical for a detailed understanding of electron scattering processes. This highlights the necessity for careful consideration when selecting a model for DCS calculations, ensuring that the chosen methodology aligns with the specific goals and

Methodology

- Static field approximation, which simplifies the electron-atom interaction by assuming a static, time-independent potential.
- To solve the Schrödinger equation, the study employs the Frobenius series expansion method.
- Using Runge-Kutta, the R studio software used to numerically solve the reduced form of mathematical model.
- 4. The results are presented graphically, illustrating the relationship between electron energy and phase shifts, as well as the variation in differential cross-sections.

Figure 1: Types of scattering



accuracy requirements. In conclusion, this work of study demonstrated the fulfillment of the two objectives outline at the beginning of the study.

Acknowledgement

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A Review on General Relativity and its Application to Cosmology

Sia Chen Yi

Universiti Malaya, Institut Sains Matematik, Malaysia



Abstract

General relativity is a crucial topic that serves as a cornerstone in the revolution of our understanding of spacetime and gravity. Concepts such as dark matter, dark energy, and black holes are fundamentally based on the principles of general relativity.

Objectives

The main goals for our research are:

Einstein's Field Equation

Einstein's Approach

From Bianchi Second Identity,

$$\nabla_{\mu}(R_{\mu\nu} - \frac{1}{2}g_{\mu\nu}R) = 0, \qquad (12)$$
$$G^{\mu\nu} = R^{\mu\nu} - \frac{1}{2}g^{\mu\nu}R. \qquad (13)$$

The Universe with Cosmological Constant

• Einstein's Static Universe,

$$a_0 = \frac{1}{\Lambda}.$$
 (34)

• De Sitter Universe with $\kappa = -1$,

- Explain the derivation of Einstein's Field Equation in general relativity;
- Apply Einstein's Field Equation to construct some basic cosmological models;
- Briefly explain the origin of cosmological constant and the notion of dark energy.

Historical Overview

The story behind the Albert Einstein's work on general relativity:

- In 1905, Albert Einstein published his paper entitled "On the Electrodynamics of Moving Bodies", which introduced his theory of special relativity;
- Albert Einstein tried to seek for the natural extension of special relativity which included the effects of gravity;
- He comes out the Einstein's Equivalence Principle and realized that the space is curved;
- He sought out his classmate at Zurich's University, Marcel Grossmann to teach him tensor calculus and Riemann geometry, which describe the movement of objects in space and time and also describe the dynamic frame of reference;
- Early development of tensor calculus are done by Gregorio Ricci-Curbastro, Tulio Levi-Civita, and Professor Luigi Bianchi;
- Einstein's student David Hilbert also attempted to derive Einstein's Field Equation before Einstein. In the end, both of

Since
$$\nabla_{\mu}G^{\mu\nu} = 0$$
 and we assume $\nabla_{\mu}T^{\mu\nu} = 0$,

$$R_{\mu\nu} - \frac{1}{2}g_{\mu\nu}R = \kappa T_{\mu\nu}.$$
 (14)

After we solving for κ ,

$$R_{\mu\nu} - \frac{1}{2}g_{\mu\nu}R = 8\pi G T_{\mu\nu}.$$
 (1)

Hilbert's Approach

Our total action gravity is as follows:

$$S = \gamma S_g + \kappa S_m, \tag{16}$$
$$S = \int \mathcal{L}(\Phi^i, \nabla_\mu \Phi^i) d^n x. \tag{17}$$

We can separate
$$S_g$$
 into three parts:

$$\delta S_g = \delta S_1 + \delta S_2 + \delta S_3,$$

$$\delta S_1 = \int d^n x \sqrt{-g} g^{\mu\nu} \delta R_{\mu\nu}, \qquad (19)$$

$$\delta S_2 = \int d^n x \sqrt{-g} R_{\mu\nu} \delta g^{\mu\nu}, \qquad (20)$$

$$\delta S_3 = \int d^n x \left(-\frac{R}{2\sqrt{-g}}\frac{\delta g}{\delta g^{\mu\nu}}\right)\delta g^{\mu\nu}.$$
 (21)

 δS_1 will equals to 0 by Stokes' Theorem. After that we will get the following:

$$\delta S_g = \int ([R_{\mu\nu} - \frac{1}{2}Rg_{\mu\nu}]\sqrt{-g}\delta g^{\mu\nu})d^n x.$$
(22)
After we get δS_g , then we substitute it into (16) to get Einstein's
Field Equation,

$$a(t) = \sqrt{\frac{3}{\Lambda}} \sinh(\sqrt{\frac{\Lambda}{3}}t). \tag{35}$$

• De Sitter Universe with $\kappa = 0$,

$$a(t) = a_0 e^{\sqrt{\frac{\Lambda}{3}}t}.$$
(36)

• De Sitter Universe with $\kappa = 1$,

$$a(t) = \sqrt{\frac{3}{\Lambda}} \cosh(\sqrt{\frac{\Lambda}{3}}t). \tag{37}$$

Now we are looking at some tools which help us to relate the non-observable parameters with observable parameters.

• Redshift,

5)

(18)

(26)

(28)

(29)

(30)

(31)

(32)

(33)

$$a(t_e) = \frac{1}{1+z}.$$
 (38)

• Instantaneous Physical Distance,

$$d_p(t) = a(t) \int_0^r \frac{dr}{\sqrt{1 - \kappa r^2}}.$$
 (39)

• Luminosity Distance Formula,

$$d_L = (1+z)a(t_0)r.$$
 (40)

them managed to derive the same equation by using different methods by November 1915;

• Einstein derived the equation by realizing Second Bianchi Identity is proportional to Noether's theorem. While Hilbert derived the equation using Lagrangian approach.

Tensor Calculus

• Covariant Tensor,

$$A_{i_1 i_2 \dots i_m} = \bar{A}_{\alpha_1 \alpha_2 \dots \alpha_m} \frac{\partial \bar{x}^{\alpha_1}}{\partial x^{i_1}} \frac{\partial \bar{x}^{\alpha_2}}{\partial x^{i_2}} \dots \frac{\partial \bar{x}^{\alpha_m}}{\partial x^{i_m}}.$$
 (1)

• Contravariant Tensor,

$$A^{j_1 j_2 \dots j_n} = \bar{A}^{\beta_1 \beta_2 \dots \beta_n} \frac{\partial x^{j_1}}{\partial \bar{x}^{\beta_1}} \frac{\partial x^{j_2}}{\partial \bar{x}^{\beta_2}} \dots \frac{\partial x^{j_n}}{\partial \bar{x}^{\beta_n}}.$$
 (2)

• Connection,

$$\Gamma^{h}_{\ ij} = x^{h}_{\ \lambda}(\bar{x}^{\lambda}_{\ ji} + \bar{\Gamma}^{\lambda}_{\ \mu\nu}\bar{x}^{\mu}_{\ i}\bar{x}^{\nu}_{\ j}). \tag{3}$$

• Christoffel Symbol,

$$\mathring{\Gamma}^{\lambda}_{\ \mu\nu} = \frac{1}{2} g^{\lambda\alpha} (\partial_{\nu} g_{\alpha\mu} + \partial_{\mu} g_{\alpha\nu} - \partial_{\alpha} g_{\mu\nu}). \tag{4}$$

• Covariant Derivative for Covariant Vector,

$$\nabla_j A_i = \partial_j A_i - \mathring{\Gamma}^l{}_{ij} A_l.$$

(5)

(6)

(8)

(9)

(10)

(11)

$$R_{\mu\nu} - \frac{1}{2} R g_{\mu\nu} = 8\pi G T_{\mu\nu}.$$
 (23)

Cosmological Constant

We

take
$$S = \gamma (S_g + S_p) + \kappa S_m$$
,
 $R_{\mu\nu} - \frac{1}{2} R g_{\mu\nu} + \Lambda g_{\mu\nu} = 8\pi G T_{\mu\nu}.$ (24)

Cosmos with only matter and radiation

• Ricci scalar of FLRW Metrics,

$$R = 6[\frac{\ddot{a}}{a} + (\frac{\dot{a}}{a})^2 + \frac{\kappa}{a^2}].$$
 (25)

Friedmann Equation,

$$(\frac{\dot{a}}{a})^2 + \frac{\kappa}{a^2} = \frac{8\pi G\rho}{3} + \frac{\Lambda}{3}.$$

Acceleration Equation,

$$\frac{\ddot{a}}{a} = \frac{\Lambda}{3} - \frac{4\pi G}{3}(\rho + 3p). \tag{27}$$

• Energy Conservation Equation,

$$\dot{\rho} = -3\frac{\dot{a}}{a}(\rho + p).$$

Conclusions

In conclusion, we have successfully derived Einstein's Field Equations by employing the Einstein-Hilbert action, a pivotal step in our journey through the realms of theoretical physics. These equations have served as the foundation for deriving several other fundamental equations, including the Friedmann Equation, Acceleration Equation, and Energy Conservation Equation, among others.

Furthermore, our exploration extended to various cosmological solutions within the framework of general relativity. We delved into the origin of the cosmological constant in Einstein's Field Equations and its reinterpretation as dark energy, based on empirical evidence such as supernova measurements. Additionally, we discussed how cosmological models are verified through the measurement of key cosmological parameters, redshift, instantaneous physical distance, and also luminosity distance formula on the intricate workings of our universe.

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• Covariant Derivative for Contravariant Vector,

 $\nabla_j A^i = \partial_j A^i + \mathring{\Gamma}^i{}_{lj} A^l.$

• Riemann-Christoffel Curvature Tensor,

 $R^{\lambda}_{\ \mu\alpha\nu} = \partial_{\alpha}\mathring{\Gamma}^{\lambda}_{\ \mu\nu} - \partial_{\nu}\mathring{\Gamma}^{\lambda}_{\ \mu\alpha} + \mathring{\Gamma}^{\lambda}_{\ \sigma\alpha}\mathring{\Gamma}^{\sigma}_{\ \mu\nu} - \mathring{\Gamma}^{\lambda}_{\ \sigma\nu}\mathring{\Gamma}^{\sigma}_{\ \mu\alpha}.$ (7)

• Ricci Tensor,

 $R_{\mu\nu} = R^{\alpha}_{\ \mu\alpha\nu},$ = $R_{\beta\mu\alpha\nu}g^{\alpha\beta}.$

 $R = R^{\nu}_{\ \nu},$

 $= R_{\mu\nu}g^{\mu\nu}.$

• Ricci Scalar,

By using Energy Conservation Equation, we do able to express energy density in terms of ω .

$$\rho = \rho_0 (\frac{a_0}{a})^{3(1+\omega)}.$$

• Einstein-de Sitter Universe,



Radiation Dominated Universe,



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IDENTIFYING SIGNIFICANT FACTORS IN HOSPITAL AND COMMUNITY-ACQUIRED MRSA USING LOGISTIC REGRESSION MODELLING



MUHAMMAD AFIQ BIN ZAINIDAR U2103651

SUPERVISORS: DR NORLI ANIDA ABDULLAH DR NUR ANISAH MOHAMED

INTRODUCTION

Methicillin-Resistant Staphylococcus aureus (MRSA) infection originates from a group of Staphylococcus bacteria that has become resistant to many antibiotics typically used to treat standard *Staphylococcus* infections. Most MRSA infections occur in people who have been hospitalized or have had contact with healthcare settings such as hospitals and dialysis centers, and are known as hospitalacquired MRSA (HA-MRSA). MRSA is also found in community clusters, such as among child care workers, athletes, and people who live in close proximity. This form is known as communityacquired MRSA (CA-MRSA).

OBJECTIVES

- 1. To investigate the association between type of gene and infection type using chi-squared test.
- 2. To model the dataset for the identification of significant factors in HA-MRSA and CA-MRSA using logistic regression (LR) model.
- 3. To evaluate the performance of the model using testing dataset.

METHODOLOGY



classified instances, while AUC assesses the model's ability to distinguish between classes.

RESULTS AND DISCUSSIONS

		HA	CA	
	sea	38	17	
	seb	17	7	
	sec	15	13	
	sed	5	5	
	see	44	18	
	pvl	32	33	
	hlα	370	153	
	hlв	354	146	
	fnbA	46	32	
	fnbB	28	11	
	tsst-1	82	33	
	sccmecl	13	2	
	sccmecIII	5	3	
	sccmecIV	342	139	
TYPE OF GENE	sccmecV	20	15	

The contingency table shows the distribution of virulence genes in HA and CA. The p-value of the chi-square test for the association between MRSA type and the presence of these 16 virulence genes is 0.01366, which is below the conventional significance level of 0.05. This indicates that there is significant evidence to suggest an association between MRSA type (HA or CA) and the presence of various virulence genes.

In Model 1, all predictors were included, yet none exhibited a satisfactory p-value, indicating insignificance. Subsequently, multicollinearity assessment using VIF revealed severe multicollinearity with SCCmecIV, registering a coefficient of 11.74. Thus, in Model 2, SCCmecIV was omitted. In Model 2, none of the predictors are statistically significant, and the chi-square goodnessof-fit test yields a very low p-value, indicating a poor model fit. No multicollinearity was detected in Model 2. Consequently, the backward elimination method was employed to improve the model's accuracy and interpretability in model 3.



From the confusion riv the model's

call: glm(formula = HaCa ~ pvl, family = binomial(link = "logit"),

There is only one

0 1	(Actual) (Actual)	0 (Predicted) 9 0	1 (Predicted) 3 42	accuracy is calculated to be 94.44%, indicating that the model is highly accurate.		Coefficients: Estimate Std. Error z value Pr(> z) (Intercept) 0.9050 0.1060 8.538 <2e-16 *** pvl -0.6173 0.2901 -2.128 0.0333 * Signif codes: 0 (*** 0.001 (** 0.01 (** 0.05 (* 0.1
01 08			sh M be	The AUC for the ROC curv nown is 0.875, which means odel 3 has an 87.5% proba of correctly distinguishin etween HA-MRSA and CA-M	ve s that ibility ig IRSA.	(Dispersion parameter for binomial family taken to be 1) Null deviance: 602.00 on 489 degrees of freedom Residual deviance: 597.61 on 488 degrees of freedom AIC: 601.61 Number of Fisher Scoring iterations: 4
				n conclusion, this study id with existing scientific res significant role of <i>pvl</i> usin	entifie earch. g logis	D NCLUSION es <i>pvl</i> as a key significant factor in CA-MRSA, consisten It provides robust statistical evidence supporting the stic regression model . This underscores the need for
	0.0 0.2	0.4 0.6	0.8 10			CA-MRSA infections

significant predictor variable in model 3. The odds ratio for pvl (Panton-Valentine · · 1 *leukocidin*) is 0.5394. This indicates the odds of pvl being detected in CA-MRSA are 53.94% higher than in HA-MRSA.

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Introduction

The fundamental theorem of finitely generated abelian group is frequently used to classify and study the different structure of abelian groups. It is a generalization of the fundamental theorem of finite abelian group proven by Leopard Kronecker (1870) and who defined a arbitrary set of elements with an operation satisfying certain laws which now known as finite abelian groups.

F.G. Frobenius and L. Stickelbelger (1879) proved the fundamental theorem of finite abelian groups with a group theoretical approach and formulated the finite abelian group theory to modern mathemat-ical views. Later, Poincare (1900) and Emmy Noether (1926) gave a proof for the fundamental theorem of finitely generated abelian groups which does not have the restriction on finite abelian groups.

We can use the theorem to generalize the structures of finitely generated modules over a principle ideal domain and use the theorem for other applications such as study the structure of the groups and finding isomorphic abelian groups of an given group.

Fundamental Theorem of Finitely Generated Abelian Groups

There are two forms for the theorem namely:

Invariant Factors Decomposition Every finitely generated abelian group G is isomorphic to a finite direct sum of cyclic groups in which the finite cyclic groups are in the order m_1, m_2, \cdots, m_t where $m_1 > 1$ and $m_1 | m_2 | \cdots | m_t$.

 $G \cong \mathbb{Z}_{m_1} \oplus \mathbb{Z}_{m_2} \oplus \cdots \oplus \mathbb{Z}_{m_t} \oplus F$

where F is a free abelian group.

Elementary Divisors Decomposition Every finitely generated abelian group is isomorphic to a finite direct sum of cyclic groups, each of which is either infinite or of order a power of a prime.

 $G \cong \mathbb{Z}_{p_1^{n_1}} \oplus \mathbb{Z}_{p_2^{n_2}} \oplus \cdots \oplus \mathbb{Z}_{p_t^{n_t}} \oplus F$

where F is a free abelian group, p_i are primes and n_i are positive integers.

Remark on the Theorem

- For a finite abelian group, the decomposition of the group does not contains the free abelian group part.
- If the group G is a free abelian group, then it is isomorphic to the group F stated in the theorem.

Classification of Abelian Group and Related Problems

Pong Ching Hin

Universiti Malaya

List of Nonisomorphic Groups

Using the theorem, we can get all the possible combinations of nonisomorphic abelian group of order n. For invariant factors:

- Find the prime factorization of n and list out all the possible m_t .
- For each m_t , work out all the possible combination of m_1, m_2, \ldots, m_t such that $m_1 |m_2| \ldots |m_t$.

For elementary divisors:

- Find the prime factorization of n and find the partition of each power of the prime factors.
- The direct sum of the p^k where k is the partition of the power of p is the nonisomorphic list for the prime factor. Last we just pair each combination of a prime factor to other factors to obtain the list of the abelian group of order n.

For a group of order $n = p_1^{\alpha_1} p_2^{\alpha_2} \cdots p_k^{\alpha_k}$, the number of nonisomorphic abelian group is the product of the number of partition for each α_i .

Maximum Order of Elements in an Abelian Group

The maximum order of elements in an abelian group can be determine by using the theorem and some tricks. If we know the decomposition of the group, we can find the maximum order of elements from the group by finding the lowest common multiple of the order of each cyclic group.

For example, the element in the group \mathbb{Z}_4 has element of order 4 but the group $\mathbb{Z}_2 \oplus \mathbb{Z}_2$ only has element of maximum order 2 even though both group are of order 4. This is a useful tools that we can use to determine the isomorphism.

Isomorphism Between 2 Abelian Groups



We can show that two groups are isomorphic or not by using the conditions above. As mention in the previous part, we can conclude that $\mathbb{Z}_4 \ncong \mathbb{Z}_2 \oplus \mathbb{Z}_2$ because they exist an element of order 4 in one group that cannot be mapped into the other group. Using this we can also match up the pair of invariant factors and elementary divisors from the list for order n.



Find the order of the group and find all the possible nonisomorphic group for the order.

> Eliminate the group that is nonisomorphic to the group such as non having elements of certain orders.

> > number of element of certain orders.

Note that for a group $\mathbb{Z}_{p^m} \oplus G$ is a subgroup of $\mathbb{Z}_{p^n} \oplus G$ for all integer $0 \leq m \leq n$ and G is a group.

Other Applications

- abelian group.
- generated abelian group.
- finitely generated abelian group.

- zahlen.
- Physikalische Klasse, 1926, 28–35.



Classification of Abelian Groups

By using the statement from before we can now determine the isomorphic group of any given abelian group by the following steps:



Finding the number of subgroups for a given finitely generated

• Finding number of elements with certain order for a given finitely

Investigate and classify the structure of the factors group of a

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ANALYZING URBAN DEVELOPMENT AND ACCESSIBILITY THROUGH FRACTAL DIMENSIONS OF ROAD NETWORKS IN PETALING JAYA, SELANGOR, MALAYSIA

NUR DANIA BINTI NOR AZMI

Supervised by: Dr. Shahizat bin Amir and Dr. Mohd Zahurin Mohamed Kamali



INTRODUCTION

Urbanization, a hallmark of modern society, necessitates sustainable planning as cities grow. In Petaling Jaya (PJ), Malaysia, a city of over 600,000 residents spanning 97.2 square kilometers, managing urban expansion and accessibility is crucial. Traditional analysis methods, focusing on metrics like road length and population density, often miss urban complexity. This study proposes using fractal dimensions to analyze Petaling Jaya's road networks, uncovering patterns and structures of urbanization. Fractal analysis offers enhanced connectivity insights and a visual understanding of urban form, aiding better infrastructure development and planning for improved accessibility and quality of life.

PROBLEM STATEMENT

Petaling Jaya city has complex connections of its road networks and growth patterns. Accessibility of a city is a crucial thing to its development, thus the road networks contribute a major role to determine the correlation. However, we need to use conventional methods to observe and statistically analyze road networks, its accessibility and the urban development

OBJECTIVES

• To investigate and observe road networks of Petaling Jaya and its accessibility through fractal dimensions.

• To investigate and observe how accessibility correlates to urban development through fractal dimensions.

METHODOLOGY

RESULTS



DISCUSSIONS

Petaling Jaya and Its Section Analysis

- Petaling Jaya has an overall fractal dimension of 1.8204, indicating high urbanization. All city sections have dimensions exceeding 1.5000, reflecting favorable urban conditions.
- Sungei-Way Subang (SS) has the highest complexity that is 1.7545, dedicating SS as most urbanized section.
- Petaling Jaya Utara (PJU) is the second highest of urbanization that is 1.7469 which highly accessible and well-developed.
- Seksyen has a moderate complexity with fractal dimension of 1.6473 and have increasing attractiveness and residential growth.
- Petaling Jaya Selatan (PJS) has the lowest complexity of that is 1.5530 which likely to be less accessible and urbanize but has the potential for future development.

CONCLUSION

Analyzing Petaling Jaya's road networks through fractal dimensions reveals insights into accessibility and urbanization, identifying areas for potential planning improvements. This study highlights the complexity of the PJ's road structure and suggests integrating fractal dimensions into urban planning to enhance mobility, land use, and infrastructure investments. These findings support sustainable urban growth and improve quality of life for the future

Accessibility

- Overall road network fractal dimension 1.7008 that is high in complexity
- PJS has the lowest road network dimension with 1.4990, indicating less efficient infrastructure of road.
- SS is the most accessible of all section, with highest connectivity to other sections.

Built Environment

- Overall fractal dimensions is 1.7871 with SS and PJU that are most well-developed with numerous amenities.
- Seksyen is close to optimal urbanization with 1.5968.
- PJS has lower complexity but steady population growth and potential for future development.

ACKNOWLEDGEMENT & REFERENCES







FORECASTING THE NUMBER OF MRSA CASES IN MALAYSIA USING TIME SERIES MODEL

MUHAMAD IKLIL HAKIMI BIN HASAN (U2103286)

SUPERVISOR: DR ANISAH BINTI MOHAMED @ RAHMAN DR NORLI ANIDA BINTI ABDULLAH

INTRODUCTION

Methicillin-resistant Staphylococcus aureus (MRSA) infections are rapidly spreading in Malaysia, and a particularly high number of cases are occurring. Forecasting of MRSA growth is essential for healthcare authorities to implement timely preventive measures.

This study focuses on predicting the future trend of MRSA infections in Malaysia using time series models, and it finds that the ARIMA model is the most suitable one. Through detailed analysis, ARIMA (1,2,1) was identified as the most effective model for this purpose.

The findings demonstrate that the ARIMA model can reliably forecast future MRSA cases, providing valuable insights for public health authorities to enhance their strategic planning and response efforts

OBJECTIVE

- Justify the selection of the ARIMA model based on its fit to the data compared to other models.
- Train the ARIMA model using MRSA case data from 2020 to 2022.
- Forecast future MRSA cases in Malaysia using the trained ARIMA model.
- Highlight key insights gained from the forecast.

RESULT & DISCUSSION



Utilizing historical MRSA case data collected by the National Institute of Health from February 2020 to March 2020, a time series plot was generated, comprising 26 observations. The training set includes monthly cases from February 2020 to October 2021, while the testing set spans from November 2019 to March 2022. The plot illustrates that MRSA infections in Malaysia exhibited a non-stationary trend and demonstrated an upward trajectory over the years.

Verify stationarity using the Augmented Dickey-Fuller (ADF) test. Ensure the p-value from the ADF test is less than 0.05. If the p-value is greater than 0.05, apply additional differencing to the data.



Analyze ACF and PACF plots to identify potential models based on data patterns.

Split the data into training and testing sets.

achieve stationarity.

 Use ACF and PACF to assess the suitability of AR (p) and MA (q) models and identify possible candidate models.

Transform the data to reduce variance and differentiate it to

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- Determine the parameters for potential models and select the most appropriate ones.
- Choose the best model based on the lowest AIC or BIC value.
- Verify that the residuals' ACF and PACF indicate white noise.
- Use the p-value from the Ljung-Box test or a normal probability plot to check if the residuals are normally distributed.
- Use the selected model to forecast the time series.
- Verify forecasting accuracy by comparing forecasted values with actual values.
- After validating the model, refit it using the entire dataset.

MODEL	AIC	AICc	BIC
1,2,1	143.63	145.23	146.46
0,2,1	144.69	145.44	146.58
1,2,0	151.16	151.91	153.05
2,2,1	145.05	147.9	148.82
0,2,0	159.07	159.3	160.01

These are the following 5 models that have been estimated from the ACF and PACF plots. Based on the table, it is observed that ARIMA (1, 2, 1) shows the minimum values of these measures.

It is observed that the number of MRSA cases in Malaysia from October 2021 to March are within the 80% and 95% confidence interval, which shows that the forecasting values are acceptable. TheME and MSE of the forecast values are -1.00 and 0.62, respectively, which arerelatively low. In general, the forecastresults are acceptable.

CONCLUSION

Data Limitations: Short Data Length:

• With only 26 data points, capturing longterm patterns and trends becomes challenging, which may impact the model's ability to make accurate predictions.

Potential Seasonal Patterns:

 The short length of the dataset may obscure potential seasonal patterns that could be better understood with a longer dataset.

Model Performance: Accuracy Metrics:

 The low ME and MASE values suggest the model performs reasonably well regarding

METHODOLOGY



We applied first-order differencing as the initial data was nonstationary, yet the data remained non-stationary. Consequently, we proceeded with second-order differencing. Upon reviewing the ACF and PACF plots for the second-order differencing, we observed no decay, indicating stationarity. Notably, the partial autocorrelation function displayed significant spikes at lags 1 and 2, suggesting a potential AR model order of 1 or 2. In contrast, the autocorrelation function exhibited a significant spike at lag 1, implying a possible MA model order of 1.



Month	forecast	actual	Lo 80	Hi 80	Lo 95	Hi 95
Nov-21	12.9	7	1.53	24.29	-4.49	30.31
Dec-21	12.9	12	0.74	26.46	-7.95	33.66
Jan-22	12.8	23	-3.83	29.41	-12.62	38.21
Feb-22	12.7	15	-6.28	31.74	-16.35	41.81
Mar-22	12.7	17	-8.75	34.09	-20.09	45.42

scaled error. Yet, the large deviations in forecast values indicate that further improvements are necessary.

Future Improvements: Data Enhancement:

• Increasing the dataset length by including more months or incorporating relevant external data (e.g., hospital policies and seasonal trends) can improve model accuracy.

Model Complexity:

• Exploring more sophisticated models such as SARIMA, Prophet, or neural networks could potentially yield better forecasting results, especially if there are underlying complex patterns not captured by simpler models.



Symmetric Teleparallel Geometries and its Applications to Cosmology

Kok Ying Hao*

Institute of Mathematical Sciences, Universiti Malaya, Malaysia yinghao3301@gmail.com



4. Symmetric Teleparallel Geometries

Symmetric Teleparallel Geometries refer to the class of spacetimes with zero torsion and curvature, which can be found by setting $T^{\lambda}_{\mu\nu} = R^{\lambda}_{\mu\alpha\nu} = 0.$

To determine the most general metric and connection under a given set of symmetries, we set the Lie derivative of the metric and connection to be 0 for each Killing vectors X associated with the spacetime symmetries: $0 = (\mathcal{L}_X g)_{\mu\nu} = (\mathcal{L}_X \Gamma)^{\lambda}_{\mu\nu}$.

For a spacetime with spherical symmetry, the most general symmetric teleparallel metric is:

$\int g_{tt}$	g_{tr}	0	0	
		\mathbf{O}	\cap	

1. Abstract and Objectives

In an attempt to obtain exact solutions to Einstein's field equations, we derive the most general metric and affine connection that respect various spacetime symmetries. Furthermore, we consider the case of symmetric teleparallel geometries by imposing vanishing torsion and curvature. Lastly, we obtained the general modified Friedmann equations for various $f(\mathbb{Q})$ gravity theories with the FLRW metric and a perfect fluid universe.

Objectives

- To determine the most general metric and connections of spacetimes with certain symmetries.
- To determine a special case where the torsion and non-metricity tensor vanishes (known as symmetric teleparallel geometries)
- To obtain the modified Friedmann equations in symmetric teleparallel FLRW universes using the $f(\mathbb{Q})$ field equations.

2. Background

Einstein's General Relativity (GR), published in 1915, is the current best description of gravity. However, GR is not the full story of gravity.

Notable issues include:

- What causes the expansion of the universe?
- Does dark matter and dark energy exist?

Potential modifications of GR include:

- $f(\mathbb{Q})$ gravity
- Loop quantum gravity and string theory
- Conformal field theories

$$g_{\mu\nu} = \begin{bmatrix} g_{rt} & g_{rr} & 0 & 0\\ 0 & 0 & g_{\theta\theta} & 0\\ 0 & 0 & 0 & g_{\theta\theta} \sin^2\theta \end{bmatrix}$$

while the most general symmetric teleparallel connection is in terms of 12 functions of t and r:

$$\Gamma^{t}_{\mu\nu} = \begin{bmatrix} C_{1} & C_{2} & 0 & 0 \\ C_{2} & C_{3} & 0 & 0 \\ 0 & 0 & C_{7} & 0 \\ 0 & 0 & 0 & C_{7} \sin^{2}\theta \end{bmatrix} \quad \Gamma^{\theta}_{\mu\nu} = \begin{bmatrix} 0 & 0 & C_{9} & -C_{11}\sin\theta \\ 0 & 0 & C_{10} & -C_{12}\sin\theta \\ C_{9} & C_{10} & 0 & 0 \\ -C_{11}\sin\theta & -C_{12}\sin\theta & 0 & -\sin\theta\cos\theta \end{bmatrix}$$

$$\Gamma^{r}_{\mu\nu} = \begin{bmatrix} C_{4} & C_{5} & 0 & 0 \\ C_{5} & C_{6} & 0 & 0 \\ 0 & 0 & C_{8} & 0 \\ 0 & 0 & 0 & C_{8} \sin^{2}\theta \end{bmatrix} \quad \Gamma^{\phi}_{\mu\nu} = \begin{bmatrix} 0 & 0 & C_{11} \csc \theta & C_{9} \\ 0 & 0 & C_{12} \csc \theta & C_{10} \\ C_{11} \csc \theta & C_{12} \csc \theta & 0 & \cot \theta \\ C_{9} & C_{10} & \cot \theta & 0 \end{bmatrix}$$

For spacetimes with spherical and translational symmetry, we obtained the FLRW metric for scale factor A(t) and lapse function N(t):

$$ds^{2} = -N^{2}(t) dt^{2} + \frac{A^{2}(t)}{1 - kr^{2}} dr^{2} + A^{2}(t) r^{2} d\theta^{2} + A^{2}(t) r^{2} \sin^{2} \theta d\phi^{2}.$$

For spacetimes with spherical, translational and time symmetry, we obtained the since disproven Einstein's static universe metric.

3. Tensor Calculus

The metric tensor $g_{\mu\nu}$ are the components of the distance formula:

 $ds^2 = g_{\mu\nu} dx^{\mu} dx^{\nu}.$

An affine connection $\Gamma^{\lambda}_{\ \mu\nu}$ satisfies the transformation rule:

$$\Gamma^{\lambda}_{\ \mu\nu} = \frac{\partial x^{\lambda}}{\partial \bar{x}^{k}} \left(\frac{\partial^{2} \bar{x}^{k}}{\partial x^{\nu} \partial x^{\mu}} + \bar{\Gamma}^{k}_{\ ij} \frac{\partial \bar{x}^{i}}{\partial x^{\mu}} \frac{\partial \bar{x}^{j}}{\partial x^{\nu}} \right).$$

The torsion, non-metricity and Riemann curvature tensor are defined respectively as follows:

$$T^{\lambda}_{\ \mu\nu} = \Gamma^{\lambda}_{\ \mu\nu} - \Gamma^{\lambda}_{\ \nu\mu}$$
$$Q_{\mu\nu\rho} = \partial_{\mu}g_{\nu\rho} - \Gamma^{\lambda}_{\ \mu\nu}g_{\lambda\rho} - \Gamma^{\lambda}_{\ \mu\rho}g_{\nu\lambda}$$
$$R^{\lambda}_{\ \mu\alpha\nu} = \partial_{\alpha}\Gamma^{\lambda}_{\ \mu\nu} - \partial_{\nu}\Gamma^{\lambda}_{\ \mu\alpha} + \Gamma^{\lambda}_{\ \sigma\alpha}\Gamma^{\sigma}_{\ \mu\nu} - \Gamma^{\lambda}_{\ \sigma\nu}\Gamma^{\sigma}_{\ \mu\alpha}$$

6. Conclusions

$$ds^{2} = g_{1}dt^{2} + g_{2}\left(\frac{dr}{1 - kr^{2}} + r^{2}d\theta^{2} + r^{2}\sin^{2}\theta d\phi^{2}\right).$$

5. Applications to Cosmology

 $f(\mathbb{Q})$ gravity is a modified theory of gravity using the non-metricity scalar \mathbb{Q} and an aribtrary function $f(\mathbb{Q})$, while requiring vanishing torsion and curvature.

Assume a perfect fluid universe with mass density ρ and pressure p with the FLRW metric,

• $f(\mathbb{Q})$ field equations (with $8\pi G = 1$ and c = 1):

$$\frac{df}{d\mathbb{Q}}G_{\mu\nu} + \frac{1}{2}g_{\mu\nu}\left(\mathbb{Q}\frac{df}{d\mathbb{Q}} - f\right) + 2\frac{d^2f}{d\mathbb{Q}^2}(\nabla_{\alpha}\mathbb{Q})P^{\alpha}_{\ \mu\nu} = T_{\mu\nu}$$

• Non-metricity scalar \mathbb{Q}

$$\mathbb{Q} = Q_{\alpha\mu\nu}P^{\alpha\mu\nu} = -6H^2 + 9HK_3 + 3K_1K_3 - 3K_3^2 + \frac{3HK_2}{A^2} - \frac{3K_1K_2}{A^2} - \frac{3K_2K_3}{A^2} - \frac{3K_2K$$

where K_1, K_2, K_3 are functions of time, $H = \frac{\dot{A}(t)}{A(t)}$ is the Hubble parameter and $P^{\alpha}_{\mu\nu}$ is the non-metricity conjugate tensor.

As a conclusion, we explored the history of Einstein's theory of General Relativity and its modifications.

To find solutions to Einstein's field equations, we determined the most general symmetric teleparallel metric and connection satisfying some desired symmetries, deriving the FLRW metric as a consequence.

Lastly, using the $f(\mathbb{Q})$ field equations, we obtained some modified Friedmann equations in the case of symmetric teleparallel FLRW universes.

• There are 4 possible cases, depending on the value of k. One such case is k = 0:



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SPATIAL ANALYSIS OF HOUSING PRICES USING R: **INSIGHTS FROM CASE** STUDIES

KISHAANNTH BALA CHANDRAN U2103226/1 SUPERVISED BY DR. DHARINI A/P PATHMANATHAN



ABSTRACT

The spatial variability of housing market determinants leads to differences in market activity and in real estate prices and values across different areas. This study analyses housing prices through spatial analysis in two different case studies: the suburbs of Perth and the towns of Boston, aiming to determine the existence of spatial autocorrelation. The Perth dataset is analysed using both non-spatial and spatial models, specifically the ordinary least squares (OLS) regression and the Manski model, respectively. The low rank spatial lag model (LSLM) with the selected spatial variables from the dataset was fitted to the Boston dataset.. The results from the first case study, Perth, showed a weak spatial autocorrelation and the data was proven to be a poor fit for the Manski model. However, the second case study, Boston house price data, showed a moderately strong spatial autocorrelation, and the low rank spatial lag model proved to be a good fit for the data. Several spatial variables were identified to have a direct impact on housing prices in the region.

INTRODUCTION

DATA DESCRIPTION

Spatial analysis has been used to study, understand and derive the insights from the spatial data either Perth geometrically or geographically. One of the crucial uses of spatial analysis is to predict the housing prices in a • Housing price dataset obtained from 2021 region. In earlier times, the method for analysing the data through a perceived spatial lens is by modelling • Consists of 19 columns the data onto the ordinary least squares (OLS) regression. As years went by, spatial models such as the Manski model and the low rank spatial lag model were developed. This study aims to conduct a spatial Boston analysis on housing prices with two different case studies, which are the suburbs of Perth and the towns of • Housing price dataset obtained from the 1978 census Boston. This study also aims to compare the two case studies to determine the existence of spatial • Well known dataset for conducting spatial analysis autocorrelation in the housing prices and spatial features which could be the factors that affect the housing • Consists of 14 columns prices in Perth and Boston. • The Manski Model **METHODOLOGY** DATA COLLECTION $Y = \rho WY + a_{l_N} + X\beta + WX\theta + u,$ Spatial Models • Moran's I Statistic • To calculate the impact of DATA PREPROCESSING the spatial variables onto $I = \frac{N}{\sum_{i=1}^{n} \sum_{i=1}^{n} w_{ii}} \times \frac{\sum_{i=1}^{n} \sum_{j=1}^{n} w_{ij} (x_i - \mu) (x_j - \mu)}{\sum_{i=1}^{n} (x_i - \mu)^2} , \text{for } i \neq j,$ $u = \lambda W u + \varepsilon$, the dependent variable • The Low Rank Spatial Lag Model • Provides the direct, EXPLORATORY DATA ANALYSIS $y_i = \beta_0 + \epsilon_i + z_i,$ indirect and total impact • Estimated based on covariance onto the dependent • Index values ranges from -1 to 1 variable $\epsilon_i \sim N(0,\sigma^2) z_i = \rho \sum_{i\neq i}^N w_{i,j} z_j + \sum_{k\neq 1}^K x_{i,k} \beta_k + u_i,$ NO • Values above O give a positive spatial autocorrelation SPATIAL ANALYSIS STOP • Values below 0 give a negative spatial autocorrelation YES The spatial weights matrices generated for these models are the Queen • Ordinary Least Squares (OLS) Regression MODEL SELECTION contiguity weights which are a combination of Rook contiguity (only common • Provides the impact of the spatial variables to the

vertices) and Bishop contiguity (only comon vertices).



Workflow of the study

- dependent variable
- Calculates adjusted R-squared value which gives the percentage of variability in the data due to the selected spatial variables

 $w_{ij} = \begin{cases} 1 & \ell_{ij} > 0 \\ 0 & \ell_{ii} = 0 \end{cases},$

APPLICATION OF DATA TO CASE STUDIES 1) PERTH **OLS REGRESSION**

MAP OF PERTH



Predictors	Estimates	Confidence Interval	p-value
(Intercept)	1833.08	1812.86 - 1853.30	<0.001
LAND AREA	-0.00	0.00 - 0.00	<0.001
NEAREST STN DIST	-0.04	-0.040.04	<0.001
CBD DIST	-0.03	-0.030.03	<0.001
NEAREST SCH DIST	8.01	0.31 - 15.70	0.041

- LAND AREA shows a negligible negative impact
- CBD DIST & NEAREST STN DIST shows a small negative impact
- NEAREST SCH DIST shows a significant positive impact
- Adjusted R-squared value is 0.241

THE MANSKI MODEL

TOTAL DIRECT INDIRECT

2) BOSTON

Variable

LOW RANK SPATIAL LAG MODEL

- The crime rate (CRIM) and proportion of owner-occupied buildings prior to 1940 (AGE) significant negative effect towards the median value prices
- The Charles River dummy variable (CHAS) and the rooms per property (RM) show a significant positive effect towards the median value prices
- The nitrous oxide concentrations (NOX) and the proportion of non-retail business acres (INDUS) show a lack of effect on the median value prices

Estimate

MAP OF BOSTON

t-value



(CE)

Standard Error

		(SE)		
(Intercept)	-14.6152	2.7742	-5.2682	$2.0972e^{-7}$
CRIM	-0.1073	0.0284	-3.7711	1.8315 <i>e</i> ⁻⁴
CHAS	1.8942	0.8897	2.1290	0.0338
RM	6.9288	0.3320	20.8720	0.0000 <i>e</i> ⁰
AGE	-0.0414	0.0121	-3.4088	$7.0827e^{-4}$
INDUS	-0.1012	0.0604	-1.6749	0.0946
NOX	-5.1642	3.8309	-1.3481	0.1783

Mean Housing Price 0 to 1,000 1,000 to 2,000 2,000 to 3,000 3,000 to 4,000

	CBD_DIST	0.018167695	-0.05751353	-0.03934583		
	NEAREST_STN_DIST	-0.016319871	-0.07852185	-0.09484173		
	LAND_AREA	-0.007572612	-0.02167637	-0.02924898		
	NEAREST_SCH_DIST	-6.113631294	358.42221512	352.30858383		
•	 CBD_DIST, NEAREST_STN_DIST & LAND_AREA show a negative and low effect on the price per square meter NEAREST_SCH_DIST shows a significant positive effect on the price per square meter 					
	• The Manski model proves to be a poor fit for the data					

CONCLUSION

The Boston house price data, which is used as benchmark data in this work, is a classic spatial data set and is comprehensive. Hence, suitable spatial models can be employed to proceed with spatial predictive modelling. However, the Perth house price data needs to be addressed further by implementing more spatial variables to proceed with spatial modelling. Future studies involving more variables that project strong spatial autocorrelation can be investigated to improve the existing spatial predictive model accuracy. It is also of interest to study the effects of various spatial weight matrices to quantify the spatial relationships that exist among the features in the dataset of interest.

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UNIVERSITI MALAYA



Name: Ikhwan bin Mohd Nizam Matric No.: U2001245 Supervisor: Dr. Shahizat bin Amir & Dr. Mohd. Zahurin bin Mohamed Kamali

for SIM3021: Mathematical Science Project

Fractal Dimension of Klang River Network: A Box–counting Method

Study Area

Klang River and its network system in Klang Valley region consist of Klang, Petaling, Ulu Langat, Gombak District and Wilayah Persekutuan Kuala Lumpur

Introduction

- Fractal dimension was coined by Benoit B. Mandelbrot in 1975.
- The dimension is not integer, but rather a fraction unlike Euclidean dimensions.
- Fractals are geometry figures & their characteristics:
 - Self similar
 - Recursive
- A fractal dimension is a ratio providing a statistical index of complexity.

Problem Statement:

(2)

The Klang River network in the Klang Valley region lacks a comprehensive assessment of its fractal properties. Understanding its fractal dimension is crucial for insights into its structure, evolution, and behaviour. This study aims to evaluate the strengths and limitations of using the box-counting method integrated with GIS to estimate the fractal dimension, providing valuable insights for environmental management, flood control, and urban planning.

Objectives:

- Understanding the fractal complexity in Klang River network in Klang Valley area using spatial analysis and mathematical computation.
- Investigate the applicability of Geographic Information System (GIS) with the box counting method for estimating the fractal dimension of Klang River network.

Methodology

$$N(L) \propto L^{-D}$$
 (1) $D = \frac{\log N(L)}{\log 1/L}$

where:

- D represents the fractal dimension.
- N(L) is the number of grid boxes of side length L needed to cover the network.
- L is the grid boxes with side length L.



This power law corresponds to a linear relationship of **Equation 1** on a logarithmic scale. A log-log plot of N(L) against 1/L is constructed with the plotted points, and the slope of the resulting line gives an estimate of the fractal dimension, D by fitting these points with linear regression using **Equation 2**.



GIS & Spatial Analysis Mathematical Computation

Grid size, L (km) N(L) Log (1/L) Log (N(L)) R-squared: 0.9934 Estimated fractal dimension (D): 1.6114 9 23580 4.3725 0.25 0.6021 0.50 0.3010 3.8713 7436 1.00 2148 3.3320 2.7938 2.00 622 -0.3010 4.00 181 -0.6021 2.2577 1 9685 6 00 93 -0 7782 8 00 56 -0.9031 1 7482 12.00 34 -1.0792 1.5315 16.00 19 -1.2041 1.2788 32.00 -1.5051 0.9542 -2 64.00 -1.8062 0.6021 Loa(Box Size

Results and Discussion

Interpretation of Results:

- The table lists grid sizes and corresponding box counts, showing an inverse relationship.
- The log-log plot of box size and box count indicates a power-law distribution, characteristic of fractals.
- The fractal dimension of 1.6114 suggests a complex river network, impacting hydrological processes like water flow and flood dynamics.

Box Counting Method Using GIS:

- QGIS automates the box-counting process, ensuring accuracy and efficiency.
- This method helps manage large datasets and provides insights into river network complexity.

Urban River Network and Impact of Urbanization:



- Urbanization can simplify river networks, reducing their capacity to manage heavy rainfall, as observed in the Taihu Region, China.
 - Understanding the Klang Valley river network's fractal properties is vital for effective water management and flood mitigation, particularly after the severe December 2021 floods.
- Sustainable urban planning should balance development with ecological preservation, using advanced technologies and community engagement.

Conclusion

Acknowledgement and References

The box-counting method effectively measures the complexity of the Klang River network, revealing a fractal dimension of 1.6114. This high degree of branching and connectivity is crucial for hydrological and ecological processes. Limited data accessibility highlights the need for more comprehensive studies and institutional support. The project also demonstrates the utility of GIS in enhancing the box-counting method, facilitating efficient and accurate spatial analysis. Further studies on Malaysian river systems are recommended to understand the impacts of urbanization. Policymakers should use GIS technologies for better river conservation, flood management, and urban planning. Data:

Scholar:



KEMENTERIAN SUMBER ASLI DAN KELESTARIAN ALAM PUSAT GEOSPATIAL NEGARA

