

Mathematical Modelling for the Effect of Malaria on the Heterozygous and Homozygous Genes

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ABSTRACT

This paper models the effect of malaria on the homozygous for the normal gene (AA), heterozygous for sickle cell gene (AS) and homozygous for sickle cell gene (SS) using the first order ordinary differential equation. The Diseases Free Equilibrium (DFE) was obtained and used to compute the basic reproduction Number R_0 . The local stability of the (DFE) was analyzed.

KEYWORDS: - *heterozygous, homozygous, sickle cell diseases, plasmodium, genotype.*

Introduction.

The sickle cell gene is caused by a single amino acid mutation (valine instead of glutamate at the 6th position) in the beta chain of the hemoglobin gene. Inheritance of this mutated gene from both parents leads to sickle cell disease and people with this disease have shorter life expectancy. On the contrary, individuals who are carriers for the sickle cell disease (with one sickle gene and one normal hemoglobin gene, also known as sickle cell trait) have some protective advantage against malaria. As a result, the frequencies of sickle cell carriers are high in malaria-endemic areas. It was found that that the sickle cell trait provides 60% protection against overall mortality. Most of this protection occurs between 2-16 months of life, before the onset of clinical immunity in areas with intense transmission of malaria.

The natural history of malaria involves cyclical infection of humans and female *Anopheles* mosquitoes. In humans, the parasites grow and multiply first in the liver cells and then in the red cells of the blood. The infected mosquito carries the disease from one human to another (acting as a “vector”), while infected humans transmit the parasite to the mosquito, in contrast

to the human host, the mosquito vector does not suffer from the presence of the parasites (CDC, 2018).

Methodology

Model Formulation.

In modelling diseases using Ordinary Differential Equation, Akinwande (2018) precludes an assumption of vertical transmission, in which those who recover from infection are returned into the Susceptible class making room for possible re-infection as in the case of malaria fever and some other diseases.

In modelling the effect of malaria on the heterozygous and homozygous genes, the total population is subdivided into four main subgroups namely the susceptible homozygous (AA) $S_1(t)$, the infected homozygous (AA) $I_1(t)$, the Susceptible heterozygous (AS) $S_2(t)$, infected heterozygous (AS) $I_2(t)$, the Susceptible homozygous (SS) $S_3(t)$, the infected homozygous (SS) $I_3(t)$, $M_1(t)$ the non-plasmodium carrier mosquito and $M_2(t)$ the plasmodium carrier mosquito. People enter the susceptible class through birth and recovery from the infected class and leave the susceptible class through infection, natural death and death due to infection. It was assumed in this work that infants are not infected with malaria from birth and that the mosquito is non-plasmodium carrier from birth; they become infected through contact with infected humans. The transmission dynamics of malaria as it affects the heterozygous and homozygous sickle cell genes is given by the equations 1-8 below

$$\frac{dS_1}{dt} = \omega_1\beta_1 + \theta_1\beta_2 + \gamma_1I_1 - \alpha_1S_1M_2 - (\mu_1 + \delta_1)S_1 \quad 1$$

$$\frac{dI_1}{dt} = \alpha_1S_1M_2 - \gamma_1I_1 - (\mu_1 + \delta_1)I_1 \quad 2$$

$$\frac{dS_2}{dt} = \omega_2\beta_1 + \theta_2\beta_2 + \rho_1\beta_3 + \gamma_2I_2 - \alpha_1S_2M_2 - (\mu_1 + \delta_2)S_2 \quad 3$$

$$\frac{dI_2}{dt} = \alpha_1S_2M_2 - \gamma_2I_2 - (\mu_1 + \delta_2)I_2 \quad 4$$

$$\frac{dS_3}{dt} = \theta_3\beta_2 + \rho_2\beta_3 + \gamma_3I_3 - \alpha_1S_3M_2 - (\mu_1 + \delta_3)S_3 \quad 5$$

$$\frac{dI_3}{dt} = \alpha_1 S_3 M_2 - \gamma_3 I_3 - (\mu_1 + \delta_3) I_3 \quad 6$$

$$\frac{dM_1}{dt} = \beta_4 - \alpha_3 M_1 (I_1 + I_2 + I_3) - \mu_2 M_1 \quad 7$$

$$\frac{dM_2}{dt} = \alpha_3 M_1 (I_1 + I_2 + I_3) - \mu_2 M_2 \quad 8$$

Notation and definition of variables and parameter

$S_1(t)$ Number of susceptible AA

$I_1(t)$ Number of Infected AA

$S_2(t)$ Number of susceptible AS

$I_2(t)$ Number of Infected AS

$S_3(t)$ Number of susceptible SS

$I_3(t)$ Number of Infected SS

$M_1(t)$ Number of non-plasmodium carrier mosquitoes

$M_2(t)$ Number of plasmodium carrier mosquitoes

β_1 normal birth rate in human (AA)

β_2 normal birth rate in human (AS)

β_3 normal birth rate in human (SS)

β_4 normal birth rate in mosquitoes

μ_1 normal death rate in human

μ_2 normal and induced death rate in mosquitoes

δ_1 death due to infection in AA

δ_2	death due to infection in AS
δ_3	death due to infection in SS
α_1	infection rate from mosquitoes to human
α_2	infection rate from infected human to mosquitoes
γ_1	recovery rate in AA
γ_2	recovery rate in AS
γ_3	recovery rate in SS
θ_1	proportion of birth of AA by AS
θ_2	proportion of birth of AS by AS
θ_3	proportion of birth of SS by AS
ω_1	proportion of birth of AA by AA
ω_2	proportion of birth of AS by AA
ρ_1	proportion of birth of AS by SS
ρ_2	proportion of birth of SS by SS

Note that i) $\omega_1 + \omega_2 = 1$ 9

ii) $\theta_1 + \theta_2 + \theta_3 = 1$ 10

iii) $\rho_1 + \rho_2 = 1$ 11

Existence of Equilibrium State of the Model

At equilibrium,

$$\frac{dS_1}{dt} = \frac{dI_1}{dt} = \frac{dS_2}{dt} = \frac{dI_2}{dt} = \frac{dS_3}{dt} = \frac{dI_3}{dt} = \frac{dM_1}{dt} = \frac{dM_2}{dt} = 0 \quad 12$$

Let

$$(S_1, S_2, S_3, I_1, I_2, I_3, M_1, M_2) = (S_1^0, S_2^0, S_3^0, I_1^0, I_2^0, I_3^0, M_1^0, M_2^0) \quad 13$$

Be arbitrary equilibrium points, therefore the system 1 – 8 becomes

$$\frac{dS_1^0}{dt} = \omega_1\beta_1 + \theta_1\beta_2 + \gamma_1I_1^0 - \alpha_1S_1^0M_2^0 - (\mu_1 + \delta_1)S_1^0 \quad 14$$

$$\frac{dI_1^0}{dt} = \alpha_1S_1^0M_2^0 - \gamma_1I_1^0 - (\mu_1 + \delta_1)I_1^0 \quad 15$$

$$\frac{dS_2^0}{dt} = \omega_2\beta_1 + \theta_2\beta_2 + \rho_1\beta_3 + \gamma_2I_2^0 - \alpha_1S_2^0M_2^0 - (\mu_1 + \delta_2)S_2^0 \quad 16$$

$$\frac{dI_2^0}{dt} = \alpha_1S_2^0M_2^0 - \gamma_2I_2^0 - (\mu_1 + \delta_2)I_2^0 \quad 17$$

$$\frac{dS_3^0}{dt} = \theta_3\beta_2 + \rho_2\beta_3 + \gamma_3I_3^0 - \alpha_1S_3^0M_2^0 - (\mu_1 + \delta_3)S_3^0 \quad 18$$

$$\frac{dI_3^0}{dt} = \alpha_1S_3^0M_2^0 - \gamma_3I_3^0 - (\mu_1 + \delta_3)I_3^0 \quad 19$$

$$\frac{dM_1^0}{dt} = \beta_4 - \alpha_3M_1^0(I_1^0 + I_2^0 + I_3^0) - \mu_2M_1^0 \quad 20$$

$$\frac{dM_2^0}{dt} = \alpha_3M_1^0(I_1^0 + I_2^0 + I_3^0) - \mu_2M_2^0 \quad 21$$

Let

$$\mu_1 + \partial_1 = A_1, \mu_1 + \partial_2 = A_2, \mu_1 + \partial_3 = A_3 \quad 22$$

$$\omega_1\beta_1 + \theta_1\beta_2 + \gamma_1I_1^0 - \alpha_1S_1^0M_2^0 - A_1S_1^0 = 0 \quad 23$$

$$\alpha_1S_1^0M_2^0 - \gamma_1I_1^0 - A_1I_1^0 = 0 \quad 24$$

$$\omega_2\beta_1 + \theta_2\beta_2 + \rho_1\beta_3 + \gamma_2I_2^0 - \alpha_1S_2^0M_2^0 - A_2S_2^0 = 0 \quad 25$$

$$\alpha_1S_2^0M_2^0 - \gamma_2I_2^0 - A_2I_2^0 = 0 \quad 26$$

$$\theta_3\beta_2 + \rho_2\beta_3 + \gamma_3I_3^0 - \alpha_1S_3^0M_2^0 - A_3S_3^0 = 0 \quad 27$$

$$\alpha_1 S_3^0 M_2^0 - \gamma_3 I_3^0 - A_3 I_3^0 = 0 \quad 28$$

$$\beta_4 - \alpha_3 M_1^0 (I_1^0 + I_2^0 + I_3^0) - \mu_2 M_1^0 = 0 \quad 29$$

$$\alpha_3 M_1^0 (I_1^0 + I_2^0 + I_3^0) - \mu_2 M_2^0 = 0 \quad 30$$

Let

$$I_1^0 = I_2^0 = I_3^0 = 0 \quad 31$$

$$\omega_1 \beta_1 + \theta_1 \beta_2 - \alpha_1 S_1^0 M_2^0 - A_1 S_1^0 = 0 \quad 32$$

$$\alpha_1 S_1^0 M_2^0 = 0 \quad 33$$

$$\omega_2 \beta_1 + \theta_2 \beta_2 + \rho_1 \beta_3 - \alpha_1 S_2^0 M_2^0 - A_2 S_2^0 = 0 \quad 34$$

$$\alpha_1 S_2^0 M_2^0 = 0 \quad 35$$

$$\theta_3 \beta_2 + \rho_2 \beta_3 - \alpha_1 S_3^0 M_2^0 - A_3 S_3^0 = 0 \quad 36$$

$$\alpha_1 S_3^0 M_2^0 = 0 \quad 37$$

$$\beta_4 - \mu_2 M_1^0 = 0 \quad 38$$

$$-\mu_2 M_2^0 = 0$$

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$$\text{From 39 } M_2^0 = 0, \quad 40$$

$$\text{From 38 } M_1^0 = \frac{\beta_4}{\mu_2} \quad 41$$

$$\text{From 32 } S_1^0 = \frac{\omega_1 \beta_1 + \theta_1 \beta_2}{A_1} \quad 42$$

$$\text{From 34 } S_2^0 = \frac{\omega_2 \beta_1 + \theta_2 \beta_2 + \rho_1 \beta_3}{A_2} \quad 43$$

$$\text{From 36 } S_3^0 = \frac{\theta_3 \beta_2 + \rho_2 \beta_3}{A_3} \quad 44$$

Thus, the DFE (E^0) exists at the points

$$E^0 = (S_1^0, S_2^0, S_3^0, I_1^0, I_2^0, I_3^0, M_1^0, M_2^0) \\ = \left(\frac{\omega_1\beta_1 + \theta_1\beta_2}{A_1}, \frac{\omega_2\beta_1 + \theta_2\beta_2 + \rho_1\beta_3}{A_2}, \frac{\theta_3\beta_2 + \rho_2\beta_3}{A_3}, 0, 0, 0, \frac{\beta_4}{\mu_2}, 0 \right)$$

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Computation of Basic Reproduction Number (R_o)

Somma et al (2017) applied next generation matrix operation to compute the Basic reproduction Number of the model as used by Diekmann et al(1990) and improved by Driessche (2002). The effective basic reproduction number is the largest Eigenvalue or spectral radius of (FV^{-1}). The basic reproduction number of the model is given as the largest eigenvalue or spectral radius of FV^{-1} .

$$FV^{-1} = \left\{ \left[\frac{\partial F_i(E^0)}{\partial x_i} \right] \left[\frac{\partial V_i}{\partial x_i} \right]^{-1} \right\}$$

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Where F_i is the rate of appearance of new infection in compartment i , V_i is the transfer of infections from one compartment i to another and E^0 is the Disease-Free Equilibrium.

$$f_i = \begin{pmatrix} f_1 \\ f_2 \\ f_3 \\ f_4 \end{pmatrix} = \begin{pmatrix} \alpha_1 S_1^0 M_2^0 \\ \alpha_1 S_2^0 M_2^0 \\ \alpha_1 S_3^0 M_2^0 \\ \alpha_3 M_1^0 (I_1^0 + I_2^0 + I_3^0) \end{pmatrix}$$

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Differentiating (47) partially with respect to time gives

$$F = \begin{bmatrix} 0 & 0 & 0 & \alpha_1 S_1^0 \\ 0 & 0 & 0 & \alpha_1 S_2^0 \\ 0 & 0 & 0 & \alpha_1 S_3^0 \\ \alpha_3 M_1^0 & \alpha_3 M_1^0 & \alpha_3 M_1^0 & 0 \end{bmatrix}$$

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$$F = \begin{bmatrix} 0 & 0 & 0 & \frac{\alpha_1(\omega_1\beta_1 + \theta_1\beta_2)}{A_1} \\ 0 & 0 & 0 & \frac{\alpha_1(\omega_2\beta_1 + \theta_2\beta_2 + \rho_1\beta_3)}{A_2} \\ 0 & 0 & 0 & \frac{\alpha_1(\theta_3\beta_2 + \rho_2\beta_3)}{A_3} \\ \frac{\alpha_3\beta_4}{\mu_2} & \frac{\alpha_3\beta_4}{\mu_2} & \frac{\alpha_3\beta_4}{\mu_2} & 0 \end{bmatrix} \quad 49$$

$$\begin{aligned} K_1 &= \omega_1\beta_1 + \theta_1\beta_2, \\ K_2 &= \omega_2\beta_1 + \theta_2\beta_2 + \rho_1\beta_3, \\ K_3 &= \theta_3\beta_2 + \rho_2\beta_3, \\ K_4 &= \frac{\beta_4}{\mu_2} \end{aligned} \quad 50$$

$$F = \begin{bmatrix} 0 & 0 & 0 & \frac{\alpha_1 K_1}{A_1} \\ 0 & 0 & 0 & \frac{\alpha_1 K_2}{A_2} \\ 0 & 0 & 0 & \frac{\alpha_1 K_3}{A_3} \\ \alpha_3 K_4 & \alpha_3 K_4 & \alpha_3 K_4 & 0 \end{bmatrix} \quad V = \begin{bmatrix} A_1 & 0 & 0 & 0 \\ 0 & A_2 & 0 & 0 \\ 0 & 0 & A_3 & 0 \\ 0 & 0 & 0 & \mu_2 \end{bmatrix} \quad 52$$

$$\det V = A_1 A_2 A_3 \mu_2 \quad 53$$

$$AdjV = \begin{bmatrix} A_2 A_3 \mu_2 & 0 & 0 & 0 \\ 0 & A_1 A_3 \mu_2 & 0 & 0 \\ 0 & 0 & A_1 A_2 \mu_2 & 0 \\ 0 & 0 & 0 & A_1 A_2 A_3 \end{bmatrix} \quad 54$$

$$V^{-1} = \frac{AdjV}{\det V} = \begin{bmatrix} \frac{1}{A_1} & 0 & 0 & 0 \\ 0 & \frac{1}{A_2} & 0 & 0 \\ 0 & 0 & \frac{1}{A_3} & 0 \\ 0 & 0 & 0 & \frac{1}{\mu_2} \end{bmatrix} \quad 55$$

$$FV^{-1} = \begin{bmatrix} 0 & 0 & 0 & \frac{K_1\alpha_1}{A_1\mu_2} \\ 0 & 0 & 0 & \frac{K_2\alpha_1}{A_2\mu_2} \\ 0 & 0 & 0 & \frac{K_3\alpha_1}{A_3\mu_2} \\ \frac{K_4\alpha_3}{A_1} & \frac{K_4\alpha_3}{A_2} & \frac{K_4\alpha_3}{A_3} & 0 \end{bmatrix} \quad 56$$

The characteristics equation of (56), gives

$$|FV^{-1} - \lambda I| = 0 \quad 57$$

$$\begin{vmatrix} -\lambda & 0 & 0 & \frac{K_1\alpha_1}{A_1\mu_2} \\ 0 & -\lambda & 0 & \frac{K_2\alpha_1}{A_2\mu_2} \\ 0 & 0 & -\lambda & \frac{K_3\alpha_1}{A_3\mu_2} \\ \frac{K_4\alpha_3}{A_1} & \frac{K_4\alpha_3}{A_2} & \frac{K_4\alpha_3}{A_3} & -\lambda \end{vmatrix} = 0 \quad 58$$

$$-\lambda \left[-\lambda(\lambda^2 - \frac{\alpha_1\alpha_3 K_3 K_4}{A_3^2 \mu_2}) \right] - 0 + \frac{\alpha_1 K_1}{A_1 \mu_2} (0) = 0 \quad 59$$

$$\lambda^2 - \frac{\alpha_1\alpha_3 K_3 K_4}{A_3^2 \mu_2} = 0 \quad 60$$

$$\lambda = \pm \sqrt{\frac{\alpha_1\alpha_3 K_3 K_4}{A_3^2 \mu_2}} \quad 61$$

$$\lambda_1 = 0, \lambda_2 = -\sqrt{\frac{\alpha_1\alpha_3 K_3 K_4}{A_3^2 \mu_2}}, \lambda_3 = +\sqrt{\frac{\alpha_1\alpha_3 K_3 K_4}{A_3^2 \mu_2}} \quad 62$$

λ_3 is the spectral radius of $\rho(FV^{-1})$

$$R_0 = \sqrt{\frac{\alpha_1\alpha_3 K_3 K_4}{A_3^2 \mu_2}} \quad 63$$

Concluding Remarks

The Disease-Free Equilibrium exists and it is equal to

$$E^0 = (S_1^0, S_2^0, S_3^0, I_1^0, I_2^0, I_3^0, M_1^0, M_2^0) \\ = \left(\frac{\omega_1 \beta_1 + \theta_1 \beta_2}{A_1}, \frac{\omega_2 \beta_1 + \theta_2 \beta_2 + \rho_1 \beta_3}{A_2}, \frac{\theta_3 \beta_2 + \rho_2 \beta_3}{A_3}, 0, 0, 0, \frac{\beta_4}{\mu_2}, 0 \right) \quad 64$$

The Disease-Free Equilibrium is locally asymptotically stable if Basic reproduction Number $R_0 \leq 1$ and unstable otherwise.

If $R_0 < 1$ then

$$\frac{\alpha_1 \alpha_3 K_3 K_4}{A_3^2 \mu_2} < 1 \quad 65$$

References

- Akinwande, N.I (1996), A mathematical model of Yellow Fever endemics. Afrika Matematika 6: 50-59.
- Akinwande N.I. (2018); Introductory notes on biomathematics a paper presented at the third Workshop on mathematical modelling, Department of mathematics, University of Nigeria, Nsukka.
- Diekmann, O, JAP Heesterbeek and JAJ metz 1990, "On the definition and the computation of the Basic Reproduction Ratio (R_0) in model for infectious Diseases in heterogeneous populations" Journal of mathematical Biology. 28(4) 365 – 382.
- <https://www.cdc.gov/malaria/about/biology/index.html> retrieved on 5th march, 2019
- Somma A.S., Akinwande N. I., Jiya M., and Abdulrahman S. Stability Analysis of Diseases Free Equilibrium (DFE) State of mathematical model of Yellow Fever incorporating Secondary Host Pacific Journal of Science and Technology 18(2)110-119 (2017)
- Van den Driessche, P; Watmough, J (2002). Reproduction numbers and sub threshold endemic equilibria for the compartmental models of disease transmission. Math. Biosci. 180:29-48