Grading Rubric - Modeling and Simulation Contest

Team: FSV

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Summary Report Document

	Code not well	2.	5 Very good description	3)
Item Description	description	Brahimes as ware Gradi	ng Guidance	graphic higs rations	Points
Completeness and quality of the report	Report is not well organized and/or is poorly written Main code not well	Pretty well organized but not well written with or other errors	Pretty well organized with a few grammar errors Very good description	Report is well organized, well written, and easy to follow	Awarde 5
Description of the main code and code listing	documented and incompletely bescribed	Documentation of code is missing significant sections	and documentation of the code, missing minor points	Excellent description and documentation of the code	Q
Demonstrated understanding of the system and model assumptions	adequately describe the system and the model assumptions	Description shows a lack of understanding of the model components or assumptions	Very good understanding of the system but missing a few of the key model assumptions	Discussion indicates full understanding of the model premises and model assumptions	30 E
Descriptions of the runs and results	Summary provides little or no evidence to support description	Summary of model runs and results lacks supporting graphics or data	sufficient data and	Excellent summary of the model runs and results including sufficient data and graphic illustrations	8
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Sage Codes	2	5	8	10	PAS
Correctness of base code used to solve the problem	Code not well documented and incompletely described	Documentation of code is missing significant sections	Very good description and documentation of the code, missing minor points	Excellent description and documentation of the code	5
Correctness of sensitivity runs ancillary codes	Summary provides little or no evidence to support description	Summary of model runs and results lacks supporting graphics or data	Very good summary of the model runs and results including sufficient data and graphic illustrations	Excellent summary of the model runs and results including sufficient data and graphic illustrations	5
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Bonus Points	2 Code not well	5 terms A of model runs	8 = wood uma sug	10 com somethy or me	P
Discussion of added component 1		Documentation of code is missing significant sections	Very good description and documentation of the code, missing minor	Excellent description and documentation of the code	5
Correctness of code for added component 1	Summary provides little or no evidence to support description	Summary of model runs and results lacks supporting graphics or data	very good summary of the model runs and results including sufficient data and graphic illustrations	Excellent summary of the model runs and results including sufficient data and graphic illustrations	5
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Discussion of added component 2	documented and incompletely described	Documentation of code is missing significant sections	and documentation of the code, missing minor	Excellent description and documentation of the code	80
Correctness of code for added component 2	Summary provides little or no evidence to support description	Summary of model runs and results lacks supporting graphics or data	very good summary of the model runs and results including sufficient data and graphic illustrations	Excellent summary of the model runs and results including sufficient data and graphic illustrations	8
	2 Code not well	5	0	10	- hPL4
Discussion of added component		Documentation of code is missing significant sections	Very good description and documentation of the code, missing minor point	Excellent description and documentation of the code	2

Correctness of code for added component 3	Summary provides little or no evidence to support description	Summary of model runs and results lacks supporting graphics or data	sufficient data and	Excellent summary of the model runs and results including sufficient data and graphic illustrations	2
Discussion of added component 4		Documentation of code is missing significant sections	Very good description and documentation of the code, missing minor points very good summary of the model runs and results including sufficient data and	10 Excellent description and	n/a
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component 4	Summary provides little or no evidence to support description	Summary of model runs and results lacks supporting graphics or data		Excellent summary of the model runs and results including sufficient data and graphic illustrations	no

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J subtitle/title XSEDE 2016 Maluria Simulation Brian Bartoldson, Eitan Lees, Ian McCann, Amirhessam Tahmassebi and Alex Townsend April 3, 2016 Malaria However, IT IN THE WAR "Malaria is a serious and sometimes fatal disease caused by a parasite that commonly infects a certain type of mosquito [that] feeds on humans" (CDC). Last year, there were 214 million Malaria cases and 438,000 Malaria deaths (WHO). Malaria is both preventable and curable, however, and Malaria death rates among at-risk populations have fallen by 60% in the last 15 years (WHO). short souteres Given that Malaria is preventable, curable, and very harmful, it's important to determine the do not floor most effective measures to combat the illness. A mathematical model of Malaria infection that includes factors like mosquito population size and mosquito bite rate will enable us to understand what drives Malaria incidence. This information can be used by public health professionals to further alleviate the effect of Malaria. 2 Model Design The details of the model for Malaria incidence were provided in "Mathematical Modeling for Malaria Control.pd". Basically, the model assumes no latent period for infection, the possibility to become immune, and therefore three populations of human villagers: sick, immune, and healthy. Mosquitos are not treated as capable of becoming immune, and thus there are two populations of mosquitos: infected and healthy. The size of each of these populations is calculated at each time step in our simulation program (attached). The size of each population is a function of other populations' sizes, and several parameters that define quantities such as the mosquito bite rate and the human birth rate. See section "4.1" Sensitivity Analysis" below for a discussion and graphical depictions of the impact of varying some use of these parameters.

The model was converted into a Dython program that was provided to us in an almost-complete state. Our role was developing functions that would define the sizes of several populations at each time step. Importantly, the definition for the size of the population of healthy villagers was provided Section left to use LaTex internal references to us, and we made changes to it for a couple of reasons: 1) healthy villager population size was not use hyperre a function of the sizes of the immune and sick villager populations, even though these two groups pachage could be giving birth to healthy villagers; 2) healthy villager population size should never be lower (Opare, 2010) W. n.+. to make These two reasons for changing the provided code for healthy villagers also influenced our definitheseactive tions of the other populations. Basically, we disallowed negative population values, and we assumed that infected organisms (mosquitos and humans) would give birth to healthy organisms in line with the literature on the topic (Opare, 2010). if not allenth populationsher to match new stort state which want to emphasize populations he added value of Thorn smoken 1 your contributions not the deficiencies c the shellton

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The system of time dependent complete equations for this model was solved with using an expirct the time stepping method. Similar to the Forward Ever method each the current timestep caps be solved for only using-

Model Solution

This model was solved using a time-stepping method to solve the system of coupled equations that made up the model, very similar to the Forward Euler method. This method is an explicit method for solving models such as this in the sense that each can be directly solved using information from the previous timesteps and does not require solving a nonlinear system with dependence on the current time at each timestep. For implementation, we used Python with packages MatPlotLib and Numpy for plotting and arrays. Numpy proved most useful in the sensitivity analysis as it was required to compute the grid for the 3D plots shown in the next section.

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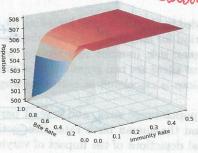
4 Results and Conclusions

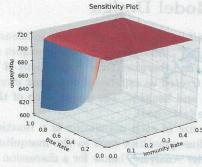
4.1 Sensitivity Analysis
We investigated the sensitivity of total population was the parameters immunity and
The immunity and the bite rate are the two parameters which were analyzed for sensitivity using bite rate. total population as a sensitivity metric. As the plot below shows, the immunity rate plays a major role in final population. Bite rate is only able to cause a population crash in the event that the immunity rate is sufficiently small, usually less than 10%. The simulation was also run out to 5000 days (nearly 14 years) in order to see how the populations would respond to varying parameters in the long term. This is the subject of the next figure. As you can see, the population will still start to erash, as expected. Though oddly the effect of bite rate, on the long-term, is not felt until the immunity rate gets very low. This may be an effect of the underlying model or the assumptions made in the model.

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shown in Fig 16. Why is it expected Sensitivity Plot elaborate 507





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(a) 200 Day Short-Term Sensitivity

(b) 5000 Day Long-Term Sensitivity

Figuret 1: Caption = add this

Model Additions

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5.1 Relapse

A major assumption of our model is that all healthy people in our population have the same probability to become sick for all time. With many diseases, once an individual has been infected and has recovered they are more likely to relapse in to a sick state. In order to incorporate the effects of relapse into our model, we had to define a new population of villagers who previously had the disease and recovered. The recovered villagers would be a subset of the healthy villagers and would lead to an increase in the growth of the sick villager population.

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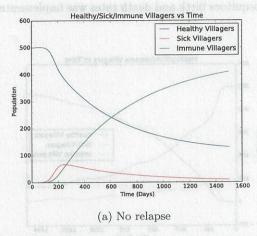
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ship is not referred to in the text stris equation needs to be specifically referred to inshe Relapse = Recovered Villagers * Relapse Rate

difference

in magnitude

Recovered Villagers = Recovered - Death - Relapse - InfectedVillagers



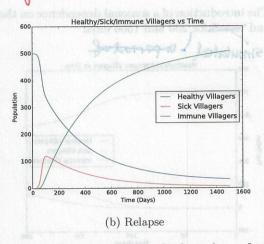


Figure 2: The introduction of relapse results in a larger initial sickness spike but also a faster immunity saturation.

We used the provided relapse roots of X assuming

A relapse rate of 17% was provided and was taken to mean that after each day there was a 17% chance for previously sick villagers to become sick again. Modification to the healthy villagers and sick villagers were made such that

SickVillagers = Infected - Recover - Immune - SickDies + Relapse

HealthyVillagers = Births + Recover - HealthyDies - Infected - Relapse.

allowing for the relapsed villagers to be removed from the healthy group and added to the sick population. Immune villager population was not altered in this modification. Left unaltered.

We see that for the case of no relapse around the 200 day mark there there is a rapid increase in the sick population and leading to a decrease in the healthy population as well as an increase in the immune population. For the introduction of a relapse model we see that this spike in sickness_sick occurs sooner around the 100 day marks and is almost twice the size in magnitude. What is the

An interesting effect was also observed with respect to the immune villager population. The only way for a villager to become immune in our current model is to become sick in a previous time step. We observed that the introduction of relapse into our model increases the number of sick villagers which in turn results in an increase in the number of immune villagers. As long as the death rate for a sick villager is not to high, the introduction of relapse seems increase the rate at which the village is saturated with an immune population. How which is too with?

Seasonal Mosquito Population

We madeled the According to the seasonal increase in the number of mosquitoes, such as a rainy season in tropical areas) we have to figure out the new rates for mosquitoes birth and death. The mosquito birth rate "mbr" and death rate "mdr" are then defined by 29 (2) as

 $mbr = \frac{1}{2}[cos(2\pi t/365 + 1) * MaxMosquitosBirthRate]$

where $mdr = \frac{1}{2}[cos(2\pi t/365 + 1)*MaxMosquitosDeathRate]$

(3)

MaxMosquitosBirthRate = 0.02

MaxMosquitosDeathRate = 0.022

The introduction of a seasonal dependence on the mosquitoes birth and death rates was implemented and tested for 200 and 1500 days.

Simulat Healthy/Sick/Immune Villagers vs Time 500 500 400 400 Healthy Villagers 300 Healthy Villagers Sick Villagers 200 Immune Villagers 200 Immune Villagers 100 100 -100 Time (Days) (a) 200 Days (b) 1500 Days

Figure 3: The introduction of seasonal change increases the rate at which the healthy population decreases.

For the 200 day seenario only the beginning of the seasonal effects can be seen. For longer run times the oscillator nature of the mosquito population is much more dramatic. As you can see, the number of healthy villagers decreases over time like the original model but variation in mosquito population lead to a less smooth transition.

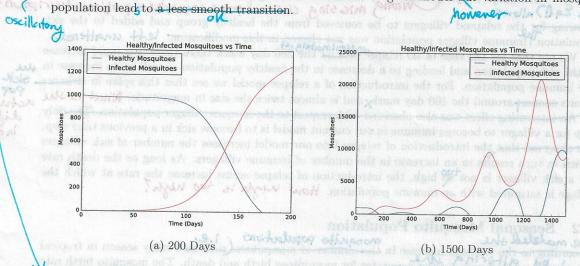
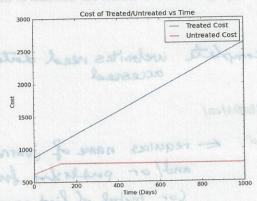


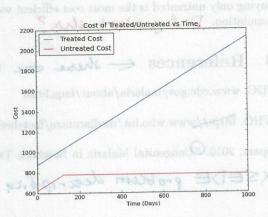
Figure 4: The introduction of seasonal change increases the rate at which the healthy mosquitoes decreases.

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5.3 Preventative Net Cost Analysis





(a) For the treated graph, treated nets were bought (b) For the treated graph, untreated nets were bought each time there were new villagers and untreated nets were bought for the untreated graph were bought for the untreated graph.

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Introduction of nets into a village is a common tool to fight against the spread of malaria. A cost analysis of using such nets was performed taking into account two possible net types. The first was a simple net with a cost of \$2.50 and the second was a treated net with a cost of \$3.50. Insecticide treated nets (ITN) reduced the incidence of uncomplicated malarial episodes by 50% compared to no nets, and 39% compared to untreated nets. For this analysis it was assumed that one net could cover two peoples.

The difference between the cost where all treated nets and all untreated nets were bought was 1845.0. The difference in the total healthy villagers was 296 72 Keeping everything the same, but making it so each new net bought is an untreated one we get a cost difference of 345.0 and healthy villager difference of 296.64 The cost difference between of buying all treated nets and half treated, half untreated is 1005.0 and the healthy villager difference is 178.25 It is clear that buying the treated net for each new villager is not worth the much higher cost in the long term.

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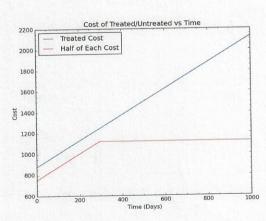


Figure 6: For the treated graph, treated nets were bought each time there were new villagers and for the half graph, originally half the nets were treated and half untreated and then untreated nets

On a short time scale, buying the villagers the more expensive net will stop the population from

explain fractional villagers

5

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decreasing immediately, but even with all untreated nets, the population will come back and increase later. I would say that originally buying a little more (maybe 60-70 percent) treated nets and then buying only untreated is the most cost efficient way of buying the nets to control the decrease of the population. Did you tost this?

References - these ove incomplete websites need date 6

CDC. www.cdc.gov/malaria/about/faqs.html

WHO. http://www.who.int/mediacentre/factsheets/fs094/en/

Opare, 2010. Congenital Malaria in Newborn Twins".

XSEDE problem description

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no note, and 39% compared to uniformed nets (100 this analysis it was assumed that one not could Cortaining where interms of infection rates or other parameters of lehorn silk the cost where all treated neigh and all unitrasted nells were bought was 1845.0. The difference in the total healthy villagers was 1967. Meroing overything the same, but

half unirested and 005.0 and the healthy villager difference is 11.2. It is clear that having the explain fractional