

Swiss TPH



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# Spatial-temporal distribution of entomological inoculation rates in Manhica DSS, Mozambique

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## Outline

- Introduction
- Objectives
- Methodology
- Modelling
- Results
- Discussion
- Acknowledgement



## Introduction

- Malaria is still a primary cause of both morbidity and mortality in Mozambique especially among the children and pregnant women
- Almost 100% of the population lives in a high transmission zones and are at risk of infection
- It is responsible for 60% of paediatric admissions and 23% of all in-hospital deaths
- The disease is predominantly caused by plasmodium falciparum, transmission is perennial with a seasonal peak in the rainy season (November-April)



## Introduction cont..

- Since 1940s, the country has been applying WHO recommended tools (IRS, ITN and provision of treatment using ACT)
- According 2009 WHO report, the country has registered a decline in overall malaria case except for children
- The relationship between malaria mortality and levels of transmission is still unclear
- EIR, one of the recommended indices for measuring transmission levels in areas of high intensity
- In 2002, INDEPTH initiated the MTIMBA project with the aim of assembling a database that can be used to address malaria transmission and mortality relationship



## Objectives

**General Objective:** develop statistical methods for analysing large DSS data in order to examine the relationship between mortality and malaria transmission

**Specific objectives:-**

- **to** identify environmental predictors for EIR
- to establish the relationship between EIR and various predictors
- to produce spatially explicit and season specific estimates of EIR



## Methodology

### Study site:

- Maniac DSS is located in southern Mozambique (at 25°24'S and 32°48'E)
- It has the natural catchment population of the District hospital
- DSS is situated in fertile lower lands and an escarpment of moderate height
- The area has two distinct seasons, warm (Nov and April) and a cool

**DSS data:** it is collected at very large number of households repeatedly over time, and thus they are correlated in space (*spatial*) and time (*temporal*).

- Common exposures (climatic), may influence mortality similarly in households of the same geographical area, introducing spatial correlation in mortality and malaria transmission



## Entomological data

- Entomological data were collected from randomly selected compounds using both LTC and HBC
- LTC were performed overnight/over 2 consecutive nights
- HBC were carried out monthly for calibration purposes against LTC
- Mosquito catches were performed between 6.00pm to 6.00am
- LTC anopheles mosquitoes were tested for plasmodium falciparum using ELISA



## Environmental data

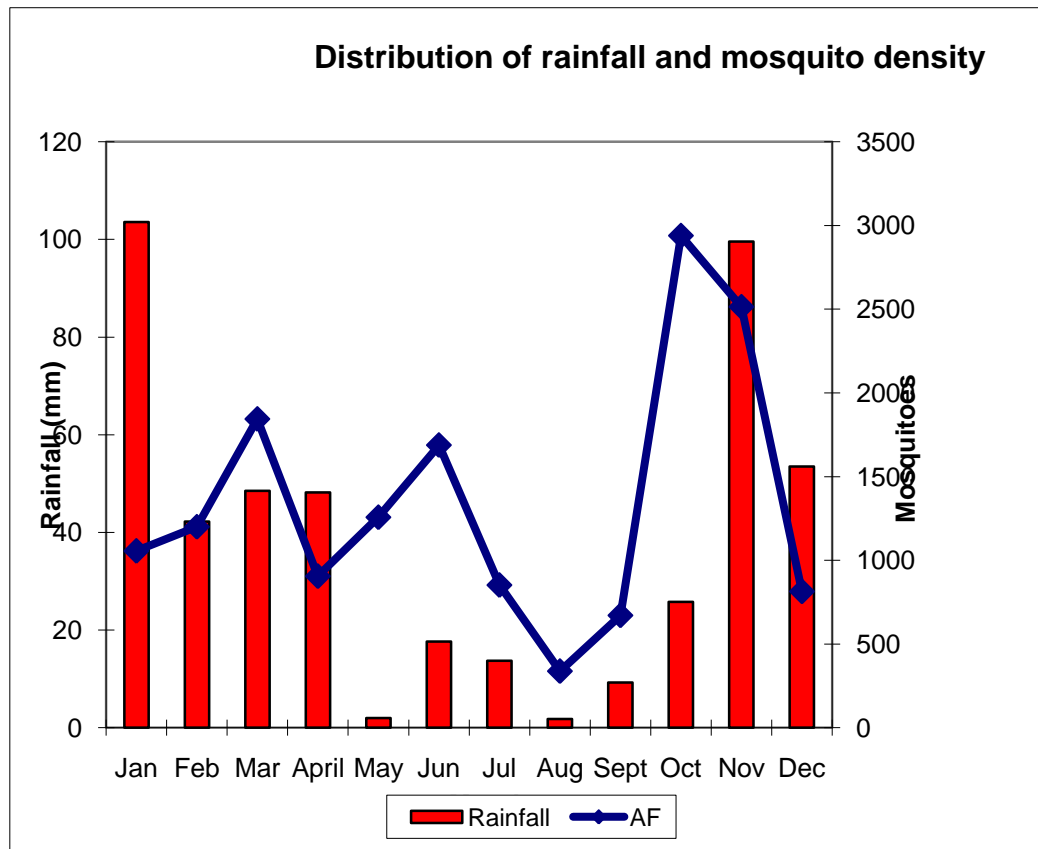
Predictor	Spatial Resolution	Temporal Resolution	Source
<b>Day land surface temperature (Day LST)</b>	1 km <sup>2</sup>	8 days	<b>MODIS</b>
<b>Night</b> land surface temperature ( <b>Night LST</b> )	1 km <sup>2</sup>	8 days	<b>MODIS</b>
<b>Normalized difference vegetation index (NDVI)</b>	250 m <sup>2</sup>	16 days	<b>MODIS</b>
<b>Enhanced Vegetation Index (EVI)</b>	250 m <sup>2</sup>	16 days	<b>MODIS</b>
<b>Rainfall estimate (RFE)</b>	8 km <sup>2</sup>	Dekadal	<b>ADDS</b>
<b>Elevation/Altitude</b>	1 km <sup>2</sup>	-	<b>USGS</b>
<b>Nearest distance to water bodies (rivers and wetlands)</b>	-	-	<b>Manhica DSS</b>



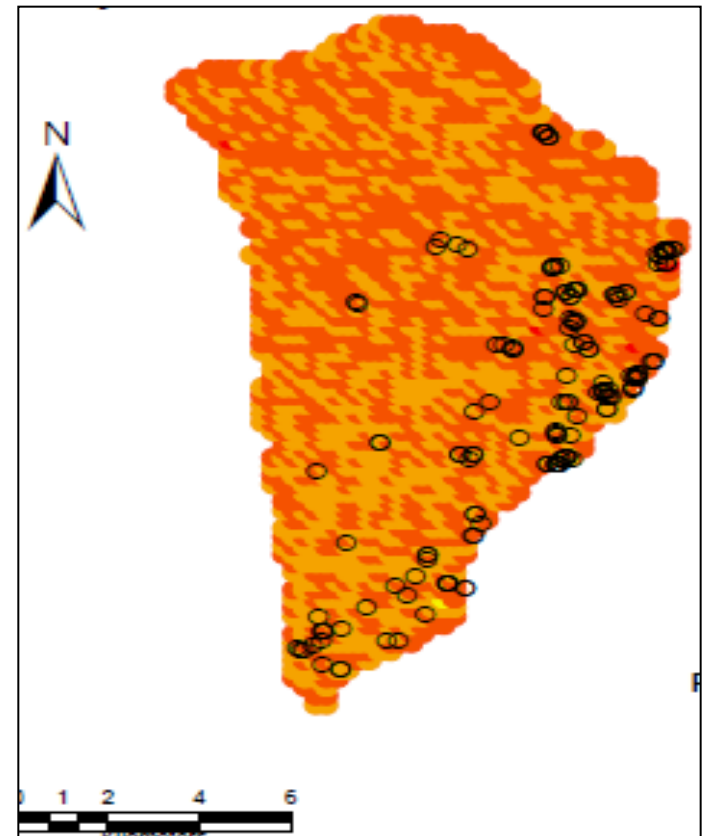


# Spatio-temporal distribution

## Temporal distribution



## Spatial distribution





## Spatio-temporal models

- Classical regression models assume independence of events which is not the case all the time
- Spatio-temporal data are correlated in space and time
- Ignoring correlation leads to under estimation of the standard error and overestimation of covariates' significance
- Spatial models incorporate spatial correlation
- Spatial models can be used to predict outcomes at new locations

## Model fit

Logistic regression model

$$Y_{it} \sim \text{Bn } N_{it}, P_{it} \quad \text{Where } N_{it} \text{ number of tested mosquitoes}$$

$P_{it}$  sporozoite rate

$$\log \text{it}(P_{it}) = X_{it}^T \beta + f(t)$$

Where  $f(t) = \varphi_1 \cos\left(\frac{2\pi t}{T}\right) + \varphi_2 \sin\left(\frac{2\pi t}{T}\right)$ , the seasonality function

$\beta$  Regression parameters



## Spatio-temporal Logistic regression model

$$\log it(P_{it}) = X_{it}^T \beta + f(t) + \phi_i + \epsilon_t$$

Where  $\phi_1, \dots, \phi_{n-1}, \phi_n \sim MVN(0, \Sigma)$

$$\Sigma_{ij} = \sigma^2 \exp -\rho d_{ij} \quad \text{and} \quad \rho > 0$$

$\sigma^2$  Spatial variance

$d_{ij}$  distance between locations i and j

$\rho$  the correlation parameter

$$\epsilon_t \sim AR(1)$$



## Model fit (2)

Negative Binomial model for count data

$Y_{it} \sim NB(p, r)$  Where  $r$  dispersion parameter

$P$  Probability of success

$$P = \frac{r}{r + \mu}$$

$$\log(\mu_{it}) = X_{it}^T \beta + f(t)$$

Where  $f(t) = \varphi_1 \cos\left(\frac{2\pi t}{T}\right) + \varphi_2 \sin\left(\frac{2\pi t}{T}\right)$ , the seasonality function

$\beta$  Regression parameters



## Spatio-temporal negative binomial

$$\log(\mu_{it}) = X_{it}^T \beta + f(t) + \phi_i + \epsilon_t$$

Where  $\phi_1, \dots, \phi_{n-1}, \phi_n \sim MVN(0, \Sigma)$

$$\Sigma_{ij} = \sigma^2 \exp -\rho d_{ij} \quad \text{and} \quad \rho > 0$$

$\sigma^2$  Spatial variance

$d_{ij}$  distance between locations i and j

$\rho$  the correlation parameter

$$\epsilon_t \sim AR(1) \quad (\text{temporal term})$$



## Data Analysis

- Lag time analysis was used in order to account for elapsing time of environmental factors on SR and MBR
- Seasonality was accounted for using dry/wet binary variable or a mixture of trigonometric functions
- Bayesian method and Markov chain Monte Carlo simulation were used to estimate model parameters.
- Due to large number of locations, the spatial process was estimated using a subset
- Analysis was performed using Stata ver 10, OpenBUGS and R softwares

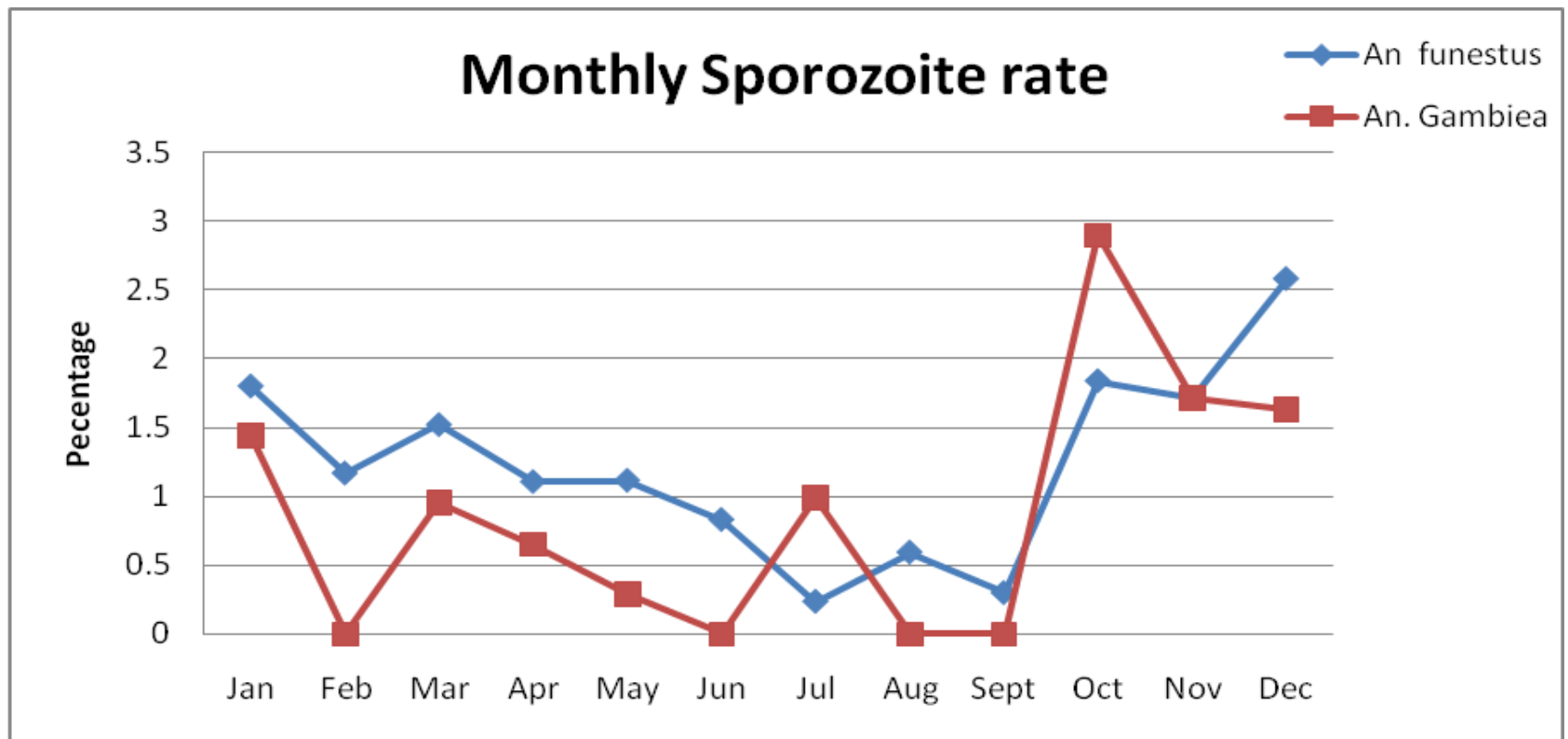


## Results

- Total number of *An. funestus* was 16078 (85%)
- Total number of *An. gambiae* was 2845 (15%)
- Overall sporozoite rate for AF was 1.4% and 1.1% (AG)

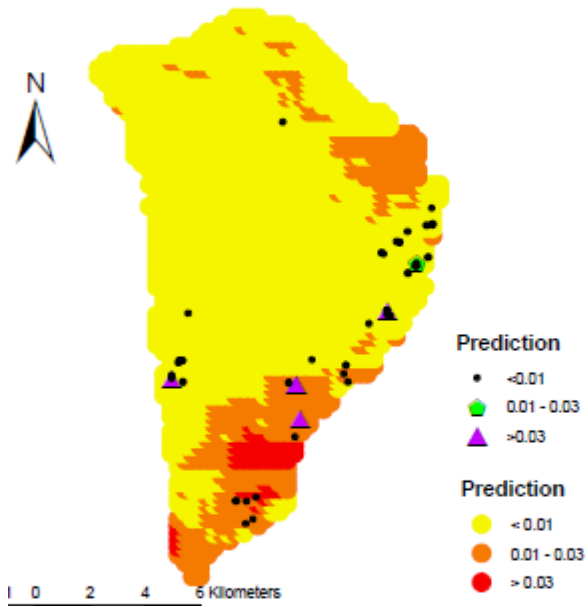


## Observed sporozoite rates

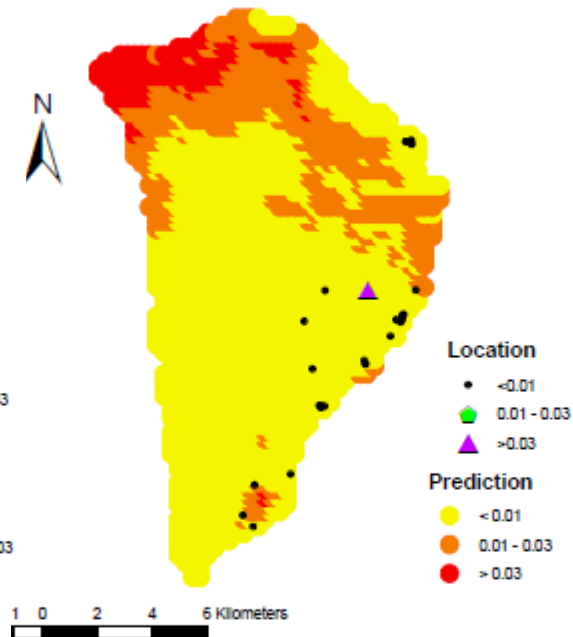


# Predicted SR using ZIB model (AG)

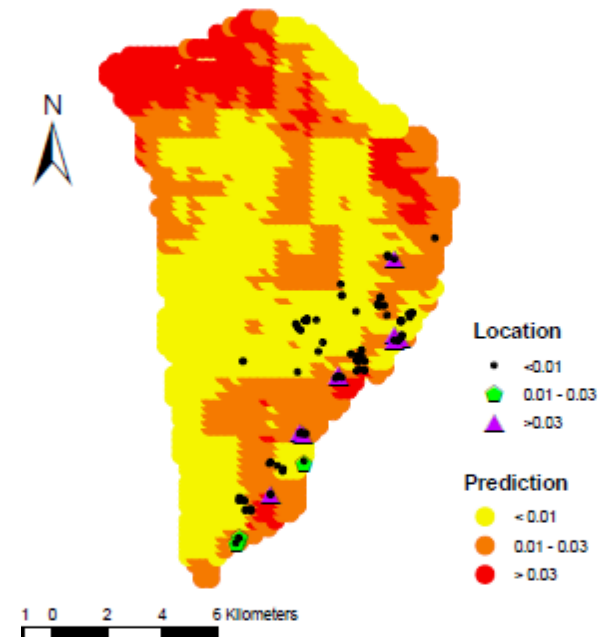
January



July

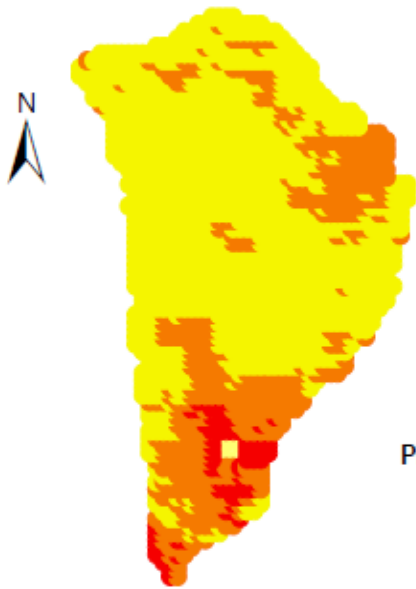


December



## Prediction errors for SR using ZIB model

January

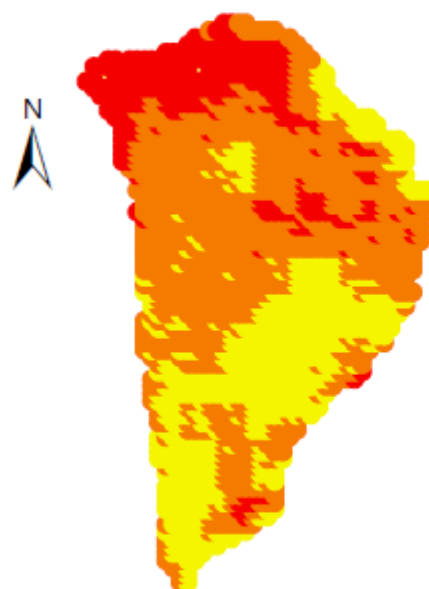


Prediction err

- < 0.01
- 0.01 - 0.0
- > 0.03

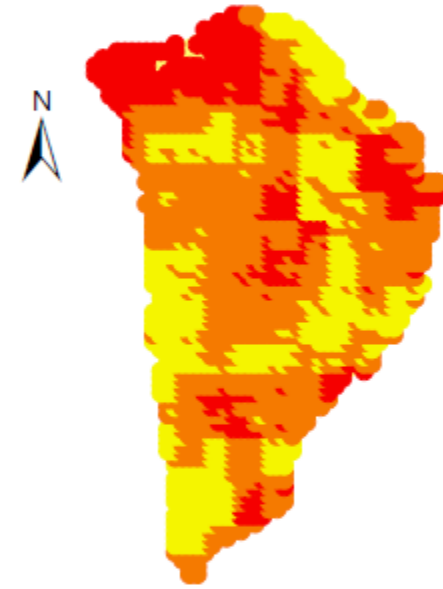
0.50 1 2 3 Kilometers

July



0.50 1 2 3 Kilometers

December



Prediction errors:

- < 0.01
- 0.01 - 0.03
- > 0.03

0.50 1 2 3 Kilometers

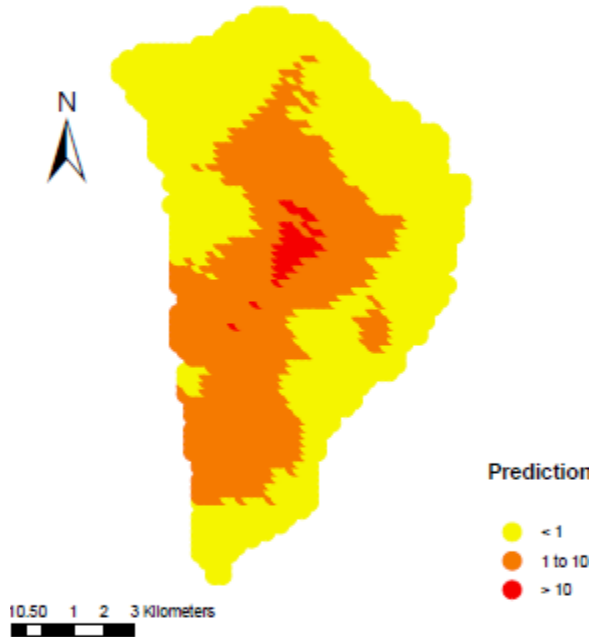
Prediction errors

- < 0.01
- 0.01 - 0.03
- > 0.03

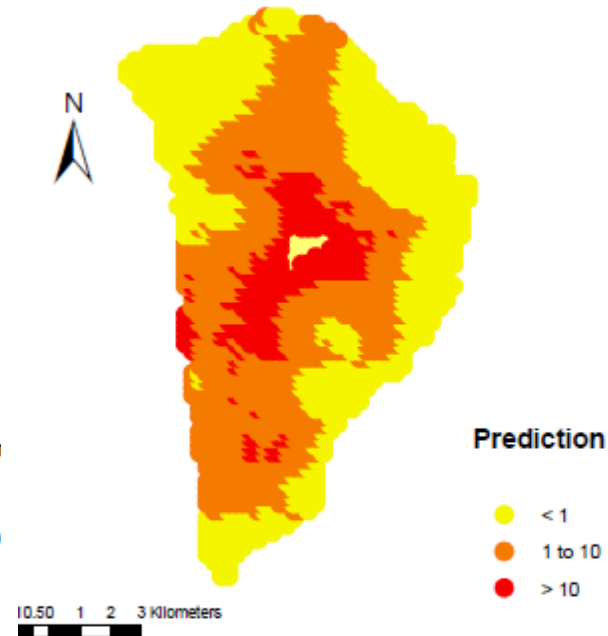


## Predicted mosquito density using ZINB (AG)

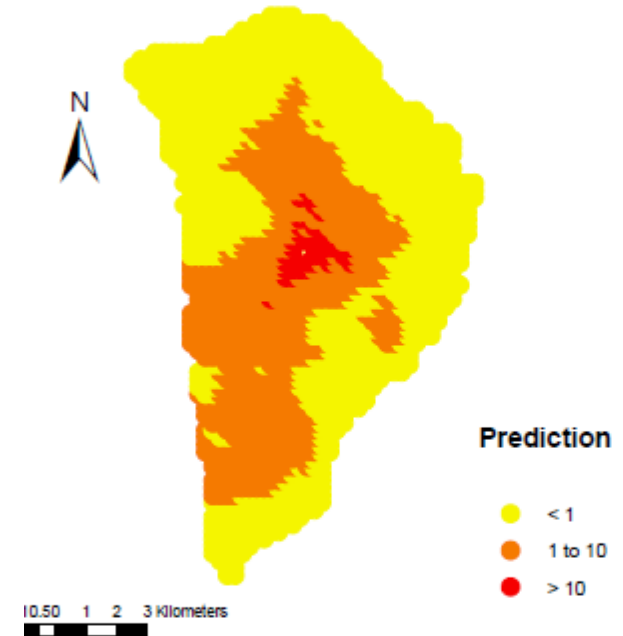
January



March

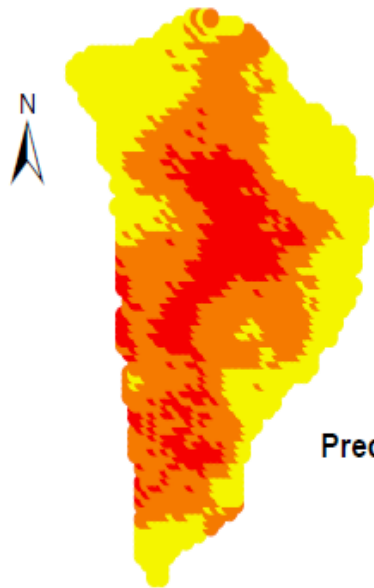


December



## Prediction errors for Density using ZIB model

January

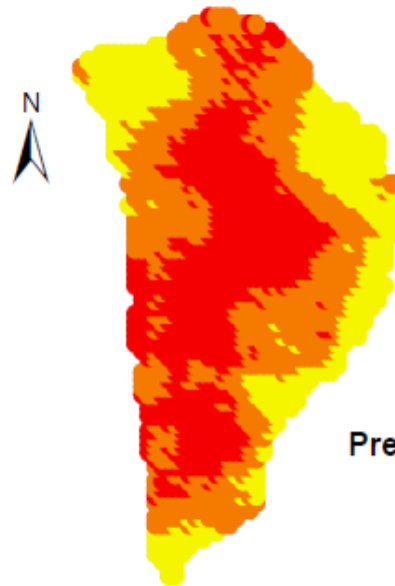


Prediction errors

- <1
- 1-10
- >10

0.50 1 2 3 Kilometers

March

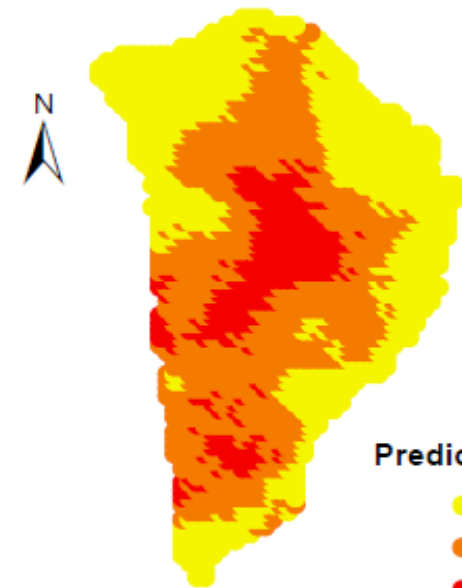


Prediction errors

- <1
- 1-10
- >10

0.50 1 2 3 Kilometers

December



Prediction error

- <1
- 1-10
- >10

0.50 1 2 3 Kilometers



## Discussion

- Accurate maps are essential tools needed in establishing the effectiveness of malaria control interventions
- Malaria transmission in sub-Saharan Africa is heterogeneous and environmentally driven. Application of spatio-temporal models is a necessary measure for developing early warning sign for a particular country.
- Entomological data is very sparse making statistical analysis complicated



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***THANK YOU !!!!***



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