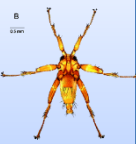
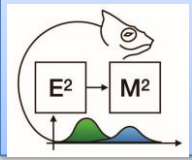


Bartonella dynamic in fruit bats and bat flies



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Background:

- *Bartonella* a hemoparasite transmitted by vector to the host;
- Little is known about *Bartonella* in Madagascar fruit bats

Questions:

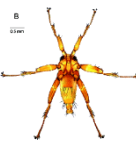
Q1-What are the factors associated with the prevalence of *Bartonella* ssp in Madagascar fruit bats?

Q2-How does batflies prevalences affect the dynamics of *Bartonella* ssp in fruit bats through time?

Acknowledgement:

All Instructors & Mentors at E^2M^2 , Herilanto Ramaroson & Nantenaina Rindraharilanto

Statistical question:



What are the factors associated with the prevalence of *Bartonella ssp* in Madagascar fruit bats?

Response variable: Presence of *Bartonella* in bats (bat.pos);

Predictor variables: morphometrics(morph), species of the flies (sp.flies), abundance of batflies (n.flies), season, site

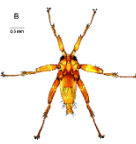
Distribution: Binomial

Link : Logit

Model : Random effect generalized linear mixte model

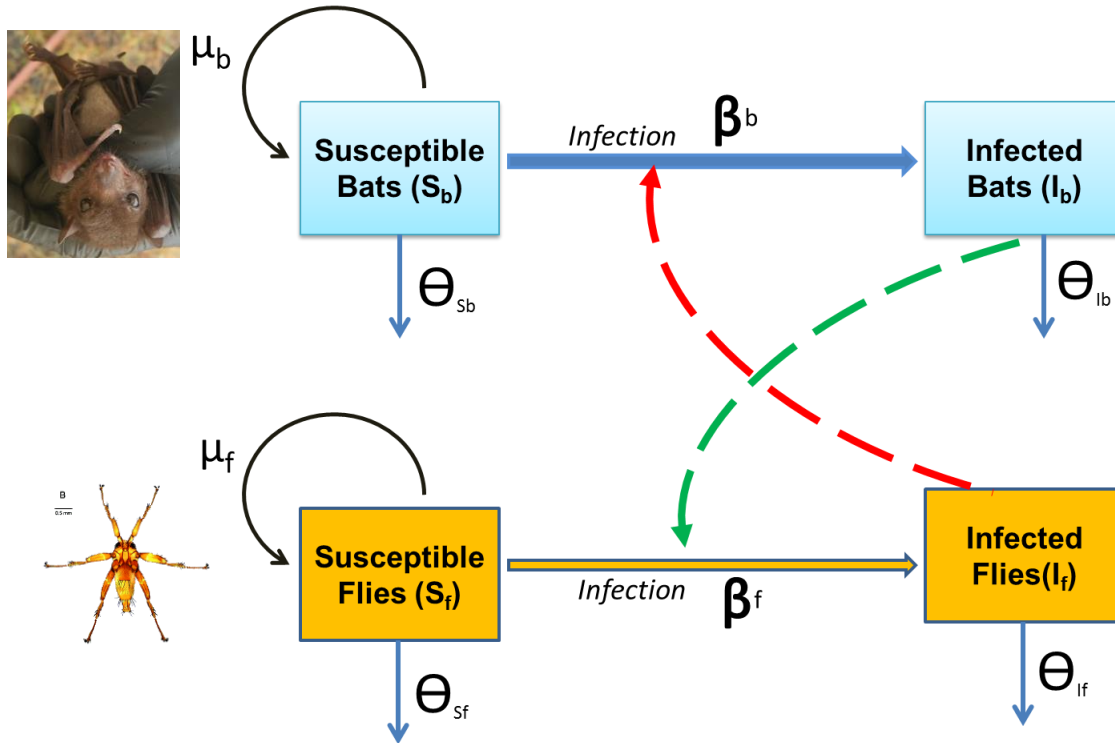
R_code:

```
> glmer(formula=bat.pos~morph+n.flies+sp.flies+season+(1 |site),  
family=« binomial »,data)
```



Dynamic question:

How does bat flies prevalences affect the dynamic of *Bartonella* ssp in Fruit bats population through time?



States:

- Susceptible bats (S_b)
- Infected bats (I_b)
- Susceptible flies (S_f)
- Infected flies (I_f)

Process:

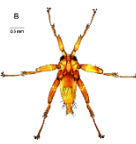
- Infection

μ_i : birth rate

Θ_i : death rate

β_i : infection rate

Next steps:



Data collection in the field and in the laboratory;



Simulate the mechanistic model and the statistical model ;



Fit the model to the data collected.