

Modeling Rift Valley fever virus transmission dynamics

Insight from micro- to macro-scale studies

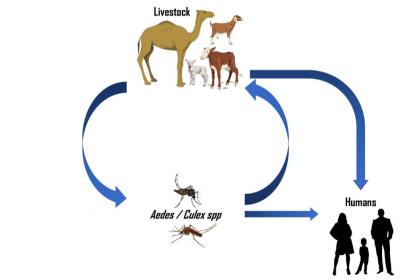
 $\frac{\partial e}{\partial t} + \frac{\partial}{\partial x}(e_{w}) = 0$

 $\frac{\partial u}{\partial t} + u \frac{\partial u}{\partial t} = -\frac{1}{6} \frac{\partial u}{\partial t}$

Hélène CECILIA

Rift Valley fever virus

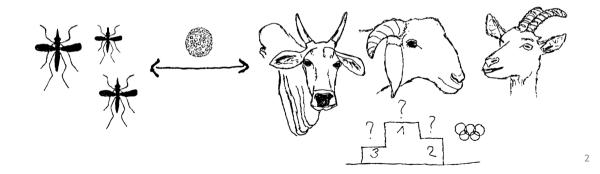




Research question

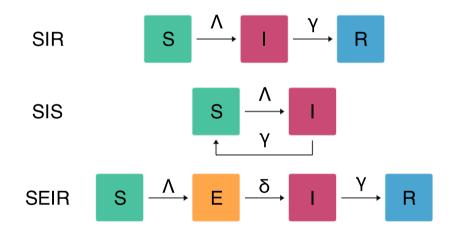


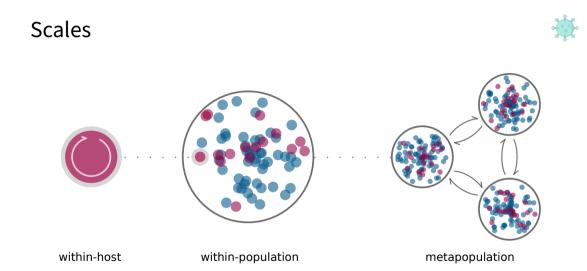
What is the relative contribution of livestock host species to RVFV transmission dynamics in Senegal?



Compartmental models







Epidemic potential in northern Senegal

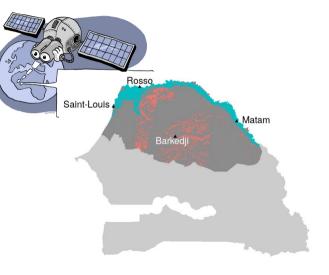
Cecilia et al. 2020 Epidemics

Objective



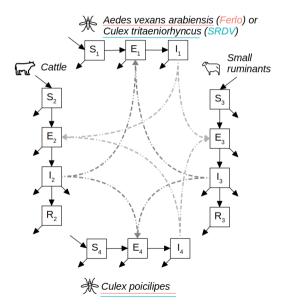
Explore RVFV suitability of two contrasted regions:

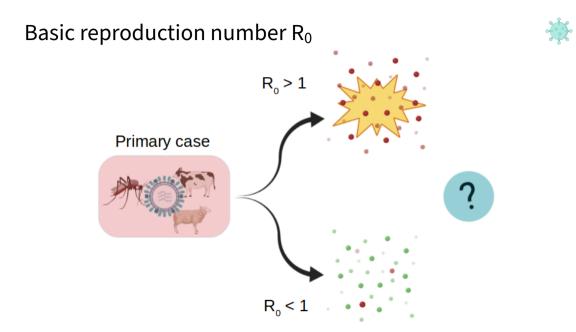
- Senegal river delta and valley (SRDV)
- ► Ferlo



Model

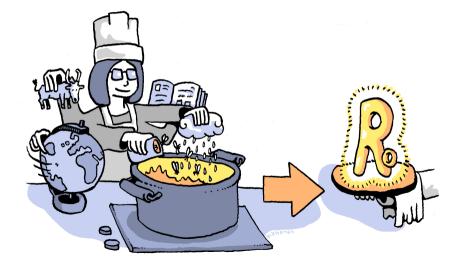






Basic reproduction number R₀





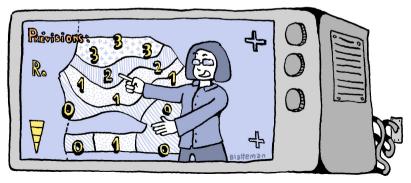
Results



September is the period of highest epidemic potential in northern Senegal

Immunity in cattle has a stronger effect on R₀ than in small ruminants

To be addressed : Hosts do not differ in their individual competence



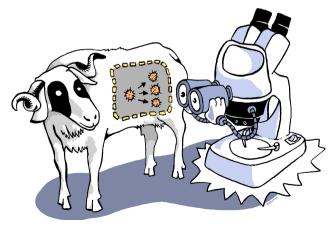
Transmission potential of livestock hosts

Cecilia et al. 2022 Plos Computational Biology

Objective

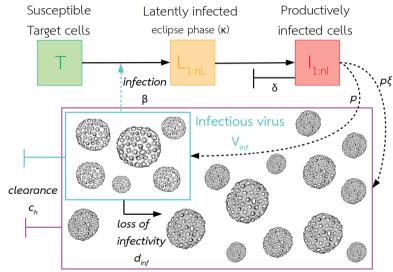


- Understand viral dynamics within RVFV livestock hosts
- Assess how efficiently livestock species can transmit RVFV to mosquitoes



Within-host model

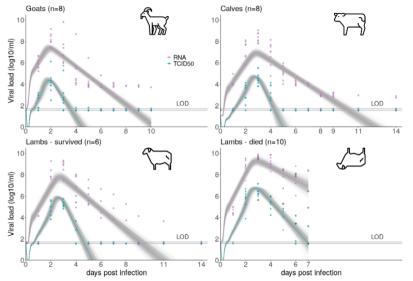




Total viral production V_{tot}

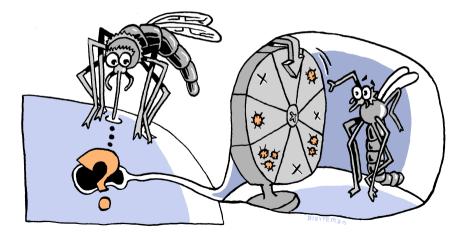
Model fit





Probability to infect mosquitoes





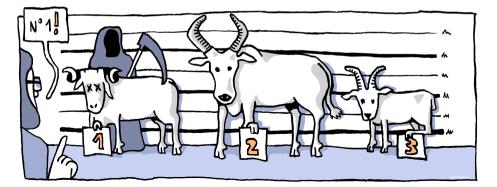
Results



Lambs are 3 to 4 times more infectious than cattle and goats

Lambs succumbing to RVF are more infectious than those less severely affected

<u>To be addressed</u> : How does this impact transmission at the population level?



Last chapter - Metapopulation model



Understand RVFV transmission dynamics at large scale,

taking into account individual heterogeneity,

in a context of seasonal, pendular animal mobility

Conclusion



Mathematical modeling is an efficient way to combine different sources of knowledge, particularly relevant to understand the complex eco-epidemiology of vector-borne, multi-host diseases

More data needed to confirm/infirm our conclusions and eventually lead to operational recommendations for RVF control in Senegal

Questions?

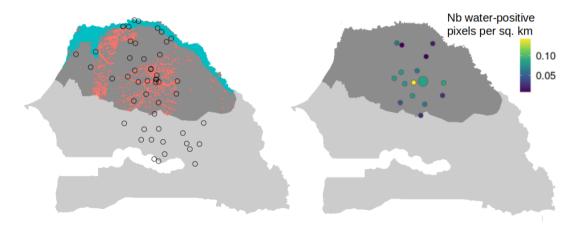
<u>My current job</u> : Postdoc New Mexico State Univ. (USA) Dengue and Zika virus in non-human primates



Metapopulation

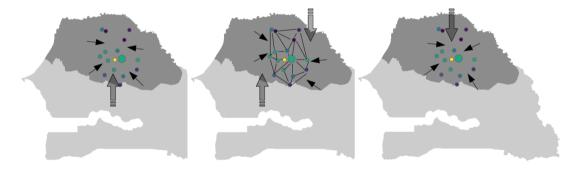
New data : field survey of 132 nomadic herders

From 4378 pixels... to 16 patches

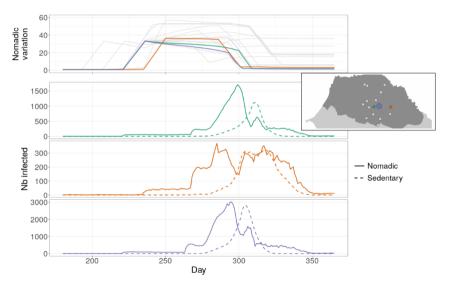


Metapopulation

Synthetic, seasonal movements during the rainy season



From nomadic to sedentary



Contribution of species

