

Model Selection and Comparison



Christian Ranaivoson

E2M2 2025 Andasibe, Madagascar

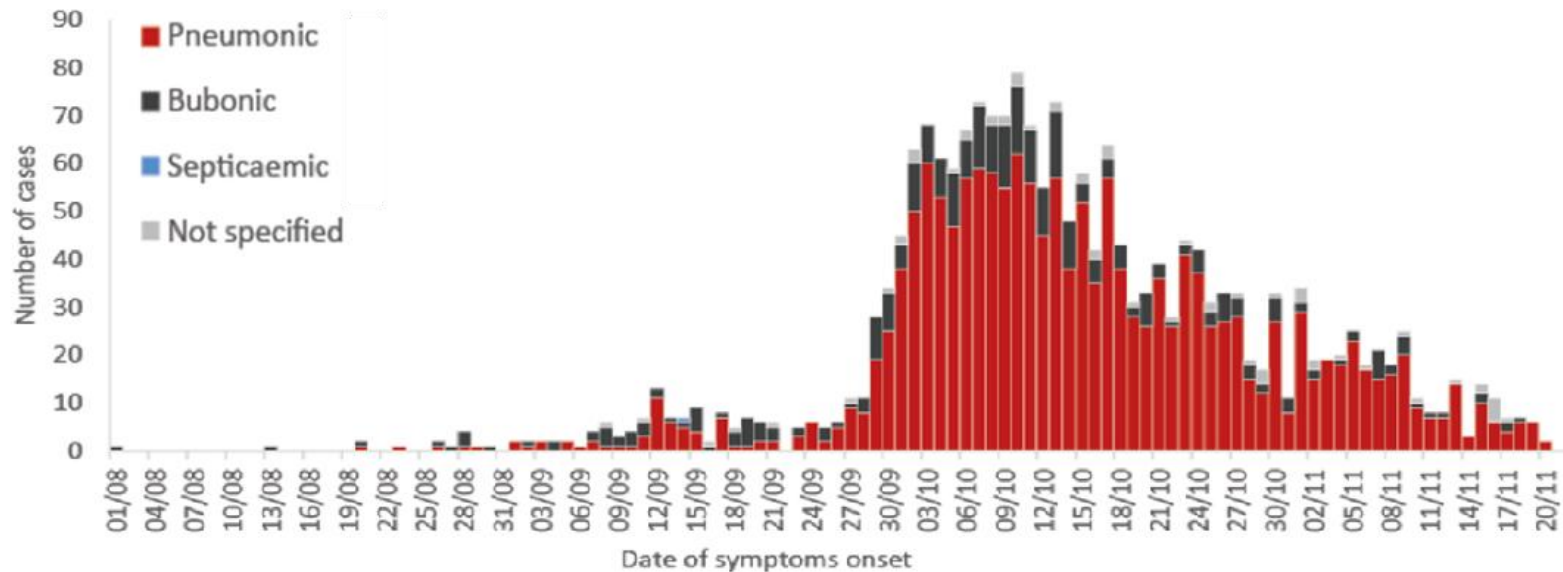
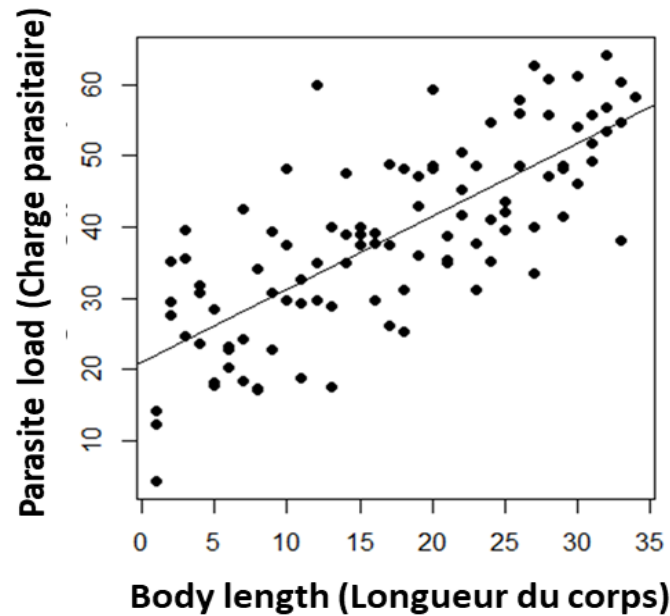
University of Chicago, USA ; Association Ekipa Fanihy; University of Antananarivo, Madagascar

Adapted from Cara Brook, Jessica Metcalf, Christian Ranaivoson E2M2 2022-2024.

Goals of the lecture

- Introduction to model selection
- Be familiar with common measures of model fit
- Tutorial on how to do it in practice (Real example)

Which model is best?



There are many statistical methods used to 'fit' models to data and there are many possible scenarios from which mechanical model can be built.

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The method best suited for your work will depend on your data and your question.

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What are some measures of model fit that you could use?

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Maximum likelihood (Maximum de vraisemblance)

(izay manakaiky indrindra ny tena izy ry reto an)

R-squared (R-carré)

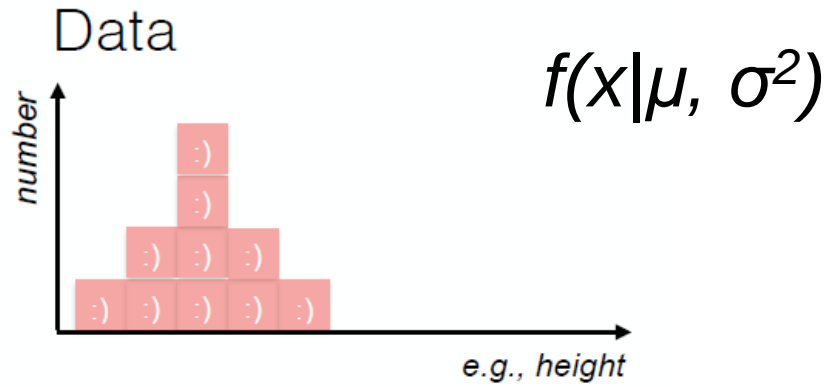
Least squares (Moindres carrés)

AIC

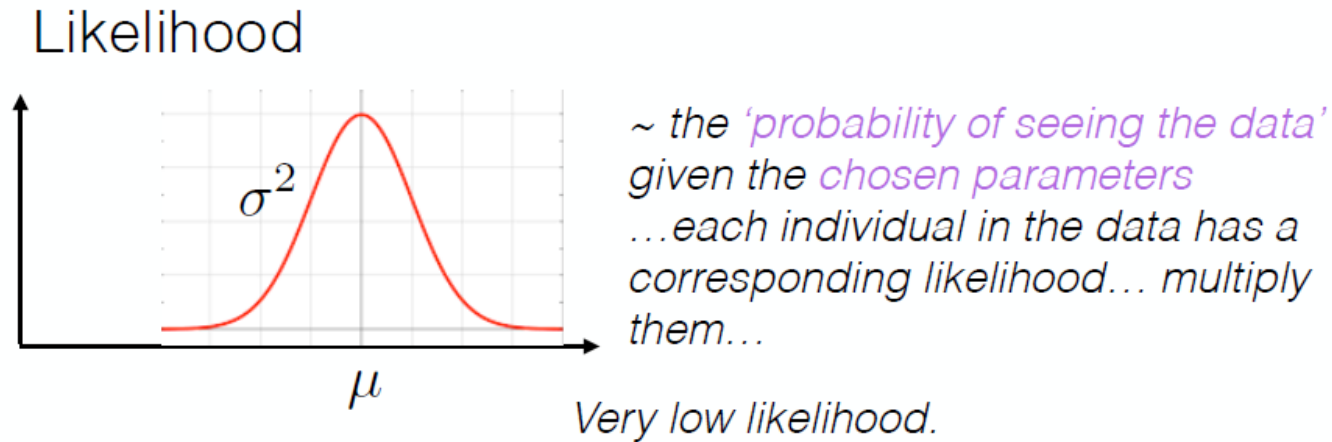
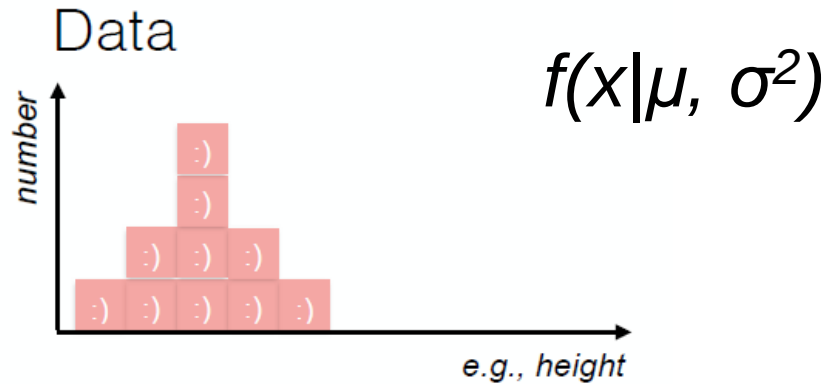
(uses least squares or log-likelihood but penalizes by number of fitted parameters)



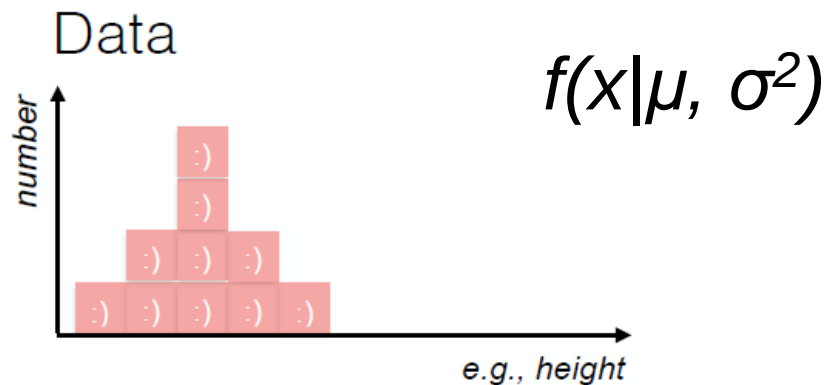
Maximum likelihood



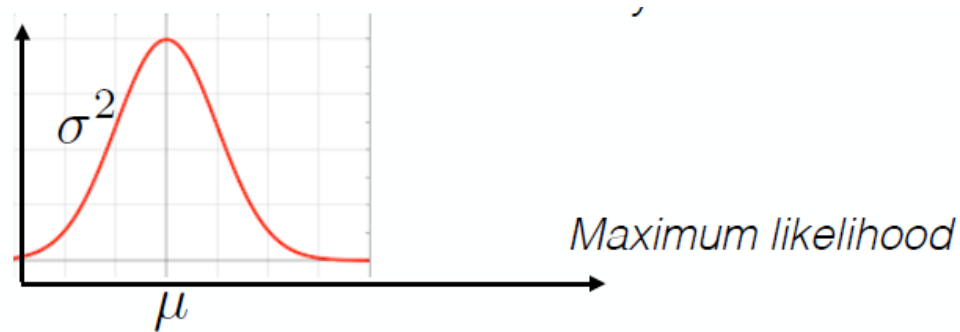
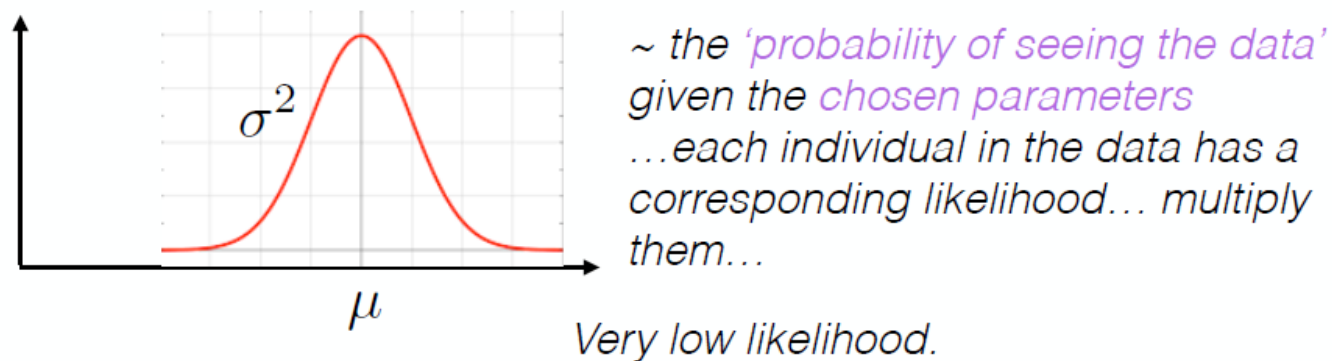
Maximum likelihood



Maximum likelihood



Likelihood



Likelihood Vraisemblance

$$L(\theta) = \prod_{i=1}^n f(x_i|\theta)$$

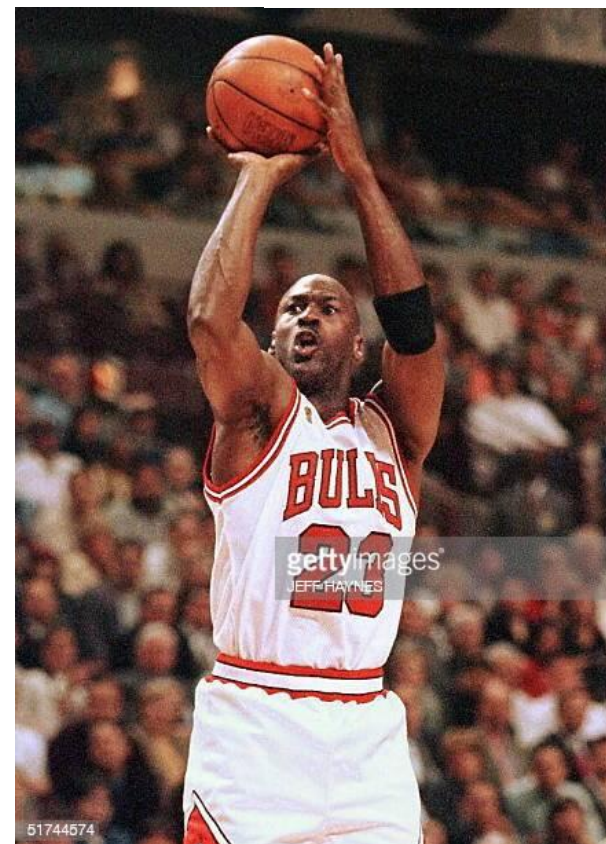
$$l(\theta|x) = \log L(\theta|x)$$

Examples:

R function : **dbinom(x, size, prob, log=T)**

➡ R function : **dbinom(8, 10, 0.835) = ?**

SUMMARY	FG3%	FT%
Career	32.7	83.5



Likelihood Vraisemblance

$$L(\theta) = \prod_{i=1}^n f(x_i|\theta)$$

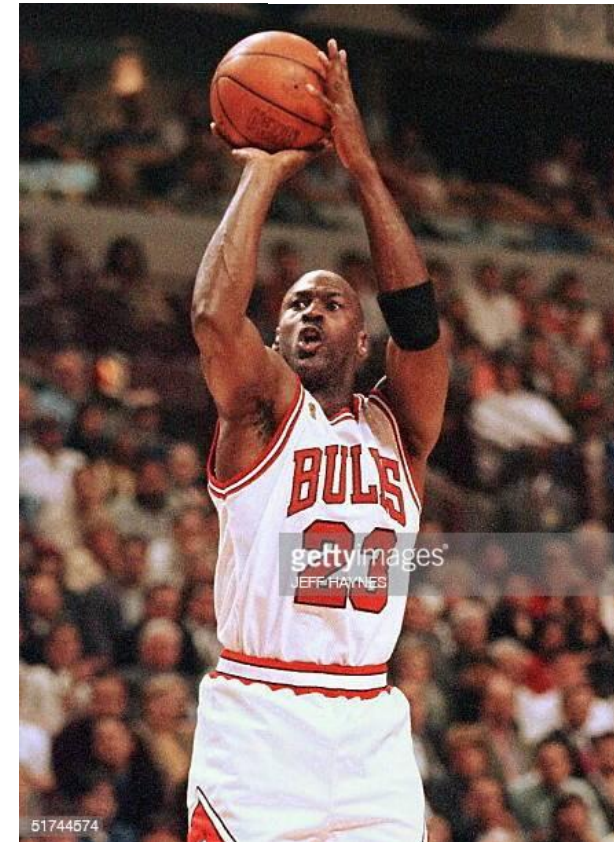
$$l(\theta|x) = \log L(\theta|x)$$

Examples:

R function : **`dbinom(x, size, prob, log=T)`**

➡ R function : **`dbinom(8, 10, 0.835) = 0.289`**

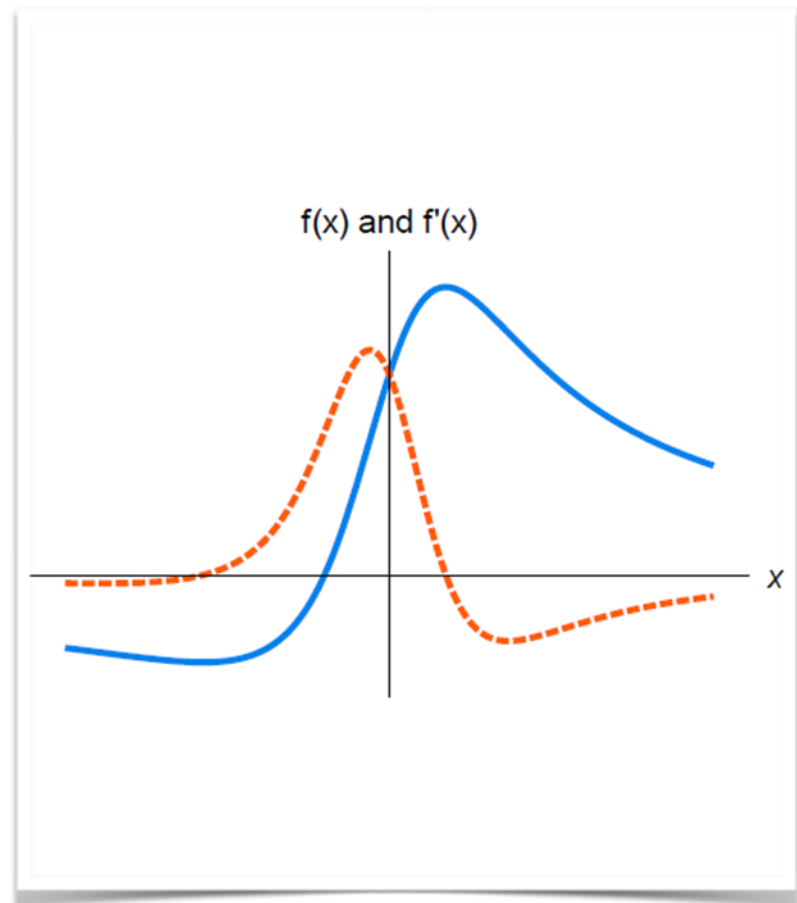
SUMMARY	FG3%	FT%
Career	32.7	83.5



Optimization/maximization

A function and its derivative

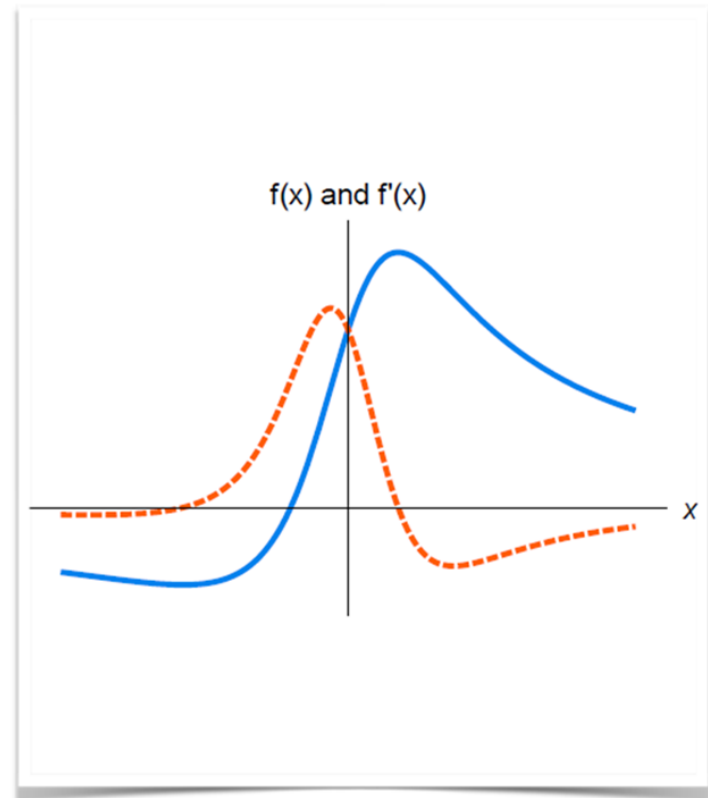
- ❖ What happens when the derivative is:
 - ❖ negative?
 - ❖ positive?
 - ❖ zero?
 - ❖ reaching a maximum (finite) value?



Optimization/maximization

A function and its derivative

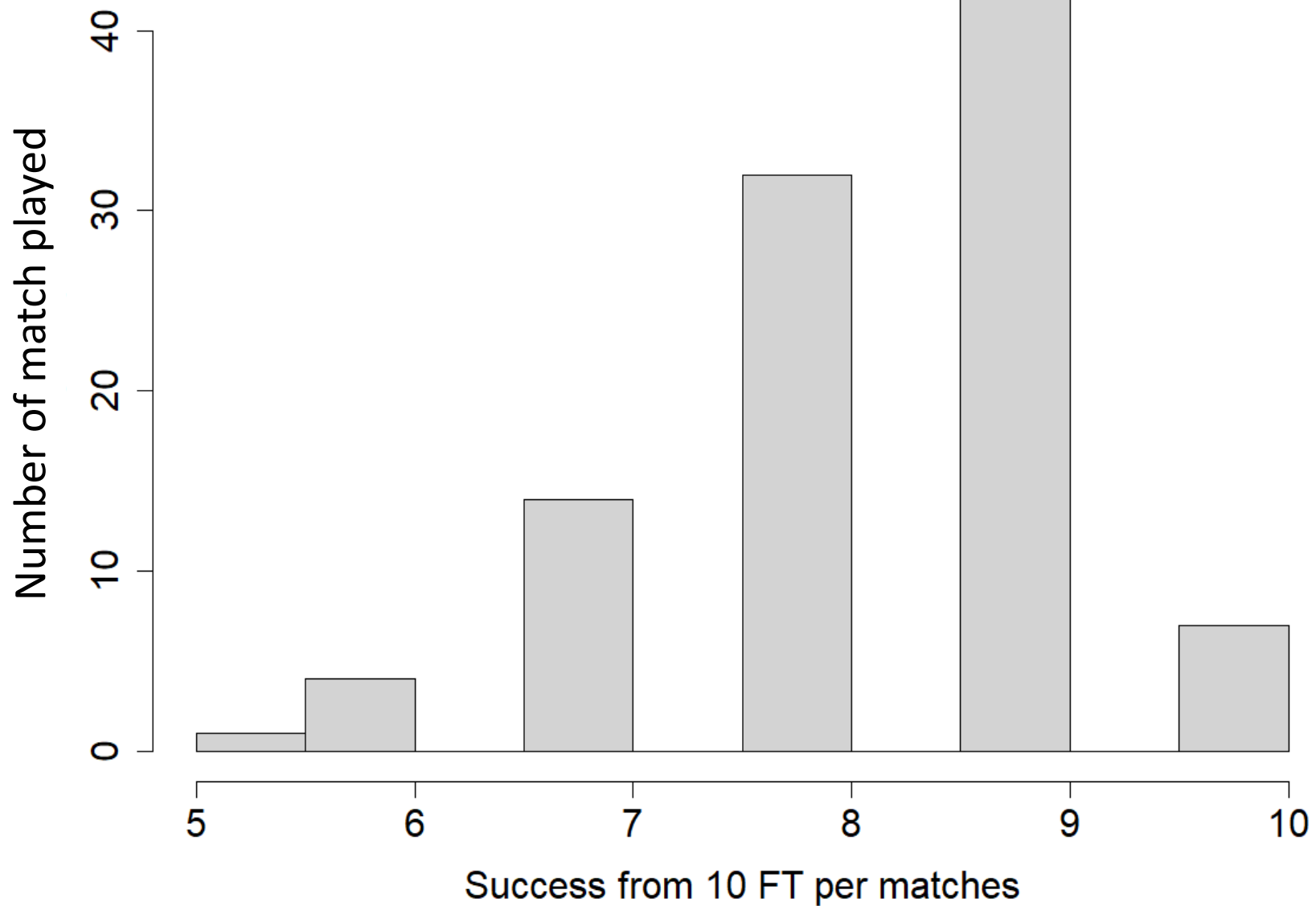
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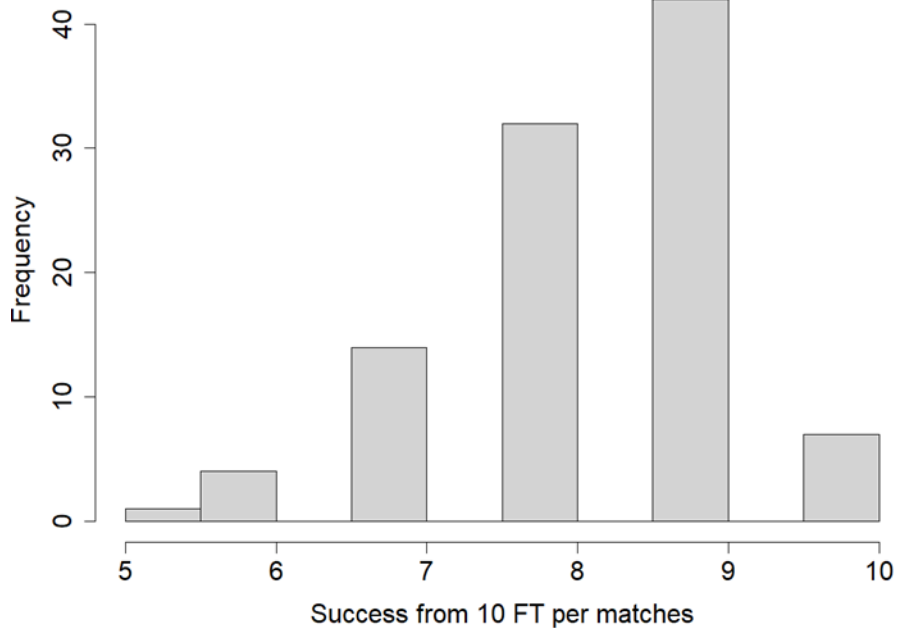
From Tanjona Ramiadantsoa

The R function 'optim' can be used to find minimum/maximum.

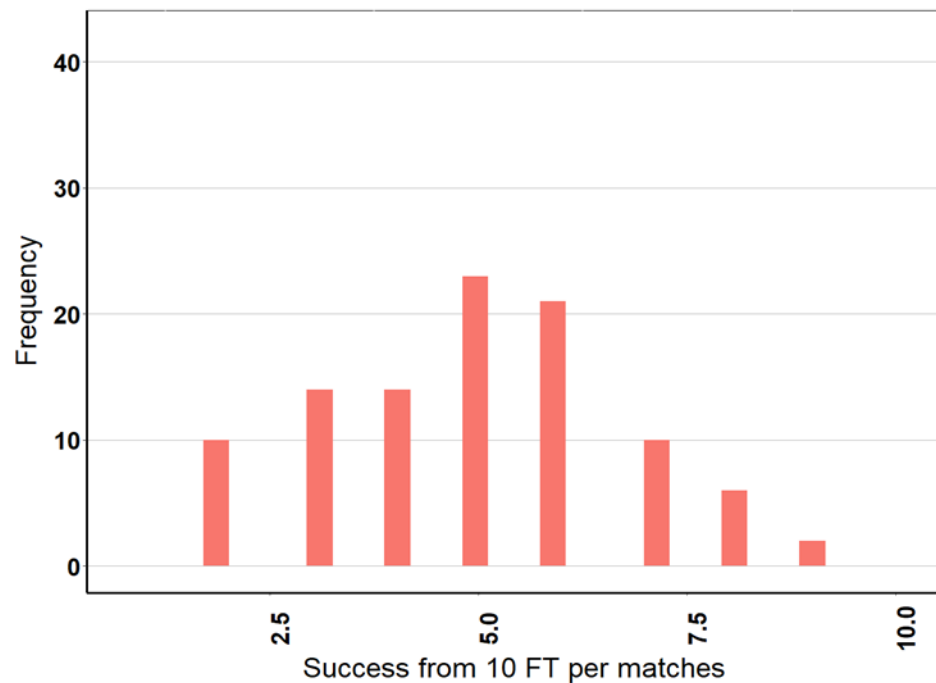
Michael Jordan's FT success in 100 match career



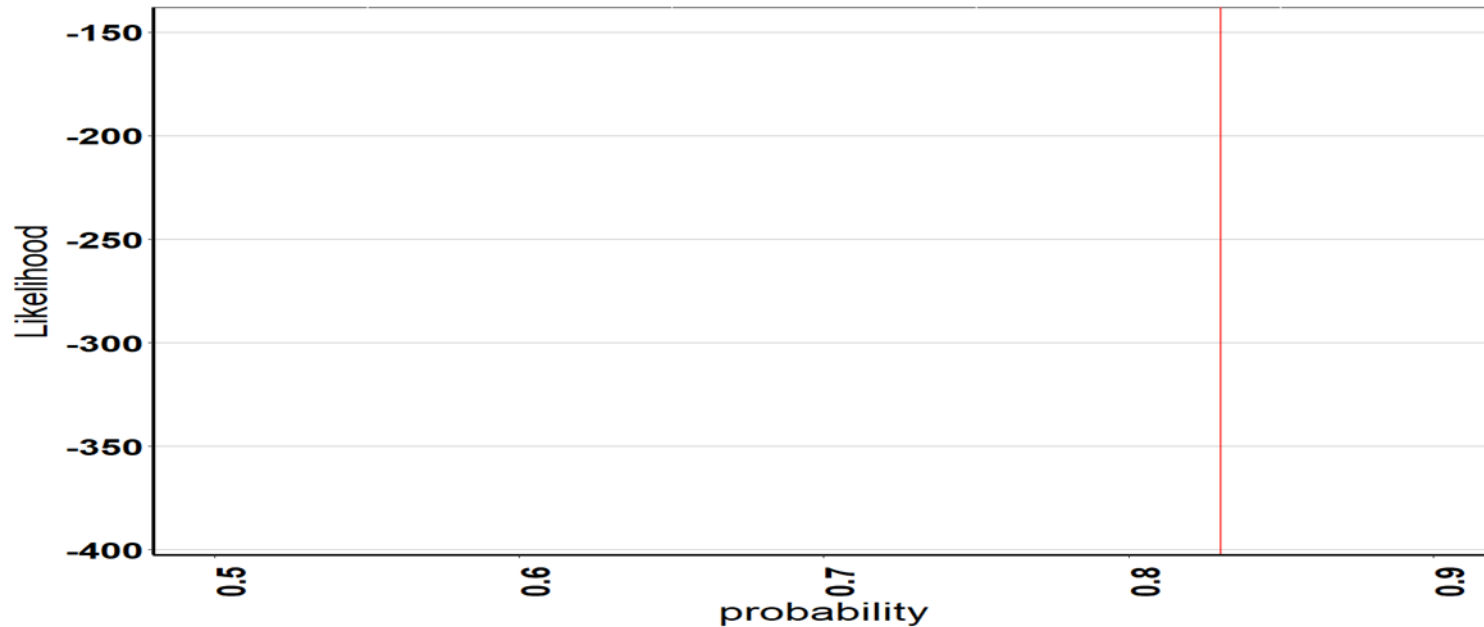
Michael Jordan's FT success in 100 match career



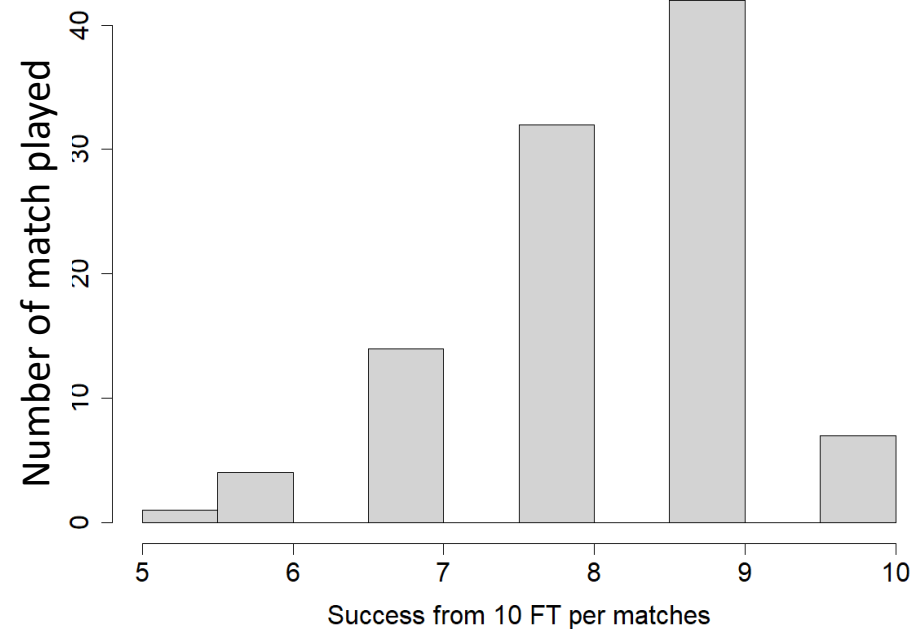
Michael Jordan's FT success in 100 match career



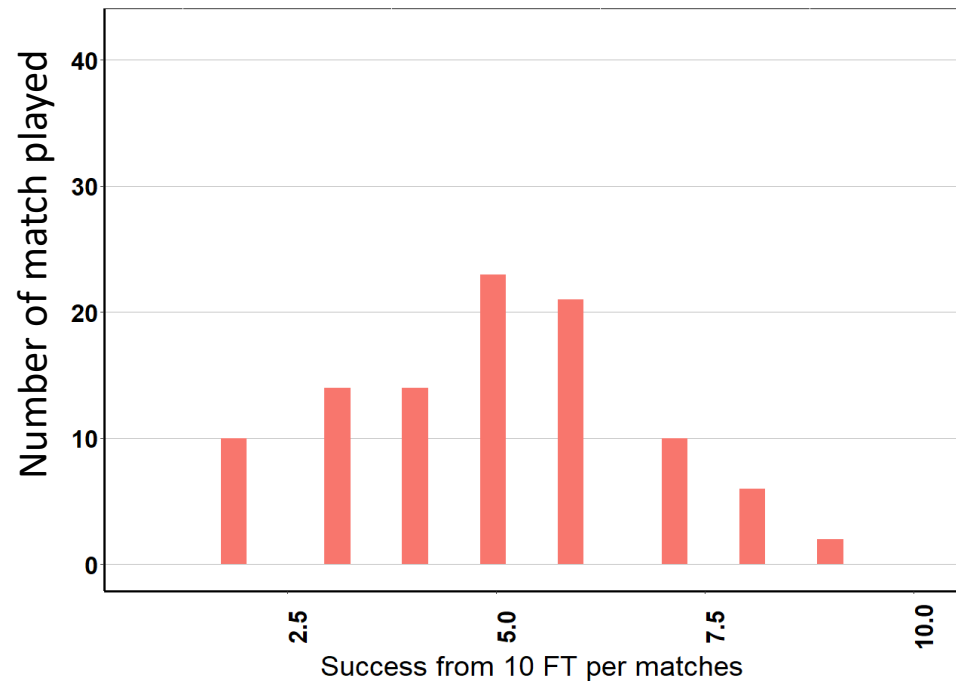
Michael Jordan's FT success in 100 match career



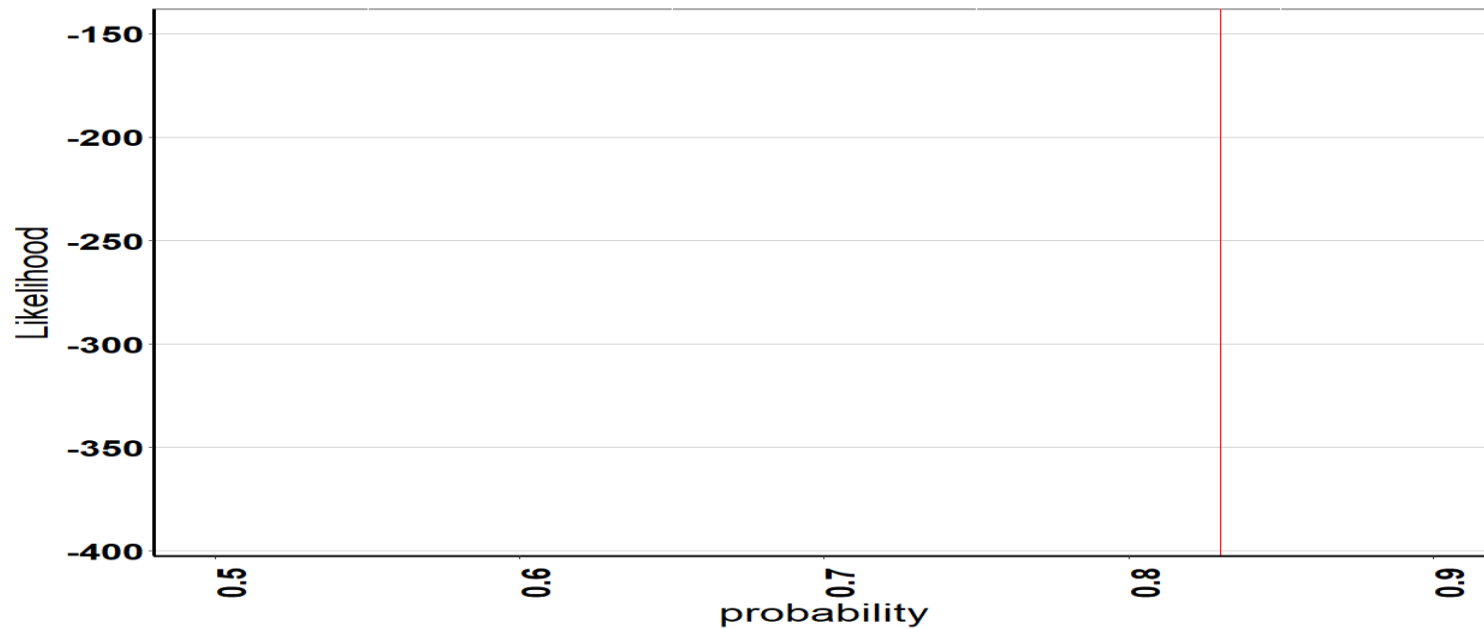
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Michael Jordan's FT success in 100 match career

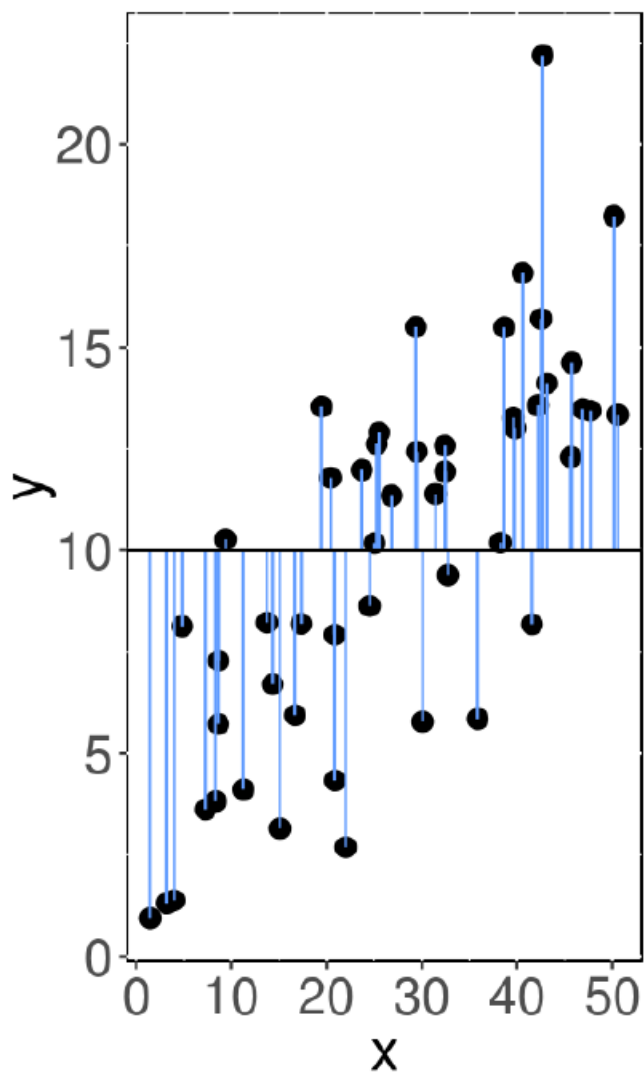


Michael Jordan's FT success in 100 match career

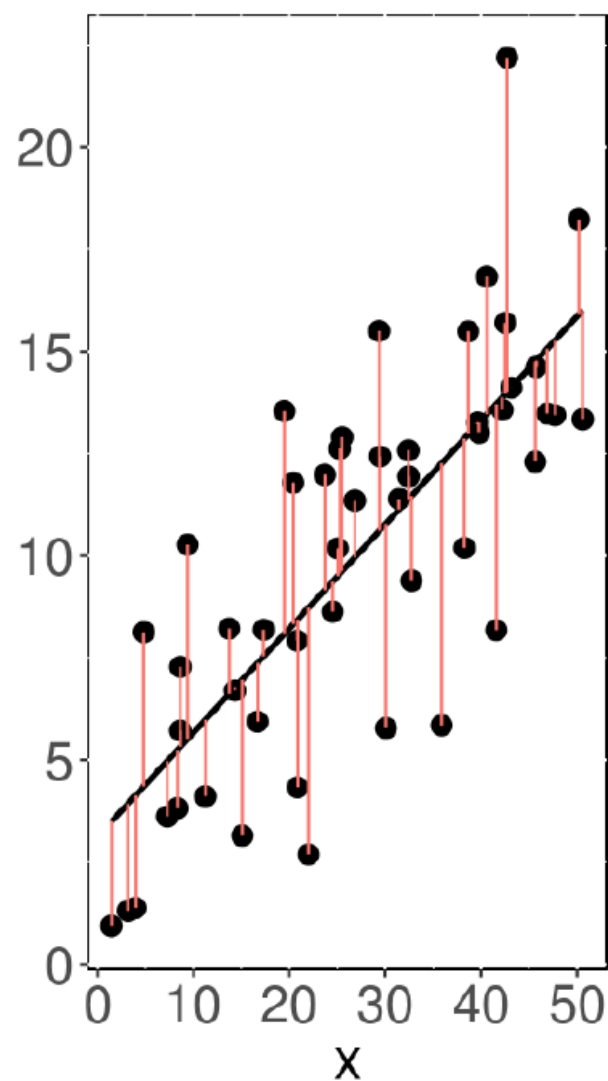


Definition r^2

SS total



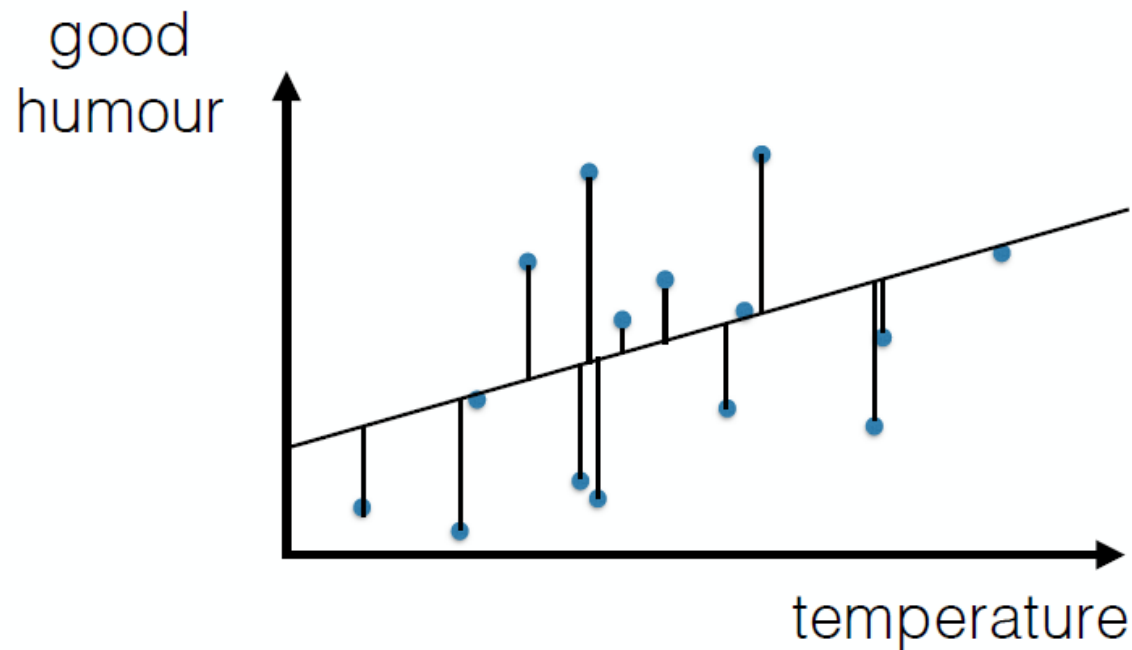
SS error



$$R^2 = 1 - \frac{SSE_p}{SST}$$

Least squares

Adding covariates and R^2

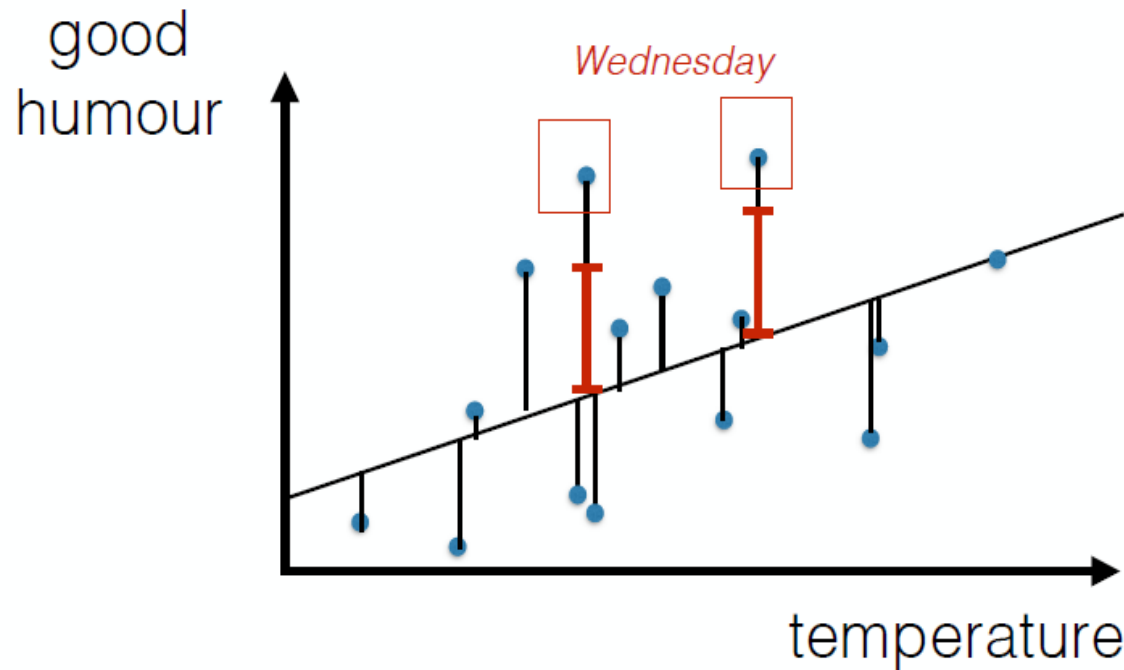


2K

$$\text{humour} = b_0 + b_1 \text{temperature} + \text{Error}$$

Least squares

Adding covariates and R^2

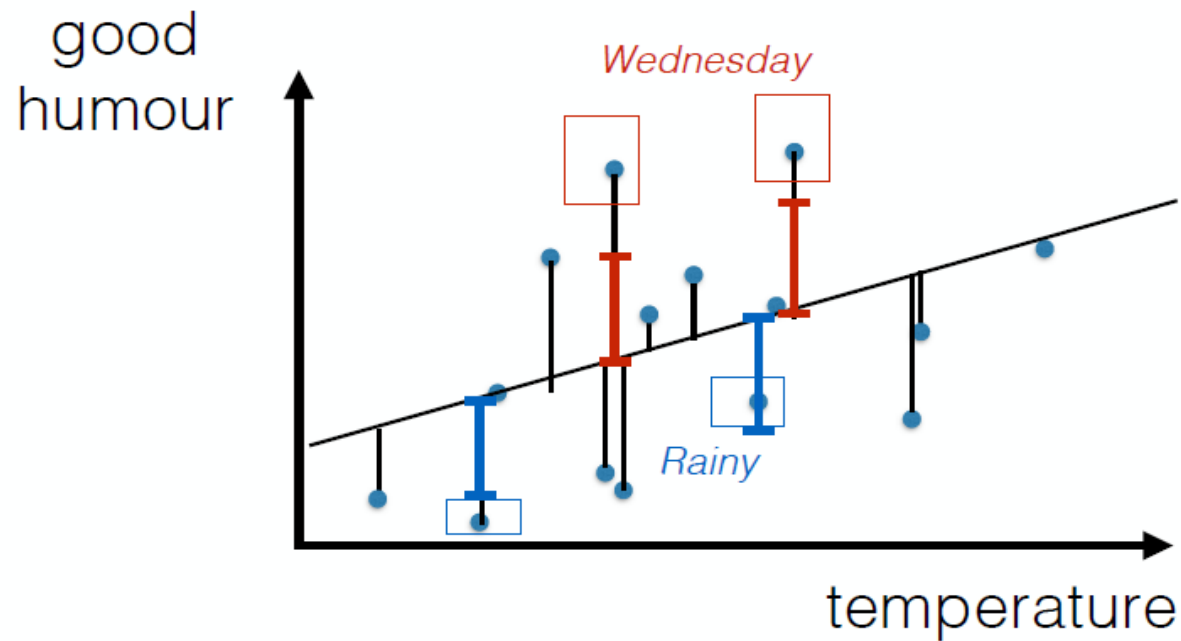


3K

$$\text{humour} = b_0 + b_1 \text{temperature} + b_2 \text{Wednesday} + \text{Error}$$

Least squares

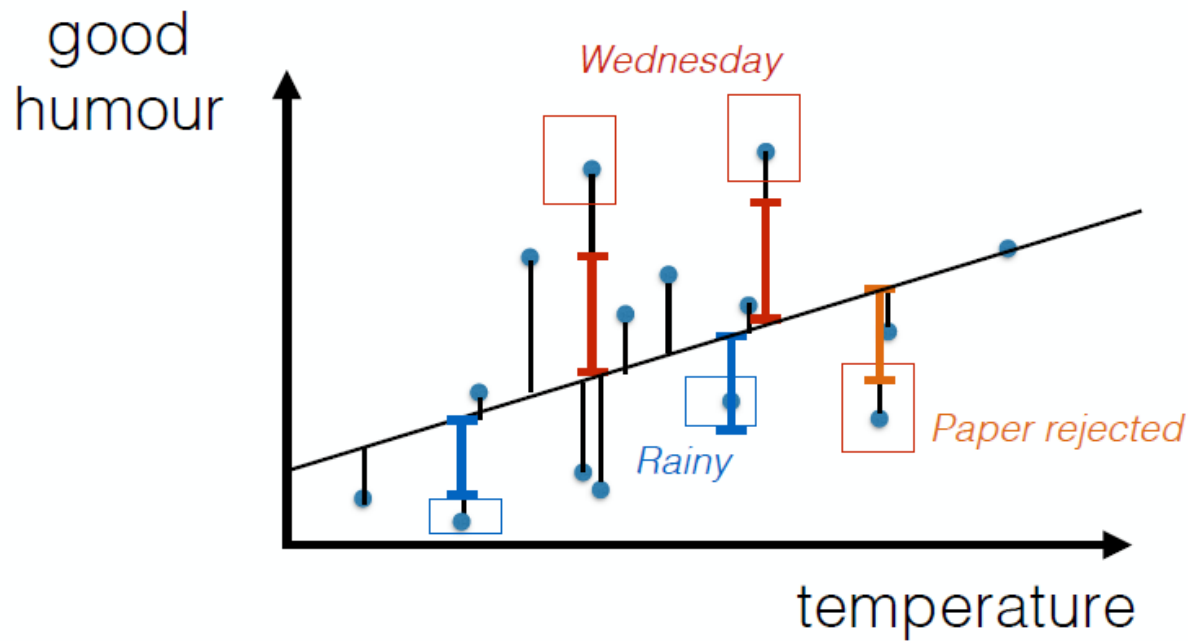
Adding covariates and R^2



$$\text{humour} = b_0 + b_1\text{temperature} + b_2\text{Wednesday} + b_3\text{rain} + \text{Error}$$

Least squares

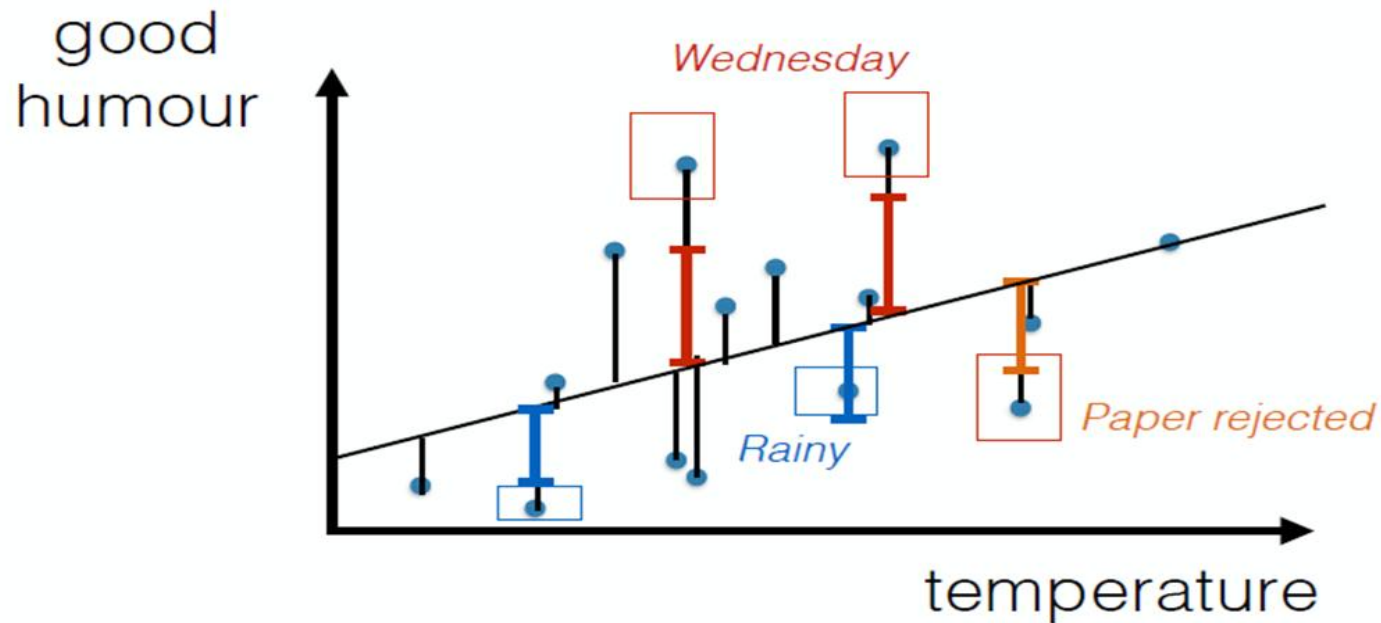
Adding covariates and R^2



$$\text{humour} = b_0 + b_1\text{temperature} + b_2\text{Wednesday} + b_3\text{rain} + b_4\text{rejection} + \text{Error}$$

Least squares

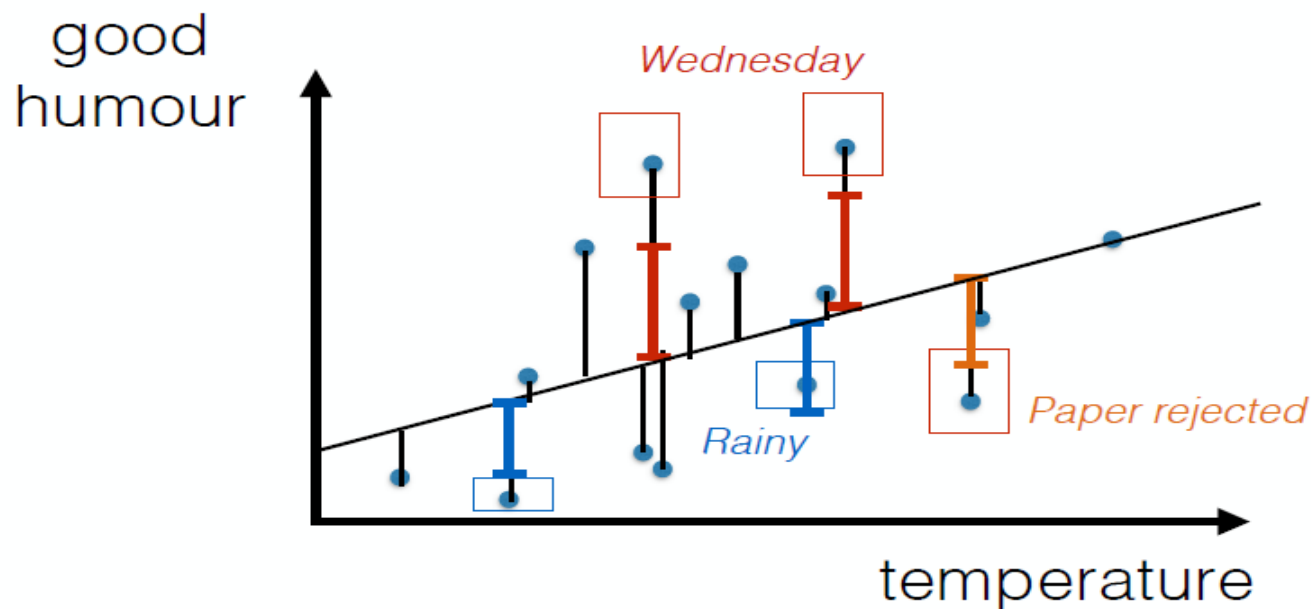
Adding covariates and R^2



Adding covariates almost always increases the R^2 .

Least squares

Adding covariates and R^2



Adding covariates almost always increases the R^2 - so a key question is when to stop.

What to choose?



Least square AIC

$$\text{AIC} = N * \ln\left(\frac{SS_e}{N}\right) + 2K$$

N: Number of observations

SS_e: Sum square of errors

K: Number of parameters

The smaller the AIC the better

Least square AIC

More parameter is not always good

$$AIC = N * \ln\left(\frac{SS_e}{N}\right) + 2K$$


N: Number of observations

SS_e: Sum square of errors

K: Number of parameters

$$(AIC = -2 \ln(L) + 2k)$$

The smaller the AIC the better

An example of model selection: *Bartonella* spp. in Madagascar rats

Epidemics 20 (2017) 56–66



Contents lists available at ScienceDirect

Epidemics

journal homepage: www.elsevier.com/locate/epidemics



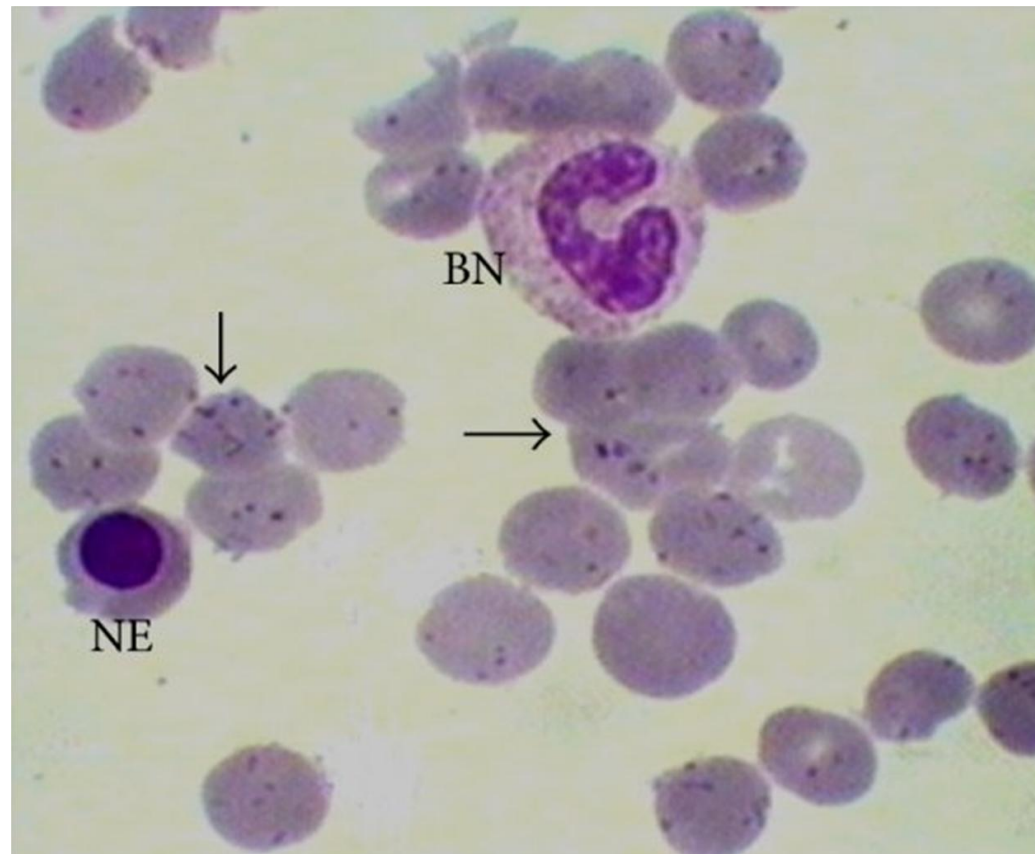
Elucidating transmission dynamics and host-parasite-vector relationships for rodent-borne *Bartonella* spp. in Madagascar



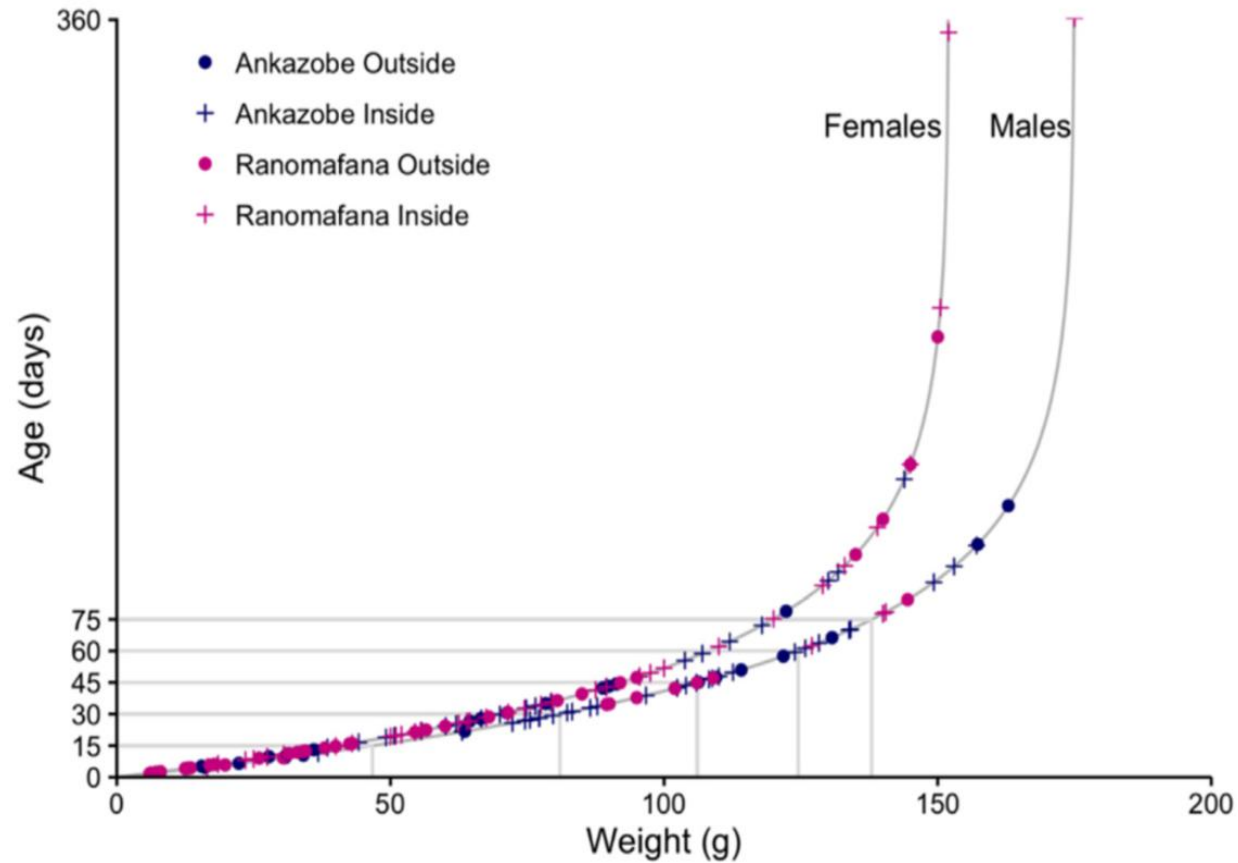
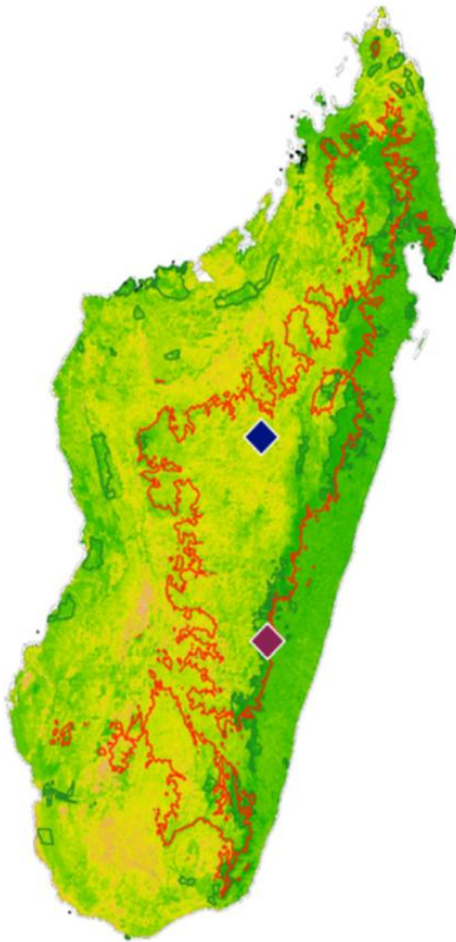
Cara E. Brook^{a,*}, Ying Bai^b, Emily O. Yu^a, Hafaliana C. Ranaivoson^{c,d}, Haewon Shin^e,
Andrew P. Dobson^a, C. Jessica E. Metcalf^{a,1}, Michael Y. Kosoy^{b,1}, Katharina Dittmar^{e,1}

Bartonella spp.

- Persistent erythrocytic bacteria that are sometimes zoonotic
- Vectored by ticks, fleas, sand flies, mosquitoes
- Some species infect humans
 - *Bartonella bacilliformis* = Carrion's disease
 - *Bartonella henselae* = cat scratch fever
 - *Bartonella quintana* = trench fever

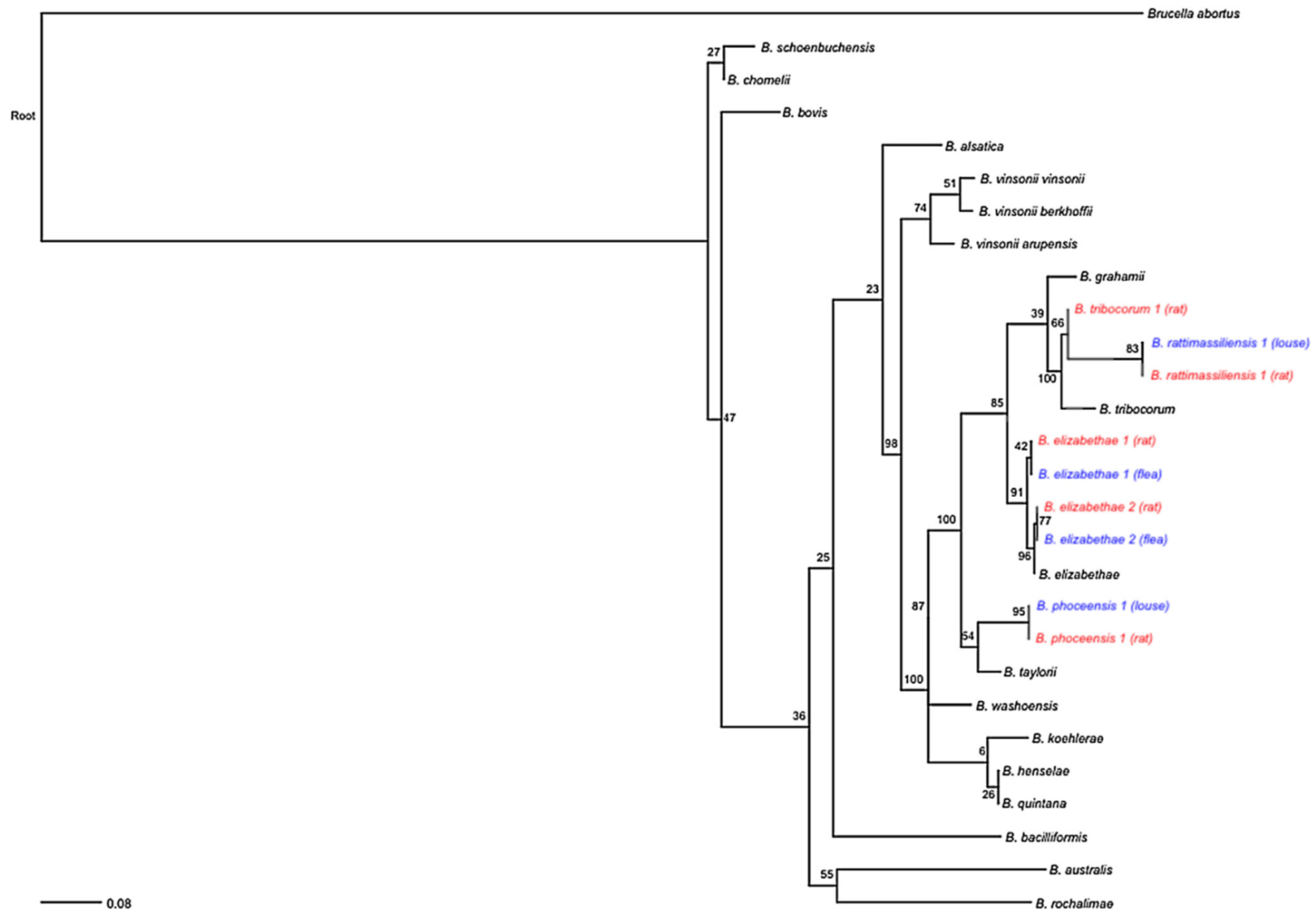


We first collected samples from rats from two sites in Madagascar.



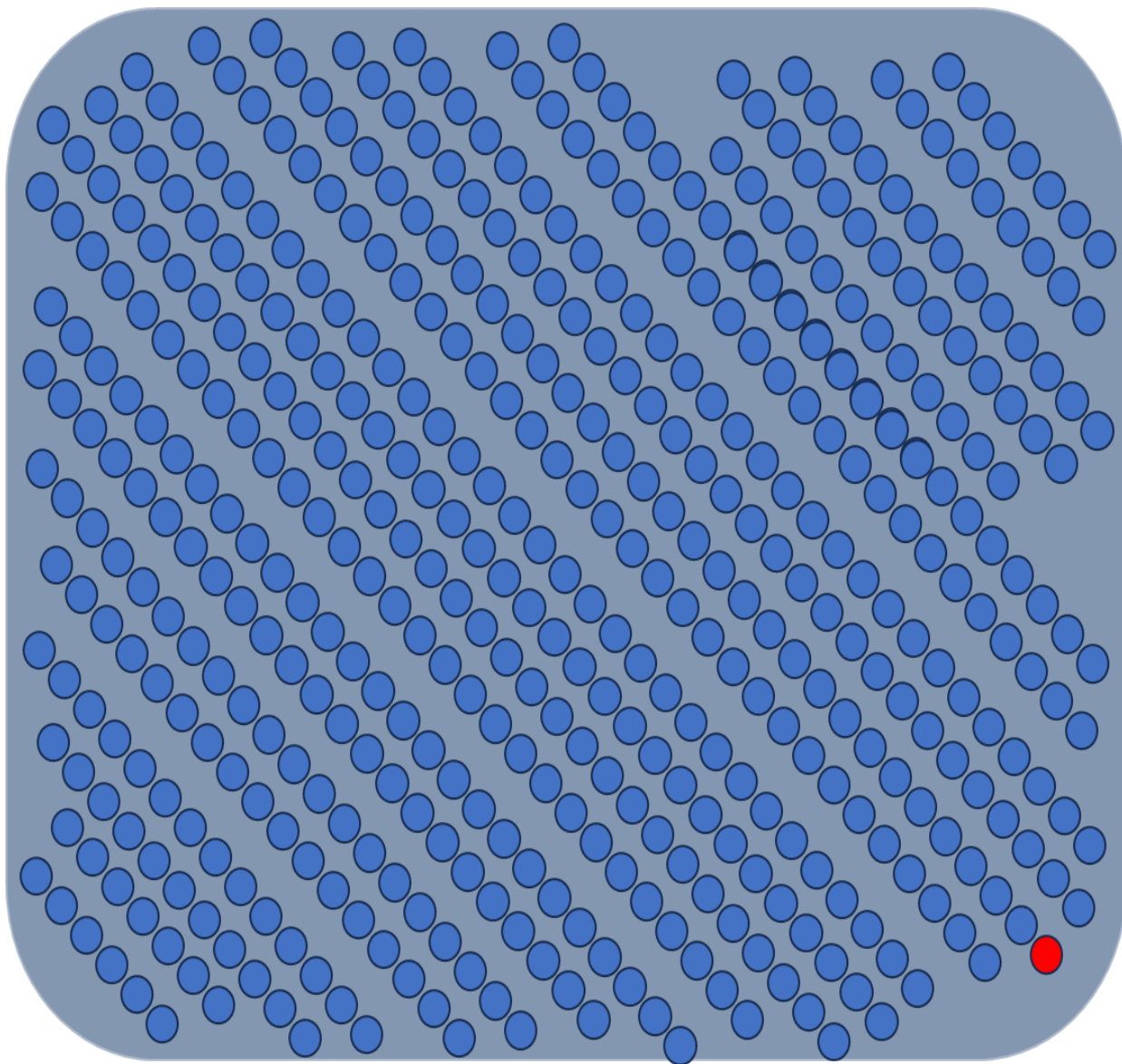
(Ricker 1979)

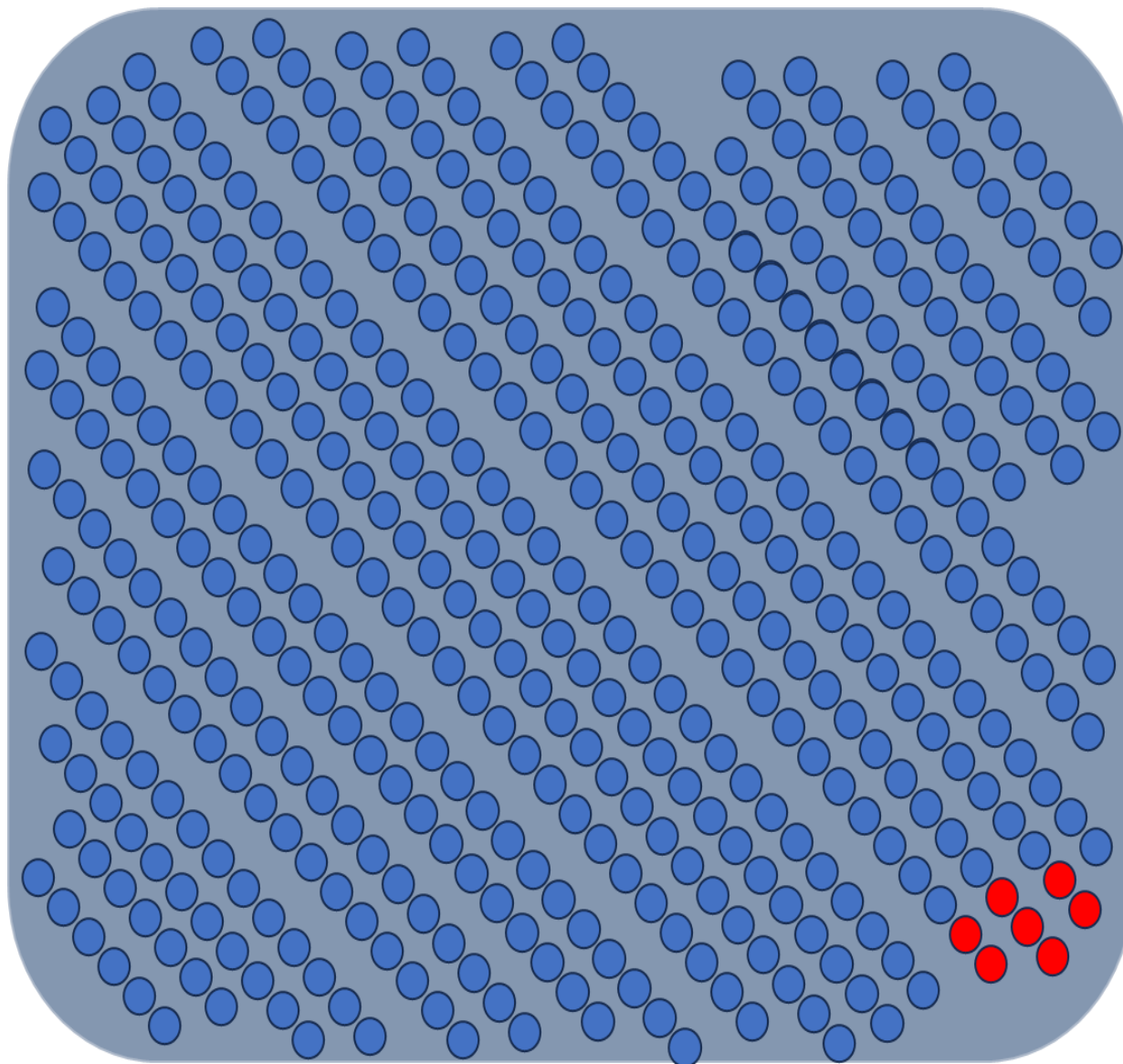
Statistically, we demonstrated an association between genotypes of *Bartonella* spp. found in rats and their ectoparasites.

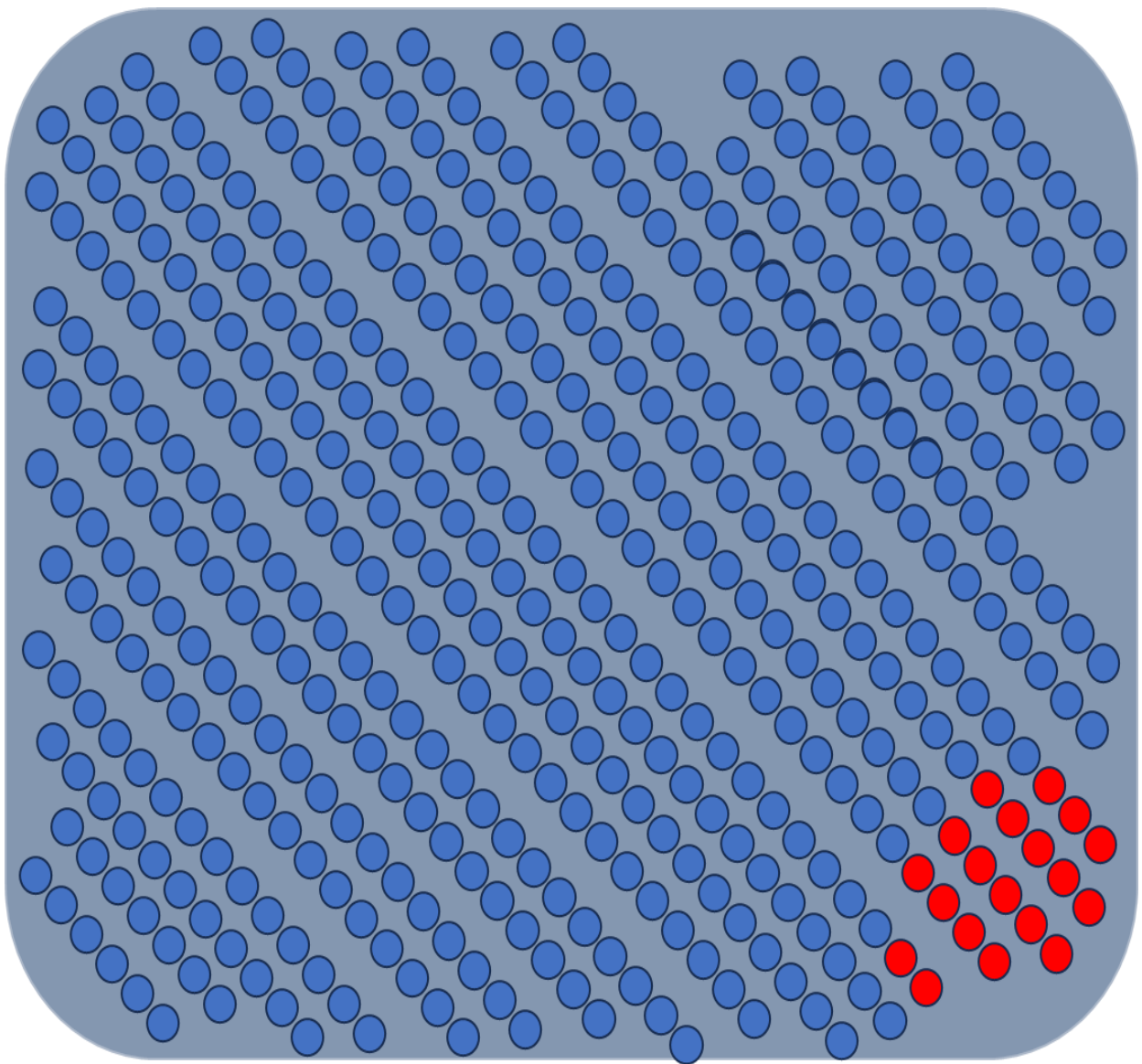


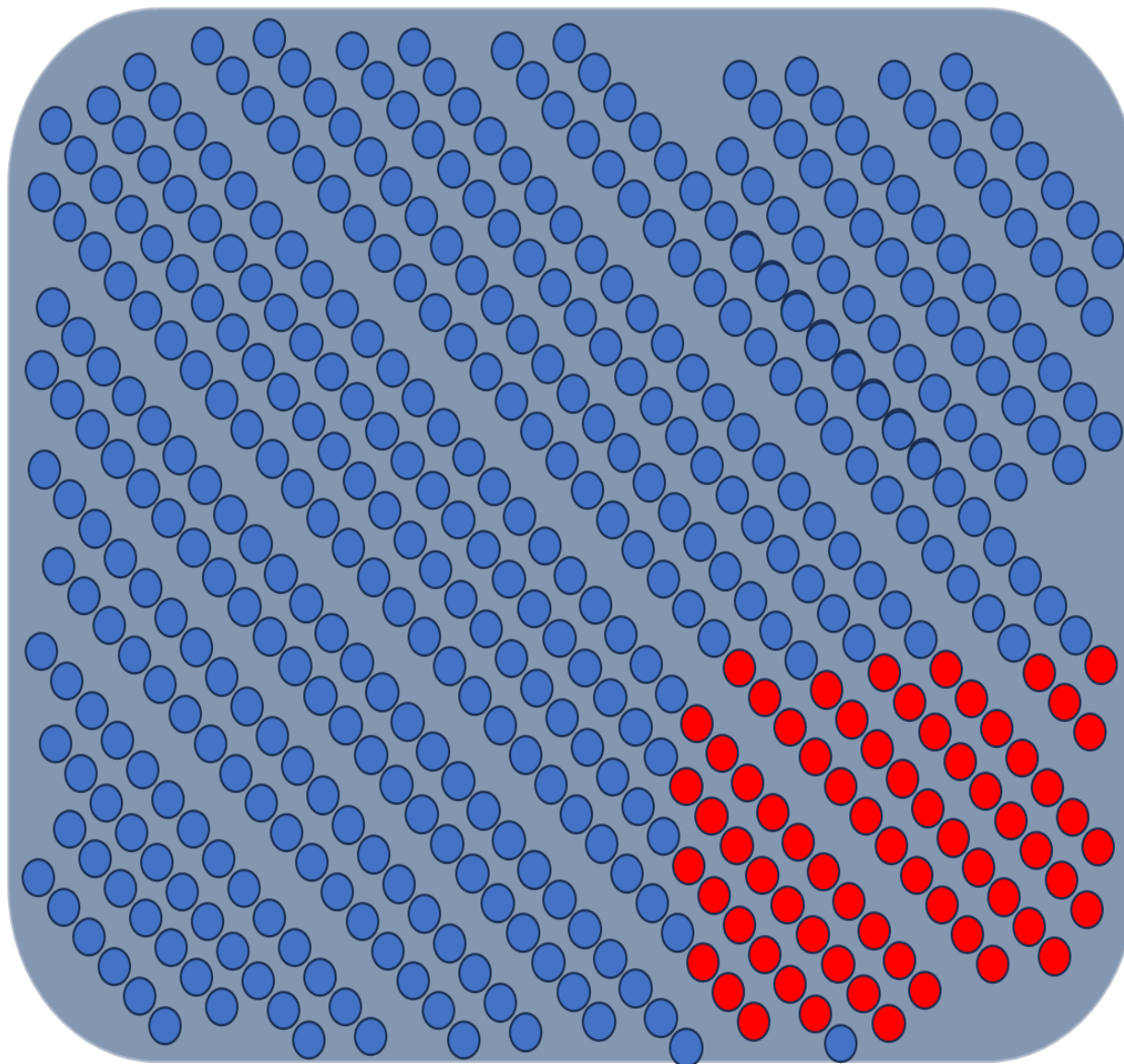
Then, we asked:
*How does the rate of becoming
infected vary with age?*

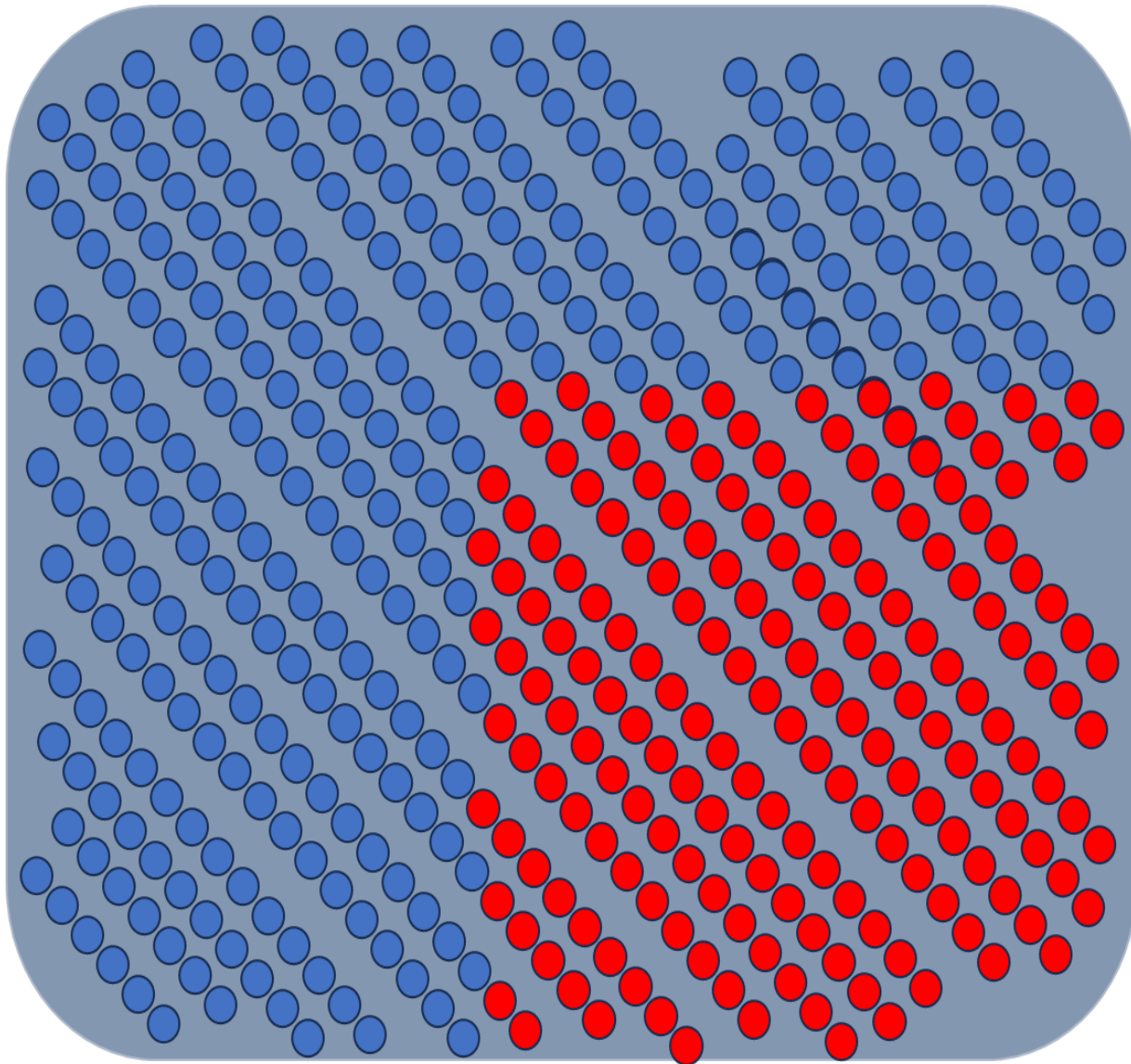
age	sick	age.class	sex
1.685375	0	1	M
2.011821	0	1	F
2.344301	0	1	F
2.611943	0	1	F
4.145697	0	1	F
4.319159	0	1	F
4.319159	0	1	F
4.493246	0	1	F
4.662362	0	1	M
4.784125	0	1	M



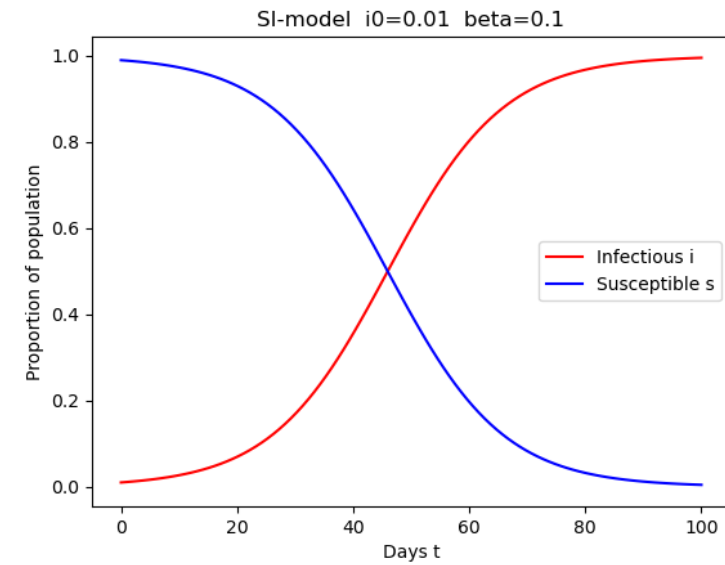
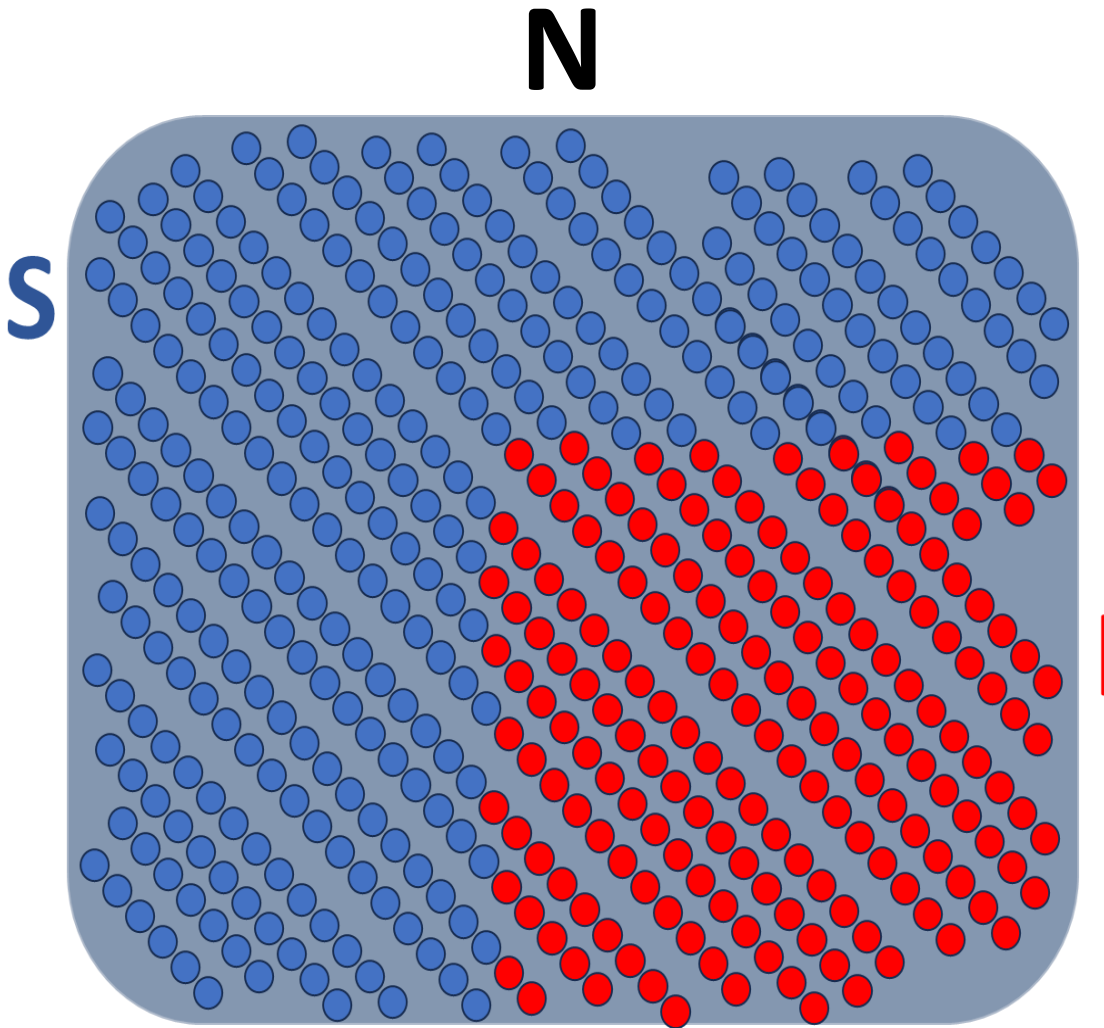




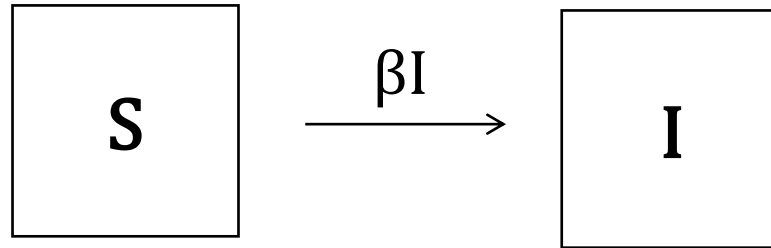




Simple SI model

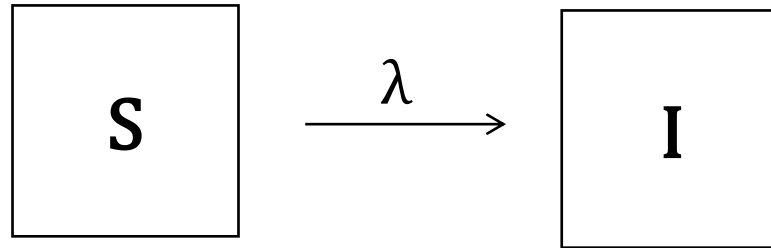


Age-prevalence data allows for powerful inference into the dynamics of pathogen transmission.



for a persistent, non-immunizing infection

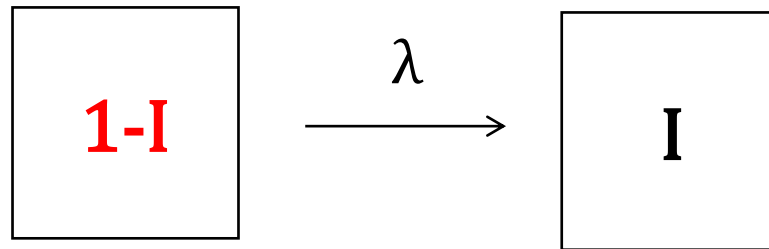
Age-prevalence data allows for powerful inference into the dynamics of pathogen transmission.



where λ , the force of infection, is the per capita rate at which susceptible hosts become infected

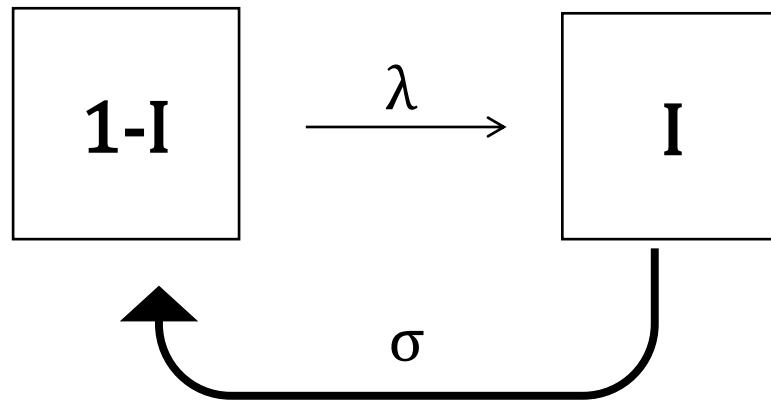
Age-prevalence data allows for powerful inference into the dynamics of pathogen transmission.

with a persistent infection,
we can assume that, if not
infected, you must be
susceptible....



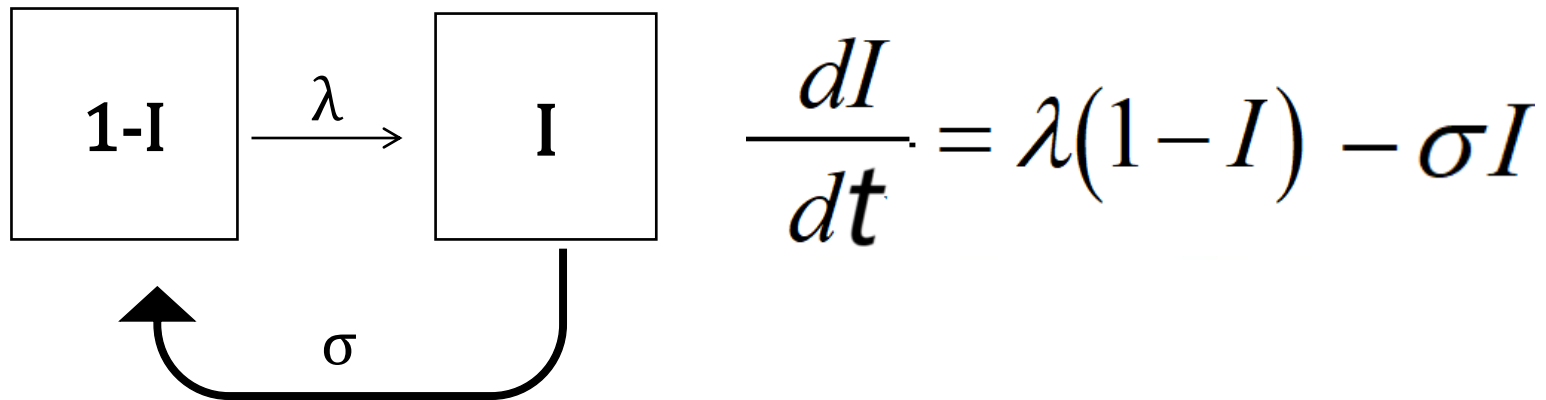
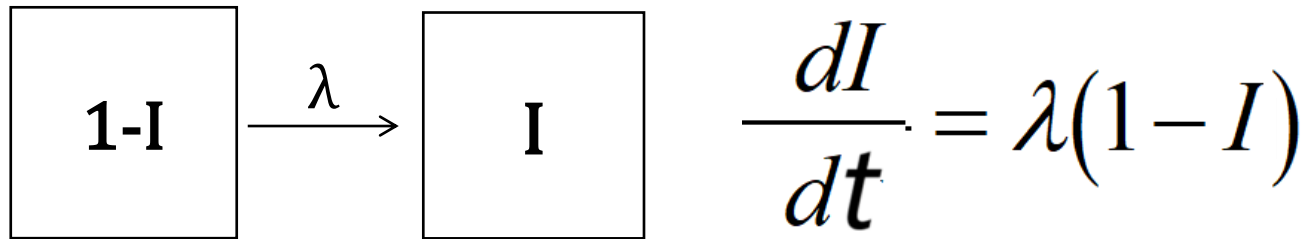
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Age-prevalence data allows for powerful inference into the dynamics of pathogen transmission.

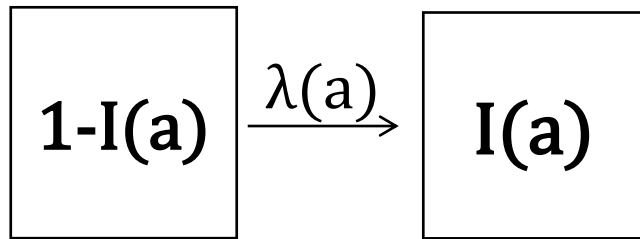


and σ is the rate of recovery from infection

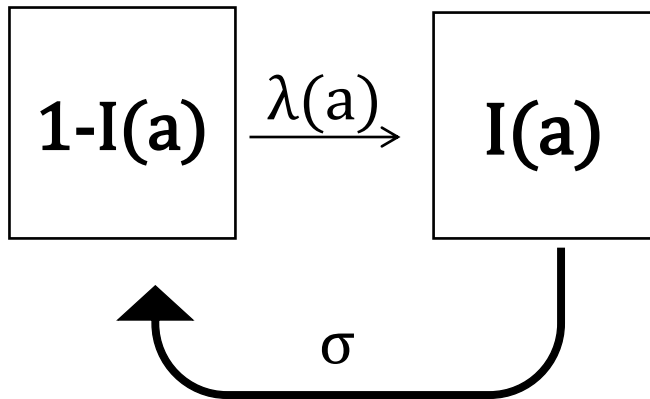
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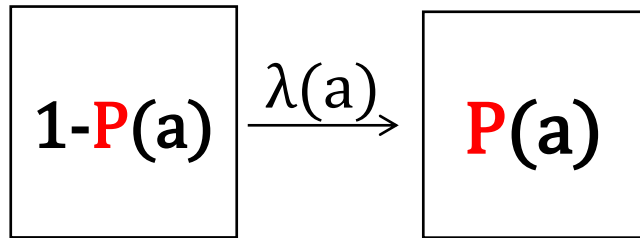


$$\frac{dI(a)}{da} = I(a)(1 - I(a))$$

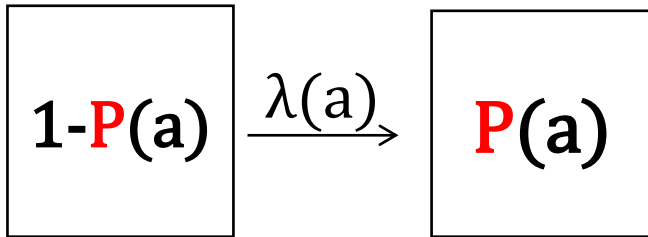


$$\frac{dI(a)}{da} = I(a)(1 - I(a)) - \sigma I(a)$$

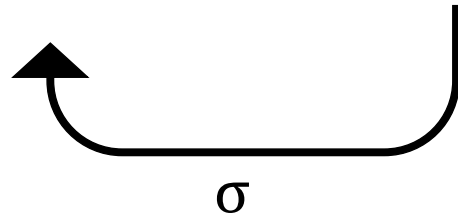
Age-prevalence data allows for powerful inference into the dynamics of pathogen transmission.



$$\frac{d P(a)}{da} = \lambda(a) (1 - P(a))$$



$$\frac{d P(a)}{da} = \lambda(a) (1 - P(a)) - \sigma(a) P(a)$$



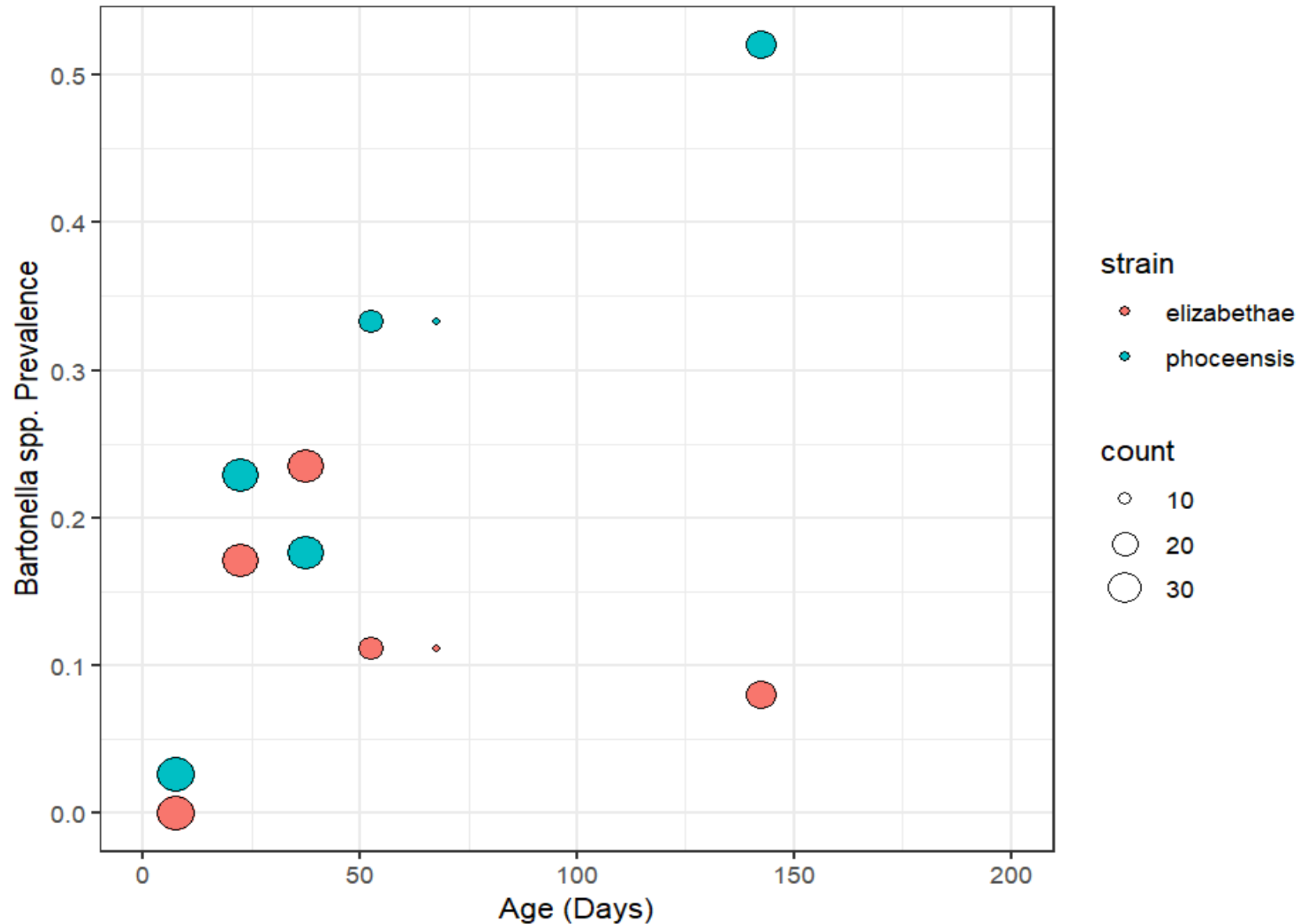
Compare using **AIC = 2K - 2ln(L)**

similar techniques can also be applied to age-seroprevalence data for immunizing infections

Let's see which model works best
for your data!

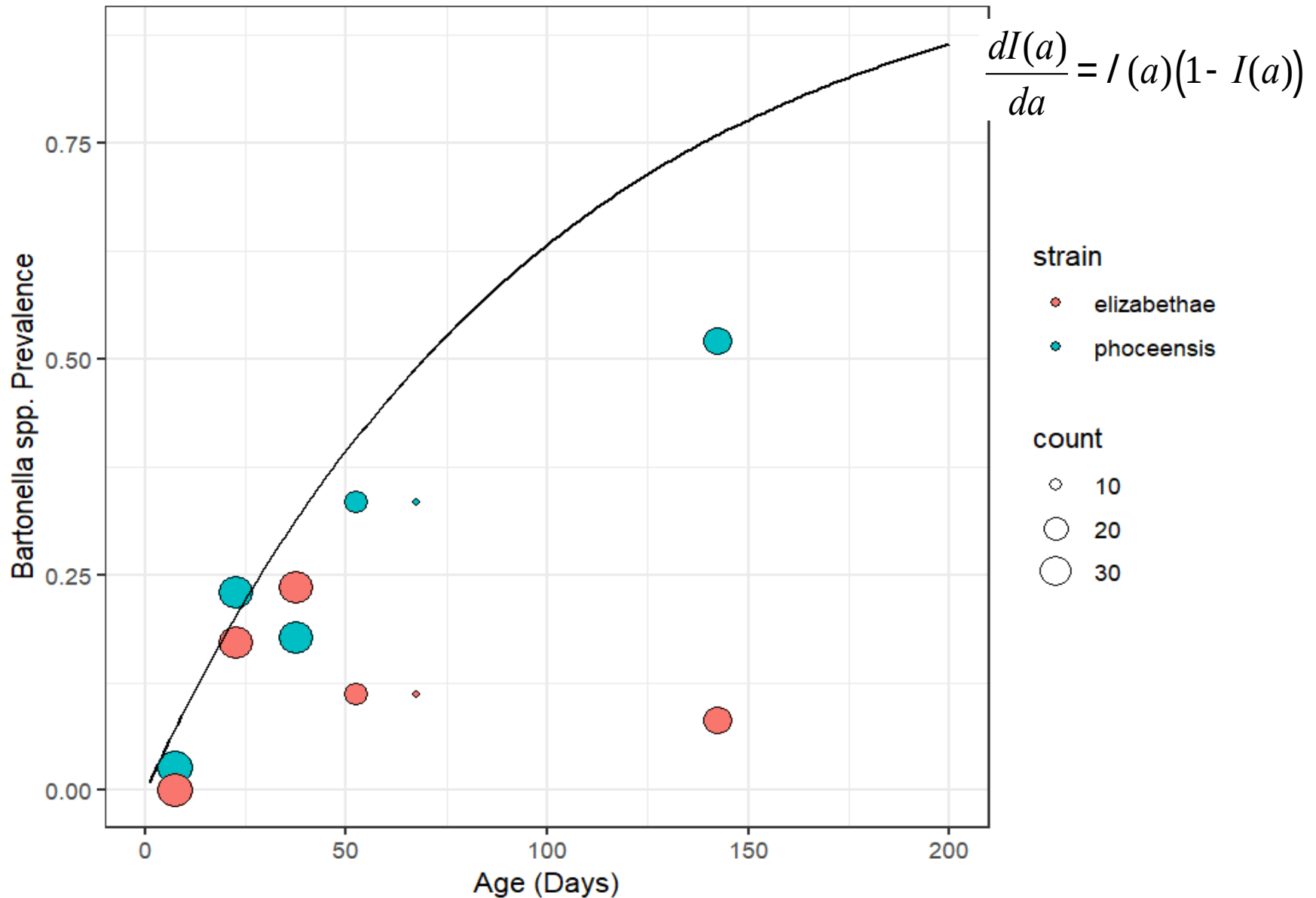
Look at the data !

Jereo aloha hoe manao ahoana



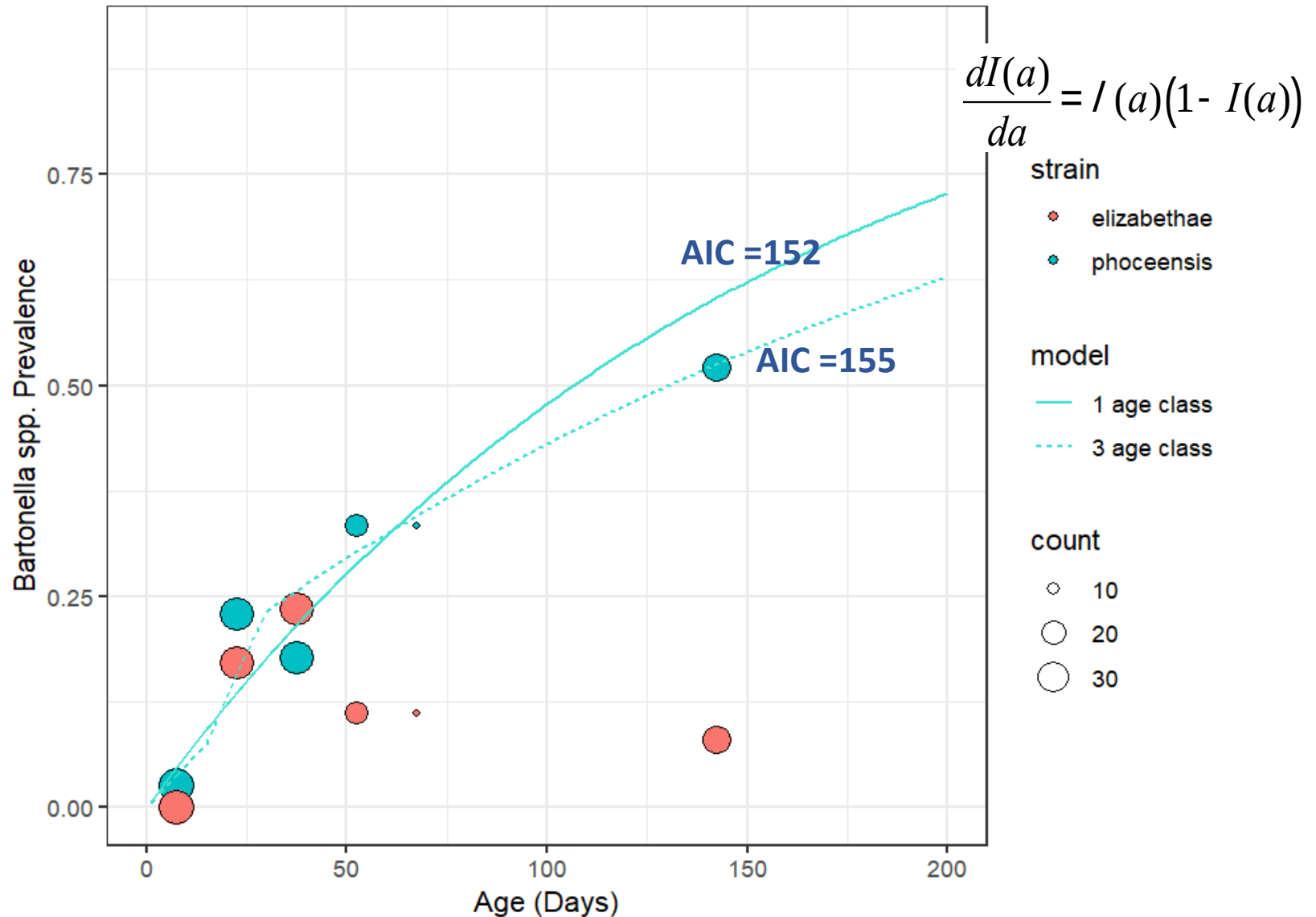
Try the model!

Andramo kely

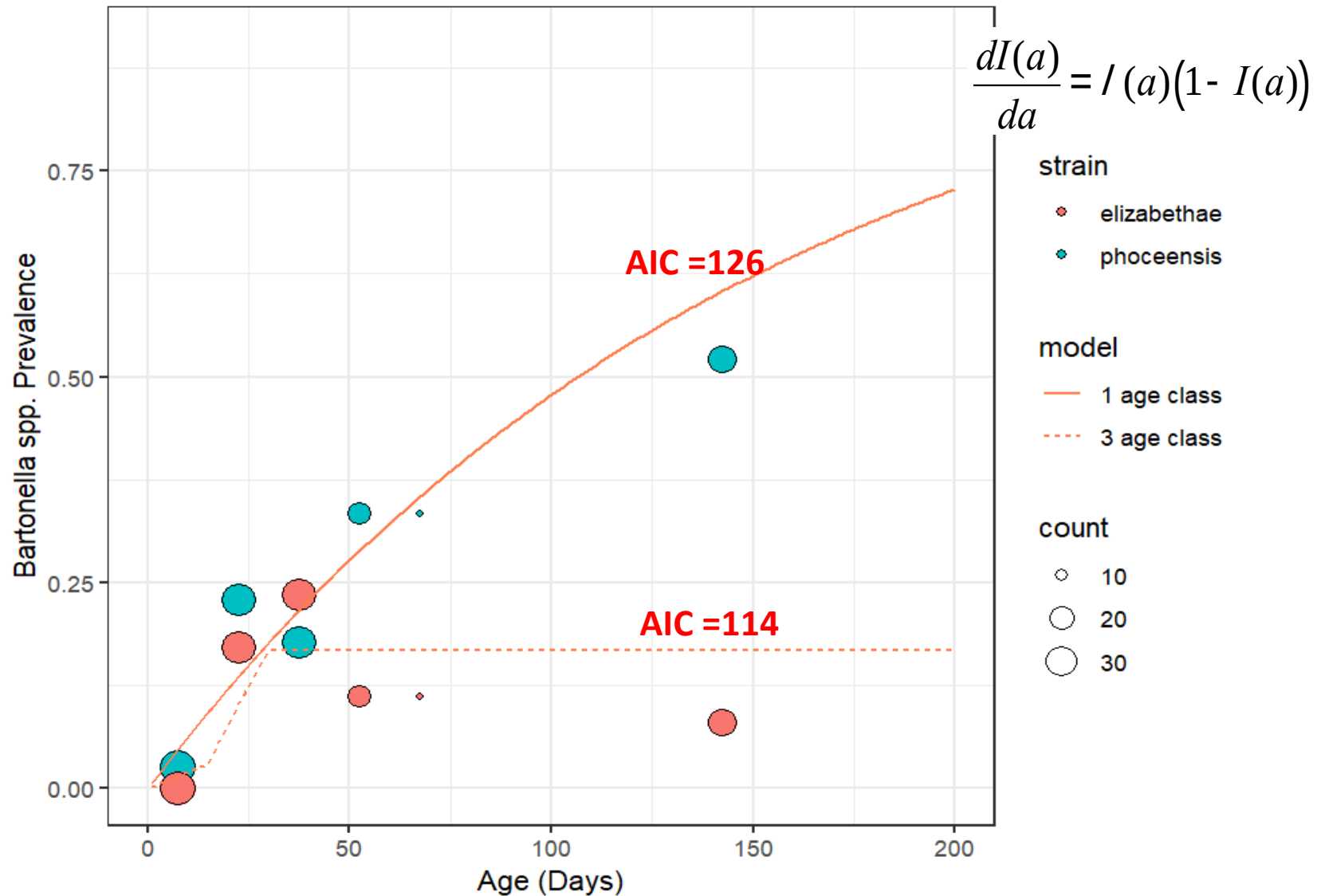


Keep trying!

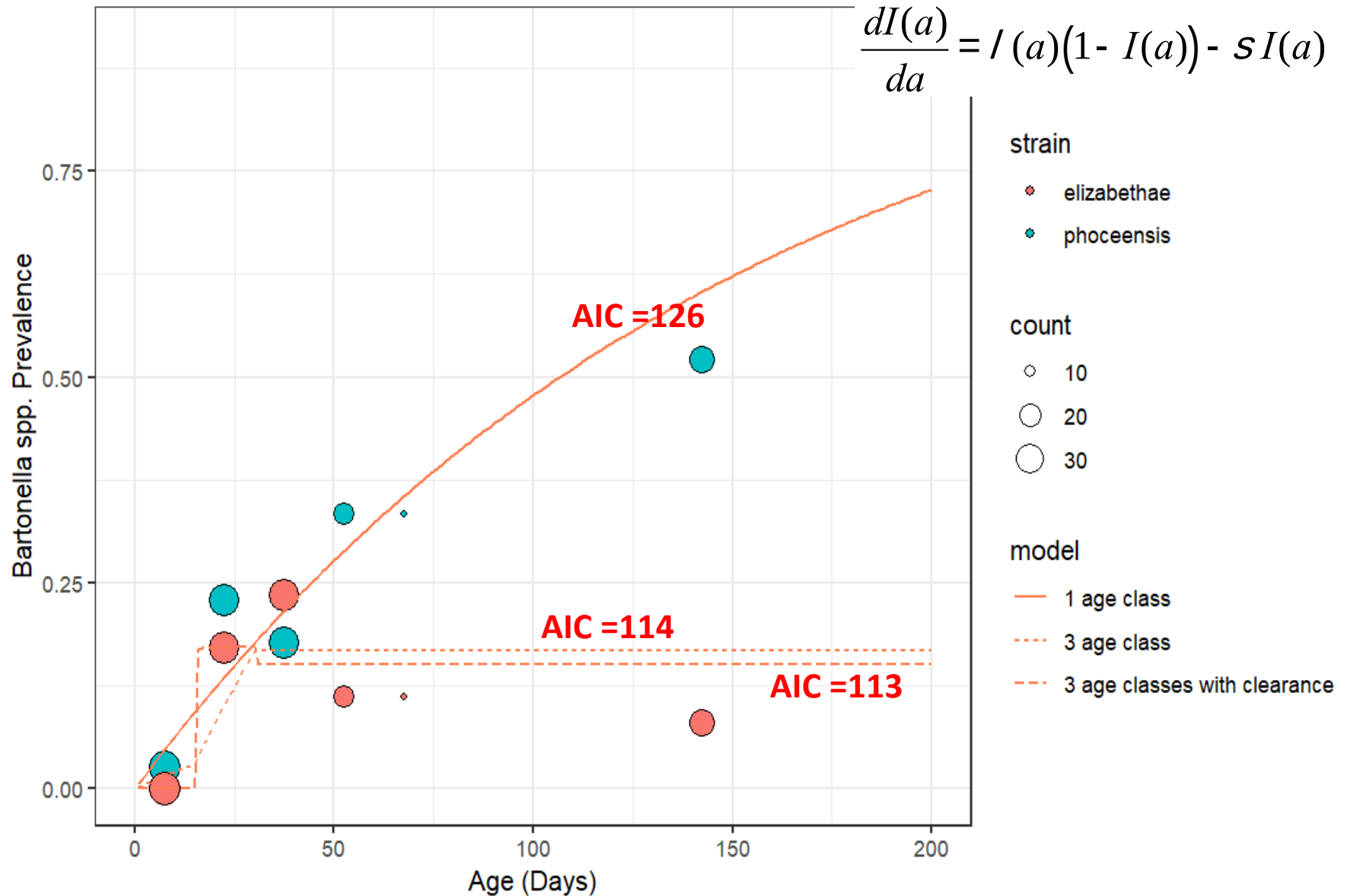
Alô fô !



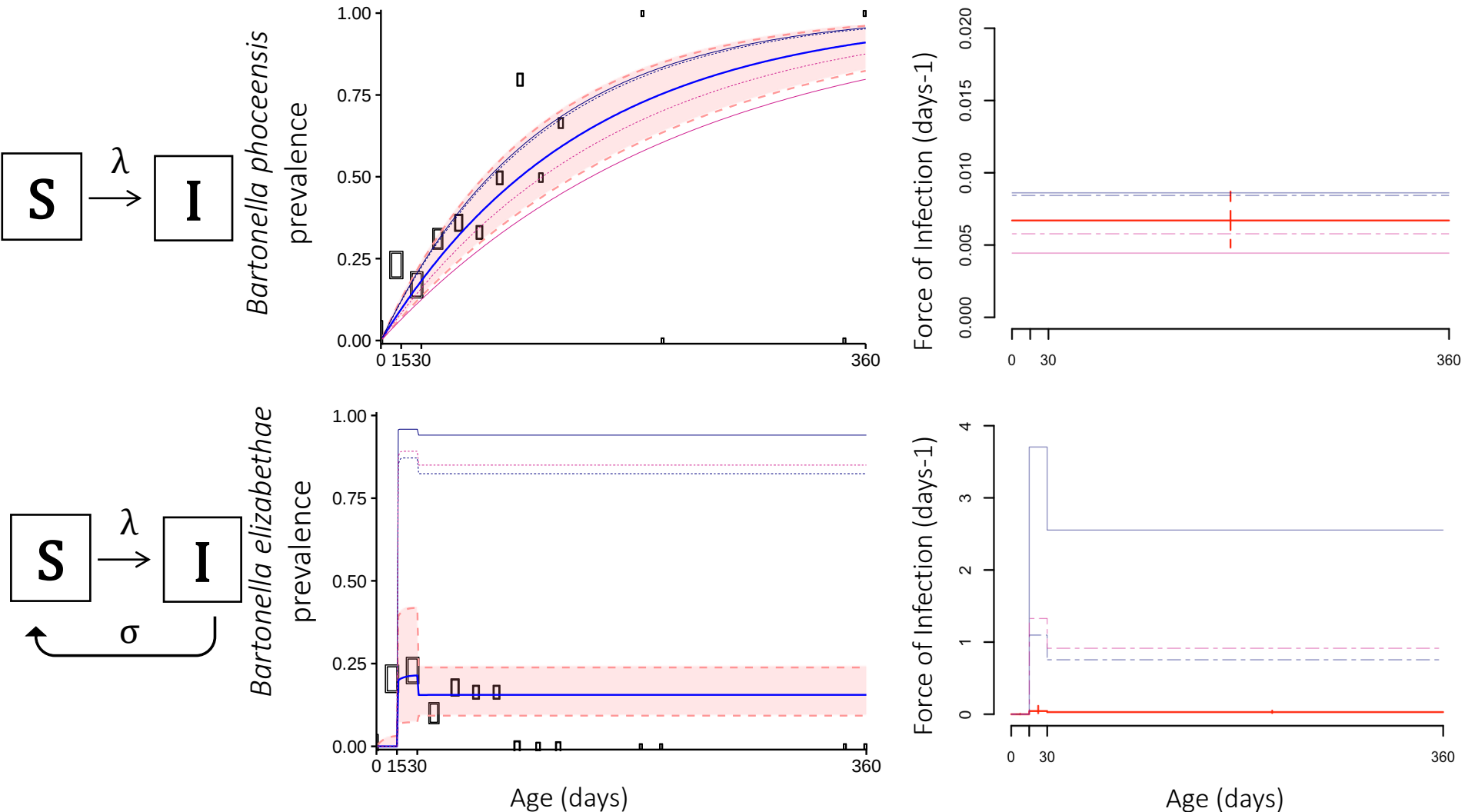
Keep trying!



Keep trying!



We found that an **SI model** offered the best fit to *B. phoceensis* data while the **SIS model** offered the best fit to the *B. elizabethae* data.



The age-structured FOI identifies age cohorts most influential in an epidemic. Juveniles showed the highest FOI.

