

# A discrete stochastic model of the transmission cycle of the tick borne encephalitis virus

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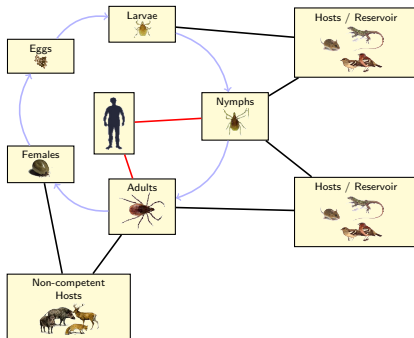
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3 September 2012



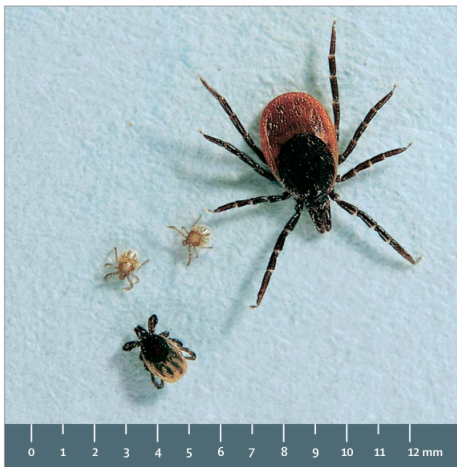
# Tick Borne Encephalitis (TBE)

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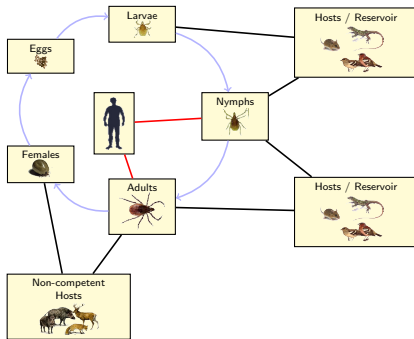


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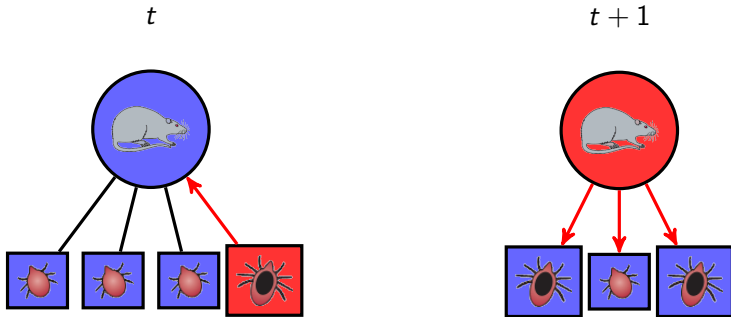


- no consequences for animals
- severe consequences for humans

# Transmissions

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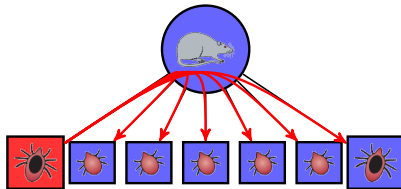
Systemic Transmission:



# Transmissions

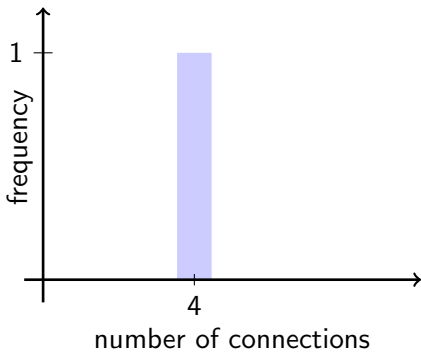
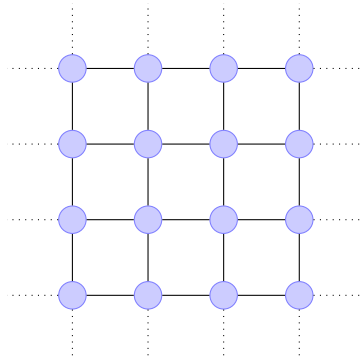
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Co-Feeding Transmission:



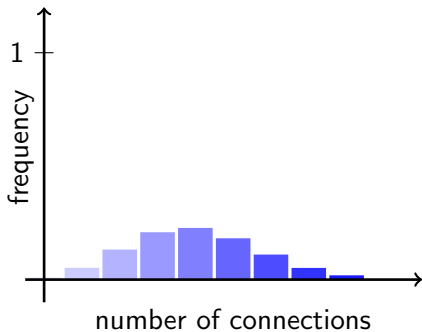
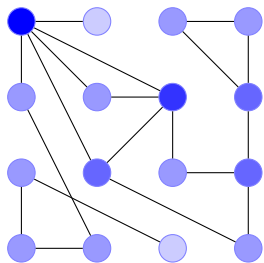
## Networks - Lattice

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## Networks - Random Graph

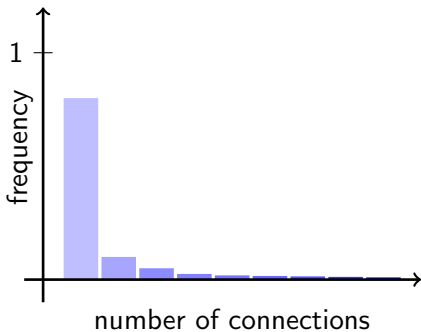
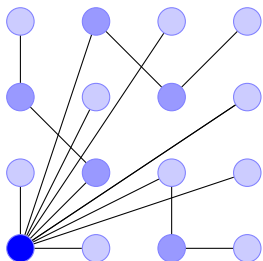
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## Networks - Scale Free Network

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# Compartmental Models on Networks

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Susceptible-Infectious-Susceptible (SIS) model:

$$\frac{d i_k}{d t} = -\mu i_k + \beta k (1 - i_k) \frac{1}{\langle k \rangle} \sum_h h p_h i_h$$

# Compartmental Models on Networks

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the epidemic threshold:

$$\frac{\beta}{\mu} > \frac{\langle k \rangle}{\langle k^2 \rangle}$$

# The model

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prevalence  $i = \frac{\text{number infected}}{\text{total population}}$

## The model

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$i_{k,y}$  prevalence for **larvae** that encounter  $k$  nymphs at the end of year  $y$

## The model

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$j_{k,y}$  prevalence for nymphs that encounter  $k$  larvae in the year  $y$

# The model

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a strong assumption(!):

$$i_{k,y} = j_{k,y+1}$$

$$i_{k,y} = 1 - (1 - A_y)^k$$

where  $A_y$  is the probability that an edge is transmitting the disease



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$$A_y = \frac{\beta}{\langle k \rangle_y^{N \rightarrow L}} \sum_h h \cdot p_{h,y}^{N \rightarrow L} \cdot i_{h,y-1}$$

therefore

$$A_y = \beta - \frac{\beta}{\langle k \rangle_y^{N \rightarrow L}} \sum_h h \cdot p_{h,y}^{N \rightarrow L} \cdot (1 - A_{y-1}^h)$$

## The Epidemic Threshold

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If we can assume that year  $y$  and year  $y - 1$  are ecologically comparable then we check when  $A_y = A_{y-1} = x$ .

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This corresponds to

$$x = F(x) = \beta - \frac{\beta}{\langle k \rangle_y^{N \rightarrow L}} \sum h \cdot p_{h,y}^{N \rightarrow L} \cdot (1 - x)^h$$

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$$F'(x) = \frac{\beta}{\langle k \rangle_y^{N \rightarrow L}} \sum h^2 p_{hy}^{N \rightarrow L} (1 - x)^{h-1}$$

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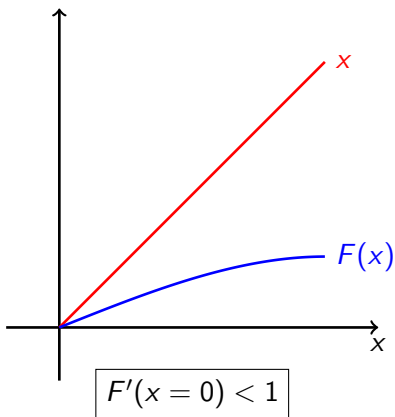
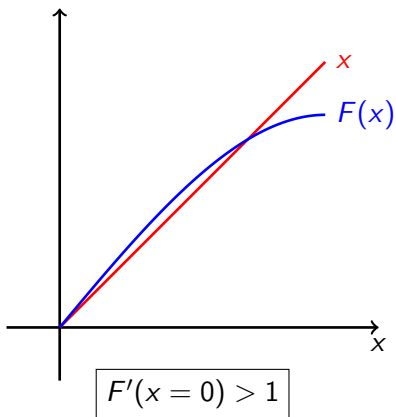
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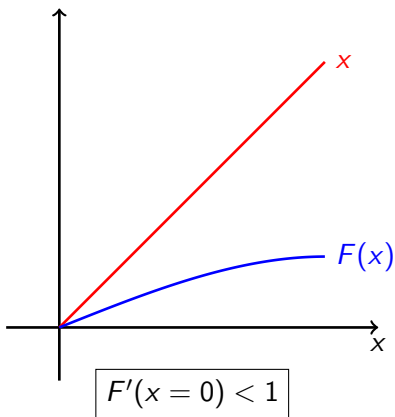
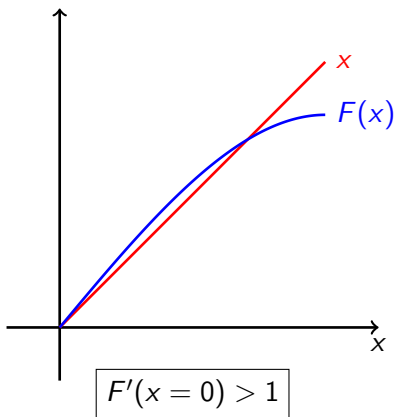
$$F'(x) = \frac{\beta}{\langle k \rangle_y^{N \rightarrow L}} \sum h^2 p_{hy}^{N \rightarrow L} (1 - x)^{h-1}$$

and

$$F''(x) = -\frac{\beta}{\langle k \rangle_y^{N \rightarrow L}} \sum h^2 (h - 1) p_{hy}^{N \rightarrow L} (1 - x)^{h-2}$$







$$\beta > \frac{\langle k \rangle_y^{N \rightarrow L}}{\langle k^2 \rangle_y^{N \rightarrow L}}$$

Some recurrent ecological cycles are reported from real world data. Hence, sometimes, instead of the assumption  $A_y = A_{y-1}$ , it could be more suitable  $A_y = A_{y-n}$ . In that case the epidemiological threshold would be:

$$\beta^n > \frac{\langle k \rangle_y^{N \rightarrow L}}{\langle k^2 \rangle_y^{N \rightarrow L}} \cdot \dots \cdot \frac{\langle k \rangle_{y-n+1}^{N \rightarrow L}}{\langle k^2 \rangle_{y-n+1}^{N \rightarrow L}}$$