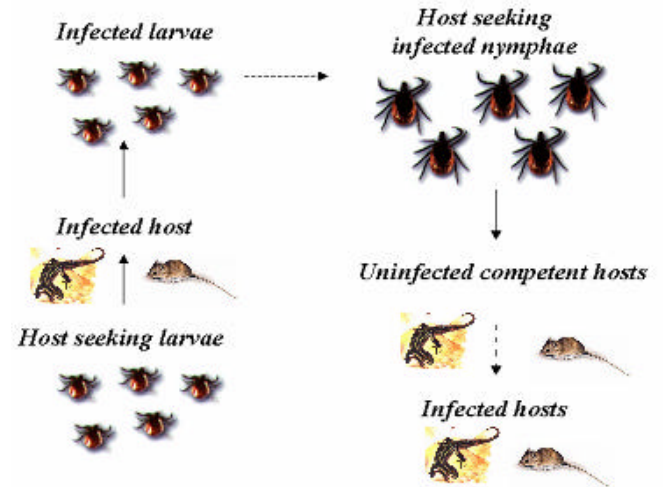


Dynamics of Borrelia burgdorferi sensu lato (s.l.) transmission between ticks and two different hosts: basic reproductive number estimation, in an Italian Natural Reserve

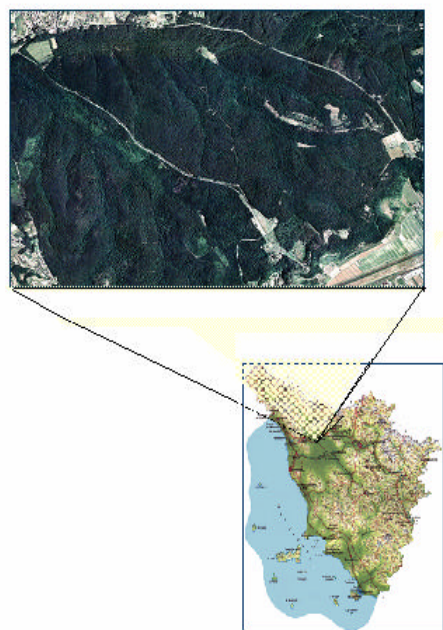
- emerging tick-borne zoonosis
- genospecies *B. burgdorferi* s.l. complex, host specie associated
- tick vector *Ixodes ricinus*
- different reservoirs hosts species

- B. lusitaniae* and *B. afzelii* in host seeking and infesting *I. ricinus* ticks,
- abundant *I. ricinus* population
- lizards (*Podarcis* spp.) and mice (*Apodemus* spp.), *B. lusitaniae* and *B. afzelii* reservoirs

Lyme Borreliosis transmission cycle



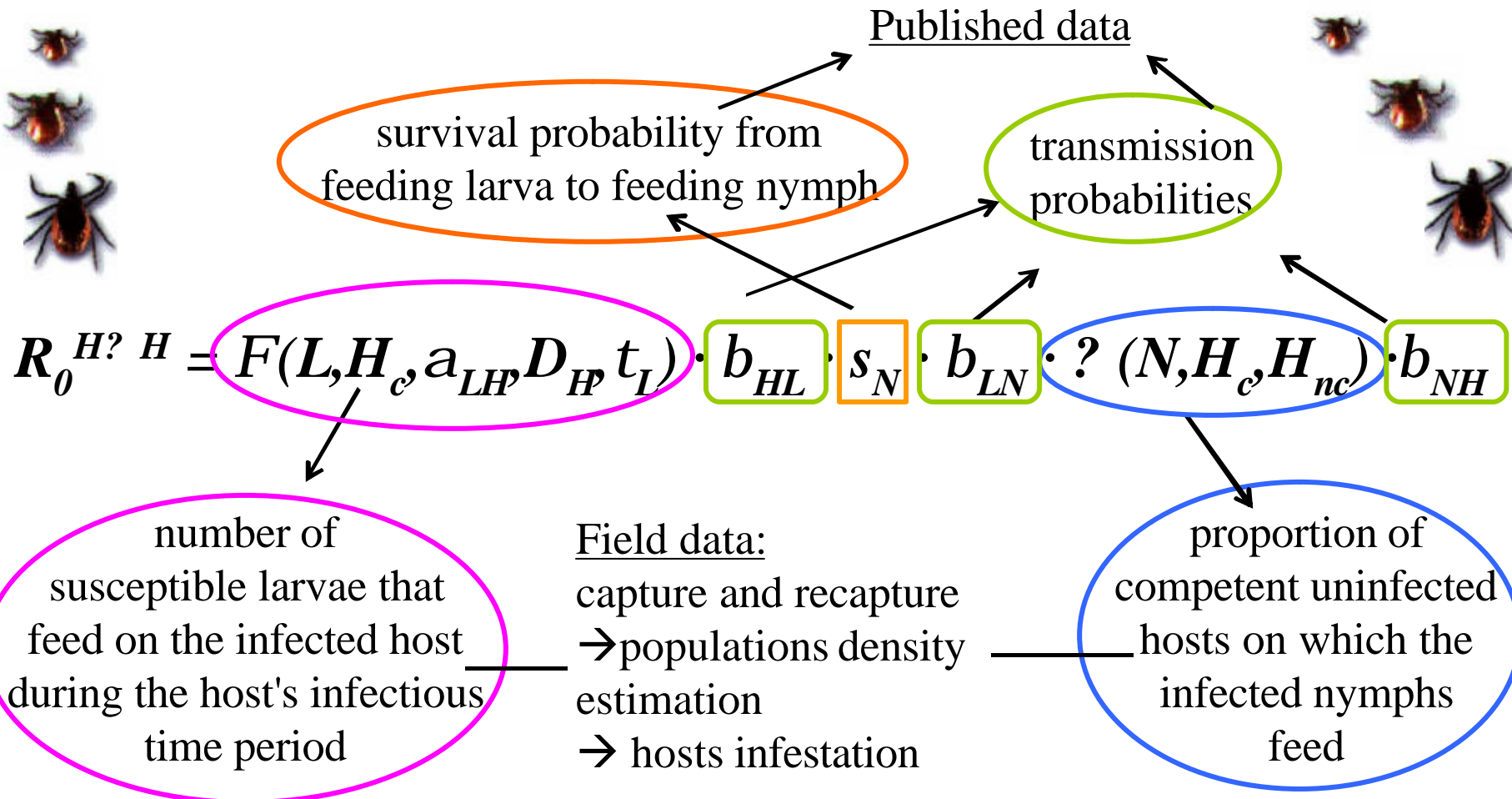
Qualitatively R_0 estimation *B. lusitaniae* and *B. afzelii* considering interactions among vectors, hosts and pathogens.





Basic Reproductive Number (R_0):

The expected number of infected hosts when a single infected host has been inserted in a susceptible population of hosts and vectors

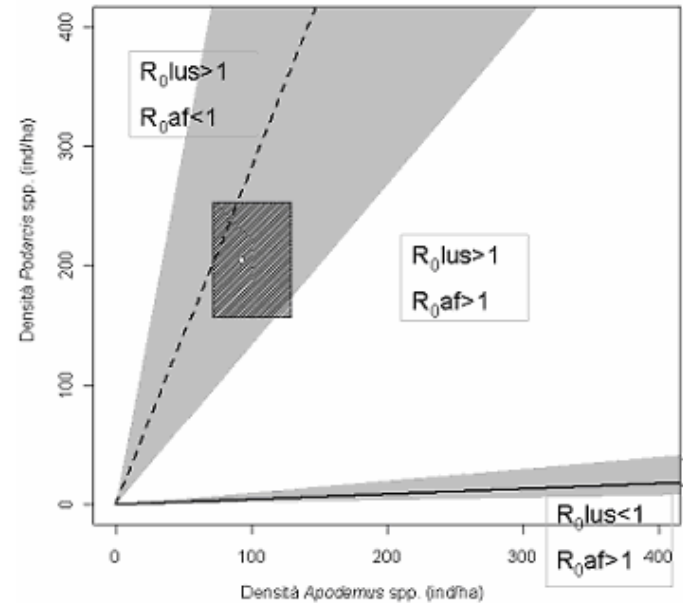




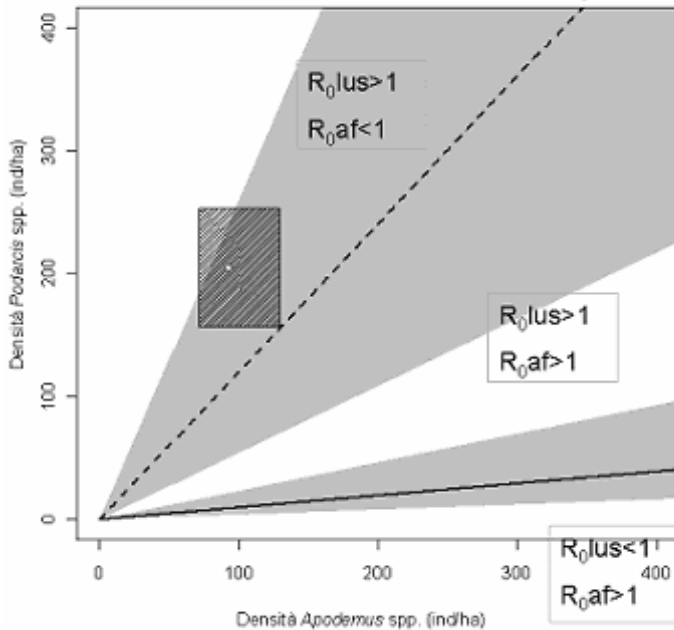
*The model suggests a R_0 greater than 1 for *B. lusitaniae* and around the epidemic threshold for *B. afzelii*:*

right combination of the two host populations sizes
 → coexistence of both spirochete genospecies

Competent hosts system



Competent and non competent hosts system



Presence of other hosts species, non competent for *B. burgdorferi* s.l.:
 → absence of influence on *B. lusitaniae* R_0 / effect on the persistence of *B. afzelii*.



Nymphs infestation, population density and diversity, and *B. burgdorferi* s.l. host species association