

A model to quantify individual tolerance and resistance to infectious diseases



DEFINITIONS

MECHANISMS

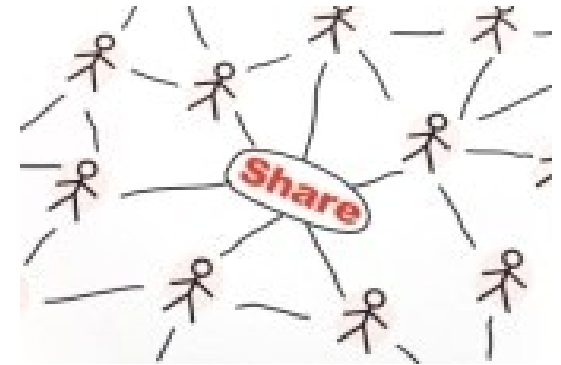
MODELS


COSTS

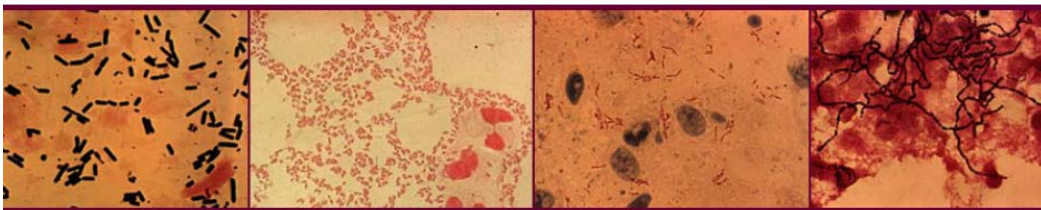
APPLICATION TO BOVINE MASTITIS

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Team work



 **VETERINARY MEDICINE**
Department of infectious and parasitic diseases
LABORATORY OF BACTERIOLOGY



Definitions



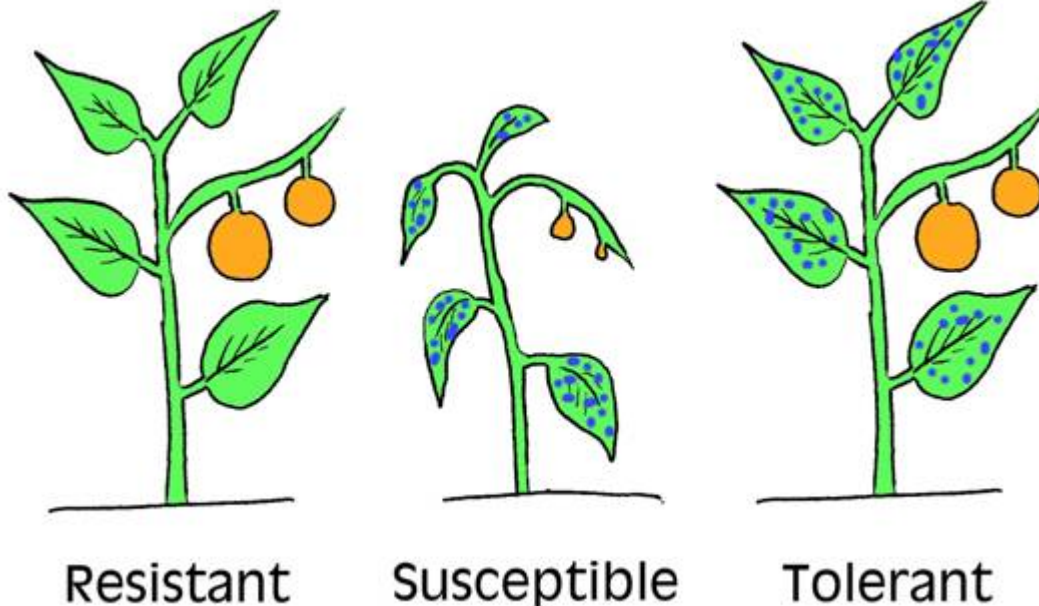
RESISTANCE
TOLERANCE

Definitions



Resistance = ability to reduce pathogen burden

Tolerance = ability to tolerate a certain concentration of pathogen



Definitions

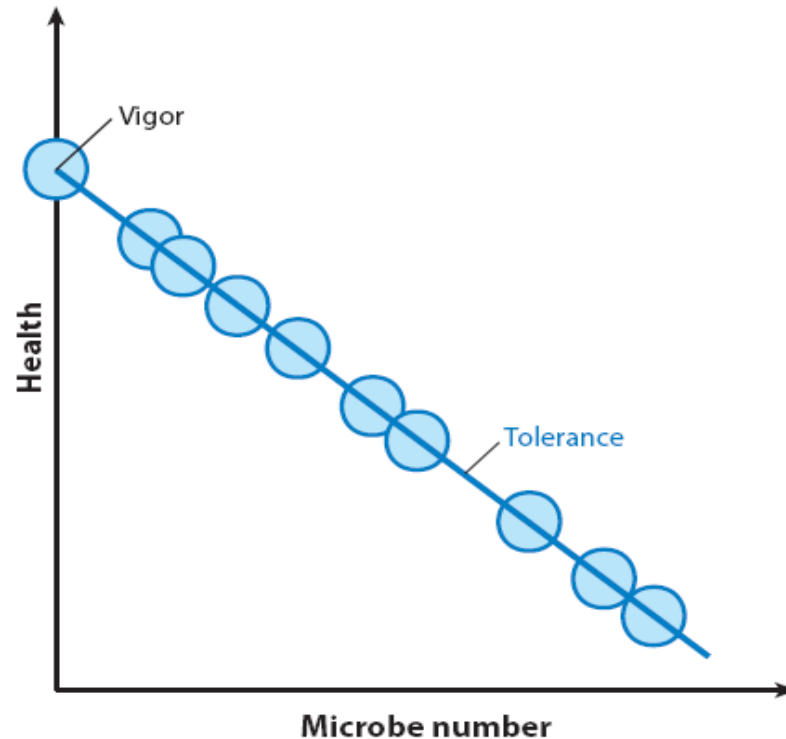


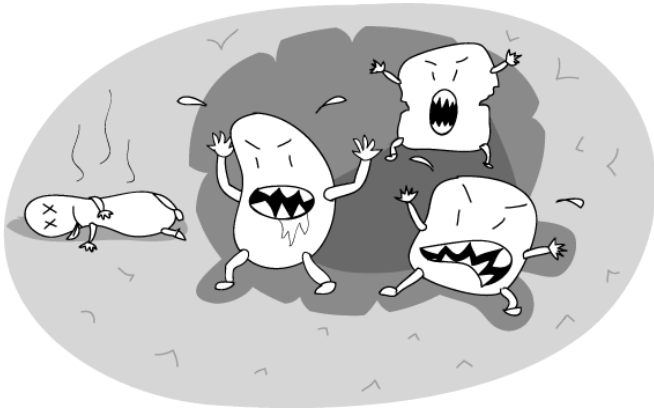
Figure 1

Graphical definitions of resistance and tolerance. Vigor is defined as health at zero parasite load; this may not be the same point that is projected as an intercept by regression (3). Resistance is defined by the inverse of the mean parasite load in the system; more resistant hosts have fewer parasites. Tolerance is defined as the slope of the health-by-microbe curve.

Resistances



Resistance to infection (avoidance)
= reduction of pathogen transmission
at contact



Resistance to disease
= reduction of pathogen growth after
infection

Tolerances



Direct tolerance

Indirect tolerance

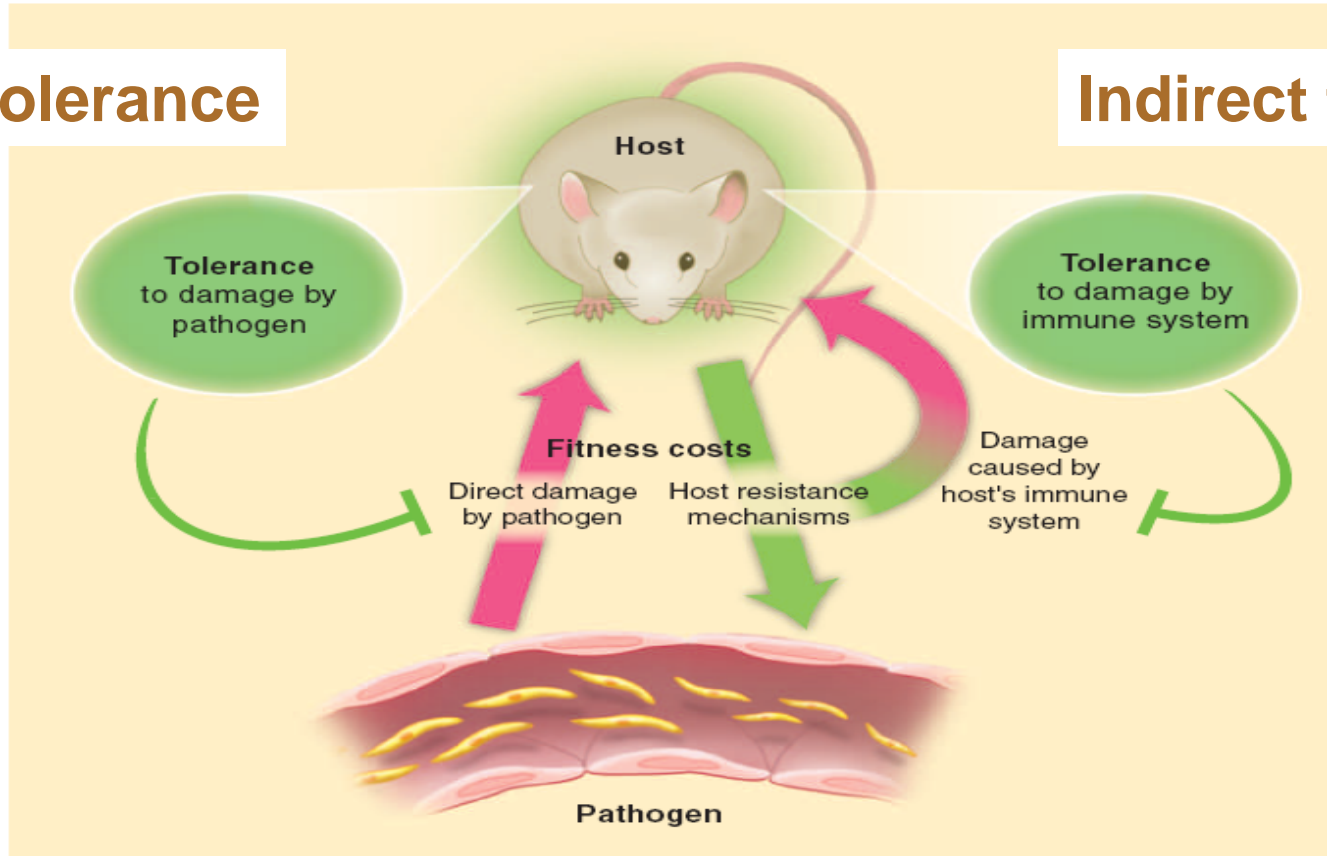


Fig. 1. Two types of fitness costs associated with infections. Pathogens can directly damage the host tissues. The immune system of the host reduces the pathogen burden through the resistance mechanism. The immune response can also damage the host tissues. The host can reduce fitness costs through tolerance mechanisms that reduce both the direct tissue damage by pathogens, and immunopathology.

Mechanisms



RESISTANCE TO INFECTION
RESISTANCE TO DISEASE
TOLERANCE TO INFECTION
TOLLERANCE TO DISEASE



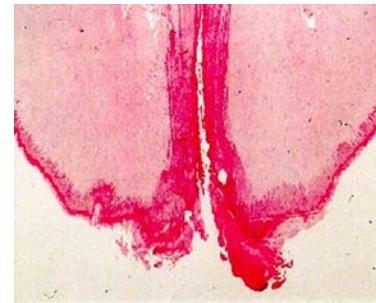
Resistance to infection

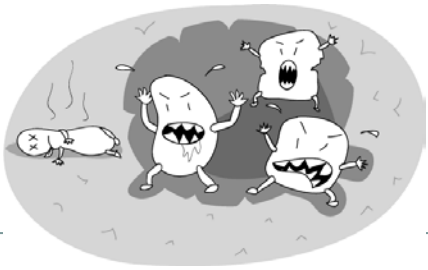


Behavioral immunity



Innate constitutive protective barriers

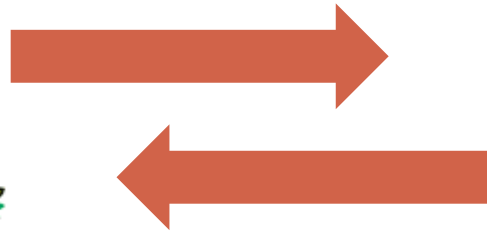
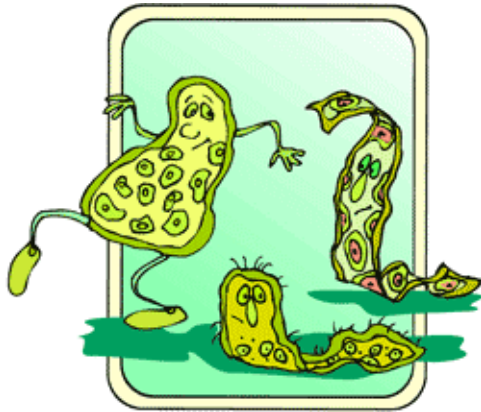




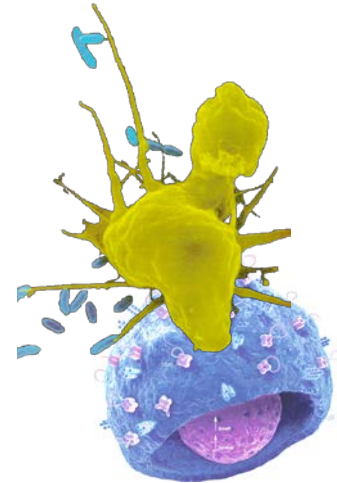
Resistance to disease



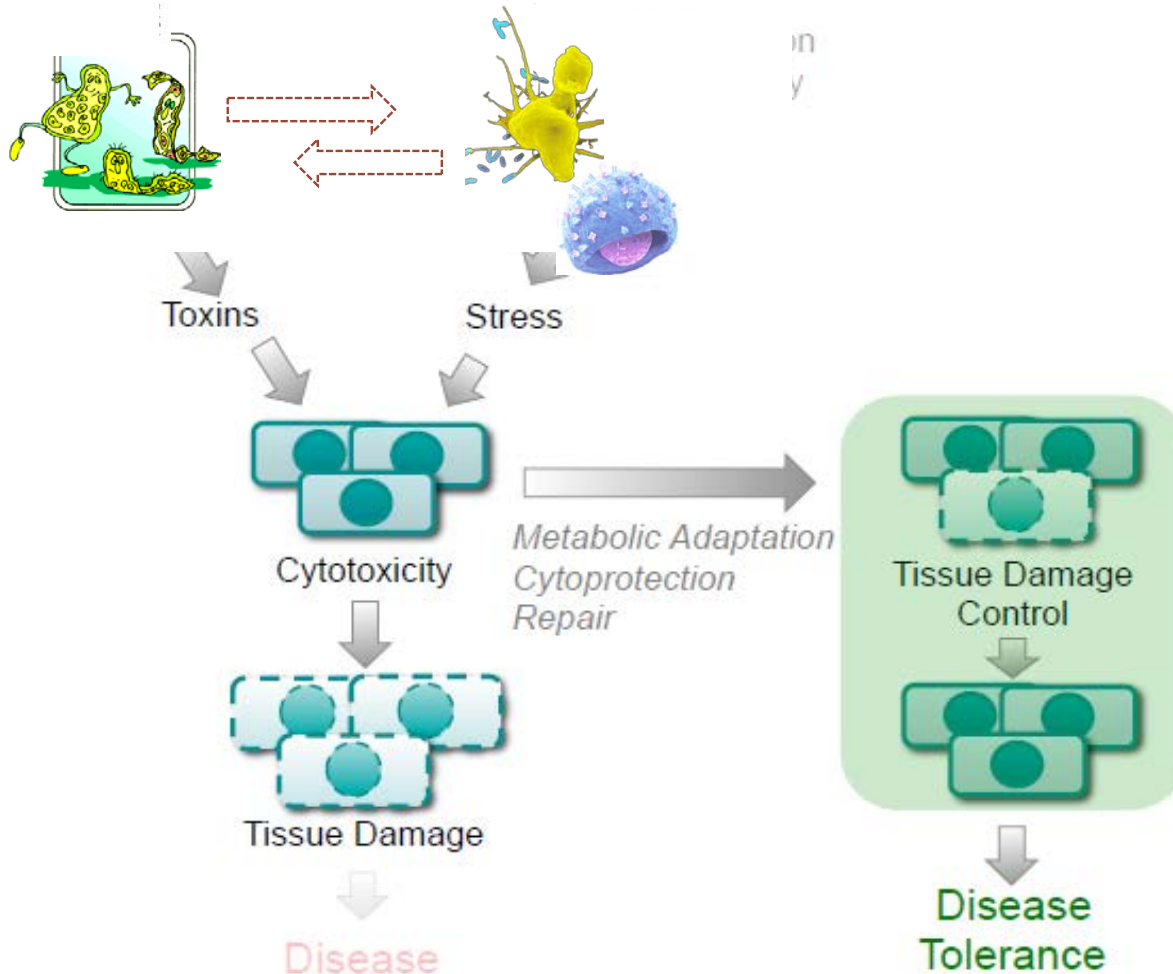
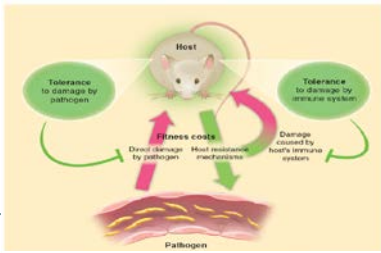
Infection



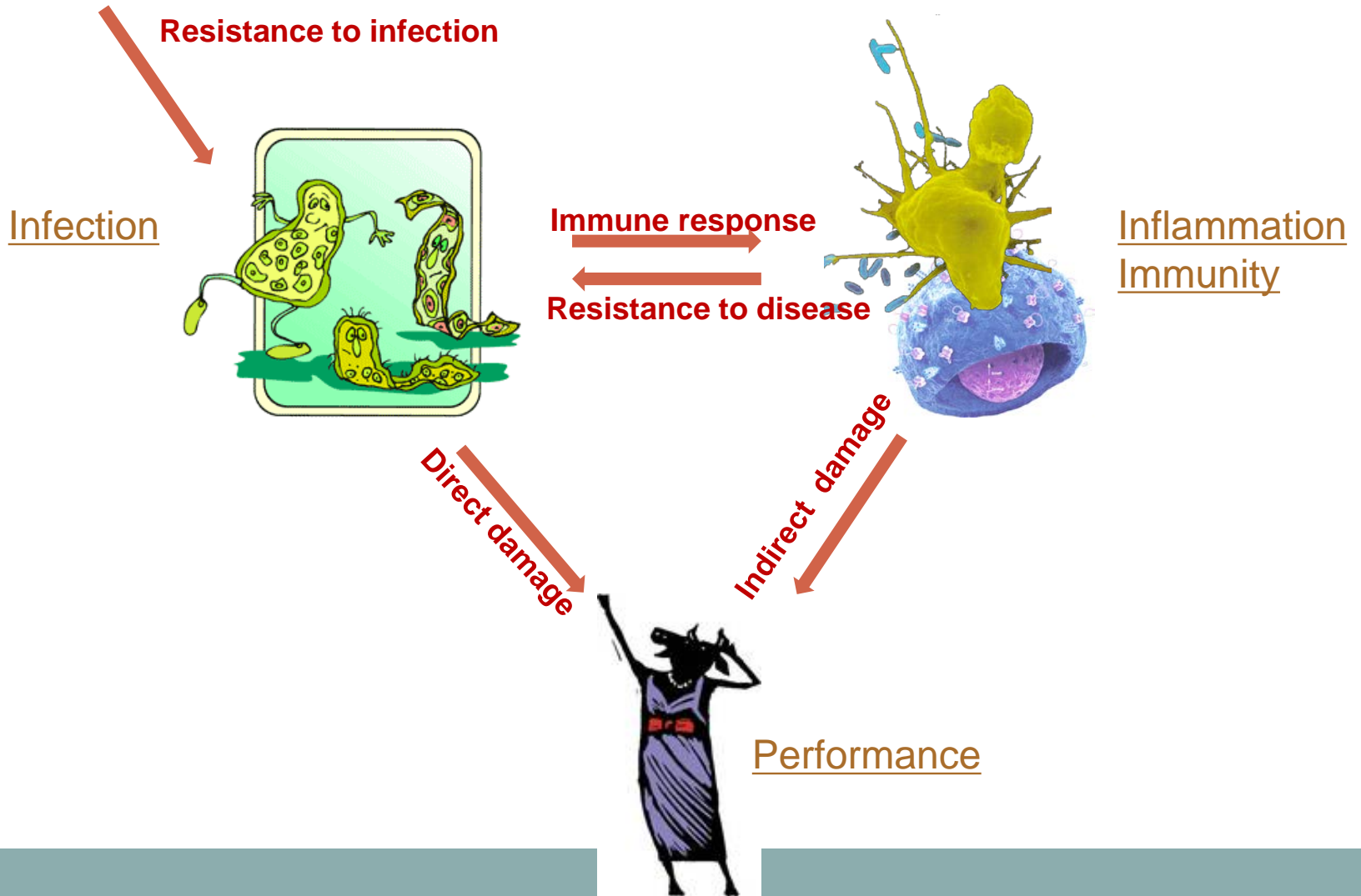
Inflammation
Immune response



Tolerance to infection and disease



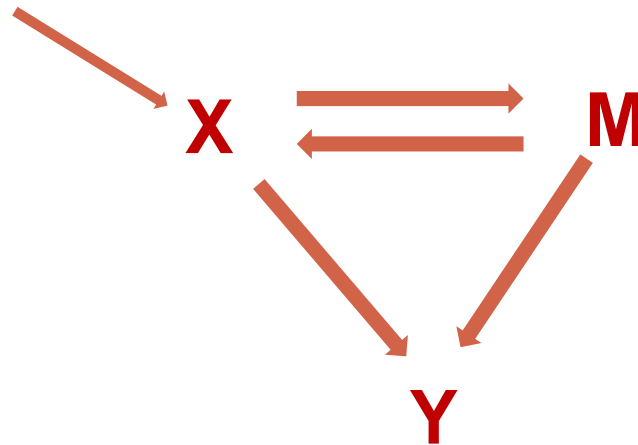
All together



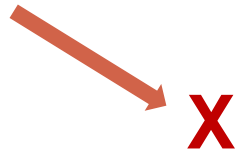
Models



X = PATHOGEN LOAD
M = IMMUNE INDEX
Y = PERFORMANCE



Resistance to infection

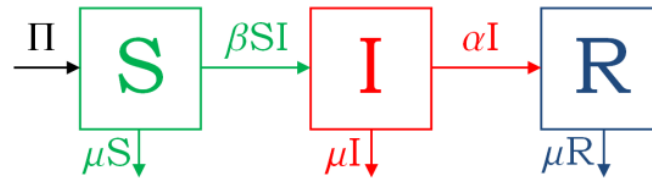


Resistance to infection



- Basic reproduction rate (R_0)

= number of new cases created by a single primary case in a susceptible population



- Individual probability to get infected

$$b_i = h^2 (1 - b_0) a_i + b_0$$

b_i = probability to become infected after contact with an infective

h^2 = heritability of the resistance to infection

a_i = degree of relationship between cows in contact

b_0 = average population transmission probability

Resistance to disease



Resistance to disease



Differential equations

$$\left(\begin{array}{l} \text{Pathogen load} \\ \text{rate of change} \end{array} \right) = \left(\begin{array}{l} \text{Pathogen} \\ \text{growth rate} \end{array} \right) - \left(\begin{array}{l} \text{Pathogen} \\ \text{killing rate} \end{array} \right)$$

$$\left(\begin{array}{l} \text{Immune index} \\ \text{rate of change} \end{array} \right) = \left(\begin{array}{l} \text{Immune index} \\ \text{entry rate} \end{array} \right) - \left(\begin{array}{l} \text{Immune index} \\ \text{exit rate} \end{array} \right)$$

Differents models for rates: Holling functional response, saturation term, logistic/exponential growth,

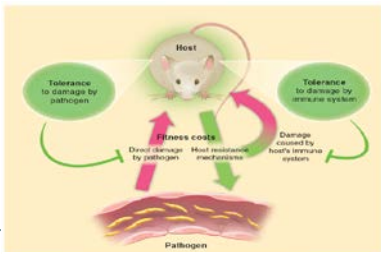
Direct tolerance



X



Y



Direct tolerance



Random regression models

$$E(Y_{ij} | b_{i0}, b_{i1}) = b_{i0} + b_{i1} X_i$$

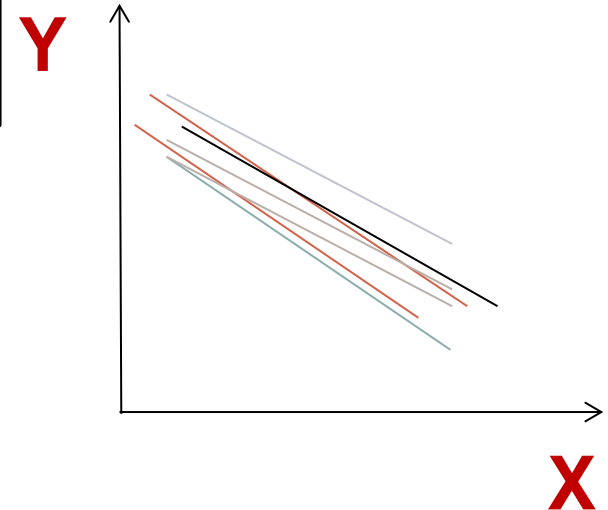
with $b_{i0} \sim N(\beta_0, \sigma_0^2)$ and $b_{i1} \sim N(\beta_1, \sigma_1^2)$

b_{i0} = vigour

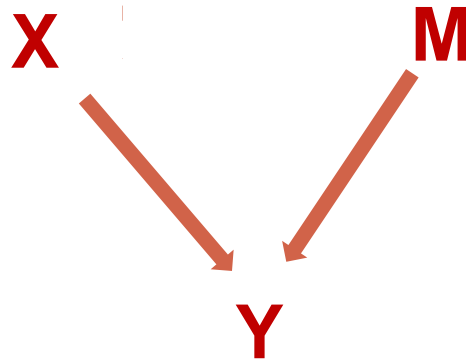
b_{i1} = tolerance

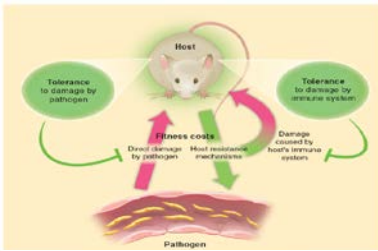
X_i = pathogen load

Y_{ij} = host infected



Direct & indirect tolerances

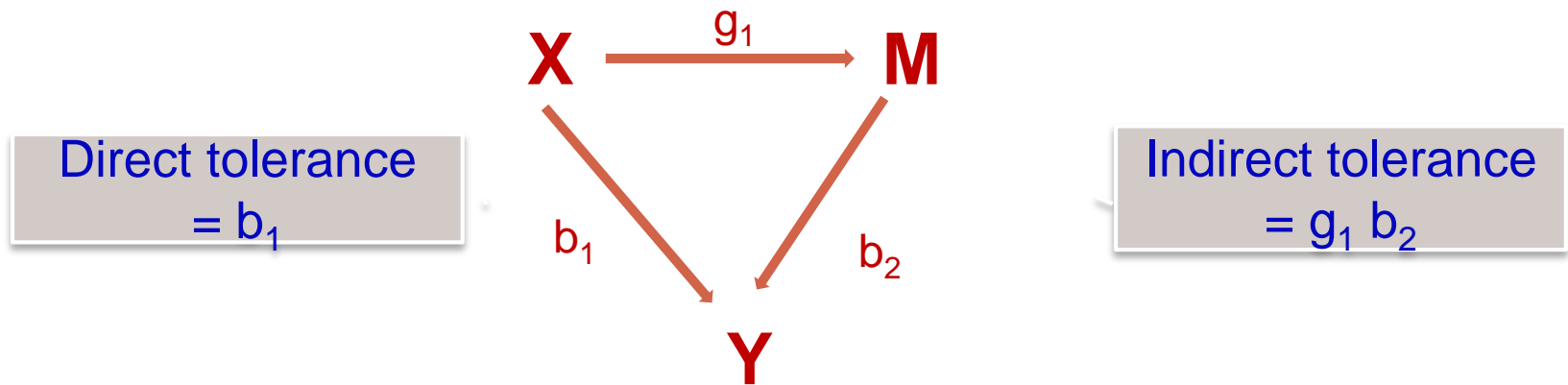




Tolerances



Structural equation models



$$E(M_{ij} | b_{i0}, b_{i1}) = g_{i0} + g_{i1} X_{ij}$$

with $g_{i0} \sim N(g_0, \tau_0^2)$ and $g_{i1} \sim N(g_1, \tau_1^2)$

$$E(Y_{ij} | b_{i0}, b_{i1}, b_{i2}) = b_{i0} + b_{i1} X_{ij} + b_{i2} M_{ij}$$

with $b_{i0} \sim N(\beta_0, \sigma_0^2)$, $b_{i1} \sim N(\beta_1, \sigma_1^2)$ and $b_{i2} \sim N(\beta_2, \sigma_2^2)$

Tolerances



Counterfactual interpretation

$Y(x,m)$ = potential performance observed in an individual with infection level set at the value x and immune index set at the value m

m_0 = level of immune index under health

m_1 = level of immune index under infection

Natural direct effect

$$E[Y(\text{infection}, m_0) - Y(\text{health}, m_0)]$$

Natural indirect effect

$$E[Y(\text{infection}, m_1) - Y(\text{infection}, m_0)]$$

Costs



RESISTANCES
TOLERANCES

Mechanisms

- **Innate constitutive protective barriers: low**

- lack of diversification process
 - low rates of cell turnover
 - small tissue mass in innate response

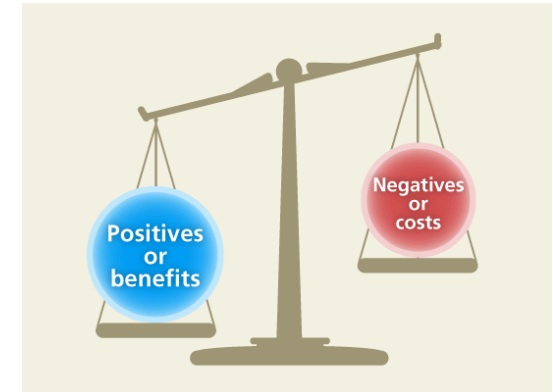
- **Inflammation: high**

- increased production of acute phase protein
 - changes in energy and nutrient metabolism
 - anorexia, fever

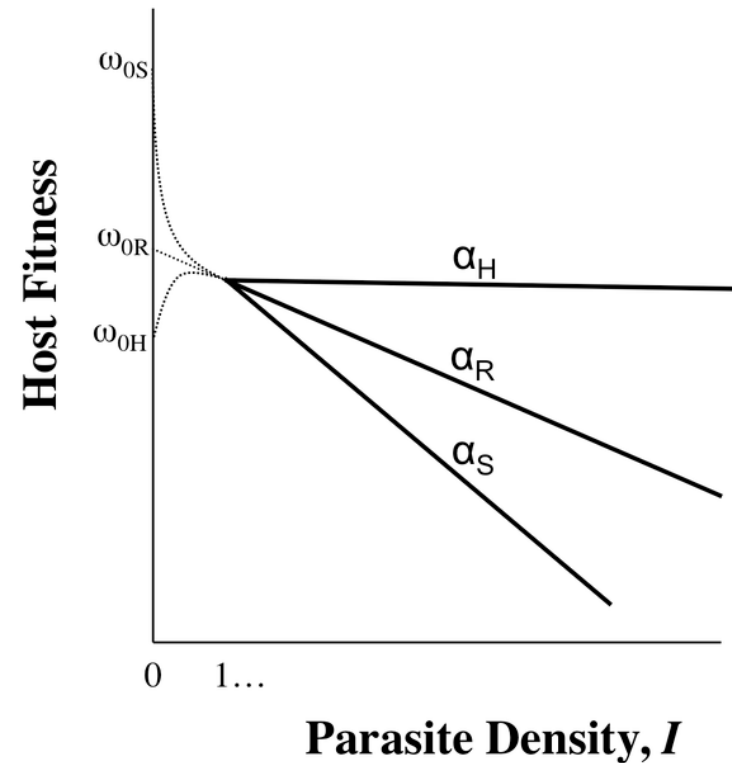
- **Immune response:**

- high for cell-mediated response, low for humoral response
 - innate more costly than adaptive response
 - reduced tolerance

- **Tissue damage repair**



Life-history traits theory

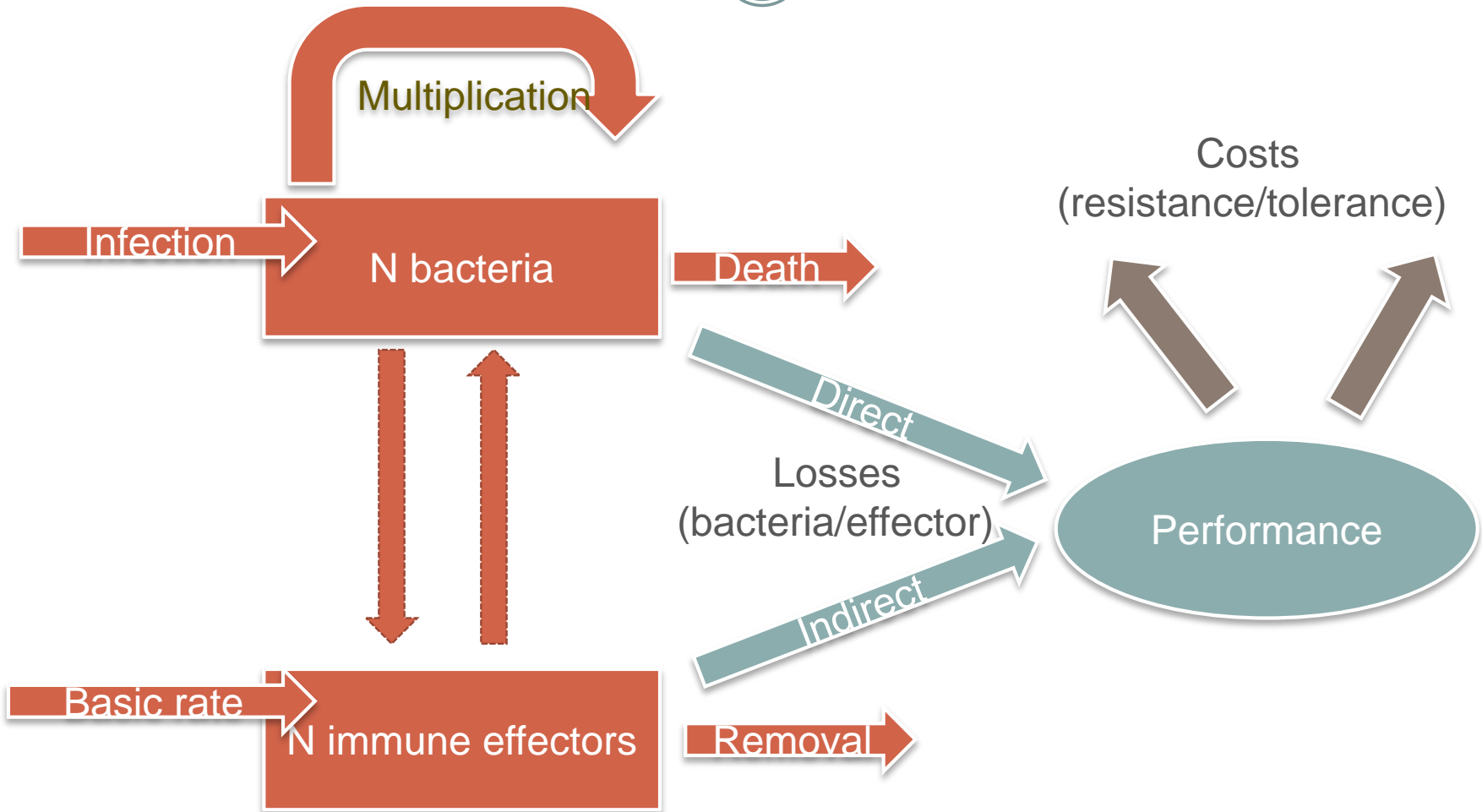


Bovine mastitis



RESISTANCES
TOLERANCES
COSTS

Model



B = N bacteria
 C = N immune effectors
 P = Performance

Contact rate

Resistance to infection

Logistic growth

Resistance to disease

Costs

Direct tolerance

Indirect tolerance

Infection: $D_{t+\Delta t} = \sum_i c_i v \beta^i B_t^i$ for $i = 1, 2, \dots, I,$

Multiplication: $N_{t+\Delta t} \sim \text{Poisson}[\gamma B_t (1 - B_t/K_B)]$

Bacteria killing: $K_{t+\Delta t} \sim \text{Poisson}[\alpha C_t B_t \rho / (1 + (\tau \alpha B_t))]$

Basic flow: $M_{t+\Delta t} \sim \text{Poisson} [\omega (C_1 - C_t)]$

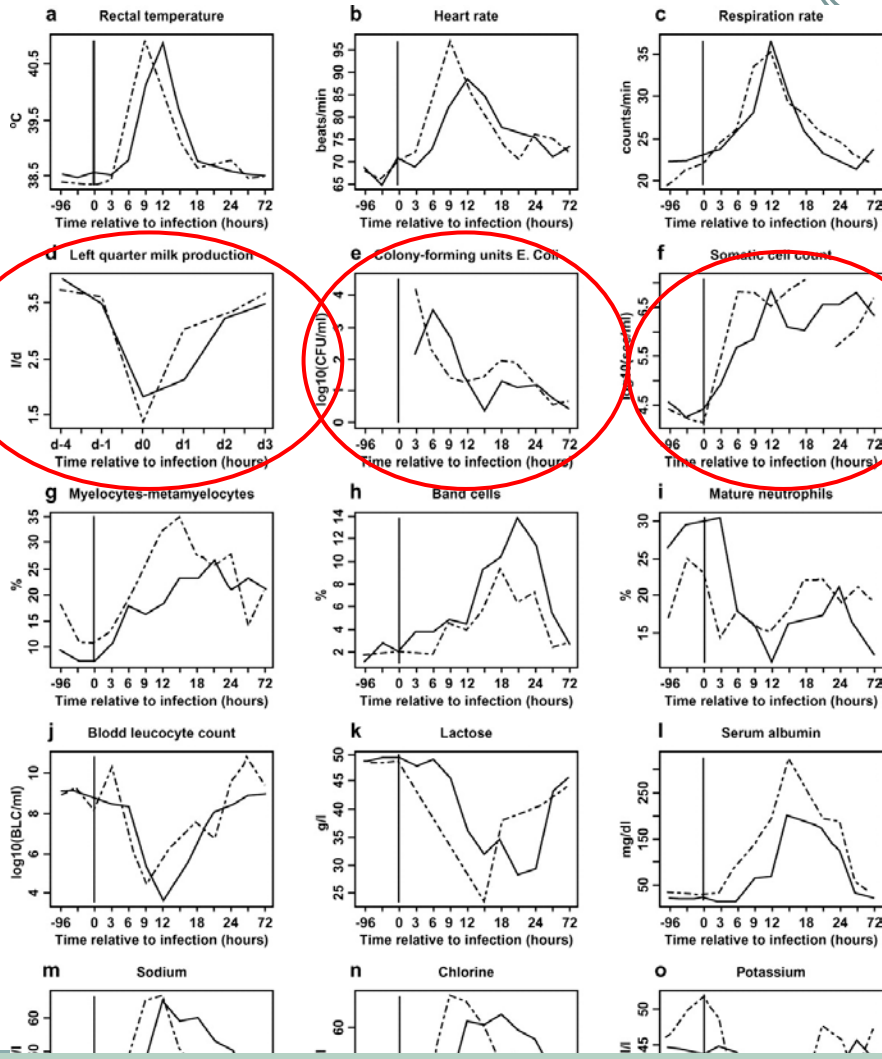
Extra-recruitment: $G_{t+\Delta t} \sim \text{Poisson} [\mu B_t C_t / (K_m + B_t)]$

Extra-removal: $S_{t+\Delta t} \sim \text{Poisson} [\alpha C_t B_t \rho / \theta (1 + (\tau \alpha B_t))]$

Costs: $P_1 = P^{\text{Max}} (1 - \rho c_\rho - \lambda_b c_b - \lambda_c c_c)$

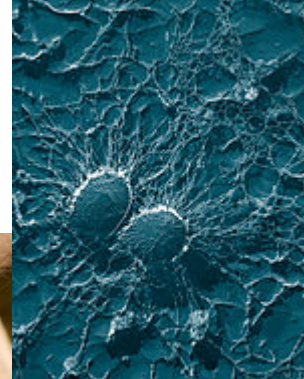
Losses: $P_t = P_1 - [B_t L_B (1 - \lambda_b) + C_t L_C (1 - \lambda_c)]$

Data



Experimental studies

Data

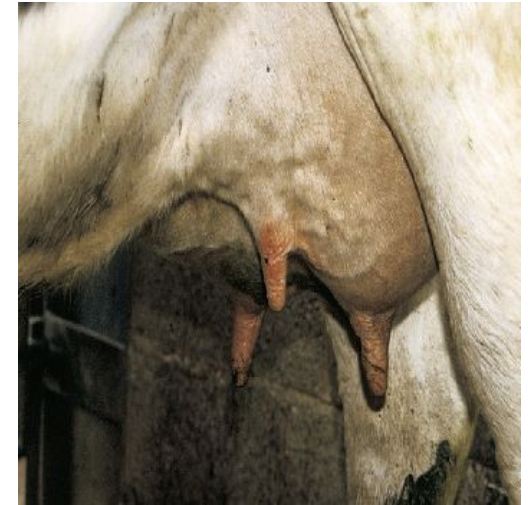


Field data

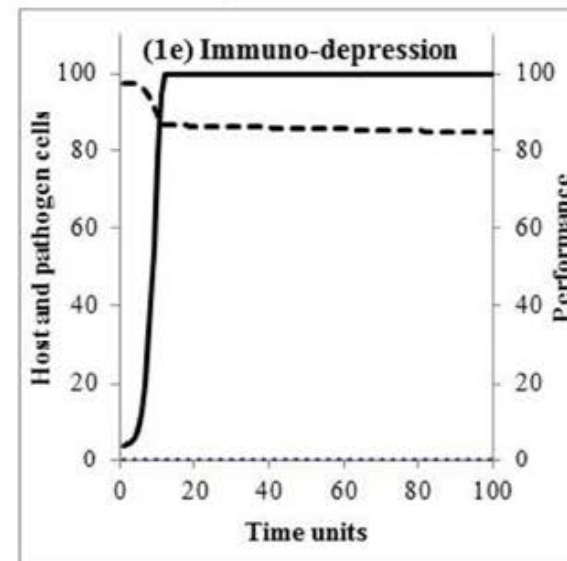
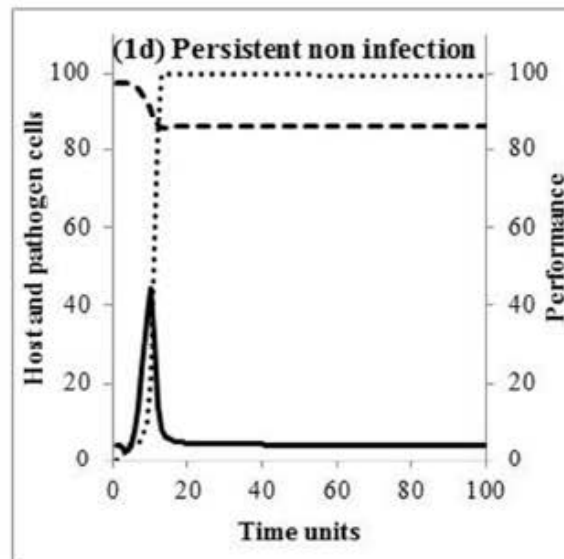
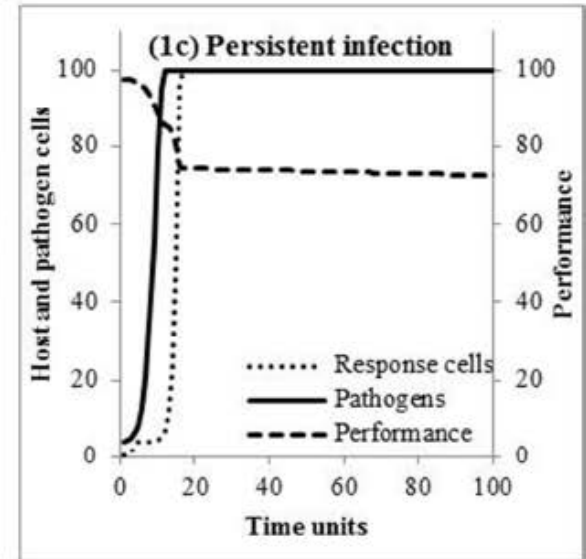
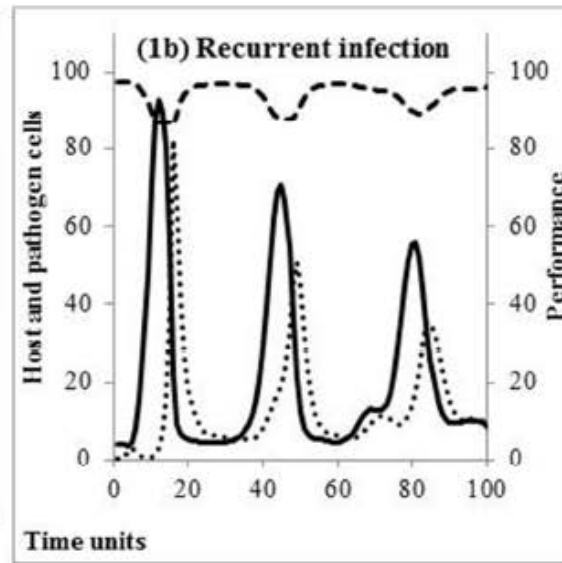
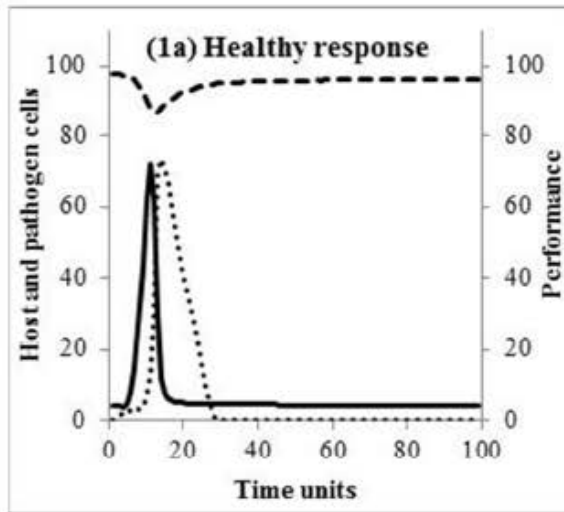
Results



- **Healthy response**
- **Recurrent infection**
- **Persistent response without infection**
- **Persistent response and infection**
- **Severe immunodeficiency**



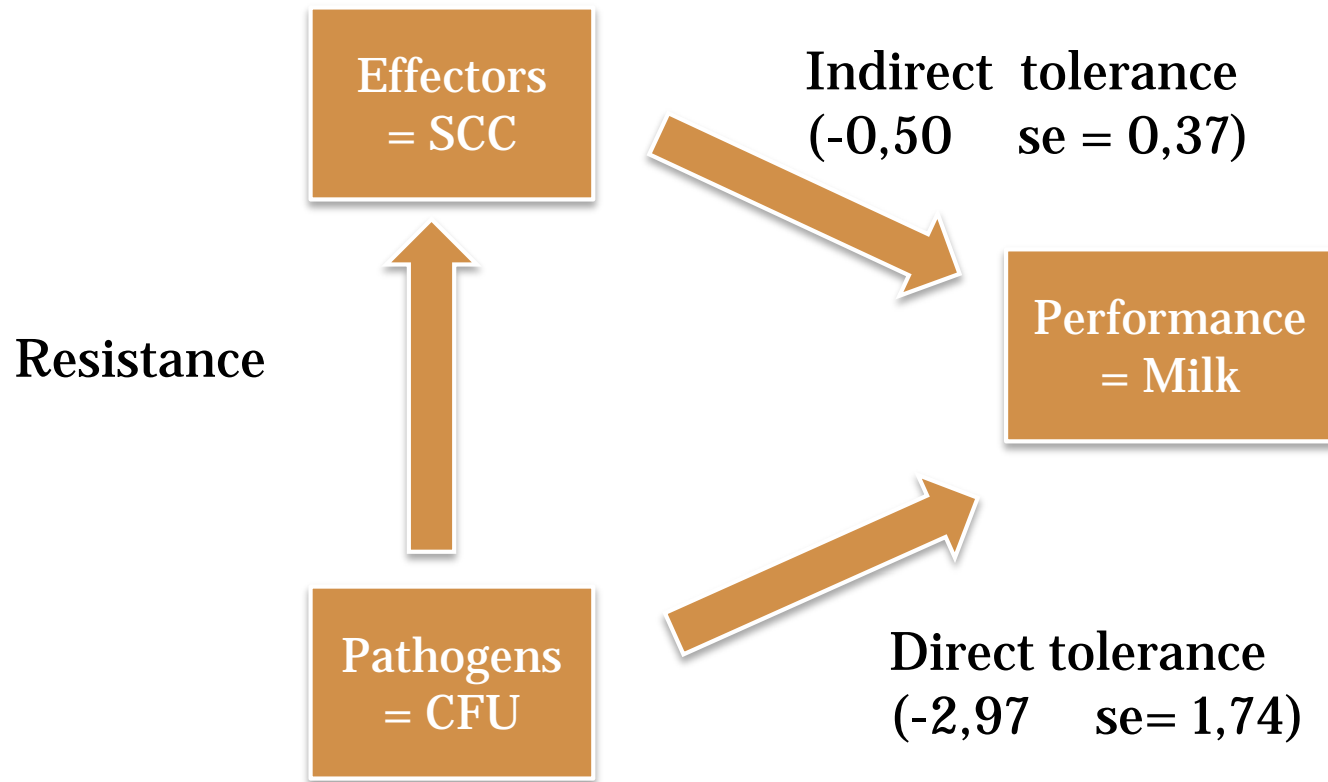
Results



Results



Structural equation model

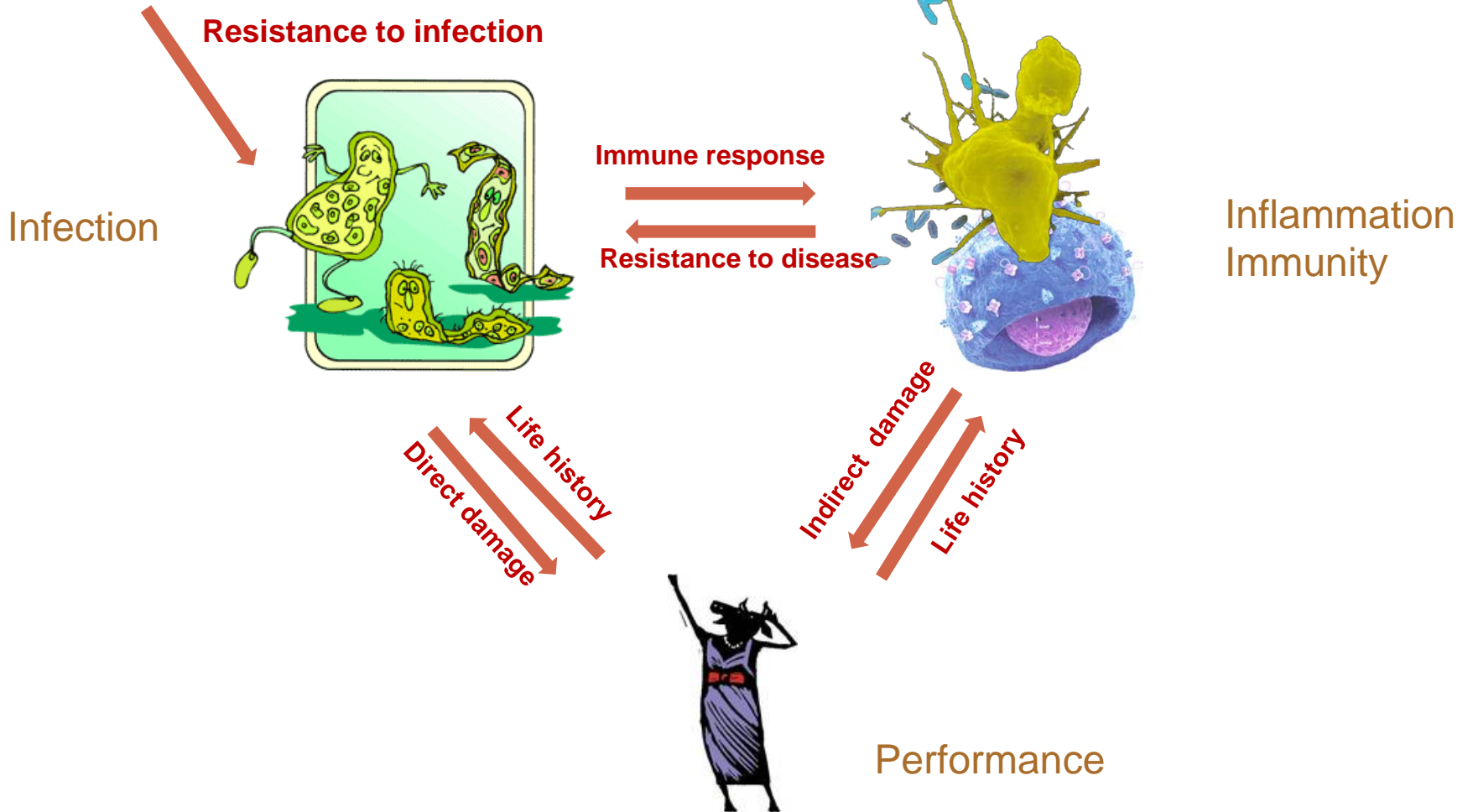


Conclusions



	Mechanisms	Costs	Models
Resistance			
- infection	Behavior Inflammation	++	R_0
- disease	Immunity		ODE
Tolerance			
- direct	Immuno-regulation	++	RRM
- indirect	Tissue damage control and repair		SEM

Further work





Results



E. coli bovine mastitis

Symbol	Signification	Values
Parameters with the same values in all simulations		
K_B	Maximum concentration of pathogens	$10^6/\mu\text{L}$
K_C	Maximum concentration of response cells	$10^7/\mu\text{L}$
P^{Max}	Maximum performance	100 units
γ	Pathogen logistic growth rate	1 pathogen / $\mu\text{L}/\text{h}$
τ	Time for a response cell to capture and kill pathogens	1 h/cell
θ	Pathogen concentration killed per response cell	10 pathogens/cell
c	Contact rate between hosts	0.1/h
Parameters for the different response scenarios		
K_M	Pathogen concentration such that response cells reach the infection site in 1 time unit Healthy response Recurrent infection	10 cells/ μL 10 000 cells/ μL
α	Pathogen clearance rate Healthy response Persistent infectious response	0.005 pathogen/cell/h 0 pathogen /cell/h
ω	Recruitment rate of response cells during health Healthy response Persistent non-infectious response	0.5 cells/h 0.01 cells/h
μ	Extra-recruitment rate of response cells during infection Healthy response Immuno-depression	2 cells/ $\mu\text{L}/\text{h}$ 0 cells/ $\mu\text{L}/\text{h}$

Resistance/Tolerance



	Parameters with uniform distributions	
β	Infectiousness	U[0;0.01]
L_C	Loss associated with each response cell	U[0; 25/ K_C]
L_B	Loss associated with each pathogen	U[0; 25/ K_B]
c_p, c_b, c_c	Resistance, direct and indirect tolerance costs	U[0; 0.1]
v	Resistance to infection Low Average High	U[0; 0.001] U[0; 0.01] U[0.009; 0.01]
ρ λ_b, λ_c	Resistance to disease Direct and indirect tolerances Low Average High	U[0; 0.1] U[0; 1] U[0.9; 1]