



# Thresholds, Dynamics, and Bifurcations

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## Objectives

- Introduce in-host models (Mathematical Immunology)
- Introduce intraguild predation models (Mathematical Ecology)
- Introduce relevant bifurcation theory (Dynamical Systems)



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Is Mathematical Biology a branch of applied maths?

What is Mathematical Biology?



# Classic In-host infection model

susceptible host cell:	$\frac{dT}{dt} = \lambda - \mu T - \beta TV$
infected host cell:	$\frac{dI}{dt} = \beta TV - \delta I$
free virus:	$\frac{dV}{dt} = pI - cV - \beta_v TV$

- Nowak and May, 2000, *Virus dynamics: mathematical principles of immunology and virology*, Oxford University Press
- Bonhoeffer et al., 1997, *Virus dynamics and drug therapy*, PNAS



# Classic In-host infection model: life-cycle

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# Classic In-host infection model: equilibria

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- Infection-free equilibrium:
- Coexistence equilibrium:



# Classic In-host infection model: stability and invasion

$$\begin{array}{ll} \text{resident: } x = (T) & \frac{dx}{dt} = f(x, y) = \lambda - \mu T - \beta TV \\ \text{invader: } y = \begin{pmatrix} I \\ V \end{pmatrix} & \frac{dy}{dt} = g(x, y) = \begin{pmatrix} \beta TV - \delta I \\ pI - cV - \beta_v TV \end{pmatrix} \end{array}$$



# Classic Immune Model

			cellular immunity
susceptible host cell:	$\frac{dT}{dt} = \lambda - \mu T - \beta TV$		
infected host cell:	$\frac{dI}{dt} = \beta TV - \delta I - pIB$		←
free virus:	$\frac{dV}{dt} = N\delta I - cV - pVB$		←
immune cells:	$\frac{dB}{dt} = (a^*I + aV)B - bB$		
	immune cell recruitment		humoral immunity

- Ciupe and Heffernan, 2017, *In-Host Modeling*, Infectious Disease Modelling
- Murase et al., 2005, J. Math. Biol.





# Classic Lotka-Volterra Predator Prey Model

- ❖<sub>1</sub> Prey grow exponentially at per capita rate  $r$  in absence of the predator.
- ❖<sub>2</sub> Prey are killed at a constant per capita rate  $a$  per predator.
- ❖<sub>3</sub> Each prey killed due to predation gives rise to  $c$  predators.
- ❖<sub>4</sub> Predator death is at a constant per capita rate  $m$ .

The model consists of a pair of differential equations. Here  $N$  and  $P$  are the prey and predator densities, respectively.

$$\frac{dN}{dt} = rN - aNP, \quad (1)$$

$$\frac{dP}{dt} = caNP - mP. \quad (2)$$



# The Volterra Model

In 1931, Volterra published an analysis of a predator-prey model with a prey carrying capacity:

$$\frac{dN}{dt} = rN \left(1 - \frac{N}{K}\right) - aNP, \quad (3)$$

$$\frac{dP}{dt} = caNP - mP. \quad (4)$$

The model can be rescaled to yield

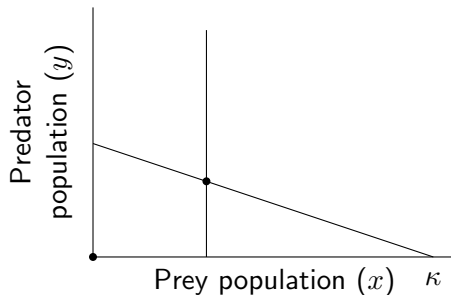
$$x' = x \left(1 - \frac{x}{\kappa}\right) - xy, \quad (5)$$

$$y' = \mu y(x - 1). \quad (6)$$



# The Volterra Model State Space

These equations represent a similar vector field to the original Lotka-Volterra model. The  $x$  nullcline is no longer horizontal, but has the equation  $y = 1 - x/\kappa$ .



Computations show the interior equilibrium to be stable, the trivial equilibrium to be a saddle, and the solutions are oscillations decaying to the interior equilibrium.



# The Rosenzweig-MacArthur Model

Rosenweig and MacArthur (1963) are credited with the addition of a hyperbolic functional response to the classic predator-prey model.

$$\frac{dN}{dt} = rN \left( 1 - \frac{N}{K} \right) - \frac{aNP}{1 + bN}, \quad (7a)$$

$$\frac{dP}{dt} = c \frac{aNP}{1 + bN} - mP. \quad (7b)$$



# The Functional Response

Turchin (2003) provides an excellent discussion of the various responses of predators to changes in prey populations.

$$\begin{aligned}\frac{dN}{dt} &= NF(N) - H(N, P)P, \\ \frac{dP}{dt} &= G(N, P)P - mP.\end{aligned}$$

The three functions are usually referred to as

- the net per capita growth rate  $F$ , which is assumed independent of predation,
- the *functional response*  $H$ , which is the rate individual predators kill prey, and
- the *numerical response*  $G$ , which models the dependence of the per capita predator growth rate on population densities.



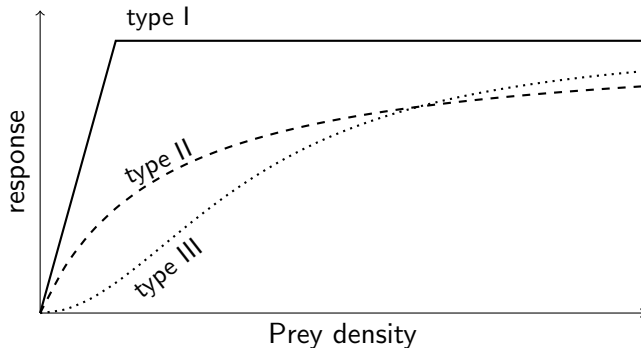
## **Predator-prey** relationships include

- Consumer-resource
- Plant-herbivore
- susceptible-infective
- host-pathogen
- host-parasite



# The Functional Response

Holling (1959) introduced a simple classification of functional responses of predators into type I, II, or III. Mathematically, type I and II are similar, and represent a saturating response.





# The Rosenzweig-MacArthur Model: equilibria

$$\frac{dN}{dt} = rN \left(1 - \frac{N}{K}\right) - \frac{aNP}{1 + bN}, \quad (8a)$$

$$\frac{dP}{dt} = c \frac{aNP}{1 + bN} - mP. \quad (8b)$$

- trivial equilibrium:  $(0, 0)$
- predator-free equilibrium:  $(0, K)$
- coexistence equilibrium:  $(\bar{N}, \bar{P})$ , if  $0 < \frac{m}{ca - mb} < K$

$$\bar{N} = \frac{m}{ca - mb},$$

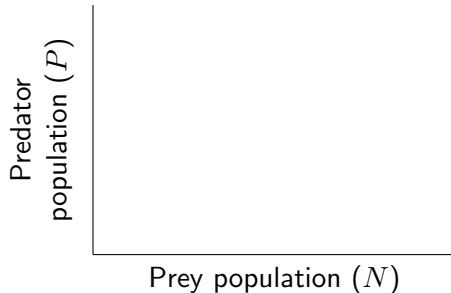
$$\bar{P} = r\bar{N}(1 + b\bar{N}) \left(1 - \frac{\bar{N}}{K}\right).$$





# The Rosenzweig-MacArthur Model: state-space

$$\begin{aligned}\frac{dN}{dt} = 0 & \quad r \left( 1 - \frac{N}{K} \right) - \frac{aP}{1 + bN} = 0, \\ \frac{dP}{dt} = 0 & \quad c \frac{aN}{1 + bN} - m = 0.\end{aligned}$$





# The Rosenzweig-MacArthur Model: Hopf-bifurcations

A stability analysis of the interior fixed point is simplified by writing the system in the *Kolmogorov form*

$$\frac{dN}{dt} = Nf(N), \quad (9a)$$

$$\frac{dP}{dt} = Pg(P), \quad (9b)$$

where

$$f(N, P) = r \left( 1 - \frac{N}{K} \right) - \frac{aP}{1 + bN}, \quad g(N, P) = c \frac{aN}{1 + bN} - m.$$

The interior equilibrium satisfies  $f(\bar{N}) = g(\bar{P}) = 0$ .



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The Jacobian at  $(\bar{N}, \bar{P})$  is

$$\begin{pmatrix} \bar{N} \bar{f}_N & \bar{N} \bar{f}_P \\ \bar{P} \bar{g}_N & \bar{P} \bar{g}_P \end{pmatrix}$$



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- $f(\bar{N}) = g(\bar{P}) = 0.$
- $\bar{g}_P = 0, \bar{f}_P < 0, \bar{g}_N > 0$



# The Rosenzweig-MacArthur Model: Hopf-bifurcations

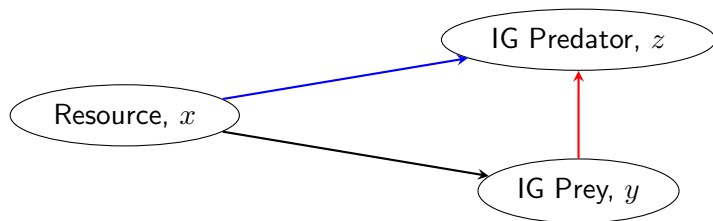
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- $f(\bar{N}) = g(\bar{P}) = 0$ .
- $\bar{g}_P = 0$ ,  $\bar{f}_P < 0$ ,  $\bar{g}_N > 0$
- The Hopf Bifurcation Theorem implies  $(\bar{N}, \bar{P})$  changes from a stable to unstable focus as  $\bar{f}_N$  changes from positive to negative.
- a stable limit cycle bifurcates from the equilibrium as the trace increases through zero.



# A Simple Intraguild Predation Model



$$\dot{x} = x(1 - x) - xy - \alpha_2 xz,$$

$$\dot{y} = \sigma_1 xy - \frac{yz}{1 + \beta_3 y} - \mu_1 y,$$

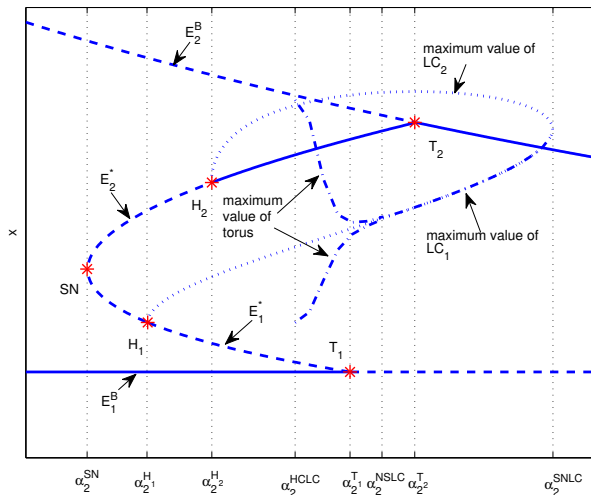
$$\dot{z} = \frac{\sigma_2 \alpha_2 xz}{1 + \beta_2 x} + \frac{\sigma_3 yz}{1 + \beta_3 y} - \mu_2 z.$$

- Type II interaction between IG prey and IG predator ( $\beta_3$ )
- Saturating numerical response of IG predator to resource ( $\alpha_2, \beta_2$ )

Xi Hu, 2014, PhD Thesis

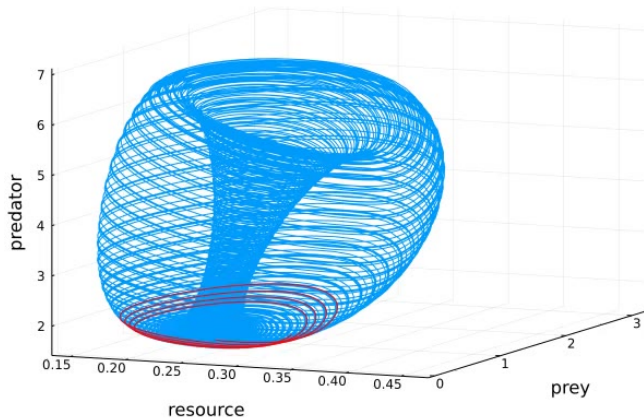


# Intraguild Predation: bifurcation diagram





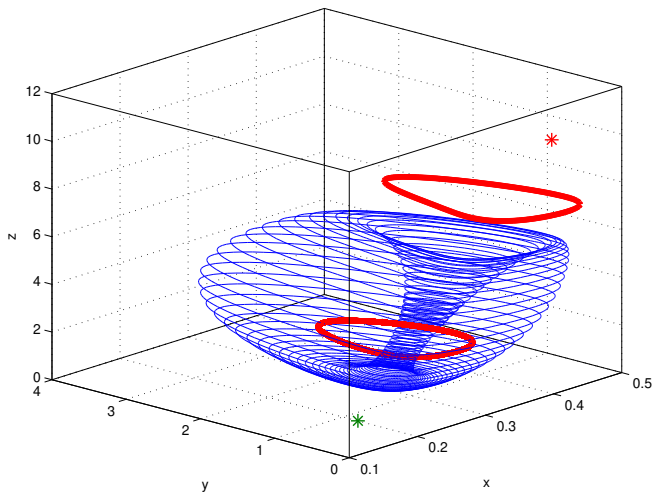
# The Stable Torus

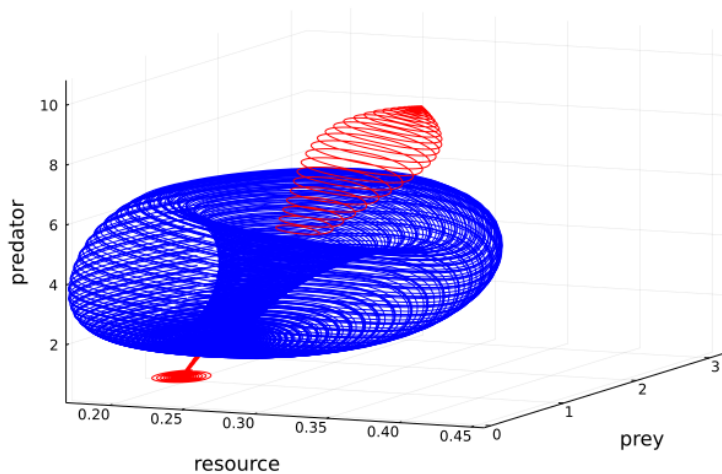






# Tristability with an Attracting Torus







# Poincaré section through torus

