

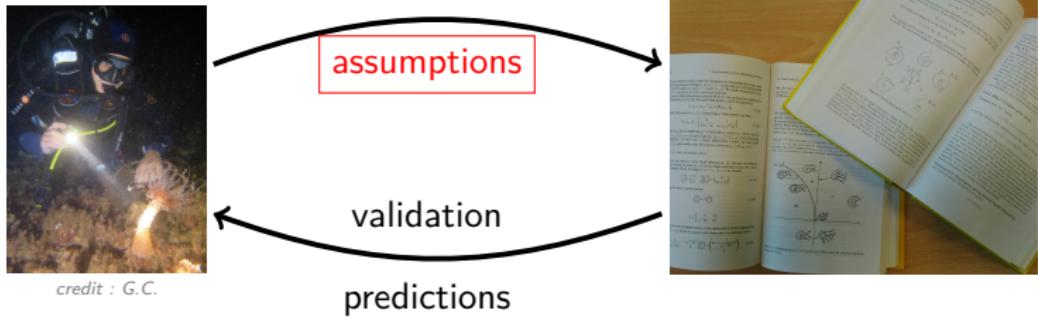
Structural sensitivity: predator-prey models to food webs, through Dynamic Energy Budget theory

Clément Aldebert¹, D. Nerini¹, M. Gauduchon¹, BW. Kooi², JC. Poggiale¹

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² Vrije Universiteit, Amsterdam, Netherlands 

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Assumptions \Rightarrow simplifications (e.g. : predation)

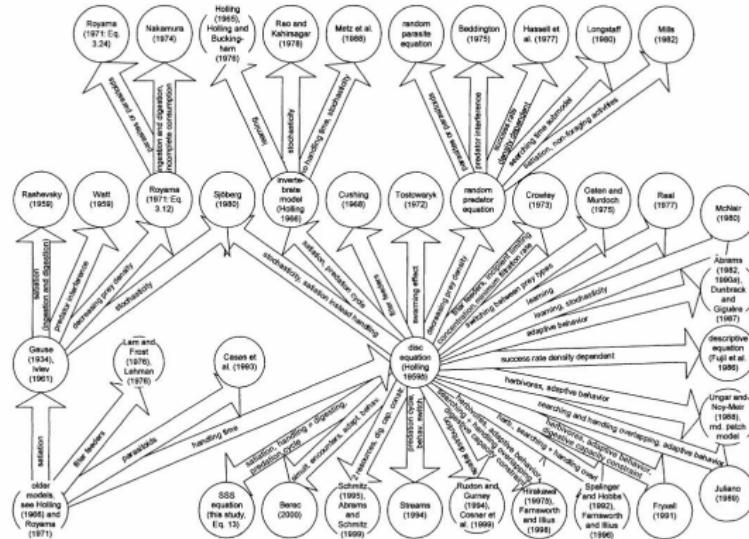
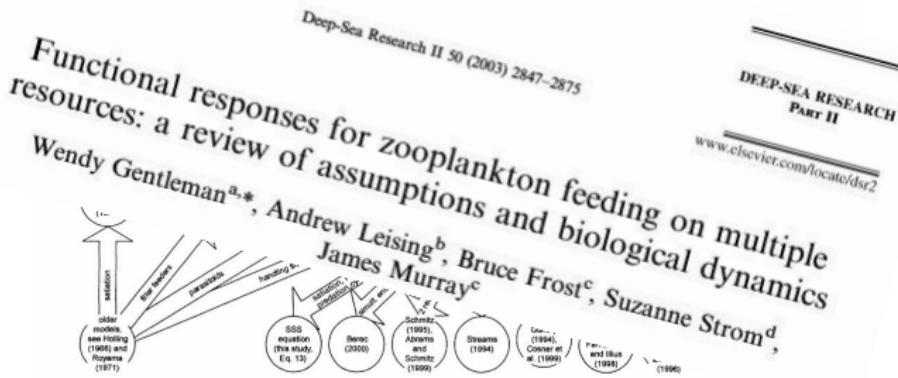
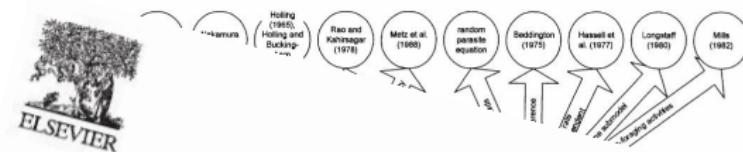


FIG. 1. A “family tree” of functional response models. (Jeschke et al., 2002)

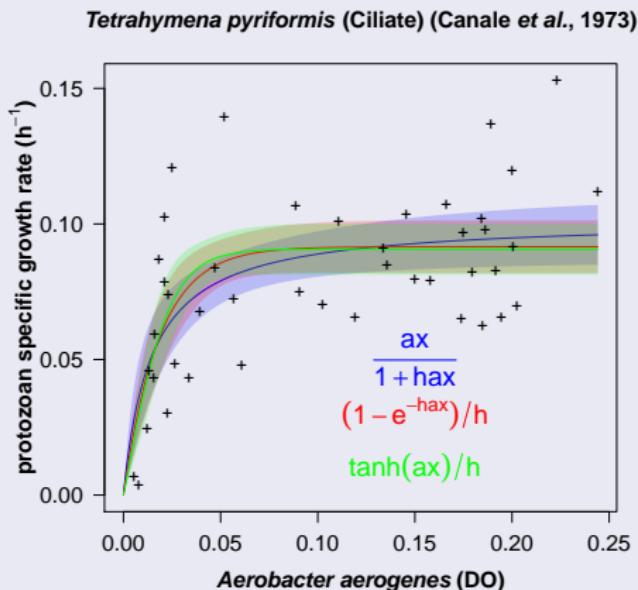
encounter / attack, handling, digestion / metabolism, spatial heterogeneity, individual variability, collective behaviour, ...

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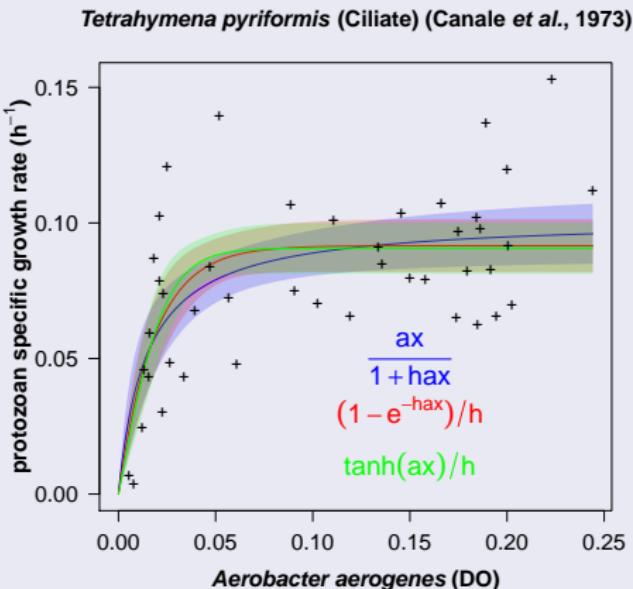
Modelling predation at population scale



functional response

- amount of prey eaten / unit of predator / time unit

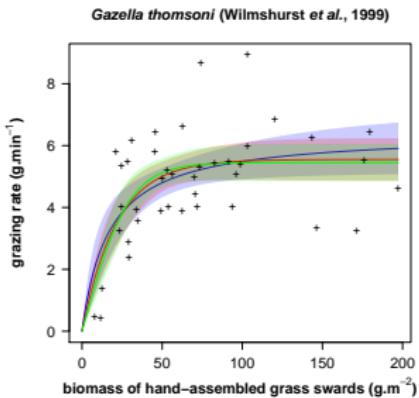
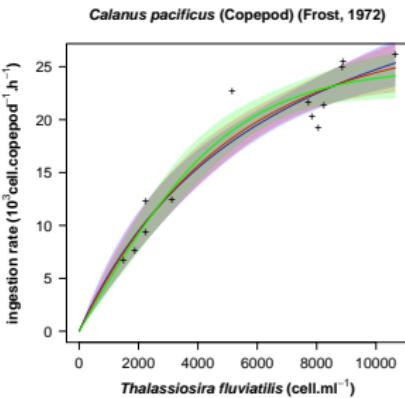
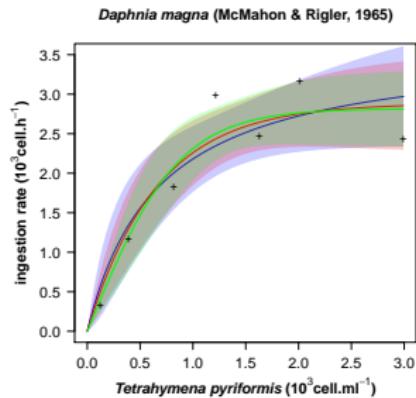
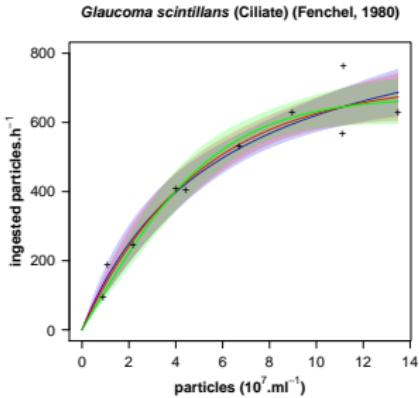
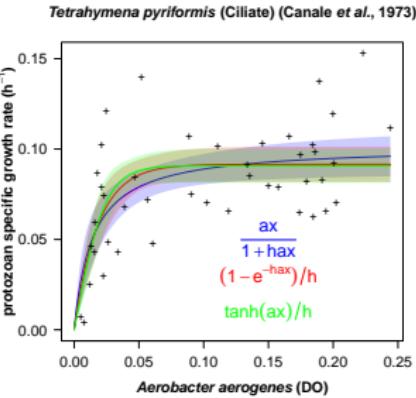
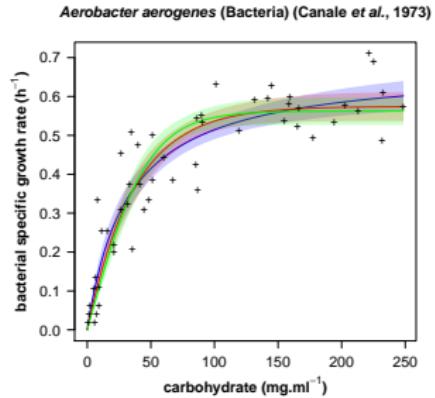
Modelling predation at population scale



functional response

- amount of prey eaten / unit of predator / time unit
- 3 functions with the same mathematical properties \Rightarrow same assumptions on **process shape**
- different assumptions on **underlying mechanisms**

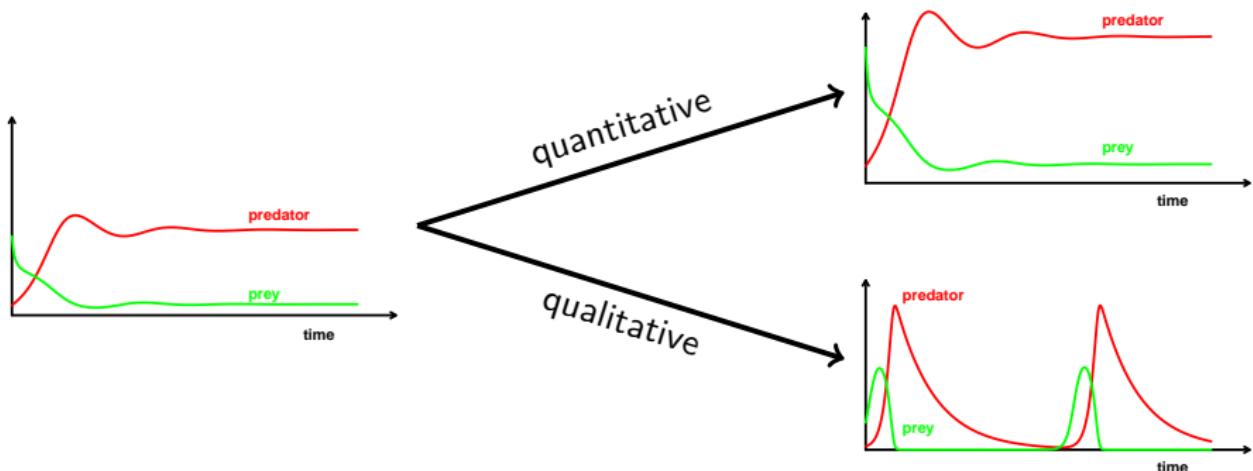
Modelling predation at population scale



Structural sensitivity

model change \Rightarrow change in predicted dynamics

- quantitative change (e.g. equilibrium value)
- qualitative change in dynamics (bifurcation) (Kuznetsov, 2004)



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- **functional response** in predator-prey, food chain and biogeochemical models

(*Myerscough et al., 1996, Gross et al., 2004, Fussmann & Blasius, 2005, Anderson et al., 2010, Cordoleani et al., 2011, M. Baklouti, pers. comm.*)

- infection in a host-pathogen model (*Wood & Thomas, 1999*)
- co-limitation in a multi-nutrient model (*Poggiale et al., 2010*)

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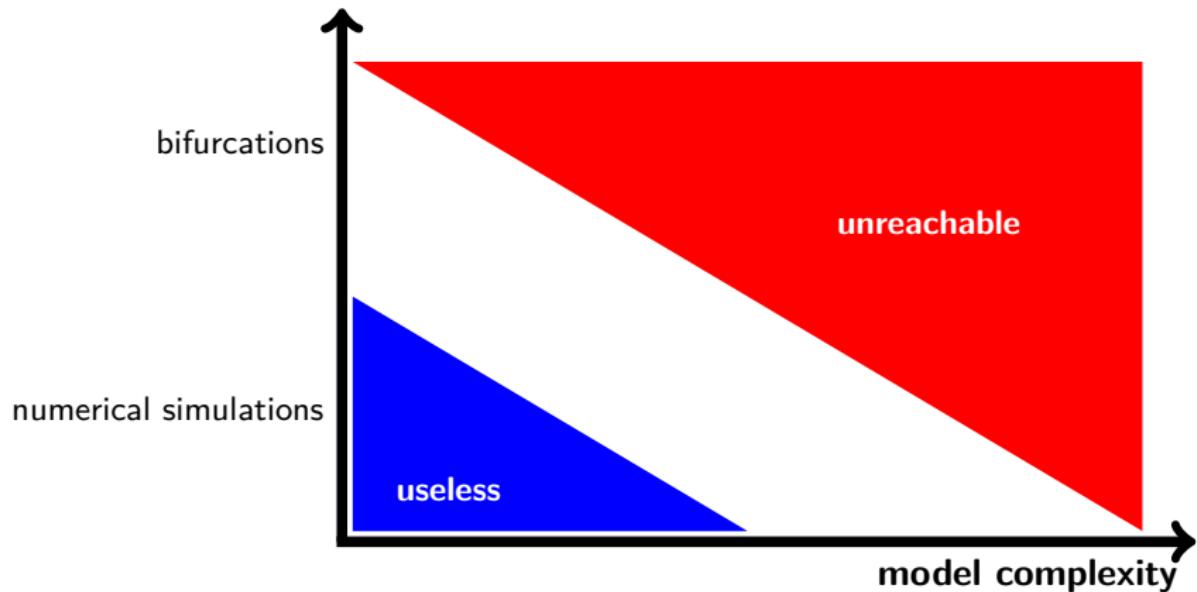
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previous studies : few state variables, “simple” dynamics (one stable state)

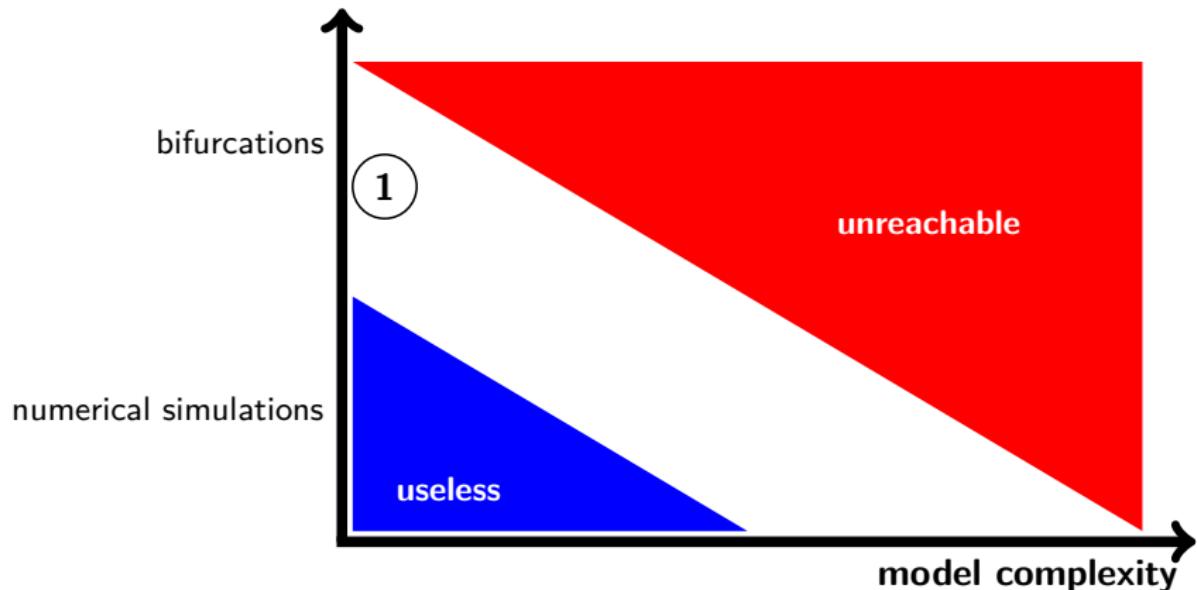
Structural sensitivity : \pm complex models

model understanding



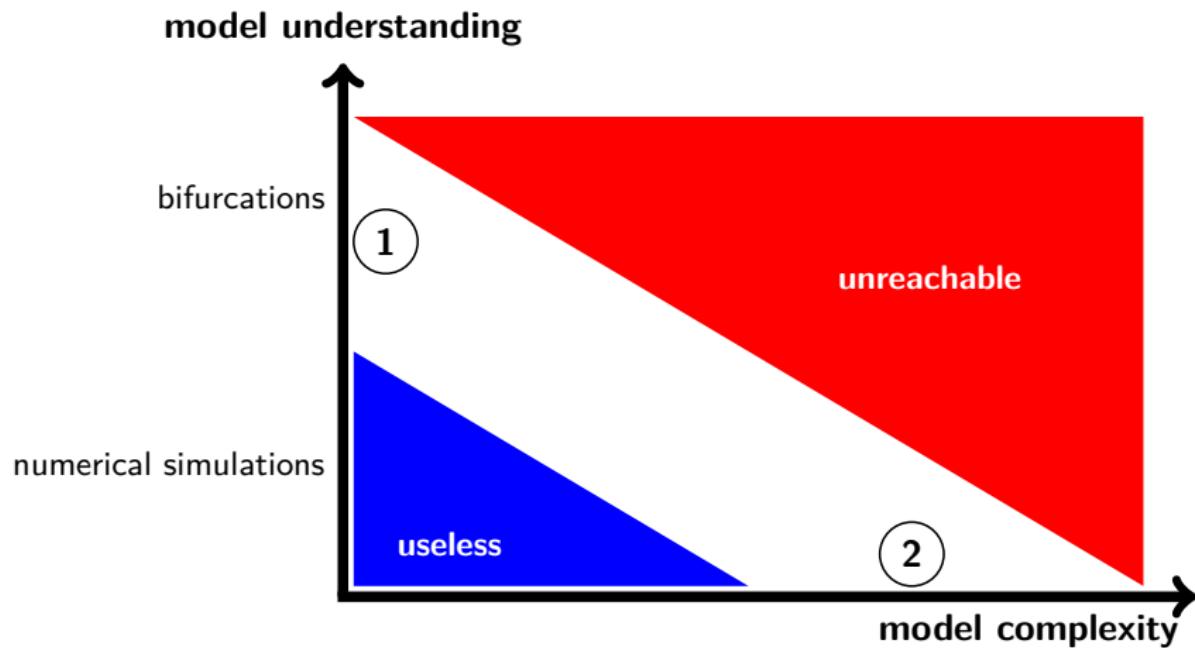
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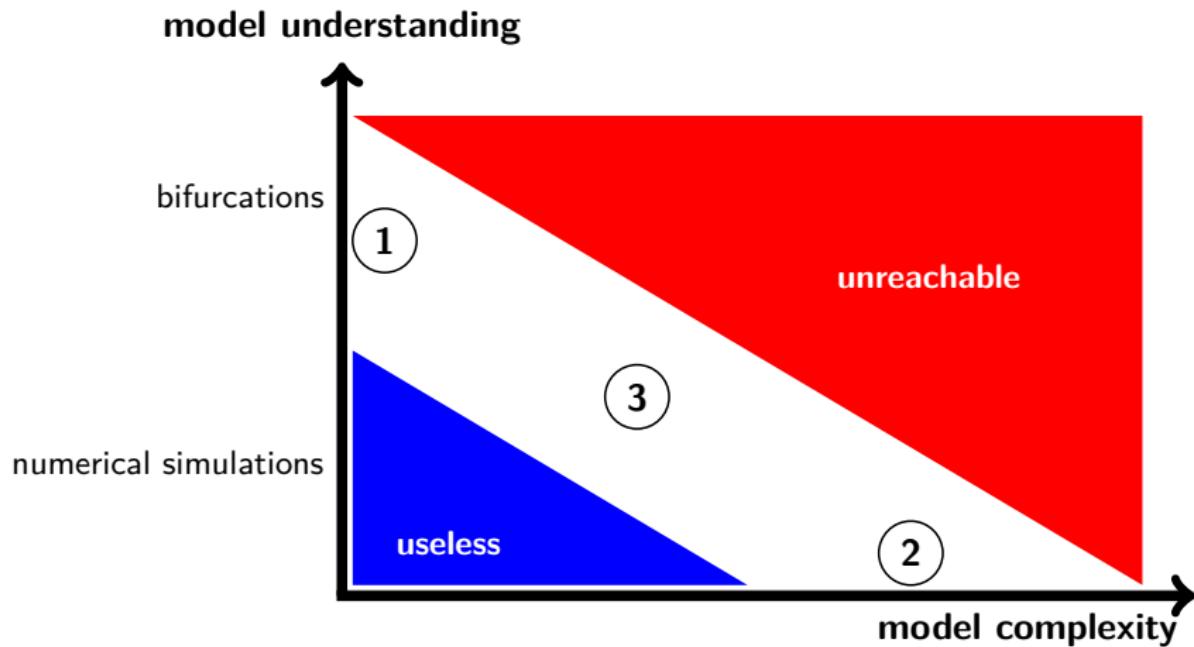
- ① predator-prey system : multiple stable states & **resilience**

Structural sensitivity : \pm complex models



- 1 predator-prey system : multiple stable states & **resilience**
- 2 food webs : predation vs **trophic complexity**

Structural sensitivity : \pm complex models



- 1 predator-prey system : multiple stable states & **resilience**
- 2 food webs : predation vs **trophic complexity**
- 3 predator-prey system : predation vs details on **metabolism**

Predator-prey model : Bazykin (1998)

$$\left\{ \begin{array}{l} \frac{dB_{prey}}{d\bar{t}} = [\lambda \bar{q}^\xi - \alpha - \beta \omega B_{prey}] B_{prey} - \bar{G}^\xi(B_{prey}) B_{pred} \left(\frac{M_{pred}}{M_{prey}} \right)^{-0.25} \\ \frac{dB_{pred}}{d\bar{t}} = [\lambda \bar{G}^\xi(B_{prey}) - \alpha - \beta B_{pred}] B_{pred} \left(\frac{M_{pred}}{M_{prey}} \right)^{-0.25} \end{array} \right.$$

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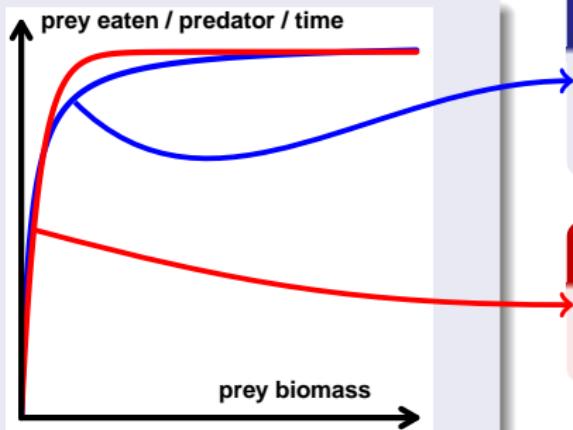
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Holling (1959) : handling

$$\bar{G}^H = \frac{a^H B_{prey}}{1 + h^H a^H B_{prey}}$$

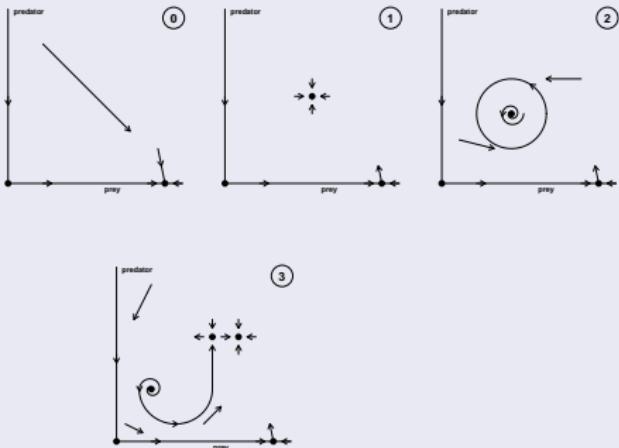
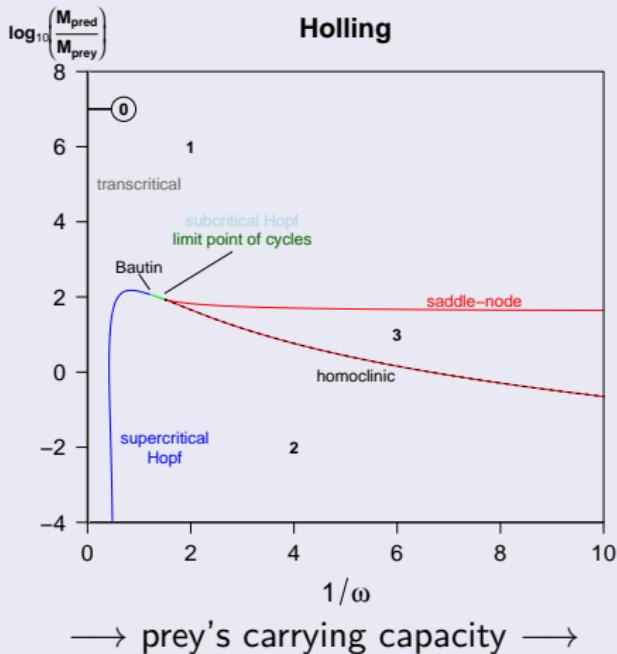
Ivlev (1955) : satiation

$$\bar{G}^I = \left(1 - e^{-h' a' B_{prey}} \right) / h'$$

(Aldebert et al., in press a)

Bifurcation diagram with Holling

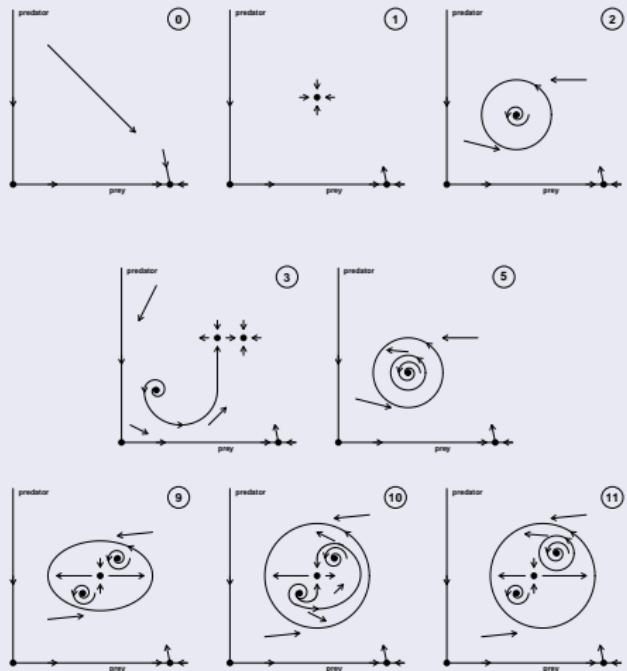
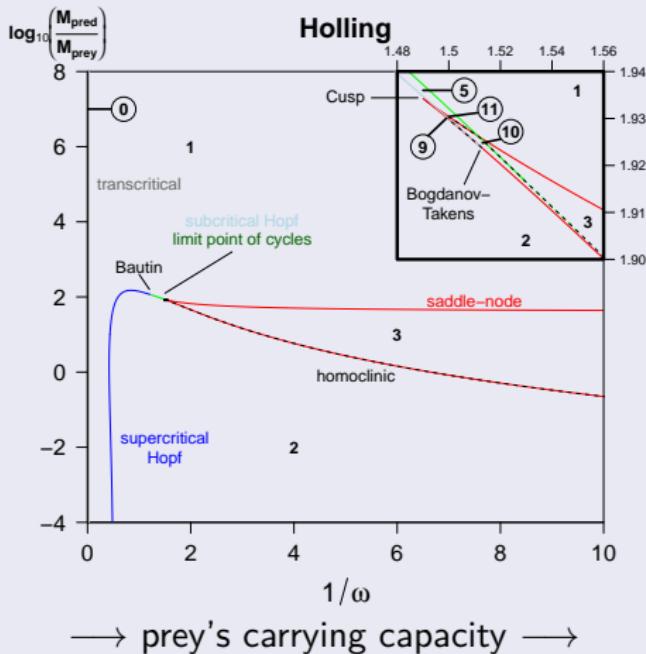
body mass ratio



(modified from Aldebert et al., in press a)

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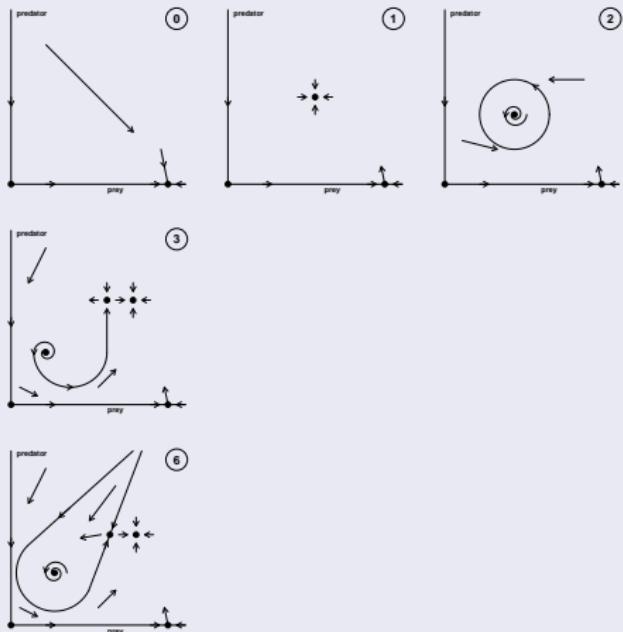
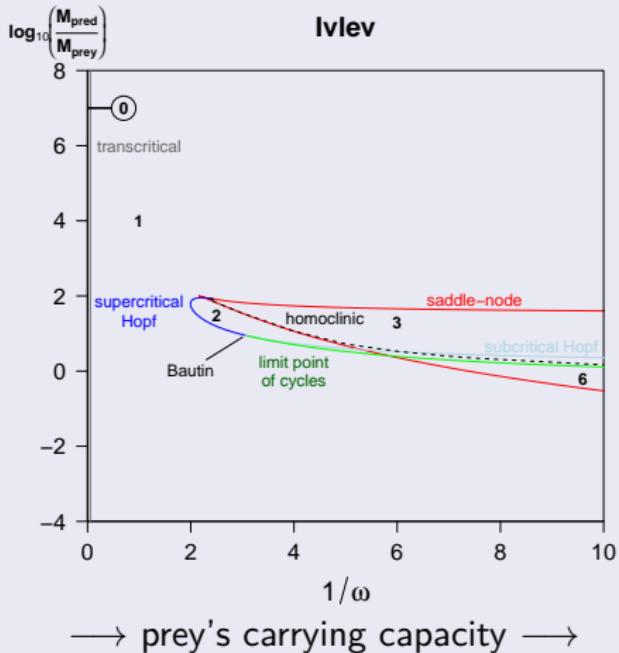
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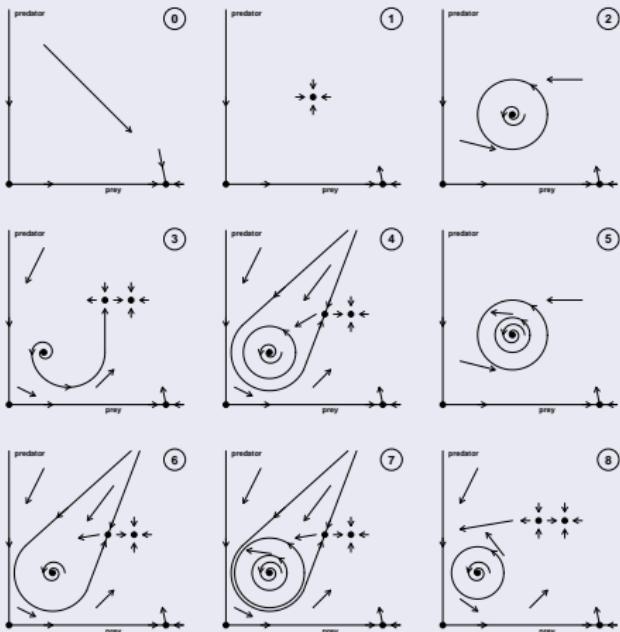
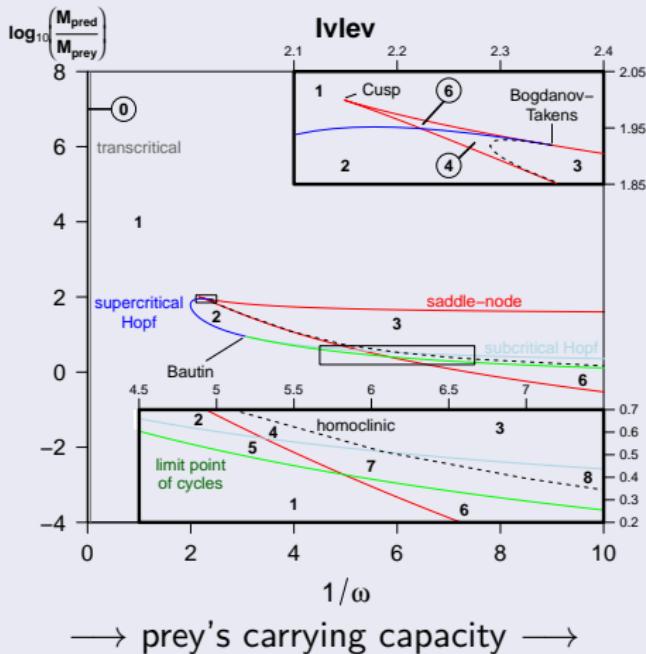
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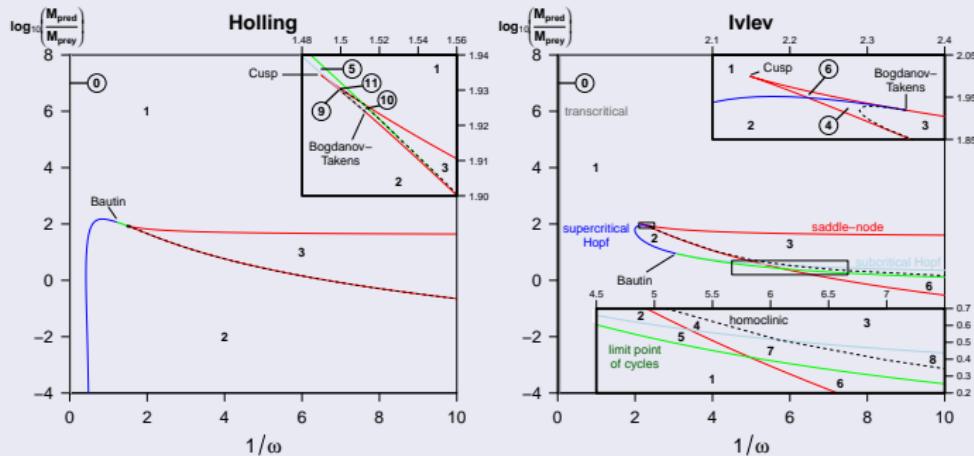
Bifurcation diagram with Ivlev

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Bifurcation diagram : Holling vs Ivlev

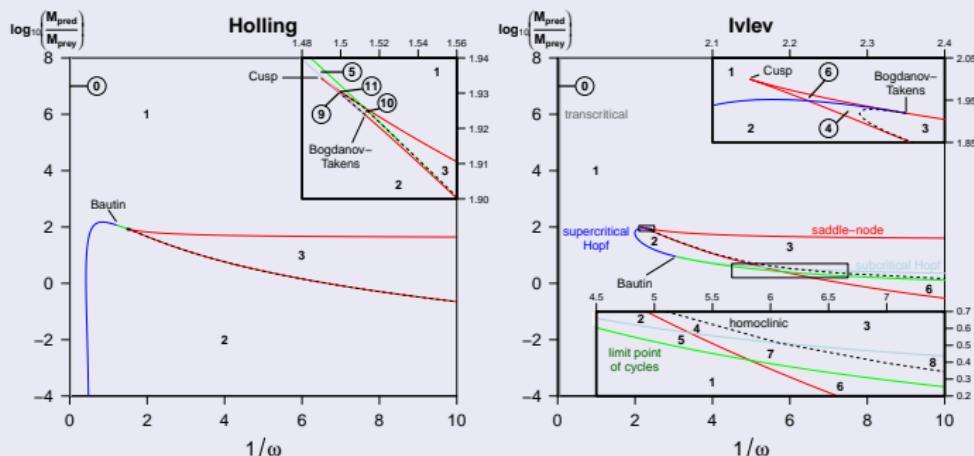


main differences

- stable equilibrium vs stable limit cycle : 26.0 % – 49.4 %

(modified from Aldebert et al., in press a, in prep. b)

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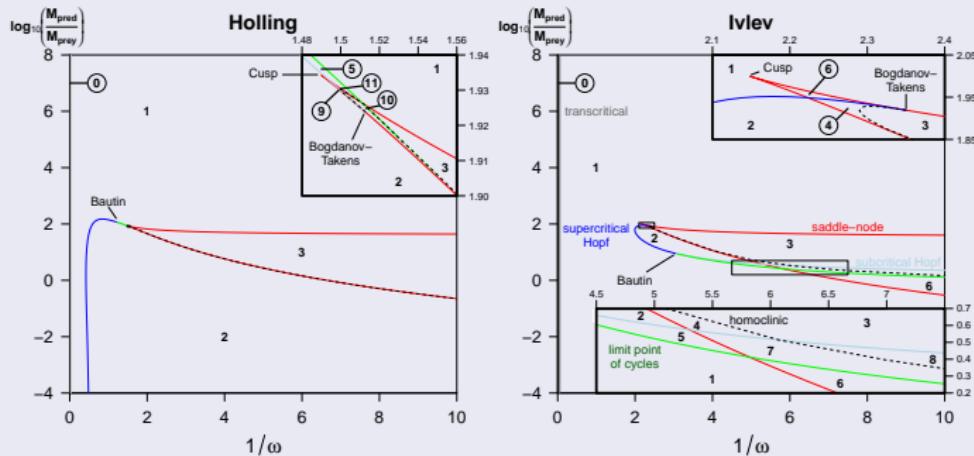
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NEW

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Bifurcation diagram : Holling vs Ivlev

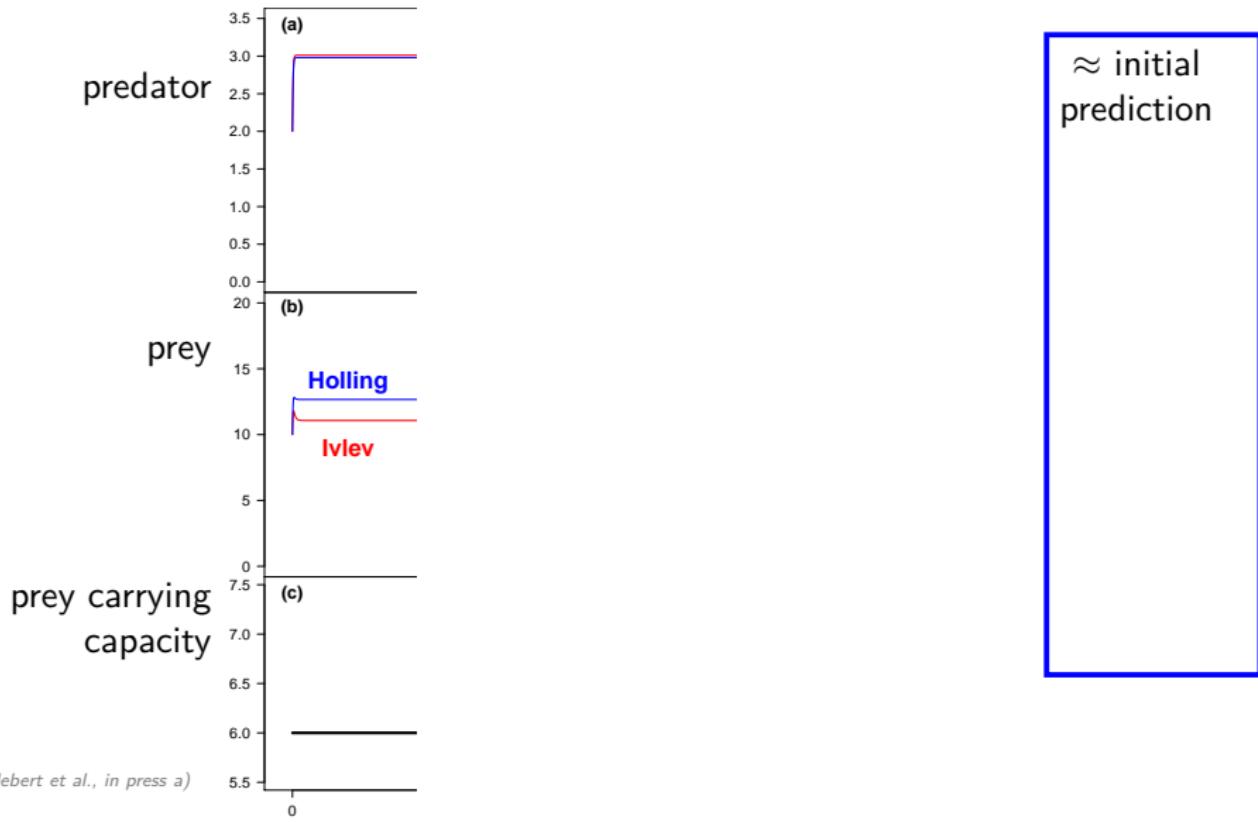


main differences

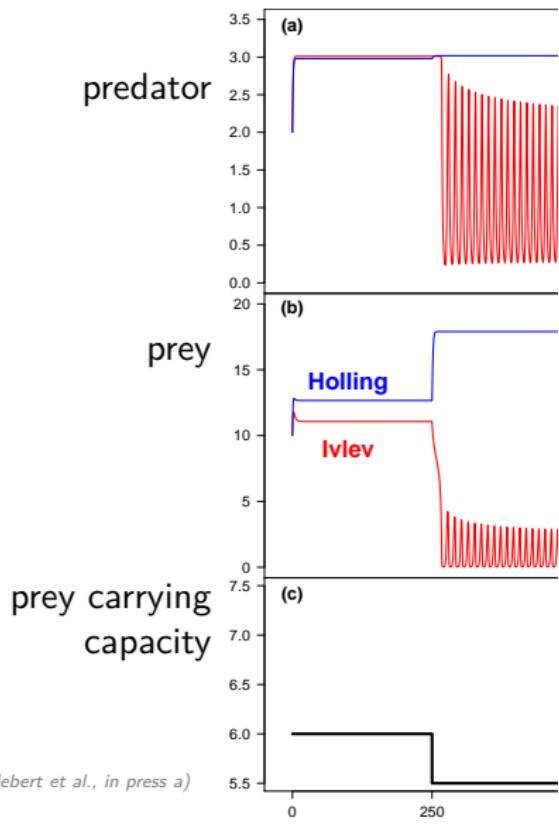
- stable equilibrium vs stable limit cycle : 26.0 % – 49.4 %
- multiple attractors (up to 3) with Ivlev : 0.1 % – 14.3 % **NEW**
- continuous switch : degenerated Bogdanov-Takens bifurcation (codim 3)

(modified from Aldebert et al., in press a, in prep. b)

Resilience in changing environmental conditions

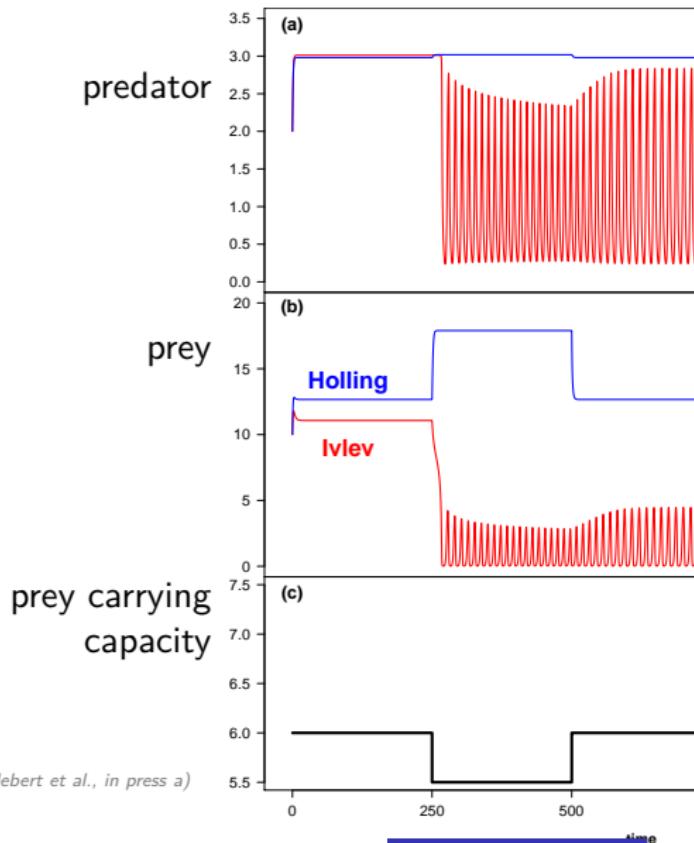


Resilience in changing environmental conditions



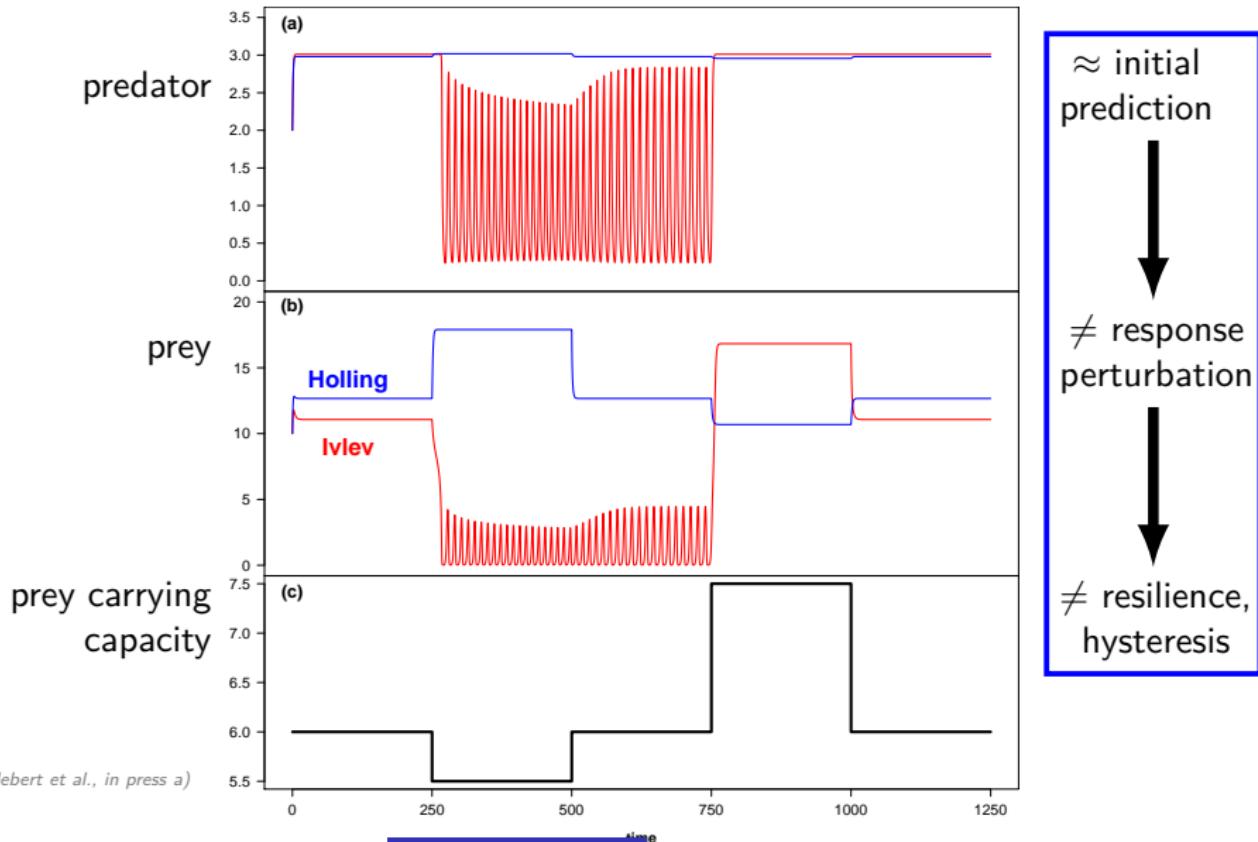
≈ initial prediction
↓
≠ response perturbation

Resilience in changing environmental conditions



≈ initial prediction
↓
≠ response perturbation
↓
≠ resilience,

Resilience in changing environmental conditions



(Aldebert et al., in press a)

Structural sensitivity in food webs

millions of theoretical networks (20-60 species, connectance : 0.1-0.3)



- randomly built using “niche model” \Rightarrow empirically consistent structural properties (*Williams & Martinez, 2000, 2004; Cattin et al., 2004; Allesina et al., 2008*)
- structure \Rightarrow build and parameterize a dynamical system

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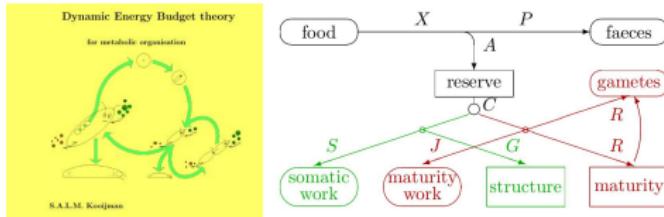
		persistence	equilib. / oscillations
trophic complexity	# species	0.76	NS
	connectance	0.37	0.36
functional response	formulation	-0.17	0.82
	maximum slope	0.12	-0.22
	maximum rate	0.07	0.07

TABLE : correlation coefficients

($\approx 6.10^7$ food webs were simulated to obtain this table)
computational effort : 3 years.processors

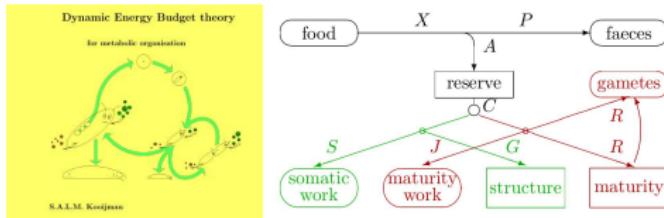
(Aldebert et al., in press b)

Structural sensitivity and metabolism : DEB theory



- focus on the individual, based on mechanistic assumptions on metabolism
- same framework for most species, metabolic classification of species

Structural sensitivity and metabolism : DEB theory



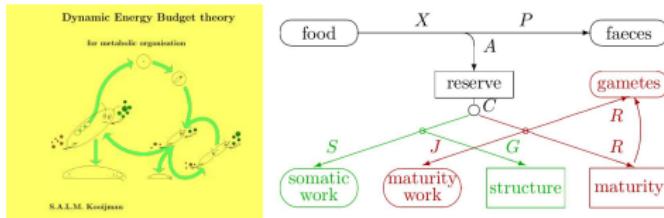
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structural sensitivity

- chemostat data \Rightarrow ± detailed models with a coherent theoretical framework
(Canale et al., 1973, Dent et al., 1976, Kooi & Kooijman, 1994b)
- multiple stable states in DEB model (reserve and maintenance)

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- multiple stable states in DEB model (reserve and maintenance)
- influence of **metabolism > functional response**
- change of functional response : no new bifurcations (and dynamics) if
minimum of biological realism (maintenance/mortality, explicit resource)

(Aldebert et al., in prep. a)

Conclusion

- predator-prey model : structural sensitivity affects the type and number of stable states → affects the **predicted resilience** of the system
- trophic complexity → food web persistence,
functional response → food web variability
- DEB : details on metabolism ↴ structural sensitivity to functional response

Concluding remarks and perspectives

Conclusion

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- trophic complexity → food web persistence,
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- DEB : details on metabolism ↘ structural sensitivity to functional response

Perspectives (Predictive Ecology group, UZH, Zürich)

- **quantification method** with multiple stable states (ongoing work)
- structural sensitivity and resilience in food webs, other networks
- dealing with **uncertainty in predictions** due to model construction

Thank you for your attention !!!

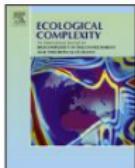


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Original Research Article

Structural sensitivity and resilience in a predator-prey model with density-dependent mortality

C. Aldebert *, D. Nerini, M. Gauduchon, J.C. Poggiale



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Does structural sensitivity alter complexity-stability relationships?

C. Aldebert *, D. Nerini, M. Gauduchon, J.C. Poggiale

EXTRA-SLIDES

$$\text{Switch Holling - Ivlev : } \bar{G}^\xi = \xi \bar{G}^H + (1 - \xi) \bar{G}^I$$

(Aldebert et al., in prep b)

Switch Holling - Ivlev : $\bar{G}^\xi = \xi \bar{G}^H + (1 - \xi) \bar{G}^I$

Bogdanov-Takens de point triple : forme normale

⇒ forme normale : Bob Kooi

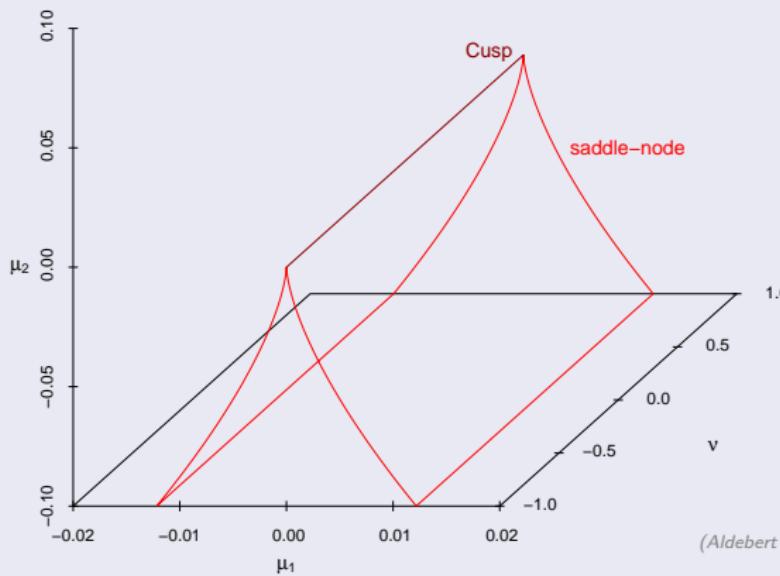
(Baer, Kooi, Kuznetsov, Thieme, 2006)

$$\begin{cases} \frac{d\xi}{dt} = \eta \\ \frac{d\eta}{dt} = -\mu_1 - \mu_2\xi + \nu\eta + \beta\xi\eta - \xi^3 - \eta\xi^2 \end{cases}$$

(Aldebert et al., in prep b)

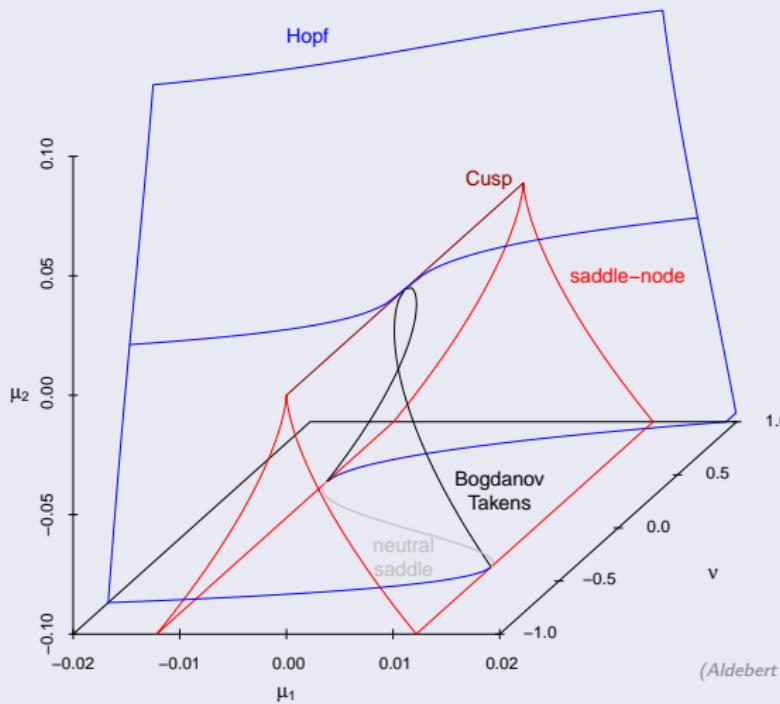
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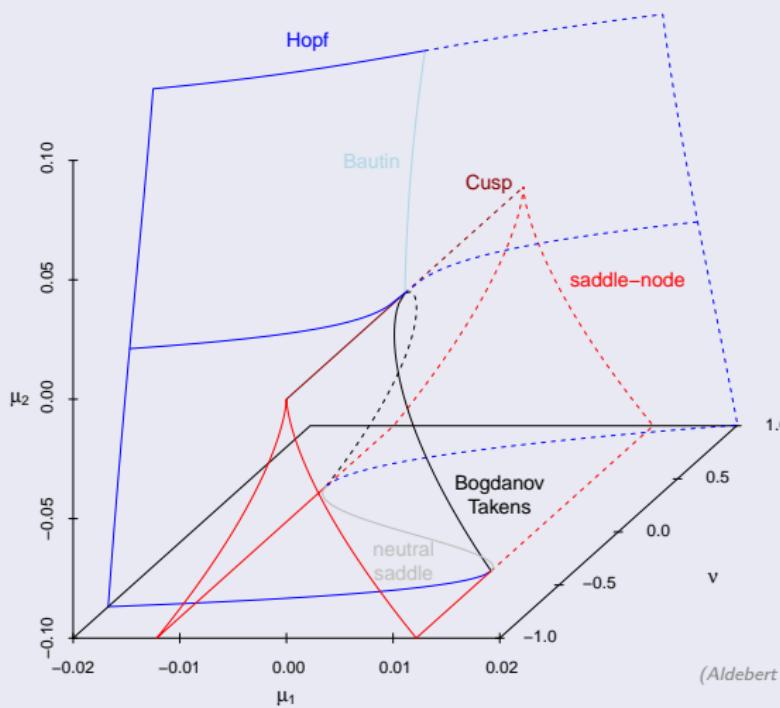
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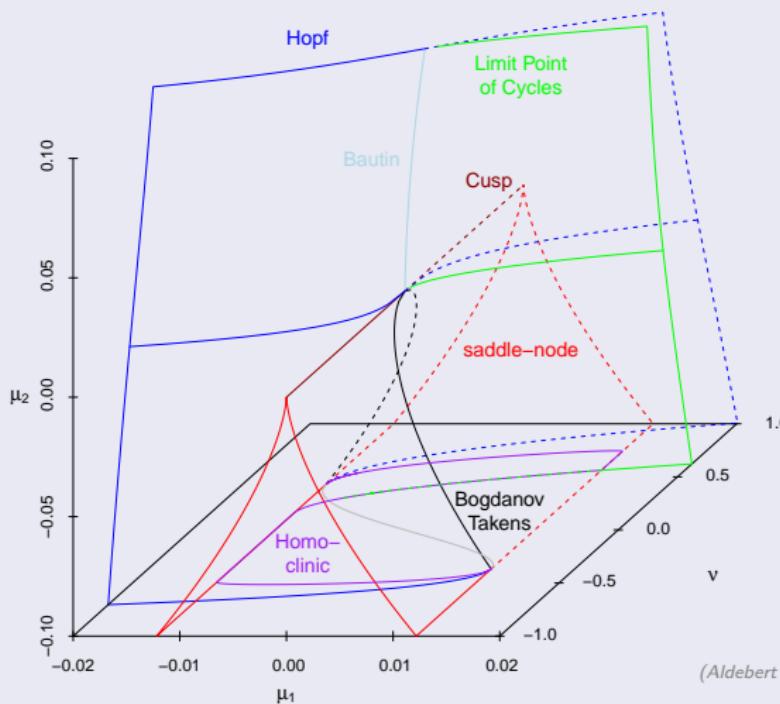
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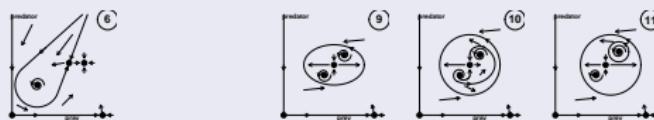
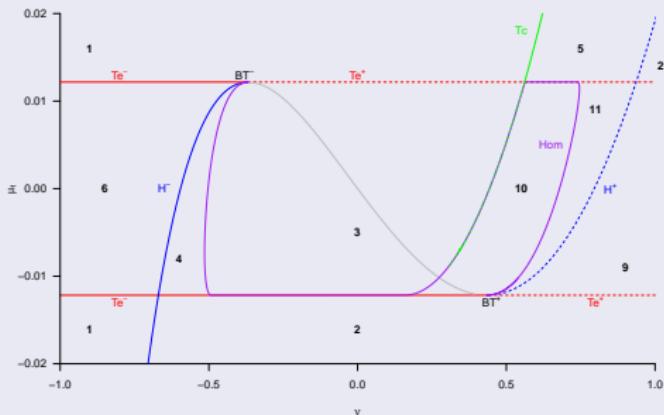
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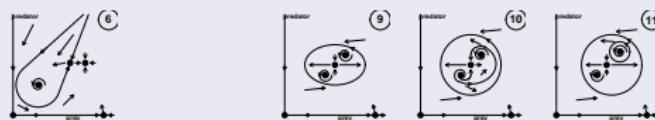
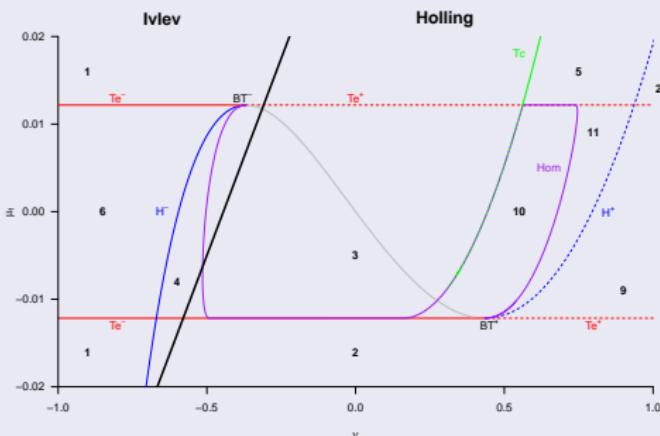
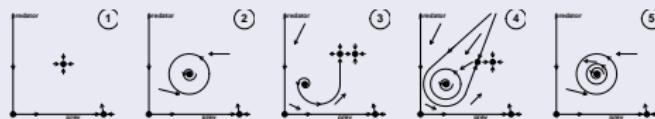
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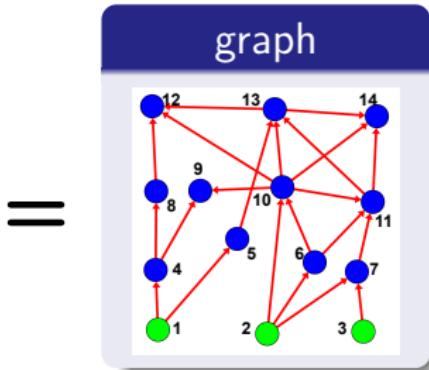
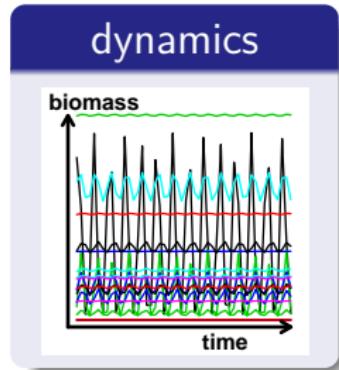
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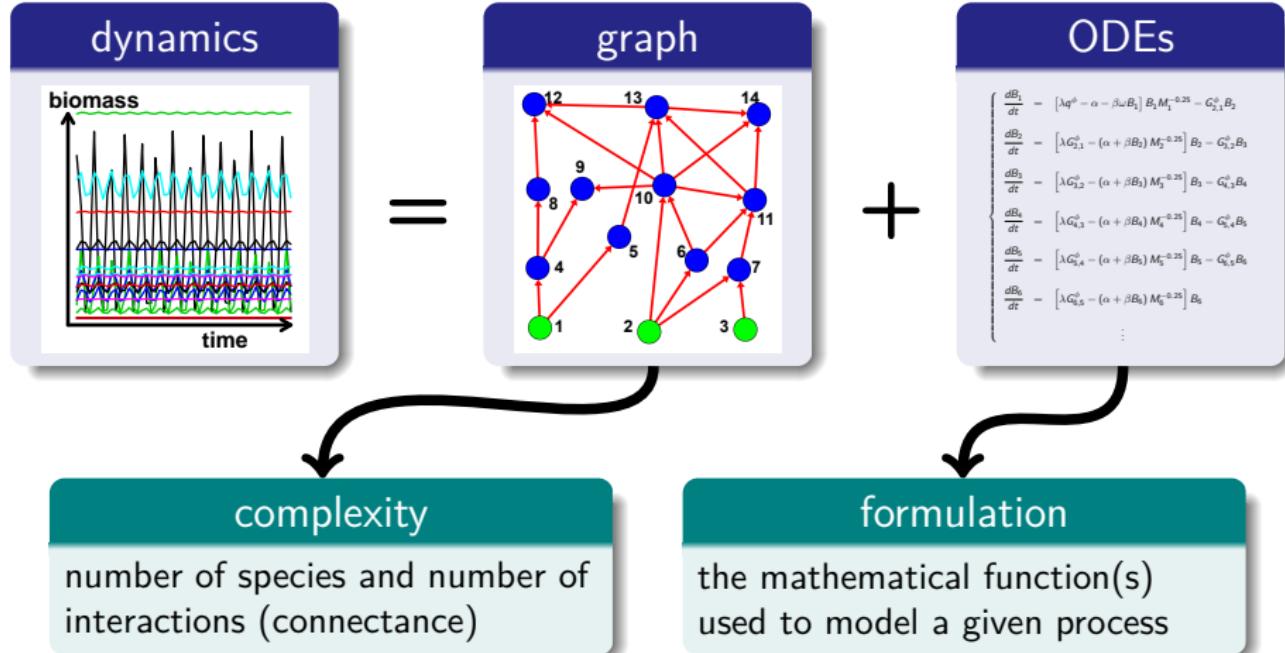
Structural sensitivity vs. complexity-stability



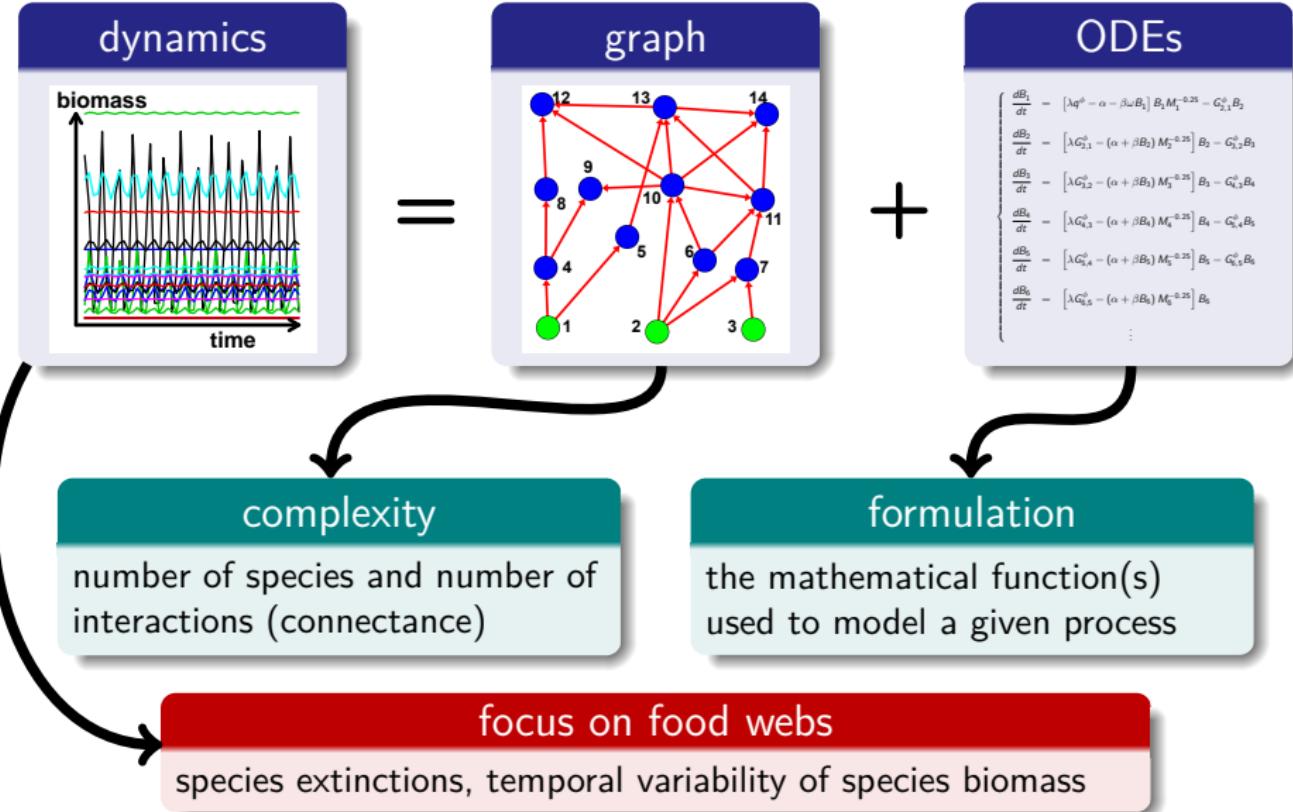
ODEs

$$\begin{cases}
 \frac{dB_1}{dt} = [\lambda q^0 - \alpha - \beta \omega B_1] B_1 M_1^{-0.25} - G_{2,1}^0 B_2 \\
 \frac{dB_2}{dt} = [\lambda G_{2,1}^0 - (\alpha + \beta B_2)] B_2 - G_{1,2}^0 B_1 \\
 \frac{dB_3}{dt} = [\lambda G_{1,2}^0 - (\alpha + \beta B_3)] B_3 - G_{4,3}^0 B_4 \\
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 \frac{dB_5}{dt} = [\lambda G_{4,5}^0 - (\alpha + \beta B_5)] B_5 - G_{5,4}^0 B_4 \\
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 \frac{dB_7}{dt} = [\lambda G_{6,7}^0 - (\alpha + \beta B_7)] B_7 - G_{7,6}^0 B_6 \\
 \vdots
 \end{cases}$$

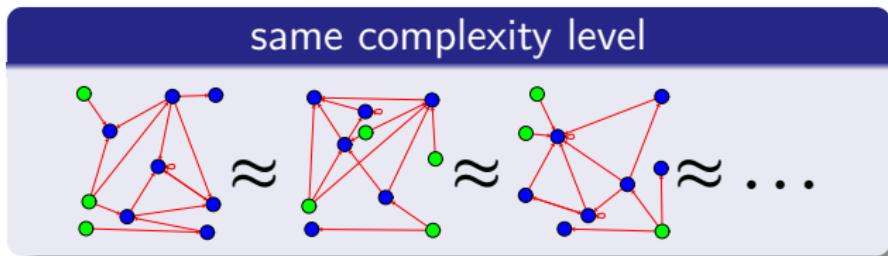
Structural sensitivity vs. complexity-stability



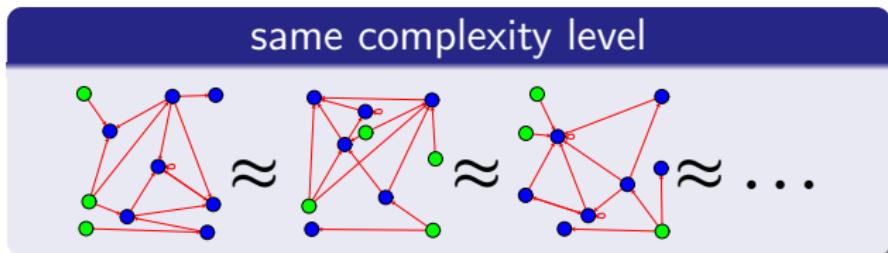
Structural sensitivity vs. complexity-stability



A statistical approach



A statistical approach



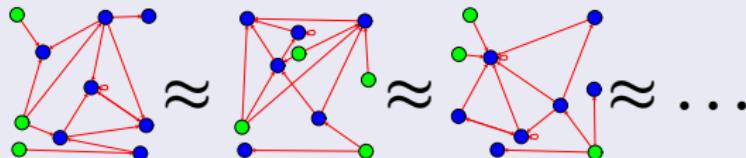
millions of “realistic” food webs



- randomly built using the niche model → structural properties that are empirically consistent (*Williams & Martinez, 2000, 2004; Cattin et al., 2004; Allesina et al., 2008*)
- use the structure to build and parameterize a dynamical system

A statistical approach

same complexity level



millions of “realistic” food webs



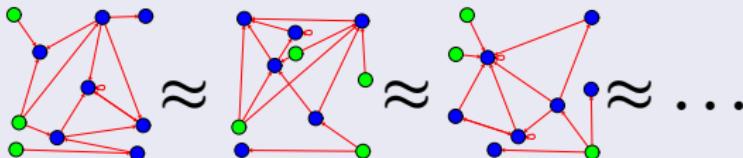
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a bio-energetic model

$$\frac{dB_i}{dt} = \underbrace{\lambda q_i^\phi B_i + \lambda \sum_{j \in R_i} G_{i,j}^\phi B_i}_{\text{gain by primary production or predation}} - \underbrace{\sum_{j \in C_i} G_{j,i}^\phi B_j}_{\text{losses by predation}} - \underbrace{\alpha_i B_i - \beta_i B_i^2}_{\text{mortality and competition}}$$

A statistical approach

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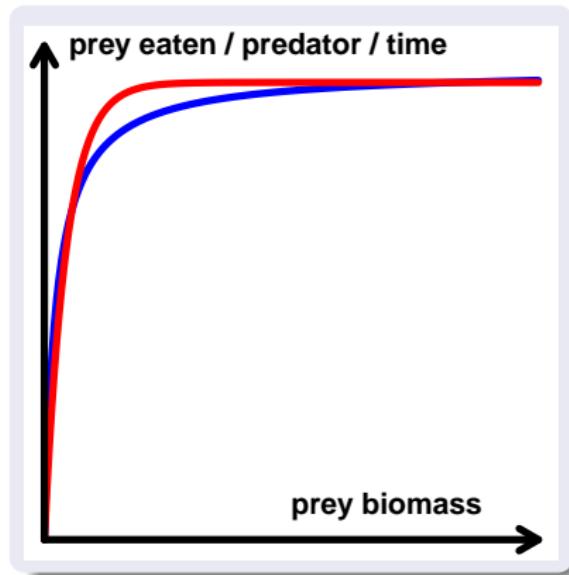


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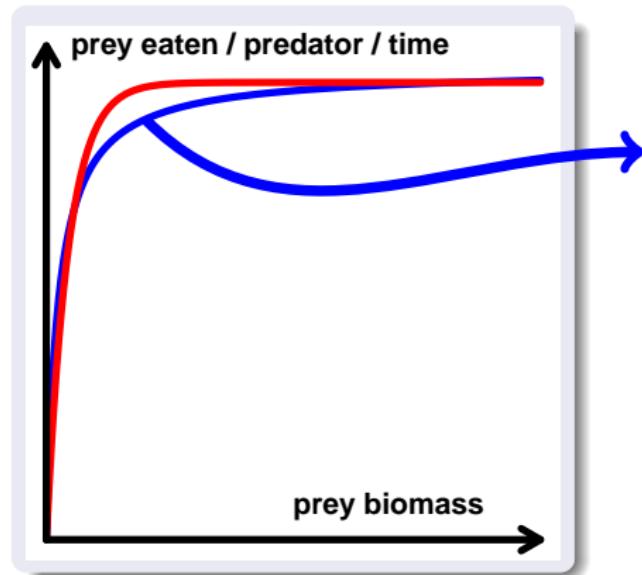
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Functional response (type II)



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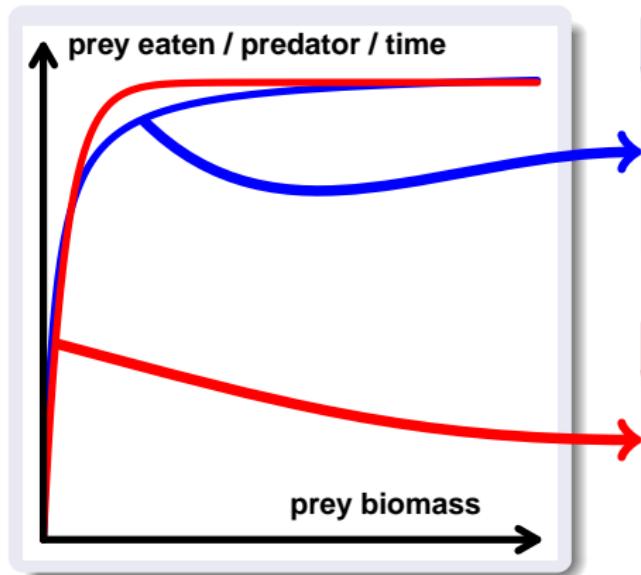
Holling (1959, 1965)

handling time

$$G_{i,j}^H = \frac{a_i^H f_{i,j} B_j}{1 + h_i^H a_i^H T_i}$$

$$T_i = \sum_{j \in R_i} f_{i,j} B_j$$

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Holling (1959, 1965)

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Ivlev (1955)

digestion

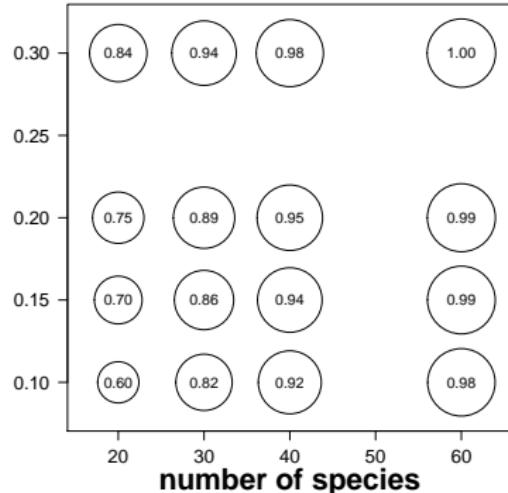
$$G_{i,j}^I = \frac{1}{h_i^I} \left(1 - e^{-h_i^I a_i^I T_i}\right) \frac{f_{i,j} B_j}{T_i}$$

$$T_i = \sum_{j \in R_i} f_{i,j} B_j$$

Fraction of food webs with extinction(s)

Holling

connectance

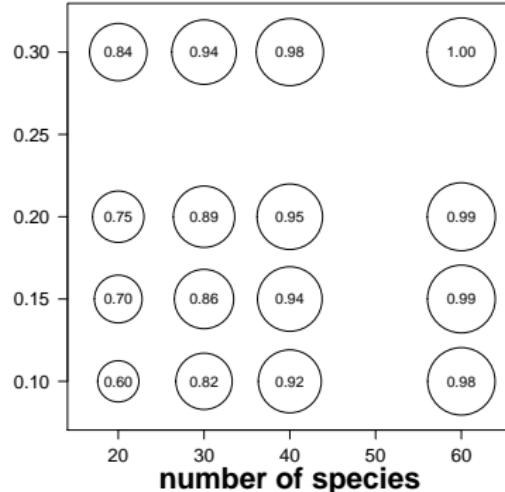


(modified from Aldebert et al., subm. rev. b)

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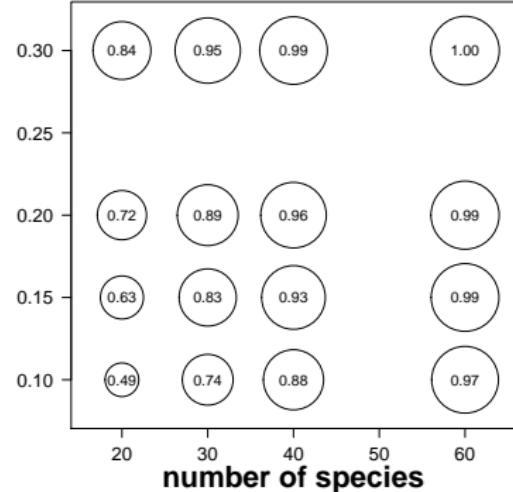
connectance



\approx

Ivlev

connectance

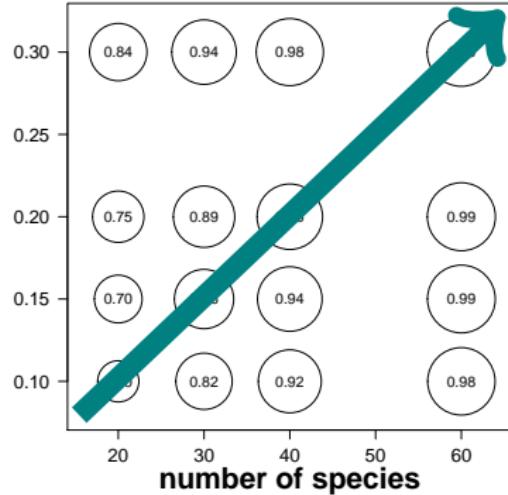


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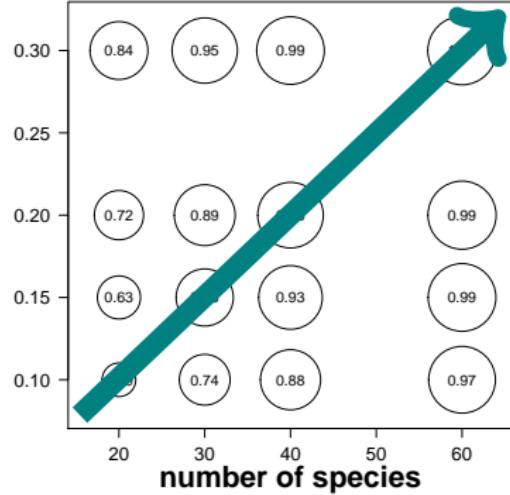
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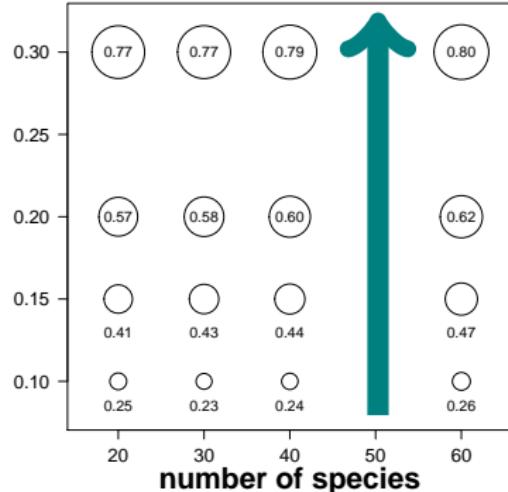
predation fluxes of similar intensity between functional responses

(modified from Aldebert et al., subm. rev. b)

Fraction of persistent food webs which reach an equilibrium

Holling

connectance

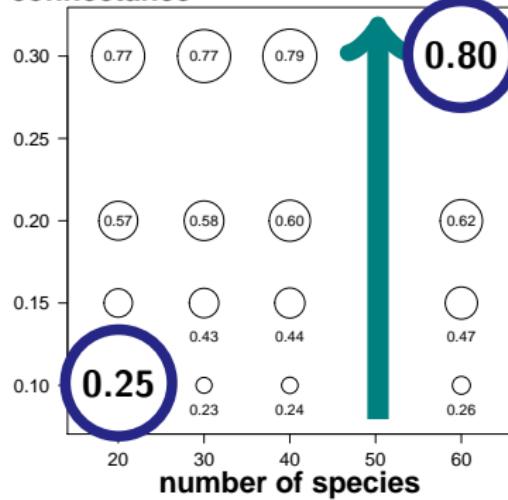


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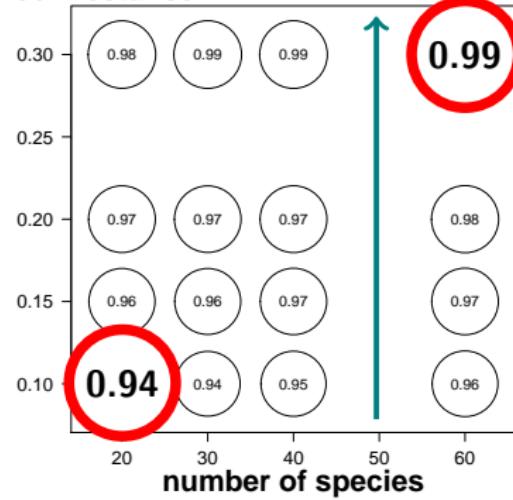
connectance



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Ivlev

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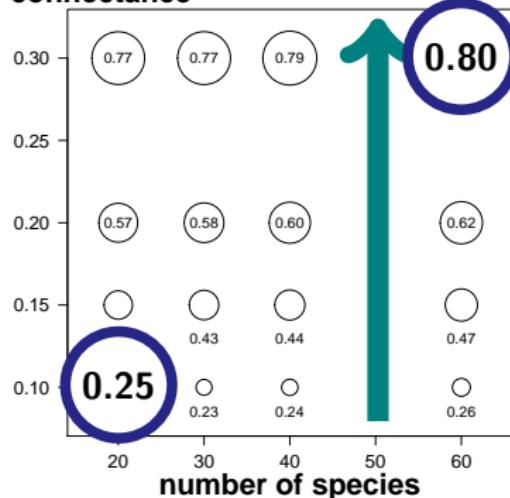


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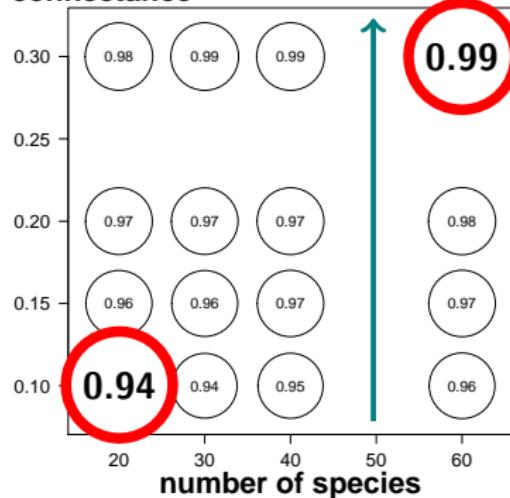
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\neq

!!! formulation > complexity !!!

(modified from Aldebert et al., subm. rev. b)

Is it a specific case ?

- functional response impact is unaffected by changes in model assumptions (primary production, cannibalism) and measure of persistence/variability
- no way to fit functional responses in order to obtain the same dynamics

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	% with extinctions	% → equilibrium
number of species	0.76	NS
connectance	0.37	0.36
maximum slope	0.12	-0.22
maximum rate	0.07	0.07
functional response	-0.17	0.82

TABLE : Correlation coefficients

($\approx 6.10^7$ food webs were simulated to obtain this table)

computational effort : 3 years.processors

Bifurcations in a predator-prey system

same model \Rightarrow the simplest food web

$$\begin{cases} \dot{B_{prey}}(t') = [\lambda q^\xi - \alpha - \beta \omega B_{prey}] B_{prey} - G^\xi M_{pred}^{0.25} B_{pred} (M_{pred}/M_{prey})^{-0.25} \\ \dot{B_{pred}}(t') = [\lambda G^\xi M_{pred}^{0.25} - \alpha - \beta B_{pred}] B_{pred} (M_{pred}/M_{prey})^{-0.25} \end{cases}$$

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- Rosenzweig-MacArthur's model with predator competition

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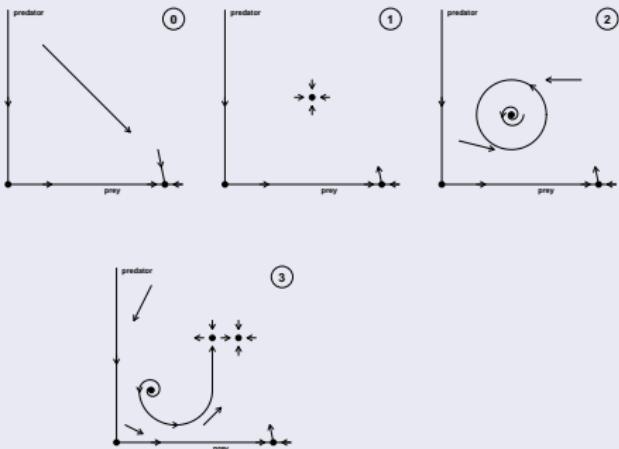
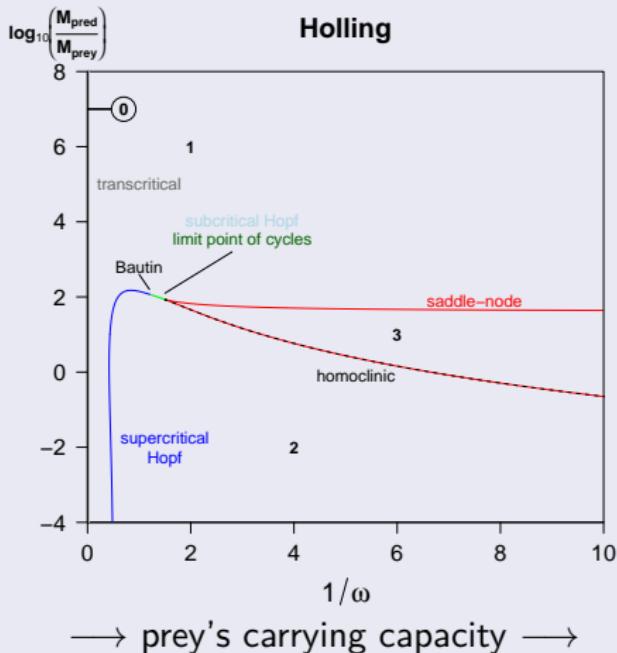
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- has up to 3 positive equilibria and 2 limit cycles (12 phase portraits)
- exhibits all possible codimension 2 bifurcations in planar systems

Bifurcation diagram with Holling

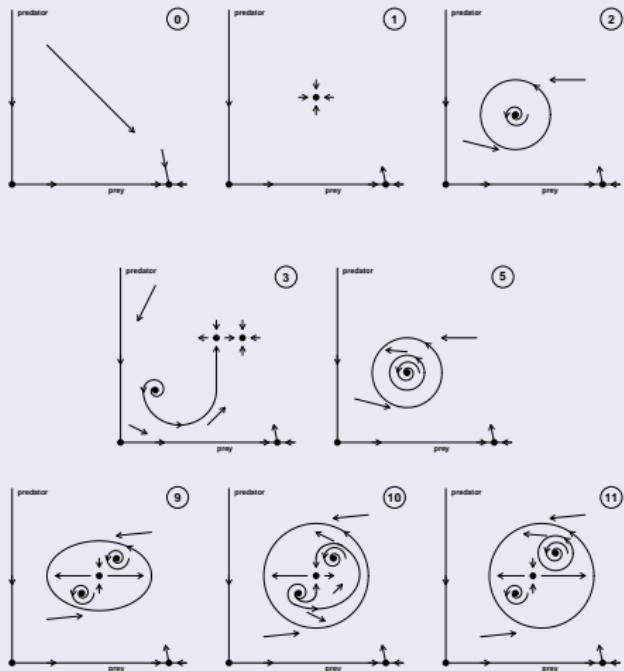
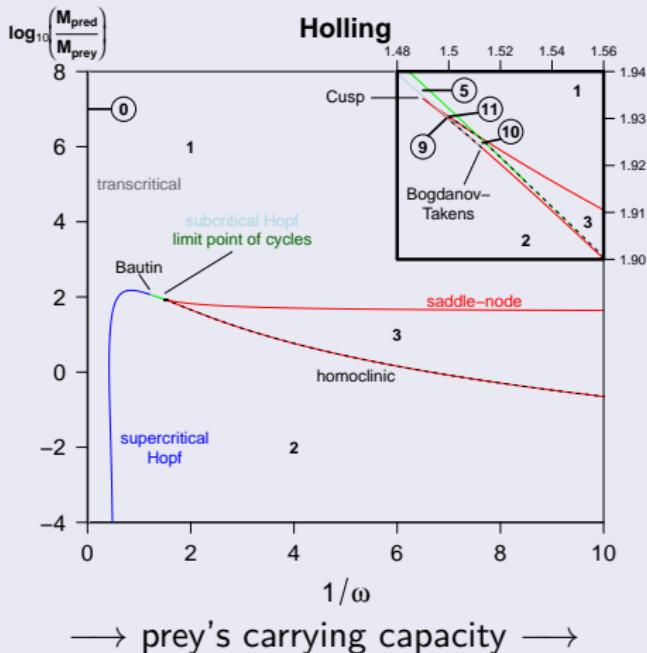
body mass ratio



(modified from Aldebert et al., in press)

Bifurcation diagram with Holling

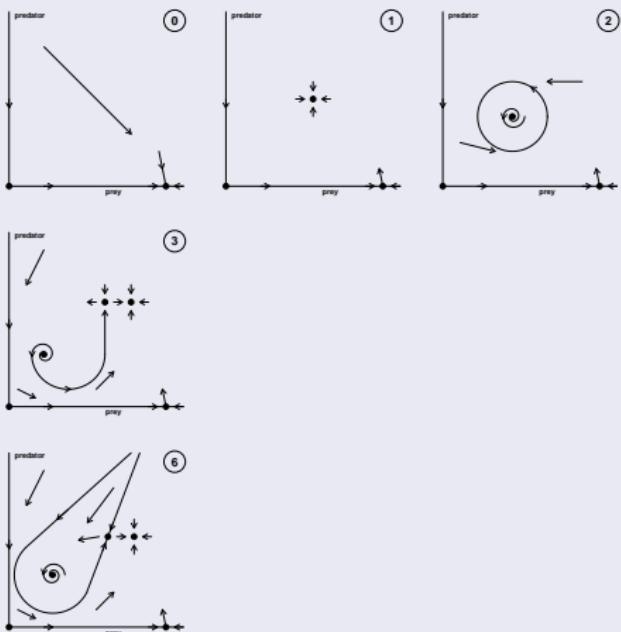
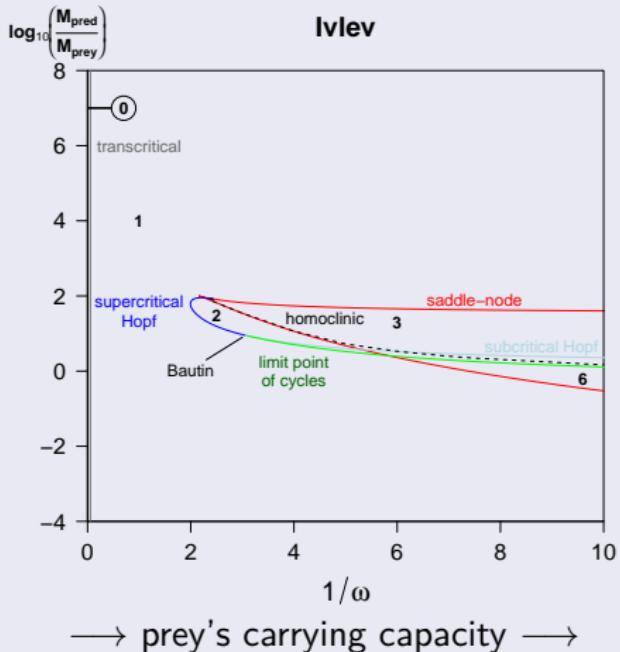
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Bifurcation diagram with Ivlev

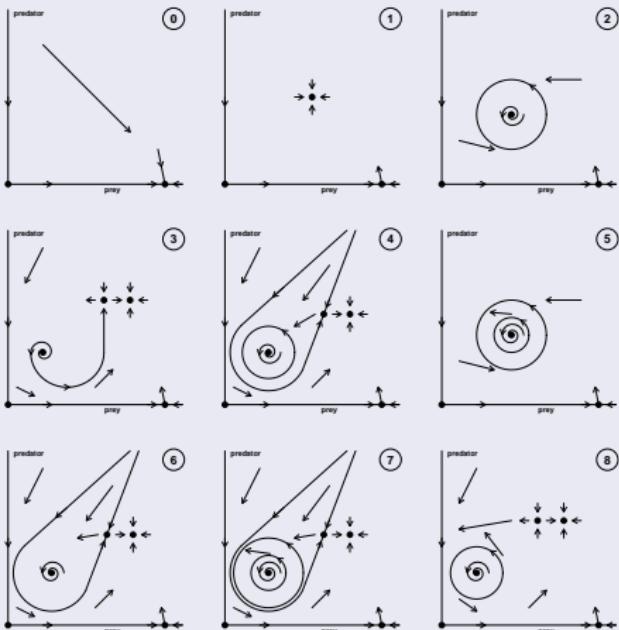
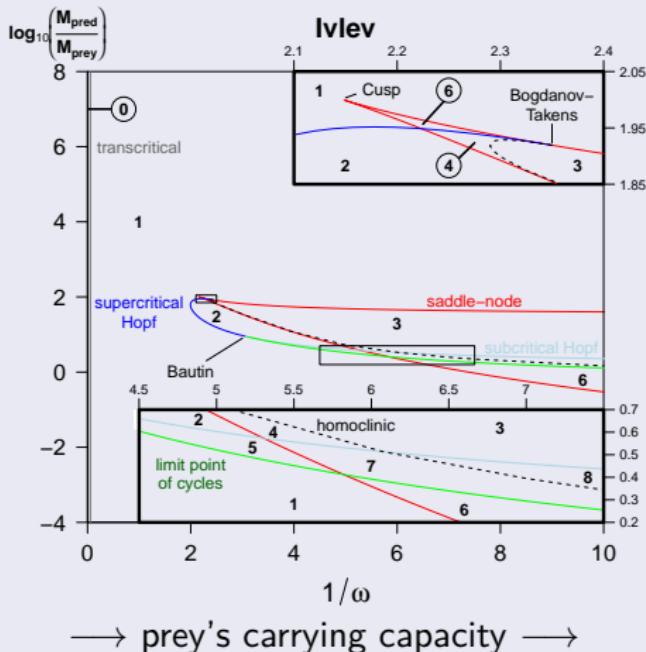
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(modified from Aldebert et al., in press)

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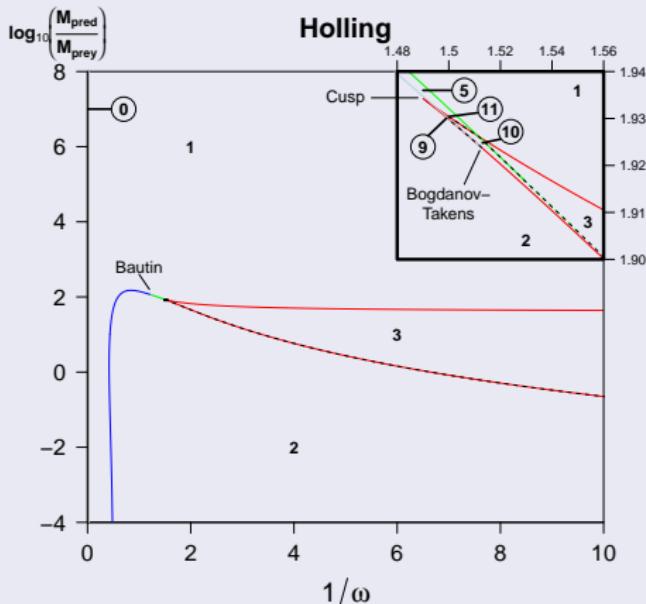
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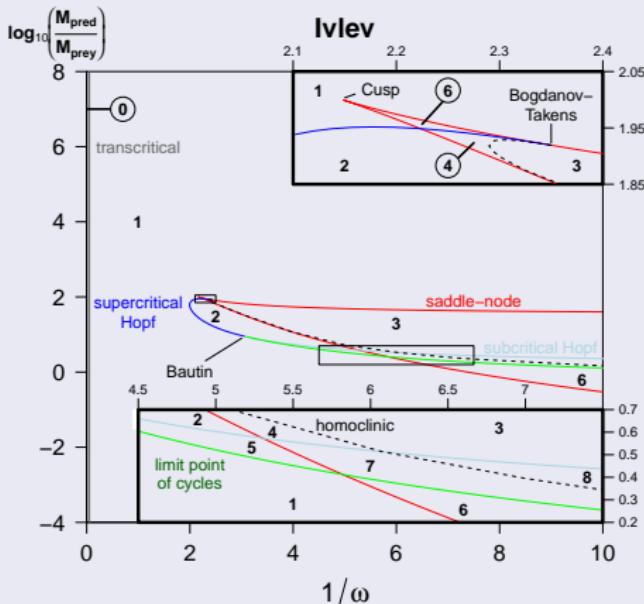
Bifurcation diagram : Holling vs. Ivlev

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→ prey's carrying capacity →

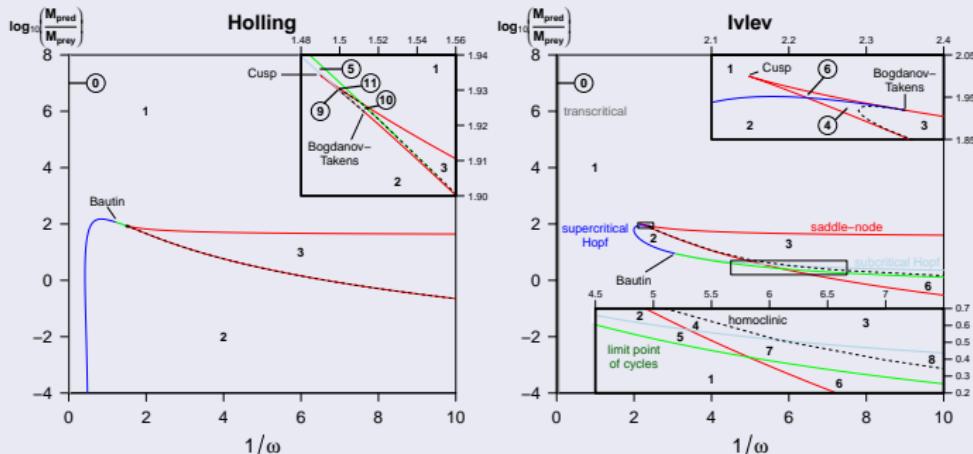
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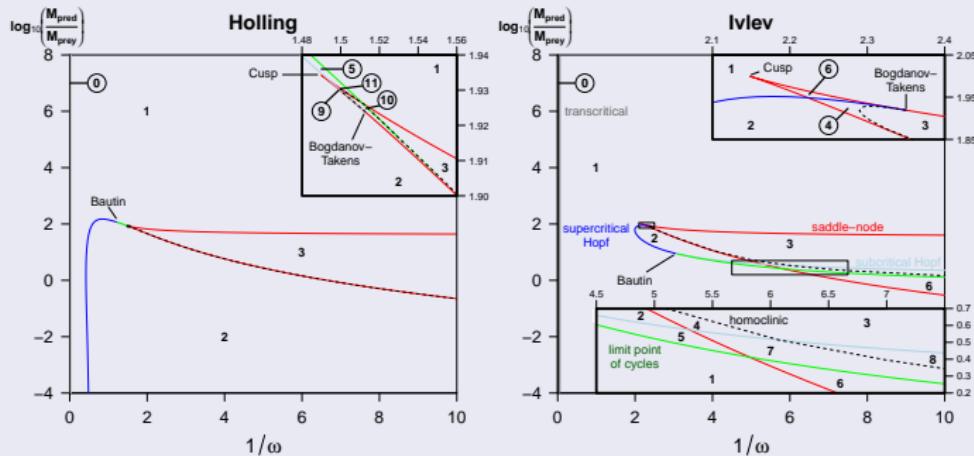


main differences

- stable equilibrium vs stable limit cycle : 26.0 % – 49.4 %
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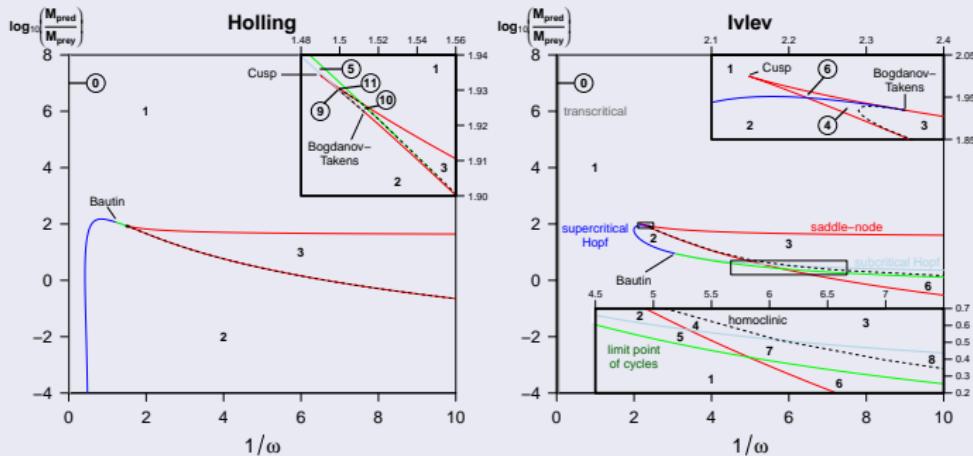


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Generalized modelling : predator-prey system

general idea

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- derive the Jacobian matrix

(modified from Aldebert et al., in press)

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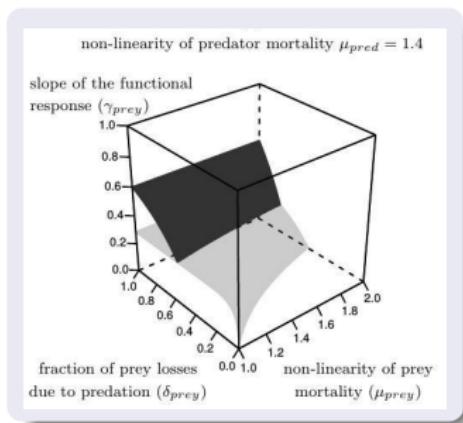
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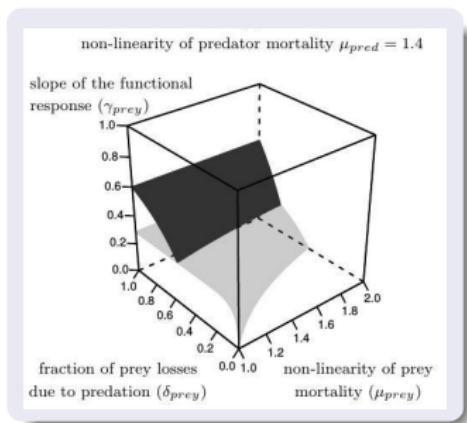


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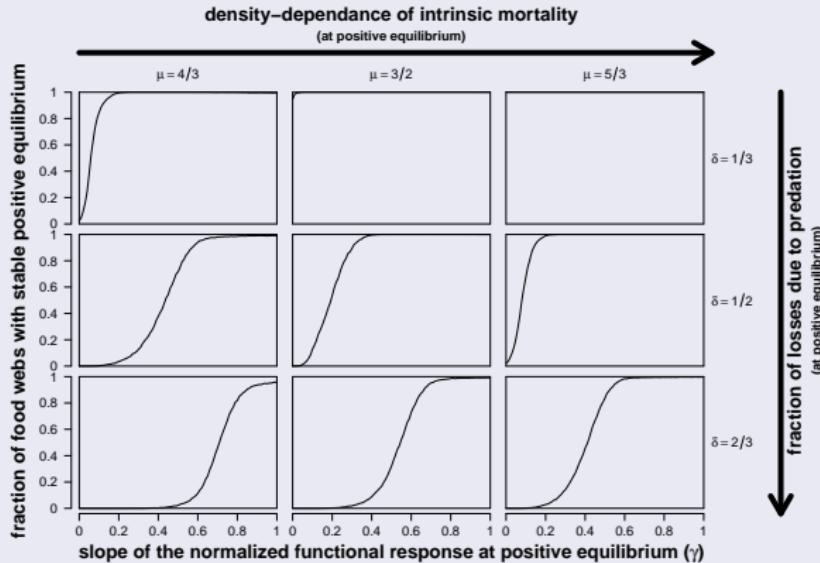


5 parameters, stabilizing factors

- high density-dependent intrinsic mortality, low losses through predation
- high slope of the functional response near equilibrium

(modified from Aldebert et al., in press)

Generalized modelling : food webs

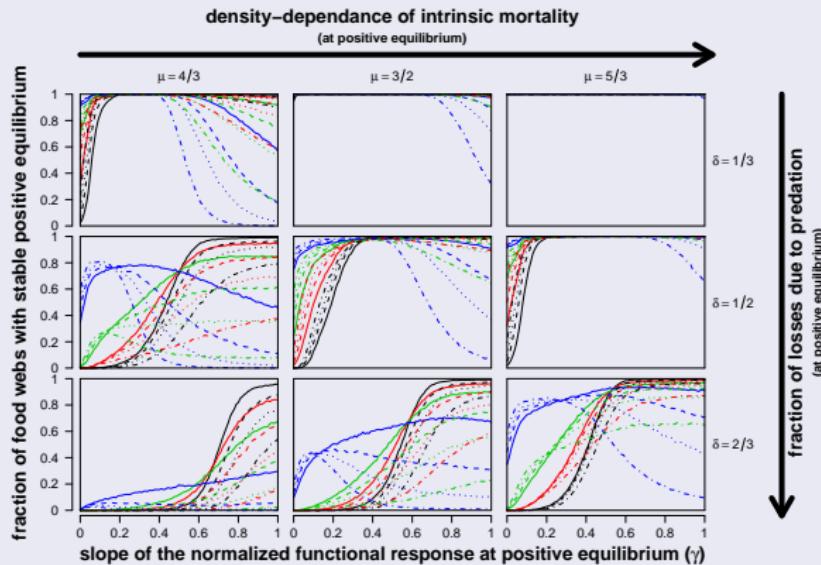


number of species :
20

connectance :
0.10

(modified from Aldebert et al., subm. rev. b)

Generalized modelling : food webs

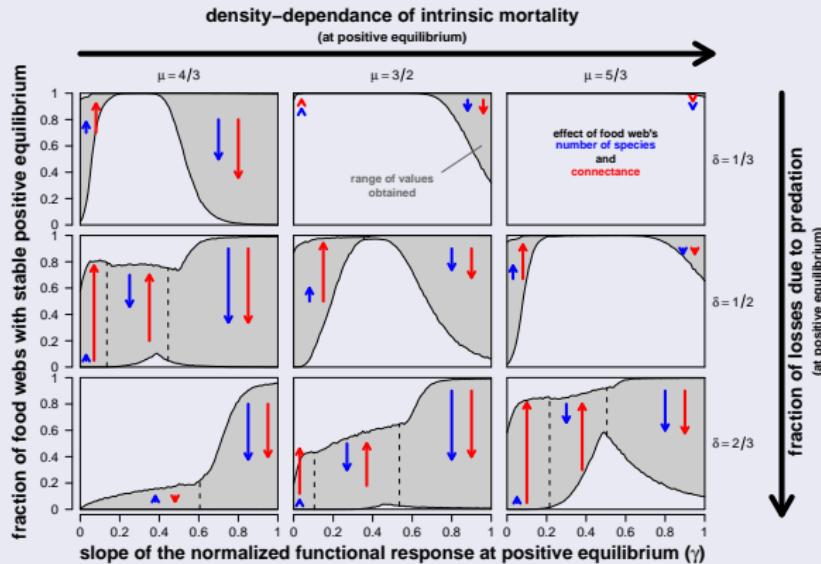


number of species :
20, 30, 40, 60
(line type)

connectance :
0.10, 0.15, 0.20, 0.30
(line color)

(modified from Aldebert et al., subm. rev. b)

Generalized modelling : food webs

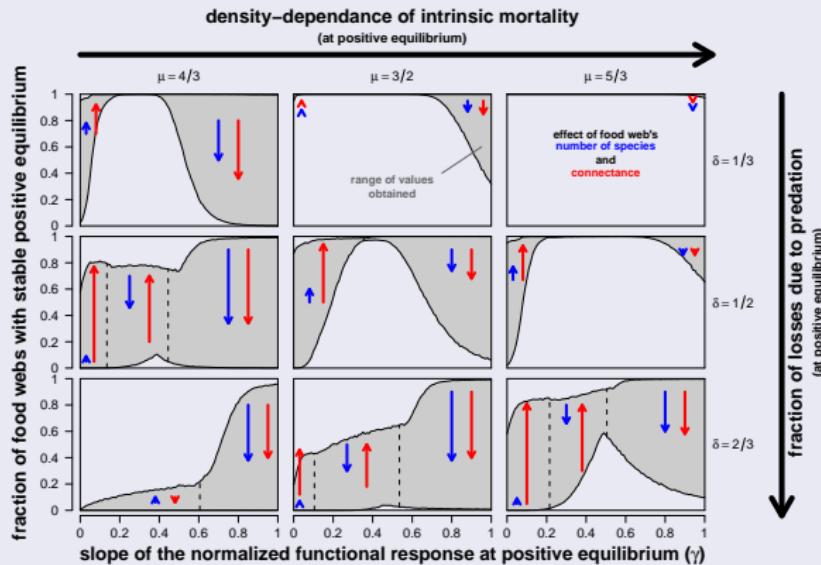


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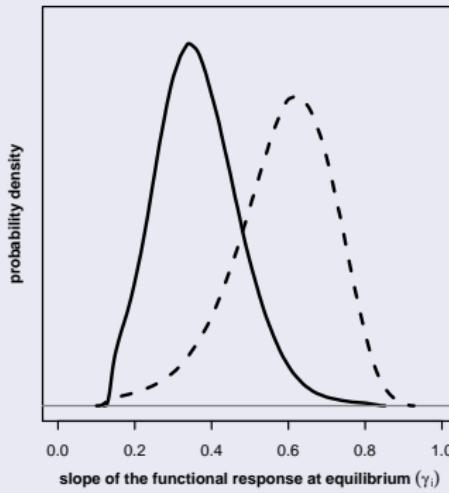
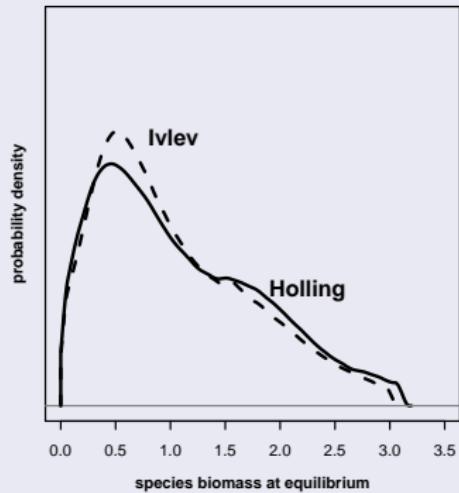
Generalized modelling : food webs



- stabilizing effect of a higher slope of the functional response
- robust to changes in model assumptions (primary production, cannibalism)

(modified from Aldebert et al., subm. rev. b)

Generalized parameters as indicators in food webs



food webs at positive equilibrium

- same species biomass and generalized parameters distributions
- except for the slope of the functional response (higher with Ivlev)

(modified from Aldebert et al., subm. rev. b)

Structural sensitivity : ongoing works

more physiological details : DEB models

- a family of bi-trophic food chain models (explicit resource) in chemostat

(Aldebert et al., in prep)

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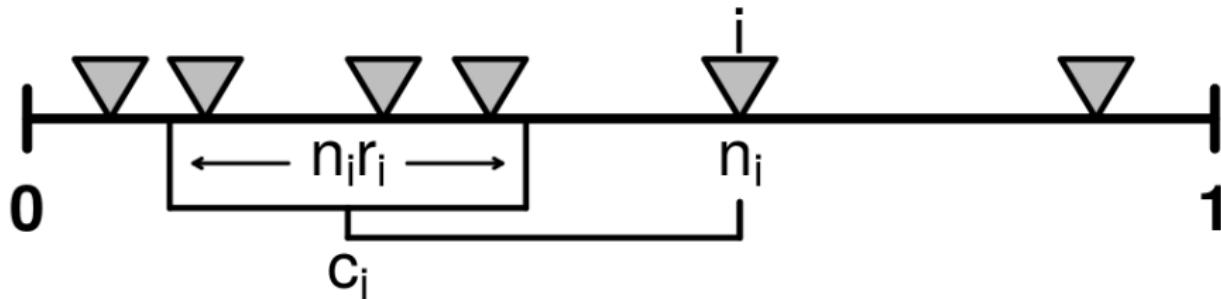
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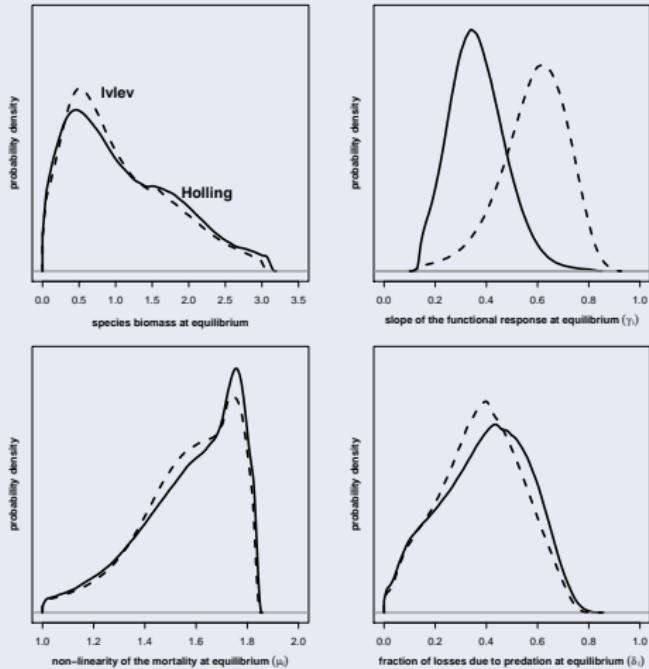
The niche model (*Williams & Martinez, 2000, 2004*)



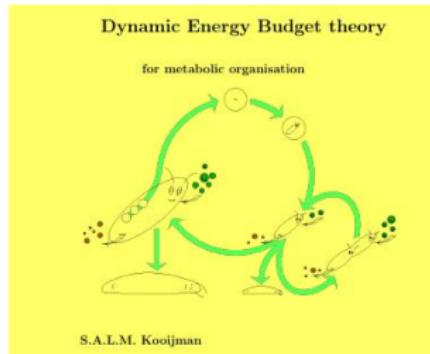
based on the principle of ecological niche (*Hutchinson, 1957*)

- the segment $[0, 1]$ summarizes the ∞ -dimensional niche space
- the niche indice n_i summarizes species i 's ecological niche
- the relative width of species i 's feeding range r_i depends on connectance
- c_i is the center of species i 's feeding range
- species i feeds on all species who belong to its feeding range $[c_i \pm \frac{n_i r_i}{2}]$
- species with an empty feeding range are defined as primary producers
- more complex models (e.g. with \pm discontinuous feeding range) are based on the niche model

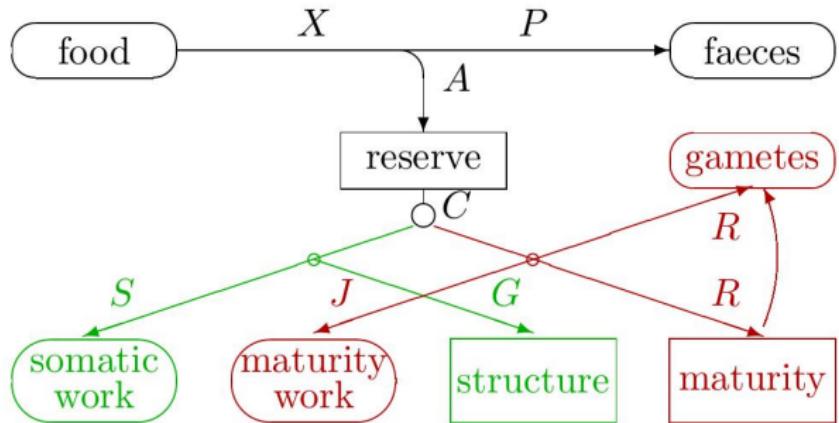
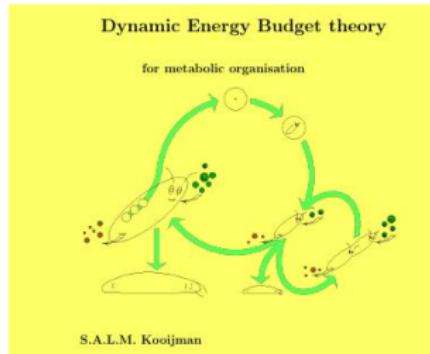
Generalized modelling : food webs



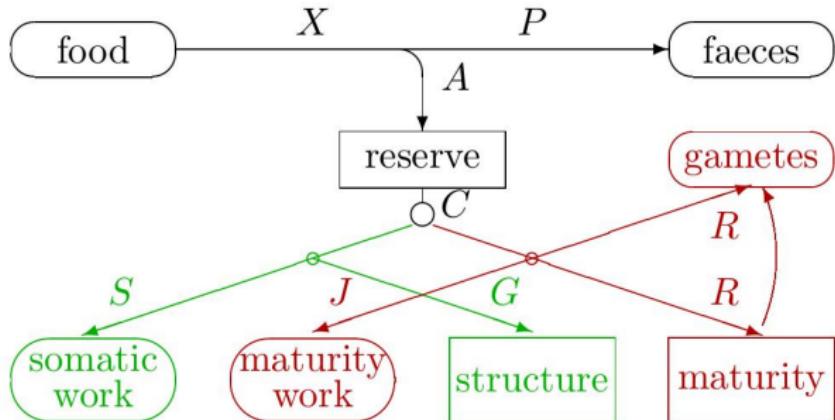
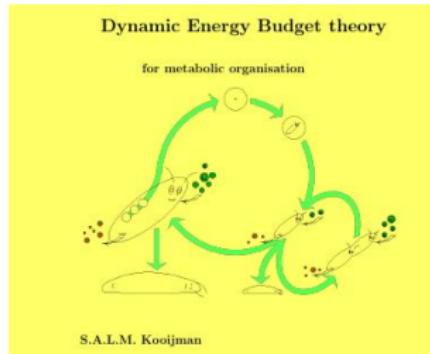
Dynamic Energy Budgets theory



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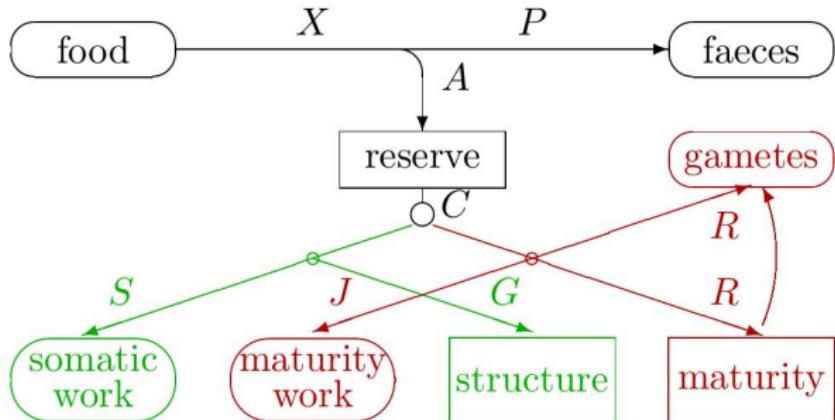
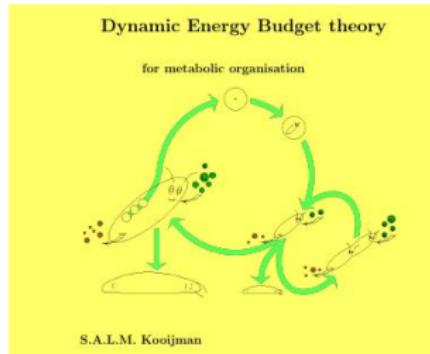


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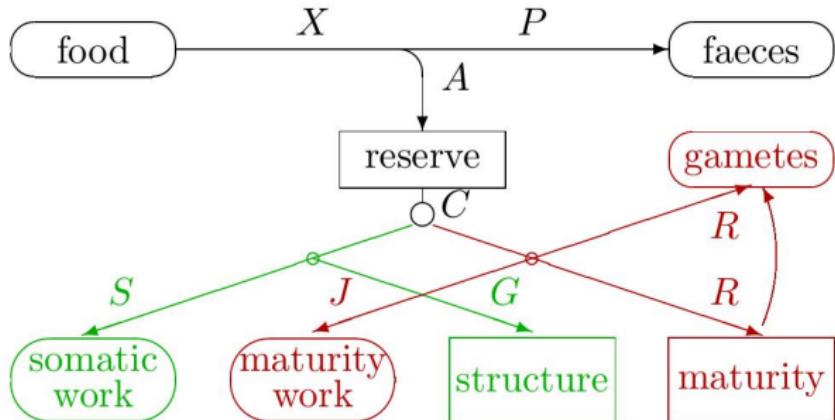
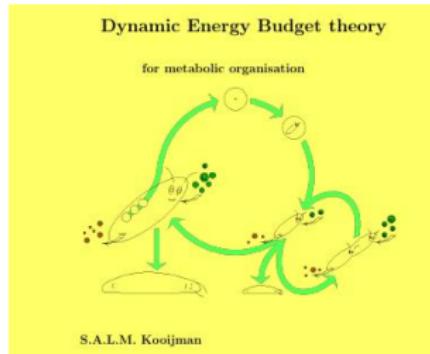
- focus on the individual, based on mechanistic assumptions on metabolism

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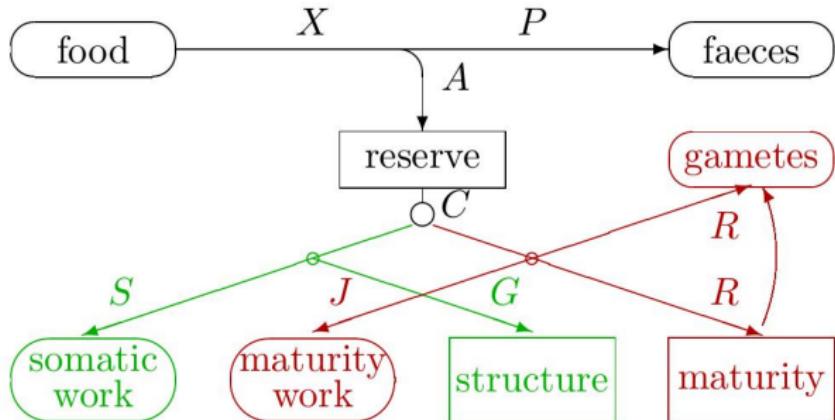
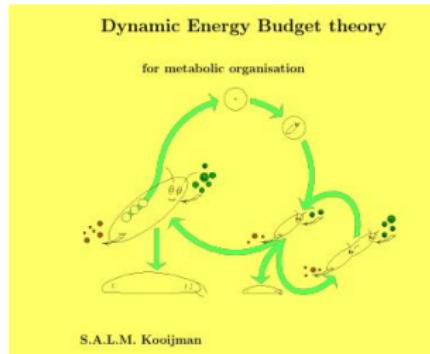
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- unicellular organisms : upscaling to population dynamics is easy

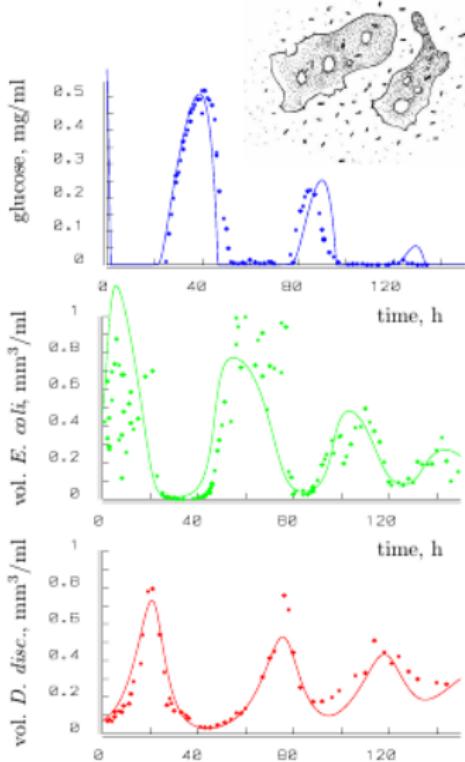
DEB model for a bitrophic food chain in chemostat

(data from Dent *et al.*, 1976; model from Kooi & Kooijman, 1994) (fig 9.15, p 358)

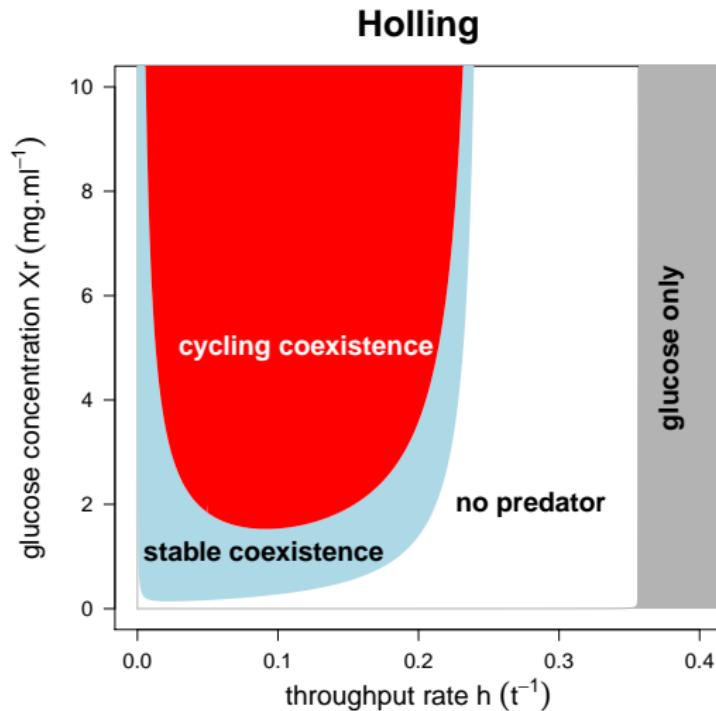
$X_0(0)$	0.433	mg ml^{-1}
$X_1(0)$	0.361	$\text{mm}^3 \text{ml}^{-1}$
$e_1(0)$	1	-
K_1	0.40	K_2
g_1	0.86	0.18
\dot{k}_M^1	0.008	$\mu\text{g ml}^{-1}, \text{mm}^3 \text{ml}^{-1}$
\dot{k}_E^1	0.67	-
j_{XAm}^1	0.65	\dot{k}_E^2
		0.16
		h^{-1}
		\dot{k}_E^2
		2.05
		h^{-1}
		j_{XAm}^2
		0.26
		$\frac{\text{mg}}{\text{mm}^3 \text{h}}, \text{h}^{-1}$

$$\begin{aligned}\frac{d}{dt}e_1 &= \dot{k}_E^1(f_1 - e_1); \quad f_1 = \frac{X_0}{K_1 + X_0} \\ \frac{d}{dt}e_2 &= \dot{k}_E^2(f_2 - e_2); \quad f_2 = \frac{X_1}{K_2 + X_1} \\ \frac{d}{dt}X_0 &= \dot{h}(X_r - X_0) - f_1 j_{XAm}^1 X_1 \\ \frac{d}{dt}X_1 &= \left(\frac{\dot{k}_E^1 e_1 - \dot{k}_M^1 g_1}{e_1 + g_1} - \dot{h} \right) X_1 - f_2 j_{XAm}^2 X_2 \\ \frac{d}{dt}X_2 &= \left(\frac{\dot{k}_E^2 e_2 - \dot{k}_M^2 g_2}{e_2 + g_2} - \dot{h} \right) X_2\end{aligned}$$

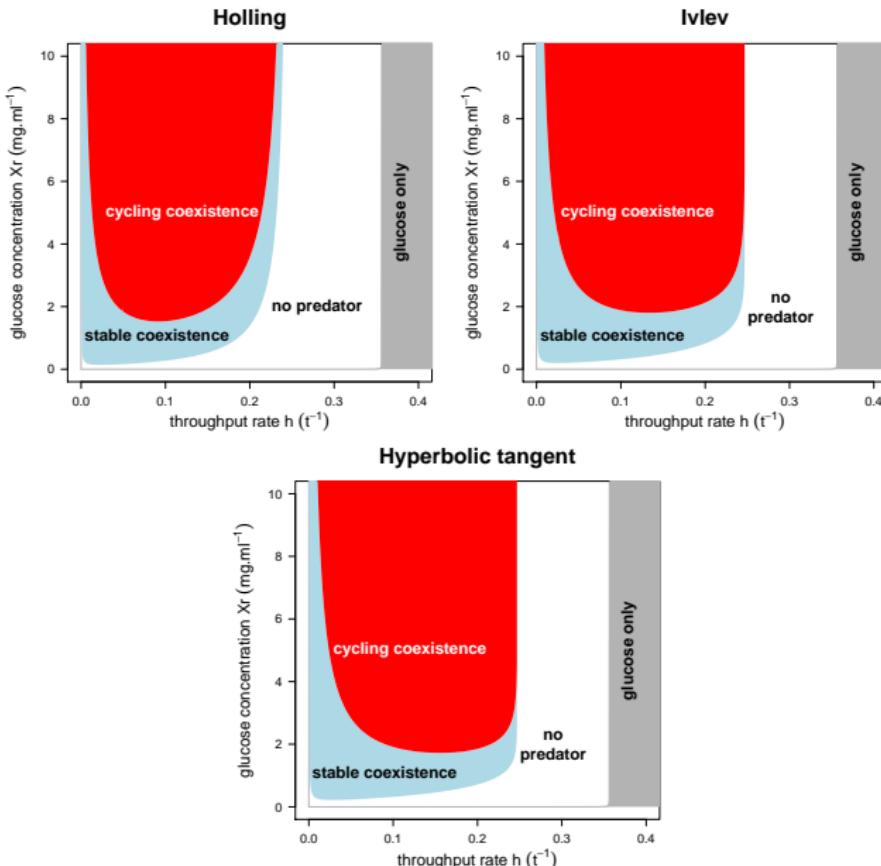
V1-morph, $\kappa \rightarrow 1$, predator feed on prey's structure, fixed environment



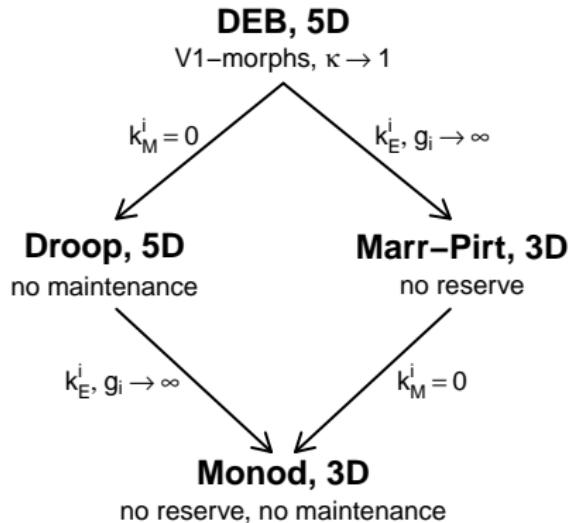
Structural sensitivity in this DEB model



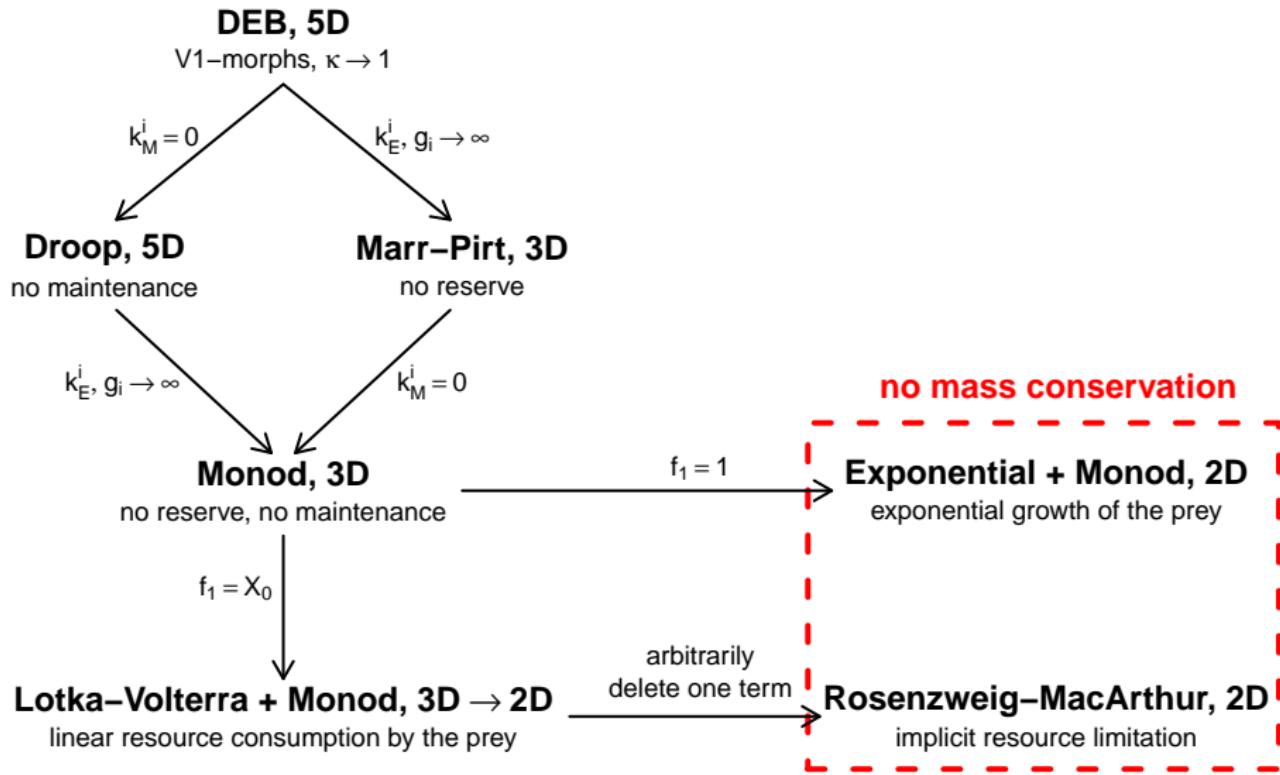
Low structural sensitivity in this DEB model



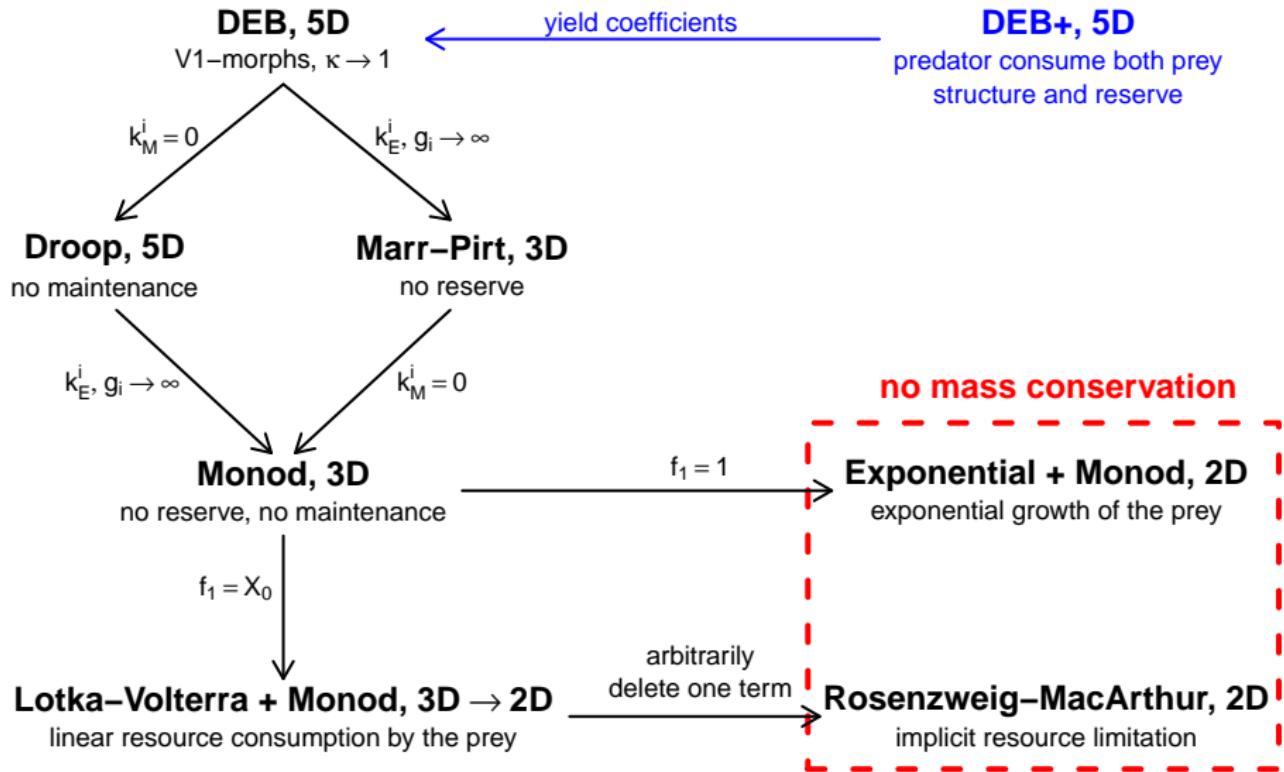
A family tree of DEB models



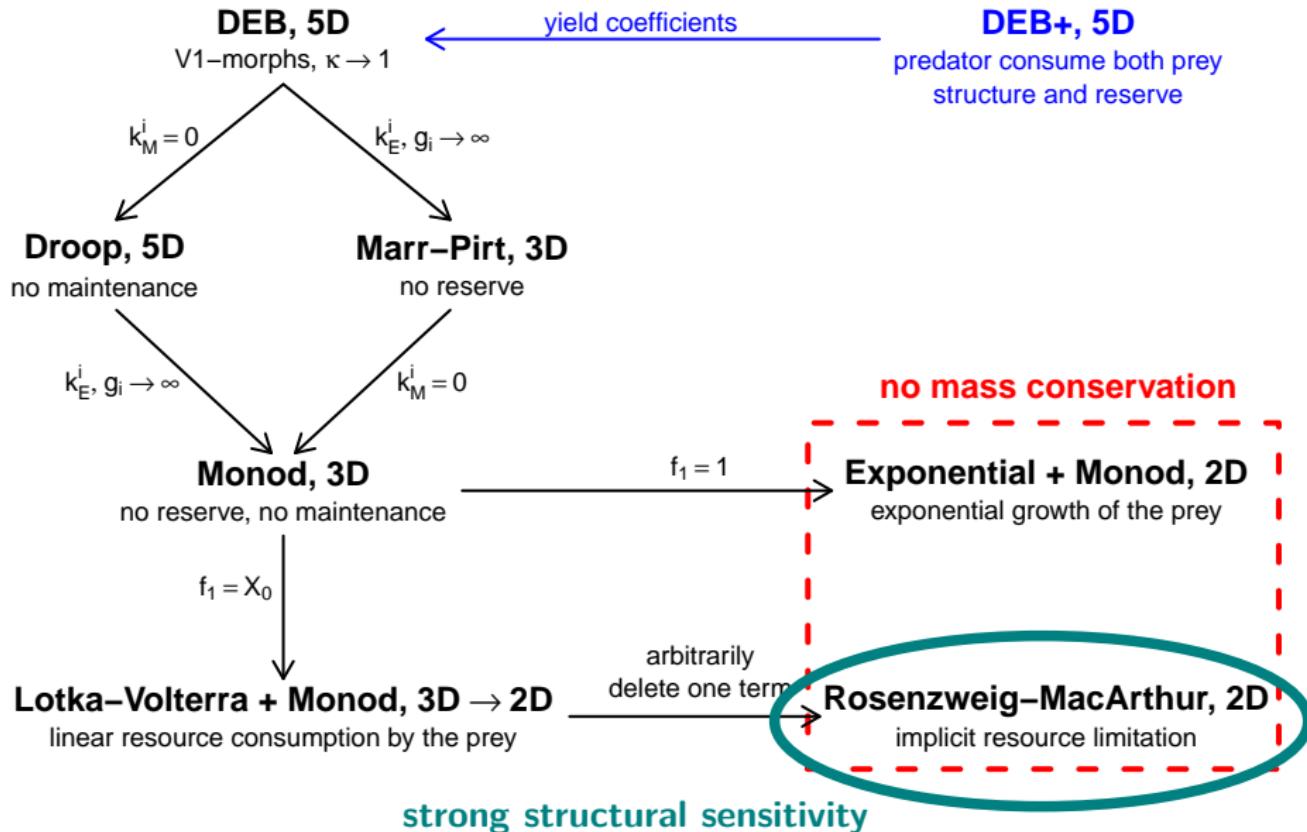
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