



Food webs



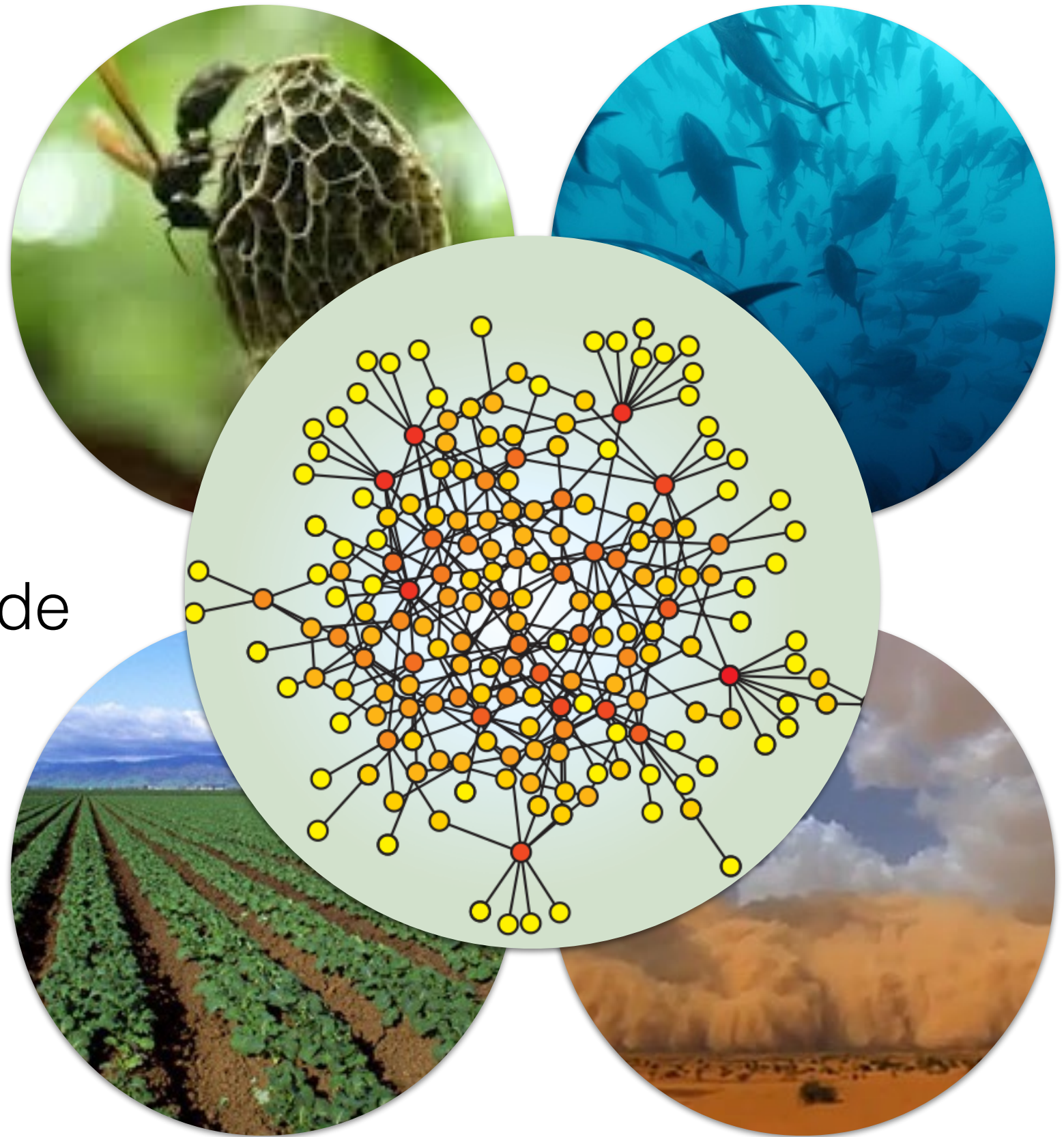
 <http://jdyeakel.github.io>

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University of California, Merced



Biological diversity
populations
species
communities
ecosystems

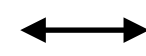
Diverse ecosystems provide
Food (e.g. fisheries)
Genetic diversity
Predictability
Beauty



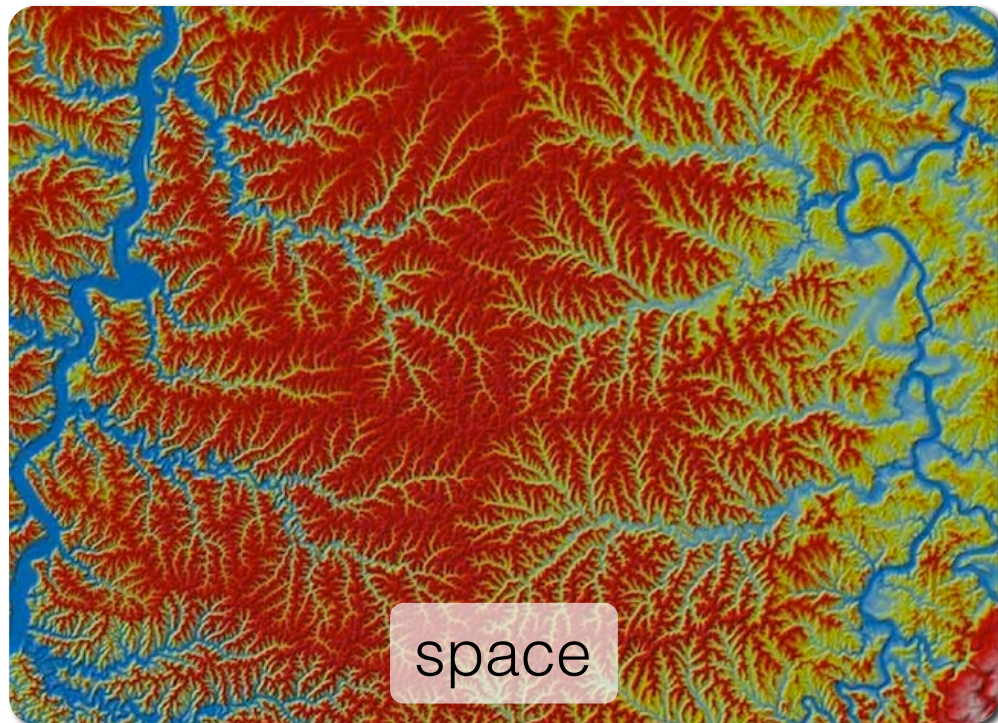
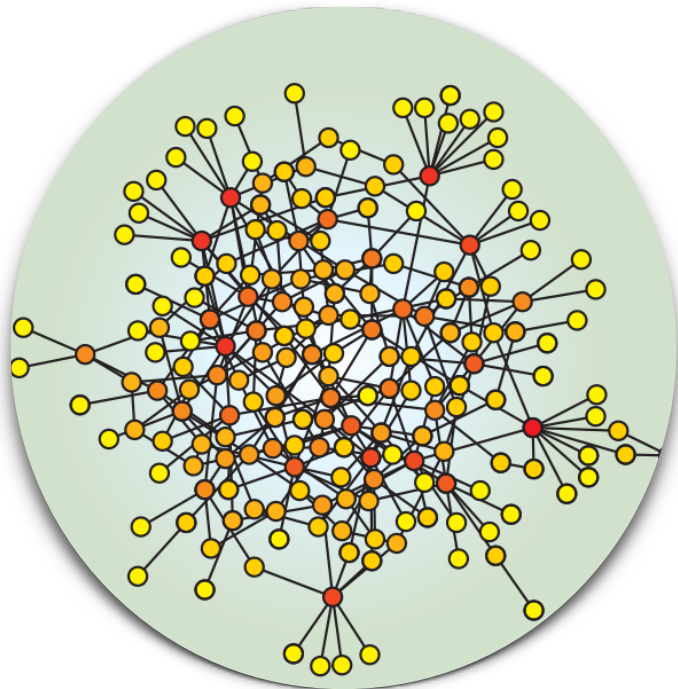
Food webs



Interactions



Behavior



space



time

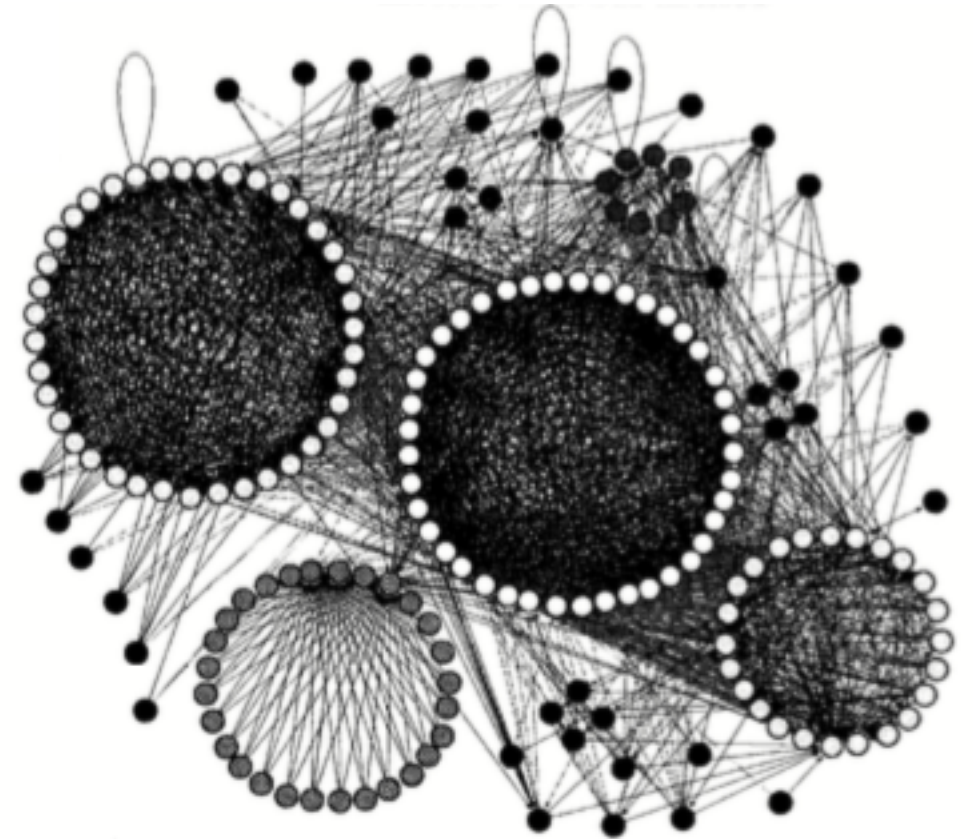
Structure of this talk:

- Empirical food web networks
 - Statistical properties

- Model food web networks
 - Structure
 - Parameterizing models from data
 - Incorporating additional biological constraints

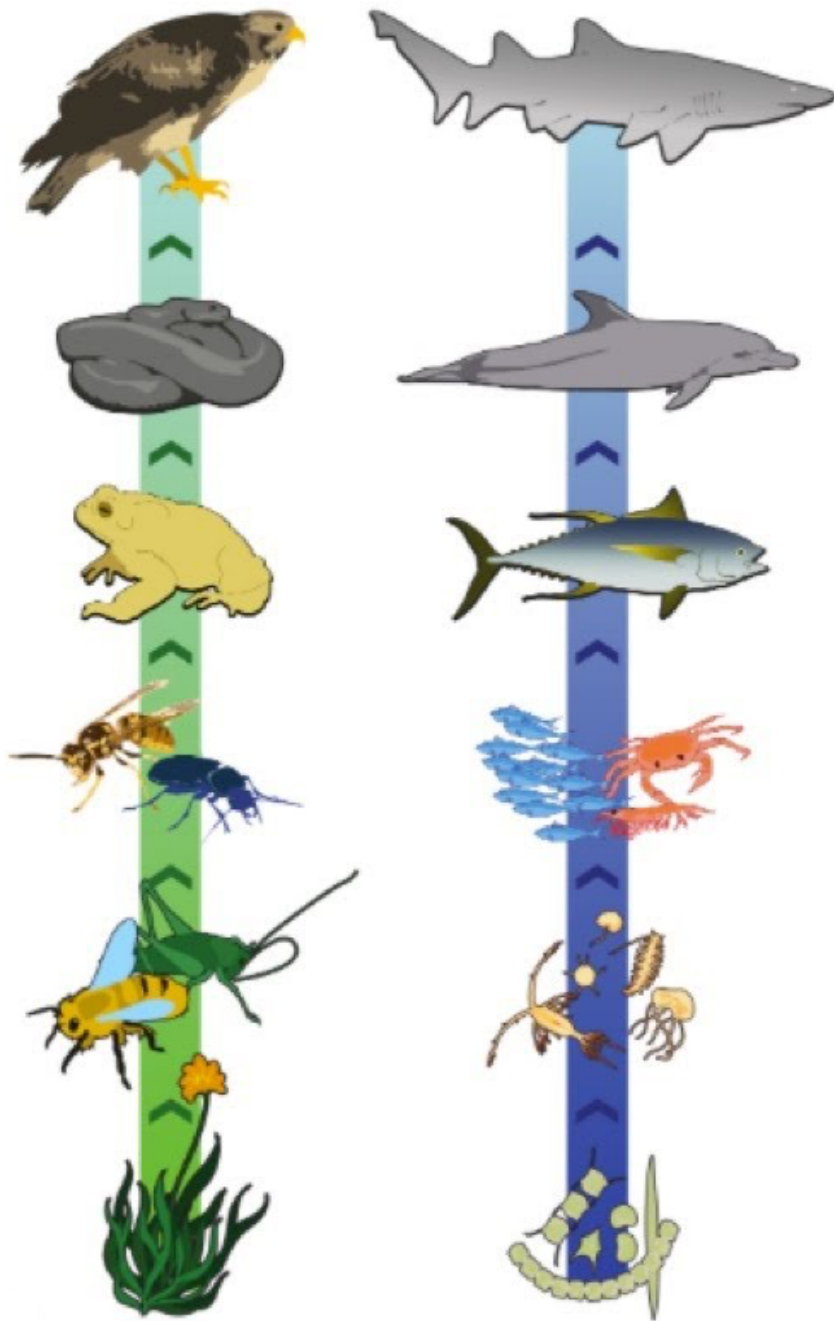
- Food web dynamics
 - Matrix models
 - Generalized models

- Food webs over time

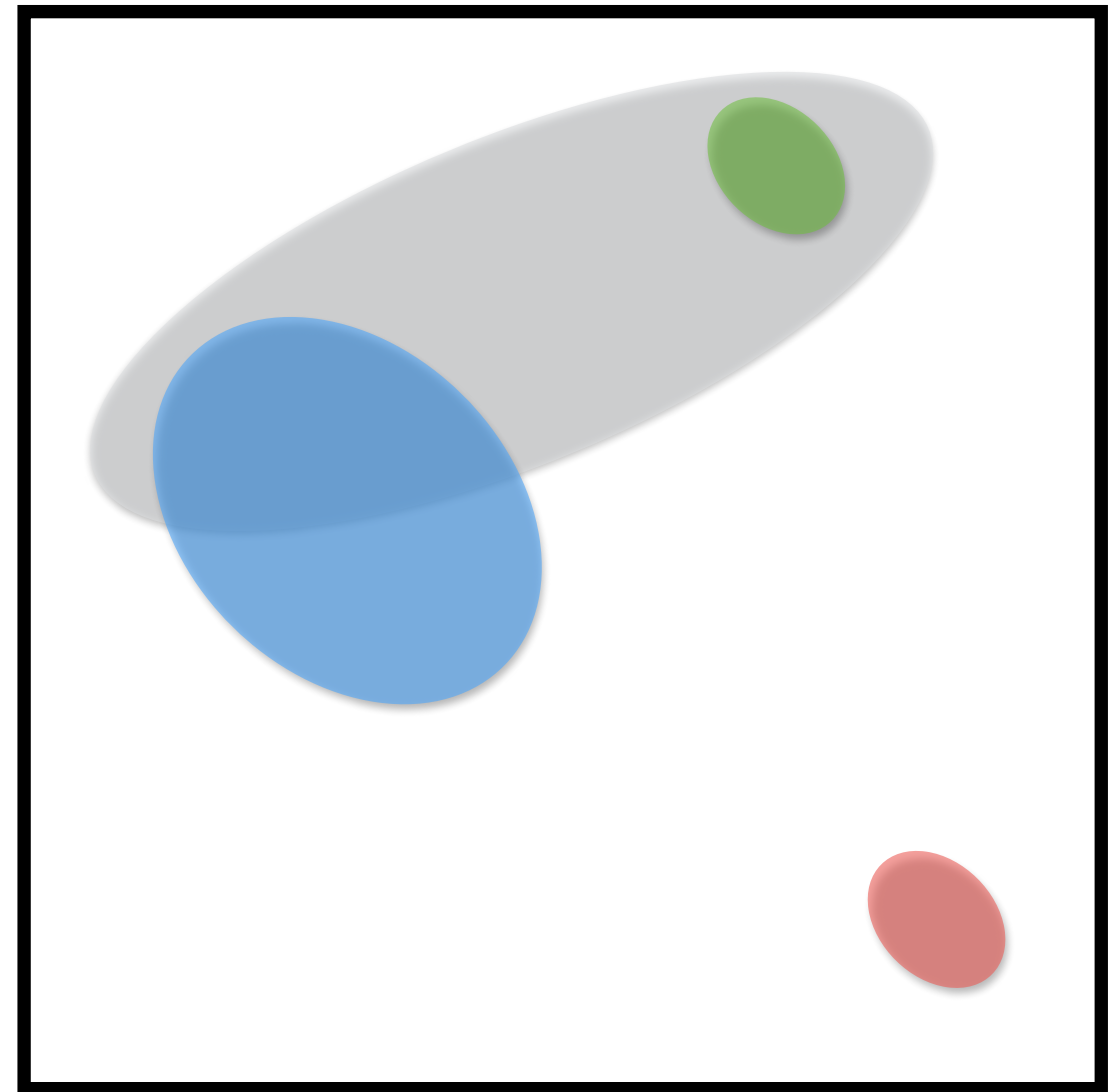


The dietary niche

The food chain

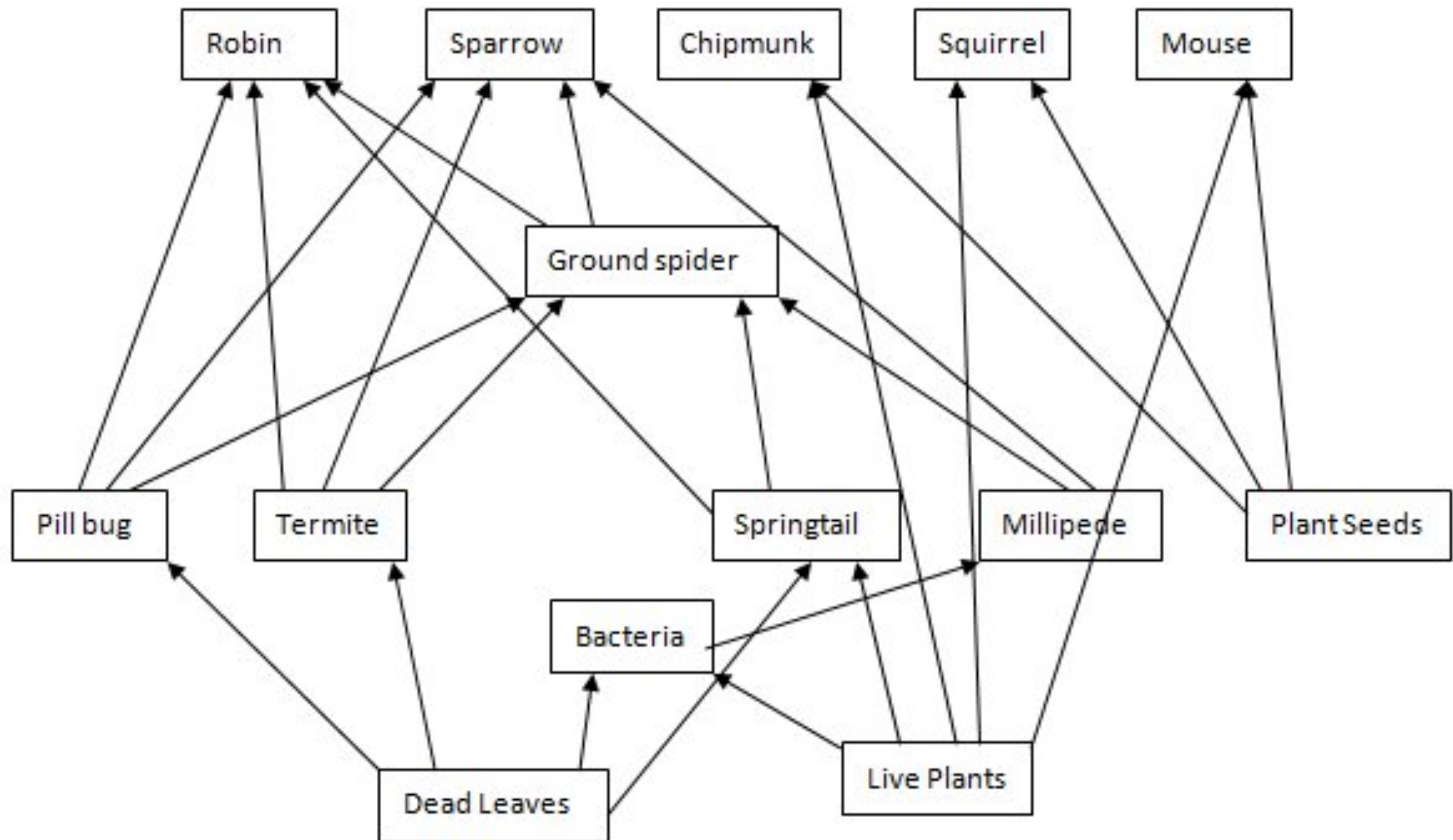


Niche axis 2



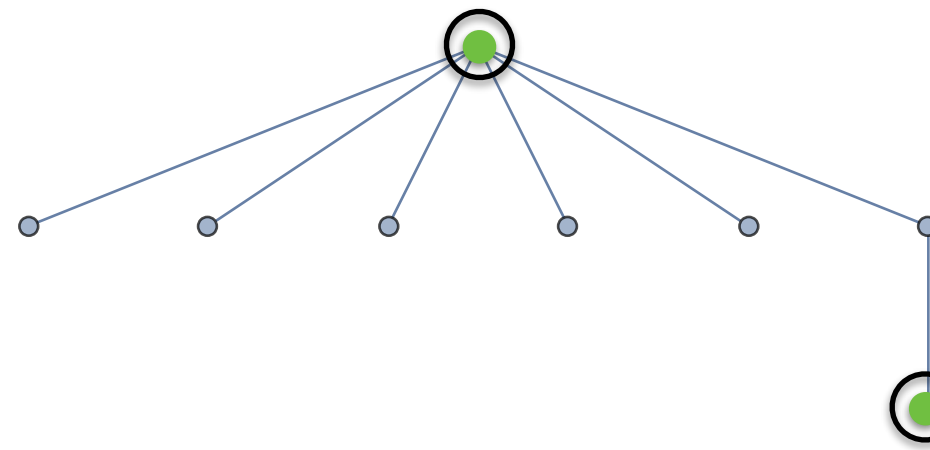
Niche axis 1

From Forbes 1876



Predation: a +/- interaction

Specialists vs. Generalists

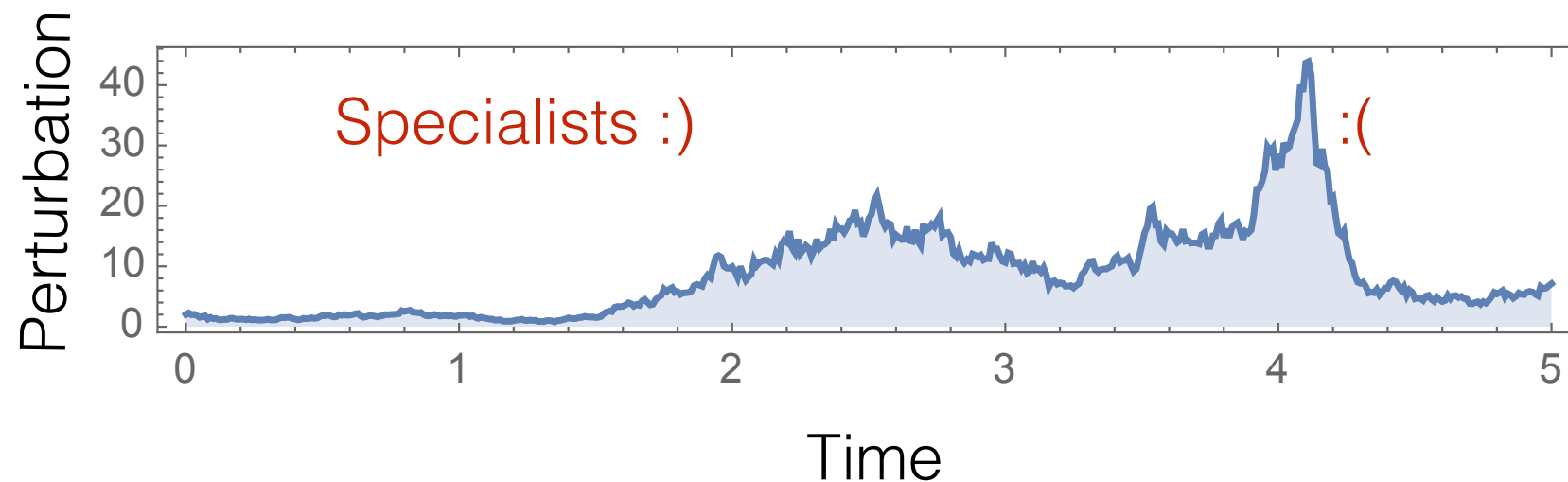


Jack of all trades

Master of one
low “asset diversity”

Ecological rate (ecological timescales)

Evolutionary rate (evolutionary timescales)

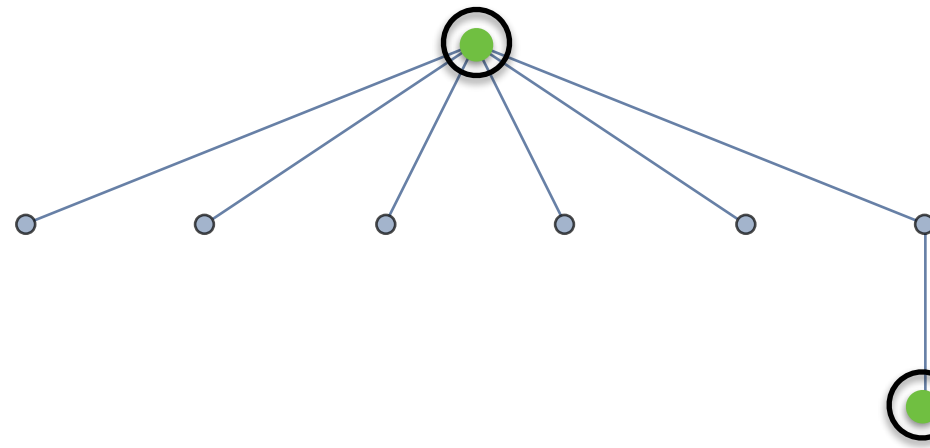


Specialists can outcompete generalists within shorter timescales

Specialists tend to lose in the long run (more sensitive to large perturbations)

Not this simple... eco-evo dynamics

Specialists vs. Generalists



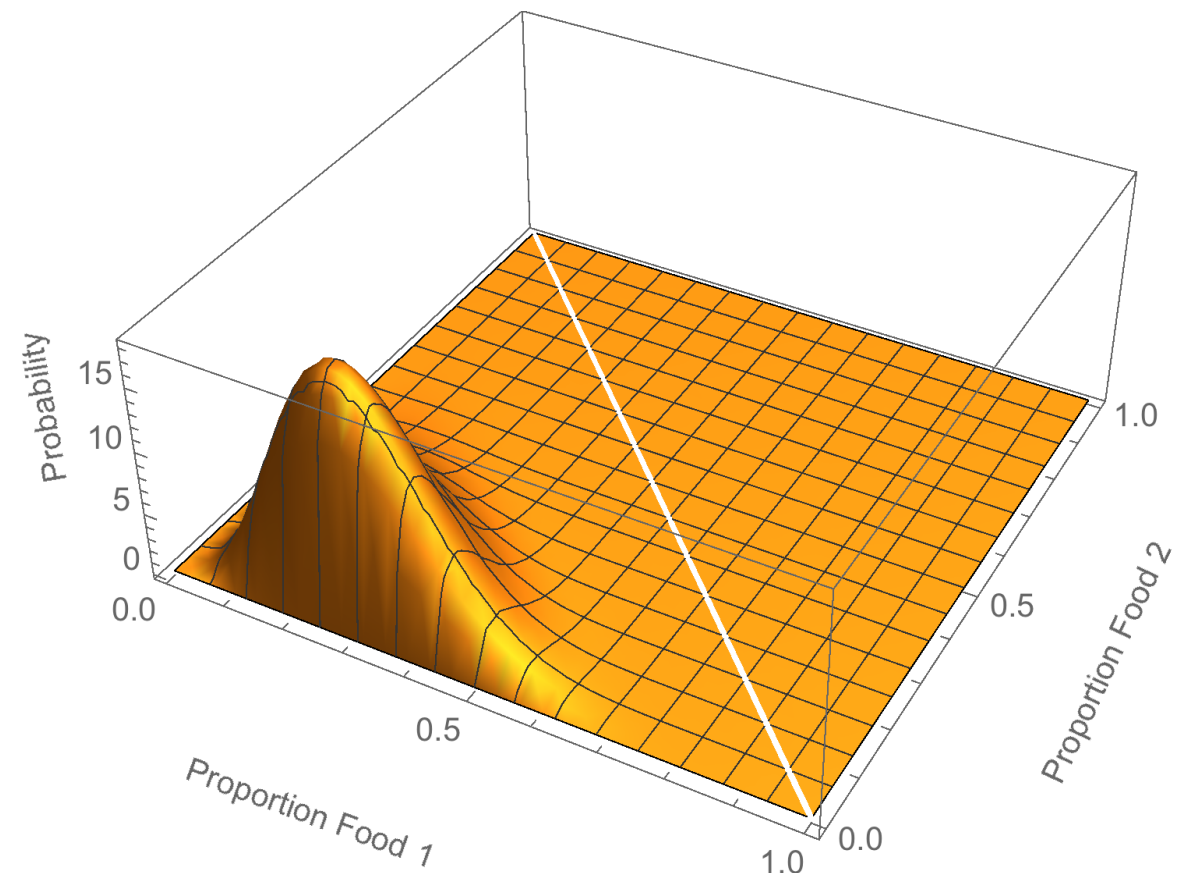
Jack of all trades

Master of one
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More complicated:
individuality
life-history



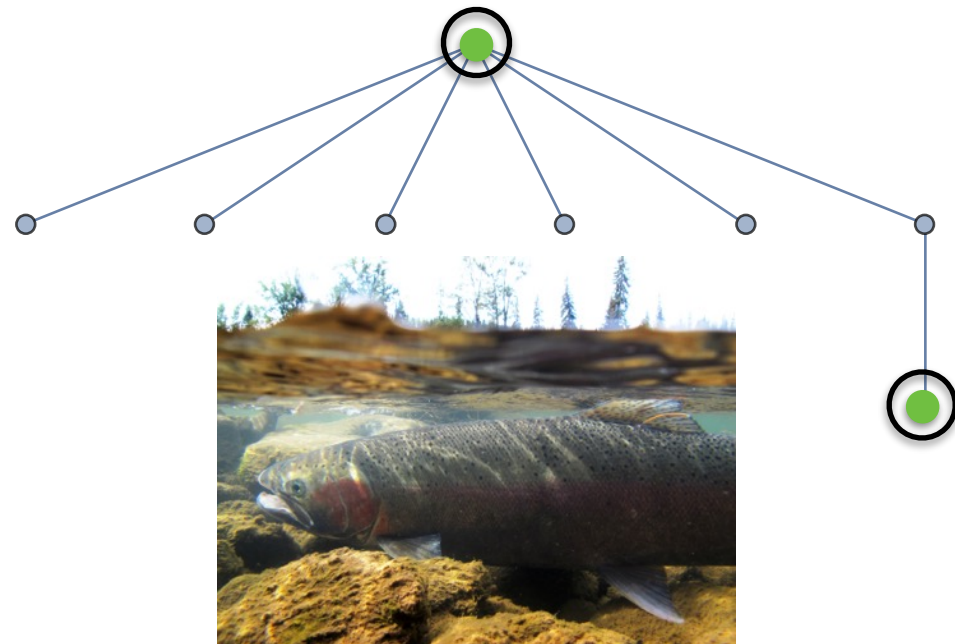
Links have probabilistic weights



Specialists vs. Generalists

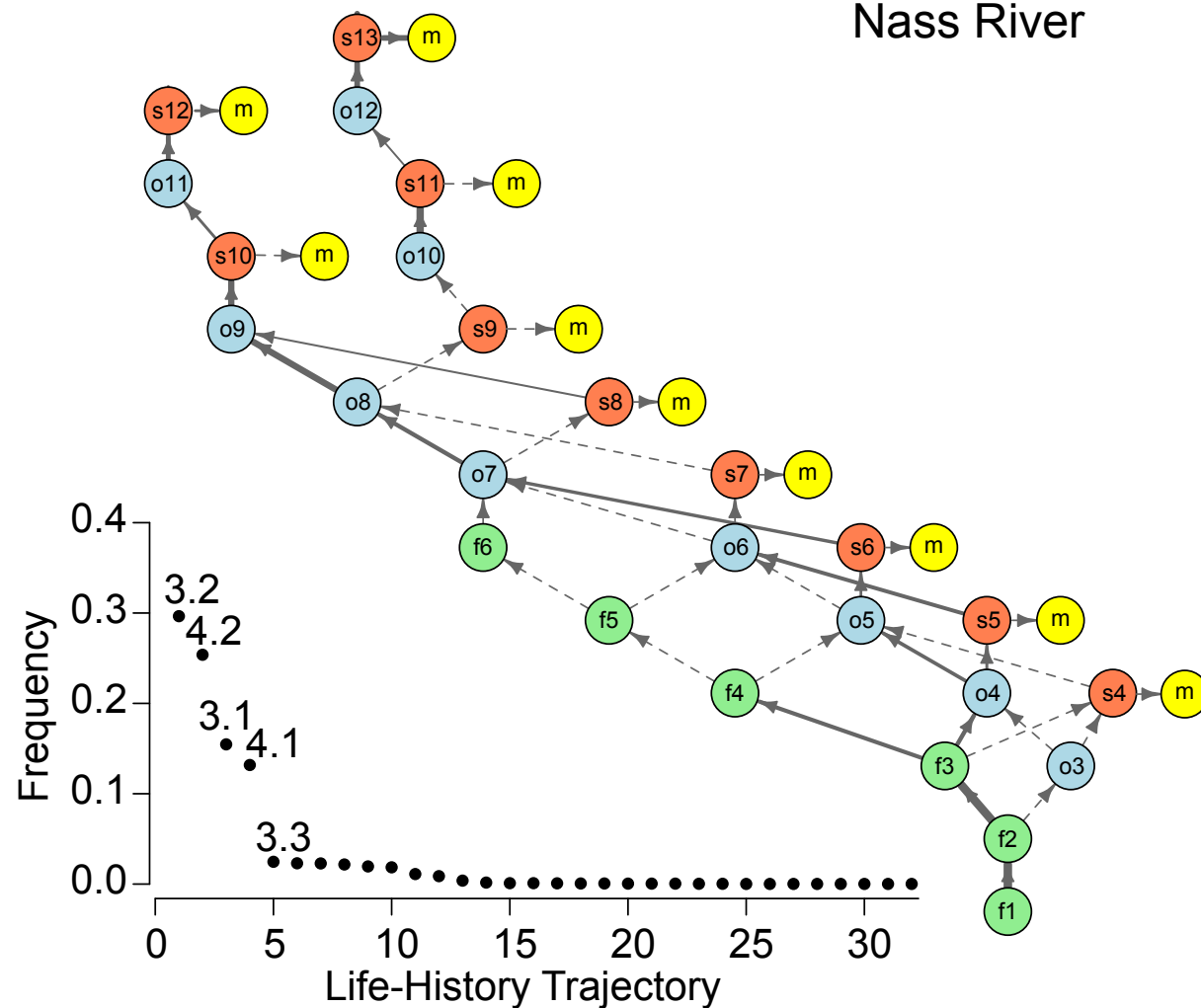
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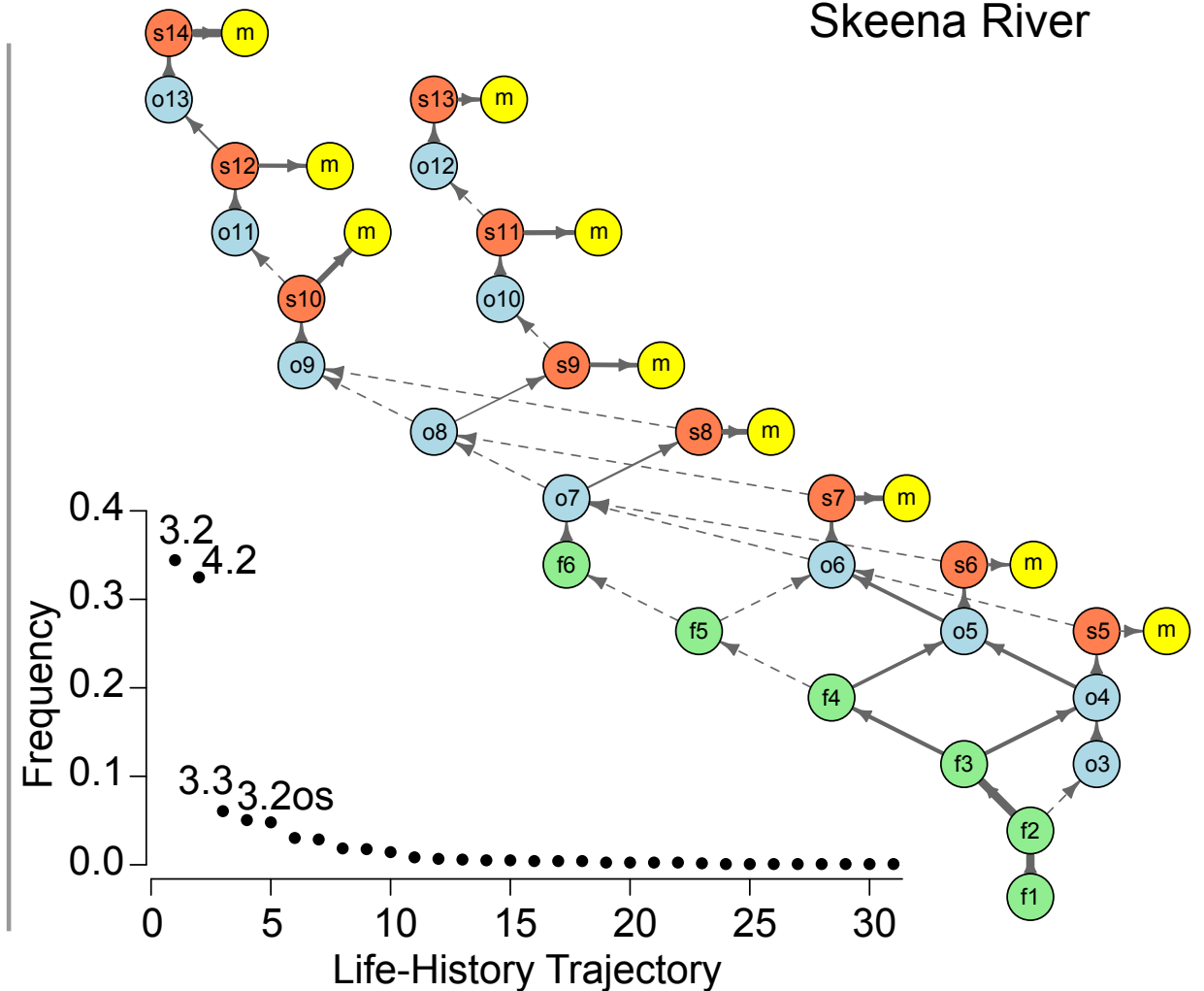


Master of one
low “asset diversity”

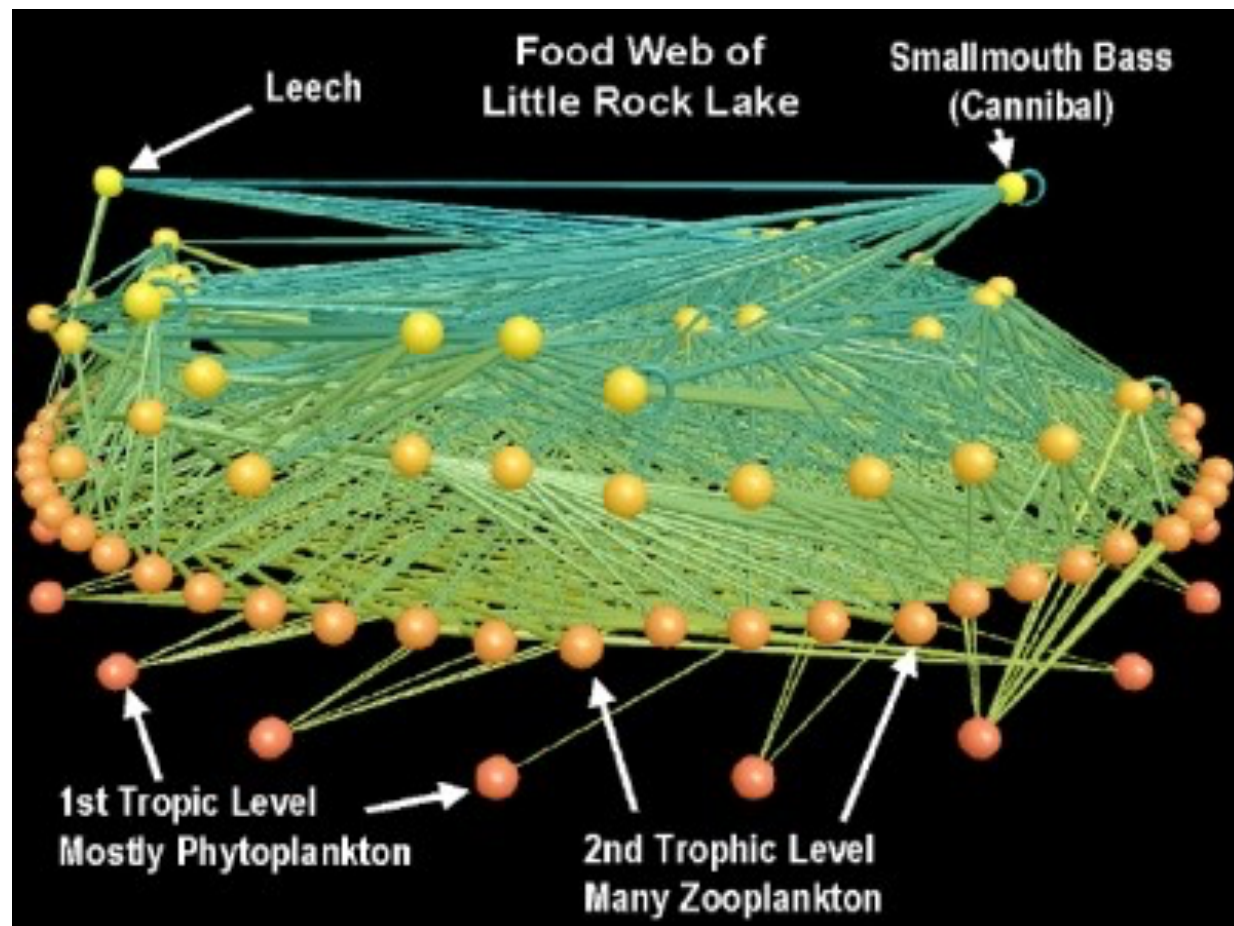
Nass River



Skeena River



The structure of trophic interactions within communities (Dunne, Stouffer)



Some

S # species
 L # Links

$C = L/S^2$ Directed connectance

$SC = L/S$ Links per species
(Avg generality)

Scale-invariant or Scale-dependent?

Scale-invariant:

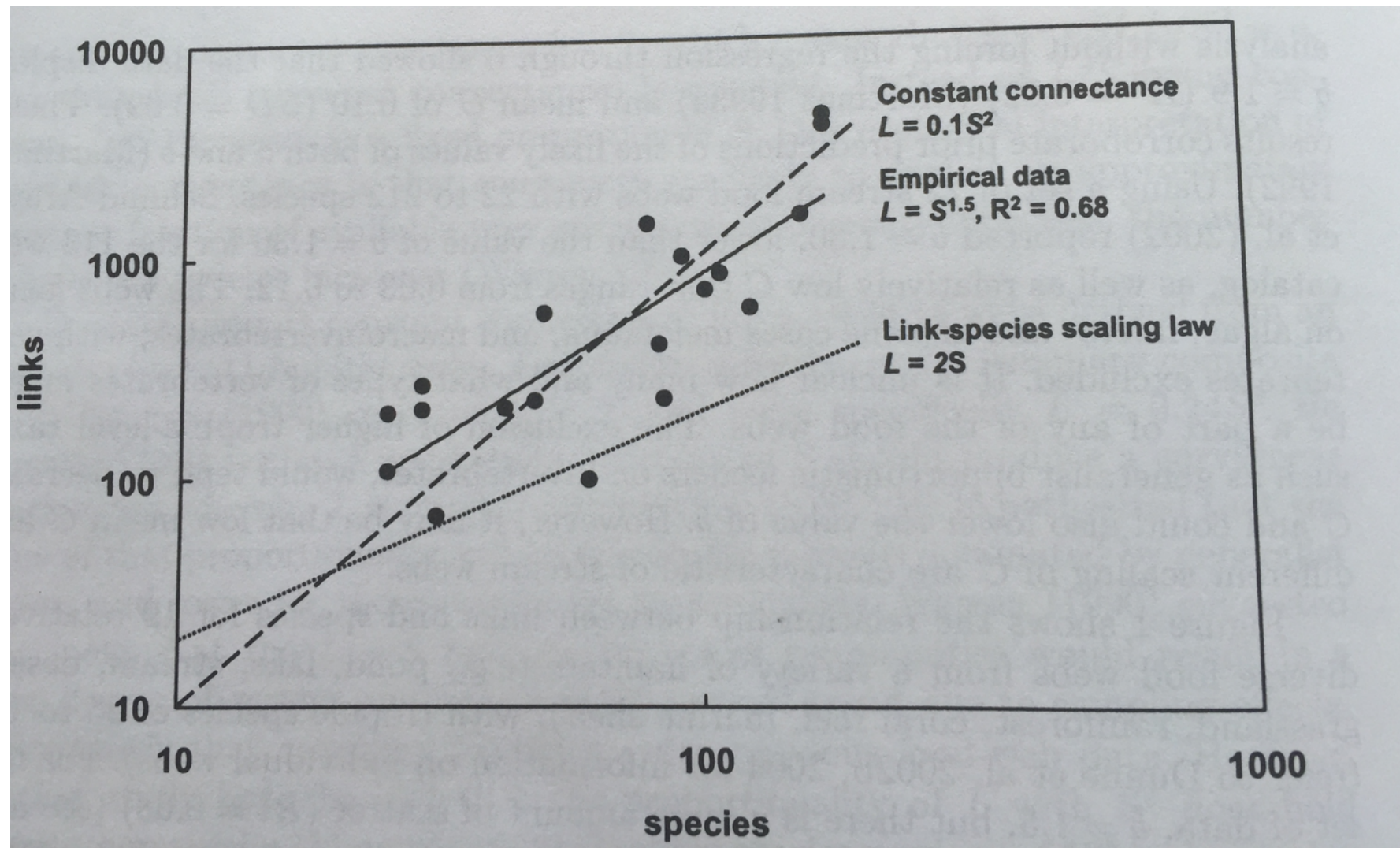
Constant link density (L/S) or pred/prey ratios, etc, as the number of species increase

i.e. increasing #links with increasing numbers of species

Scale-dependent:

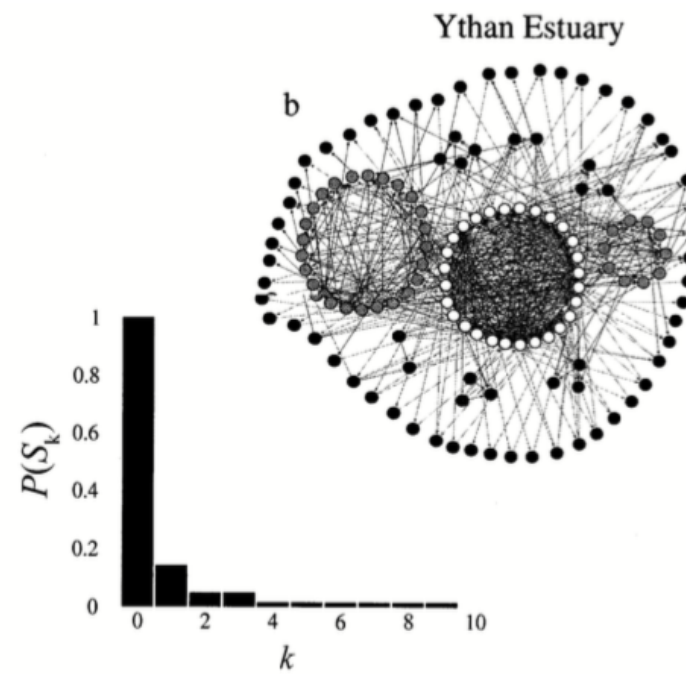
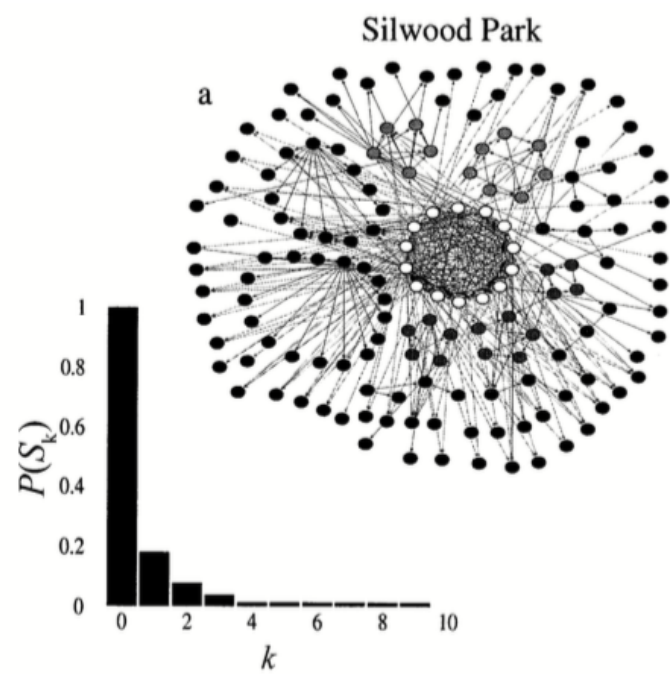
Different link densities (L/S) as well as other structural features for webs of different sizes

There is no universal functional form that describes degree distributions for food web networks

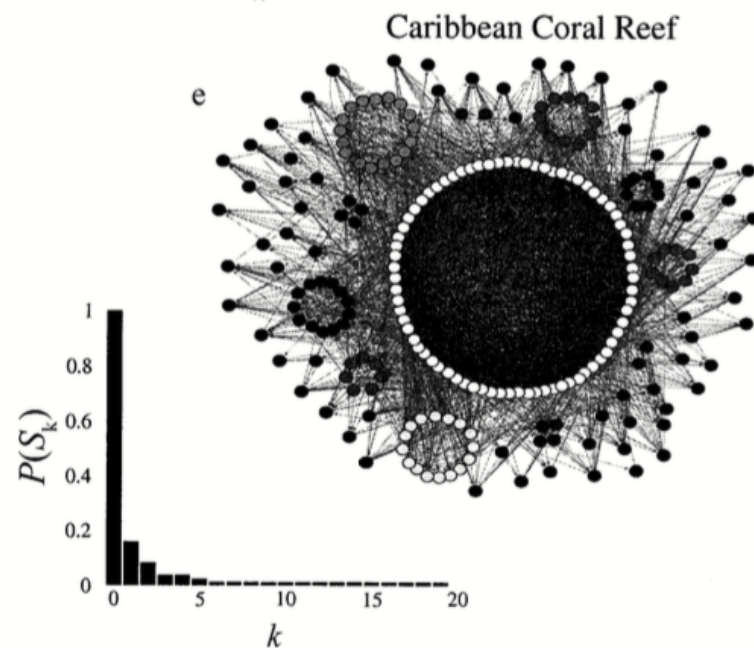
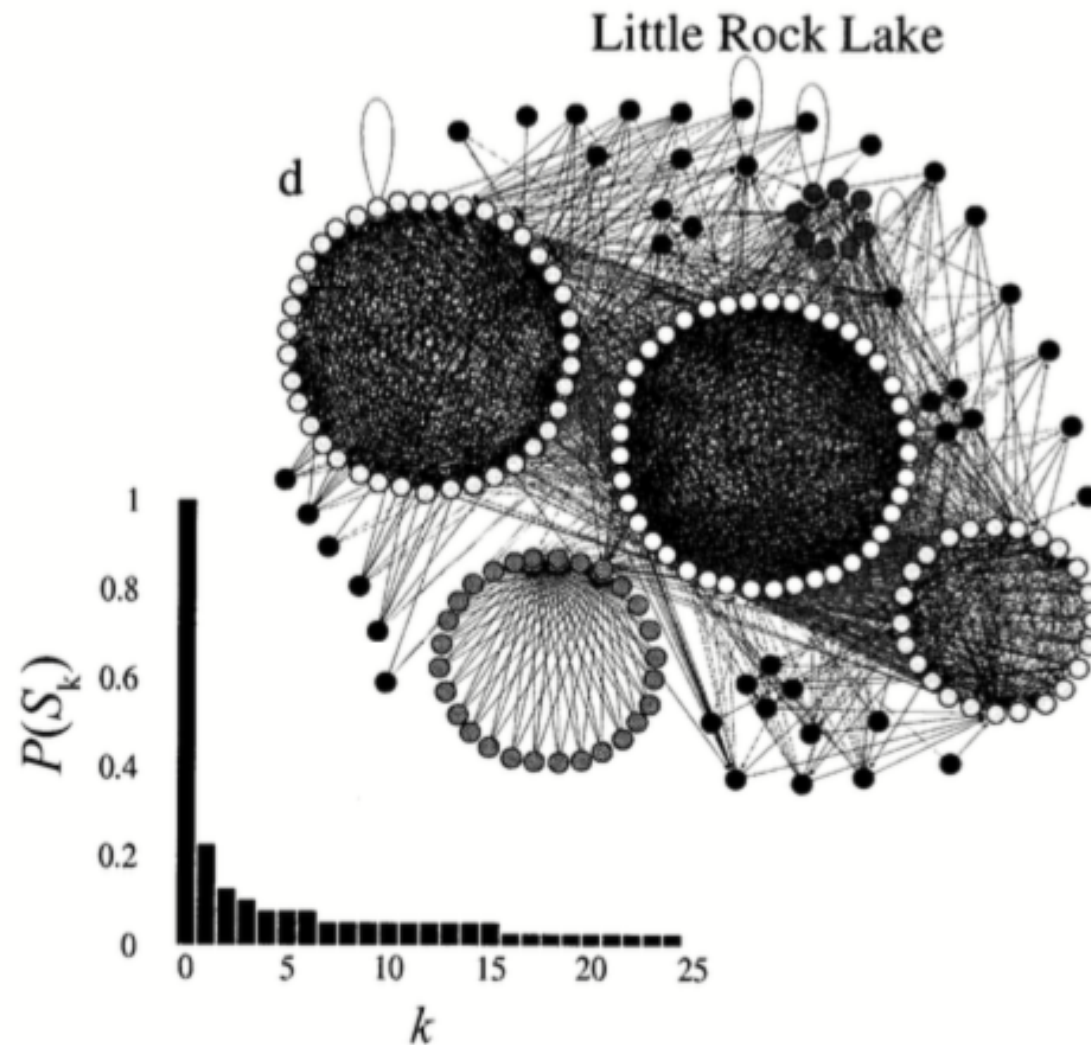
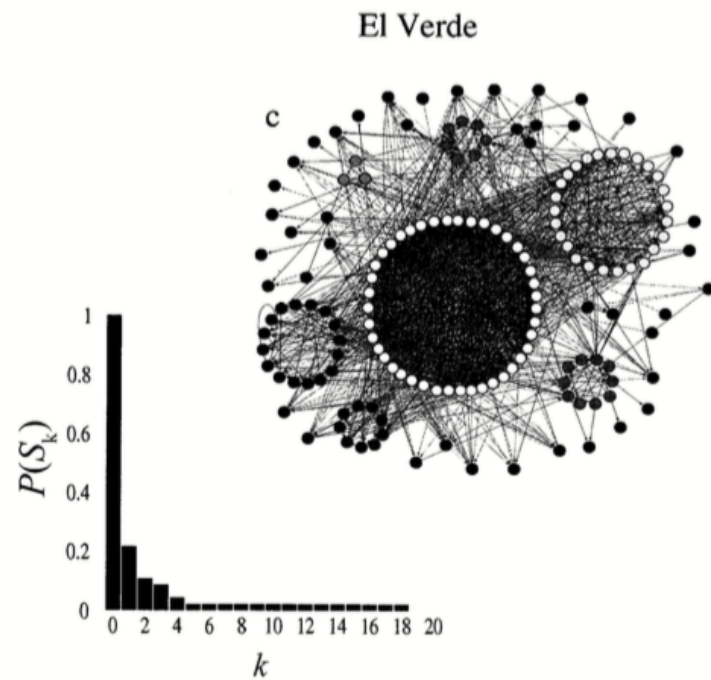


Not a scaling law...

but these are also summary statistics



CDFs for k-subweb frequencies

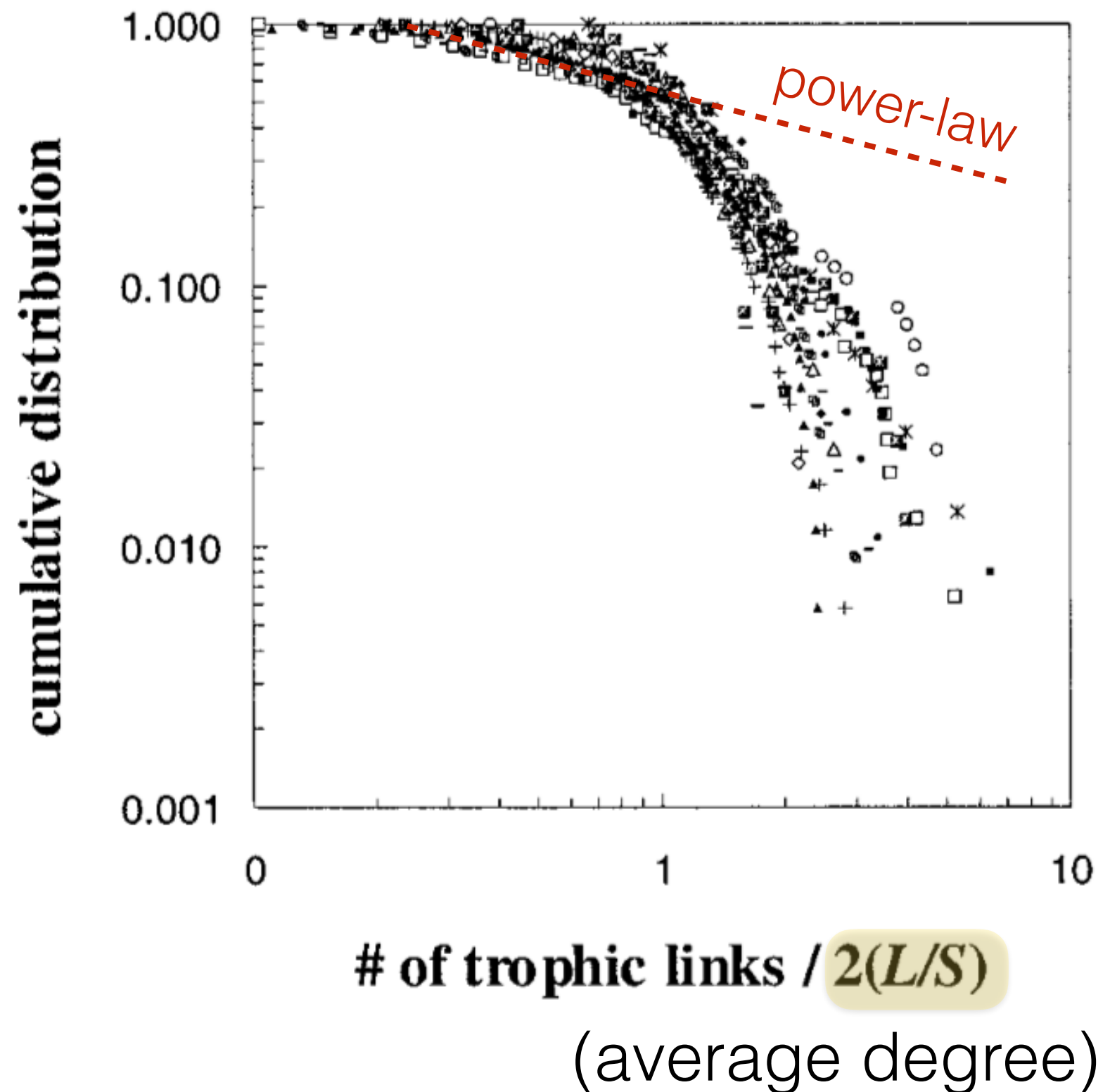


Melian & Bascompte 2004

The structure of trophic interactions within communities

Scaling the CDFs
to $1/2(L/S)$...
i.e. controlling for
scale dependence

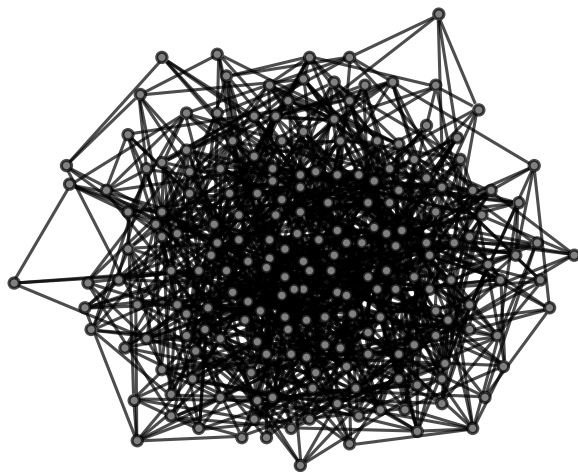
*Distribution tails fall
off more quickly than
for scale-free nets



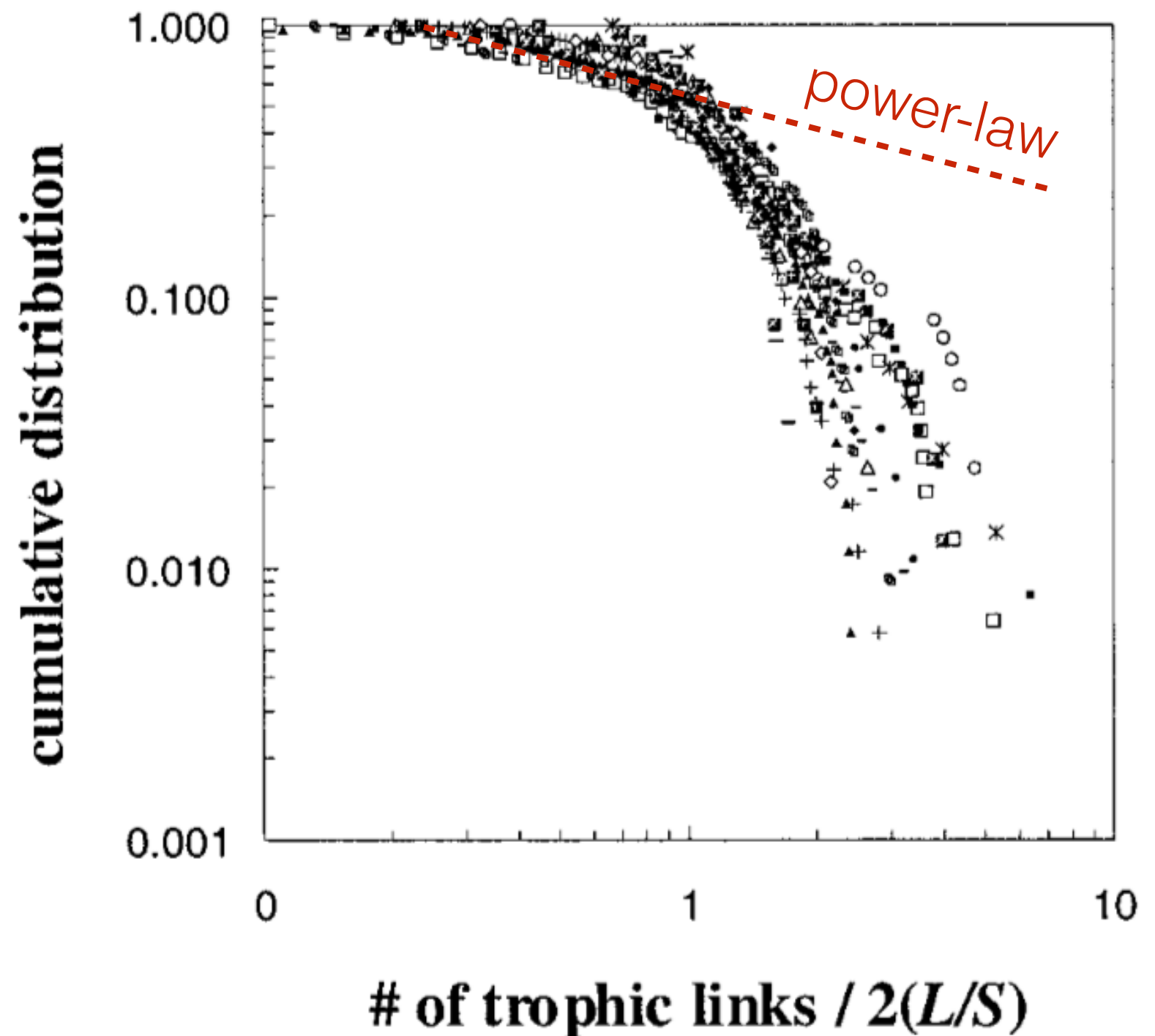
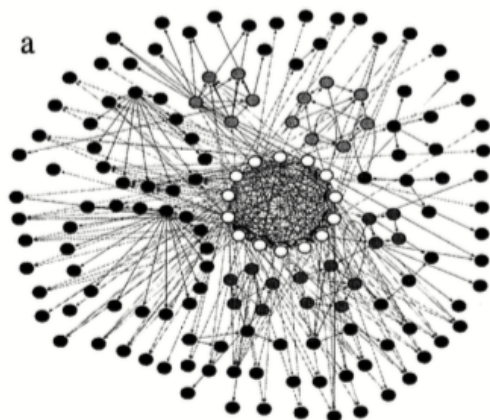
Dunne 2002

The structure of trophic interactions within communities

Assembly differences?
preferential attachment
vs.
ecological assembly

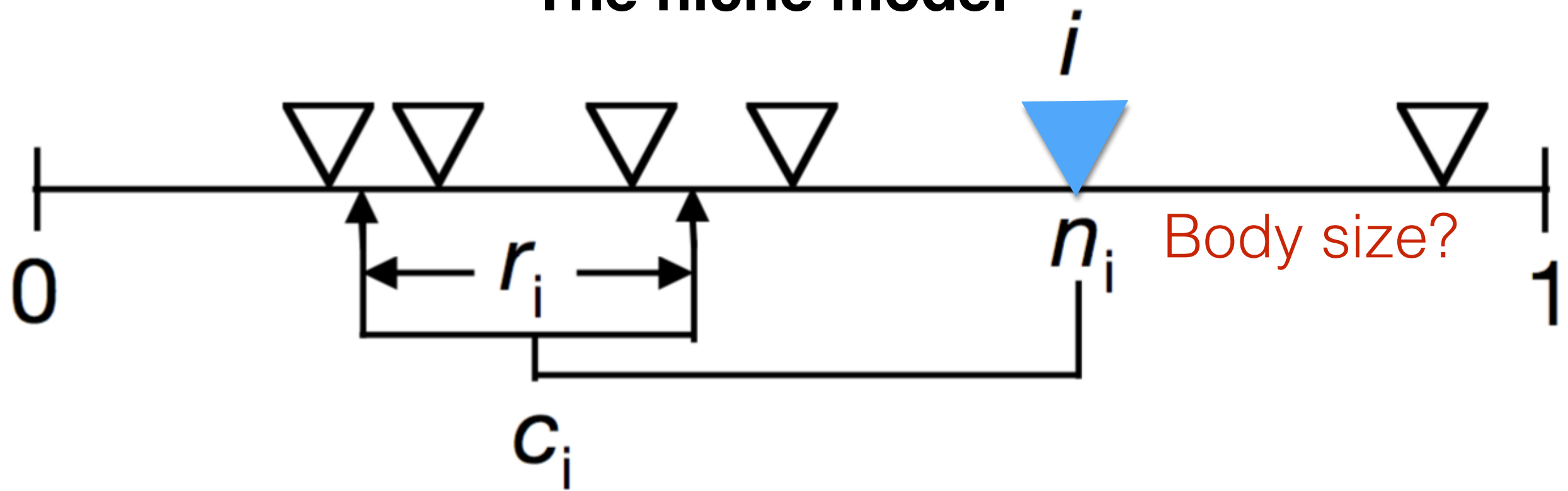


vs.



A statistical model for food web structure

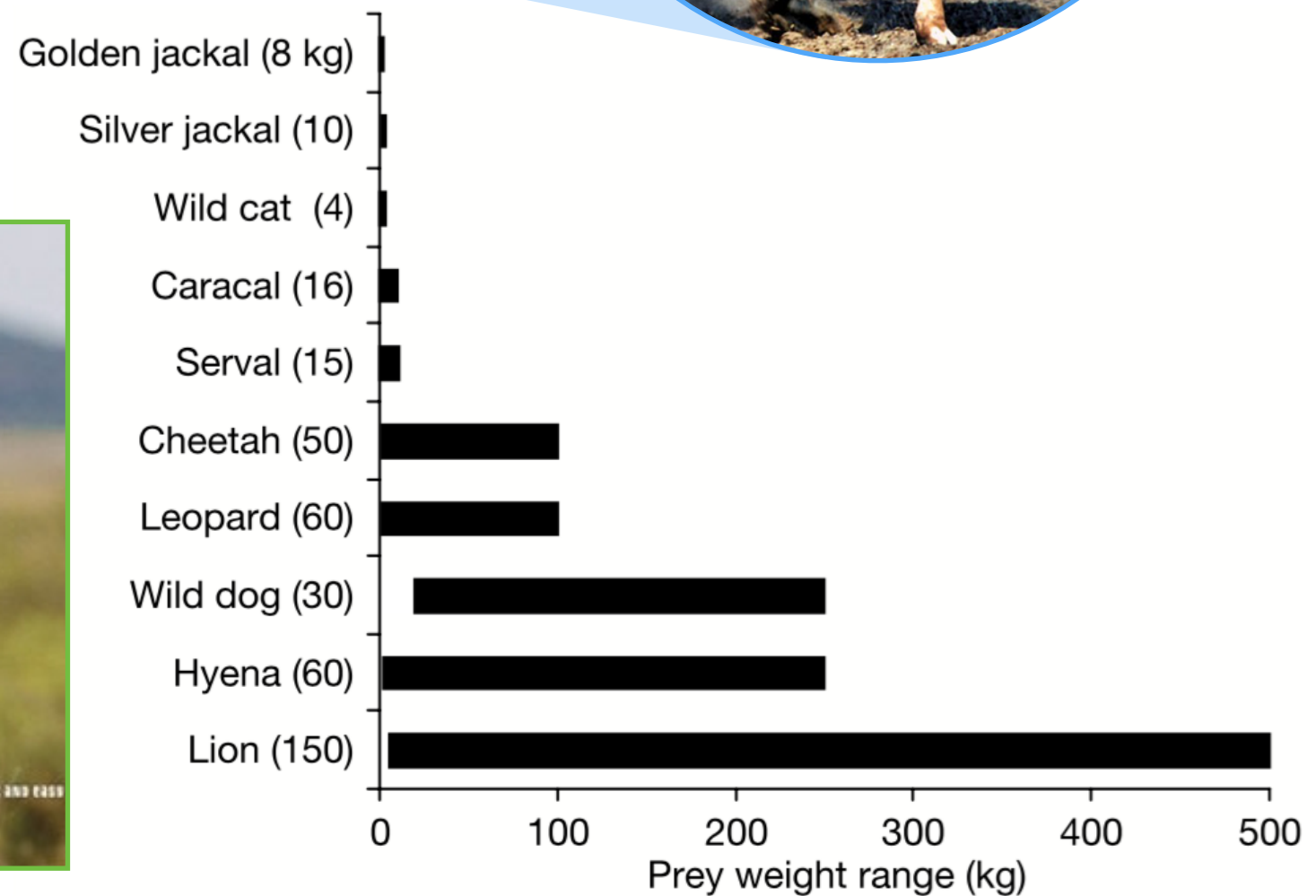
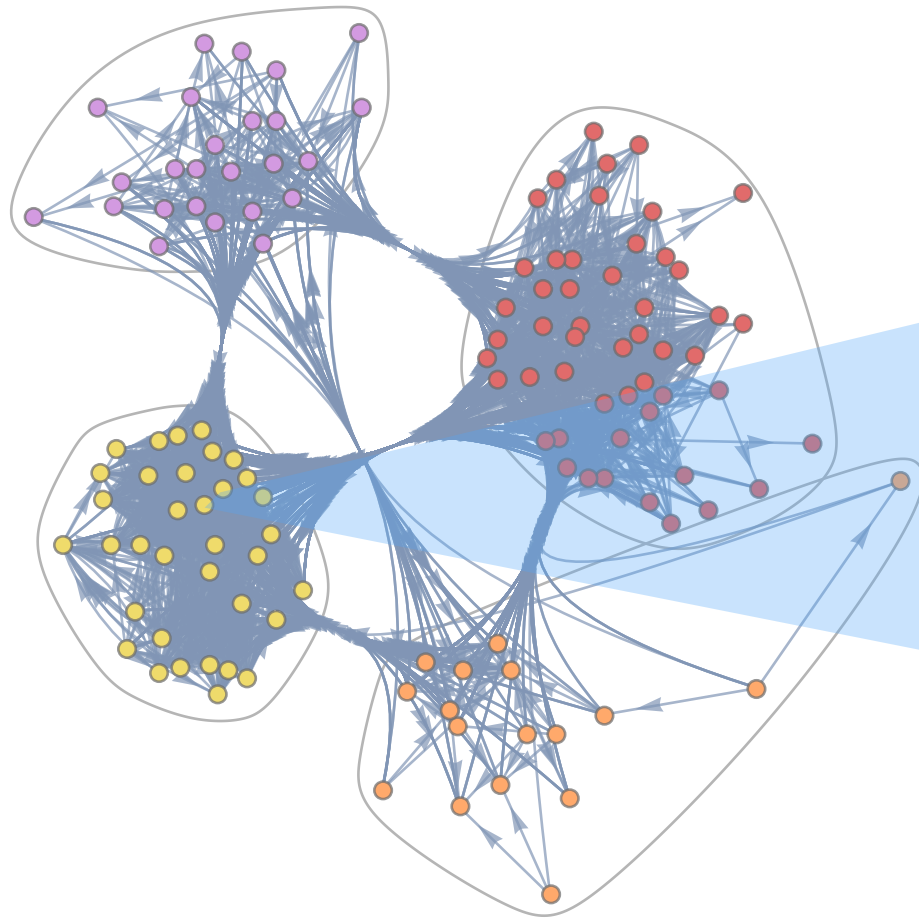
“The niche model”



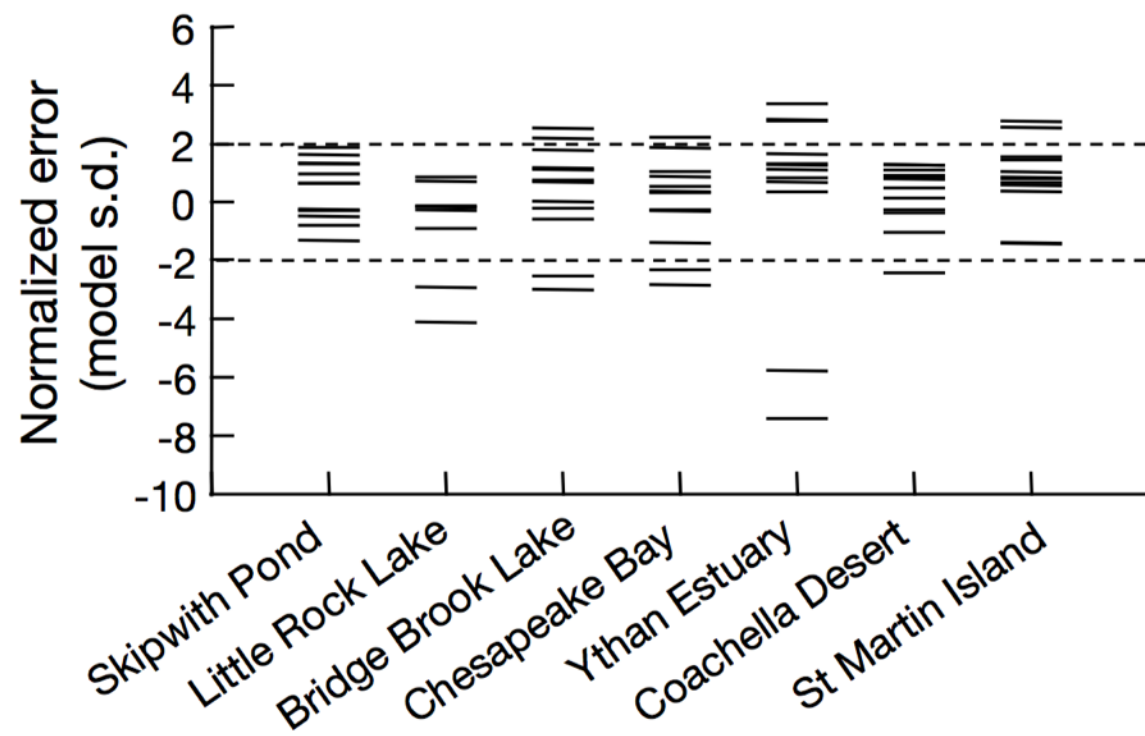
- 1) Feeding interaction distributed across a niche axis
- 2) Dietary generality/specialization determined by range
 $r_i = n_i * x_i$ where $x_i \sim \text{Beta}(a=1, b)$ where b is chosen such that system matches observed connectance
- 3) $c_i \sim \text{Uniform}(r_i/2, n_i)$

Generality increases with n_i !

Generality increases with n_i !

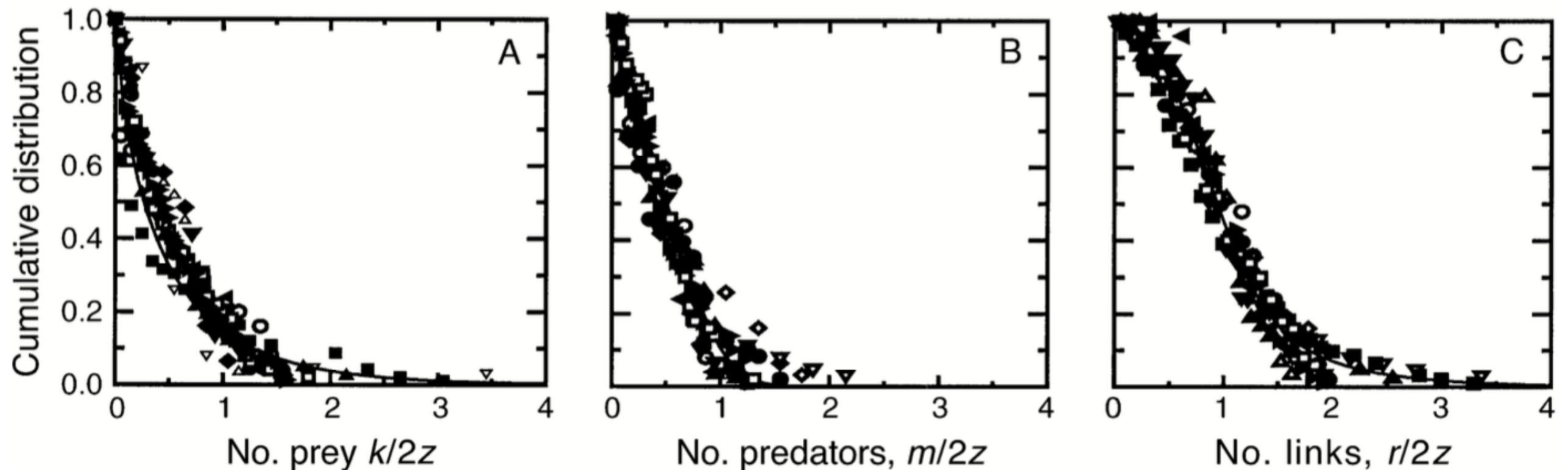


Sinclair et al. 2010

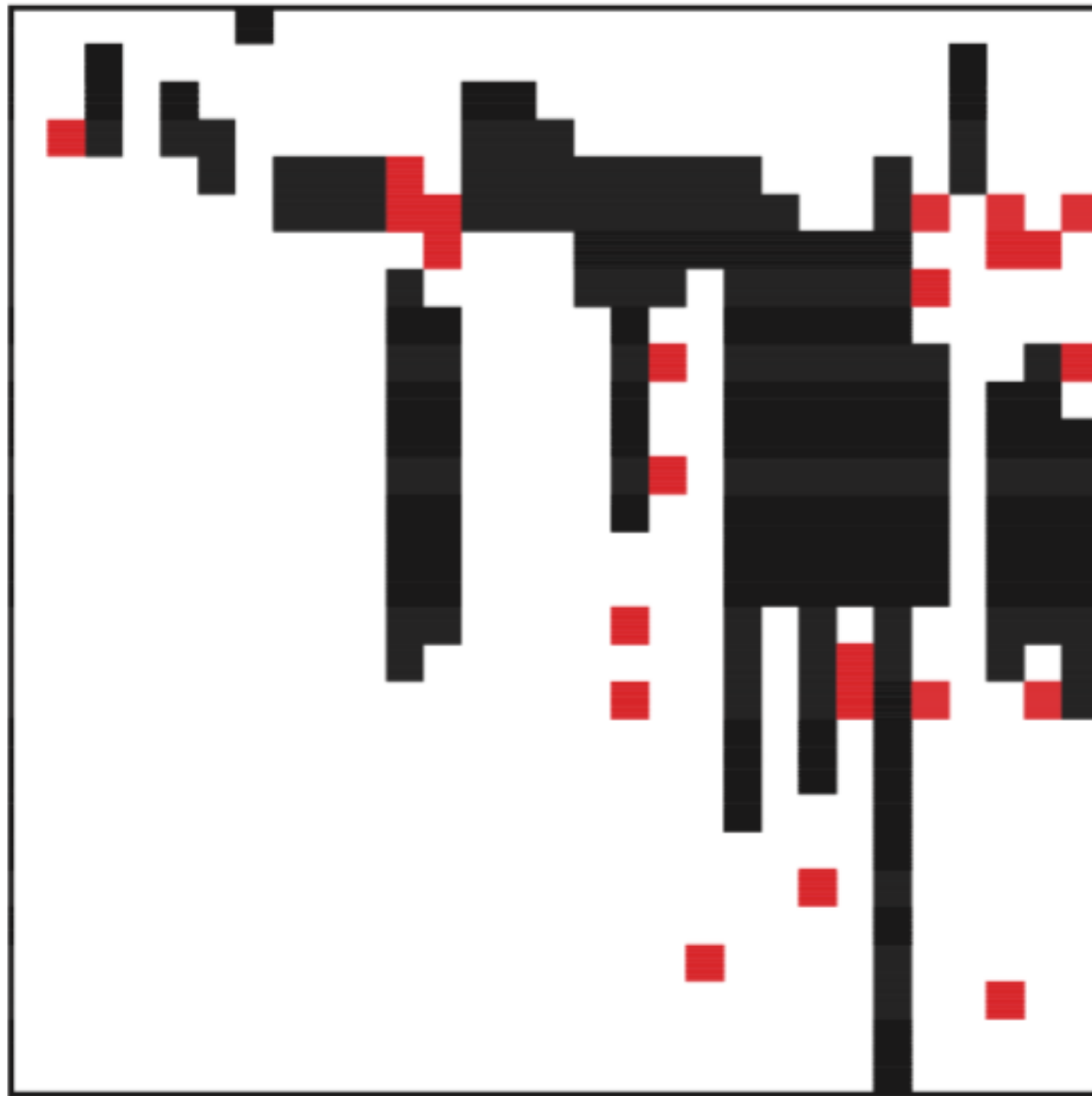


Testing the niche model's ability to replicate various food web properties

Analytical predictions of the niche model vs. data



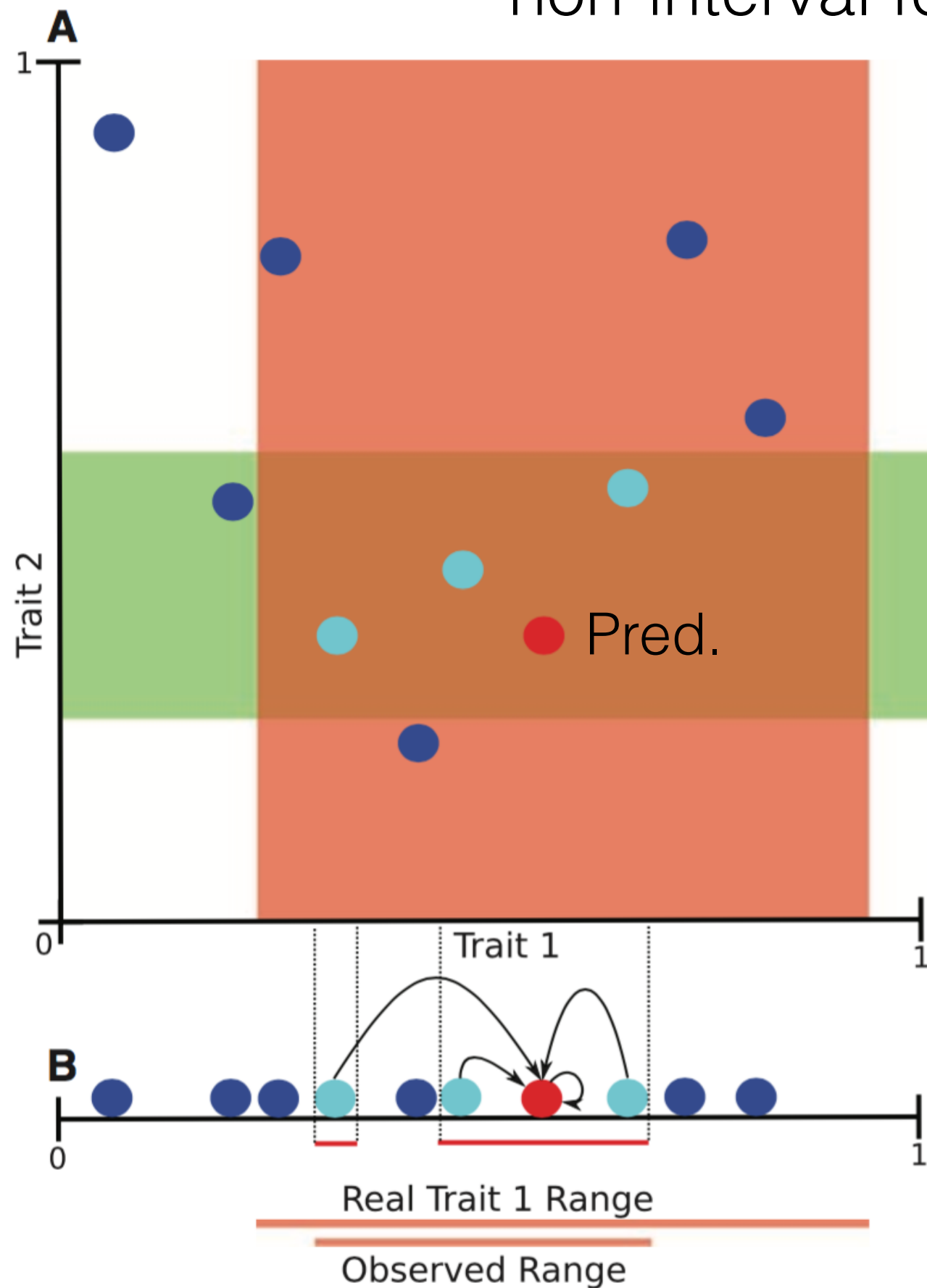
The problem with interval assumptions



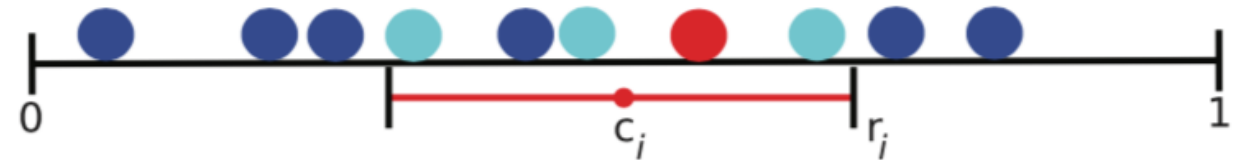
Red = interactions incompatible with niche-type models

What do we gain from multiple niche dimensions?

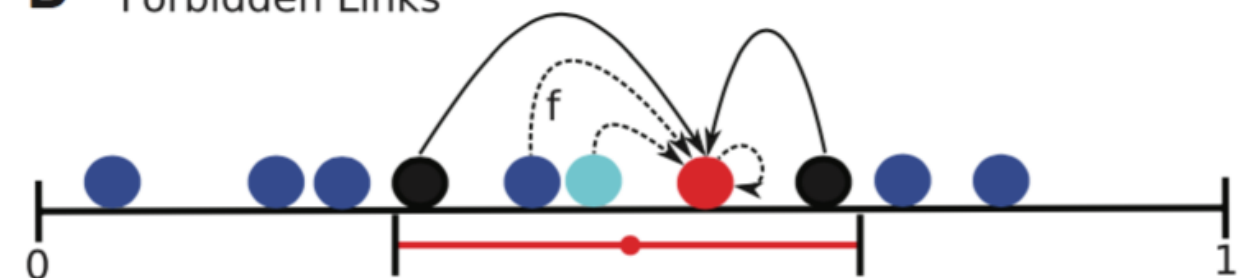
non-interval feedings ranges



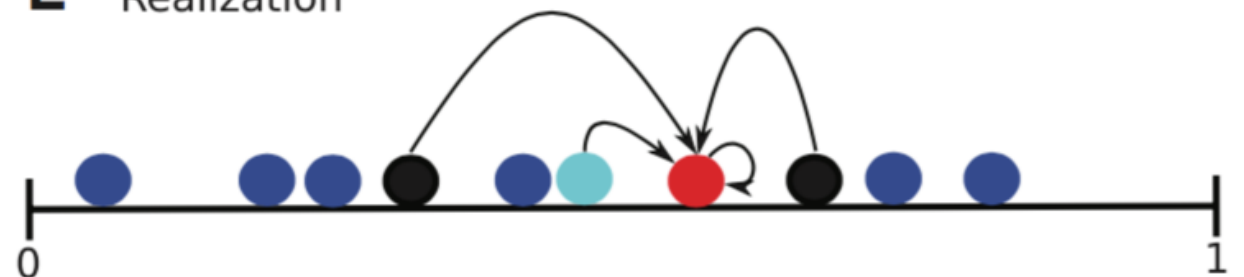
C Potential Range



D Forbidden Links



E Realization

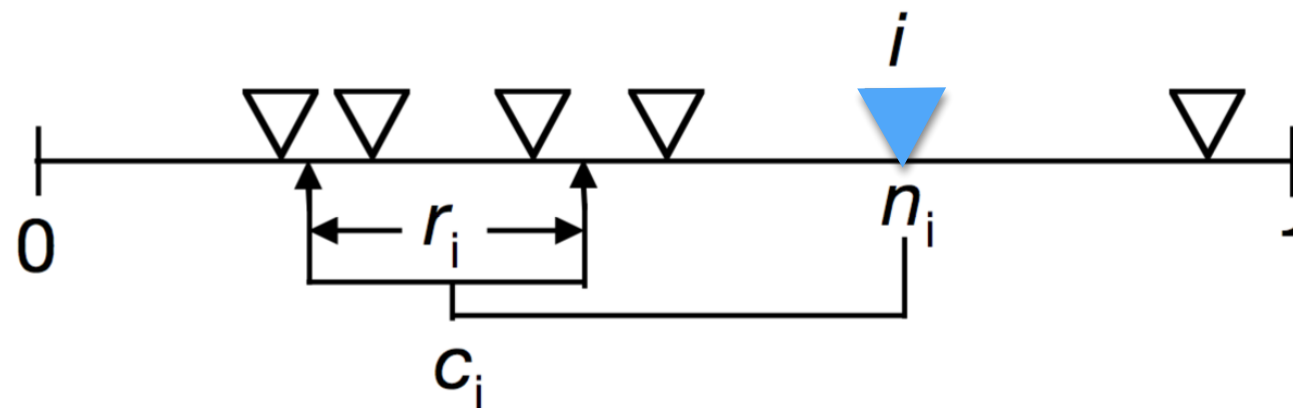


Fitting models to data...

The probabilistic niche model (PNM)

Release assumptions about:

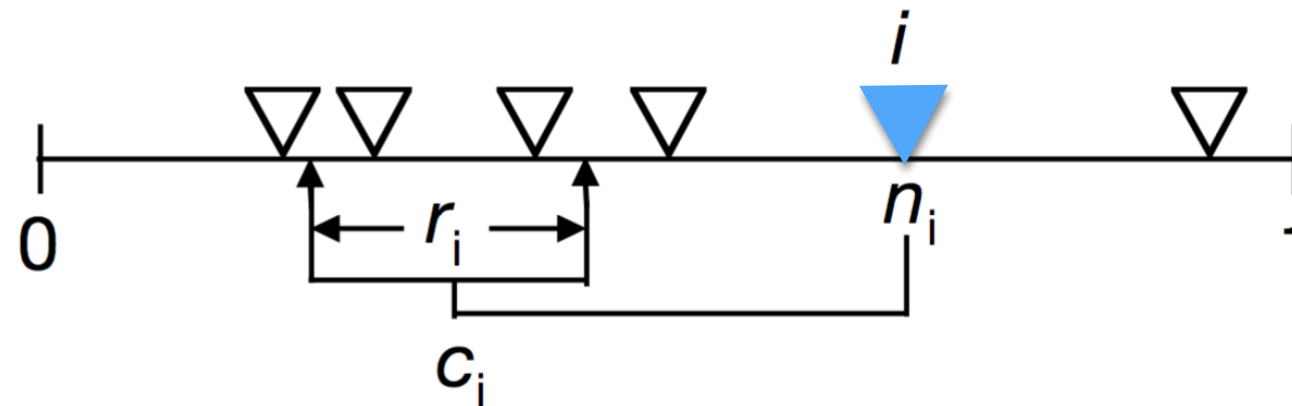
- 1) distribution of species across n
- 2) dimensionality of n
- 3) c_i the center of the dietary range for each species
- 4) r_i the dietary generalization of each species



Randomly assign $\theta = \{n_{d,i} \ c_{d,i} \ r_{d,i}\} \sim \text{Uniform}(0,1)$

Fitting models to data...

The probabilistic niche model (PNM)



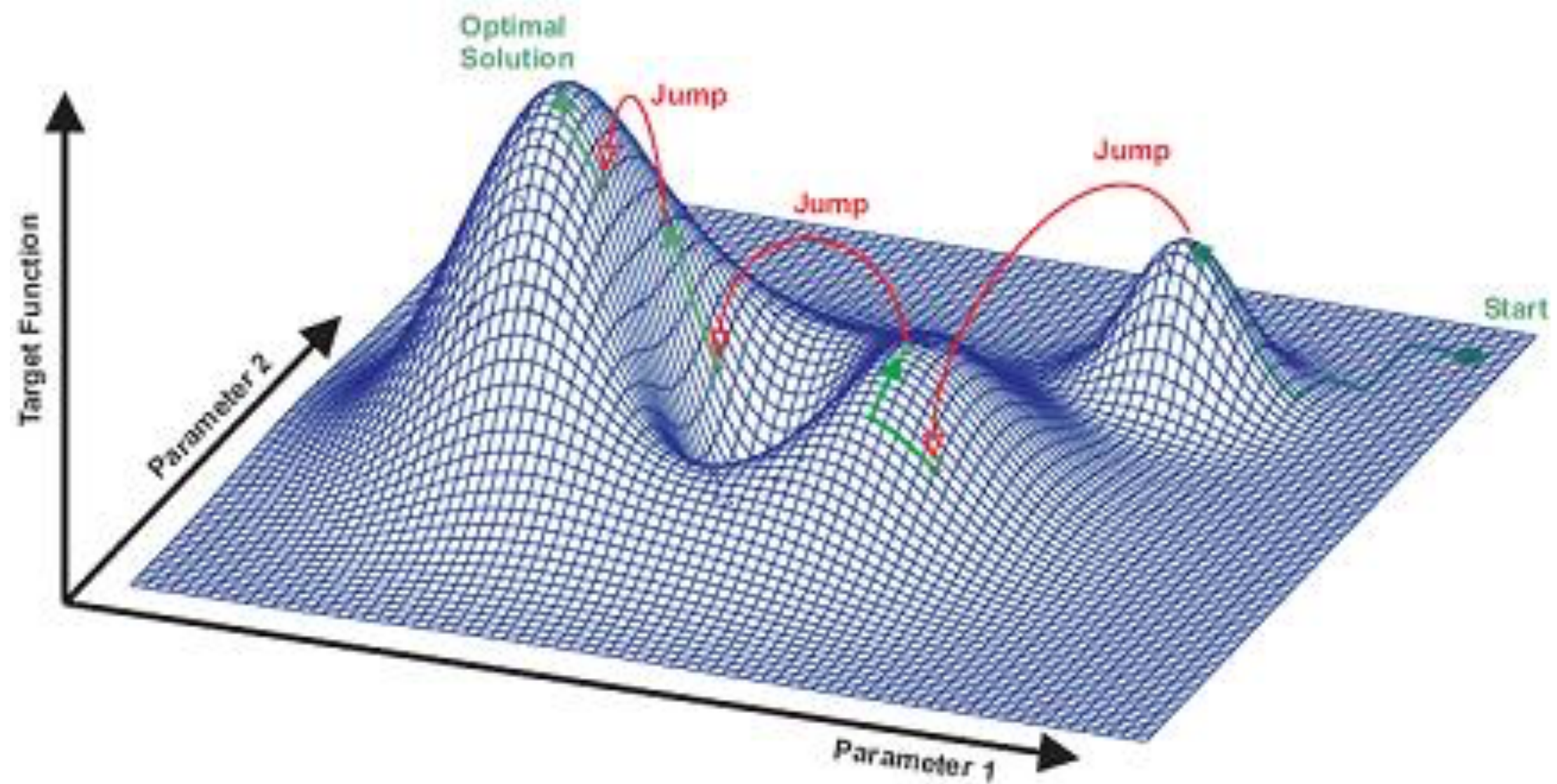
Randomly assign $\theta = \{n_{d,i} \ c_{d,i} \ r_{d,i}\} \sim \text{Uniform}(0,1)$

Use simulated annealing algorithm to find Maximum Likelihood Estimate given observed set of feeding relationships **A**

$$L(\theta|\mathbf{A}) = \sum_i \sum_j \ln \left\{ \begin{array}{ll} P(i, j|\theta) & \text{if } a_{ij} = 1 \\ 1 - P(i, j|\theta) & \text{if } a_{ij} = 0 \end{array} \right\}.$$

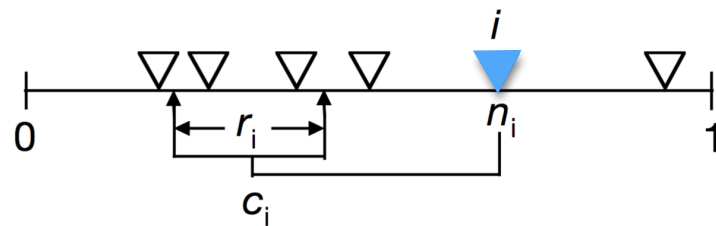
$$\text{given } P(i, j|\theta) = \alpha \prod_{d=1}^D \exp \left(- \left| \frac{n_{d,j} - c_{d,i}}{r_{d,i}/2} \right|^e \right)$$

Simulated Annealing

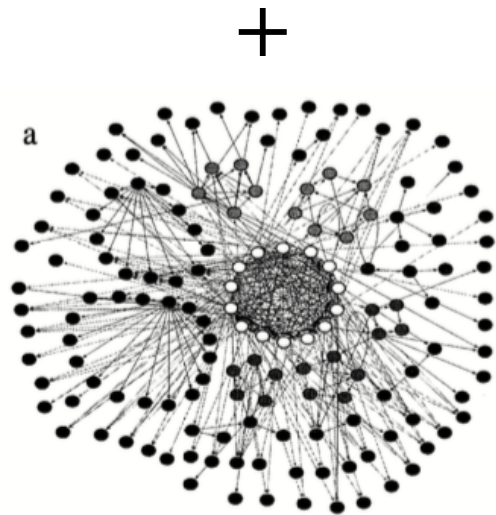


Fitting models to data...

The probabilistic niche model

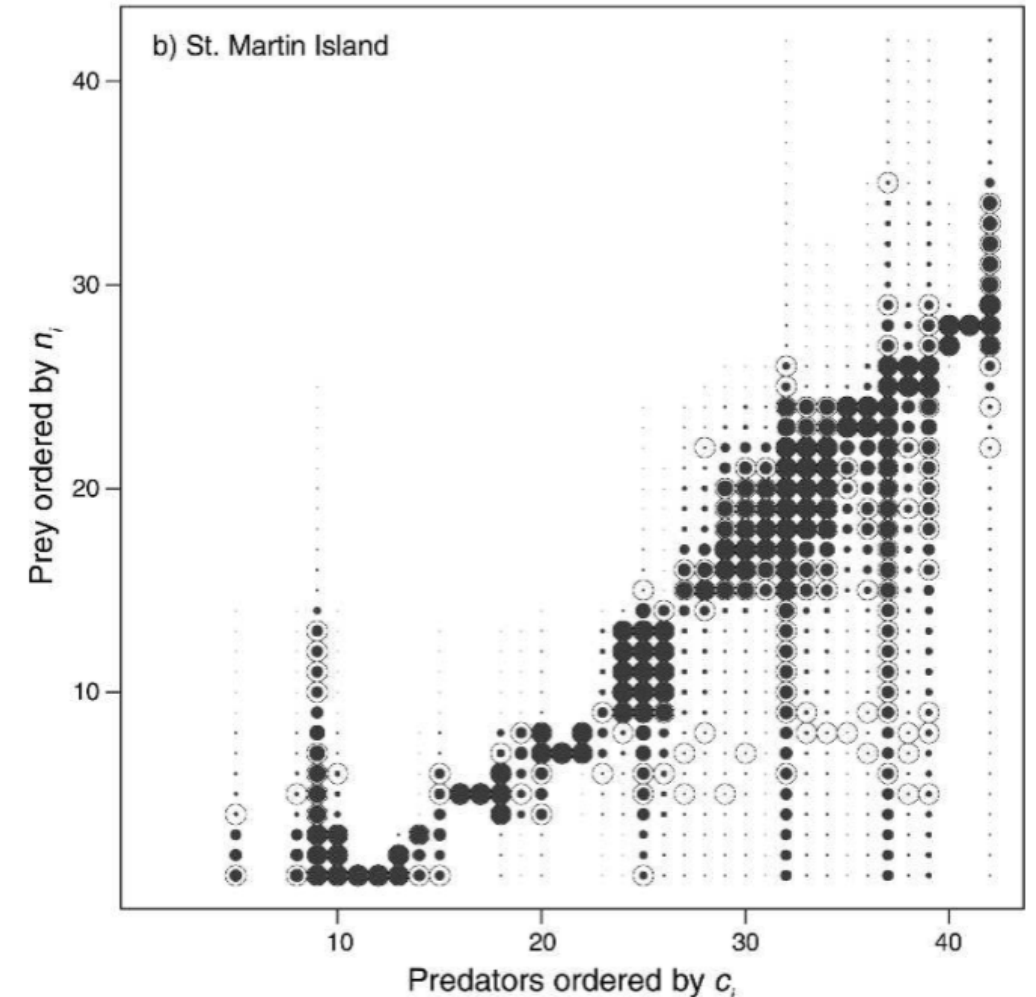
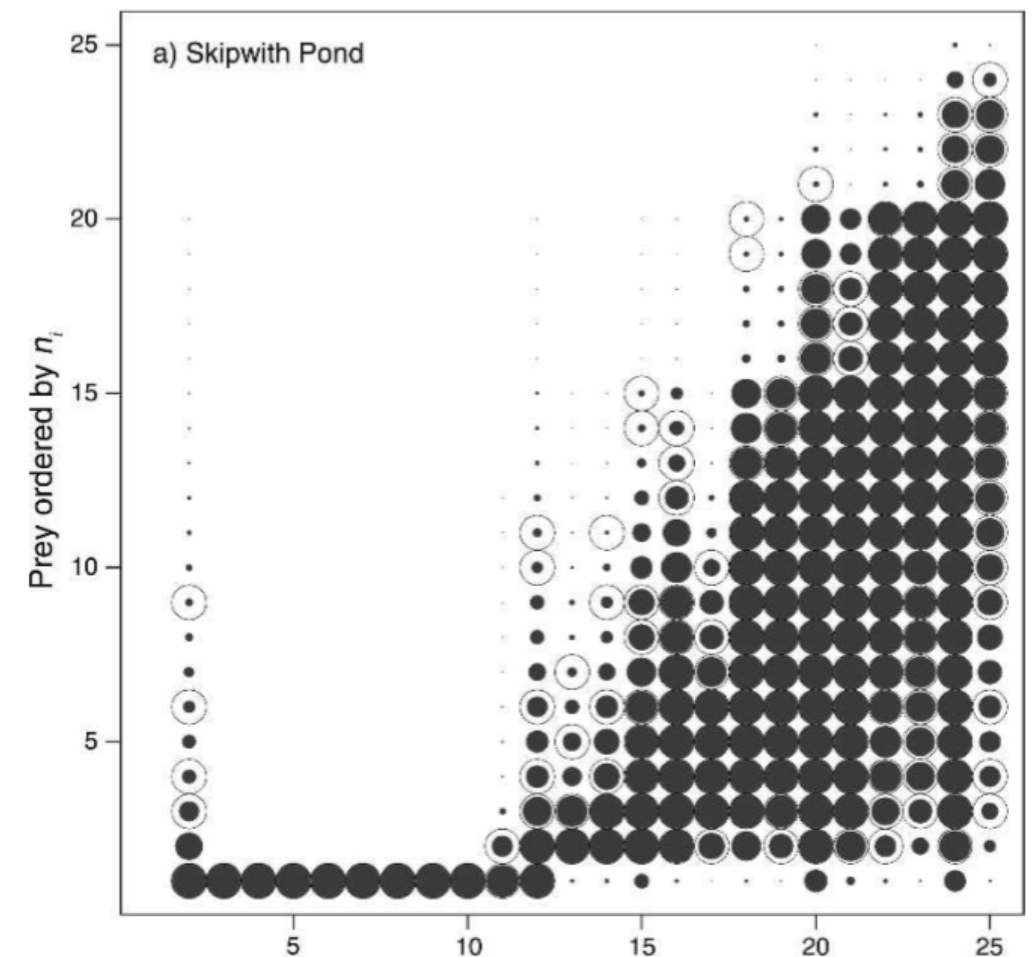


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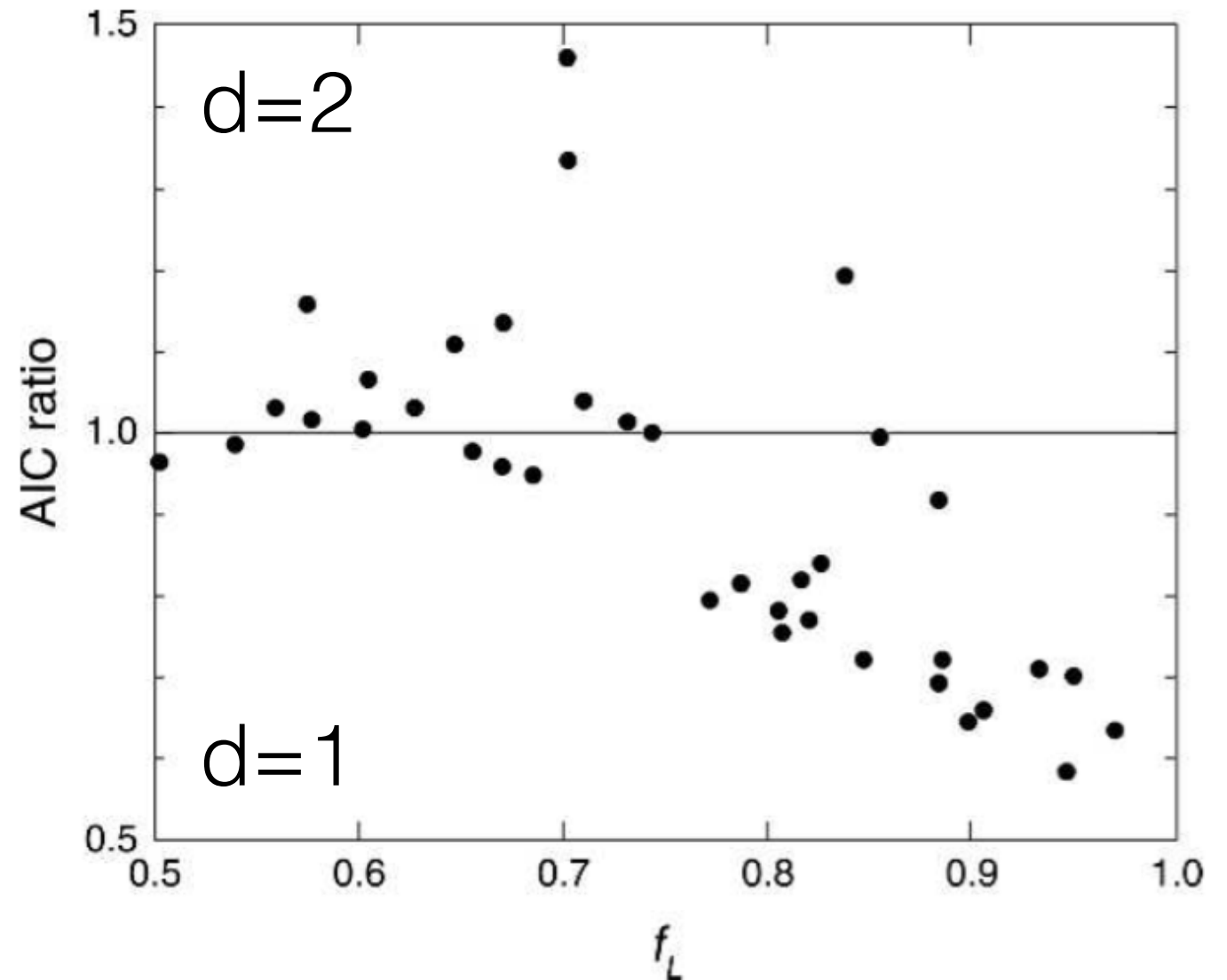
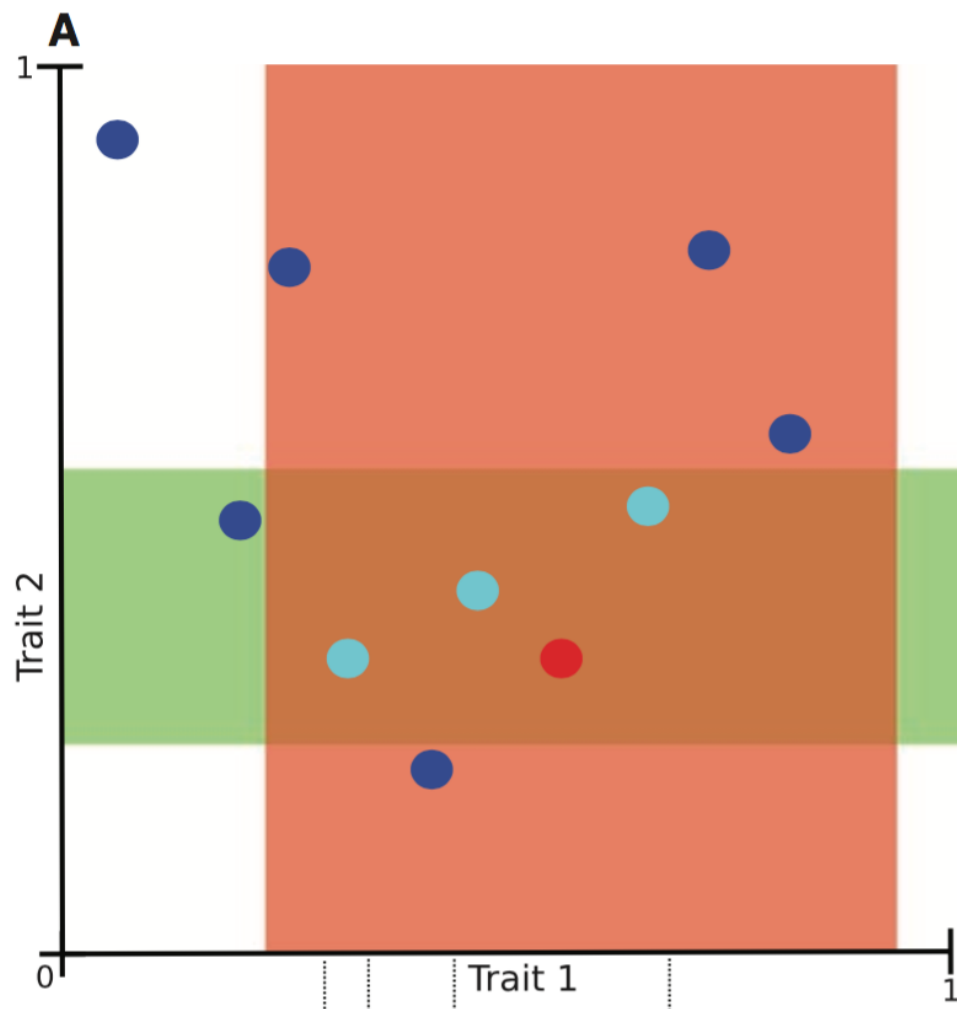


Estimated link probability ●

Measured interaction ○



How many dimensions are required?



Fraction of links correctly predicted

tend to be
larger webs

tend to be
smaller webs

Fitting models to data...

The Log-Ratio Model (LRM)

Better results, particularly for systems with strong body size constraints (large mammal communities, marine food webs)

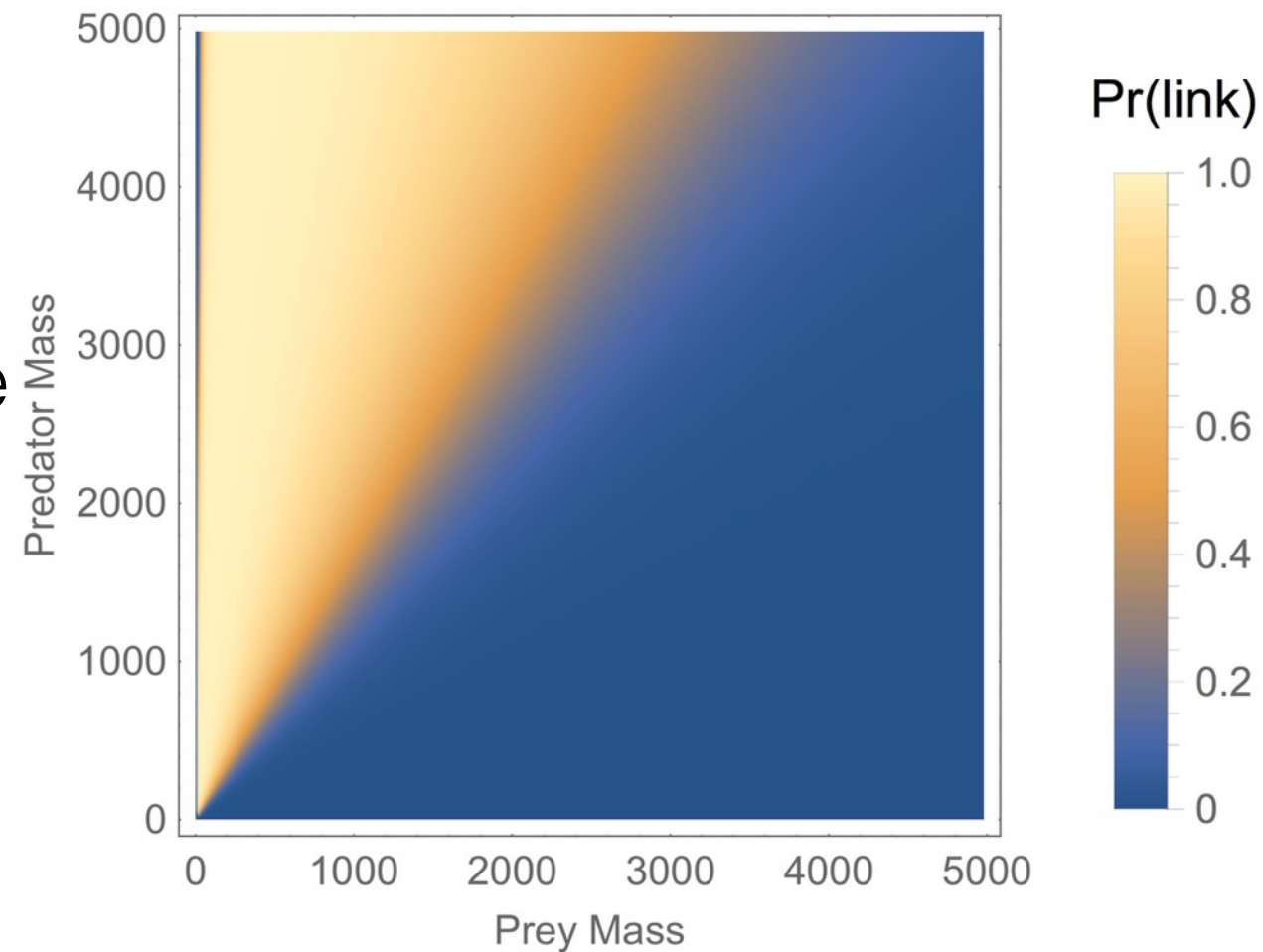
m_i = predator mass

m_j = prey mass

$$\log \left[\frac{P(a_{ij} = 1)}{P(a_{ij} = 0)} \right] = \alpha + \beta \log \left(\frac{m_i}{m_j} \right) + \gamma \log^2 \left(\frac{m_i}{m_j} \right),$$

Interaction probabilities modeled as a Logit regression

Quadratic term allows interaction probabilities to have a Gaussian-like shape



What about the parasites???



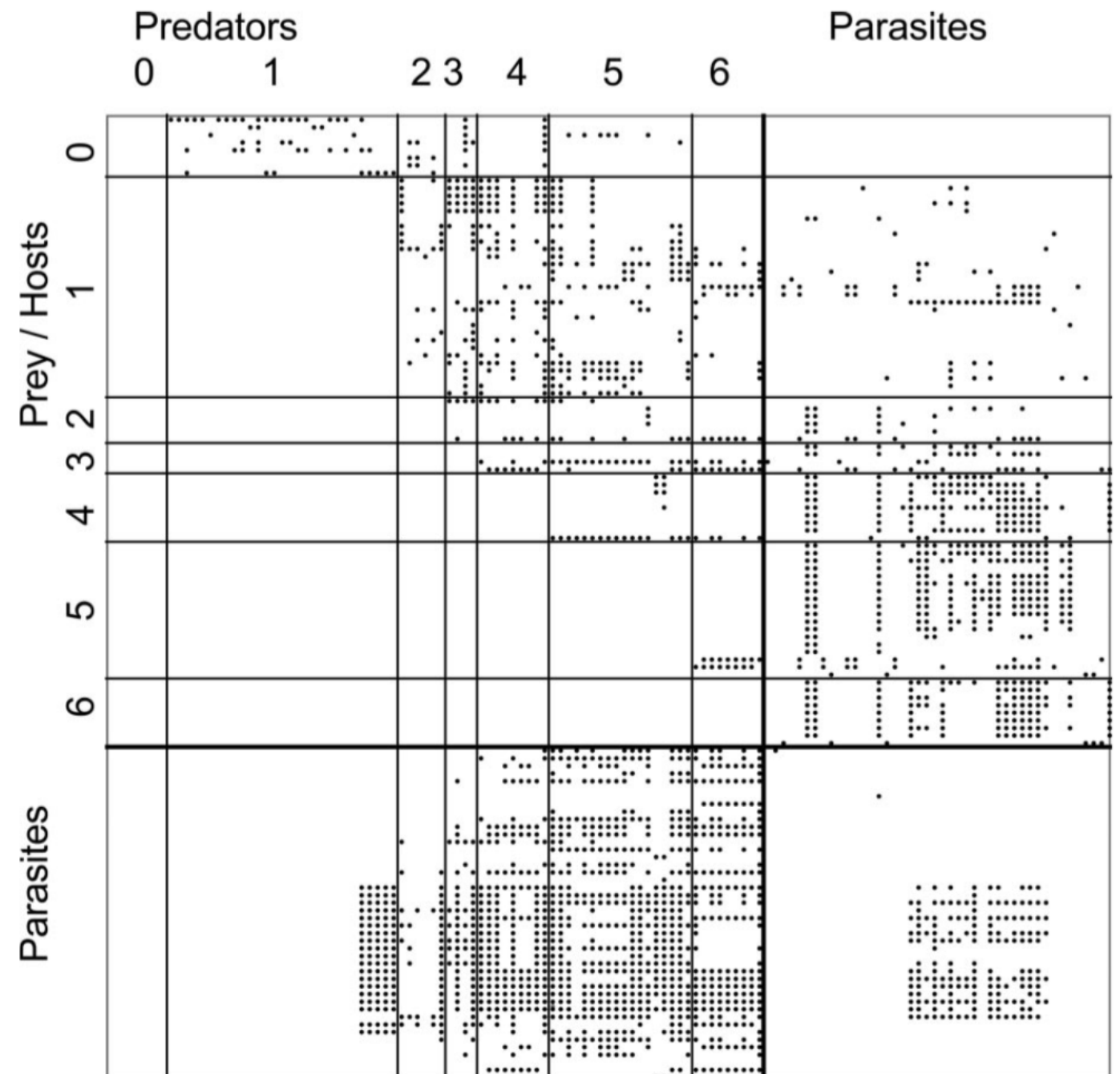
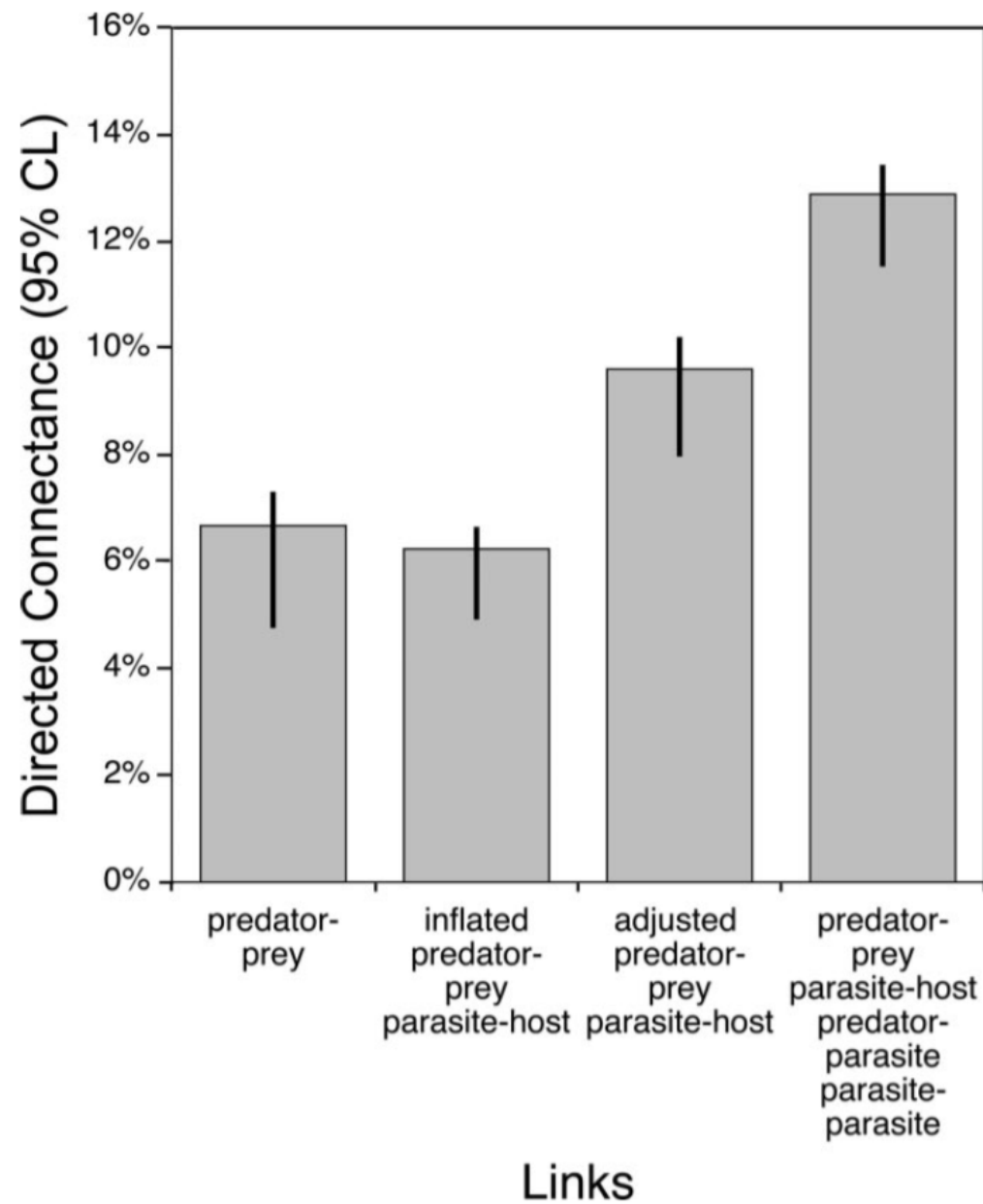
A qualitatively different
(+/-) interaction



What about the parasites???

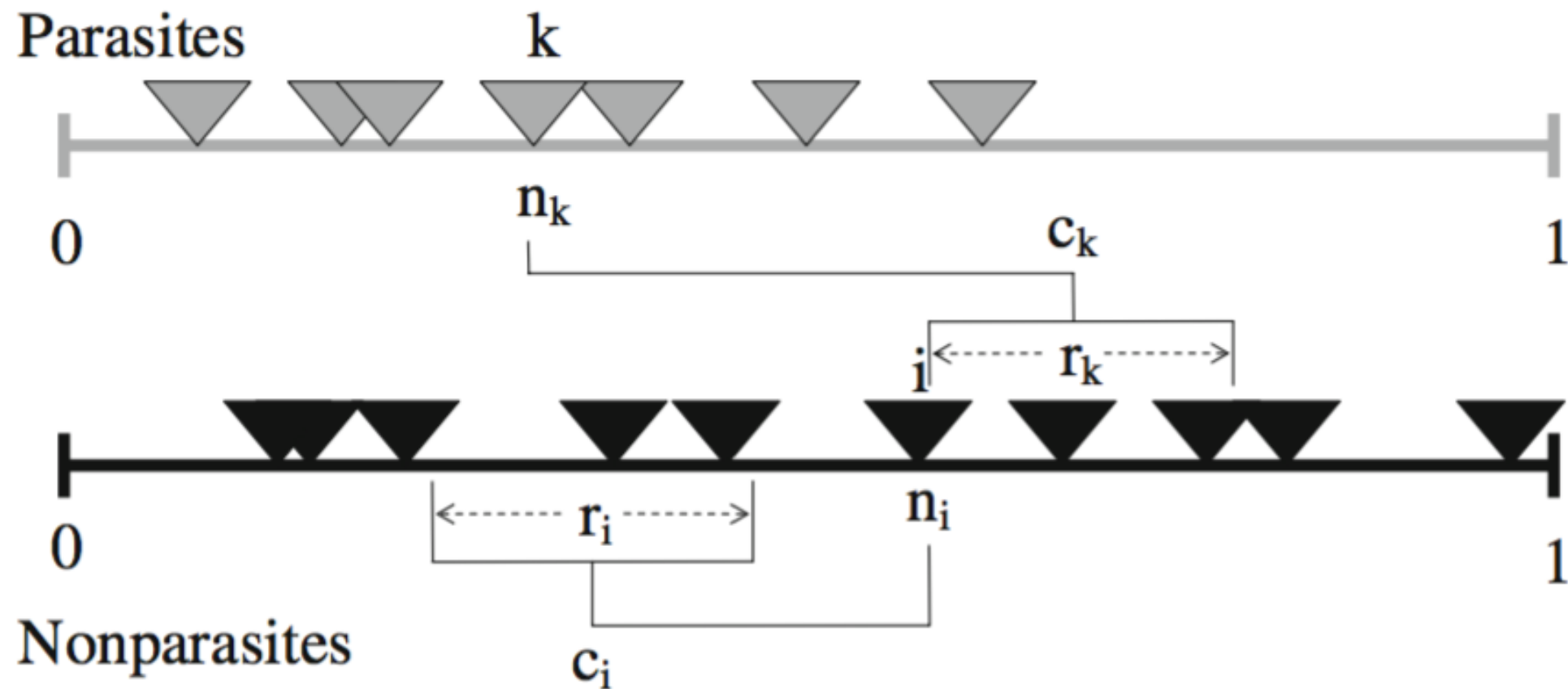


Parasites may dominate food web links



Lafferty et al. 2006

What about the parasites???



Higher trophic level parasites have **greater** specialization

“...the inclusion of parasites pushes the overall food web structure away from the niche model’s expectation.”

Warren et al. 2010

Dynamics on food webs

Linear Stability

$$\frac{d}{dt}x = f(x), \quad x^* \text{ is fixed point}$$

$$x(t) = x^* + \eta(t)$$

$$\eta(t) = x(t) - x^*$$

$$\frac{d}{dt}\eta(t) = \frac{d}{dt}(x(t) - x^*) = \frac{d}{dt}x(t)$$

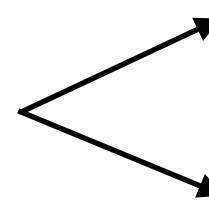
$$\frac{d}{dt}\eta(t) = \frac{d}{dt}x(t) = f(x) = f(x^* + \eta)$$

$$f(x^* + \eta) = f(x^*) + \eta f'(x^*) + \text{h.o.t.}$$

$$\frac{d}{dt}\eta = \eta \left. \frac{\partial f}{\partial x} \right|_{x^*}$$

$$\eta = \exp \left(\left. \frac{\partial f}{\partial x} \right|_{x^*} t \right)$$

$$\lambda = \left. \frac{\partial f}{\partial x} \right|_{x^*}$$

 >0 unstable
 <0 stable

Dynamics on food webs

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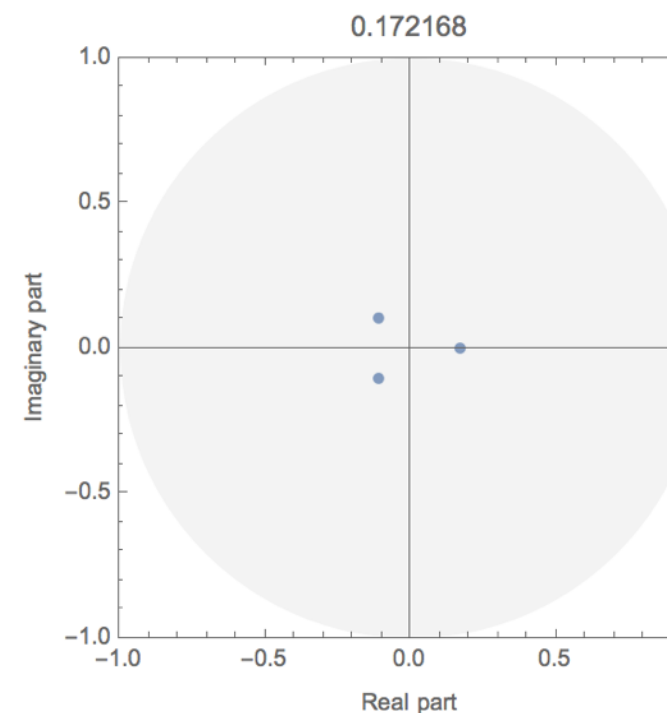
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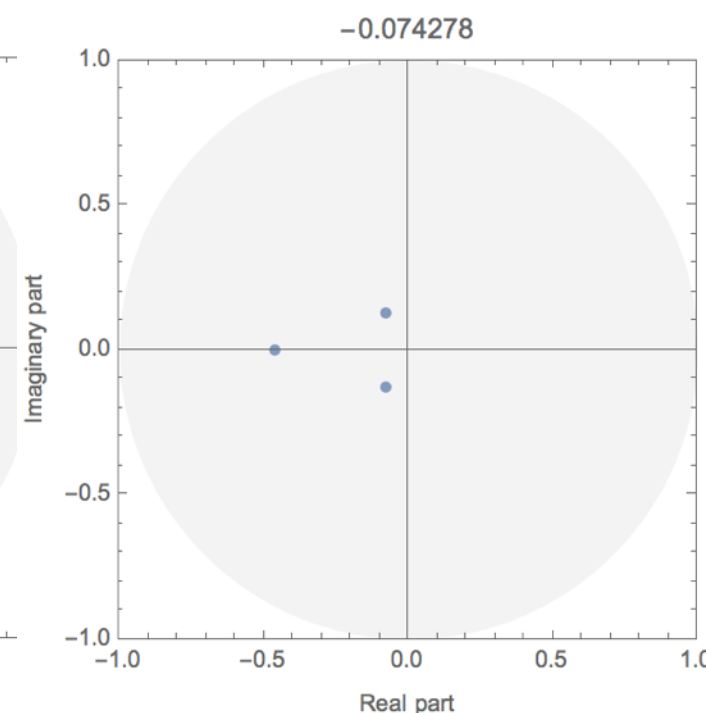
>0 unstable
 <0 stable

Multi-dimensional Systems

$$\mathbf{J}|_* = \begin{pmatrix} \left. \frac{\partial F}{\partial X} \right|_* & \left. \frac{\partial F}{\partial Y} \right|_* & \left. \frac{\partial F}{\partial Z} \right|_* \\ \left. \frac{\partial G}{\partial X} \right|_* & \left. \frac{\partial G}{\partial Y} \right|_* & \left. \frac{\partial G}{\partial Z} \right|_* \\ \left. \frac{\partial H}{\partial X} \right|_* & \left. \frac{\partial H}{\partial Y} \right|_* & \left. \frac{\partial H}{\partial Z} \right|_* \end{pmatrix}$$



unstable

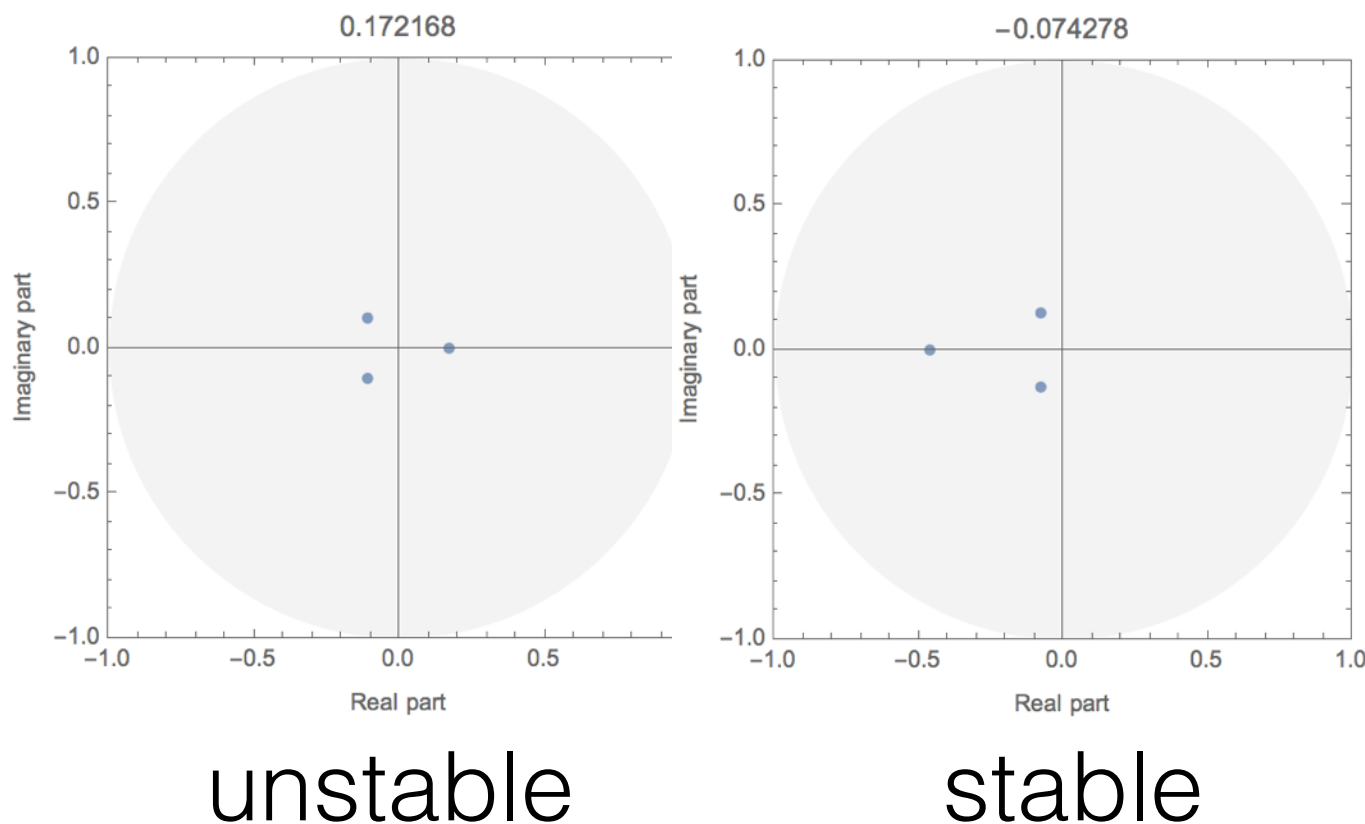


stable

Dynamics on food webs

Multi-dimensional Systems

$$\mathbf{J}|_* = \begin{pmatrix} \left. \frac{\partial F}{\partial X} \right|_* & \left. \frac{\partial F}{\partial Y} \right|_* & \left. \frac{\partial F}{\partial Z} \right|_* \\ \left. \frac{\partial G}{\partial X} \right|_* & \left. \frac{\partial G}{\partial Y} \right|_* & \left. \frac{\partial G}{\partial Z} \right|_* \\ \left. \frac{\partial H}{\partial X} \right|_* & \left. \frac{\partial H}{\partial Y} \right|_* & \left. \frac{\partial H}{\partial Z} \right|_* \end{pmatrix}$$



Trophic interactions on random graphs

‘Will a large complex system be stable?’

(May 1972)

M = SxS adjacency matrix

Diag(**M**) = -1

offDiag(**M**) ~ Norm(0, σ^2) w/ Pr(C)

offDiag(**M**) = 0 w/ Pr(1-C)

Big finding:

Pr(Stability) \rightarrow 0 when

$$\sigma \sqrt{SC} > 1$$

big, complex ecosystems shouldn't exist

Different types of interactions in the food web

Trophic interactions on
random graphs

‘Will a large complex
system be stable?’

(May 1972)

\mathbf{M} = $S \times S$ adjacency matrix

$\text{Diag}(\mathbf{M}) = -1$

$\text{offDiag}(\mathbf{M}) \sim \text{Norm}(0, \sigma^2)$ w/ $\text{Pr}(C)$

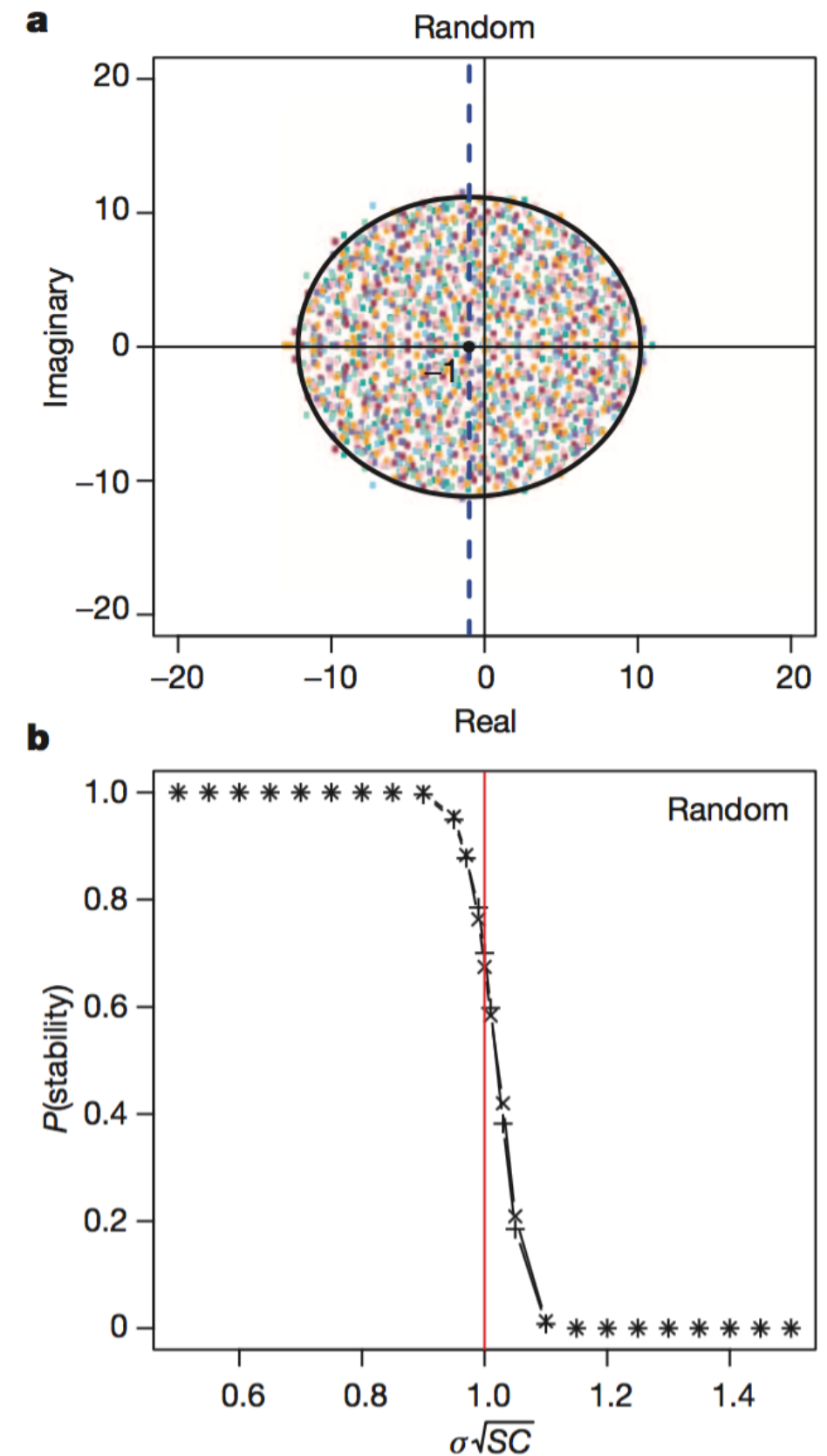
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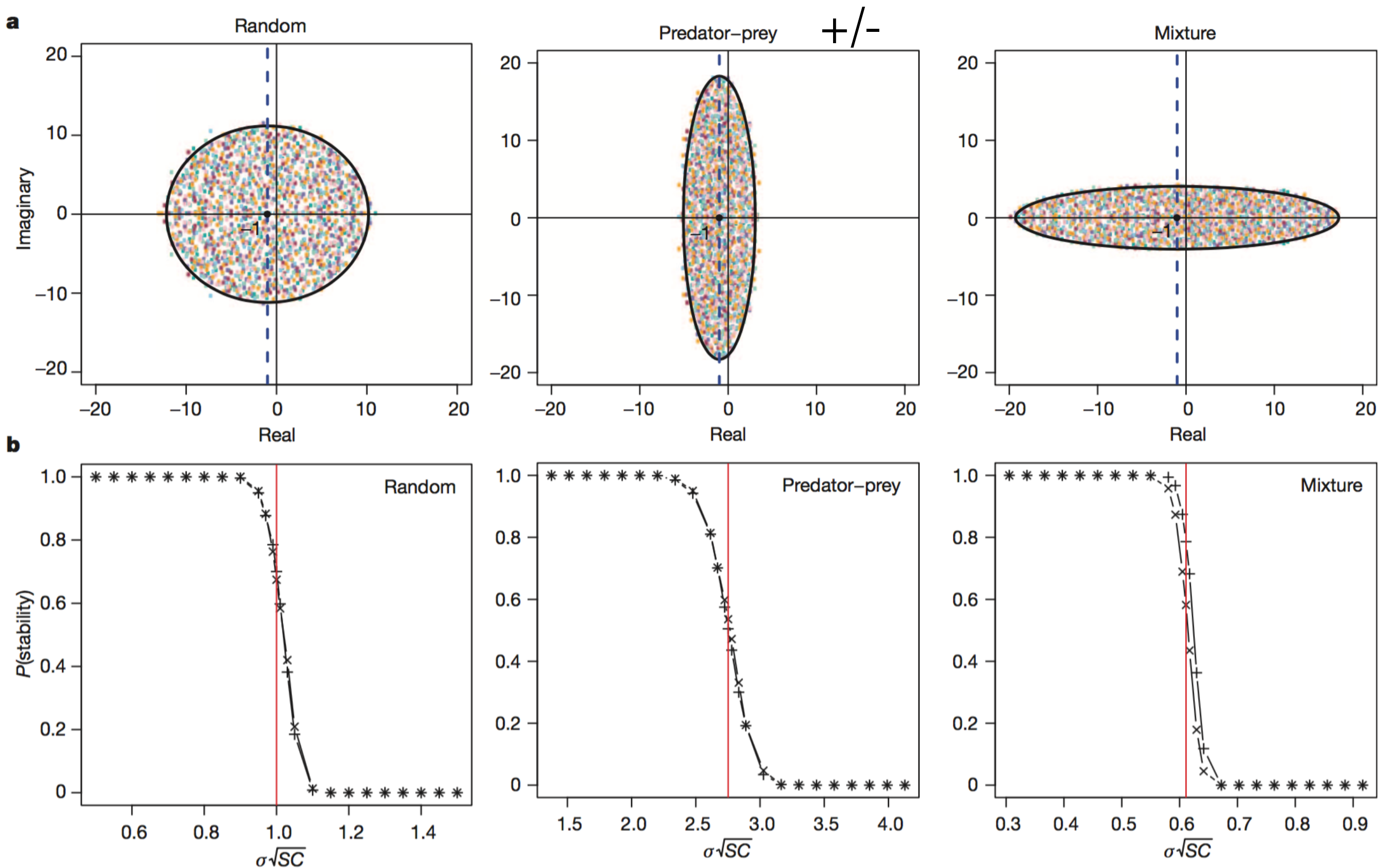
$$\sigma \sqrt{SC} > 1$$

**big, complex ecosystems
shouldn't exist**

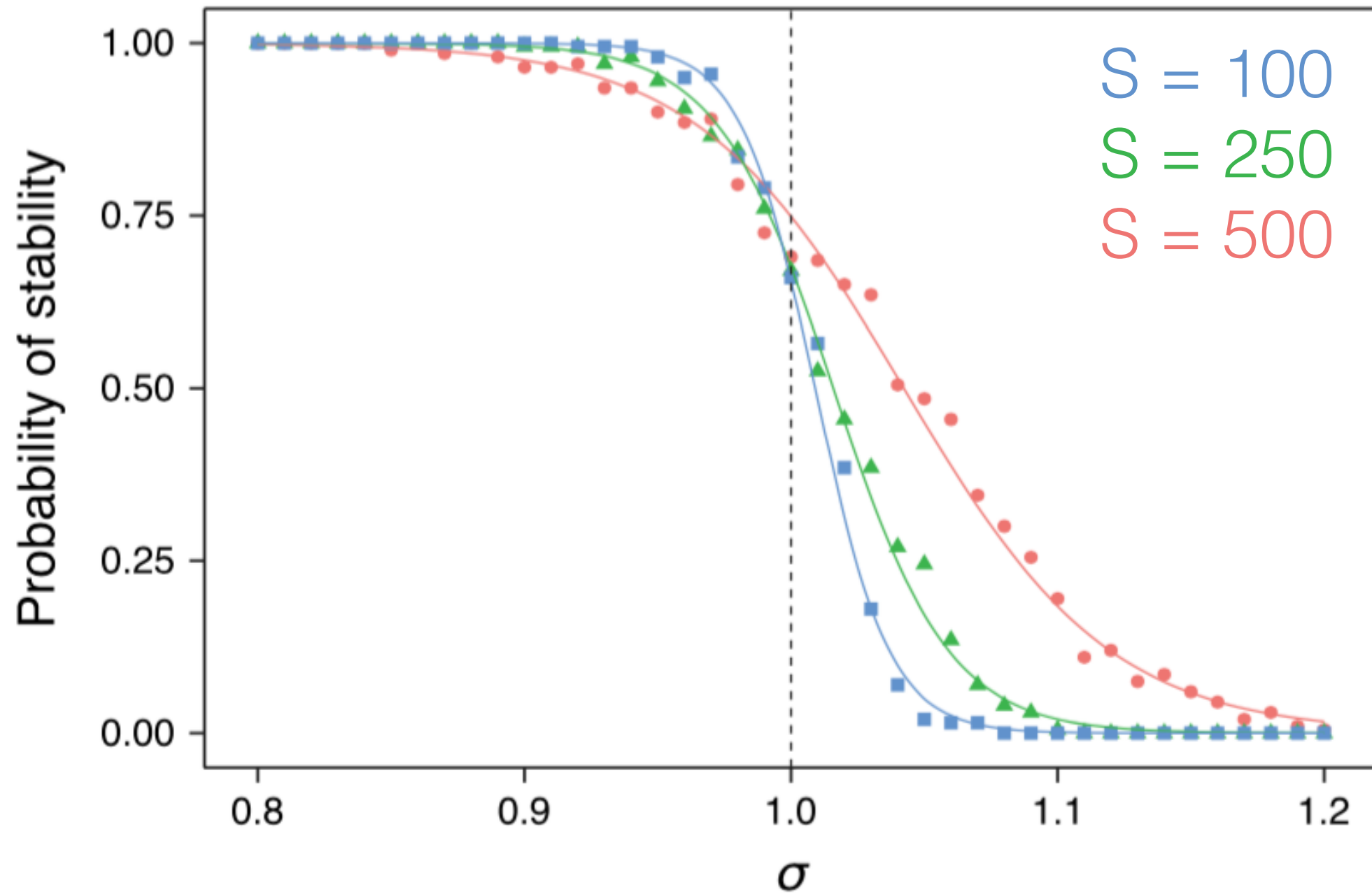


Different types of interactions in the food web

predator/prey: +/-
mutualism: +/+



Across different values of σ
(interaction strength variability)



Random matrix approaches to food webs
may be missing some important constraints

Dynamic modeling via simulations to
explore things like **adaptive foraging**

Energetics:
Optimal/Adaptive foraging

search



acquire



consume



— gains vs. costs —

Adaptive foraging and food webs

Realism vs. Complexity

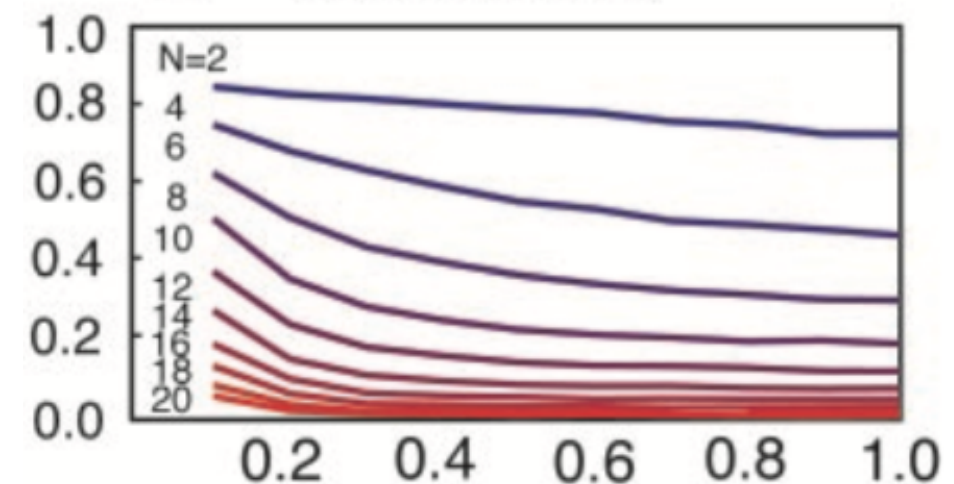
$$\frac{d}{dt}X_i = X_i \left(\underset{\substack{\uparrow \\ \text{growth rate}}}{r_i} - \underset{\substack{\uparrow \\ \text{self-regulation intensity}}}{s_i}X_i + \sum_{j \in \text{res.}} \underset{\substack{\uparrow \\ \text{met. rate}}}{e_{ij}} \underset{\substack{\uparrow \\ \text{efficiency}}}{f_{ij}} \underset{\substack{\uparrow \\ \text{foraging effort of consumer } j \text{ on resource } i}}{a_{ij}}X_j - \sum_{j \in \text{cons.}} f_{ji}a_{ji}X_j \right)$$

foraging effort of consumer j on resource i

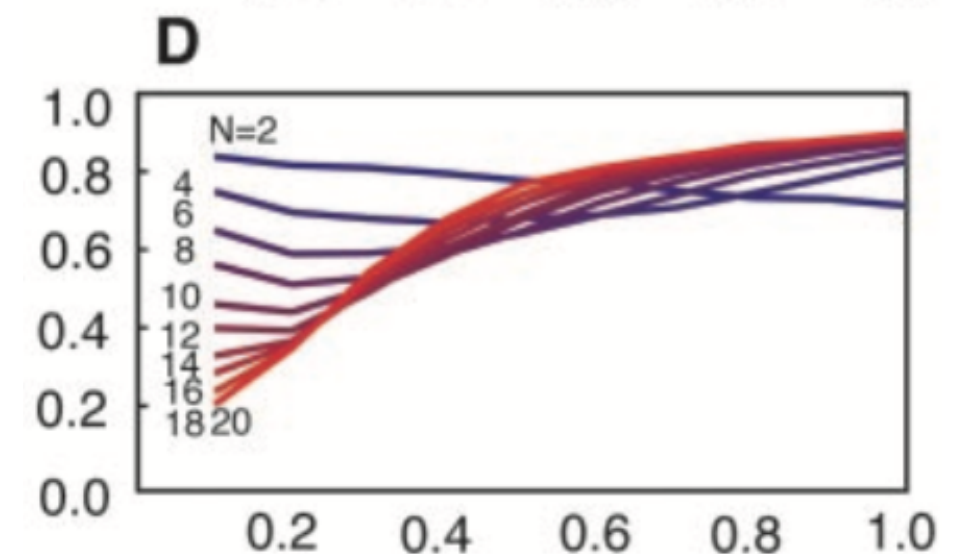
The effort is itself dynamic, and changes in response to changing resource densities

If resource profitability (i.e. energy gain per unit effort) is higher than an 'average profitability', the effort increases

w/o a



w/ a



Kondoh 2003

Realism vs. Complexity

And intermediate between
specific models and random matrix theory

Generalized modeling

$$\frac{d}{dt}B = \overset{\text{(Source)}}{S(B)} - \overset{\text{(Drain)}}{D(B)}$$

Recruitment Mortality

Keep it GENERAL...

More 'accurate' wrt our knowledge

Can't simulate or solve for the f.p.

Gross et al. 2006

Yeakel et al. 2011

Establish B^* as a variable representing all *internal* equilibria of the system.

Build a new set of parameters representing the normalized variables of the generalized system:

$$\frac{d}{dt}B = S(B) - D(B)$$

$$b := \frac{B}{B^*}, \quad s(b) := \frac{S(B)}{S(B^*)}, \quad \text{and} \quad d(b) := \frac{D(B)}{D(B^*)}$$

$$\frac{d}{dt}b = \frac{S^*}{B^*}s(b) - \frac{D^*}{B^*}d(b)$$

Rewrite in terms of b

$$\gamma = \frac{S^*}{B^*} = \frac{D^*}{B^*}$$

$$\frac{d}{dt}b = \gamma(s(b) - d(b)) \quad \textbf{PERTURB}$$

↻ biomass turnover rate

Gross et al. 2006

Yeakel et al. 2011

Stability of the generalized model

$$\frac{\partial}{\partial b} \dot{b} = \lambda = \frac{\partial}{\partial b} \gamma (s(b) - d(b))$$

$$\frac{\partial}{\partial b} s(b) = s_b = \left. \frac{\partial \log S}{\partial \log B} \right|_* \sim \frac{\% \text{ change in } S}{\% \text{ change in } B}$$

Functional elasticities are the logarithmic derivatives

**Provides a nonlinear measure for the sensitivity of the function to variations in biomass

$$\lambda = \underset{\text{Elasticity of growth}}{\gamma} \left(\underset{\text{Elasticity of mortality}}{s_b} - d_b \right)$$

Gross et al. 2006

Yeakel et al. 2011

Elasticities characterize whole families of functional forms

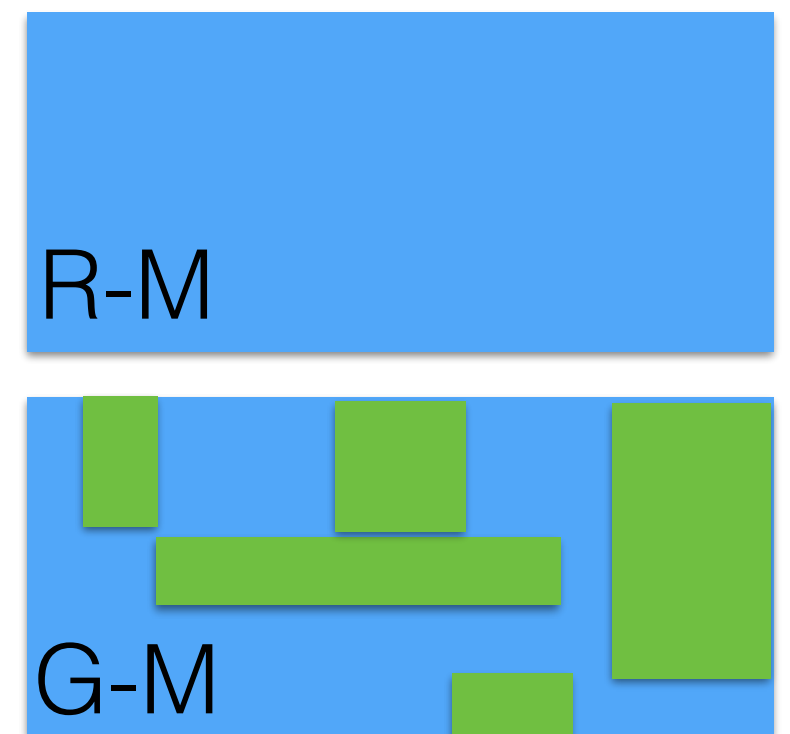
$F(X) = c_1$	$\frac{\partial f(x)}{\partial x} = 0$	Constant	(independent gain / loss)
$F(X) = c_2 X$	$\frac{\partial f(x)}{\partial x} = 1$	Linear	(intrinsic growth)
$F(X) = -c_3 X^2$	$\frac{\partial f(x)}{\partial x} = 2$	Superlinear	(self limitation)

Additional benefits

- Sets realistic bounds to parameters that will go into the Jacobian**
- Depend on the state of the system at the time a measurement is made
- Error tolerant

More complex functions have elasticities as functions of the steady state

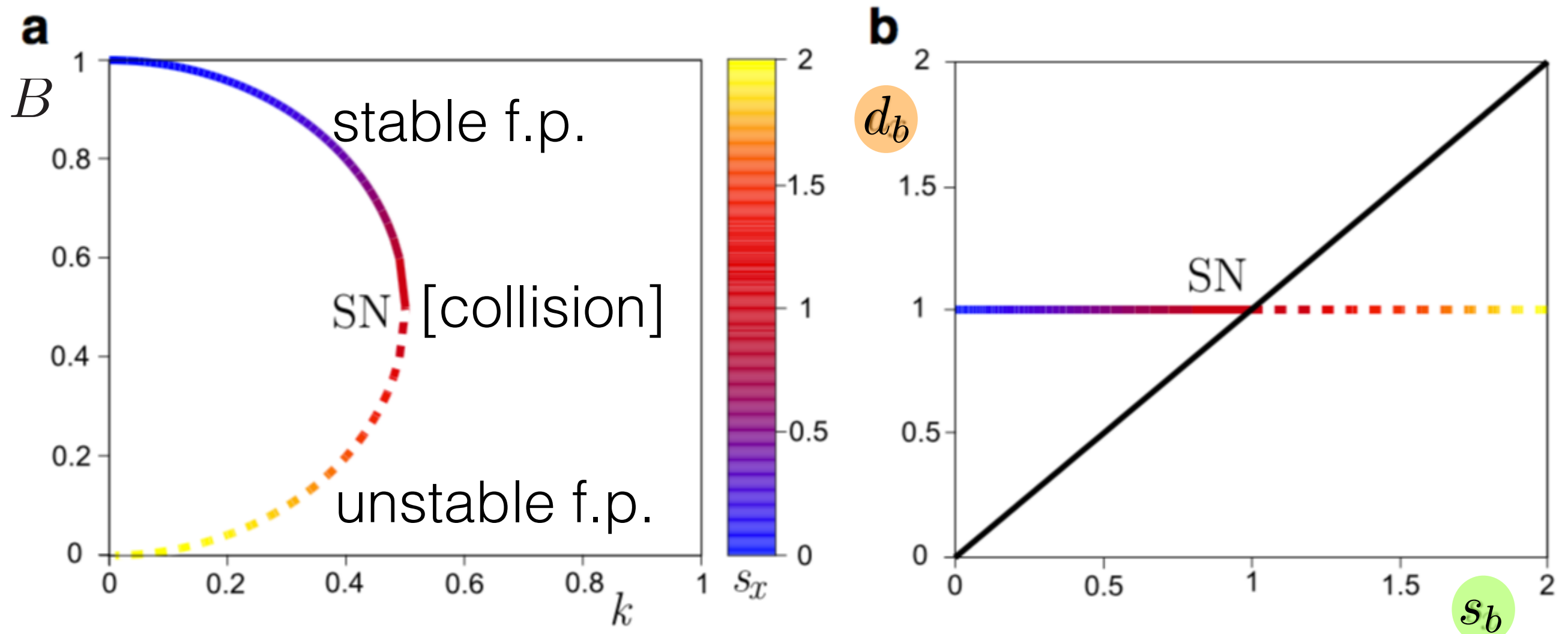
Gross et al. 2006
Yeakel et al. 2011



$$\frac{d}{dt}B = \frac{aB^2}{k^2 + B^2} - mB \quad \frac{d}{dt}B = S(B) - D(B)$$

k = half-saturation value of growth

$$\lambda = \frac{\partial}{\partial b} \gamma (s(b) - d(b)) = \gamma (s_b - d_b)$$



Realism vs. Complexity

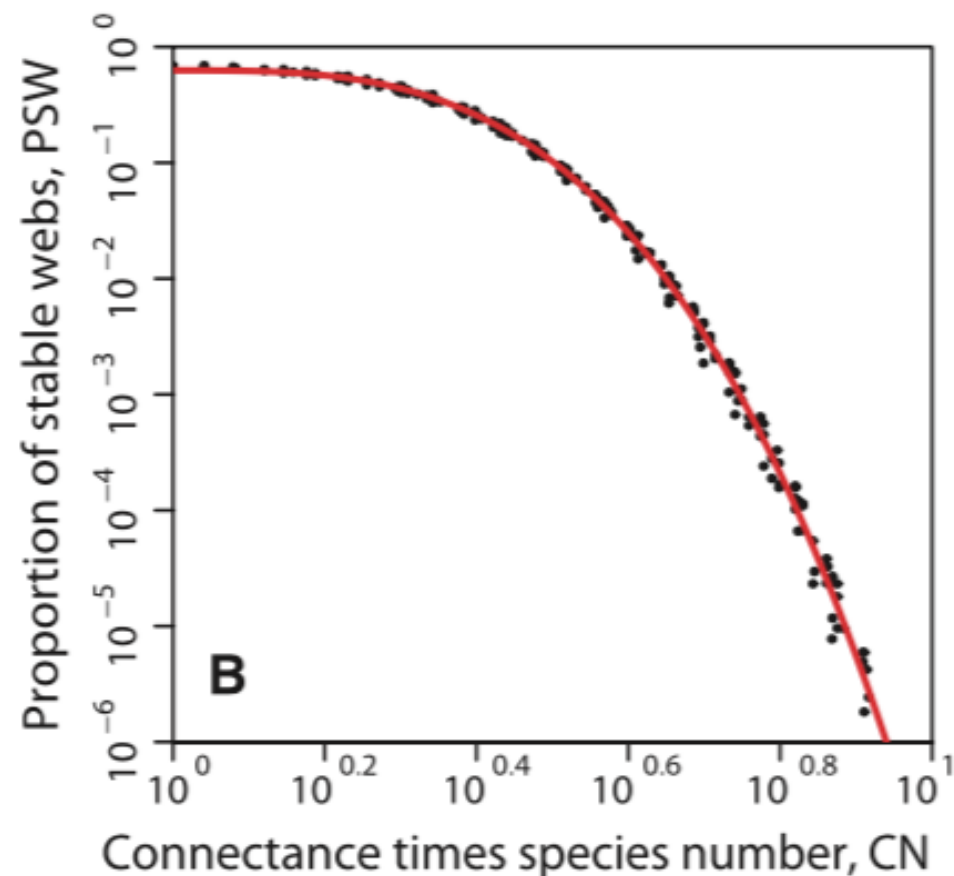
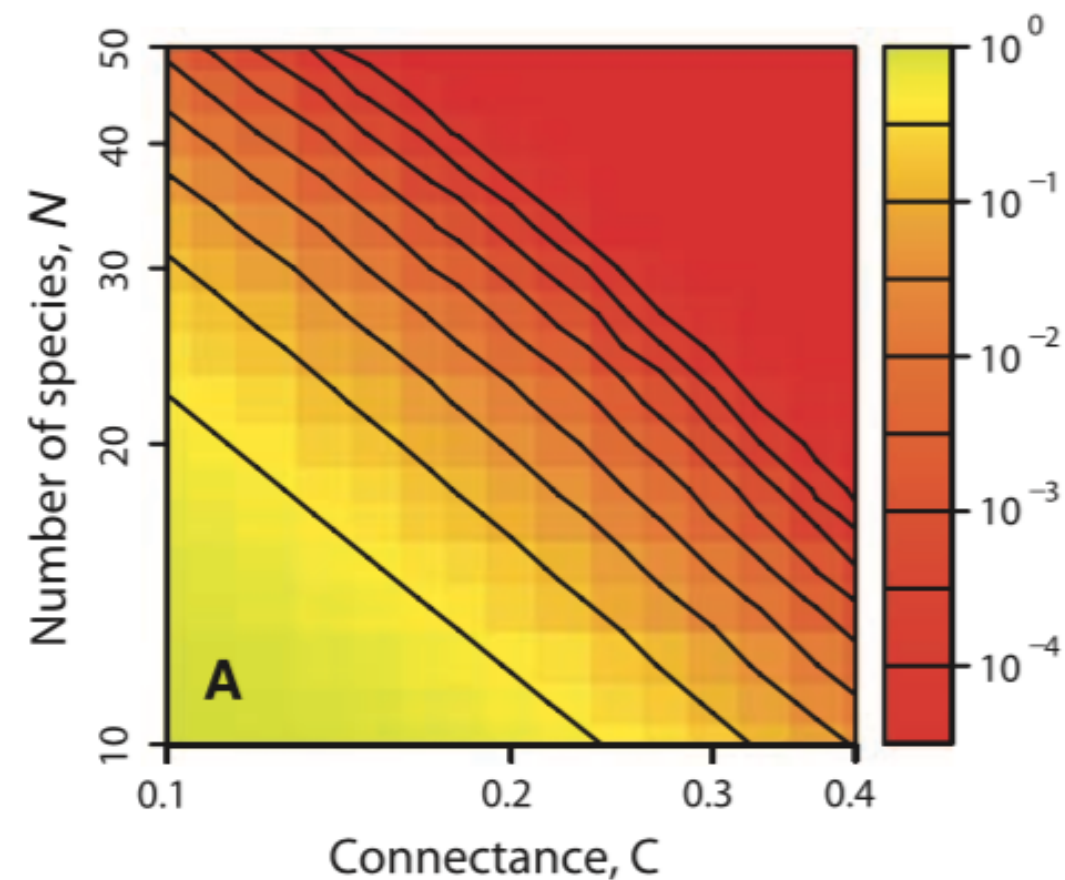
And intermediate between
specific models and random matrix theory

Generalized modeling on a larger scale

$$\dot{X}_i = S_i(X_i) + F_i(X_1, \dots, X_N) - M_i(X_i) - \sum_{j=1}^N G_{ij}(X_1, \dots, X_N)$$

Growth from 1° prod. Intrinsic mort. Mort. due to predation

Growth from pred.



Realism vs. Complexity

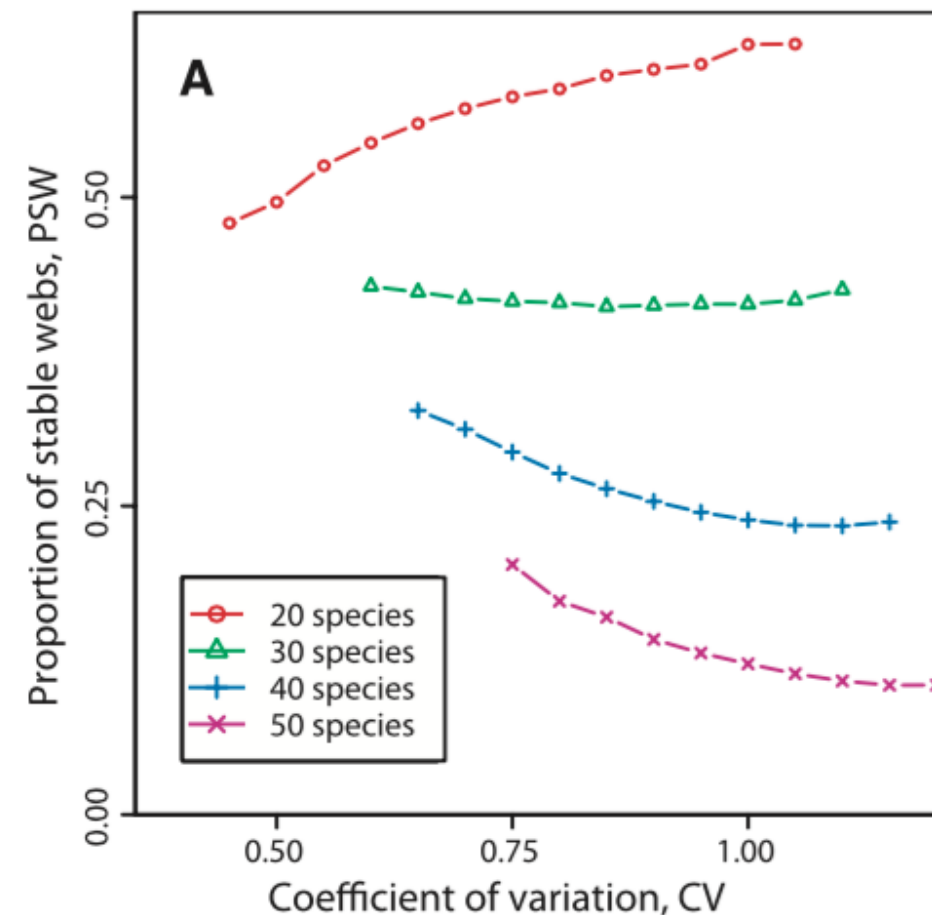
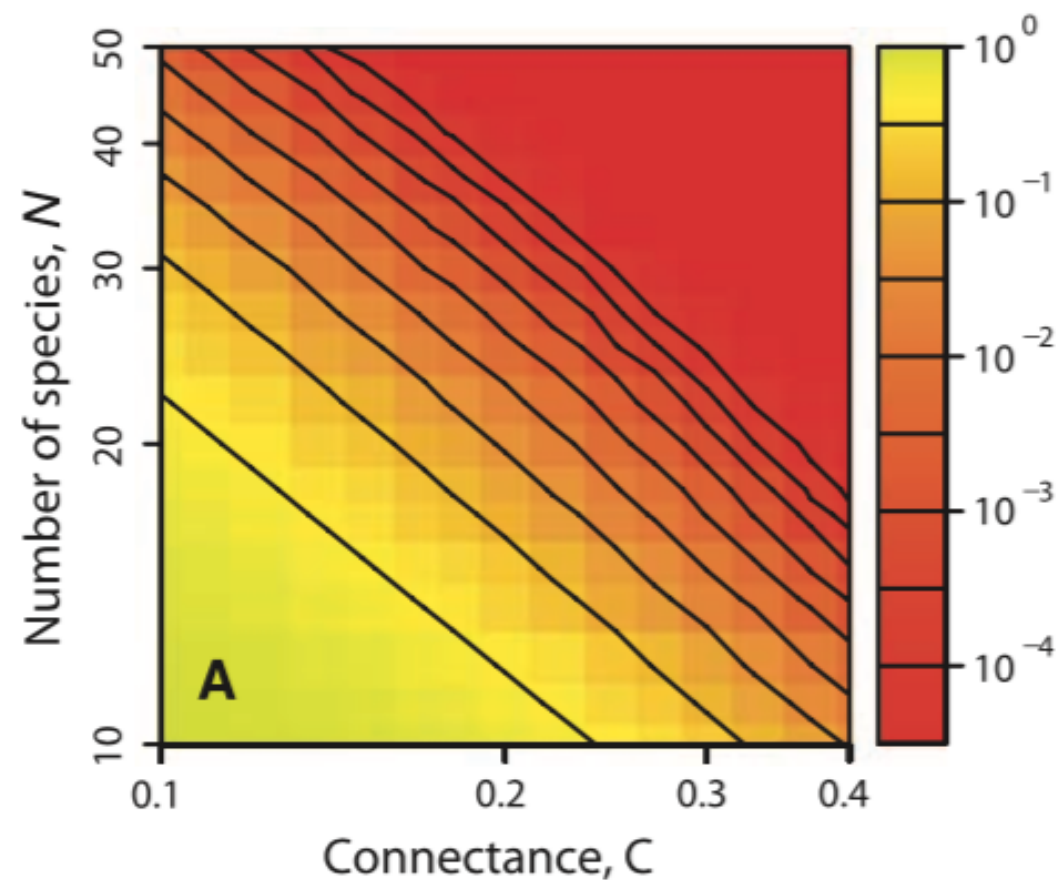
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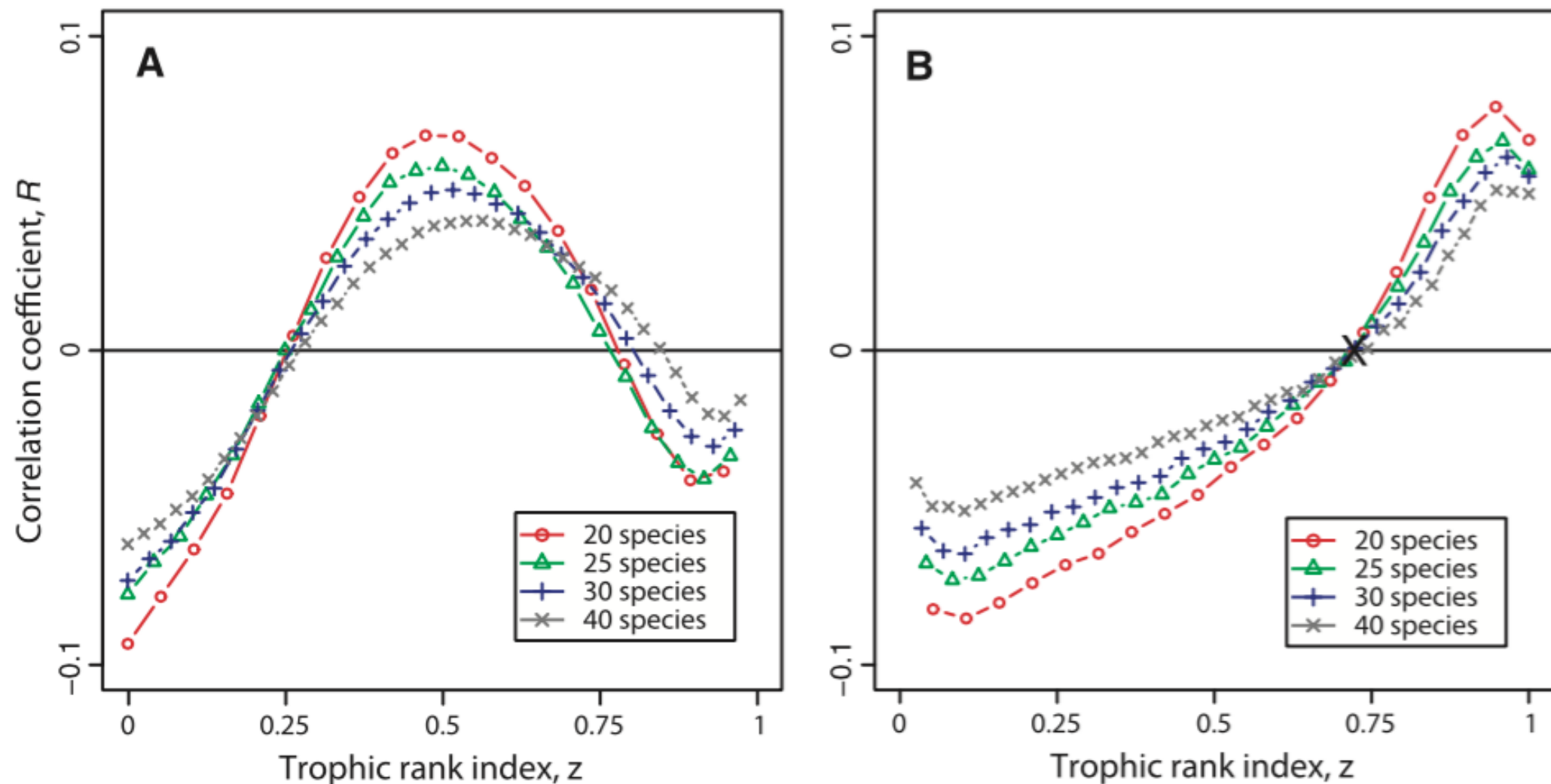
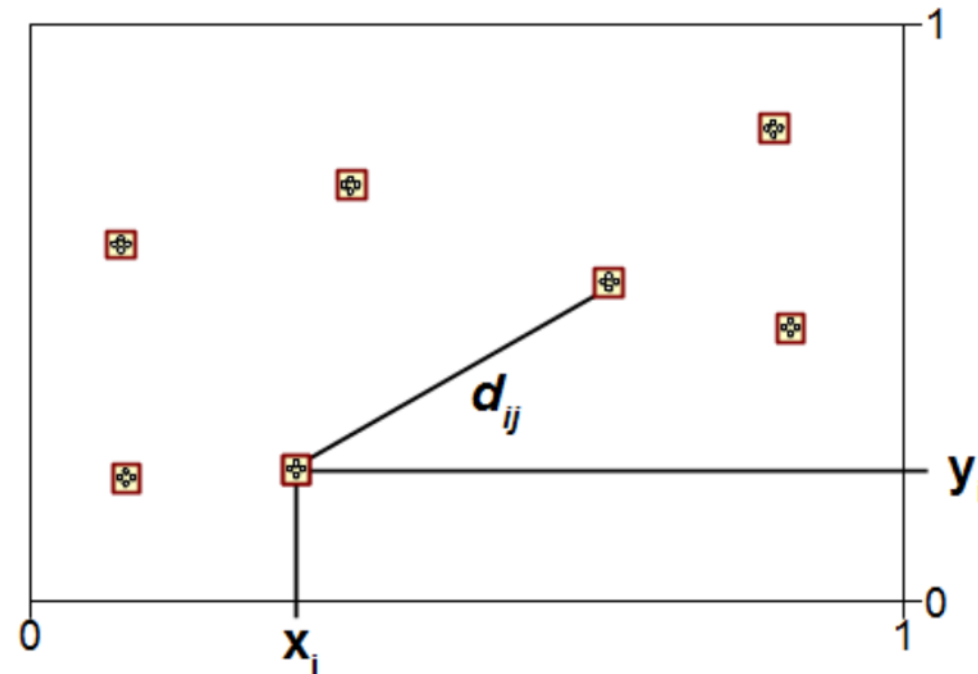


Fig. 4. Dependence of food-web stability on the distribution of links. **(A)** Correlation of stability with the number of predator species preying on a focal species, in dependence on the trophic position of the focal species as measured by its trophic-rank index z . Stability is enhanced if most species prey upon intermediate species, which are characterized by indices around $z = 0.5$. **(B)** Correlation of stability with the number of prey species predated upon by a focal species, in dependence on the trophic position of the focal species. Stability is enhanced if apical predators are generalists, whereas intermediate predators are specialists.

Food webs in SPACE

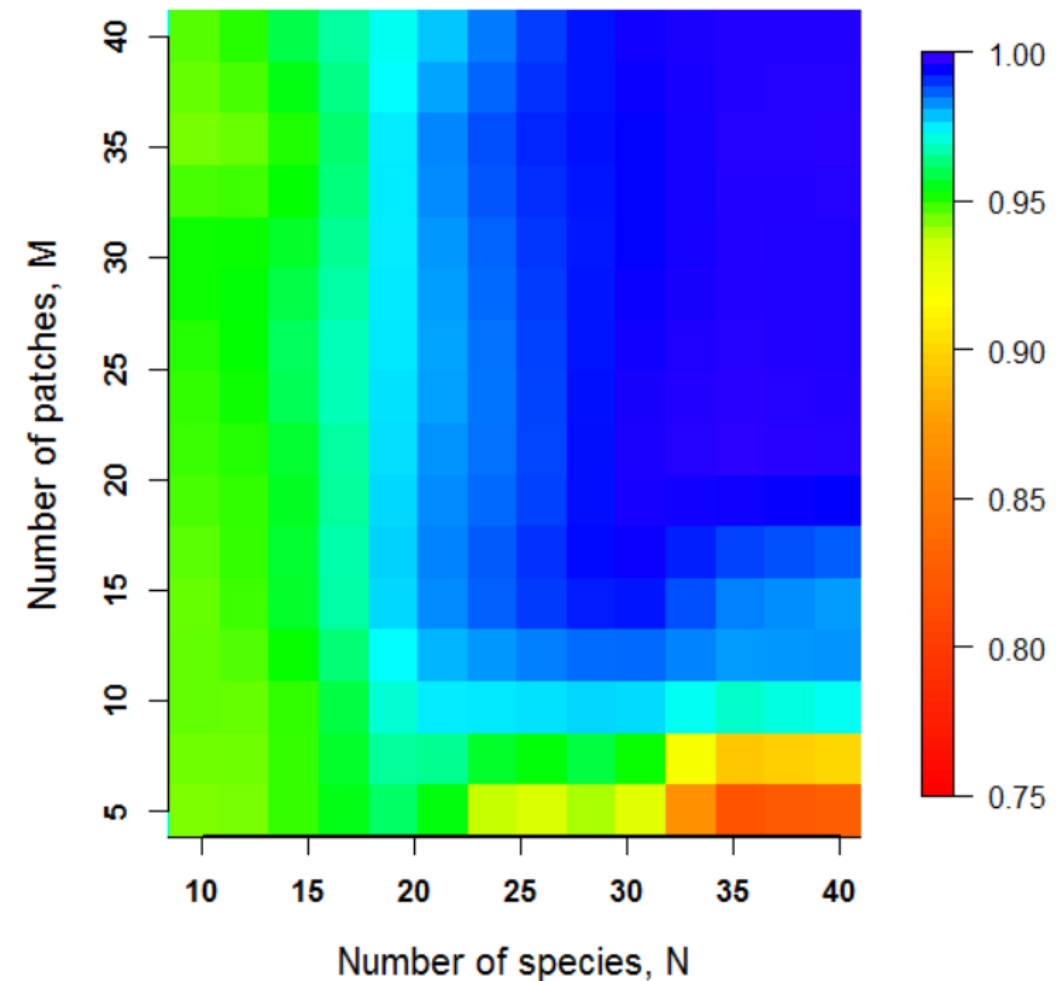
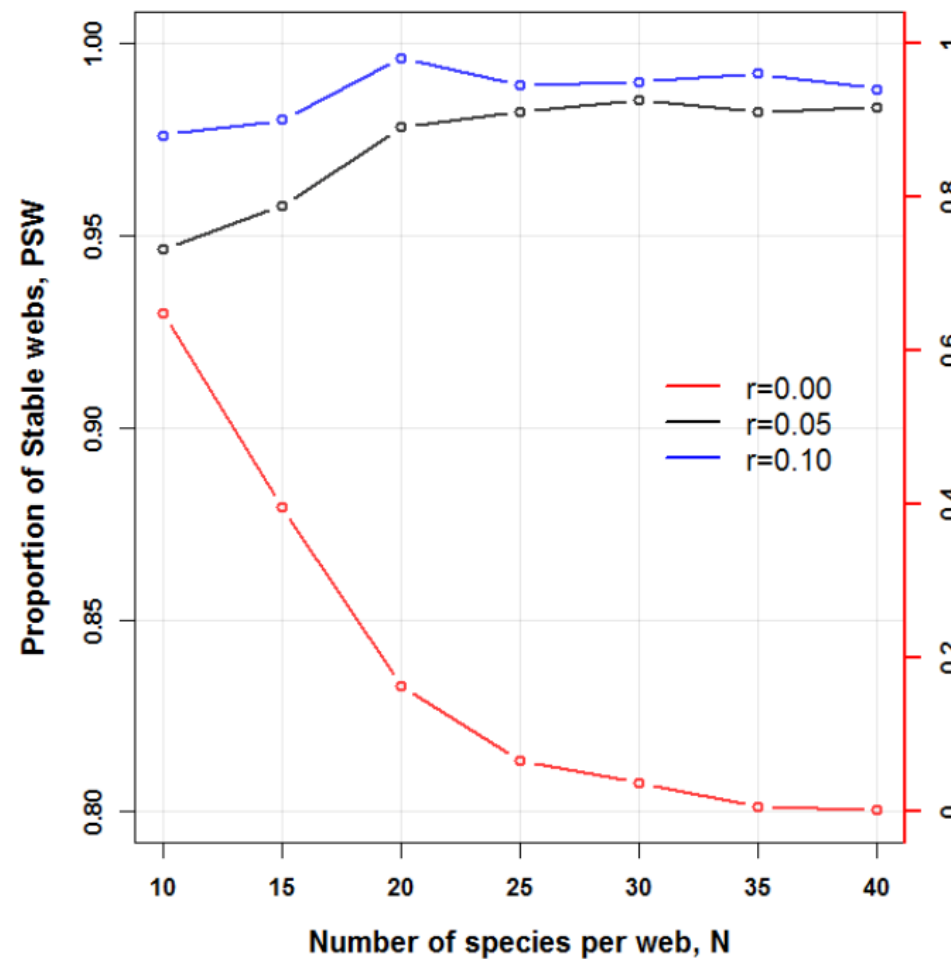


Lars
Rudolf



Thilo Gross

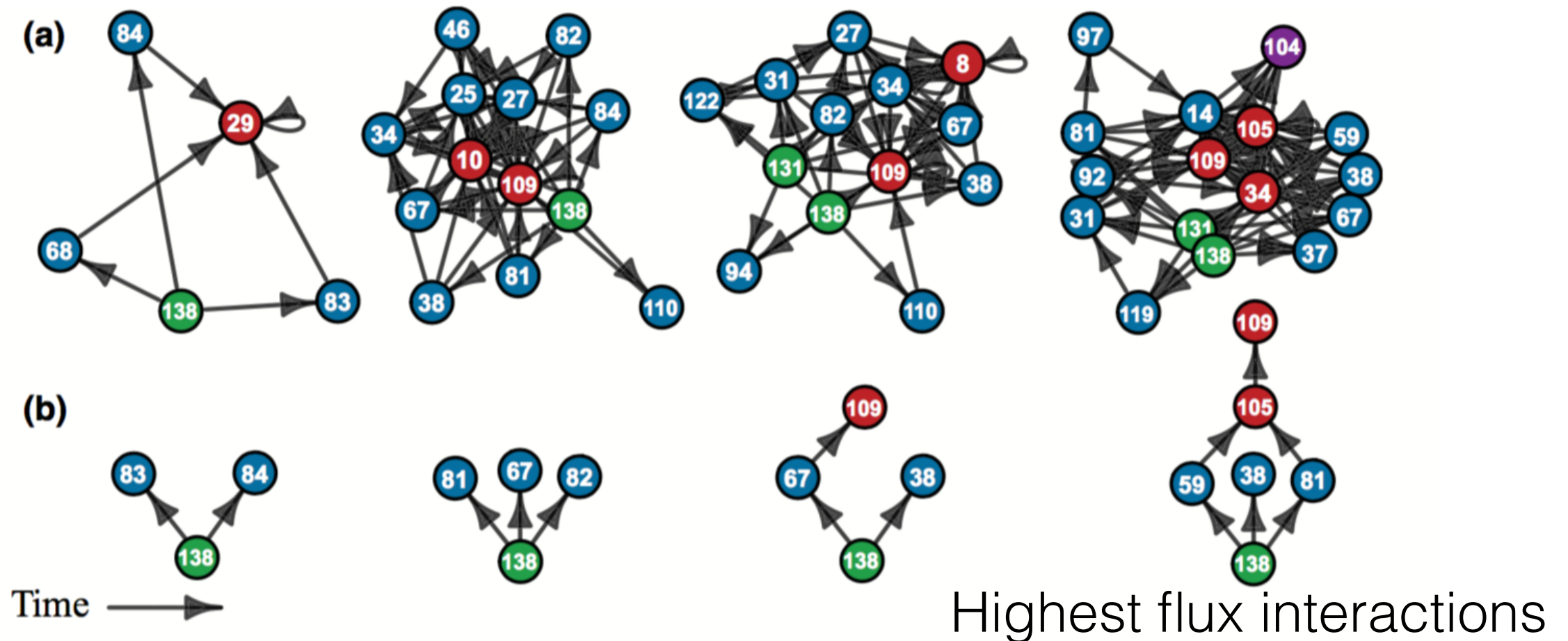
$r = \text{fraction of gain by immigration}$



PSW

Food webs in TIME

Exploring assembly from microcosm experiments



“Webs experiencing different colonisation rates had stable topologies despite significant species turnover, suggesting that some features of network architecture emerge early and change little through assembly. But webs experiencing low colonisation rates showed less variation in the magnitudes of trophic fluxes, and were less likely to develop coupled fast and slow resource channels – a common feature of published webs. **Our results reveal that food web structure develops according to repeatable trajectories that are strongly influenced by colonisation rate.**”

