

UQAR – Canada Research Chair in Biogeography and
metacommunity ecology

Ecological networks in the anthropocene

Dominique Gravel

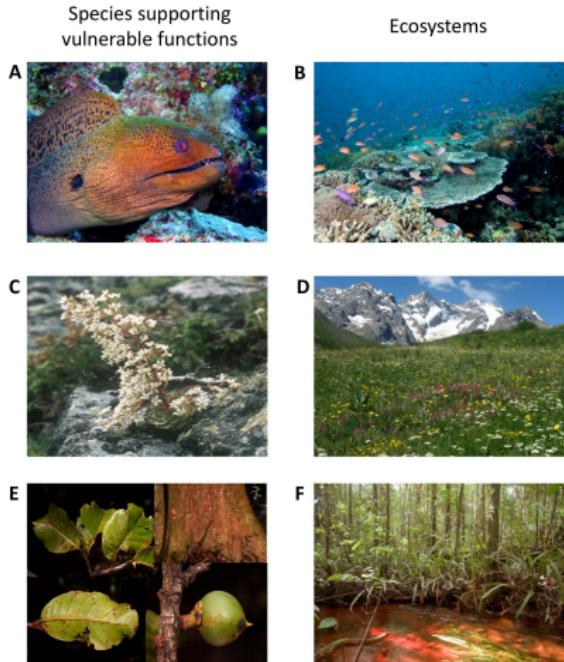
<http://www.chaire-eec.uqar.ca/>

April 9, 2014

**We need a novel approach to quantify biodiversity taking
into account the diversity of ecological interactions**



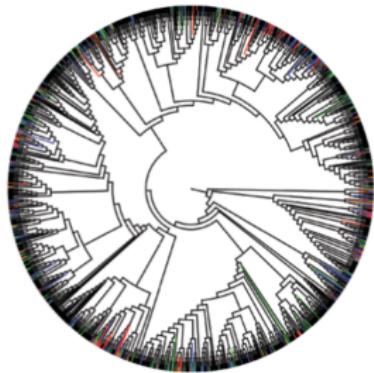
Trends in biogeography



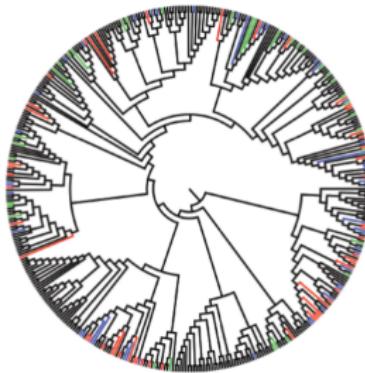
Mouillot et al. (2013)

Trends in biogeography

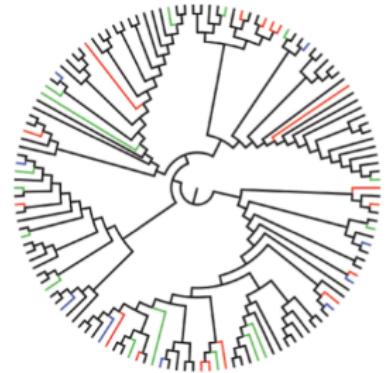
Plants



Birds

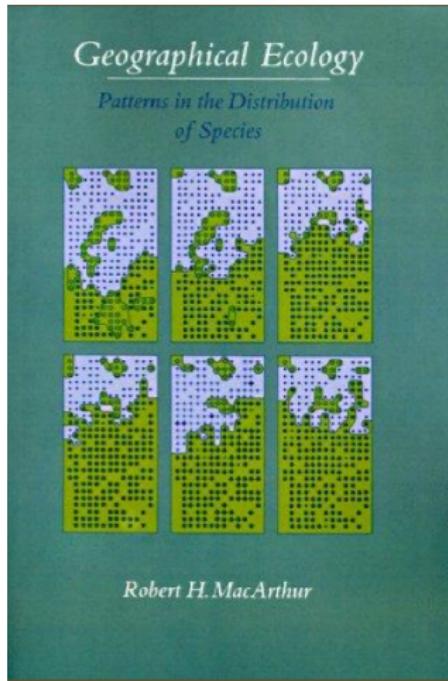


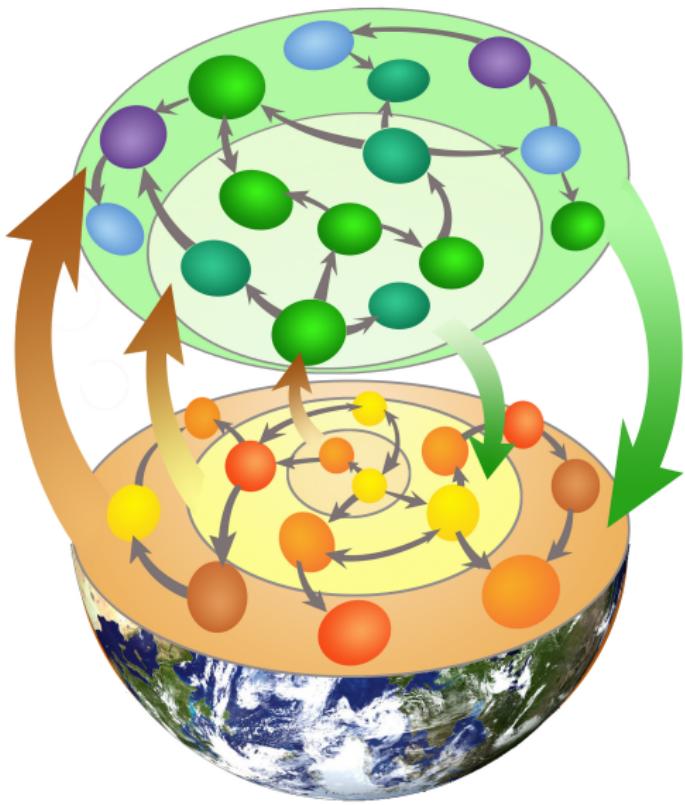
Mammals



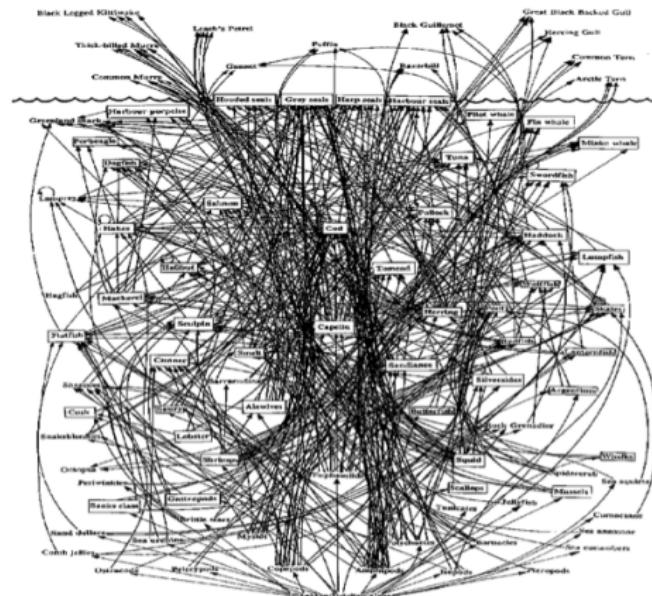
Thuiller et al. (2011)
Mazel et al. (2014)

The rich history of community ecology

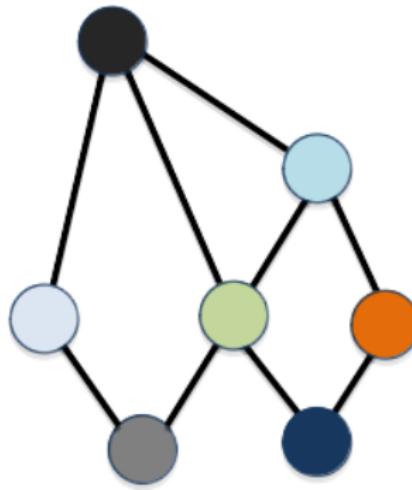




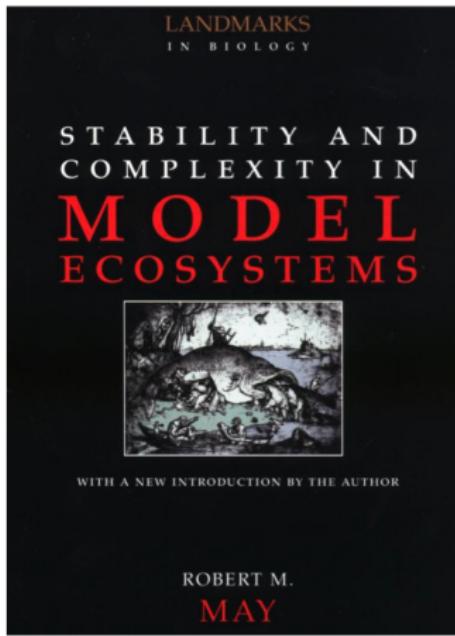
Communities are messy



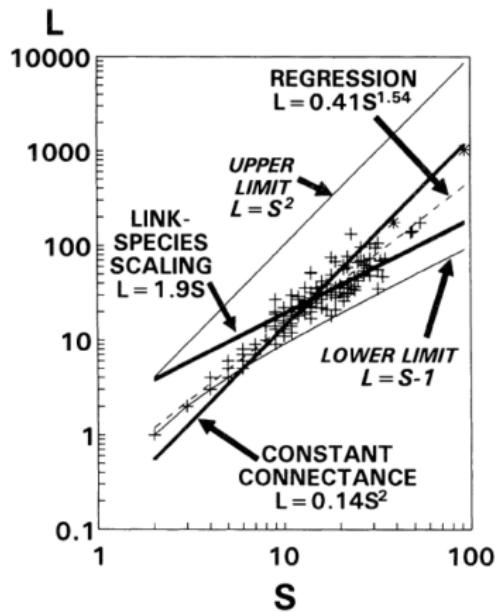
Compressing information



A successfull program

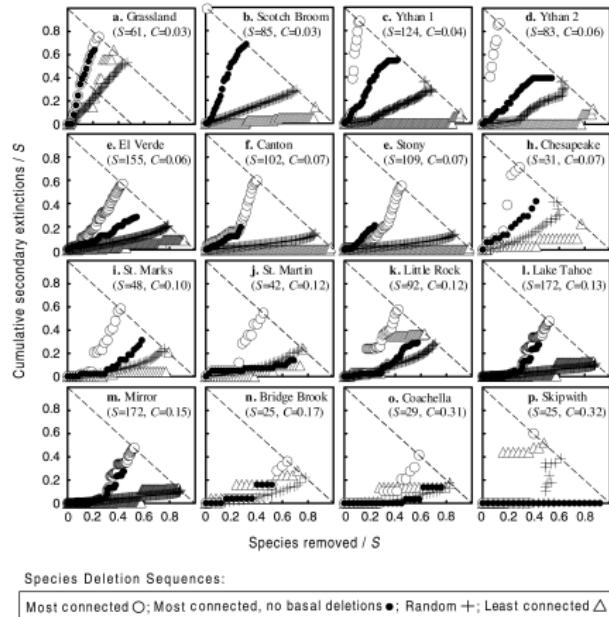


Compressing information



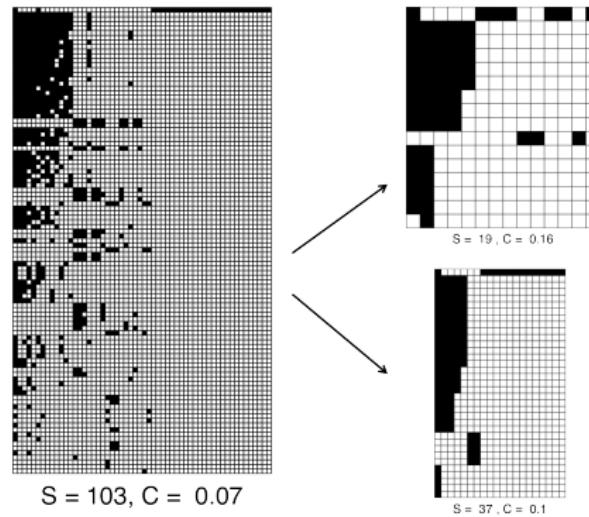
Martinez (1992)

Compressing information



Dunne et al. (2002)

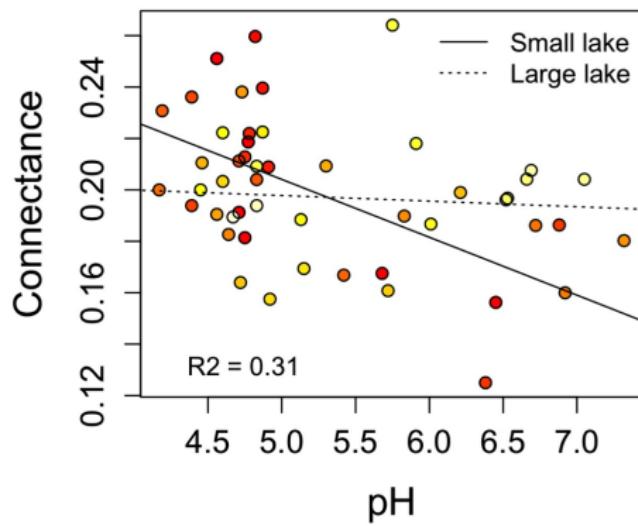
But...



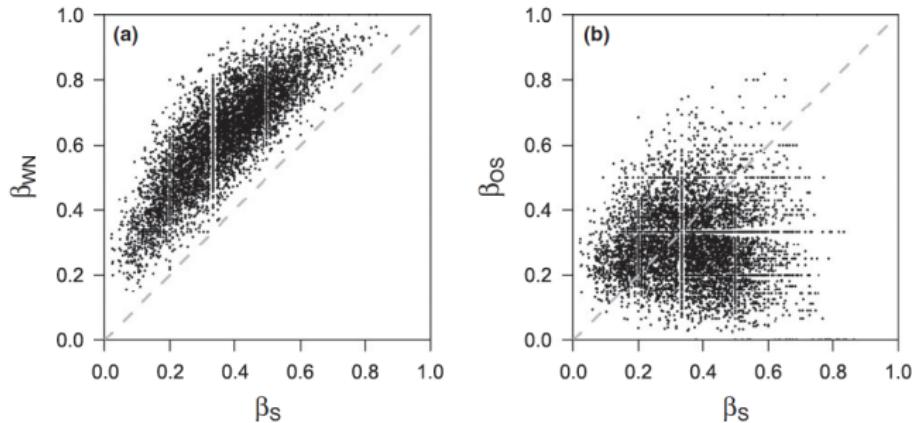
Gravel et al. (2011). *Ecol. Lett.*

What are the processes explaining how we move from a regional network (a metaweb) to a local community?

Biogeography of ecological interactions



Spatial variation of interaction networks



Poisot et al. (2012). *Ecol. Lett.*

Networks do vary in space because of:

- Species turnover;
- Link turnover;

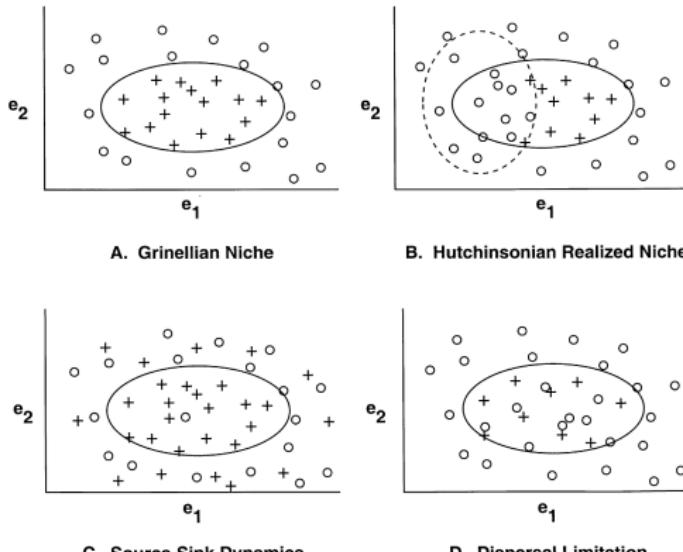
The variation of network structure is independent of the variation in species composition

- Random networks
- Cascade & Generalized cascade
- Niche model, Minimal potential niche model & Probabilistic niche model
- Nested hierarchy
- *Neutral networks*

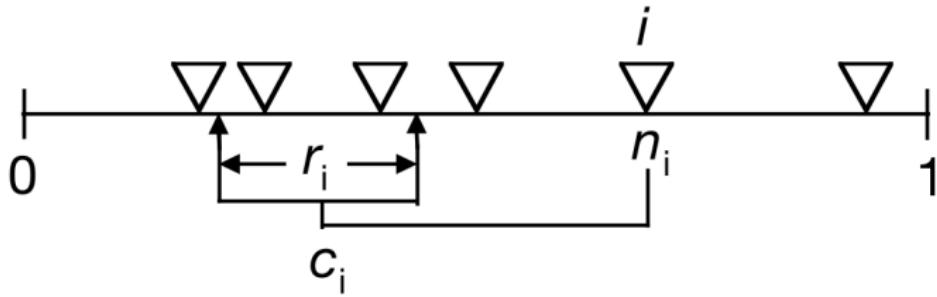
[The niche is] the joint description of the environmental conditions that allow a species to satisfy its minimum requirements so that the birth rate of a local population is equal or greater than its death rate along with the set of per capita effects of that species on these environmental conditions.

(Chase and Leibold, 2003)

The Grinnellian niche

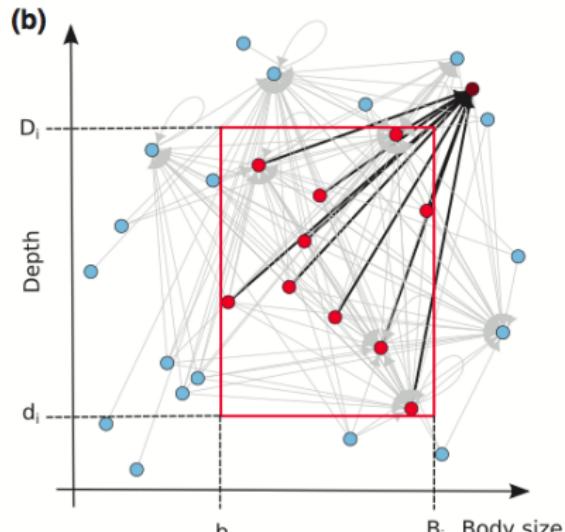


Pulliam (2000)



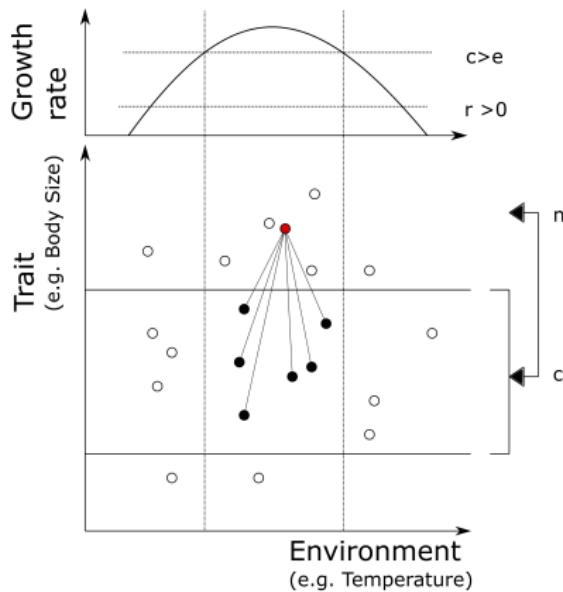
Williams & Martinez (2000)

Mapping ecological networks



Eklof et al (2013)

Mapping an integrated niche



Formulating network sampling as a stochastic process

Define the stochastic variable X_{ix} representing the occurrence of species i in a local community x.

And the variable L_{ijx} representing the occurrence of an interaction between species i and j at location x.

We are looking for the probability that an interaction occurs:

$$P(L_{ijx} = 1, X_{ix} = 1, X_{jx} = 1)$$

Obtained from the product rule we get:

$$P(L_{ijx}, X_{ix}, X_{jx}) = P(L_{ijx} | X_{ix}, X_{jx})P(X_{ix}, X_{jx})$$

Where:

$P(L_{ijx} | X_{ix}, X_{jx})$ is the metaweb (the Eltonian niche)

$P(X_{ix}, X_{jx})$ is the co-occurrence matrix (the Grinnellian niche)

We could refine the model with the addition of a condition on the environment, such as:

$$\begin{aligned} P(L_{ijx} | X_{ix}, X_{jx}, E_x) \\ \text{and} \\ P(X_{ix}, X_{jx} | E_x) \end{aligned}$$

We could also specify neutral dynamics underlying community assembly:

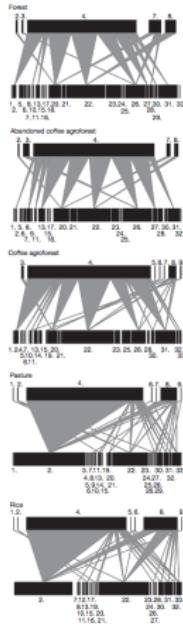
$$P(X_{ix}, X_{jx} | E_x) = P(X_{ix} | E_x)P(X_{jx} | E_x)$$

Example

Tropical cavity-nesting bees and wasps and their parasitoids

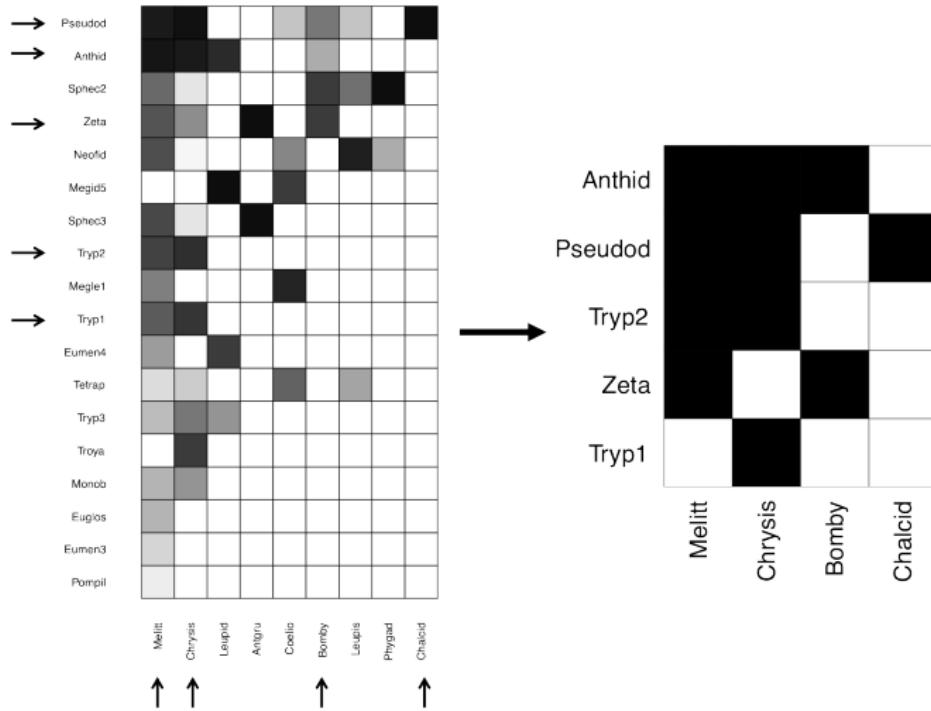
From Tylianakis et al. 2007. *Habitat modification alters the structure of tropical host-parasitoid food webs*. *Nature*, 44:202-205

- 48 networks, 4 090 recorded interactions
- 5 habitats: forest, abandoned coffee agroforest, coffee agroforest, pasture, rice
- 33 species of bees and wasps (Hymenoptera: Apidae, Megachilidae, Mutilidae, Pompilidae, Sphecidae, Vespidae)
- 9 parasitoid and kleptoparasite (Hymenoptera: Eulophidae, Ichneumonidae, Leucospidae, Megachilidae and Chrysidae; Diptera: Bombyliidae)



Example

Network #34, Pasture



Example

Observed vs Predicted

Interpretation	Model	$L(H D)$
<i>Basic model</i>		
No effect of E	$P(L_{ijx} X_{ix}, X_{jx})P(X_{ix}, X_{jx})$	-19.36
<i>Metaweb definitions</i>		
Conditional on E	$P(L_{ijx} X_{ix}, X_{jx} E_x)P(X_{ix}, X_{jx})$	-10.37
Aggregated metaweb	$P(L_{ijx}^* X_{ix}, X_{jx})P(X_{ix}, X_{jx})$	-27.96
<i>Co-occurrence</i>		
Neutral assembly	$P(L_{ijx} X_{ix}, X_{jx})P(X_{ix})P(X_{jx})$	-20.15
Conditional on E	$P(L_{ijx} X_{ix}, X_{jx})P(X_{ix} E_x)P(X_{jx} E_x)$	-13.30

Summary of the key ideas

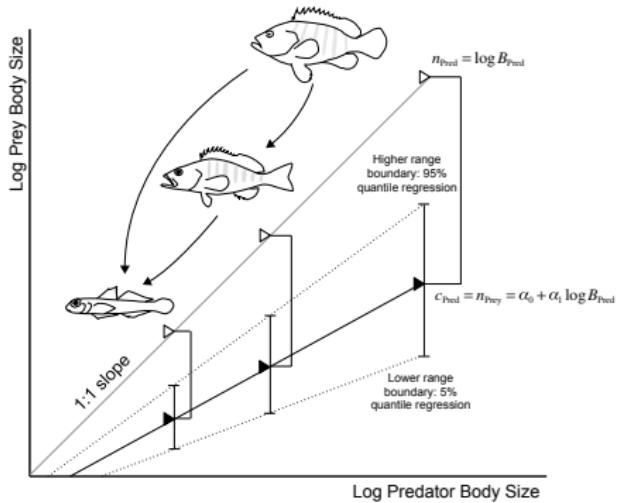
- Networks are not deterministic
- Links do change with the environment
- The environment also affect species turnover and thus network turnover
- Very weak effect of non-random species associations on the variation of network structure

Inferring the metaweb from traits

The problem: inferring interactions from species that never co-occurred and will be found in novel assemblages

Application

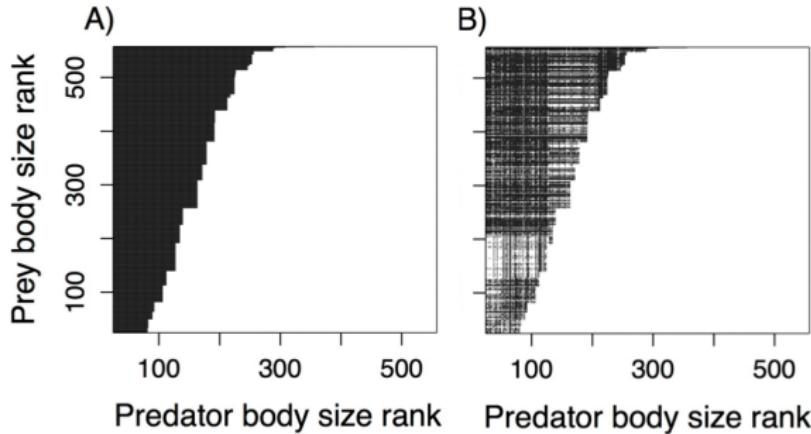
Inferring the metaweb from traits



Gravel et al. (2013). *Meth. Ecol. Evol.*

Without information on the probability of interactions, we considered a metweb *à la Havens* and neutral co-occurrence:

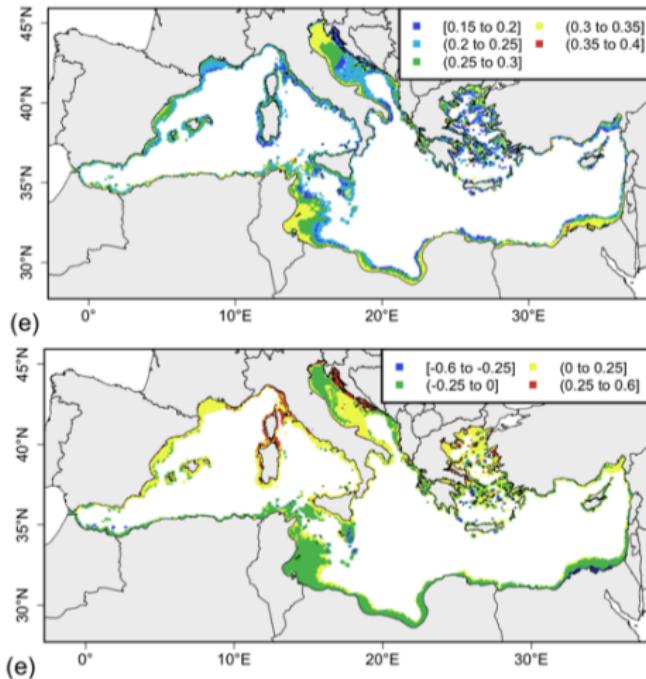
$$P(L_{ijx}, X_{ix}, X_{jx} | E_x) = P(L_{ijx} | X_{ix}, X_{jx})P(X_{ix} | E_x)P(X_{jx} | E_x)$$



Gravel et al. (2013)

Application

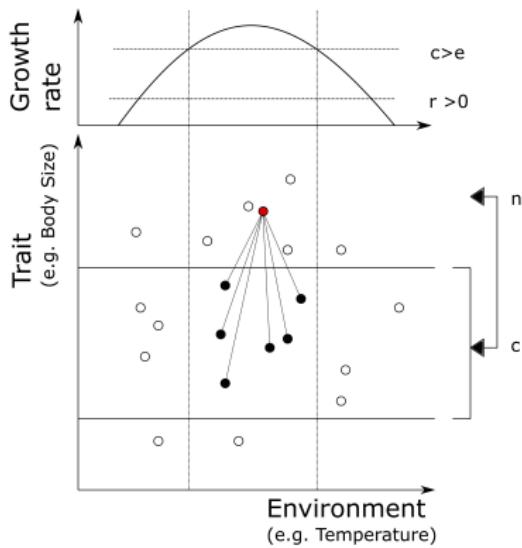
Pelagic food web of the Mediterranean sea



Albouy et al. (2014)

Conclusion

A new perspective to community structure



$$P(L_{ijx}, X_{ix}, X_{jx}) = P(L_{ijx} | X_{ix}, X_{jx})P(X_{ix}, X_{jx})$$

Given a theory for species co-occurrence, we could re-arrange information:

$$\begin{aligned} P(L_{ijx}, X_{ix}, X_{jx} | E_x) &= P(L_{ijx} | X_{ix}, X_{jx}, E_x)P(X_{ix}, X_{jx} | E_x) \\ &= P(L_{ijx} | X_{ix}, X_{jx}, E_x)P(X_{ix} | X_{jx}, E_x)P(X_{jx} | E_x) \end{aligned}$$

And invert it to derive a probability of species occurrence, accounting for the environment and interactions:

$$P(X_{jx} | E_x) = \frac{P(L_{ijx} | X_{ix}, X_{jx}, E_x)P(X_{ix} | X_{jx}, E_x)}{P(L_{ijx}, X_{ix}, X_{jx} | E_x)}$$

Acknowledgements

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