

# Introduction to the Study of Ecological Networks

Alyssa Cirtwill

Workshop NZES

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# Community 1 – a food web

Sanak Island is one of the Aleutian Islands, off the coast of Alaska. Until 1828 the island was inhabited by the Aleut people but is currently uninhabited.

In the intertidal zone, crabs and small fishes forage on barnacles, mussels, and limpets. The barnacles are filter-feeders while the limpets graze on algae and detritus. Littorinid snails compete with the limpets for algae but are able to escape predation by emerging onto shore. Large crabs occasionally prey upon smaller ones, while the small fish are eaten by large fish. Take 10 minutes to draw a food web representing this system.

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# Community 1 – Questions

1. How many species (nodes) are in your food web? How many interactions (links)?
2. What proportion of the possible links in the food web actually occur? What does this tell you about the community?
3. What's missing from this food web?



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## Community 2 – a plant-pollinator network

Now consider a meadow that is heavily used by commercial honeybees as a foraging site. In this meadow are flowers including borage, clover, and viper's bugloss that are favorite forage species for the bees. Surrounding the meadow are several patches of manuka, which the bees also visit frequently. The bees also visit forget-me-nots, gorse, and thistles, but less frequently. A few native alpine flowers also grow in the meadow (including alpine daisies, eyebrights, and orchids), but these are usually pollinated by native flies rather than the introduced honeybees. Take 10 minutes to draw a network describing the role of honeybees in this meadow.

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# Community 2 – Questions

1. This is an egocentric (or ‘sink’) web focused on the honeybees. What happens when you add non-honeybee pollinators?
2. How does this network differ from the Sanak Island food web?



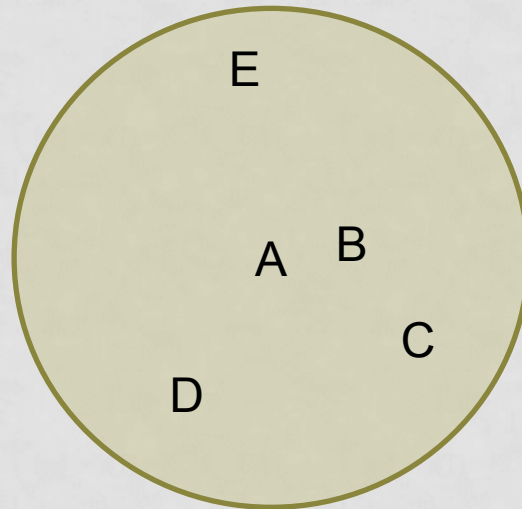


# Nestedness

Camille Coux & Marilia P. Gaiarsa

Introduction to the study of ecological networks  
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From island biogeography to explain spp occurrences in metacommunities.

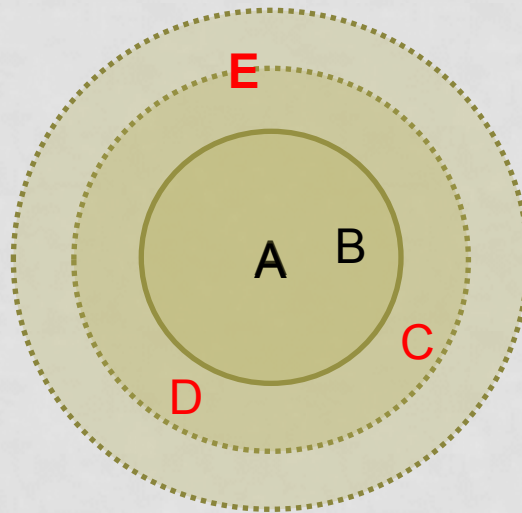


# From island biogeography to explain spp occurrences in metacommunities.

Large scale disturbance,

e.g. climate

→ Habitat reduction and fragmentation



- Archipelago of “islands”
- “Sp comprising a depauperate fauna should constitute a proper subset of those in richer fauna” (Darlington 1957)



- Archipelago of “islands”
- “Sp comprising a depauperate fauna should constitute a proper subset of those in richer fauna” (Darlington 1957)

Range	Montane mammal species*	Richness
1	ABCDEFGHIJ KLMNOPQRST UVWXYZ	26
4	ABCDEFGHIJ KLMNOPQRST UVWX	24
3	ABCDEFGHIJ KLMNO+QRST UVWX	23
2	ABCDEFGHIJ KLMNOPQRST U+	21
5	ABCDEFGHIJ KLM+OPQRST +	19
8	ABCDE+GHIJ K++NOP +	13
9	ABCDE+G+IJ K++NO+Q+ V	13
6	ABCDE++HIJ KL+NO+ +	12
24	ABCDEFGHI+KL++ P +	11
10	ABCDEFGHI+IJ K+++ + +	10
11	ABCDEFGHIJ +++ + +	10
14	ABCDEFGHI+++ +M+ R	9
23	ABCDEFGHI+H++ L++ P	9
7	ABC+E+ IJ + N	7
13	ABCDEF I +	7
15	ABCDEF +M	7
17	ABCDE+G +M	7
22	ABCDEF H +	7
25	ABCDEF H +	7
12	ABCDEF +	6
21	ABC D+F L	6
16	ABCDE +	5
20	ABC++F L	5
27	AB+DE +	4
26	AB+ F†	3
28	AB +	2
18	C	1
19	C	1

Patterson & Atmar 1986

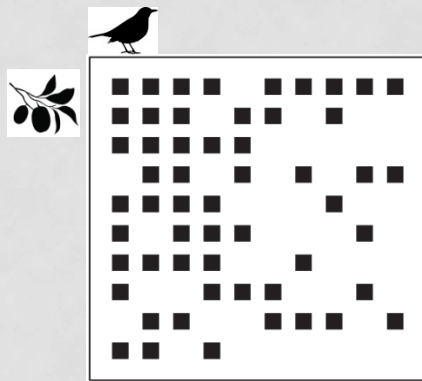
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5	ABCDEFGHIJ KLM+OPQRST +	19
8	ABCDE+GHIJ K++NOP	13
9	ABCDE+G+IJ K++NO Q+ V	13
6	ABCDE++HIJ KL+O+ +	12
24	ABCDEFGH++K++ P +	11
10	ABCDEFG+IJ K+++ + +	10
11	ABCDEFGH++ + + +	10
14	ABCDEFG++ +M+ R	9
23	ABCDEFG++ L++ P	9
7	ABC+E+ IJ + N	7
13	ABCDE+ I +	7
15	ABCDE+ +M	7
17	ABCD+G +M	7
22	ABCDEF H +	7
25	ABCDEF H +	7
12	ABCDEF +	6
21	ABCDEF L	6
16	ABCDE+ +	5
20	AB++F L	5
27	ABDE+ +	4
26	AB+ F†	3
28	AB+ +	2
18	C	1
19	C	1

Patterson & Atmar 1986

## Nestedness of interactions:

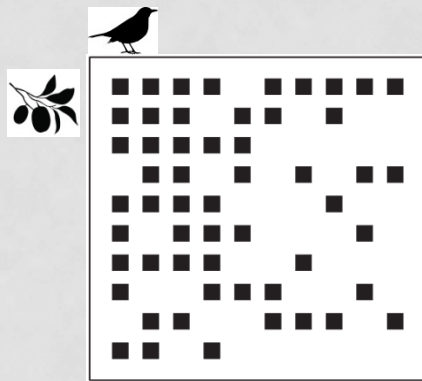
Each sp interacts with subsets of sp interacting with more generalist species.



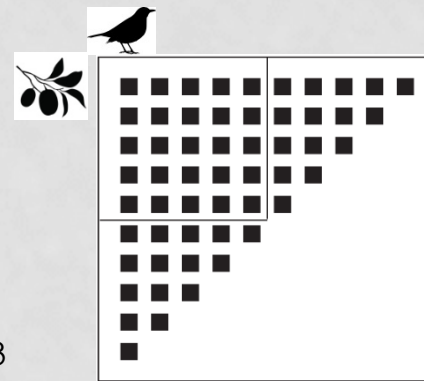
Bascompte et al., 2003

## Nestedness of interactions:

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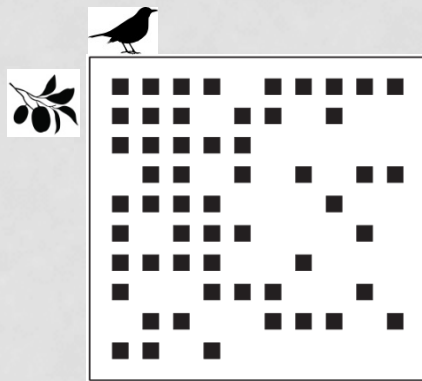
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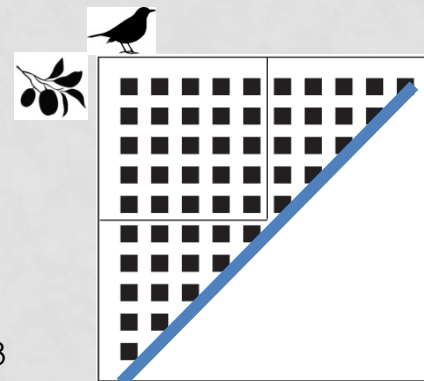
Nestedness = nonrandom pattern beyond connectedness

## Nestedness of interactions:

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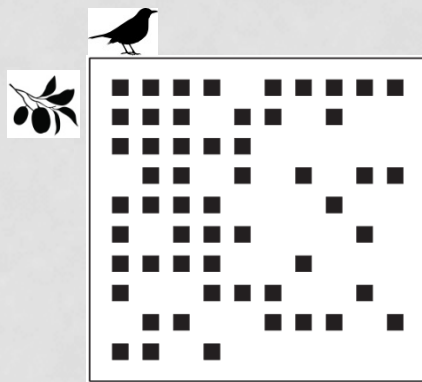
Bascompte et al., 2003



Nestedness = nonrandom pattern beyond connectedness

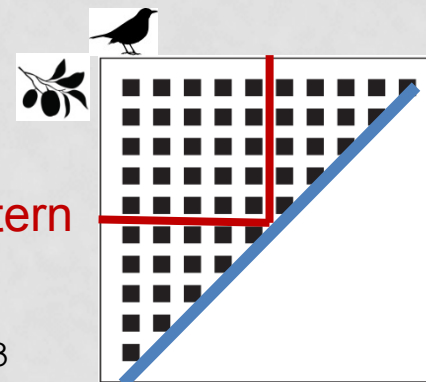
## Nestedness of interactions:

Each sp interacts with subsets of sp interacting with more generalist species.



Cohesive pattern

Bascompte et al., 2003

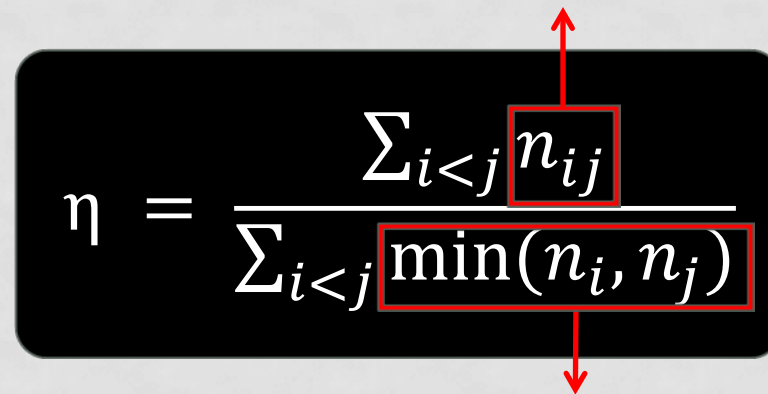


Nestedness = nonrandom pattern beyond connectedness

## How is nestedness calculated?

Unweighted nestedness from Bastollos et al., 2009:

*# of interactions common to  
both plants  $i$  and  $j$*


$$\eta = \frac{\sum_{i < j} n_{ij}}{\sum_{i < j} \min(n_i, n_j)}$$

The diagram shows the formula for unweighted nestedness. The numerator is the sum of interactions  $n_{ij}$  for all pairs  $i < j$ . The denominator is the sum of the minimum of the number of interactions for each plant,  $\min(n_i, n_j)$ . Red boxes highlight  $n_{ij}$  in the numerator and  $\min(n_i, n_j)$  in the denominator. A red arrow points from the box around  $n_{ij}$  to the text above, and another red arrow points from the box around  $\min(n_i, n_j)$  to the text below.

*minimum # of interactions between  $n_i$  and  $n_j$*

0 = random; 1=perfect nestedness

# Nestedness

# of interactions common to both plants  $i$  and  $j$

$$\eta = \frac{\sum_{i < j} n_{ij}}{\sum_{i < j} \min(n_i, n_j)}$$

minimum # of interactions between  $n_i$  and  $n_j$

↓	↓		
1	1	1	1
1	1	1	0
1	1	0	0
1	0	0	0

$N =$  \_\_\_\_\_

# Nestedness

# of interactions common to both plants  $i$  and  $j$

$$\eta = \frac{\sum_{i < j} n_{ij}}{\sum_{i < j} \min(n_i, n_j)}$$

minimum # of interactions between  $n_i$  and  $n_j$

↓	↓		
1	1	1	1
1	1	1	0
1	1	0	0
1	0	0	0

$$N = \frac{3}{\quad}$$

# Nestedness

# of interactions common to both plants  $i$  and  $j$

$$\eta = \frac{\sum_{i < j} n_{ij}}{\sum_{i < j} \min(n_i, n_j)}$$

minimum # of interactions between  $n_i$  and  $n_j$

↓	↓		
1	1	1	1
1	1	1	0
1	1	0	0
1	0	0	0

4 < 3

$$N = \frac{3}{3}$$

# Nestedness

# of interactions common to both plants  $i$  and  $j$

$$\eta = \frac{\sum_{i < j} n_{ij}}{\sum_{i < j} \min(n_i, n_j)}$$

minimum # of interactions between  $n_i$  and  $n_j$

↓	↓		
1	1	1	1
1	1	1	0
1	1	0	0
1	0	0	0

4 < 3

$$N = \frac{3 + 2 + 1 + 2 + 1 + 1}{3 + 2 + 1 + 2 + 1 + 1} = 1$$

# Nestedness

# of interactions common to both plants  $i$  and  $j$

$$\eta = \frac{\sum_{i < j} n_{ij}}{\sum_{i < j} \min(n_i, n_j)}$$

minimum # of interactions between  $n_i$  and  $n_j$

↓	↓		
1	1	0	1
1	0	1	0
1	1	0	0
0	0	1	0

$$\frac{2 + 1 + 1 + 0 + 1 + 0}{2 + 2 + 1 + 2 + 1 + 1} = 0.55$$

# Nestedness

# of interactions common to both plants  $i$  and  $j$

$$\eta = \frac{\sum_{i < j} n_{ij}}{\sum_{i < j} \min(n_i, n_j)}$$

minimum # of interactions between  $n_i$  and  $n_j$

↓	↓		
1	1	0	1
1	0	1	0
1	1	0	0
0	0	1	0
<hr/>			
3	<	2	

$$\frac{2 + 1 + 1 + 0 + 1 + 0}{2 + 2 + 1 + 2 + 1 + 1} = 0.55$$

# Nestedness

# of interactions common to both plants  $i$  and  $j$

$$\eta = \frac{\sum_{i < j} n_{ij}}{\sum_{i < j} \min(n_i, n_j)}$$

minimum # of interactions between  $n_i$  and  $n_j$

↓		↓	
1	1	0	1
<b>1</b>	0	<b>1</b>	0
1	1	0	0
0	0	1	0
<hr/>			
3	<	2	

$$\frac{2 + \textcolor{red}{1} + 1 + 0 + 1 + 0}{2 + \textcolor{brown}{2} + 1 + 2 + 1 + 1} = 0.55$$

## Many other calculations:

- Nestedness temperature : deviation from isocline (Atmar & Patterson 1993)

0=cold, i.e. perfect N, 100=hot, i.e.chaos

- NODF: Node Overlap and Decreasing Fill (Almeida-Neto et al. 2008)

high values = high nestedness

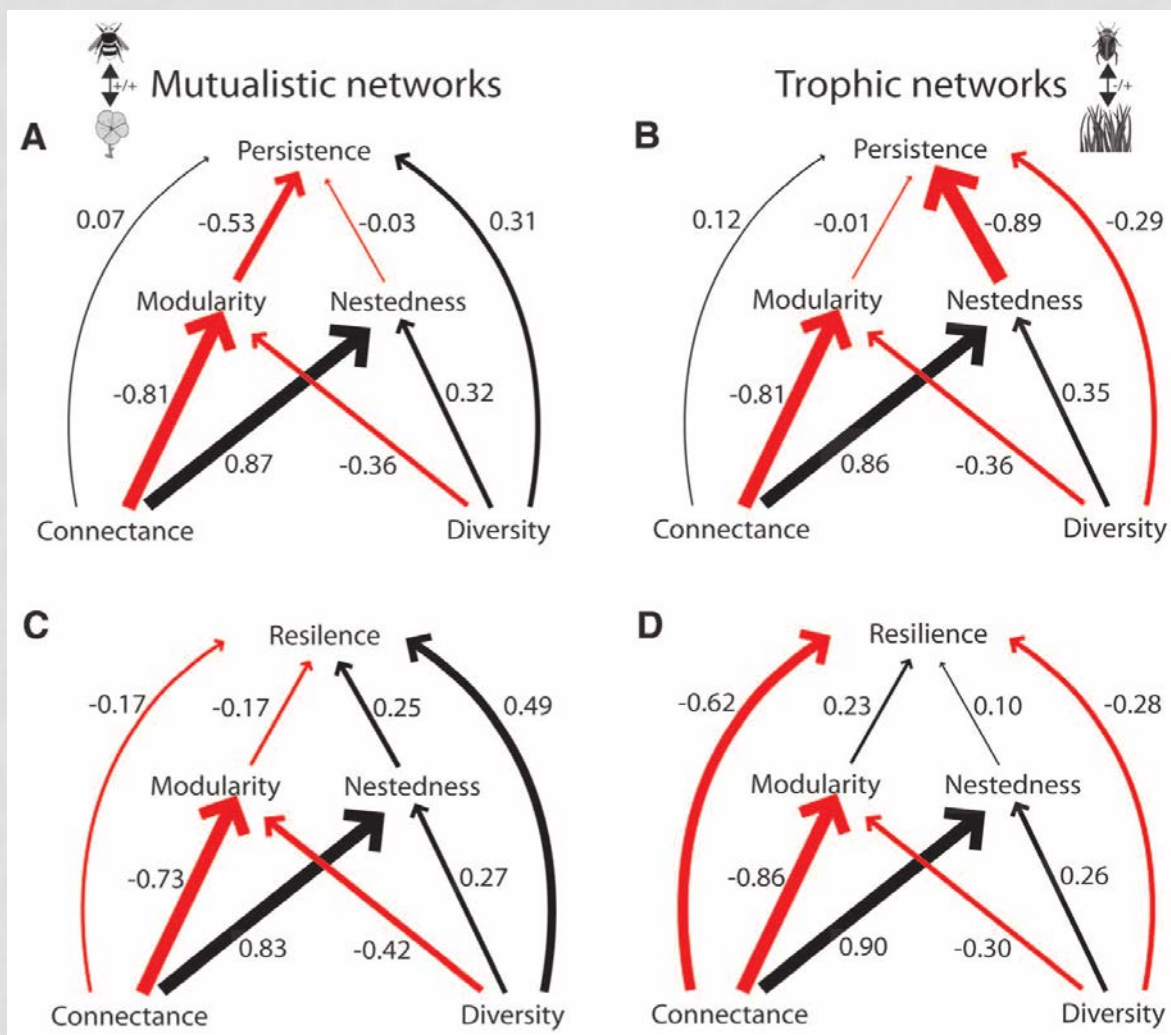
- Weighted version of NODF

(Almeida-Neto et al. 2010)

## Implications: consequences for stability

- Generalist core with rest of the community attached to it
- Asymmetric specialisation: specialists interact with generalists, not with other specialists

# Implications: consequences for stability

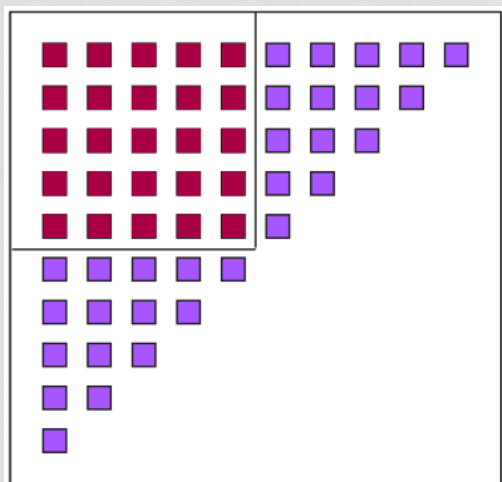


## Implications: consequences for (co)evolution

- In mutualistic networks, cohesive core of generalist species hypothesised to act as a “coevolutionary vortex” (Bascompte et al. 2003; Thompson 2005; Guimaraes et al. 2007)

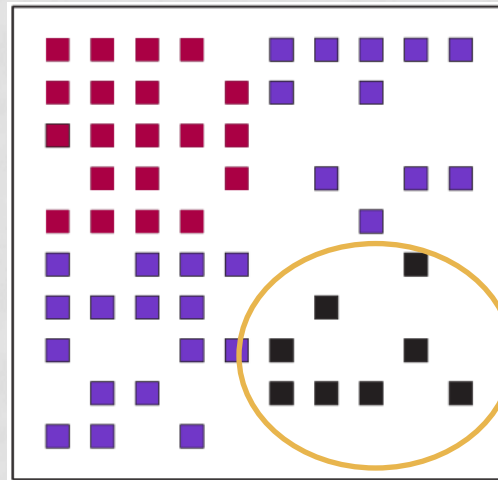
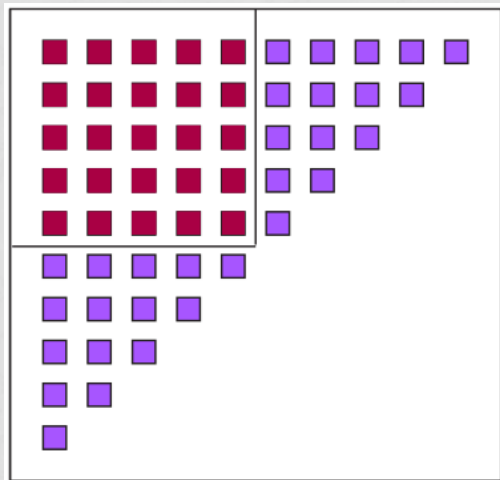
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## Implications: consequences for (co)evolution

- In mutualistic networks, cohesive core of generalist species hypothesised to act as a “coevolutionary vortex” (Bascompte et al. 2003; Thompson 2005; Guimaraes et al. 2007)



Not influenced by  
the core

# Modularity

Marilia P. Gaiarsa & Camille Coux

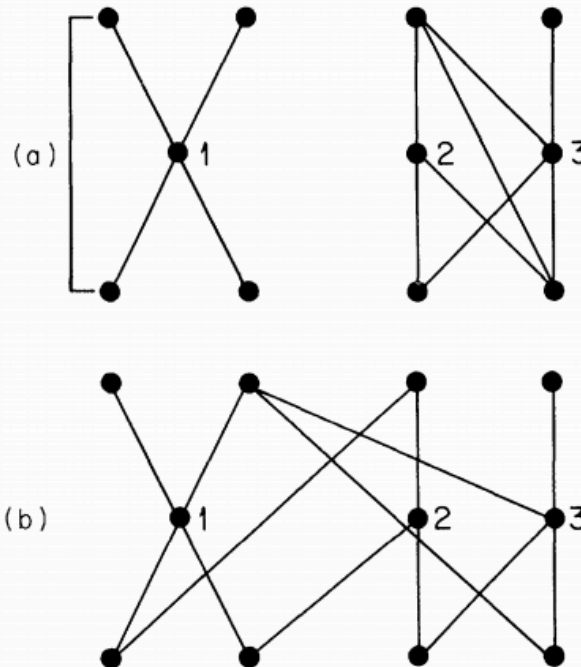
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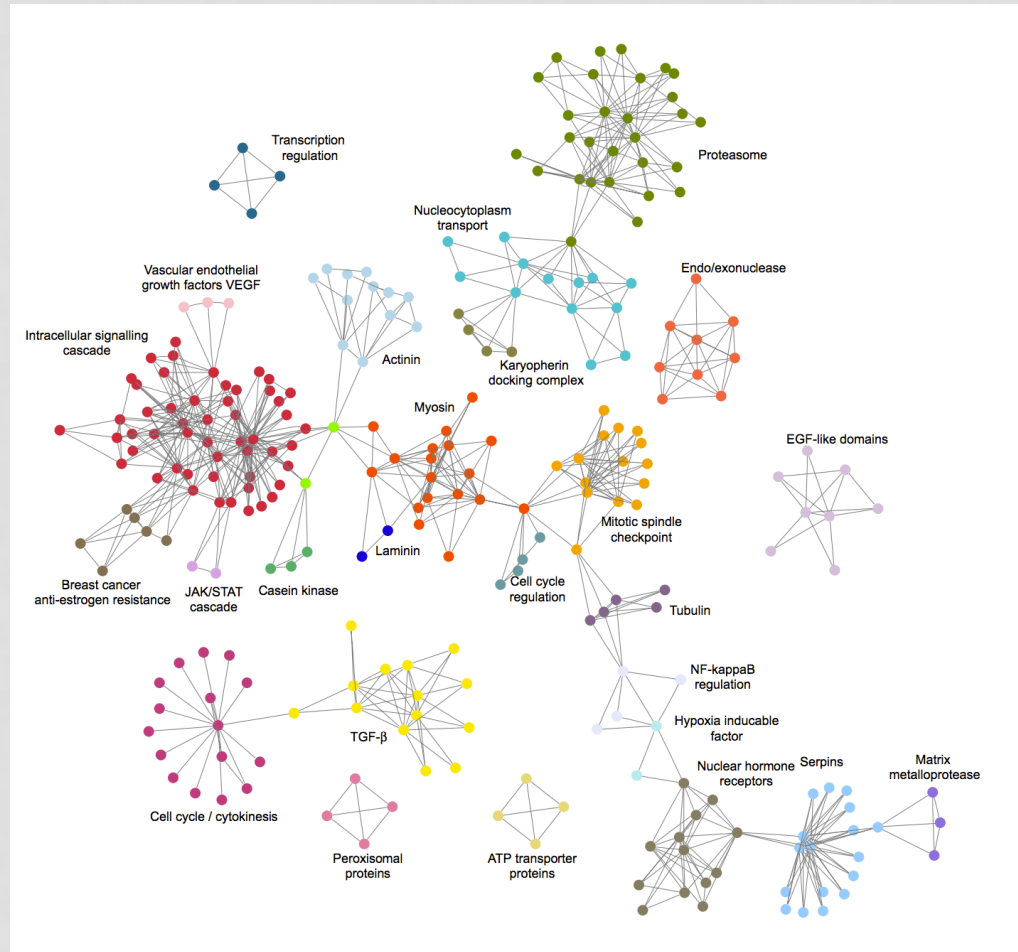
*Journal of Animal Ecology* (1980), **49**, 879–898

# ARE FOOD WEBS DIVIDED INTO COMPARTMENTS?\*

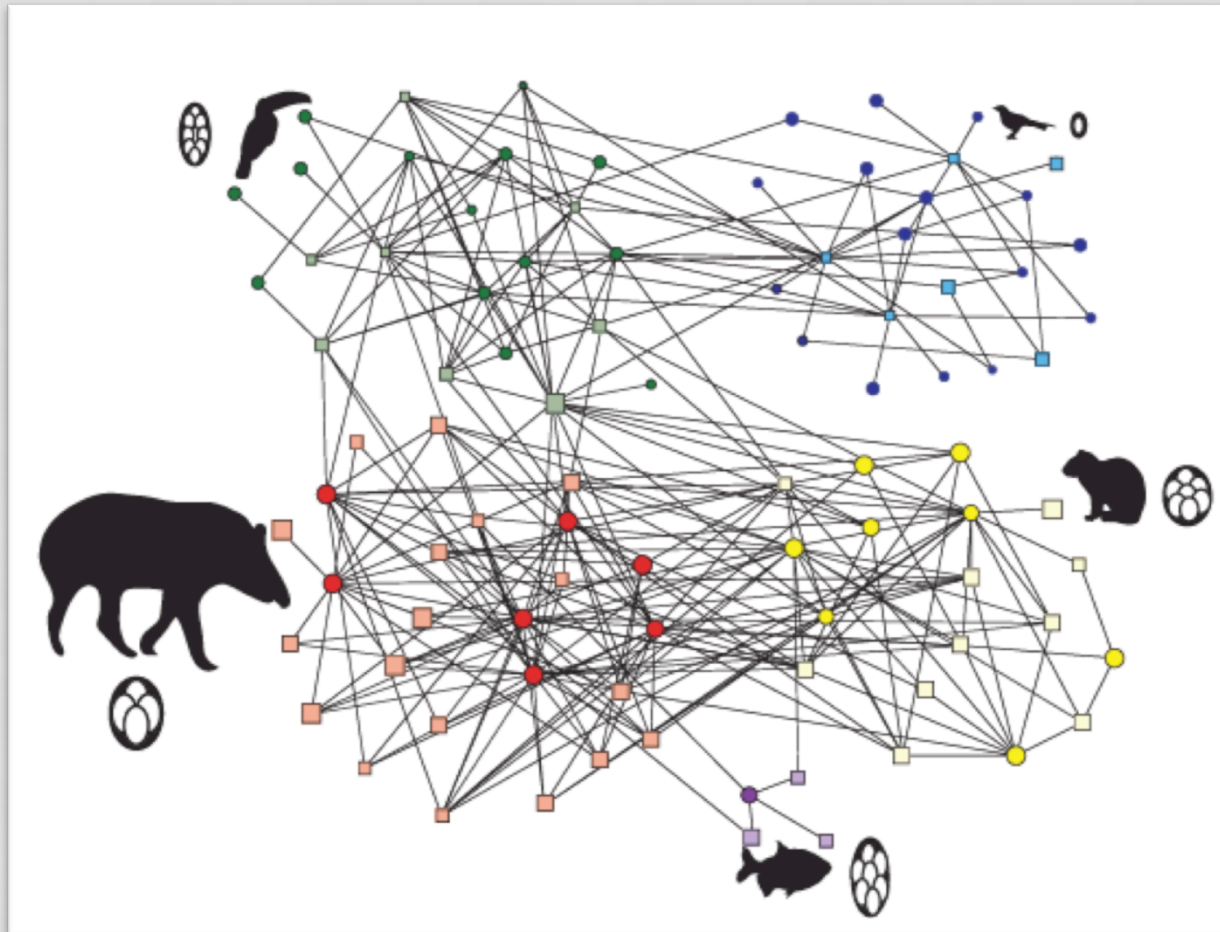
BY STUART L. PIMM† AND JOHN H. LAWTON

*Are food webs compartmented?*

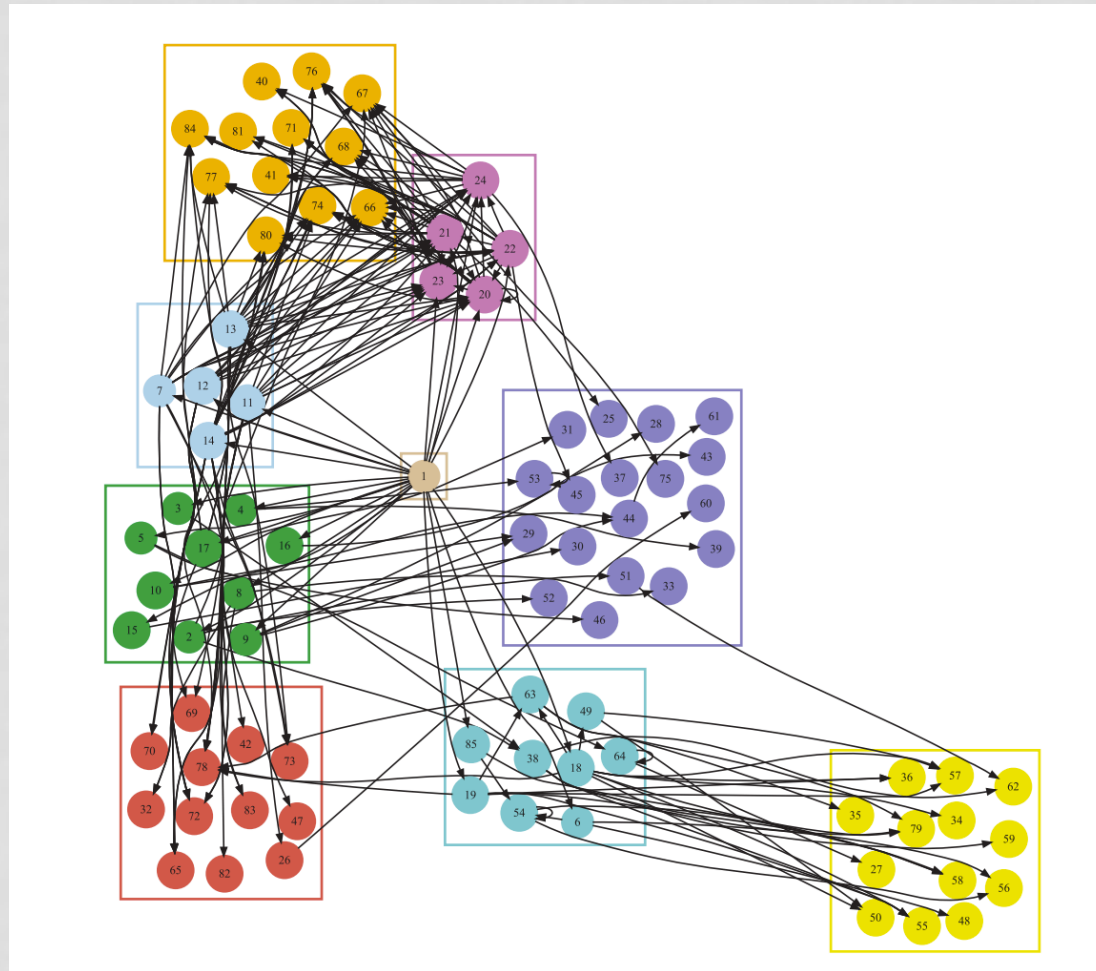




Jonsson et al. 2006, Fortunato 2010

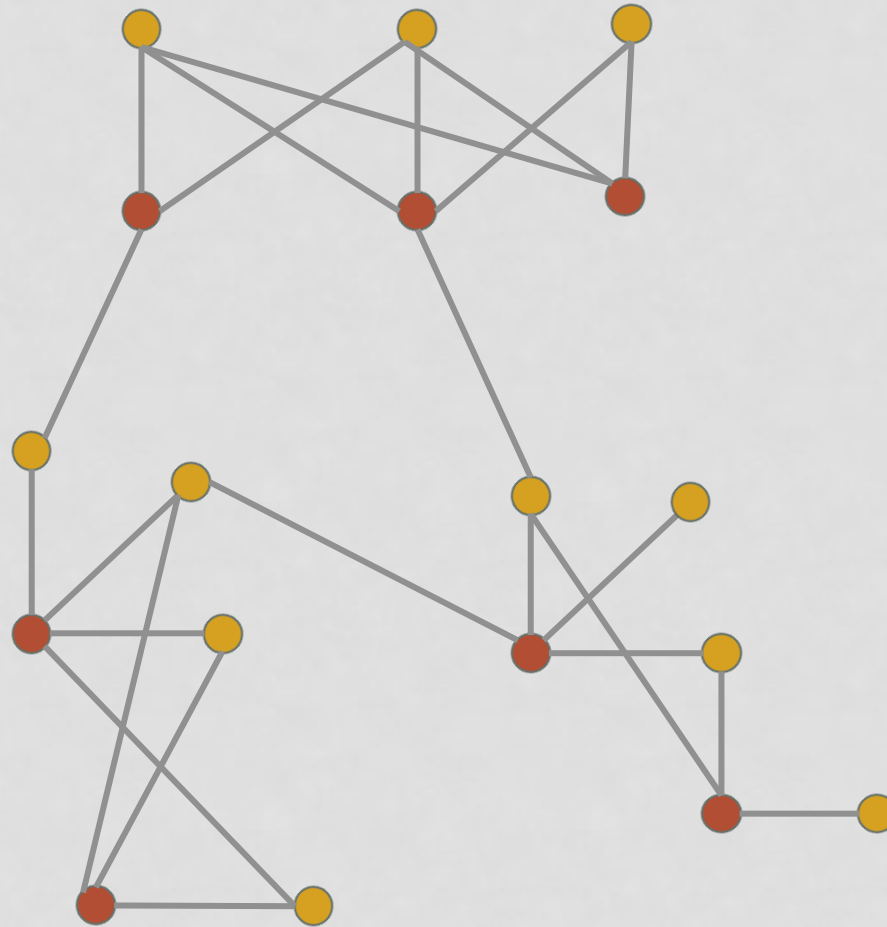


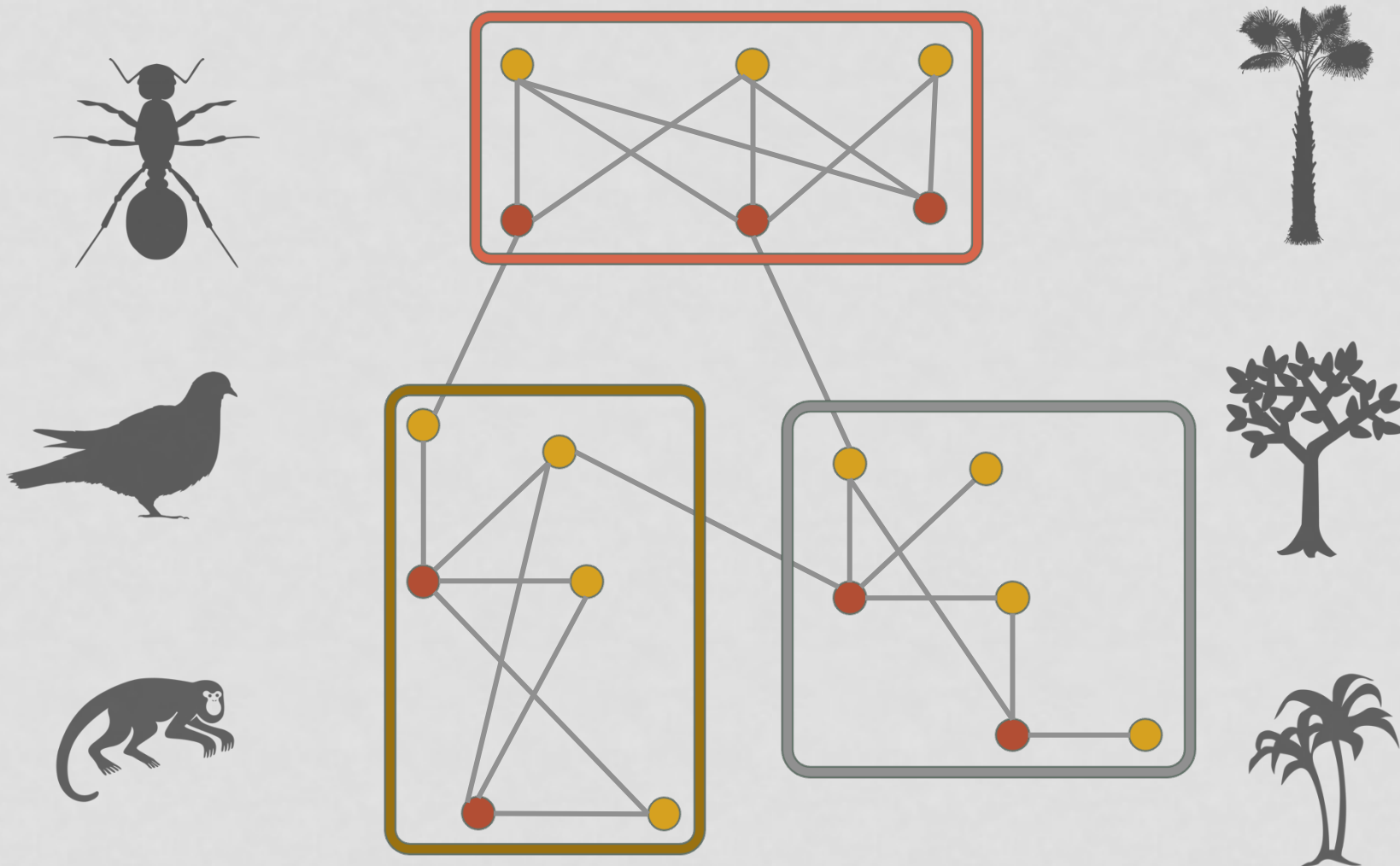
Donatti et al. 2011

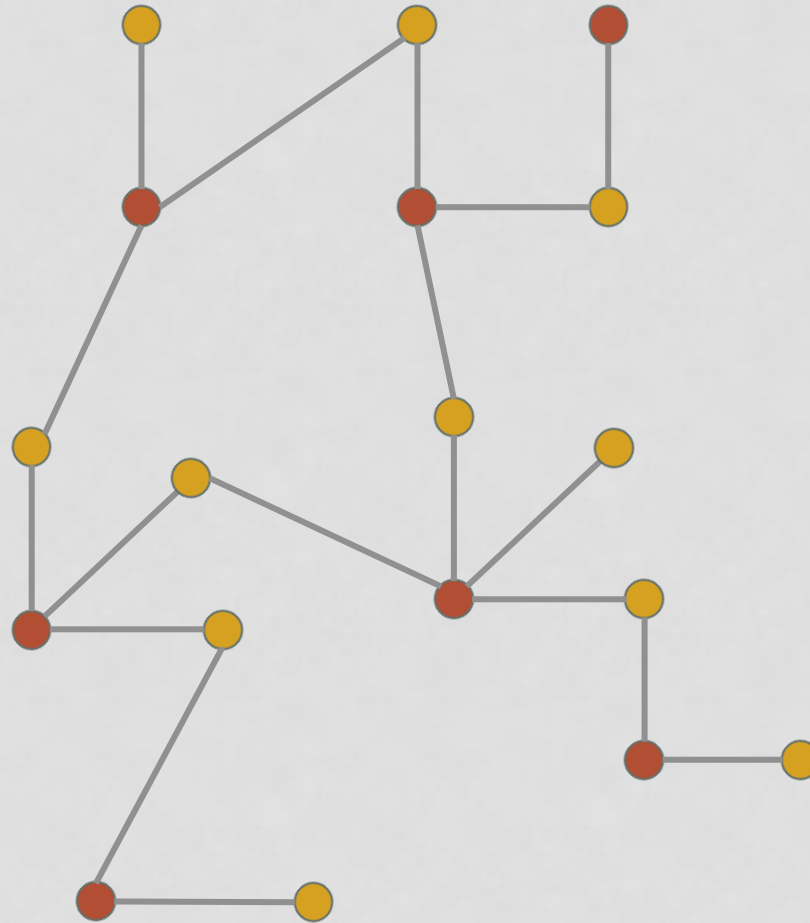


Allesina & Pascual 2009

“Modularity is the tendency where species within a module tend to interact with a much higher frequency among them than they do with species from other modules” (Bascompte & Jordano 2014)





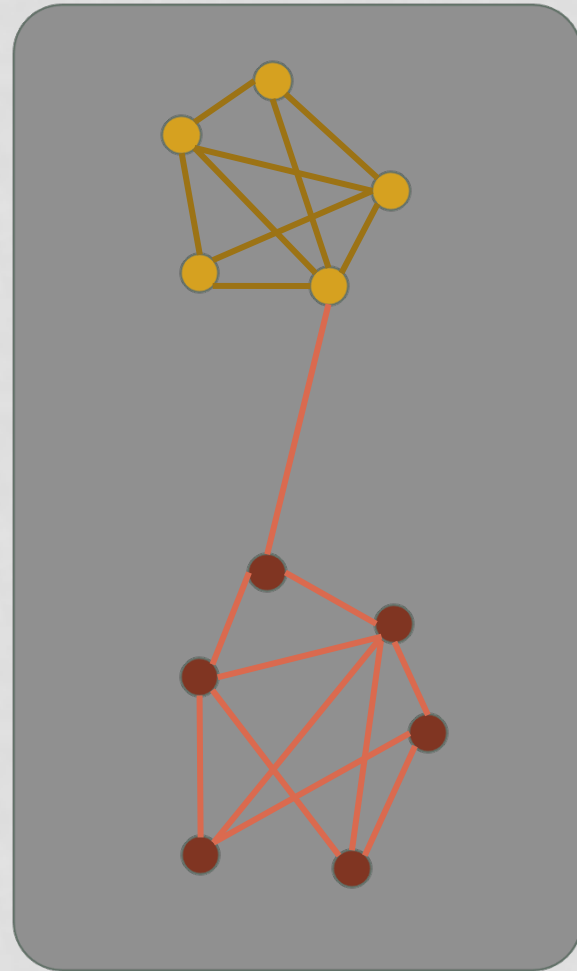


# How to characterize groups of interactions?

## The metric M

$$M = \sum_{\text{all modules } s} \left( \frac{l_s}{L} - \frac{d_s^P}{L} \frac{d_s^A}{L} \right)$$

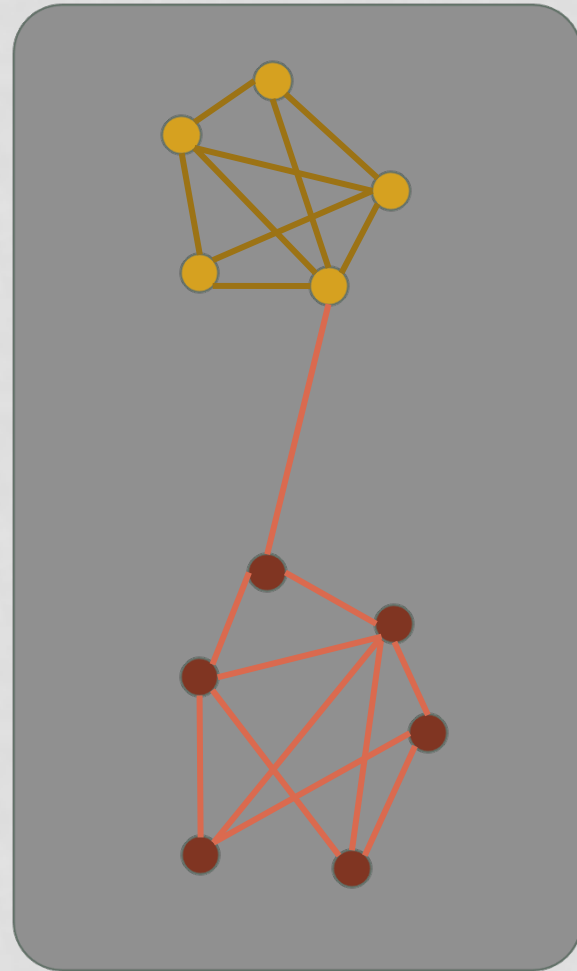
Barber 2007, Guimerà et al. 2007



*# of interactions inside module s*

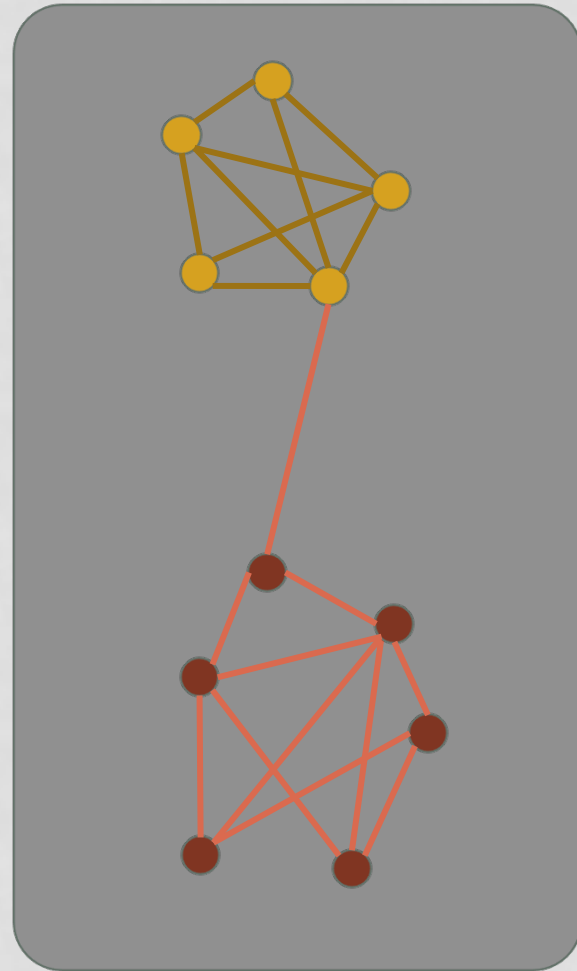
$$M = \sum_{\text{all modules } s} \left( \frac{l_s}{L} - \frac{d_s^P}{L} \frac{d_s^A}{L} \right)$$

*# of interactions in the whole network*



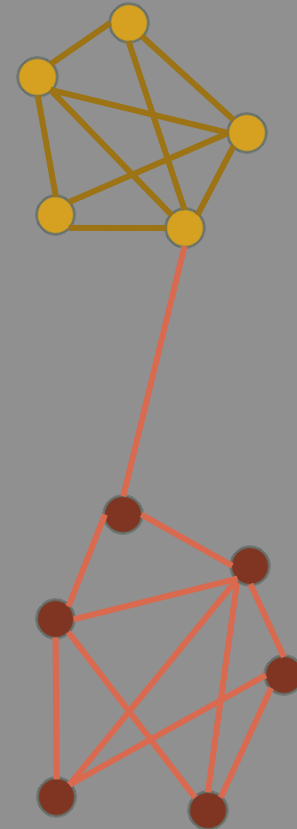
*Sum of the plants' degree inside module s*

$$M = \sum_{\text{all modules } s} \left( \frac{l_s}{L} - \frac{d_s^P}{L} \frac{d_s^A}{L} \right)$$



*Sum of the animals' degree inside module s*

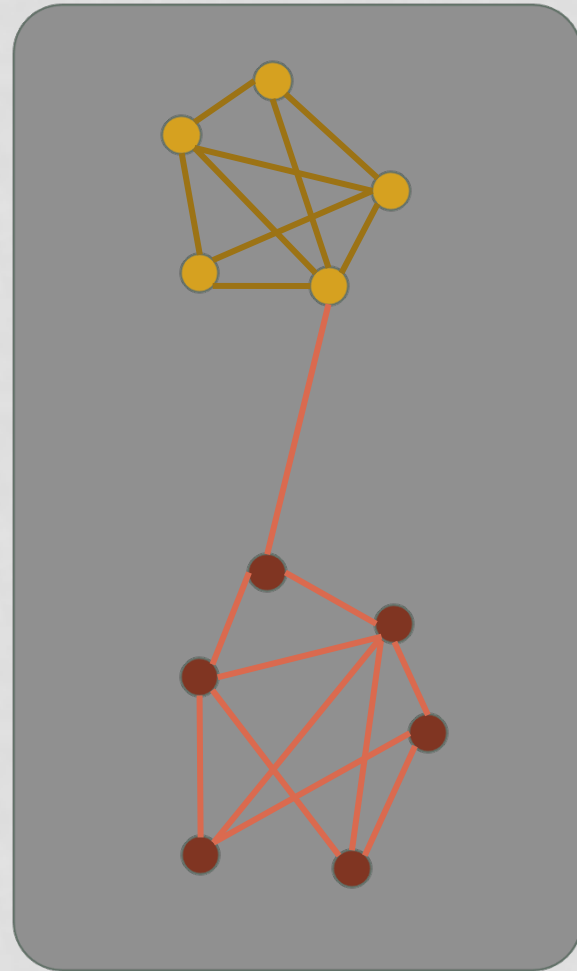
$$M = \sum_{\text{all modules } s} \left( \frac{l_s}{L} - \frac{d_s^P}{L} \frac{d_s^A}{L} \right)$$



The metric M

$$M = \sum_{\text{all modules } s} \left[ \frac{l_s}{L} - \left( \frac{d_s}{2L} \right)^2 \right]$$

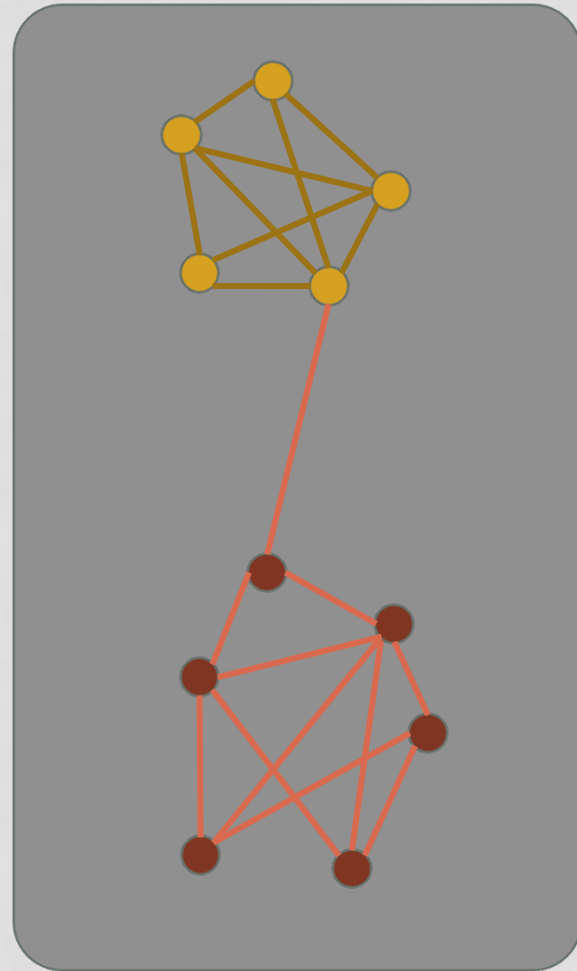
Newman & Girvan 2004



*# of interactions inside module s*

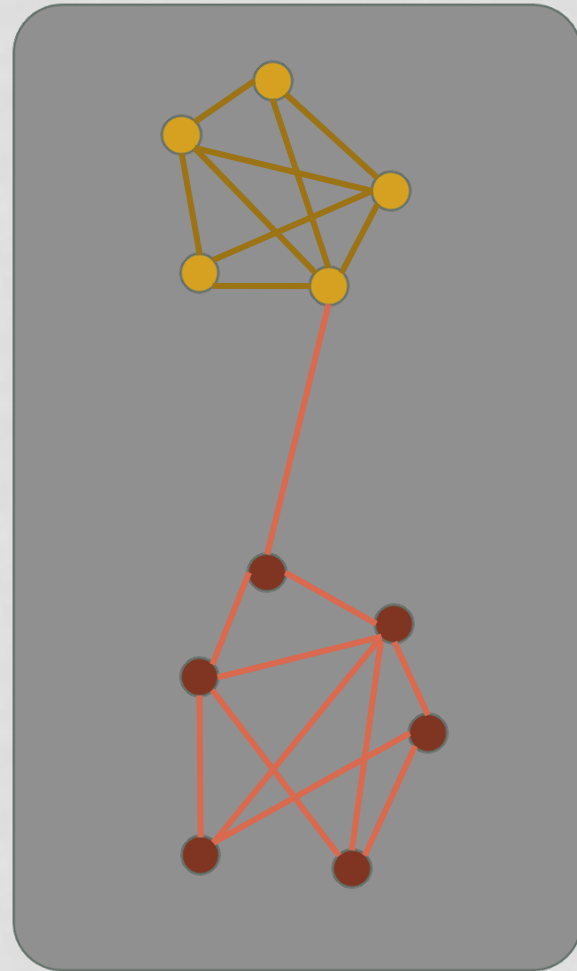
$$M = \sum_{\text{all modules } s} \left[ \frac{l_s}{L} - \left( \frac{d_s}{2L} \right)^2 \right]$$

*# of interactions in the whole network*



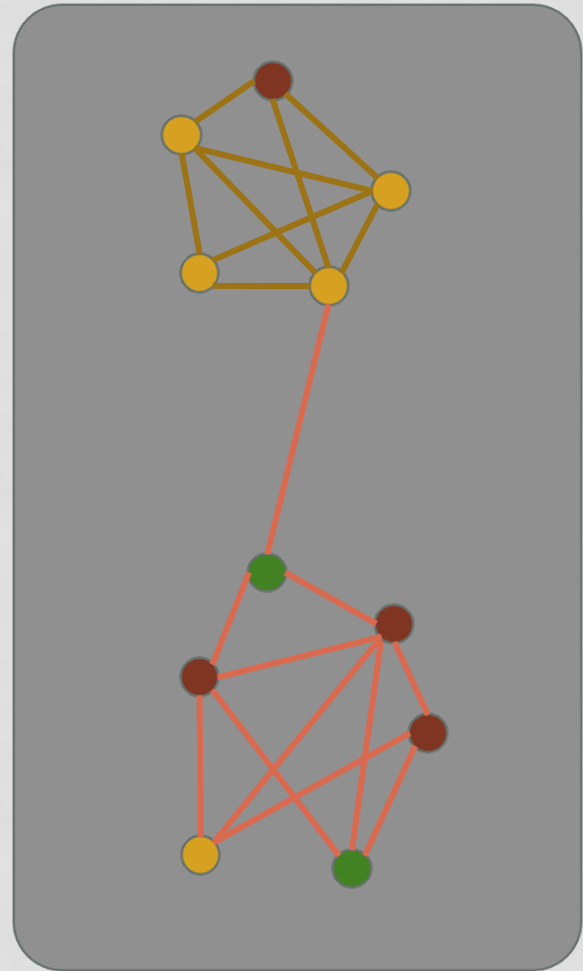
*Sum of the species'  
degree inside module s*

$$M = \sum_{\text{all modules } s} \left[ \frac{l_s}{L} - \left( \frac{d_s}{2L} \right)^2 \right]$$



How to find the modules?

$$M = \sum_{\text{all modules } s} \left[ \frac{l_s}{L} - \left( \frac{d_s}{2L} \right)^2 \right]$$



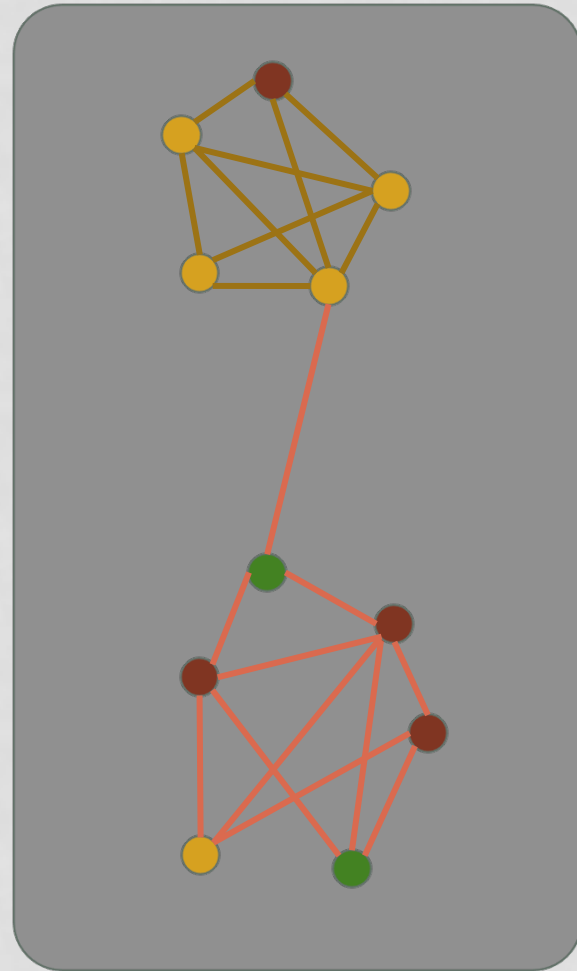
How to find the modules?

$$M = \sum_{\text{all modules } s} \left[ \frac{l_s}{L} - \left( \frac{d_s}{2L} \right)^2 \right]$$

●  $(2/20) - (15/40)^2$

●  $(6/20) - (18/40)^2$

●  $(0/20) - (6/40)^2$



How to find the modules?

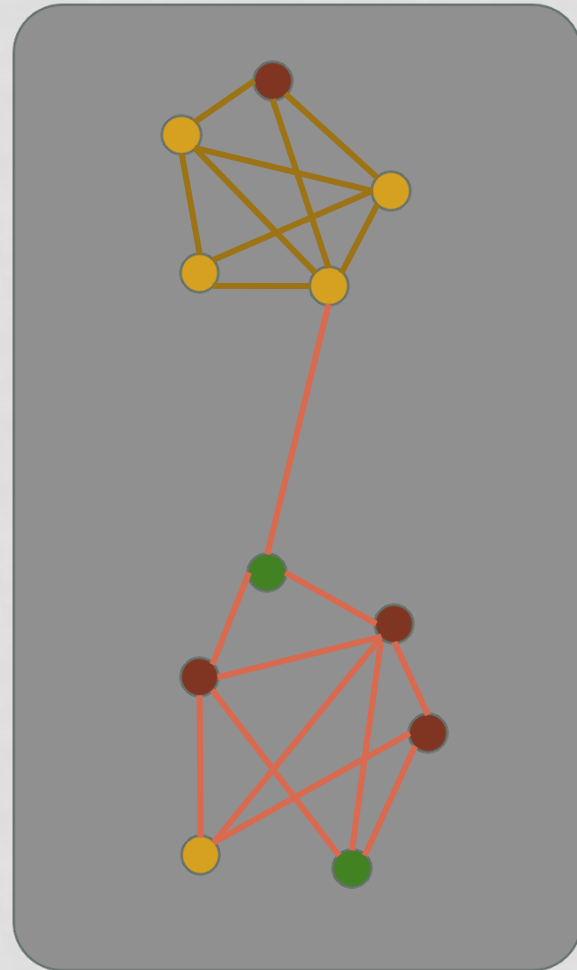
$$M = \sum_{\text{all modules } s} \left[ \frac{l_s}{L} - \left( \frac{d_s}{2L} \right)^2 \right]$$

● - 0.04

● 0.10

● - 0.02

M = 0.04



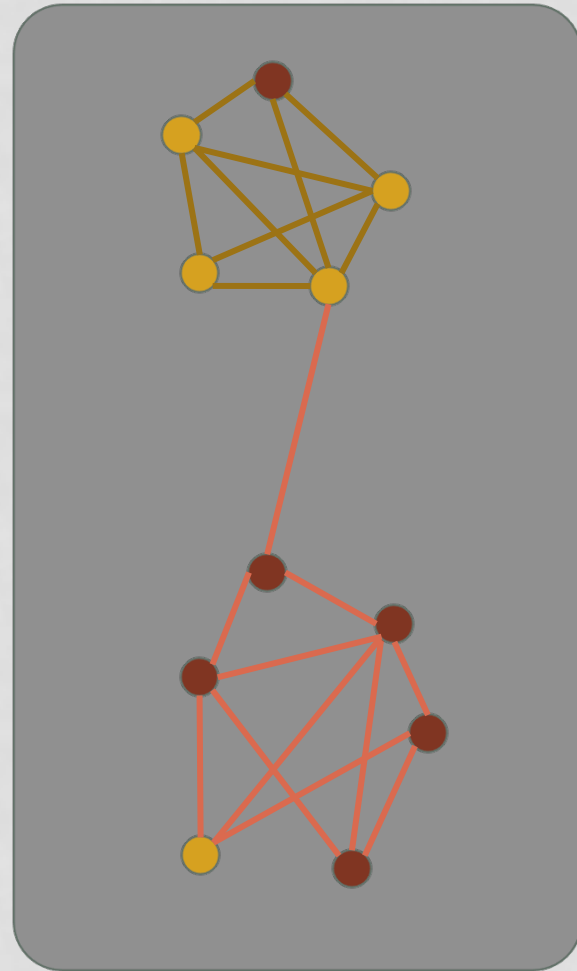
How to find the modules?

$$M = \sum_{\text{all modules } s} \left[ \frac{l_s}{L} - \left( \frac{d_s}{2L} \right)^2 \right]$$

● 0.01

● 0.15

$M = 0.16$



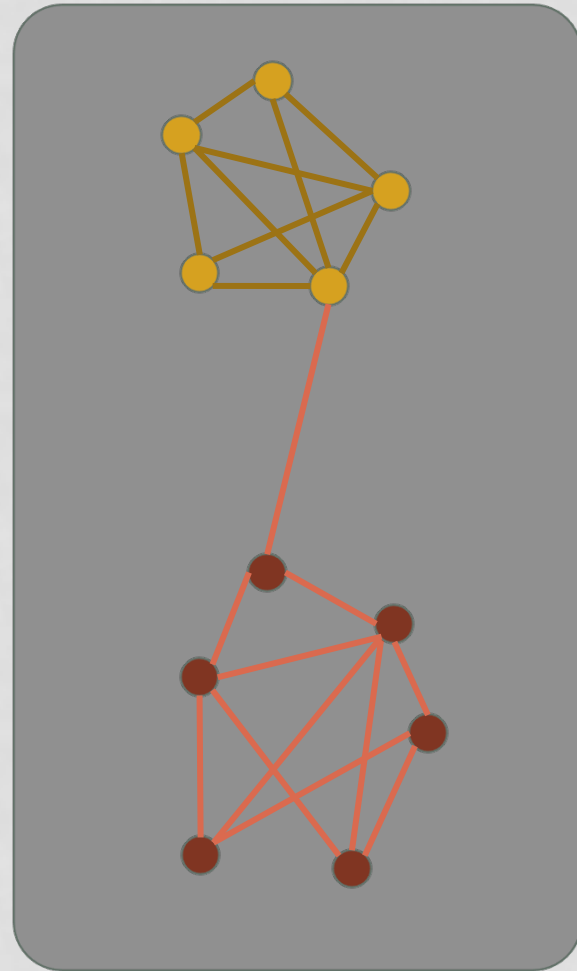
How to find the modules?

$$M = \sum_{\text{all modules } s} \left[ \frac{l_s}{L} - \left( \frac{d_s}{2L} \right)^2 \right]$$

● 0.22

● 0.25

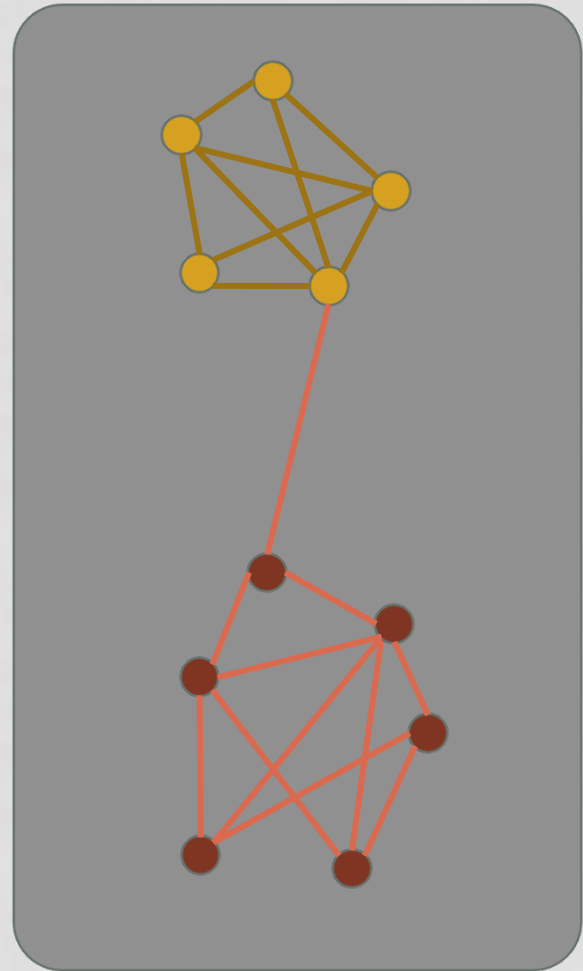
$M = 0.47$



11 species: 1 – 11 modules;

Different sizes;

How to optimize?

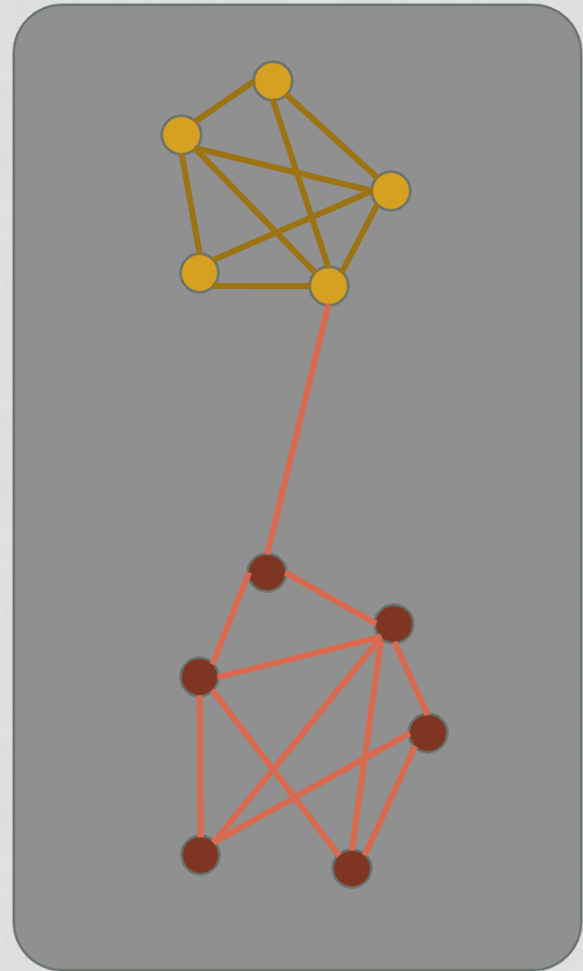


11 species: 1 – 11 modules;

Different sizes;

How to optimize?

**Simulated annealing**



11 species: 1 – 11 modules;

Different sizes;

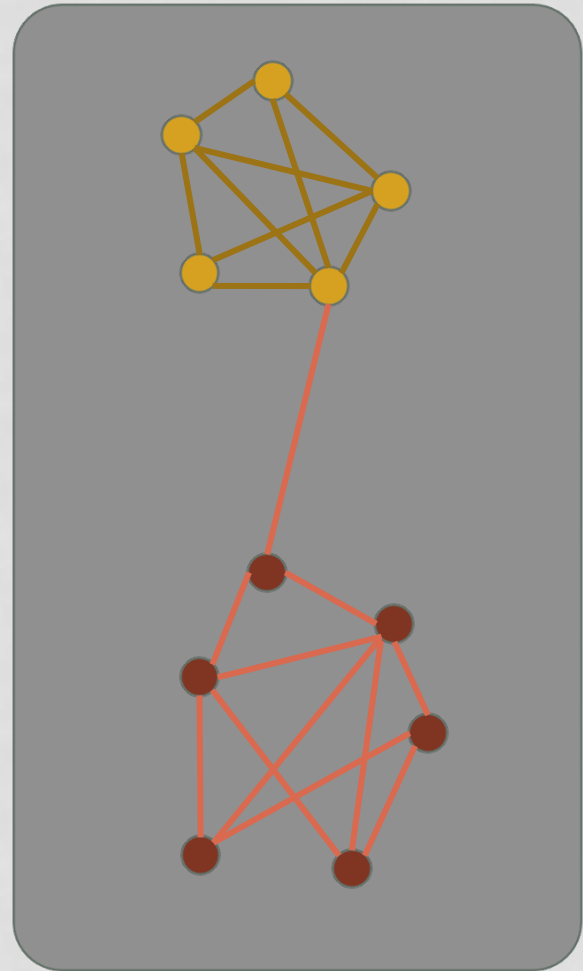
How to optimize?

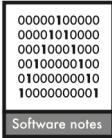
Simulated annealing

**Netcarto** (Guimera & Amaral)

**Rnetcarto** (Doulcier)

**Modular** (Marquitti et al.)





## MODULAR: software for the autonomous computation of modularity in large network sets

Flavia Maria Darcie Marquitti, Paulo Roberto Guimarães Jr, Mathias Mistretta Pires and Luiz Fernando Bittencourt

**Ecography 37: 221–224, 2014**

doi: 10.1111/j.1600-0587.2013.00506.x

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Subject Editor: Thiago Rangel. Accepted 18 October 2013

A screenshot of the MODULAR software interface, which is a graphical user interface for computing modularity in large network sets. The interface is organized into several sections: 

- Networks:** Contains two radio buttons: 'Bipartite' (selected) and 'Unipartite'.
- Input File:** Contains two radio buttons: 'Matrix' (selected) and 'UCINET'.
- Metric to optimize:** Contains two radio buttons: 'Newman & Girvan 2004' (selected) and 'Barber 2007'.
- Method:** Contains five radio buttons: 'Simulated Annealing' (selected), 'Spectral Partitioning', 'FastGreedy', 'Hybrid1: Spectral Partitioning + Simulated Annealing', and 'Hybrid2: Fast Greedy + Simulated Annealing'.
- Simulated annealing parameters:** A section with three input fields: 'Initial temperature factor (2.0):' with the value '2', 'Cooling factor (1.01 to 1.1):' with the value '1.01', and 'No. of iterations factor (1):' with the value '1'.
- Null model options:** A section with a checkbox 'Run Null Models?' and a text input field 'How many times:'.
- Networks folder:** A section with a 'Select folder ...' button and an 'Open Folder' button.
- Bottom controls:** A section with a stick figure icon, a 'Run' button, and a 'Stop' button.

File	N.modules	Modularity
q_modelo 1 1 40 .txt	2	0.500000
q_modelo 1 10 50 .txt	4	0.533504
q_modelo 1 100 42 .txt	2	0.500000
q_modelo 1 11 38 .txt	4	0.645329
q_modelo 1 12 38 .txt	4	0.530181
q_modelo 1 13 47 .txt	2	0.500000
q_modelo 1 14 55 .txt	4	0.595577
q_modelo 1 15 40 .txt	4	0.548201
q_modelo 1 16 23 .txt	2	0.500000
q_modelo 1 17 44 .txt	3	0.501417
q_modelo 1 18 39 .txt	4	0.627704
q_modelo 1 19 35 .txt	4	0.509849
q_modelo 1 2 50 .txt	2	0.500000
q_modelo 1 20 44 .txt	4	0.607114
q_modelo 1 21 40 .txt	4	0.672936
q_modelo 1 22 48 .txt	4	0.523752
q_modelo 1 23 41 .txt	2	0.500000
q_modelo 1 24 27 .txt	4	0.506483
q_modelo 1 25 27 .txt	4	0.514467
q_modelo 1 26 46 .txt	4	0.524399
q_modelo 1 27 44 .txt	2	0.500000
q_modelo 1 28 43 .txt	4	0.540542
q_modelo 1 29 54 .txt	2	0.500000
q_modelo 1 3 27 .txt	4	0.526535
q_modelo 1 30 41 .txt	2	0.500000
q_modelo 1 31 48 .txt	3	0.501357
q_modelo 1 32 45 .txt	4	0.588084
q_modelo 1 33 40 .txt	4	0.555611
q_modelo 1 34 40 .txt	2	0.500000

Node	Module
R1	2
C1	5
C5	0
C7	1
C8	4
C13	4
C16	2
C17	2
C19	2
C22	0
C23	2
C24	0
C25	2
C28	1
C29	0
C33	0
C39	5
C44	5
C46	2
C47	2
R2	0
C18	0
C26	0
R3	2
C9	3
C12	1
C30	2
C31	4

## Package ‘rnetcarto’

November 12, 2015

**Type** Package

**Title** Fast Network Modularity and Roles Computation by Simulated Annealing (Rgraph C Library Wrapper for R)

**Version** 0.2.4

**Date** 2015-11-11

**Maintainer** Guilhem Doulcier <guilhem.doulcier@ens.fr>

**Description** It provides functions to compute the modularity and modularity-related roles in networks. It is a wrapper around the rgraph library (Guimera & Amaral, 2005, doi:10.1038/nature03288).

## Package 'rnetcarto'

```
## [[1]]
##   name module connectivity participation      role
## 8    h      0   -1.4142136      0.0000000 Ultra peripheral
## 5    d      0    0.7071068      0.0000000 Ultra peripheral
## 4    c      0    0.7071068      0.6400000      Connector
## 2    b      1   -0.7071068      0.5000000      Peripheral
## 6    f      1   -0.7071068      0.6666667      Connector
## 9    i      1    1.4142136      0.0000000 Ultra peripheral
## 1    a      2   -0.7071068      0.0000000 Ultra peripheral
## 7    g      2   -0.7071068      0.5000000      Peripheral
## 3    b      2    1.4142136      0.4444444      Peripheral
##
## [[2]]
## [1] 0.2024793
```

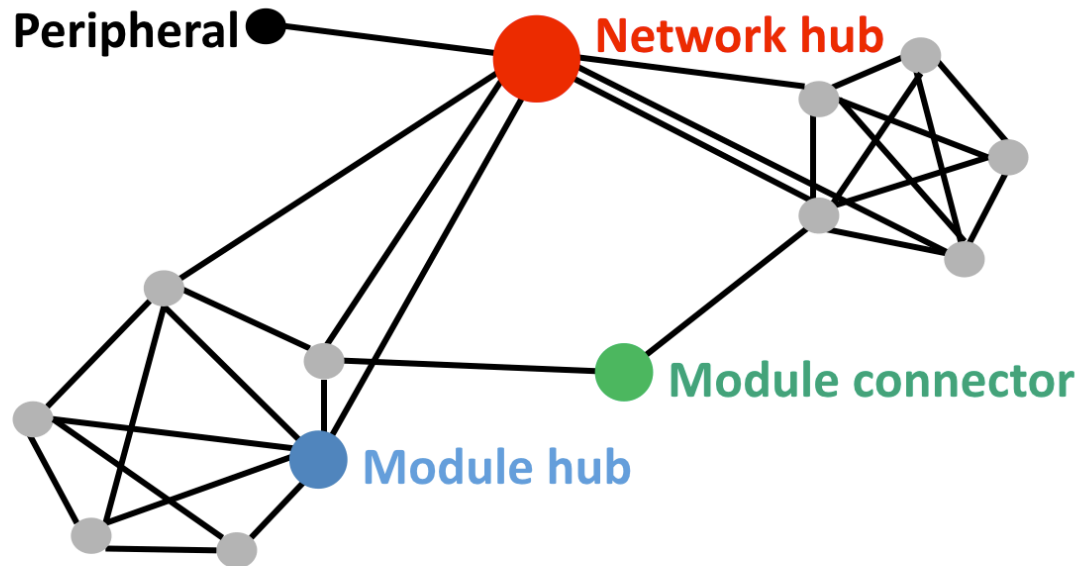
## Functional cartography of complex metabolic networks

**Roger Guimerà and Luís A. Nunes Amaral**

*NICO and Department of Chemical and Biological Engineering, Northwestern University, Evanston, Illinois 60208, USA*

## The modularity of pollination networks

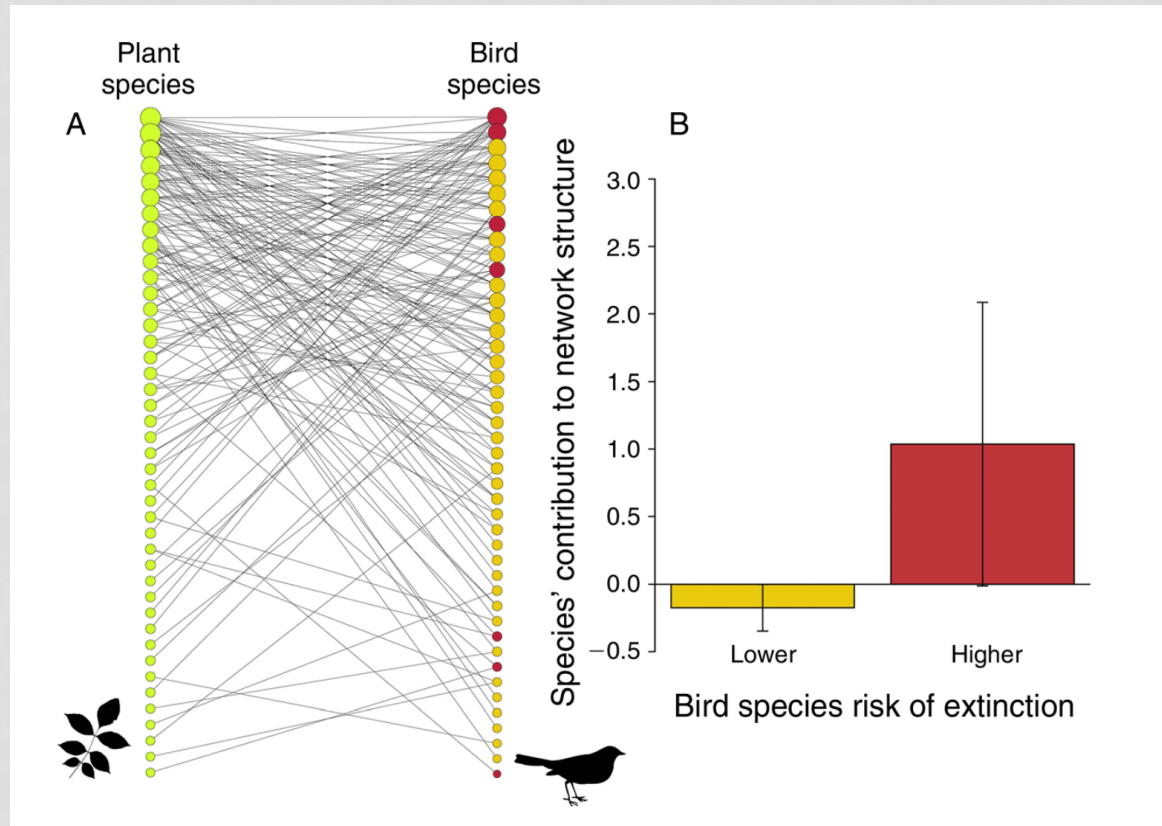
Jens M. Olesen<sup>\*†</sup>, Jordi Bascompte<sup>‡</sup>, Yoko L. Dupont<sup>\*</sup>, and Pedro Jordano<sup>‡</sup>



*Ecology*, 95(12), 2014, pp. 3440–3447  
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## Frugivores at higher risk of extinction are the key elements of a mutualistic network

MARIANA M. VIDAL,<sup>1</sup> ERICA HASUI,<sup>2</sup> MARCO A. PIZO,<sup>3</sup> JORGE Y. TAMASHIRO,<sup>4</sup> WESLEY R. SILVA,<sup>5</sup>  
AND PAULO R. GUIMARÃES, JR.<sup>1,6</sup>

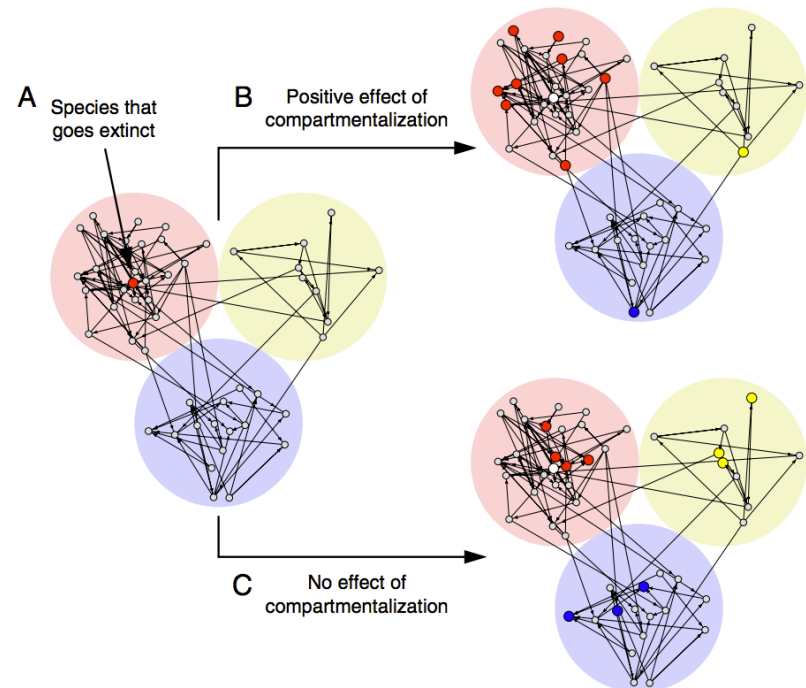


# Strong contributors to network persistence are the most vulnerable to extinction

Serguei Saavedra<sup>1,2,3\*</sup>, Daniel B. Stouffer<sup>4,5\*</sup>, Brian Uzzi<sup>1,2</sup> & Jordi Bascompte<sup>4</sup>

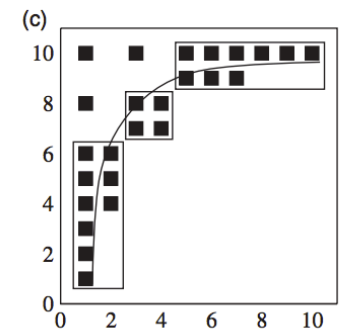
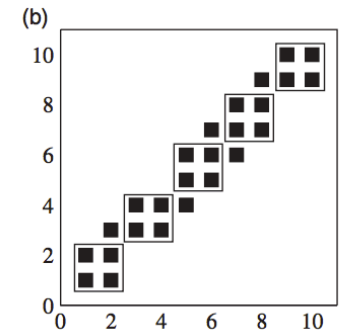
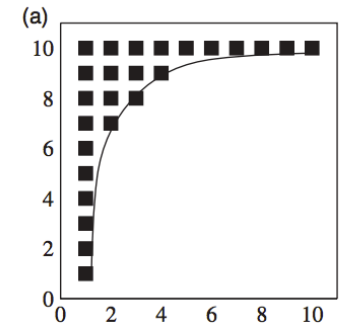
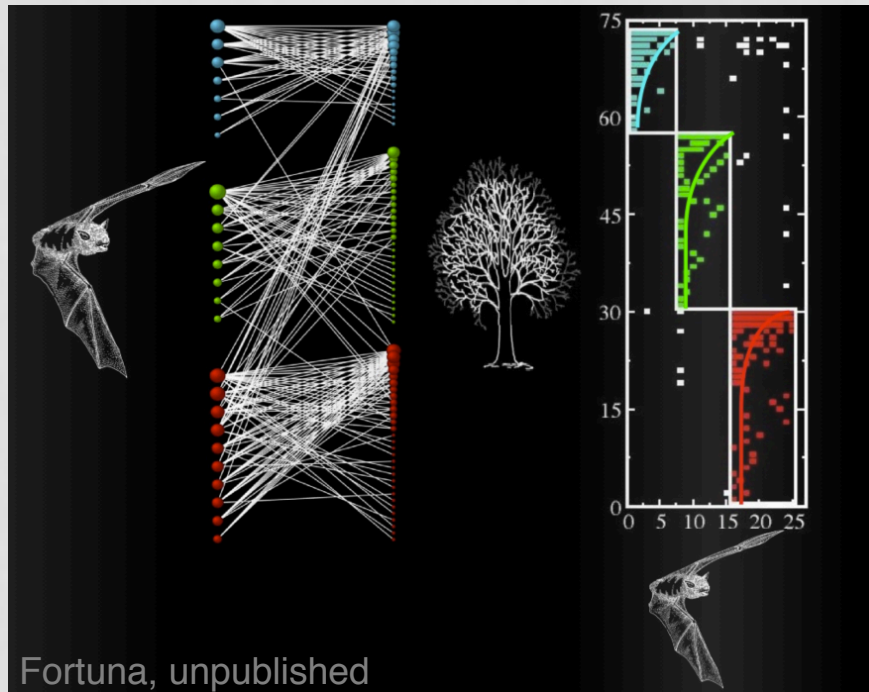
## Compartmentalization increases food-web persistence

Daniel B. Stouffer<sup>1</sup> and Jordi Bascompte



# Nestedness versus modularity in ecological networks: two sides of the same coin?

Miguel A. Fortuna<sup>1\*</sup>, Daniel B. Stouffer<sup>1</sup>, Jens M. Olesen<sup>2</sup>, Pedro Jordano<sup>1</sup>, David Mouillot<sup>3</sup>, Boris R. Krasnov<sup>4</sup>, Robert Poulin<sup>5</sup> and Jordi Bascompte<sup>1</sup>



**Plant-Seed Disperser**

<b>Network</b>	<b>Size</b>	<b>Connectance</b>	<b>Nestedness</b>	<b>Modularity</b>
1	28	0.085	* 0.763	0.311
2	58	0.106	** 0.944	0.312
3	78	0.026	** 0.842	0.308
4	26	0.264	* 0.847	0.121

**Plant-Pollinator**

<b>Network</b>	<b>Size</b>	<b>Connectance</b>	<b>Nestedness</b>	<b>Modularity</b>
23	61	0.090	** 0.925 *	0.591 **
24	185	0.043	** 0.960	0.516 **
25	107	0.071	** 0.907	0.519 **
26	90	0.098	** 0.811 *	** 0.569 **

**Host-Parasite**

<b>Network</b>	<b>Size</b>	<b>Connectance</b>	<b>Nestedness</b>	<b>Modularity</b>
57	35	0.247	** 0.819	** 0.516 **
58	36	0.384	** 0.662	0.268
59	45	0.217	** 0.783	** 0.437 **
60	46	0.191	** 0.749	0.312