

Ecosystems 2010: Global Progress on Ecosystem-based Fisheries Management
2010 Wakefield Fishery Symposium, Anchorage, USA

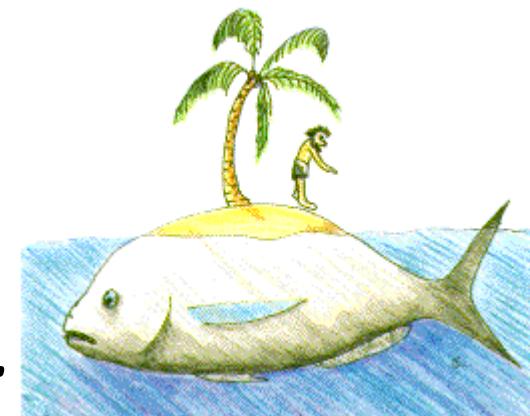
A New Topological Approach to Quantify the Interaction Structure of Food Web Network

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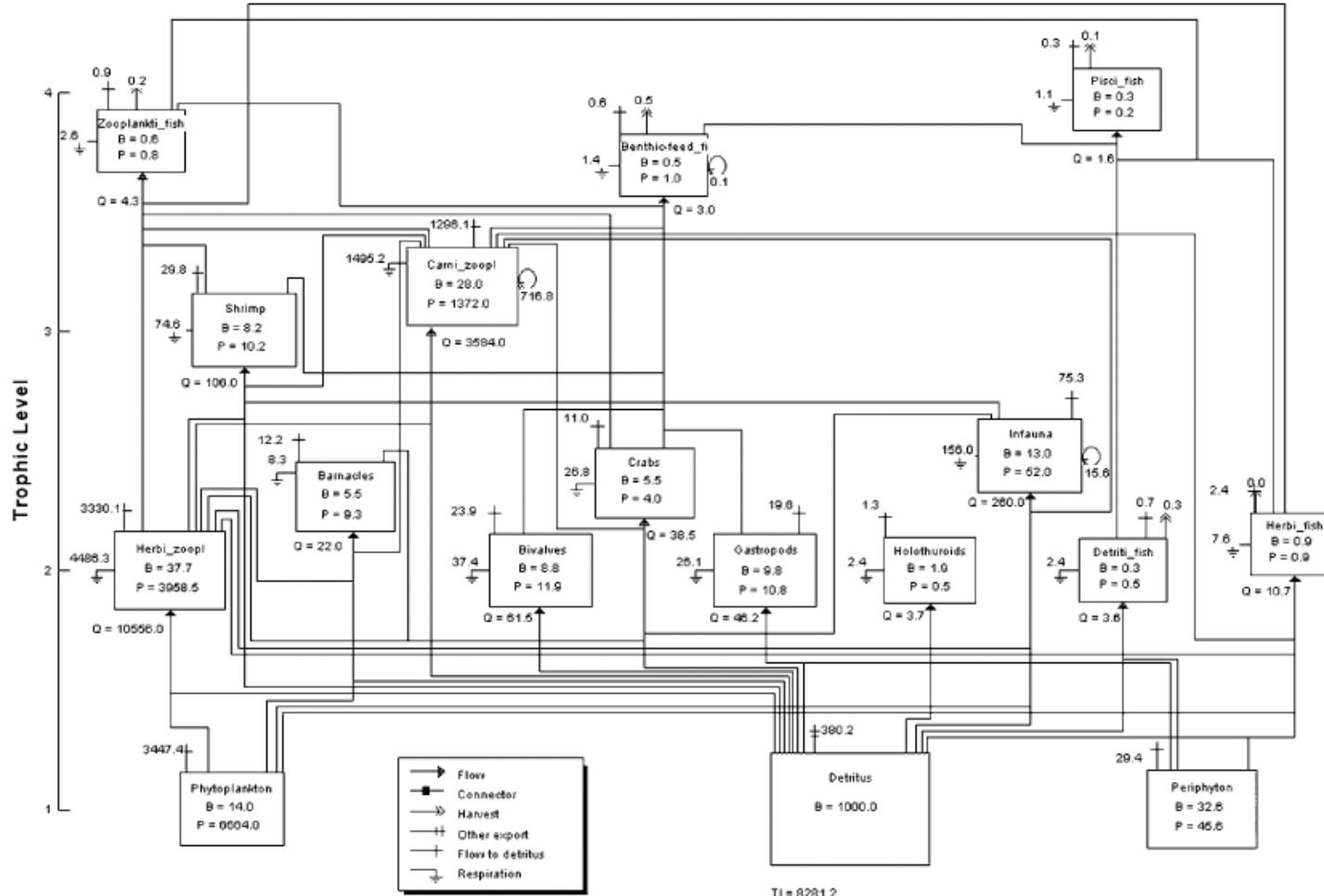
Ecosystem-Based Approach to me...

- 1. Every component in the system is interacting with all others;**
- 2. Catching a fish is not only about the fish you caught, but also effecting its preys, predators, competitors...**



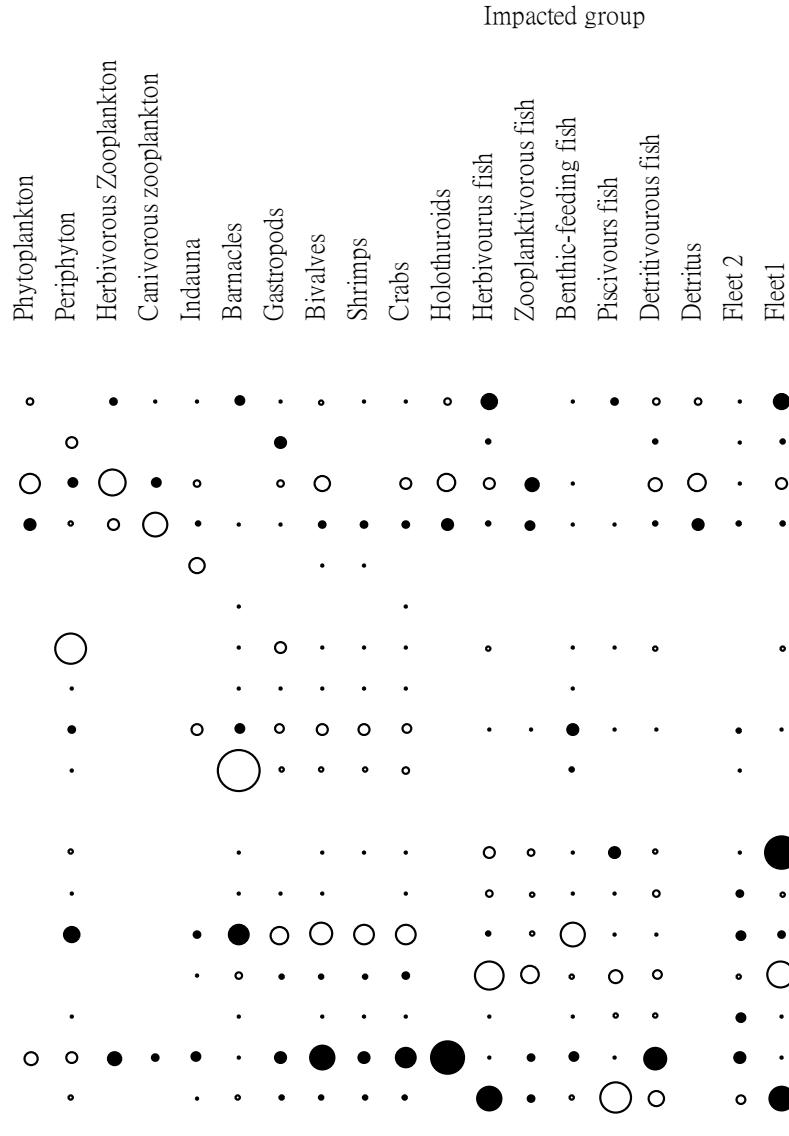
No fish is an island...

Mass-balanced food web modeling (i.e. Ecopath with Ecosim)



Mixed Trophic Index, MTI

(EwE v6: Network Analysis)



Based on Lin et al. 2004

- Positive
- Negative

- Impacting group
- Phytoplankton
 - Periphyton
 - Herbivorous Zooplankton
 - Canivorous zooplankton
 - Indauna
 - Barnacles
 - Gastropods
 - Bivalves
 - Shrimps
 - Crabs
 - Holothuroids
 - Herbivourus fish
 - Zooplanktivorous fish
 - Benthic-feeding fish
 - Piscivours fish
 - Detritivourous fish
 - Detritus
 - Fleet 2
 - Fleet1

However, Ecopath modeling do require a lot of Parameterizations

Biomass (B, g C m⁻²)

Production (P, g C m⁻² d⁻¹)

Consumption (Q, g C m⁻² d⁻¹)

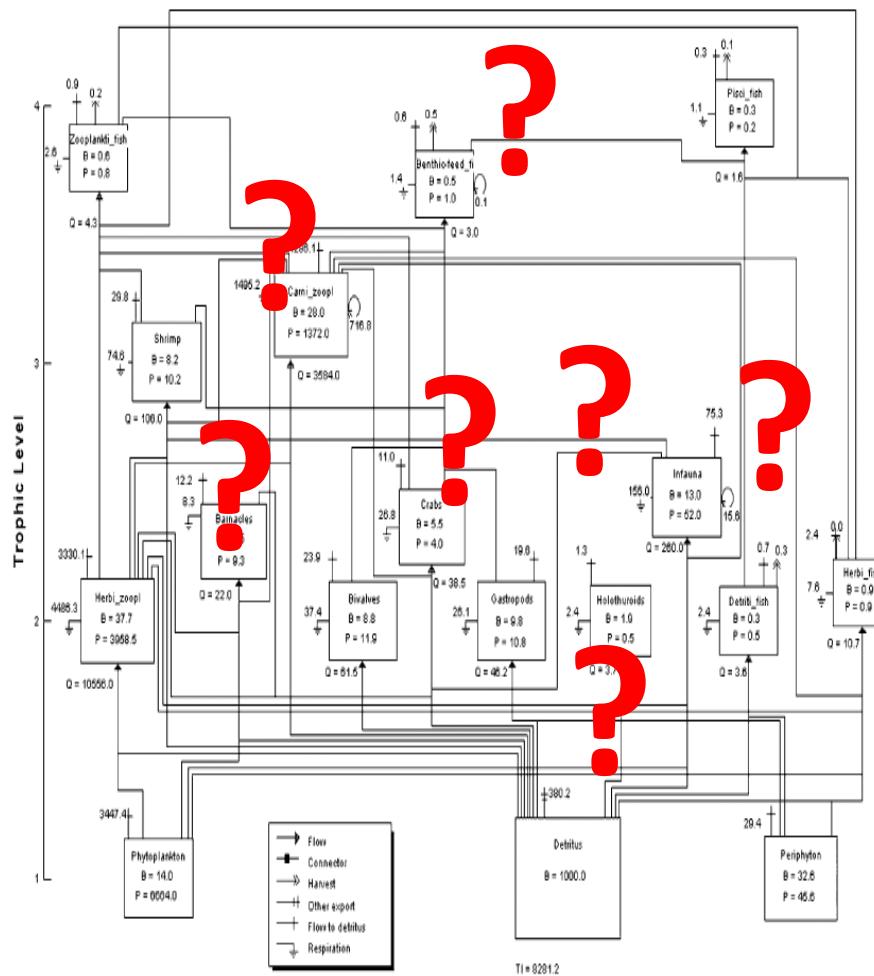
Catch

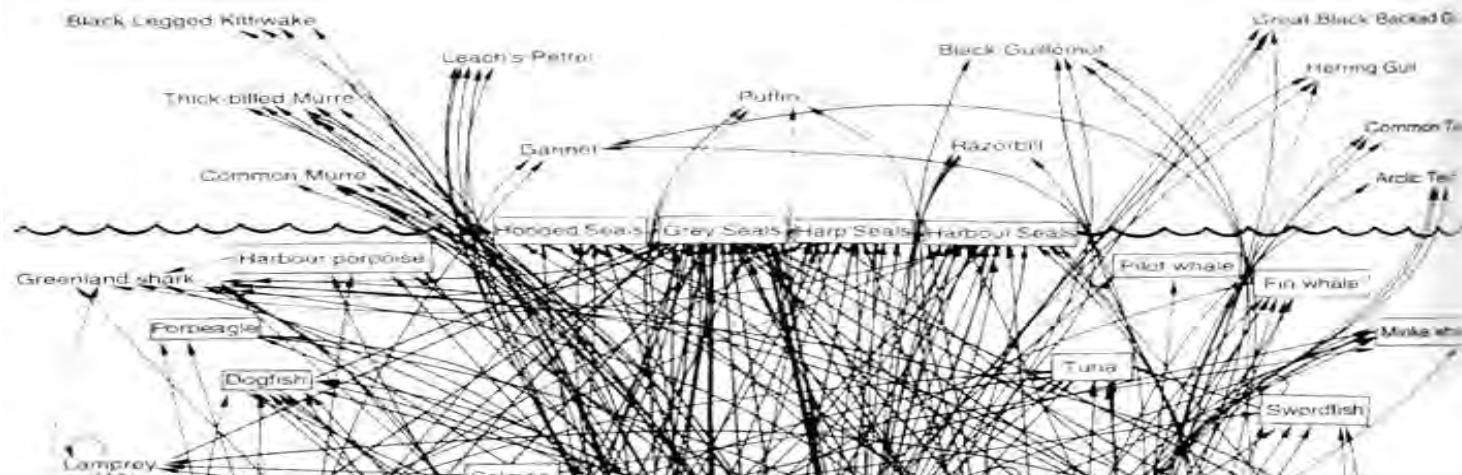
Ecotrophic Efficiency (EE)

Diet composition

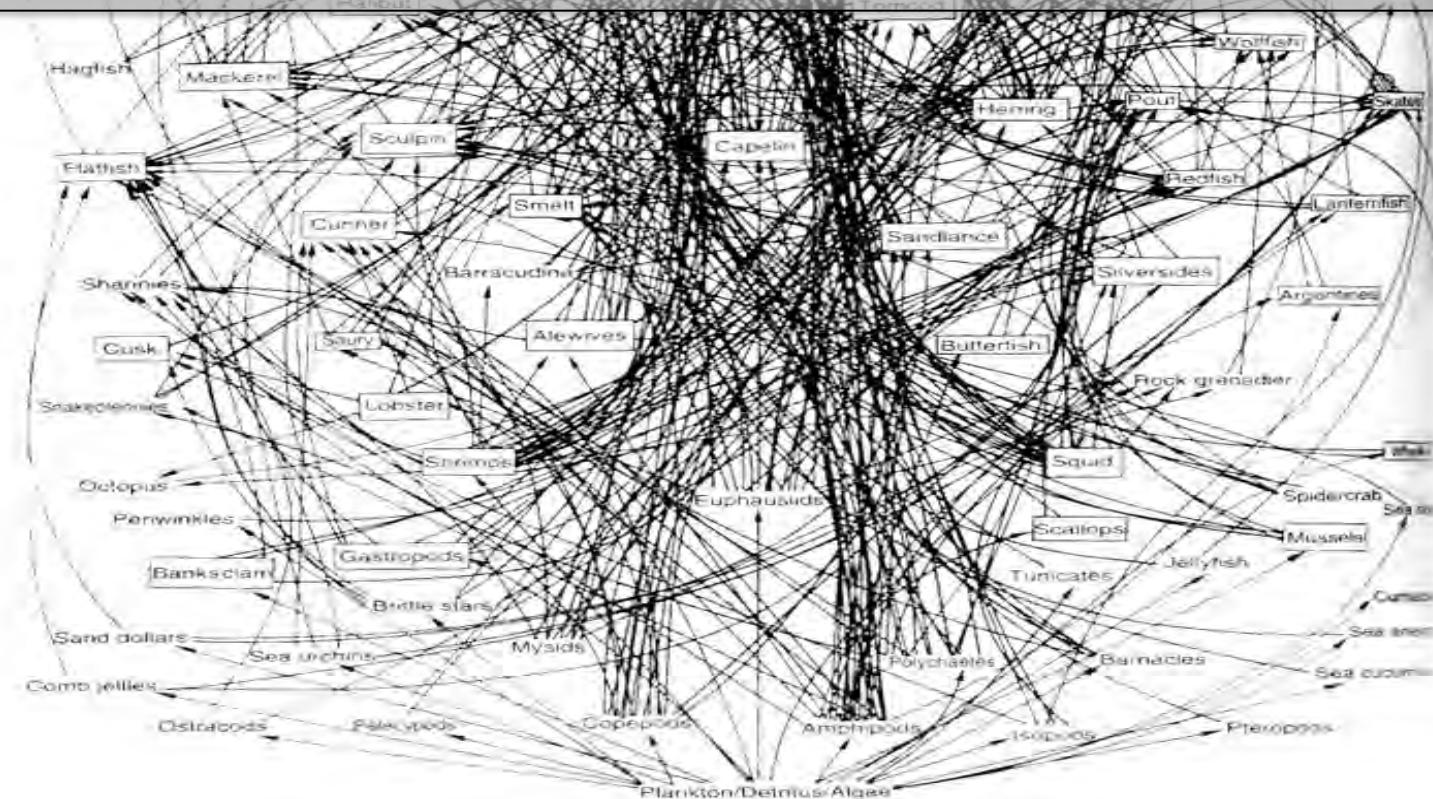
Organic detritus (g C m⁻²)

.....

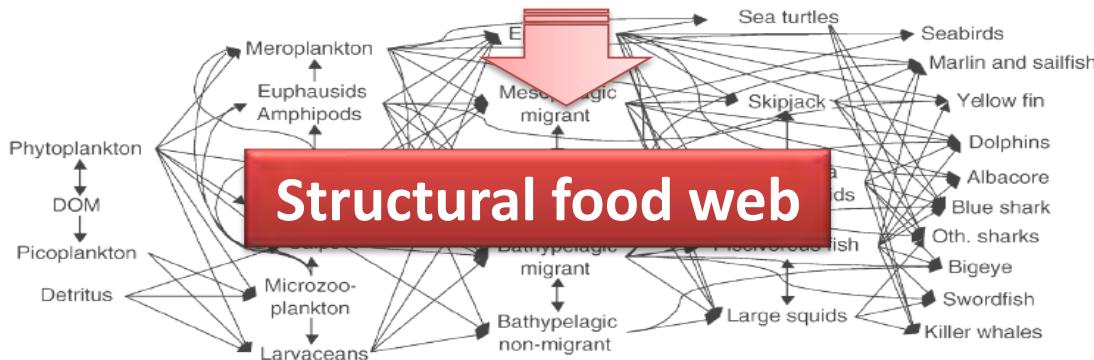




Too complex to parameterization?

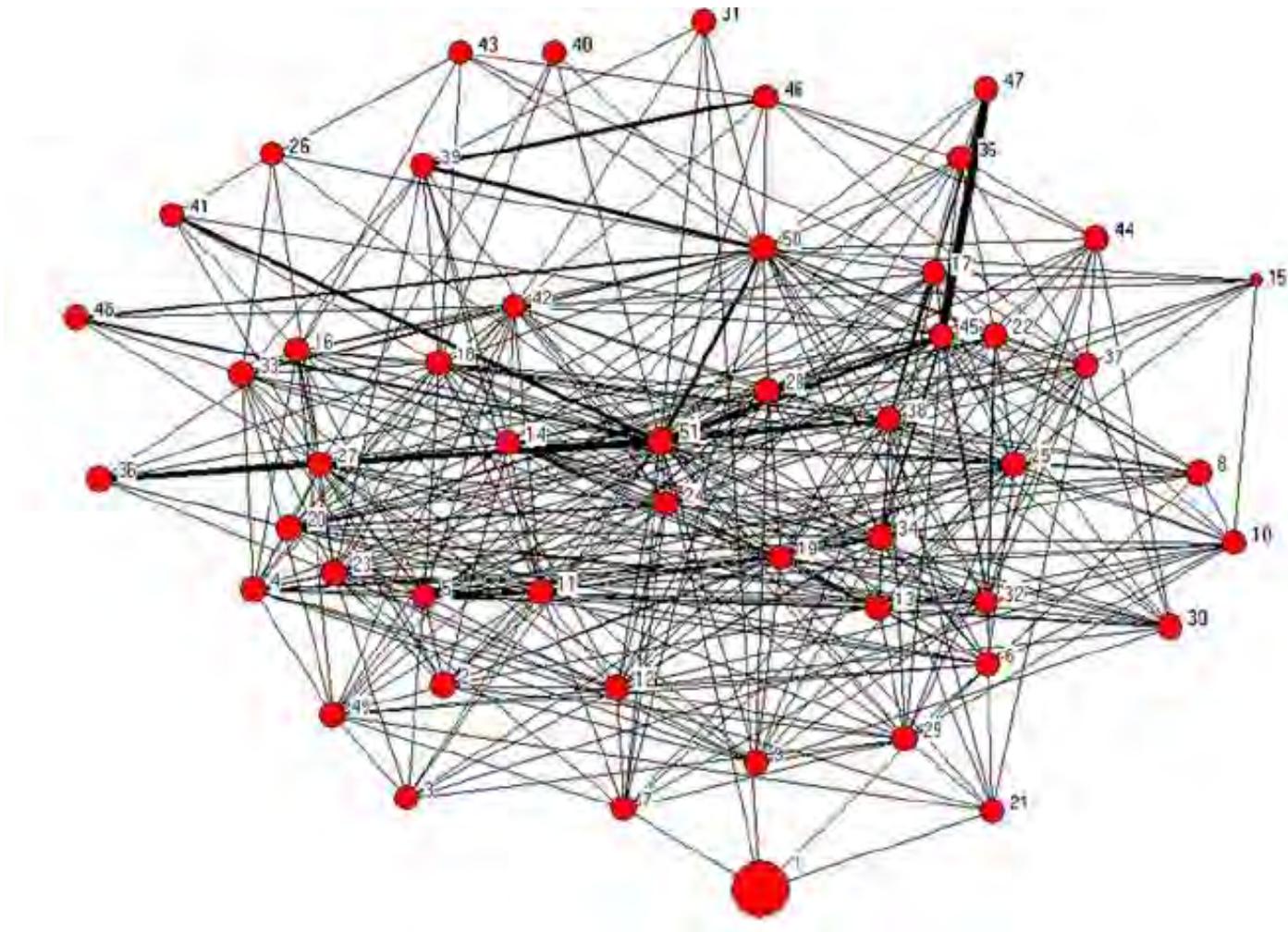


Simply knowing only Who Eats Whom should be much easier...



Ex: Building structural food web using DNA barcoding

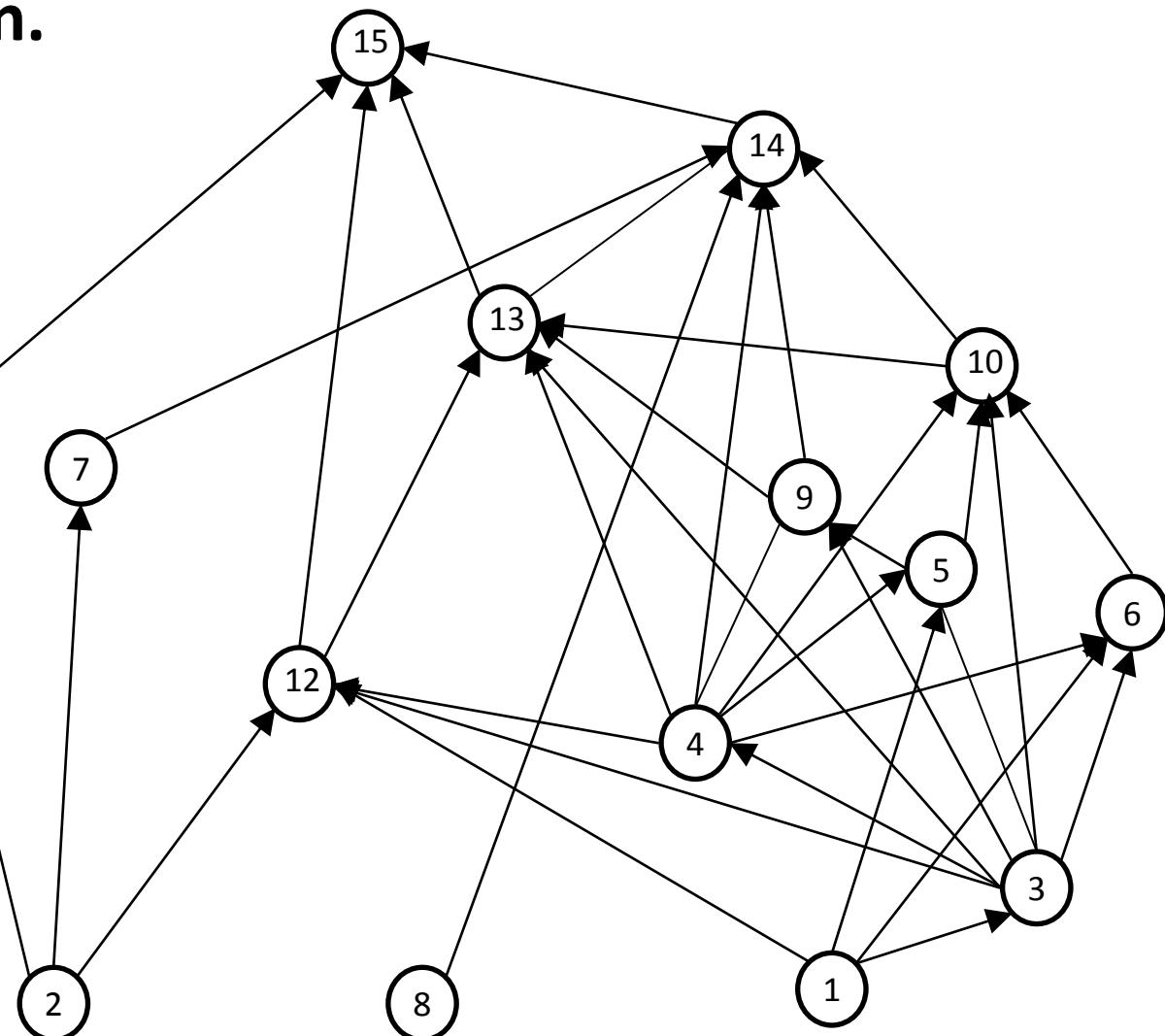
What can we learn from the structural food web along?





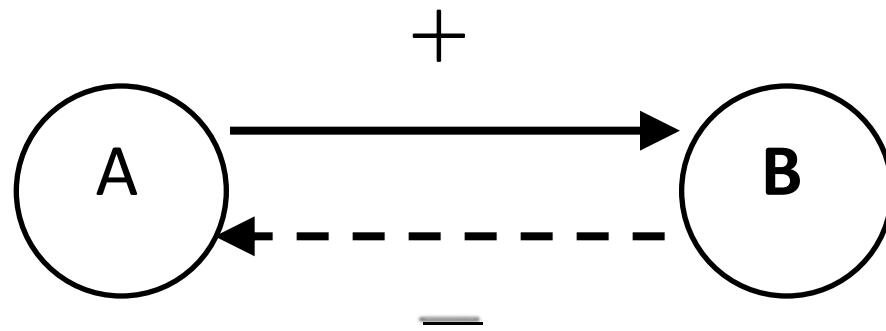
Consider a simple 15 species food web of a costal ecosystem; Kuoshen Bay in NE Taiwan.

- (1) phytoplankton,
- (2) periphyton,
- (3) herbivorous zooplankton,
- (4) carnivorous zooplankton,
- (5) infauna,
- (6) barnacles,
- (7) gastropods,
- (8) bivalves,
- (9) shrimp,
- (10) crabs,
- (11) detritivorous fish,
- (12) herbivorous fish,
- (13) zooplanktivorous fish,
- (14) benthic-feeding fish, and
- (15) piscivorous fish.



Based on Lin et al. 2004 and after Jordan et al. 2009

1. Trophic interactions are directed and signed:



Simultaneously, every trophic interaction occurs with both positive and negative effect

2a. The total magnitude of **direct effect** on any focal species = 1

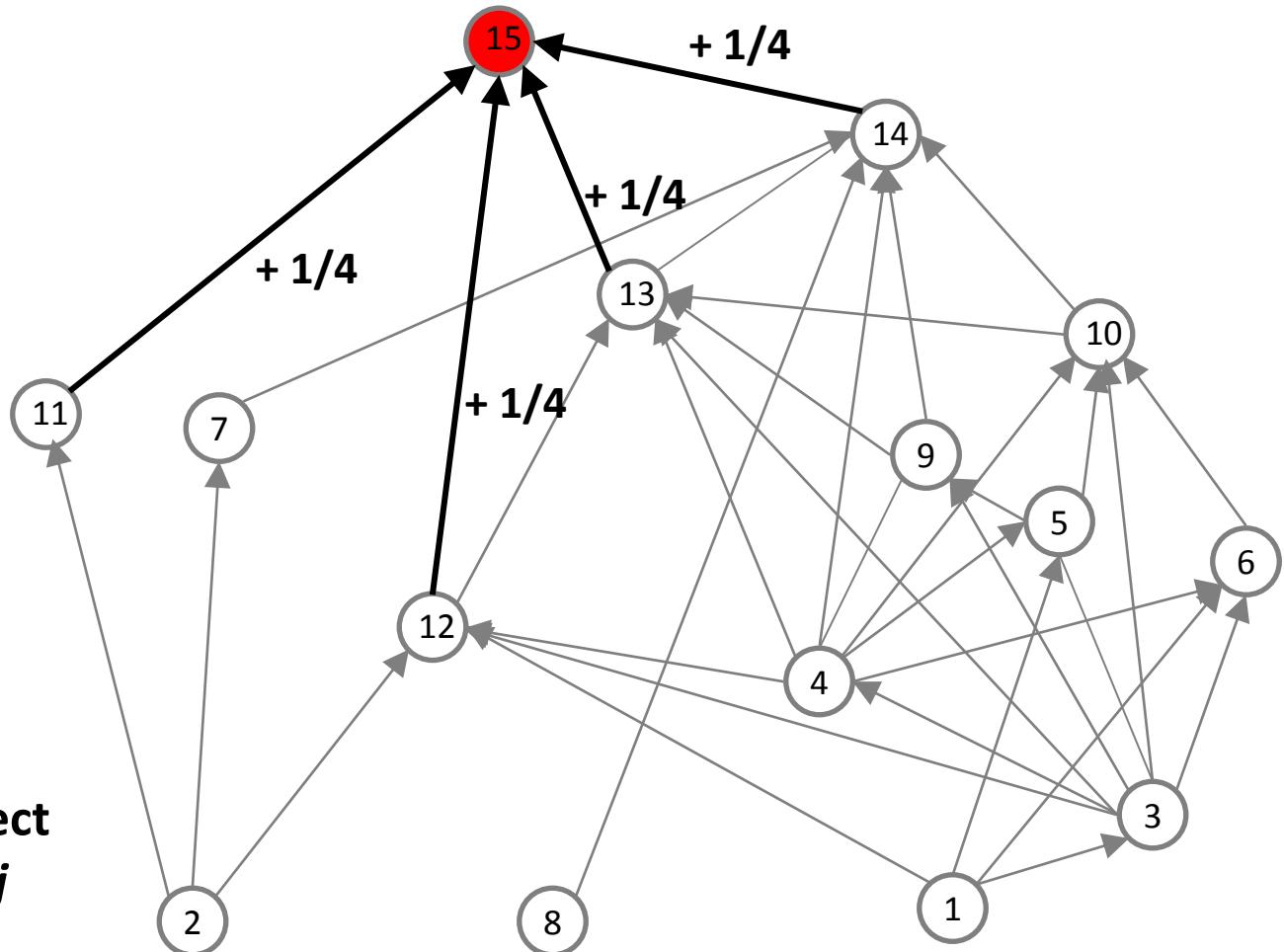
$$\sum_{j=1}^m a_{j \rightarrow i^s} = 1$$

If not weighted,

$$a_{j \rightarrow i^s} = \frac{1}{m}$$

a : magnitude of effect

m : no. neighbor of *j*



2b. The magnitudes of **direct effects** from a focal species depends on how many neighbors the effect receiver has

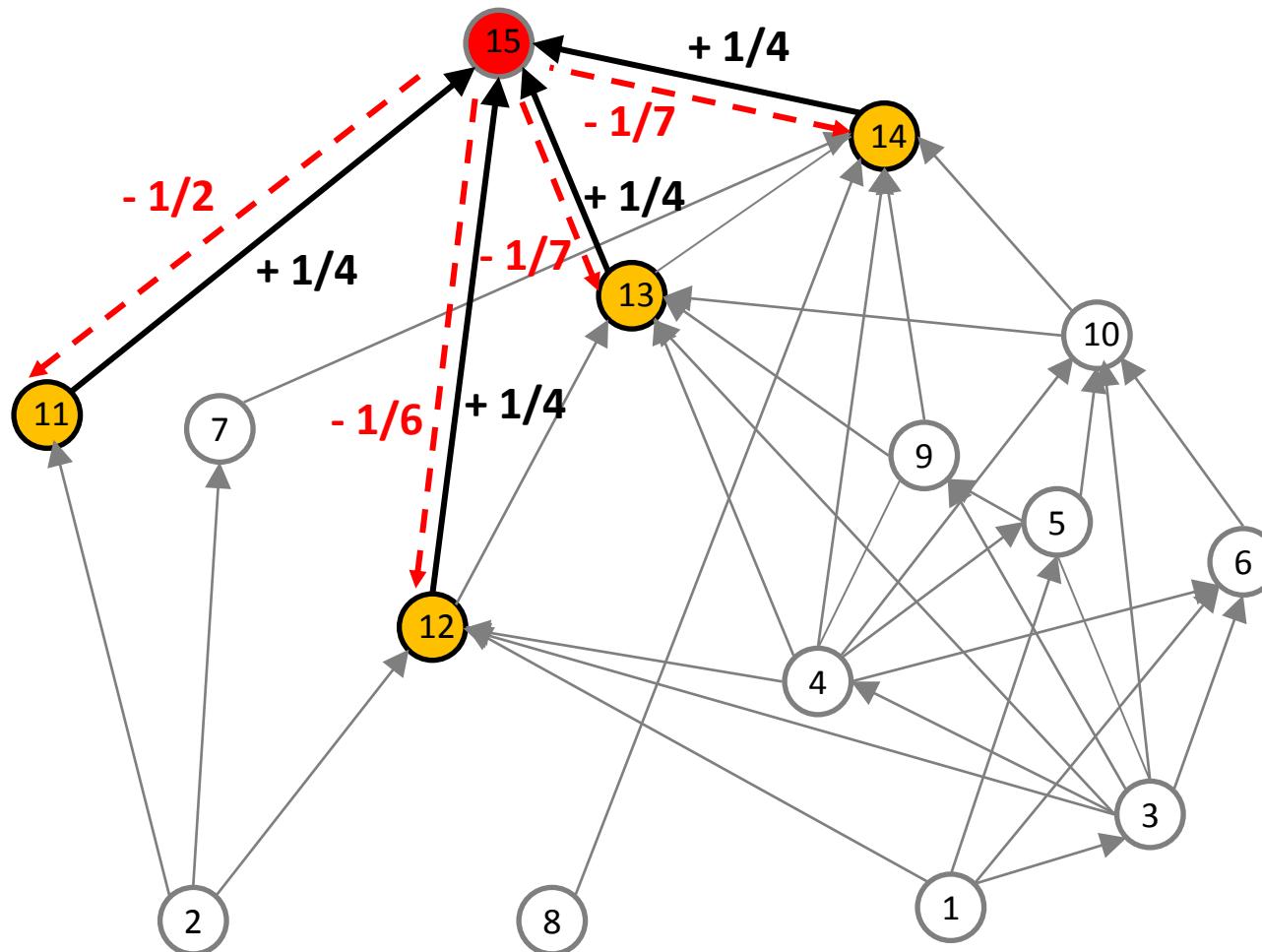


Table of 1-step positive (+) direct effects

	1	10	11	12	13	14	15	2	3	4	5	6	7	8	9
1	0.000	0.000	0.000	0.167	0.000	0.000	0.000	0.000	0.125	0.000	0.200	0.250	0.000	0.000	0.000
10	0.000	0.000	0.000	0.000	0.143	0.143	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
11	0.000	0.000	0.000	0.000	0.000	0.000	0.250	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
12	0.000	0.000	0.000	0.000	0.143	0.000	0.250	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
13	0.000	0.000	0.000	0.000	0.000	0.143	0.250	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
14	0.000	0.000	0.000	0.000	0.000	0.000	0.250	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
15	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
2	0.000	0.000	0.500	0.167	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.500	0.000	0.000
3	0.000	0.167	0.000	0.167	0.143	0.000	0.000	0.000	0.000	0.125	0.200	0.250	0.000	0.000	0.200
4	0.000	0.167	0.000	0.167	0.143	0.143	0.000	0.000	0.000	0.000	0.200	0.250	0.000	0.000	0.200
5	0.000	0.167	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.200
6	0.000	0.167	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
7	0.000	0.000	0.000	0.000	0.000	0.143	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
8	0.000	0.000	0.000	0.000	0.000	0.143	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
9	0.000	0.000	0.000	0.000	0.143	0.143	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000

Table of 1-step negative (-) direct effects

	1	10	11	12	13	14	15	2	3	4	5	6	7	8	9	Row Sum
1	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	
10	0.000	0.000	0.000	0.000	0.000	0.000	0.000	-0.125	-0.125	-0.200	-0.250	0.000	0.000	0.000	-0.700	
11	0.000	0.000	0.000	0.000	0.000	0.000	-0.333	0.000	0.000	0.000	0.000	0.000	0.000	0.000	-0.333	
12	-0.250	0.000	0.000	0.000	0.000	0.000	-0.333	-0.125	-0.125	0.000	0.000	0.000	0.000	0.000	-0.833	
13	0.000	-0.167	0.000	-0.167	0.000	0.000	0.000	0.000	-0.125	-0.125	0.000	0.000	0.000	-0.200	-0.783	
14	0.000	-0.167	0.000	0.000	-0.143	0.000	0.000	0.000	0.000	-0.125	0.000	0.000	-0.500	-1.000	-0.200	-2.133
15	0.000	0.000	-0.500	-0.167	-0.143	-0.143	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	-0.950	
2	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	
3	-0.250	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	-0.250	
4	0.000	0.000	0.000	0.000	0.000	0.000	0.000	-0.125	0.000	0.000	0.000	0.000	0.000	0.000	-0.125	
5	-0.250	0.000	0.000	0.000	0.000	0.000	0.000	0.000	-0.125	-0.125	0.000	0.000	0.000	0.000	-0.500	
6	-0.250	0.000	0.000	0.000	0.000	0.000	0.000	0.000	-0.125	-0.125	0.000	0.000	0.000	0.000	-0.500	
7	0.000	0.000	0.000	0.000	0.000	0.000	-0.333	0.000	0.000	0.000	0.000	0.000	0.000	0.000	-0.333	
8	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	
9	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	-0.125	-0.125	-0.200	0.000	0.000	0.000	-0.450	

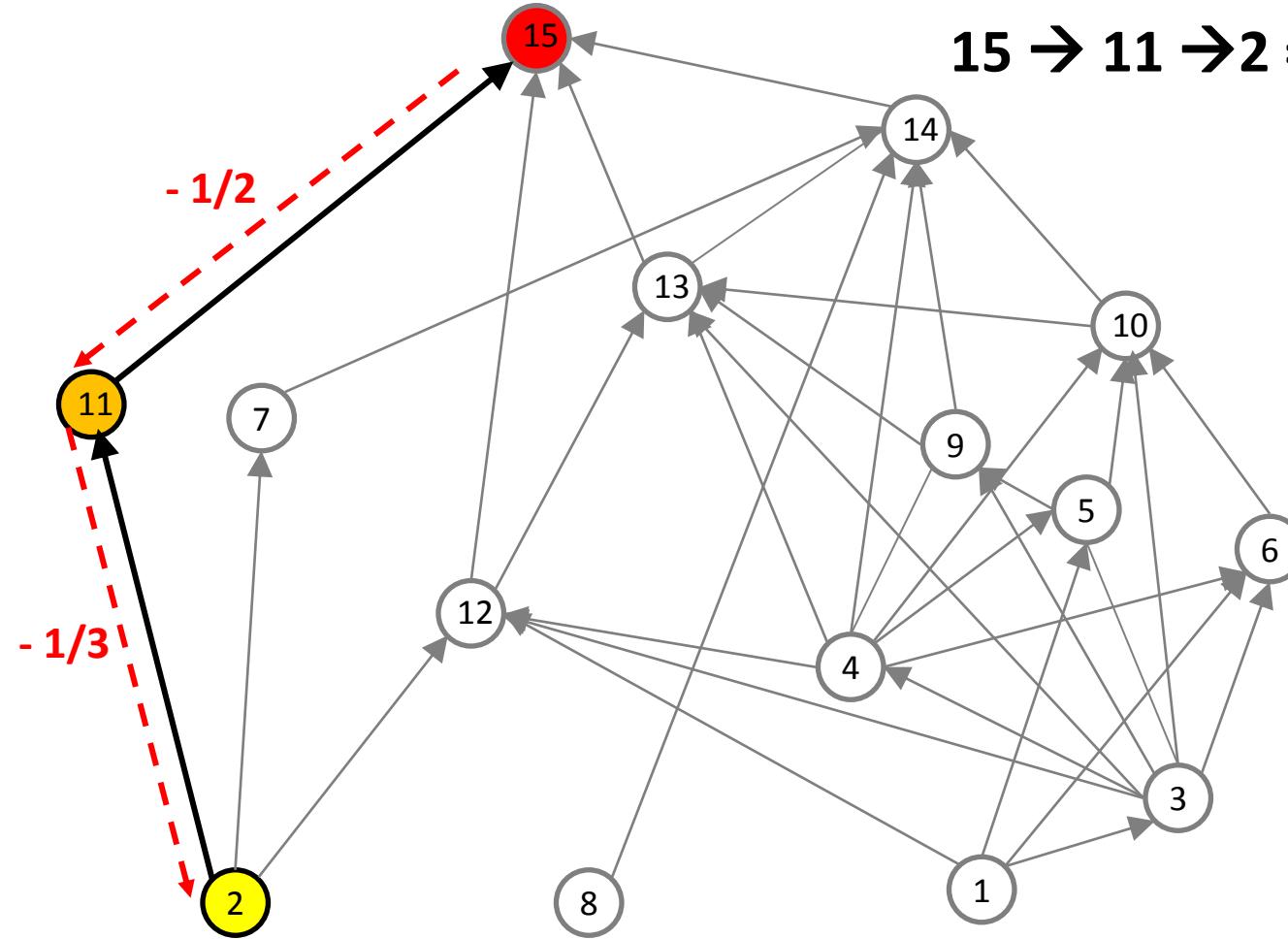
Table of 1-step net direct effects

	1	10	11	12	13	14	15	2	3	4	5	6	7	8	9	Row Sum
1	0.000	0.000	0.000	0.167	0.000	0.000	0.000	0.000	0.125	0.000	0.200	0.250	0.000	0.000	0.000	0.743
10	0.000	0.000	0.000	0.000	0.143	0.143	0.000	0.000	-0.125	-0.125	-0.200	-0.250	0.000	0.000	0.000	-0.414
11	0.000	0.000	0.000	0.000	0.000	0.000	0.250	-0.333	0.000	0.000	0.000	0.000	0.000	0.000	0.000	-0.083
12	-0.250	0.000	0.000	0.000	0.143	0.000	0.250	-0.333	-0.125	-0.125	0.000	0.000	0.000	0.000	0.000	-0.444
13	0.000	-0.167	0.000	-0.167	0.000	0.143	0.250	0.000	-0.125	-0.125	0.000	0.000	0.000	0.000	-0.200	-0.393
14	0.000	-0.167	0.000	0.000	-0.143	0.000	0.250	0.000	0.000	-0.125	0.000	0.000	-0.500	-1.000	-0.200	-1.883
15	0.000	0.000	-0.500	-0.167	-0.143	-0.143	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	-0.953
2	0.000	0.000	0.500	0.167	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.500	0.000	0.000	1.167
3	-0.250	0.167	0.000	0.167	0.143	0.000	0.000	0.000	0.000	0.125	0.200	0.250	0.000	0.200	1.000	
4	0.000	0.167	0.000	0.167	0.143	0.143	0.000	0.000	-0.125	0.000	0.200	0.250	0.000	0.200	1.143	
5	-0.250	0.167	0.000	0.000	0.000	0.000	0.000	0.000	-0.125	-0.125	0.000	0.000	0.000	0.200	-0.133	
6	-0.250	0.167	0.000	0.000	0.000	0.000	0.000	0.000	-0.125	-0.125	0.000	0.000	0.000	0.000	-0.333	
7	0.000	0.000	0.000	0.000	0.000	0.143	0.000	-0.333	0.000	0.000	0.000	0.000	0.000	0.000	-0.190	
8	0.000	0.000	0.000	0.000	0.000	0.143	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.143	
9	0.000	0.000	0.000	0.000	0.143	0.143	0.000	0.000	-0.125	-0.125	-0.200	0.000	0.000	0.000	-0.167	

3a. The magnitude of n-step **indirect effect** is simply the product of the direct effects along the given n-step path

i.e. a 2-step effect Node 15 → 2 via 11

$$\begin{aligned}15 \rightarrow 11 \rightarrow 2 &= (15 \rightarrow 11) \times (11 \rightarrow 2) \\&= (-1/2) \times (-1/3) \\&= 1/6\end{aligned}$$



3b. The total magnitude of n-step indirect effect from i to j is simply **the sum of all n-step indirect effects** between $i \rightarrow j$

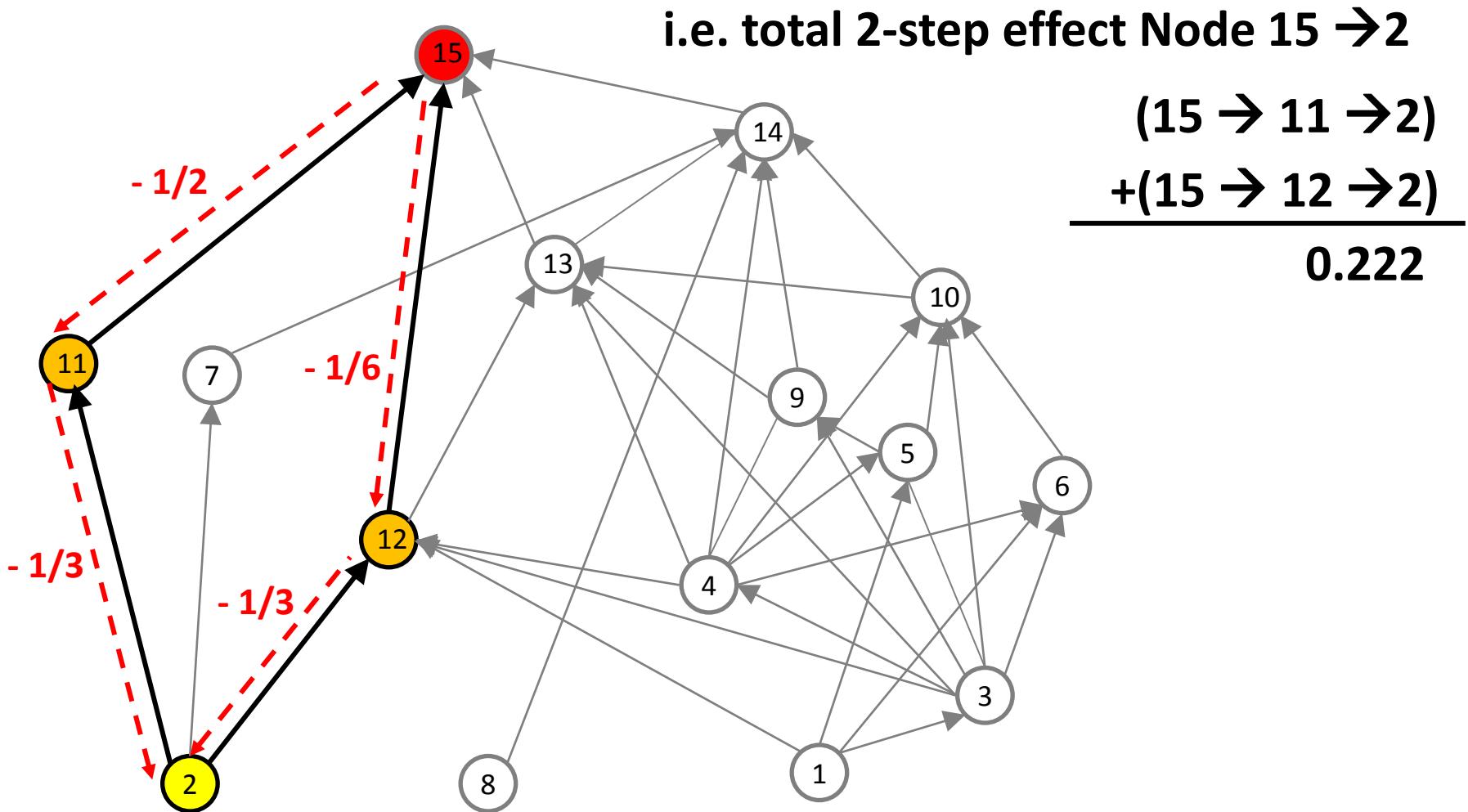
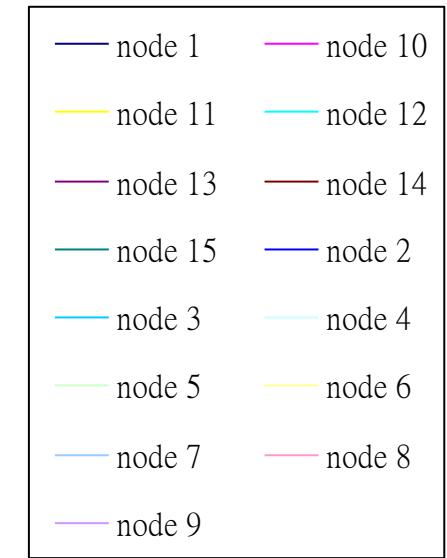
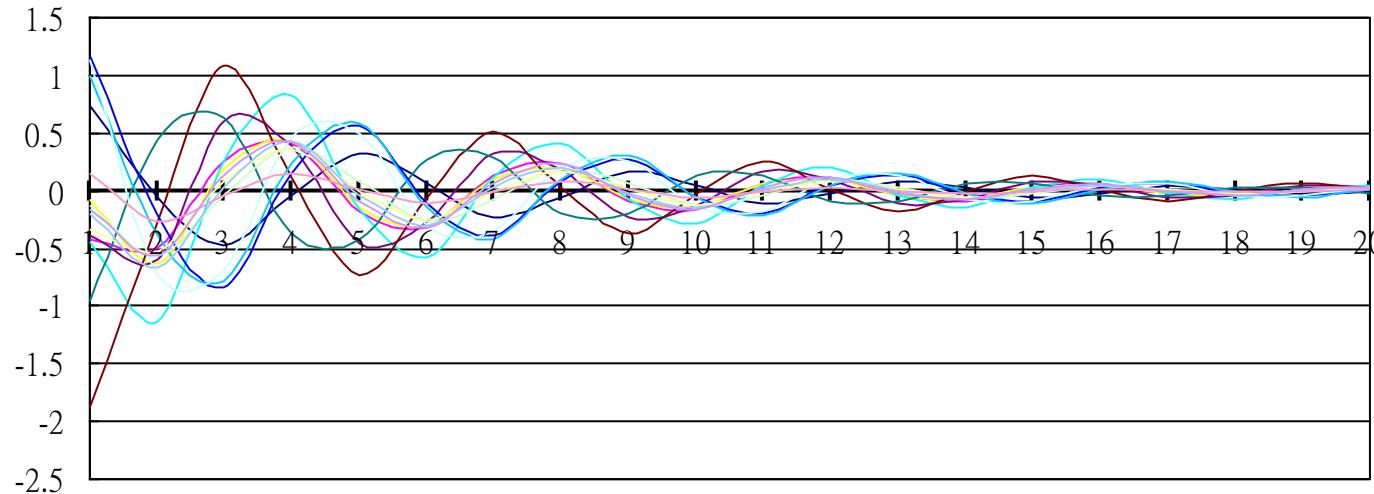


Table of 2-step net in-direct effect

	1	10	11	12	13	14	15	2	3	4	5	6	7	8	9
1	-0.185	0.096	0	0.021	0.042	0	0.042	-0.056	-0.077	-0.061	0.025	0.031	0	0	0.065
10	0.144	-0.164	0	-0.065	-0.056	0.002	0.071	0	0.054	0.005	-0.05	-0.063	-0.071	-0.143	-0.147
11	0	0	-0.292	-0.097	-0.036	-0.036	0	0	0	0	0	0	-0.167	0	0
12	0.031	-0.065	-0.292	-0.204	-0.071	-0.034	0.036	0	-0.033	-0.033	-0.1	-0.125	-0.167	0	-0.079
13	0.073	-0.065	-0.125	-0.083	-0.168	-0.106	-0.006	0.056	0.082	0.034	0.023	-0.021	-0.071	-0.143	-0.079
14	0	0.003	-0.125	-0.039	-0.106	-0.341	-0.036	0.167	0.079	0.064	0.048	0.011	0	0	0.004
15	0.042	0.048	0	0.024	-0.004	-0.02	-0.238	0.222	0.039	0.057	0	0	0.071	0.143	0.057
2	-0.042	0	0	0	0.024	0.071	0.167	-0.389	-0.021	-0.021	0	0	0	0	0
3	-0.154	0.072	0	-0.044	0.094	0.091	0.077	-0.056	-0.188	-0.141	-0.098	-0.073	0	0	0.036
4	-0.123	0.007	0	-0.045	0.038	0.073	0.113	-0.056	-0.141	-0.174	-0.098	-0.073	-0.071	-0.143	-0.042
5	0.031	-0.042	0	-0.083	0.016	0.034	0	0	-0.061	-0.061	-0.173	-0.167	0	0	-0.05
6	0.031	-0.042	0	-0.083	-0.012	0.006	0	0	-0.036	-0.036	-0.133	-0.167	0	0	-0.05
7	0	-0.024	-0.167	-0.056	-0.02	0	0.036	0	0	-0.018	0	0	-0.238	-0.143	-0.029
8	0	-0.024	0	0	-0.02	0	0.036	0	0	-0.018	0	0	-0.071	-0.143	-0.029
9	0.081	-0.123	0	-0.065	-0.056	0.002	0.071	0	0.023	-0.026	-0.05	-0.063	-0.071	-0.143	-0.147

The cumulative net effects will converge...

Net effects generated by each node from step1 to 20 in length



Cumulative effects generated by each node from step1 to 20 in length

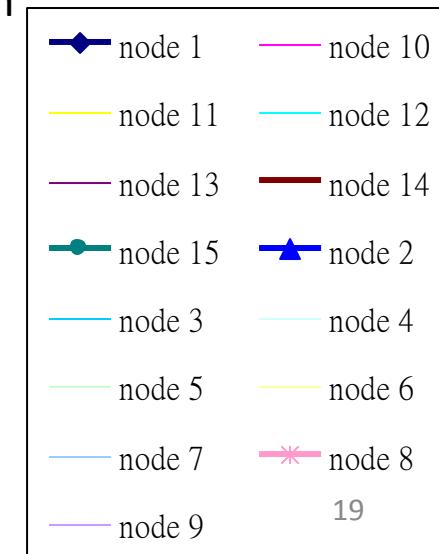
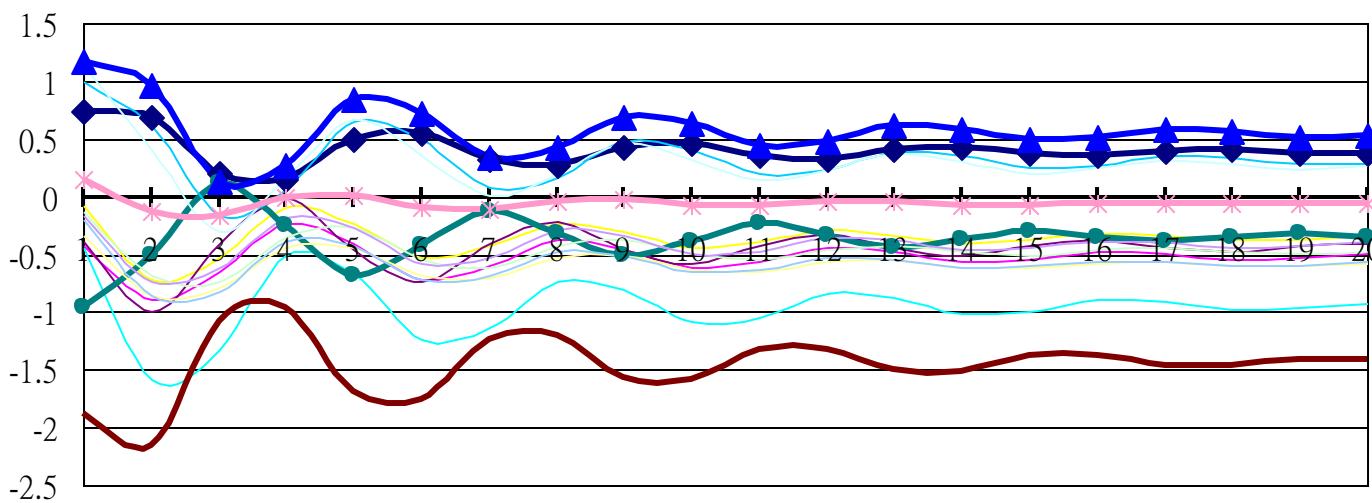
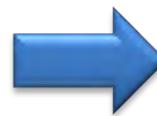
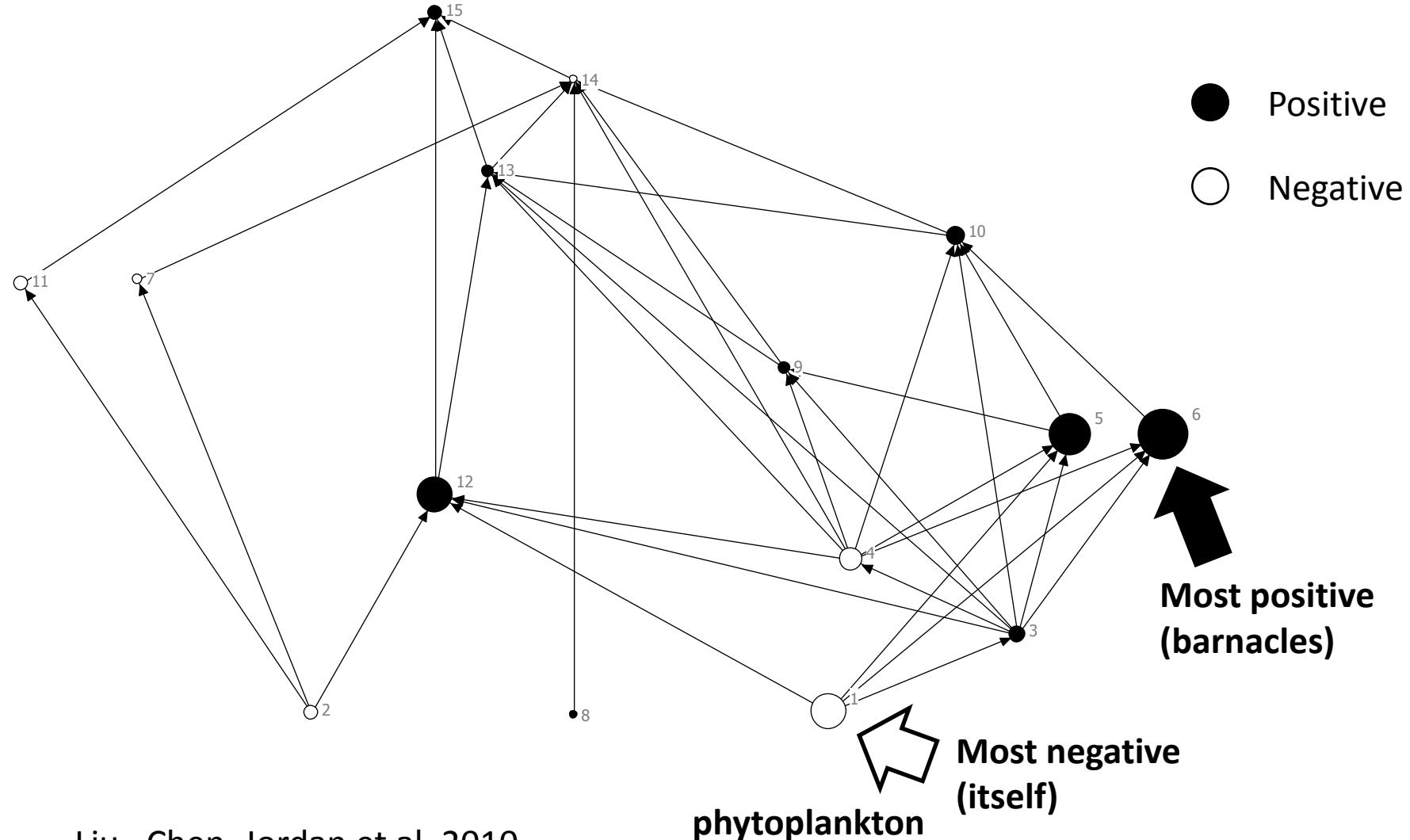


Table of the cumulative net effects from 1 to 20 steps

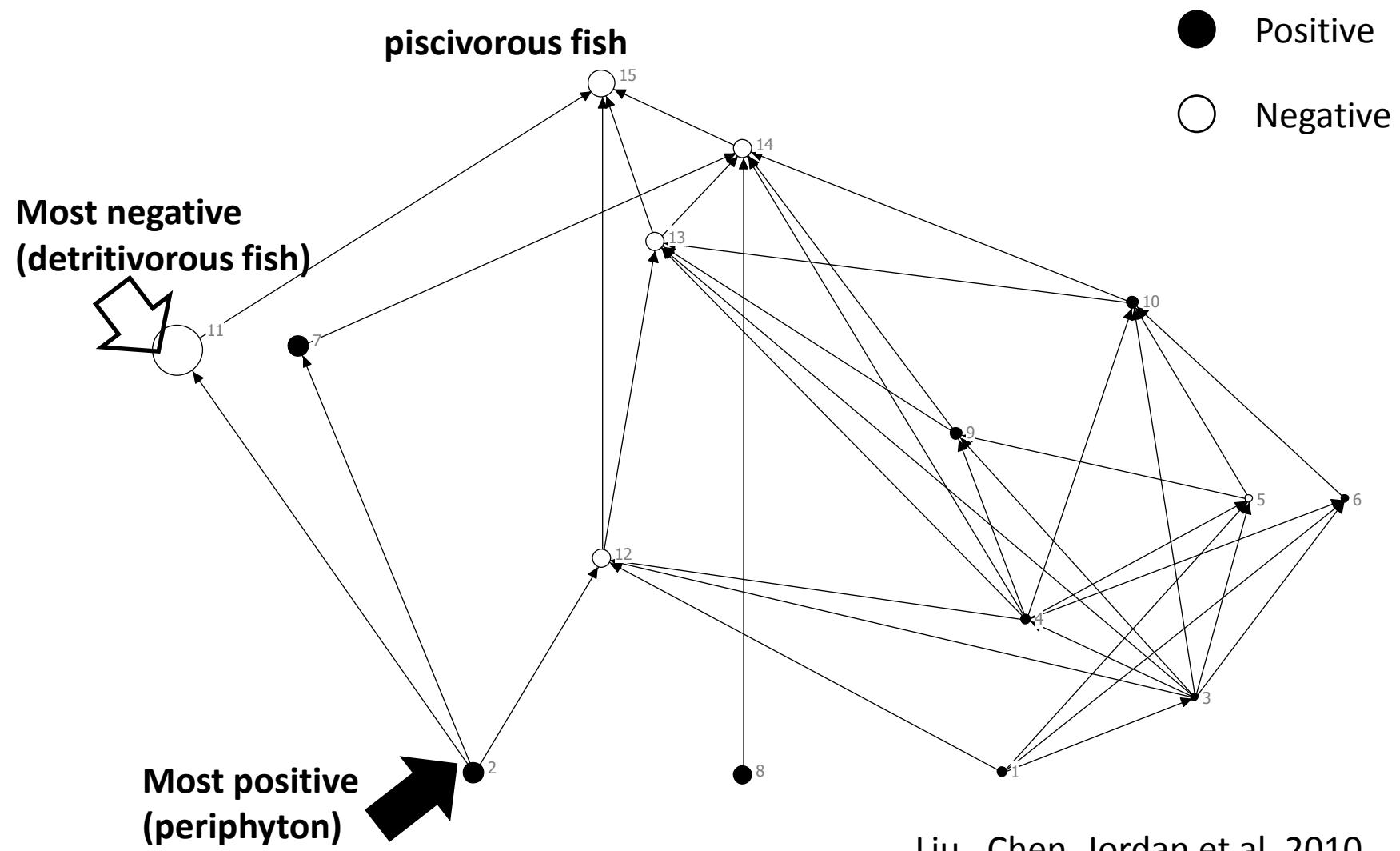
	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
1	-0.13	-0.03	0.04	-0.07	0.15	0.2	-0.02	0	0.02	0.05	-0.03	0.13	0.02	0	0.03
2	-0.02	-0.25	-0.01	-0.01	-0.01	-0.01	0.35	-0.03	-0.01	-0.01	0.31	0.1	-0.01	0.02	0.11
3	-0.28	0	-0.12	0.02	0.06	0.12	-0.03	-0.05	0.15	0.14	-0.04	0.06	0.16	0.05	0.06
4	-0.03	0.01	-0.19	-0.12	0.09	0.14	-0.07	-0.13	0.1	0.11	-0.04	0.08	0.11	0.13	0.07
5	-0.15	0.02	-0.11	-0.13	-0.12	-0.12	0	-0.01	0.13	0.09	0.01	-0.06	-0.01	0.01	-0.01
6	-0.16	0.01	-0.1	-0.11	-0.08	-0.12	0.01	0.01	-0.05	0.1	0.01	-0.05	-0.03	-0.01	-0.02
7	0	-0.23	0.01	0	0.01	0.01	-0.16	-0.1	-0.01	-0.01	-0.11	-0.03	-0.02	0.1	-0.02
8	0	0.02	0.01	-0.01	0.01	0	-0.05	-0.11	-0.02	-0.01	0	0	-0.02	0.11	0.02
9	0.07	0.03	-0.06	-0.1	-0.18	0	-0.03	-0.09	-0.1	-0.08	0	-0.03	0.07	0.09	0.03
10	0.11	0.02	-0.04	-0.07	-0.16	-0.22	-0.03	-0.09	-0.09	-0.11	-0.01	-0.01	0.08	0.09	0.03
11	0.01	-0.22	0.01	0.01	0	0.01	-0.09	0.03	0.02	0.01	-0.19	-0.06	-0.02	-0.03	0.17
12	-0.16	-0.2	-0.09	-0.09	-0.05	-0.07	-0.07	0.05	-0.05	-0.05	-0.18	-0.13	0.06	-0.05	0.17
13	0.03	0.07	-0.04	-0.06	0.06	0.02	0.02	-0.03	-0.19	-0.16	-0.05	-0.17	-0.12	0.04	0.17
14	-0.03	0.11	0.04	-0.05	0.04	0.02	-0.32	-0.76	-0.11	-0.09	-0.01	0.02	-0.16	-0.23	0.15
15	0.02	0.11	0.01	0.03	-0.01	0.01	0.11	0.09	0.05	0.04	-0.36	-0.09	-0.1	-0.09	-0.16

The matrix of the signed interaction
 magnitude (SIM) based on food web
topology along

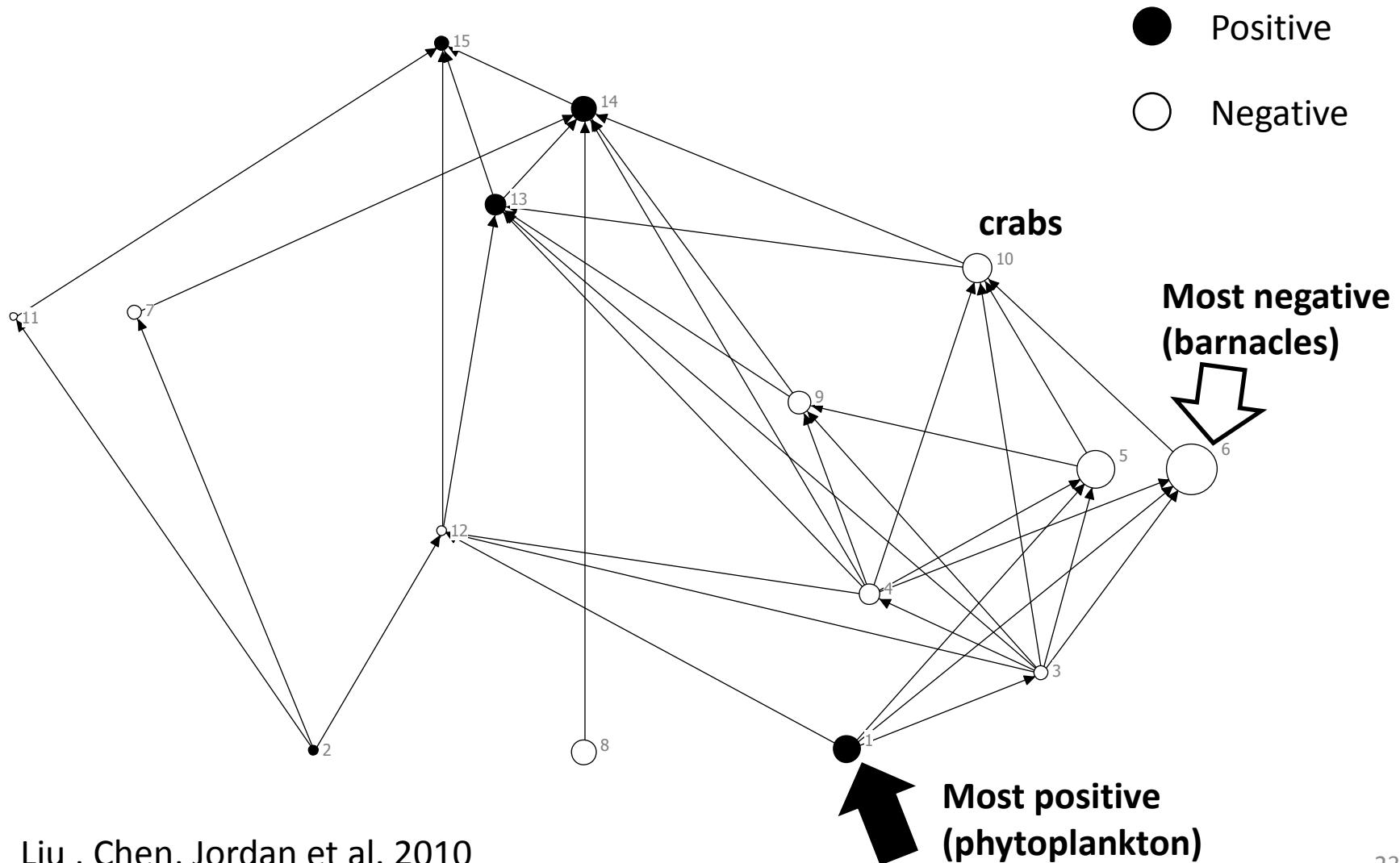
The effects from a basal species (phytoplankton, node 1)



The effects from a top species (piscivorous fish, node 15)

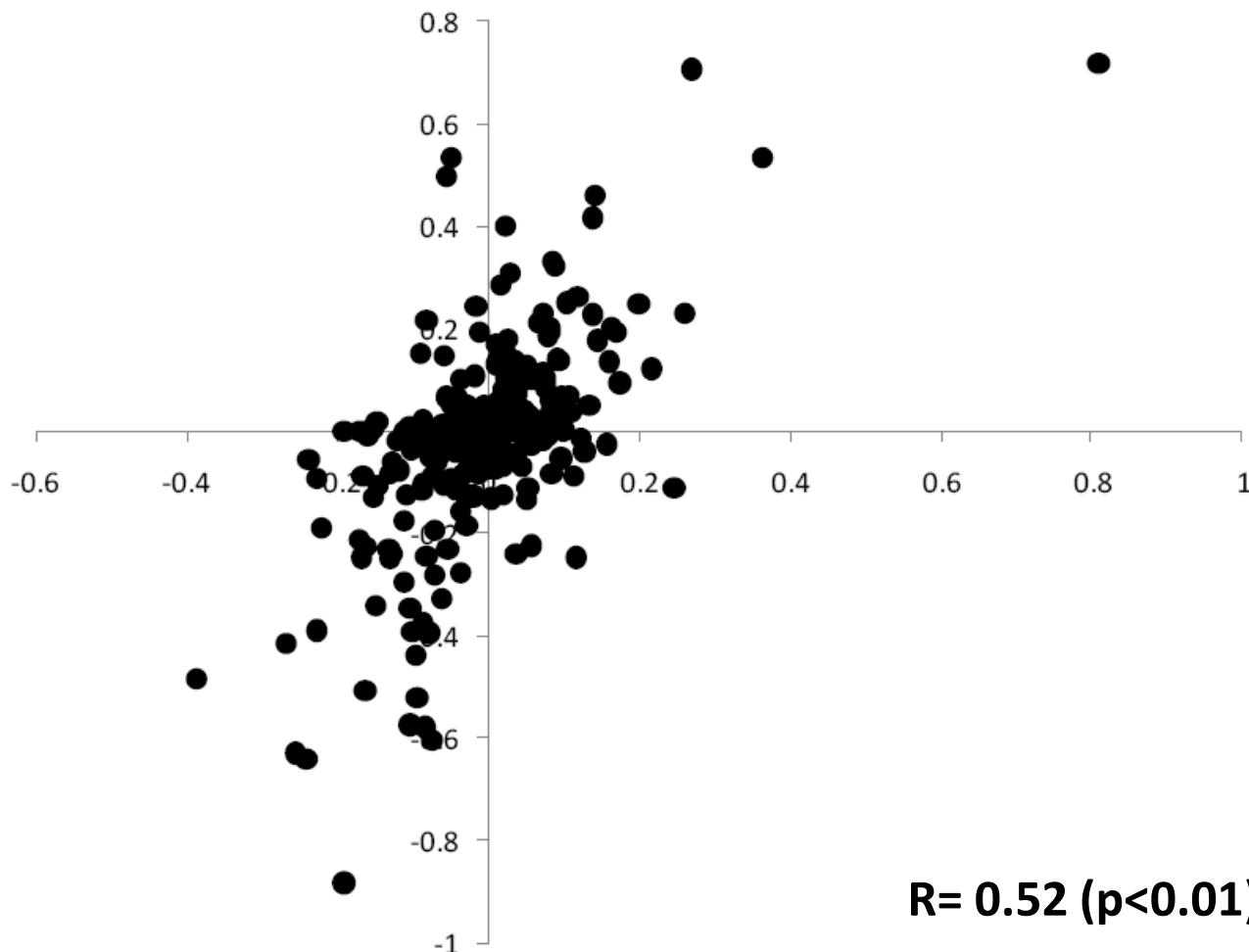


The effects from a middle species (crabs, node 10)



MTI vs. signed interaction magnitude (SIM)

Effect of node i on node j basing on the MTI approach

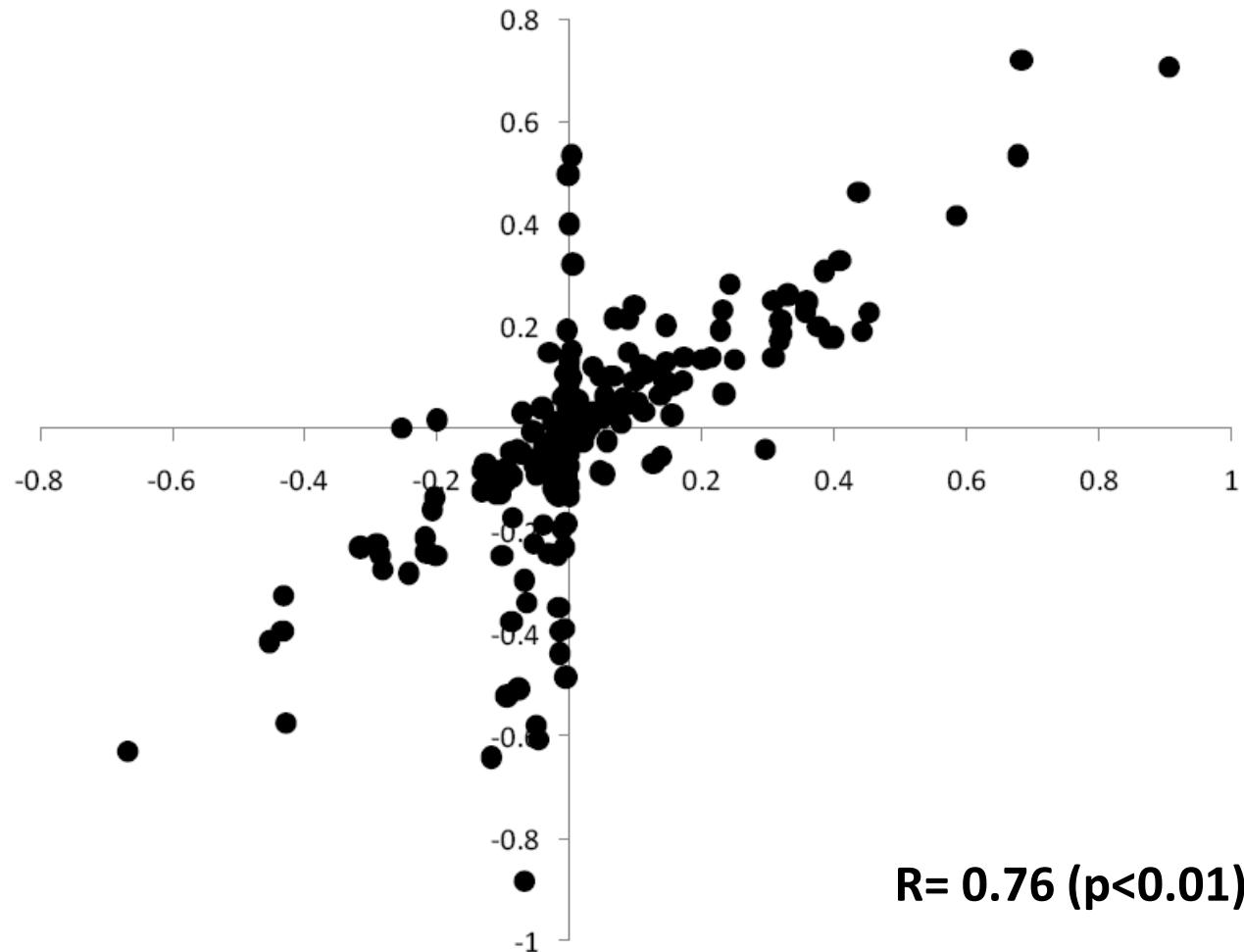


Cumulative net effect of node i on node j for un-weighted signed diagraph

Liu , Chen, Jordan et al. 2010

MTI vs. weighted SIM

Effect of node i on node j basing on the MTI approach



Cumulative net effect of node i on node j for weighted signed diagram

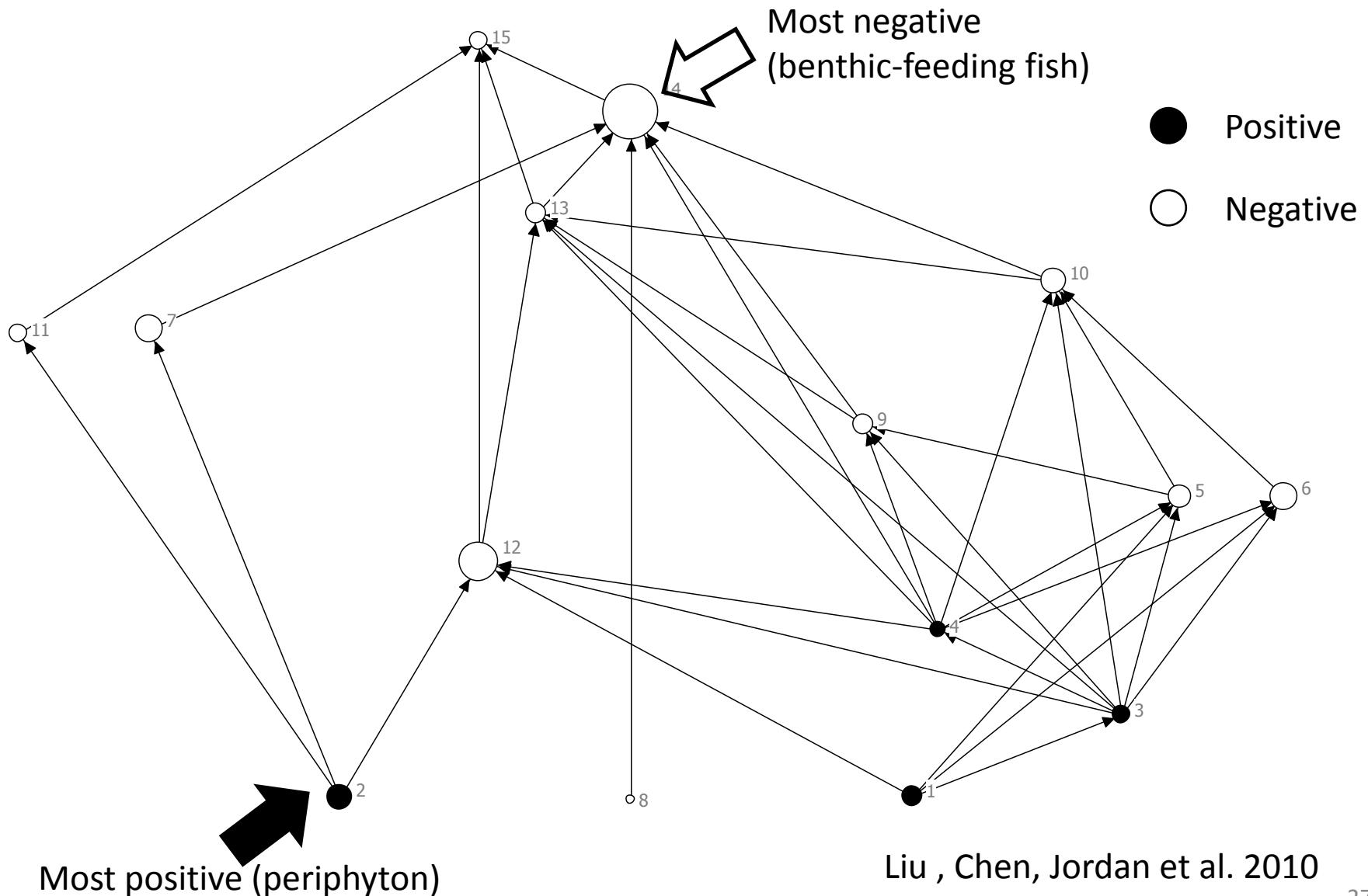
Liu , Chen, Jordan et al. 2010

The overall effect from a node to system (1-20 step)

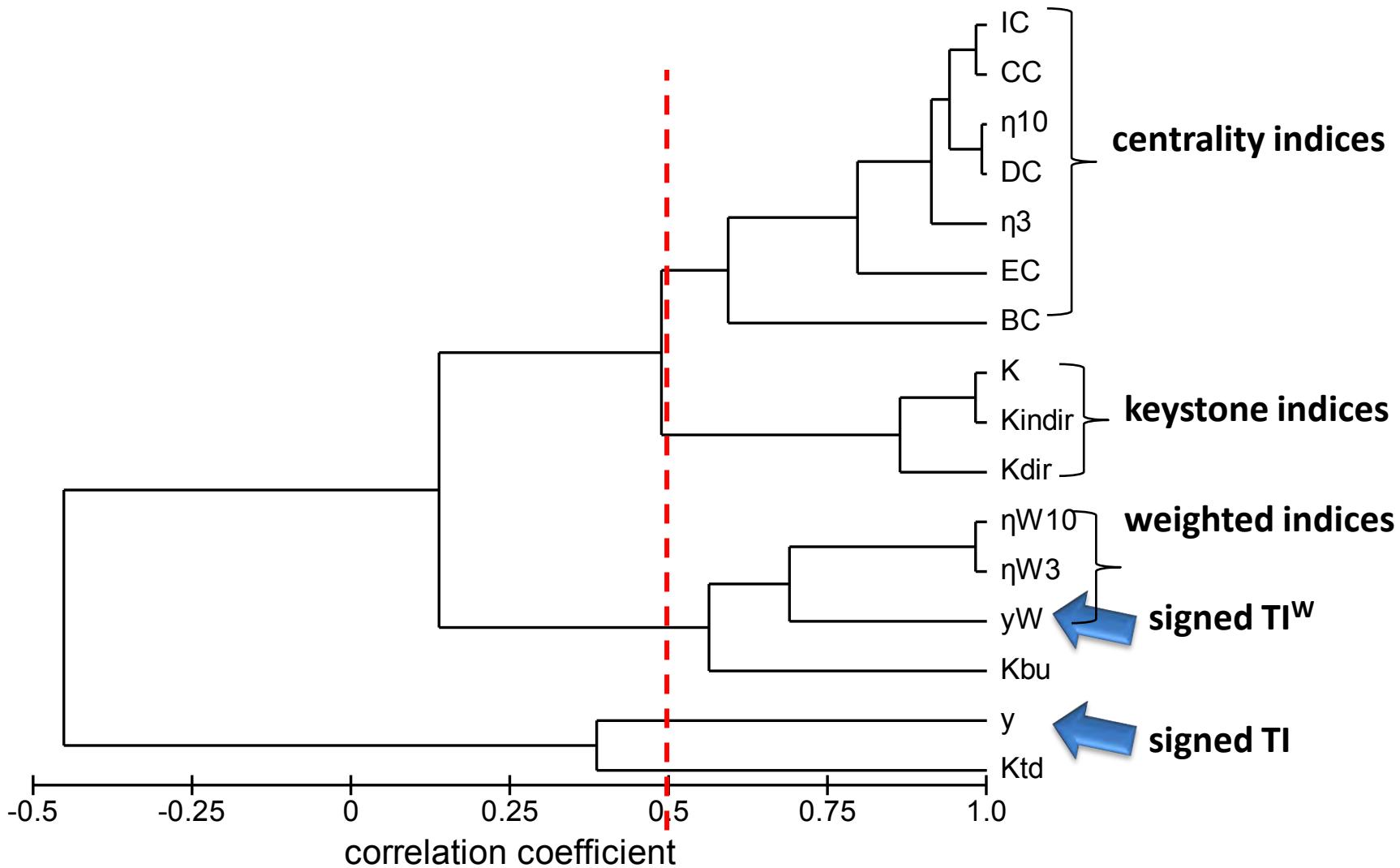
	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	Row Sum
1	-0.13	-0.03	0.04	-0.07	0.15	0.2	-0.02	0	0.02	0.05	-0.03	0.13	0.02	-0.13	-0.02	0.37
2	-0.02	-0.25	-0.01	-0.01	-0.01	-0.01	0.35	-0.03	-0.01	-0.01	0.31	0.1	-0.01	-0.02	-0.53	
3	-0.28	0	-0.12	0.02	0.06	0.12	-0.03	-0.05	0.15	0.14	-0.04	0.06	0.16	-0.02	-0.29	
4	-0.03	0.01	-0.19	-0.12	0.09	0.14	-0.07	-0.13	0.1	0.11	-0.04	0.08	0.11	-0.02	-0.27	
5	-0.15	0.02	-0.11	-0.13	-0.12	-0.12	0	-0.01	0.13	0.09	0.01	-0.06	-0.01	-0.02	-0.47	
6	-0.16	0.01	-0.1	-0.11	-0.08	-0.12	0.01	0.01	-0.05	0.1	0.01	-0.05	-0.03	-0.02	-0.58	
7	0	-0.23	0.01	0	0.01	0.01	-0.16	-0.1	-0.01	-0.01	-0.11	-0.03	-0.02	-0.02	-0.56	
8	0	0.02	0.01	-0.01	0.01	0	-0.05	-0.11	-0.02	-0.01	0	0	-0.02	-0.02	-0.06	
9	0.07	0.03	-0.06	-0.1	-0.18	0	-0.03	-0.09	-0.1	-0.08	0	-0.03	0.07	-0.02	-0.39	
10	0.11	0.02	-0.04	-0.07	-0.16	-0.22	-0.03	-0.09	-0.09	-0.11	-0.01	-0.01	0.08	-0.02	-0.49	
11	0.01	-0.22	0.01	0.01	0	0.01	-0.09	0.03	0.02	0.01	-0.19	-0.06	-0.02	-0.02	-0.34	
12	-0.16	-0.2	-0.09	-0.09	-0.05	-0.07	-0.07	0.05	-0.05	-0.05	-0.18	-0.13	0.06	-0.02	-0.92	
13	0.03	0.07	-0.04	-0.06	0.06	0.02	0.02	-0.03	-0.19	-0.16	-0.05	-0.17	-0.12	-0.02	-0.41	
14	-0.03	0.11	0.04	-0.05	0.04	0.02	-0.32	-0.76	-0.11	-0.09	-0.01	0.02	-0.16	-0.02	-1.4	
15	0.02	0.11	0.01	0.03	-0.01	0.01	0.11	0.09	0.05	0.04	-0.36	-0.09	-0.1	-0.02	-0.35	

A new food web topological index
(signed TI)

Effects exerted by species (signed TI) in food web



Signed TI vs. other food web network indices



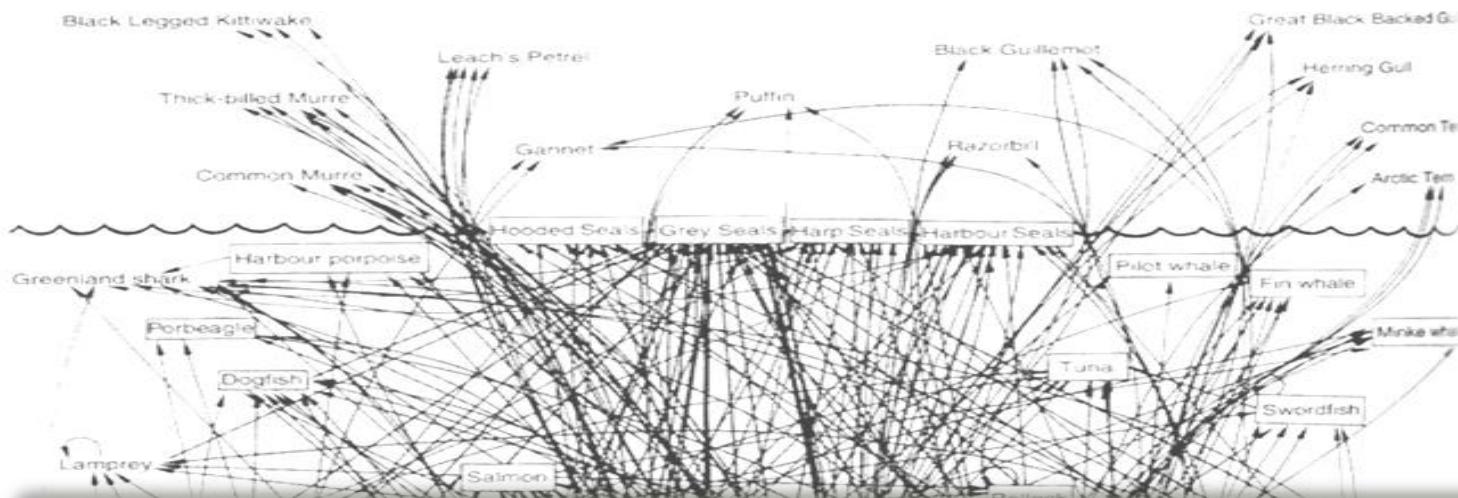
Conclusion Remarks

- The signed diagraph method provides a simple, intuitive way to reckon the species interaction structure based on the structural food web along.
- The matrix of signed interaction magnitude (SIM) seem comparable with the mixed trophic index (MTI), which is based on the mass-balanced model.
- While the dynamics nature of complex ecosystem can not be fully represented by simple static food web structure,
- our approach do provide a handy alternative means to assess species interaction and potential management consequences in species-rich, reticulate food webs.

ACKNOWLEDGEMENTS

- My coauthors: Dr. Wei-Chung Liu, Chester Wei-Jen Liu, and Wen-Shen Lin from Institute of Statistical Science, Academia Sinica. Dr. Ferenc Jordan, The Microsoft Research, University of Trento, Centre for Computational and Systems Biology, Italy.
- My supervisors: Dr. Kwang-Tsao Shao, Biodiversity Research Center, Academia Sinica. Professor Shou-Hsien Li, Dept. of Life Science, National Taiwan Normal University.
- Food web data used here is published and provided by Dr. Hsing-Juh Lin, National Chung-Hsin University.
- I am also grateful for the travel fund provided by Academia Sinica for this meeting and the food web research grant to KHS from National Science Council, Taiwan.
- A pack of programs to generate the species interaction matrix based on food web topologies presented here can be requested via emails made to me (chen7@sinica.edu.tw) or Dr. W.-C. Liu (wliu56@gate.sinica.edu.tw).
- The research presented here is part of my doctoral research conducted together with W.-C. Liu, W.-S. Lin and F. Jordan. Also, part of this research has published recently and can be cited as “Liu, W.-C., H.-W. Chen, F. Jordan, W.-H. Lin, and C. W.-J. Liu. 2010. Quantifying the interaction structure and the topological importance of species in food webs: A signed digraph approach. Journal of Theoretical Biology 267:355-362. ”





*Thank you for your attention,
Questions?*

