

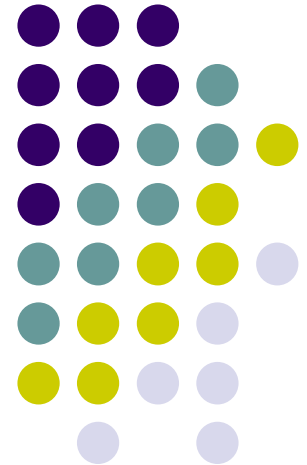


The Hypernetwork Model of Complex Systems

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Beijing Institute of Mathematical Sciences and
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Yau Mathematical Sciences Center, Tsinghua
University



Nobel Prizes 2021



Physics: Awarded “for groundbreaking contributions to our understanding of complex systems.”

Economics: Awarded “for methodological contributions to the analysis of causal relationships.”

- A complex system is composed of many components that interact with each other in a nonlinear manner.
- Interactions may be directional (casual), signed, and weighted.
- How to infer causal relationships of complex systems: Physics marries economics.

Journal of Physics: Complexity

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EDITORIAL



CrossMark

Complex systems in the spotlight: next steps after the 2021 Nobel Prize in Physics

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COMPLEX SYSTEMS | BLOG

Celebrating the complexity Nobel prize with perspectives on the future of the field

20 Jan 2023 Hamish Johnston

Inferring causal relationships of complex systems has become one of the hottest and most promising topics of research in biology, medicine, engineering, economics, and physics.

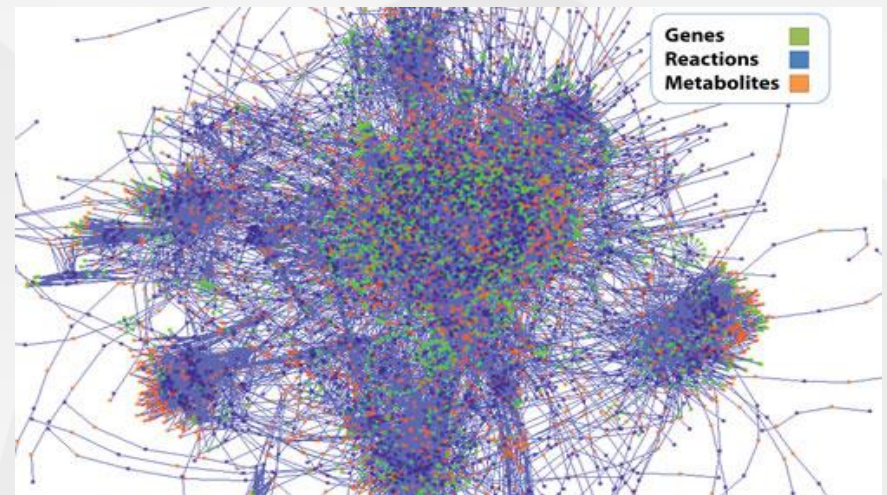
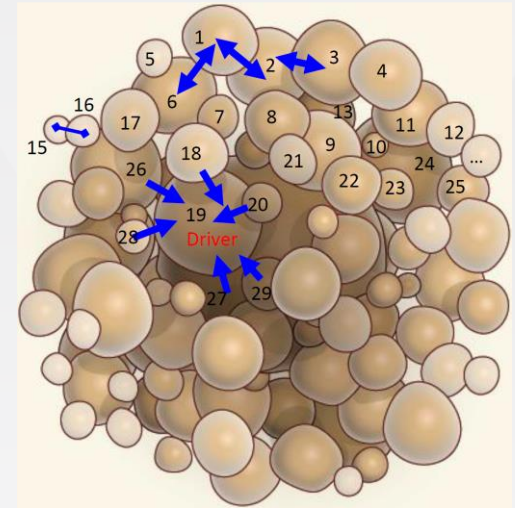


Study Complex Systems: Challenges and Opportunities

- Complex systems are “complex” in terms of the number of components and interactions
- Automatic high-throughput measurement techniques make it possible to monitor any systems.
- How can we extract causal relationships underlying complex systems using these big data sets?

Tumor →

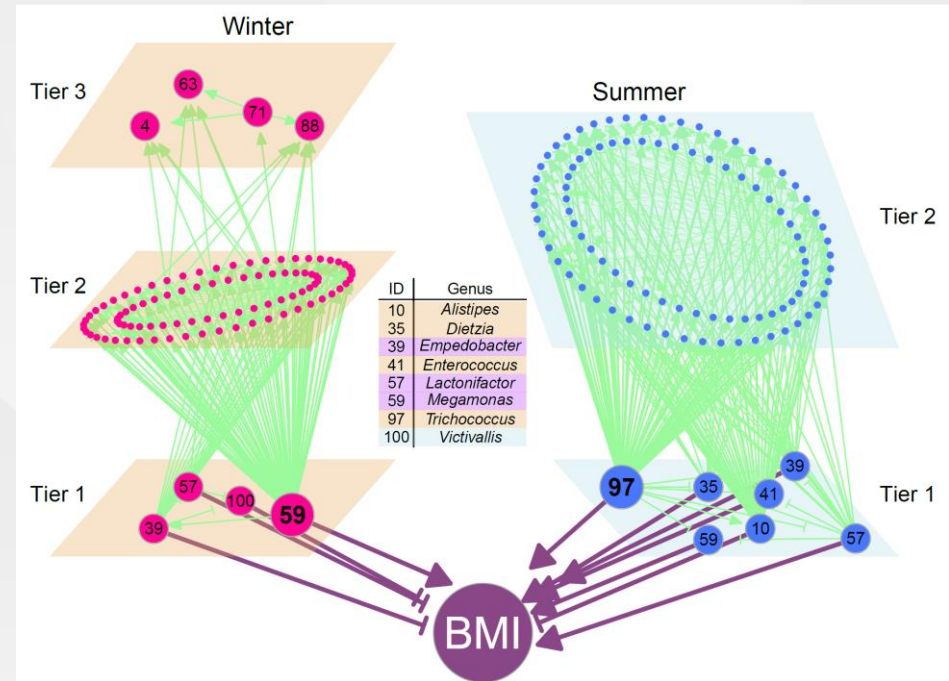
Cell



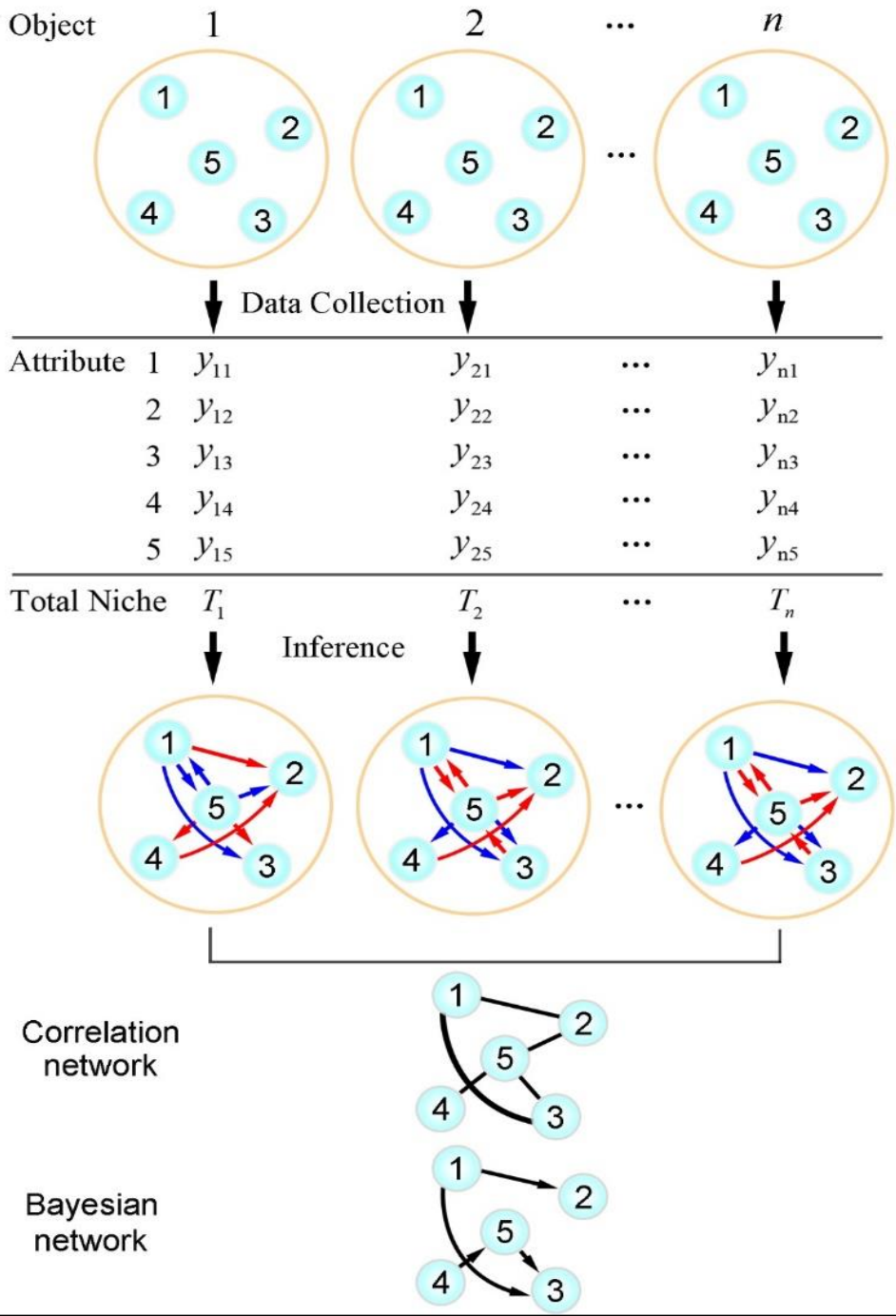
Network Modeling of Complex Systems

- All components affect system behavior through direct and/or indirect pathways.
- Network models of complex systems capture direct and indirect effects.
- Networks chart a roadmap of how each element flows its signal of influence within the whole system.

The Gut Microbiota



[Reanalysis of Davenport et al.'s \(PLoS ONE, 2015\) data](#)



- Consider n objects (e.g., cells, individuals)
- Measure p (i.e., 5) different attributes (elements) of each object
- Form an $(n \times p)$ data matrix

Current approaches can only infer an overall, less informative network.

We want to infer **sophisticated networks** from this data:

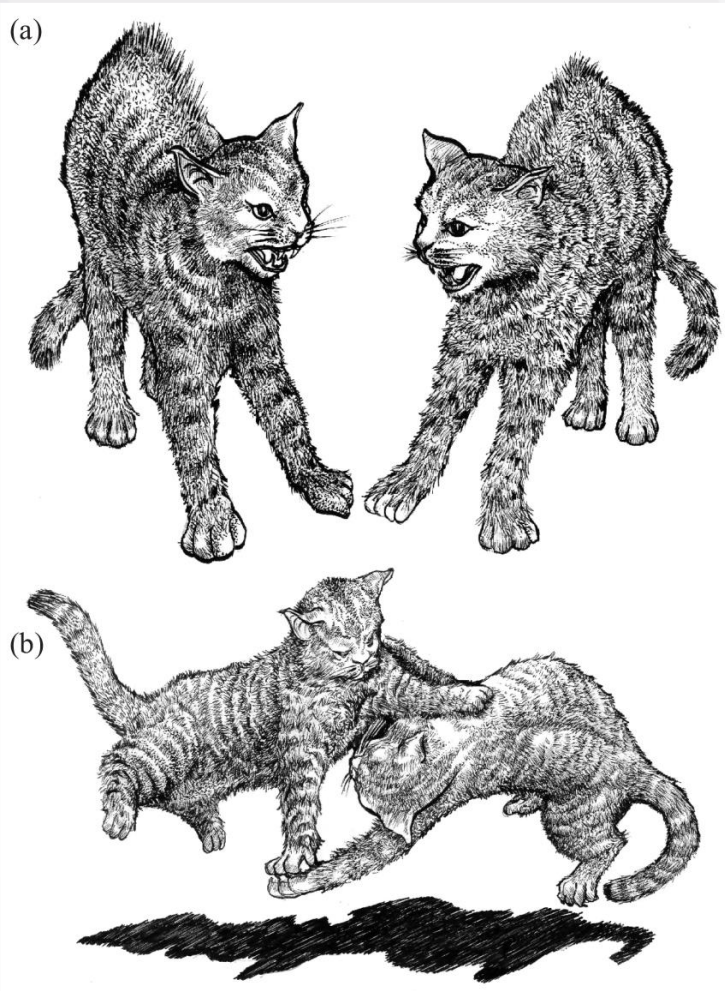
- Networks are object-specific (individualized networks)
- Networks are context-specific (e.g., diseased vs. control)
- **Bidirectional, signed, and weighted interactions** (fully informative)

How to infer such sophisticated networks?

We introduce evolutionary game theory
by viewing inter-element interactions as a game

Game Theory

The Nash equilibrium and a tit-for-tat strategy



For an evolving system, this game will occur repeatedly, expressed as

Left Cat CCCCCCCCCDCCCCCCCCCD
Right Cat CCCCCCCCCDCCCCCCCCCR

Cooperative strategy (C)
Dangerous strategy (D)
Retreats (R)



The Nash equilibrium (Nash 1950, PNAS)
↓
Evolutionarily stable strategy (ESS) (Smith and Price 1973, Nature)

Quantitative decision theory (Wu et al. 2021)

- (1) The bigger cat chooses to cooperate with the smaller cat, when their strength ratio is beyond 0.61 (golden dissection ratio)
- (2) The smaller cat would cheat the bigger cat, when their strength ratio is 0.38 to 0.61 (Fibonacci Retracement)

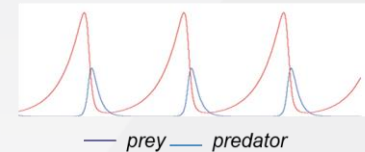
Lotka-Volterra (LV) predator-prey representation of ESS

Payoff of one cat (player) is determined by **its own strategy** and **the strategy of the other cat**



Alfred J. Lotka
(1880-1949)

Vito Volterra
(1860-1940)



$$dP_1/dt = Q_1(P_1) + Q_{1\leftarrow 2}(P_2)$$

$$dP_2/dt = Q_2(P_2) + Q_{2\leftarrow 1}(P_1)$$

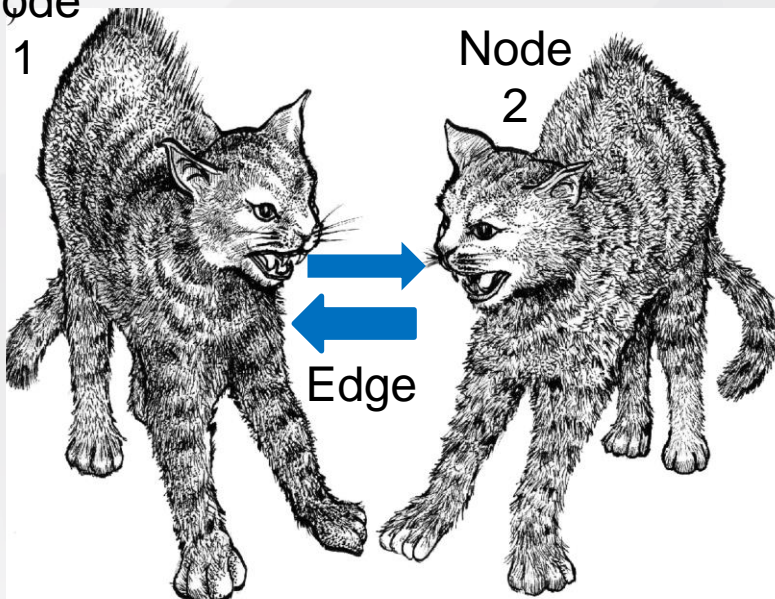
Independent Dependent

$Q_{1\leftarrow 2}(P_2)$

$Q_{2\leftarrow 1}(P_1)$

Node
1

Node
2



Mutualism	+	+
Antagonism	-	-
Aggression	+	-
Altruism	-	+

Bidirectional, signed, and weighted interactions, outperforming traditional models

Johnson, C.A., Smith, G.P., Yule, K. *et al.* Coevolutionary transitions from antagonism to mutualism explained by the Co-Opted Antagonist Hypothesis. *Nat Commun* **12**, 2867 (2021).


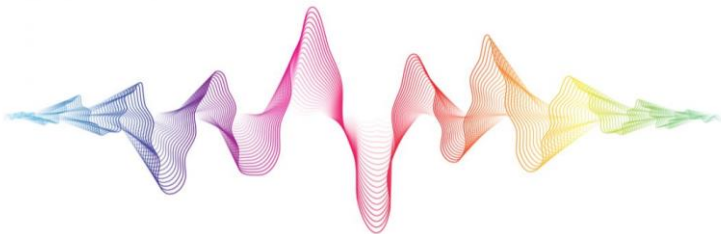
Dynamic Systems and Dynamic Models

- Differential equations:
 - Ordinary differential equations (ODE)–simplest
 - Delay differential equations (DDE)
 - Hybrid differential equations (HDE)
 - Partial differential equations (PDE)
 - Stochastic differential equations (SDE)
- Difference equations and state-space models
- Stochastic processes models: branching process etc.
- Agent-based models and cellular automata

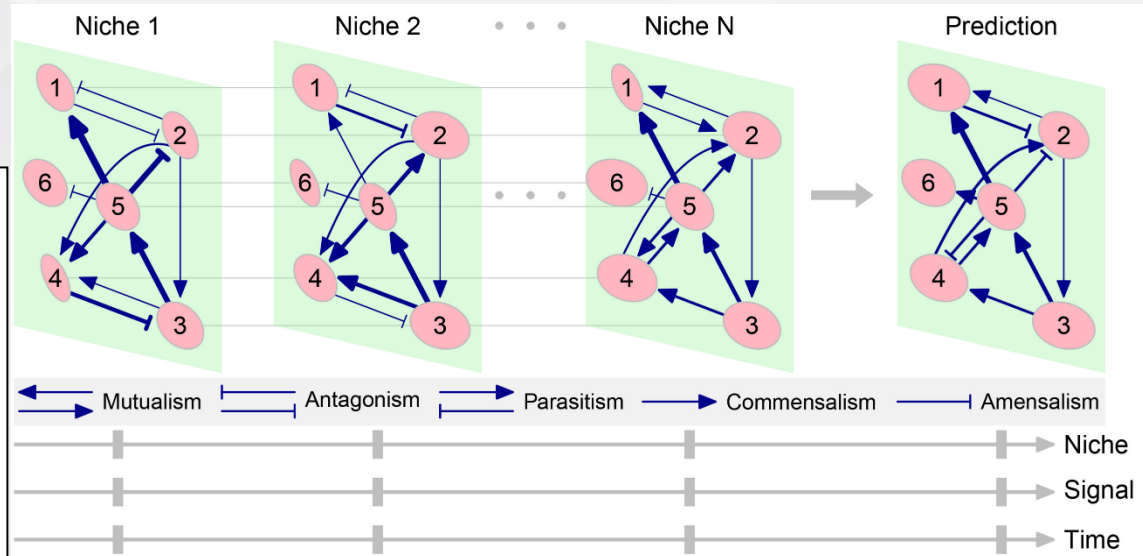
Graph Theory



Dynamic Modeling

Dynamic Measurements



Networks beyond Dyadic Interactions

- **Graph theory: pairwise interactions**
- **Hypergraph theory: high-order interactions**

Taking high-order interactions into account

- Enhance our modeling capacities
- Help to more precisely characterize complex systems

REVIEW

doi:10.1038/nature22898

Beyond pairwise mechanisms of species coexistence in complex communities

Jonathan M. Levine¹, Jordi Bascompte², Peter B. Adler³ & Stefano Allesina⁴

The tremendous diversity of species in ecological communities has motivated a century of research into the mechanisms that maintain biodiversity. However, much of this work examines the coexistence of just pairs of competitors. This approach ignores those mechanisms of coexistence that emerge only in diverse competitive networks. Despite the potential for these mechanisms to create conditions under which the loss of one competitor triggers the loss of others, we lack the knowledge needed to judge their importance for coexistence in nature. Progress requires borrowing insight from the study of multitrophic interaction networks, and coupling empirical data to models of competition.

Physics Reports 874 (2020) 1–92



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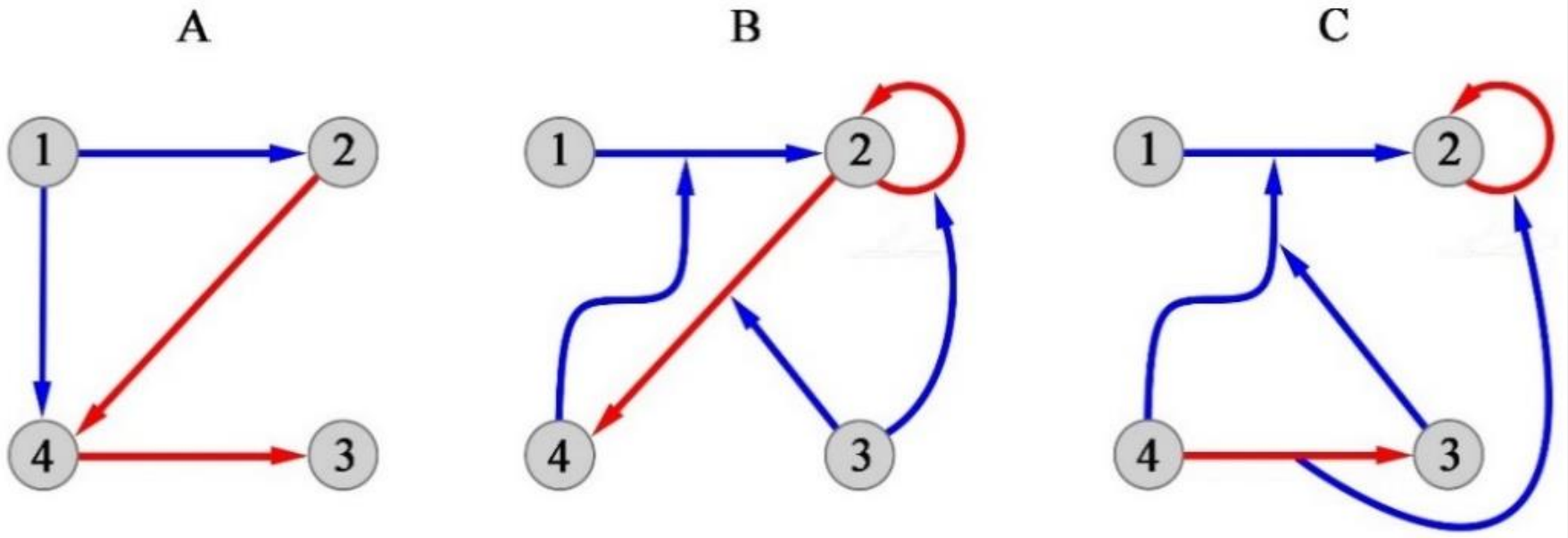
Physics Reports

journal homepage: www.elsevier.com/locate/physrep

Networks beyond pairwise interactions: Structure and dynamics

Federico Battiston^{a,*}, Giulia Cencetti^b, Iacopo Iacopini^{c,d}, Vito Latora^{c,e,f,g}, Maxime Lucas^{h,i,j}, Alice Patania^k, Jean-Gabriel Young^l, Giovanni Petri^{m,n}

Paradigm Shift: from Networks to Hypernetworks



- A node affects the interaction between two other nodes
- Interactions between two nodes is affected by a third node

A Hypernetwork Theory: Modeling High-order Interaction Networks

Consider an m -dimensional biosystem:

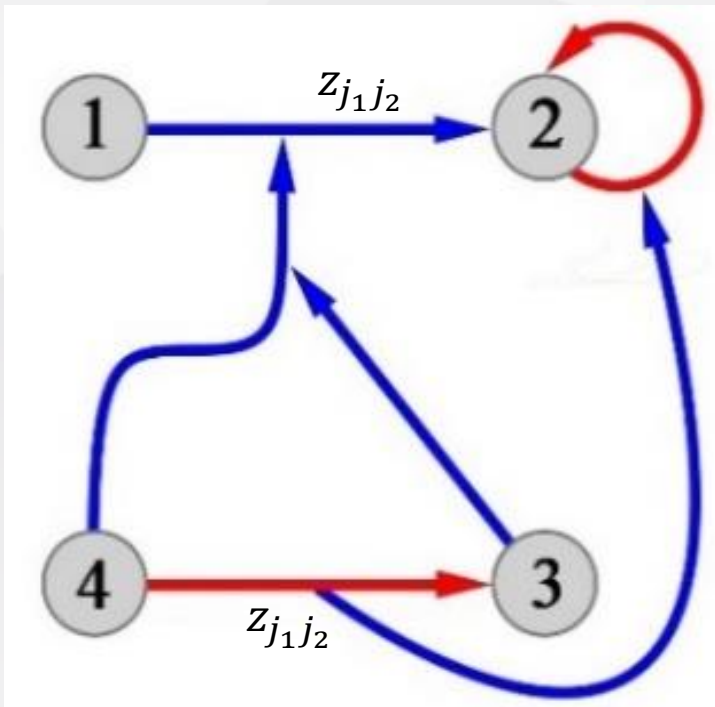
$$\begin{aligned}
 \frac{dg_j(t)}{dt} = & Q_j(g_j(t): \Theta_j) && \text{Independent} \\
 & + \sum_{j'=1}^m Q_{j \leftarrow j'}(g_{j'}(t): \Theta_{jj'}) && \text{First-order} \\
 & + \sum_{j_1=1}^m \sum_{j_2=1}^m Q_{j \leftarrow j_1 j_2}(z_{j_1 j_2}(t): \Theta_{jj_1 j_2}) && \text{Second-order} \\
 & + \sum_{j_1=1}^m \sum_{j_2=1}^m \sum_{j_3=1}^m Q_{j \leftarrow j_1 j_2 j_3}(z_{j_1 j_2 j_3}(t): \Theta_{jj_1 j_2 j_3}) && \text{Third-order} \\
 & + \\
 & \dots \quad j' \neq j = 1, \dots, m; \quad j_1 < j_2 < j_3 = 1, \dots, m
 \end{aligned}$$

- It may include more terms that describe four and higher-way interactions if needed.
- Different from pairwise network modeling, it treats predictors as interacting networks at different orders.

What is the interaction $z_{j_1 j_2}$?

$$\begin{aligned} \frac{dg_j(t)}{dt} = & Q_j(g_j(t): \theta_j) \\ & + \sum_{j'=1}^m Q_{j \leftarrow j'}(g_{j'}(t): \theta_{jj'}) \\ & + \sum_{j_1=1}^m \sum_{j_2=1}^m Q_{j \leftarrow j_1 j_2}(z_{j_1 j_2}(t): \theta_{jj_1 j_2}) \\ & \quad j' \neq j = 1, \dots, m; j_1 < j_2 = 1, \dots, m \end{aligned}$$

Independent
First-order
Second-order



Mutualism	1 ↔ 2
Antagonism	1 ↔ 2
Aggression	1 → 2
Altruism	1 → 2

How to define and quantify player-player interactions?

We introduce behavioral ecology theory



goose flying and fish schooling



A quantitative decision theory of animal conflict

(Wu and Jiang et al. 2021, *Heliyon* 7(7): e07621)



Mathematical descriptors of interactions



		Player Y	
		Cooperative	Competitive
Player X	Coop	Mutualism	Altruism
	Comp	Aggression	Antagonism

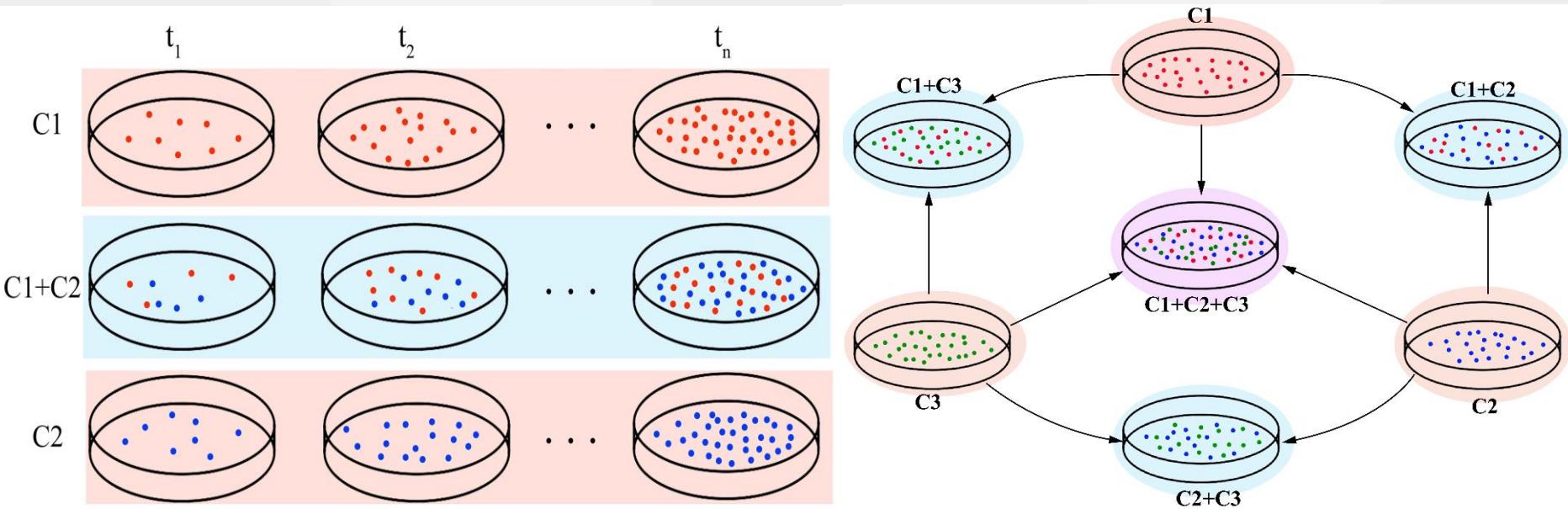
		Player Y	
		Cooperative	Competitive
Player X	Coop	$z_{mu} = xy/(x-y)$	$z_{al} = (x-y)/x$
	Comp	$z_{ag} = x/y$	$z_{an} = 1/[xy(x-y)]$

Mutualism-based hypernetworks, z_{mu}
Altruism-based hypernetworks, z_{al}
Aggression-based hypernetworks, z_{ag}
Antagonism-based hypernetworks, z_{an}

Experimental validation by Jiang et al. 2019, 2020; Wang et al. 2019, He et al. 2021

Experimental Validation of the Hypernetwork Theory

Randomly choose different lung cancer cell types and cultivate them in monoculture, co-culture, and tri-culture



By Shawn Rice, Penn State College of Medicine

Fundamental Core of the Hypernetwork Theory: Evolutionary Game Dissection

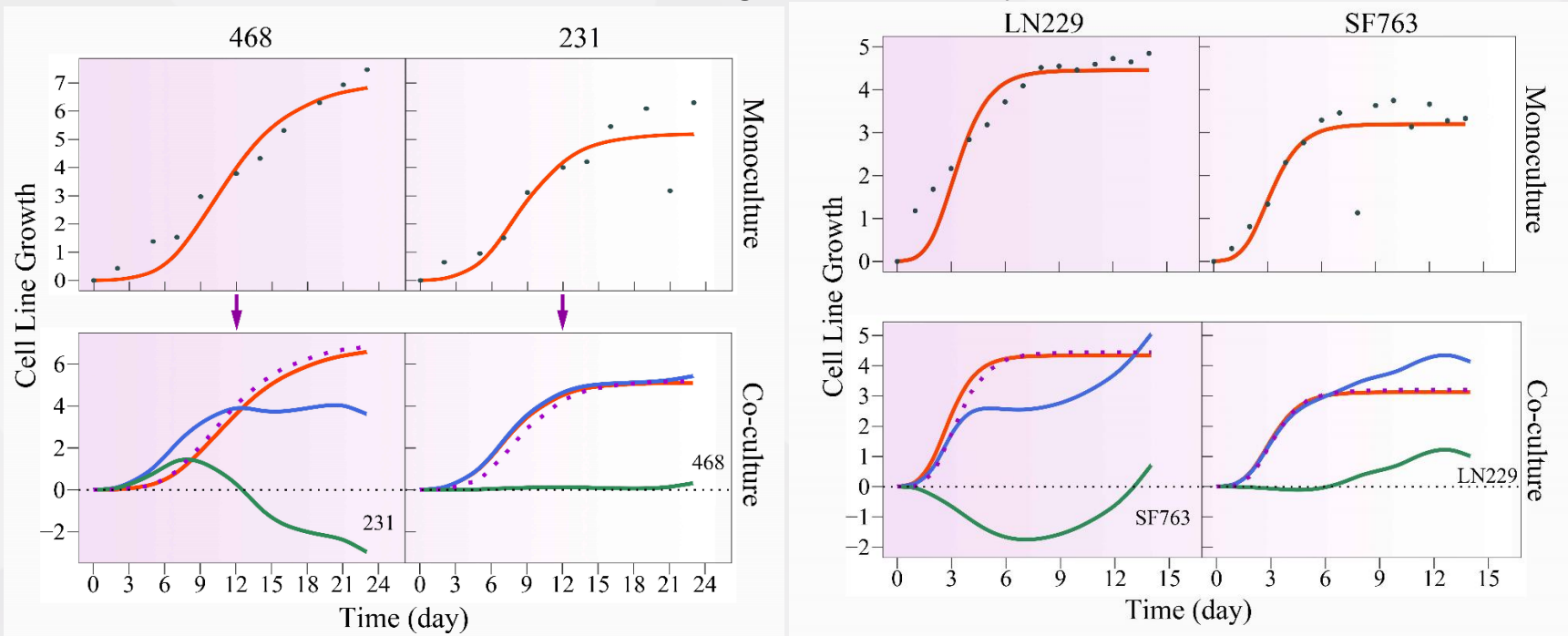
In a socialized environment, the payoff of any player is decomposed in a way like this

$$\begin{aligned} \frac{dg_j(t)}{dt} = & Q_j(g_j(t): \Theta_j) && \text{Independent} \\ & + \sum_{j'=1}^m Q_{j \leftarrow j'}(g_{j'}(t): \Theta_{jj'}) && \text{First-order} \\ & + \sum_{j_1=1}^m \sum_{j_2=1}^m Q_{j \leftarrow j_1 j_2}(z_{j_1 j_2}(t): \Theta_{jj_1 j_2}) && \text{Second-order} \\ & j' \neq j = 1, \dots, m; j_1 < j_2 = 1, \dots, m \end{aligned}$$

- The independent component occurs when this player **is assumed** to be in isolation.
- If the estimated independent component value is consistent with that value obtained from its monoculture, then this suggests that the dissection theory works.

Experimental Validation from *in vitro* Cultural Experiment

Four lung cancer cell types



Amensalism (偏害)

$$\begin{aligned} dP_1/dt &= Q_1(P_1) + Q_{1 \leftarrow 2}(P_2) \\ dP_2/dt &= Q_2(P_2) + Q_{2 \leftarrow 1}(P_1) \end{aligned}$$

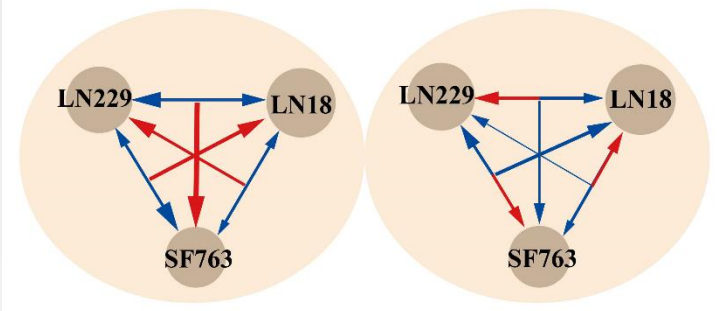
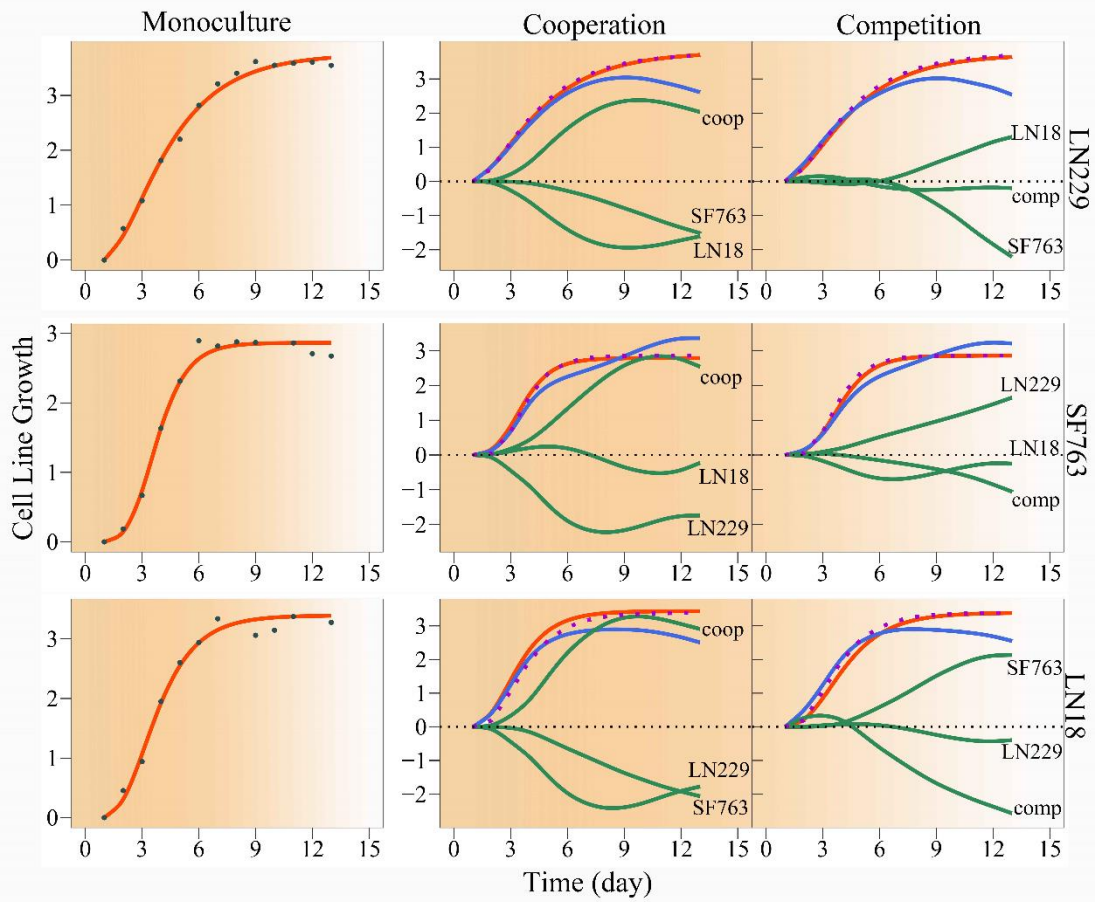
The model works!

Altruism/Aggression (利他/侵害)

Analyzed by Li Feng



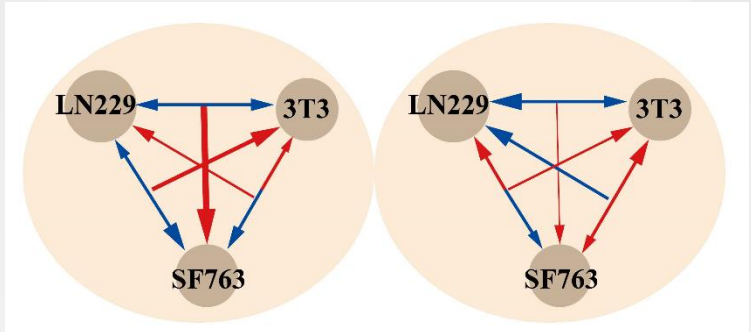
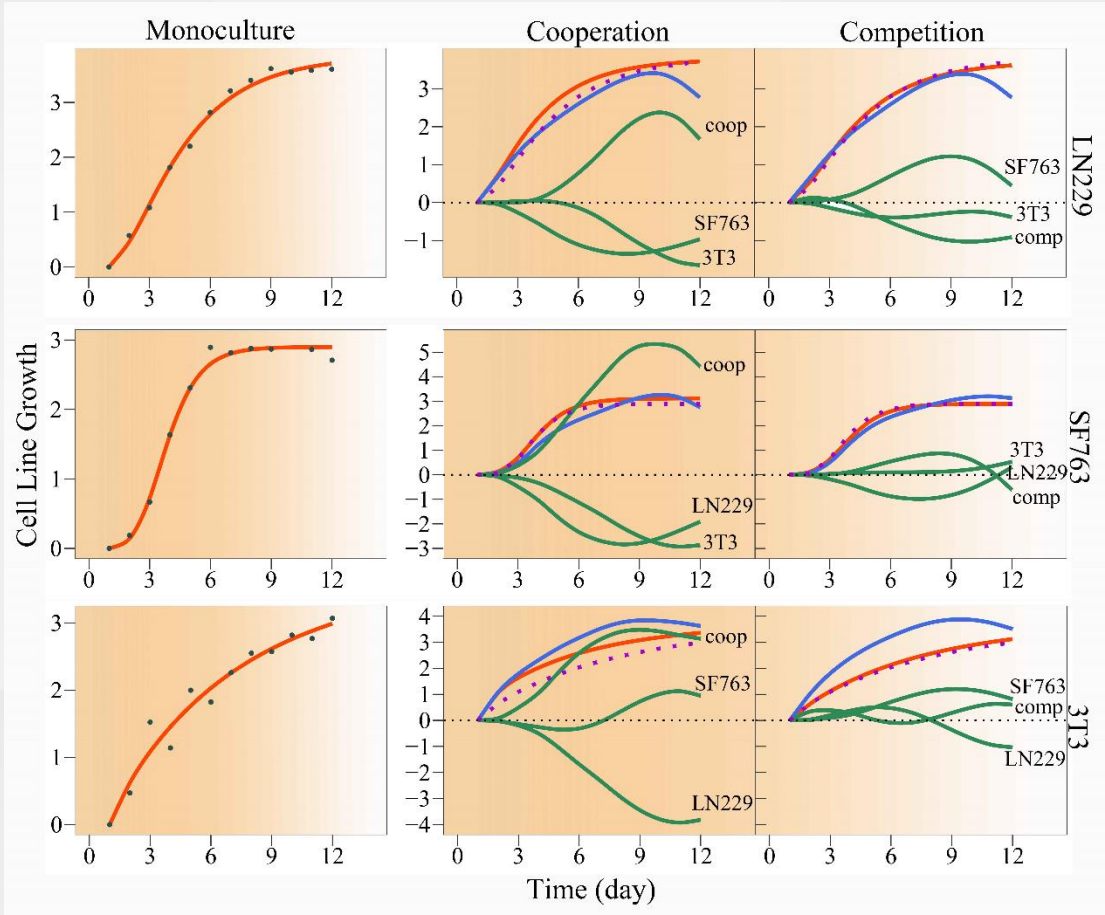
Tri-culture: Experiment 1 – LN229, SF763, LN18



LN229 = Independent
 - SF763
 - LN18
 + Cooperation

Uncouple their cooperation, making cell growth inhibited

Tri-culture: Experiment 2 – LN229, SF763, 3T3



LN229 = Independent
 - SF763
 - 3T3
 + Cooperation

Uncouple their cooperation, making cell growth inhibited

Disadvantages of Dynamic Models

Unavailability of high-density time-series measurements

- Impossible to collect
- Ethically impermissible

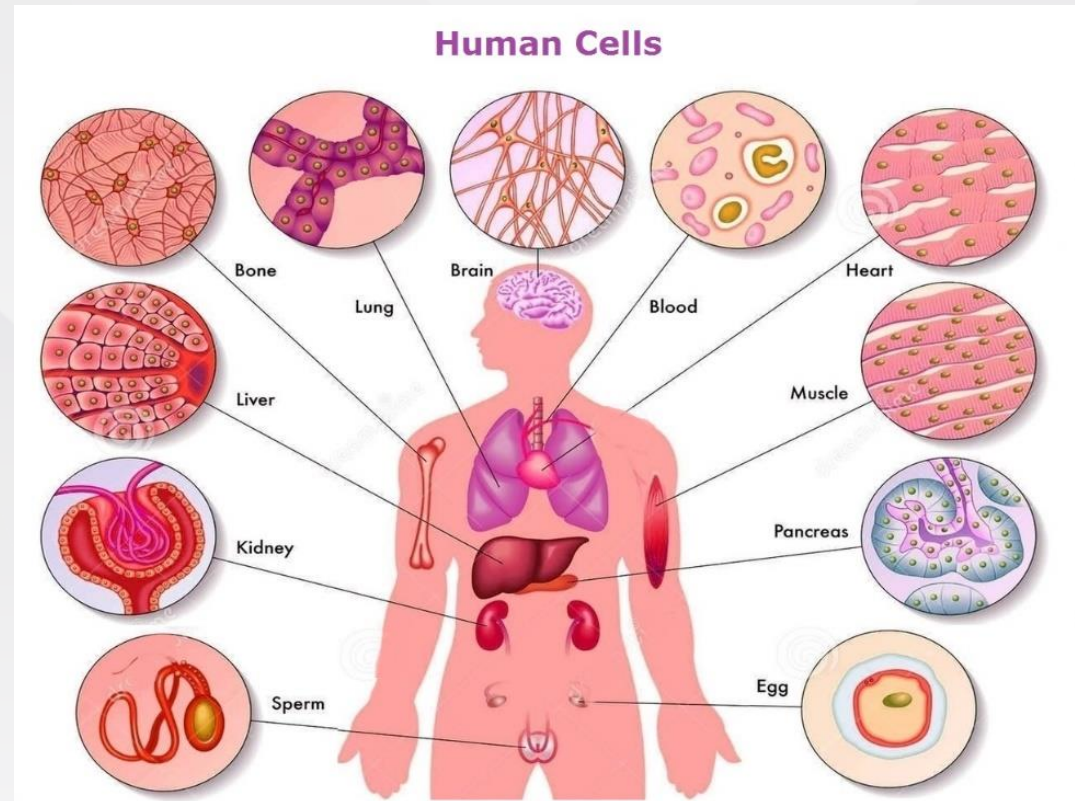
Limitations to curve fitting

- Time stable
- Resilient to perturbations

The GTEx (Genotype-Tissue Expression) Project: transcriptome measured only once for one donor

<https://commonfund.nih.gov/gtex>

Cohesive coordination of multiple tissues → Human health

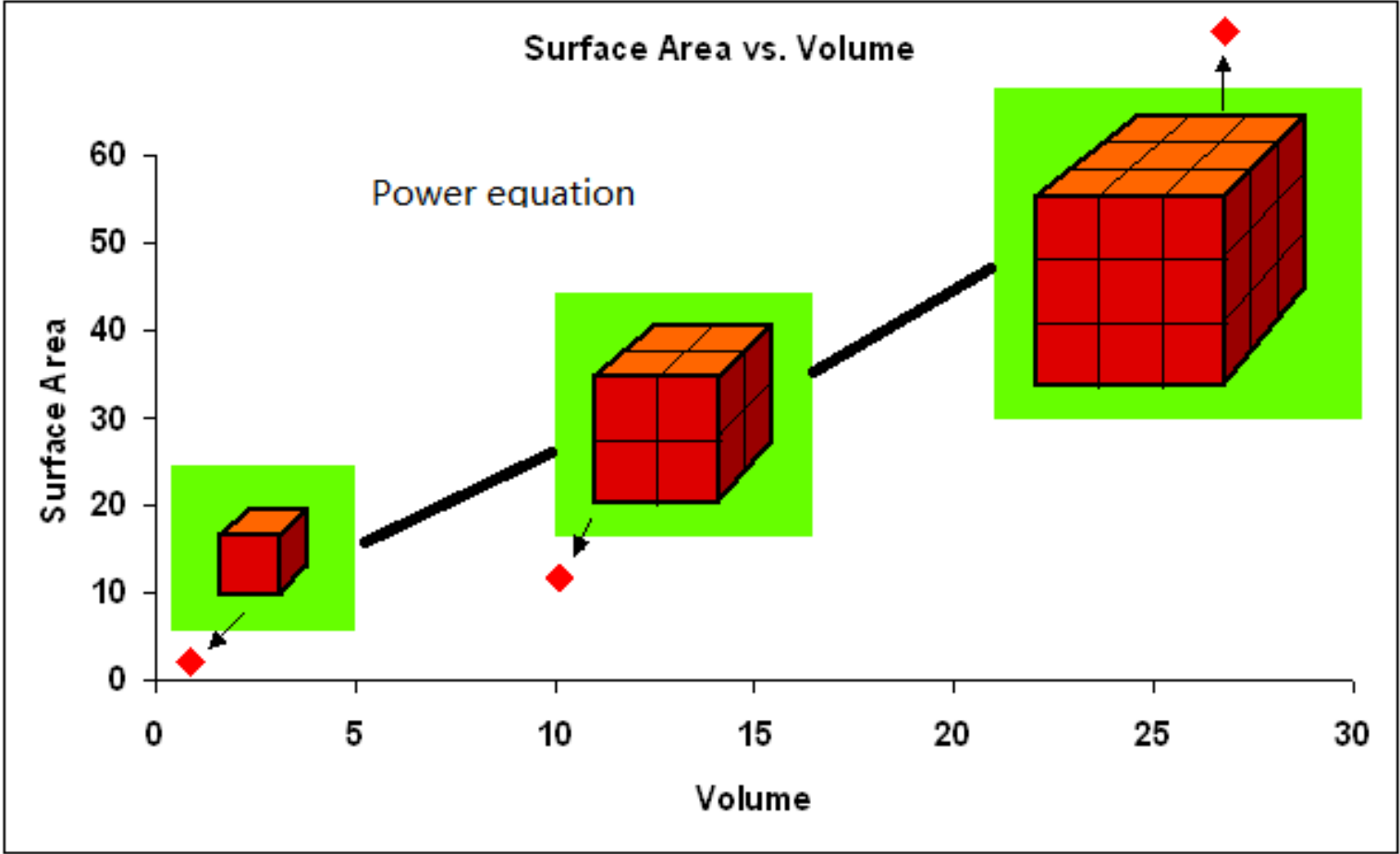


How to convert static data into their quasi-dynamic representation?

We implement allometric scaling law

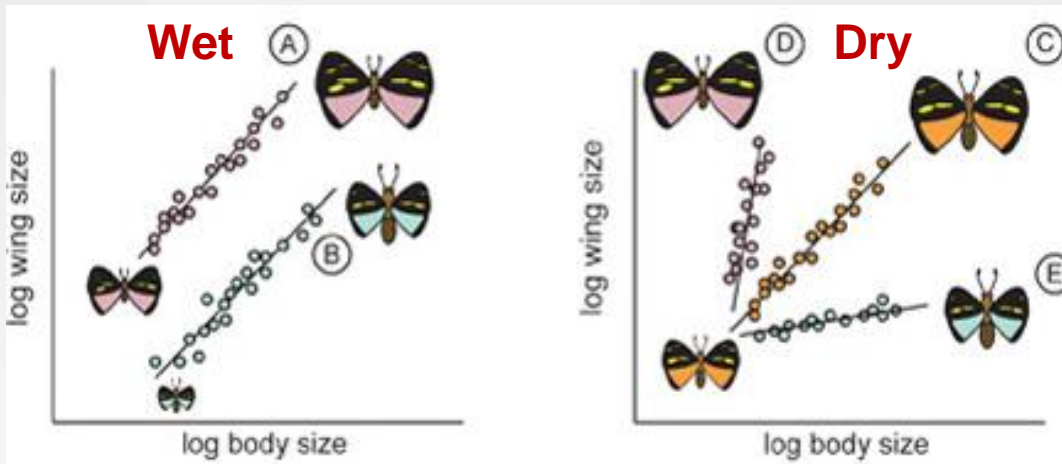
Allometric scaling law is a Physical Law:

Part-whole relationship



Allometric Scaling Law is a Biological Law

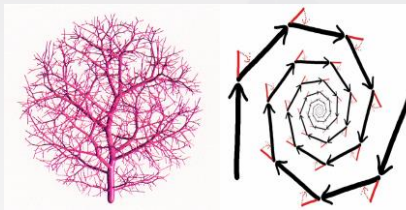
Individual elements vs. their whole system is the part-whole relationship.



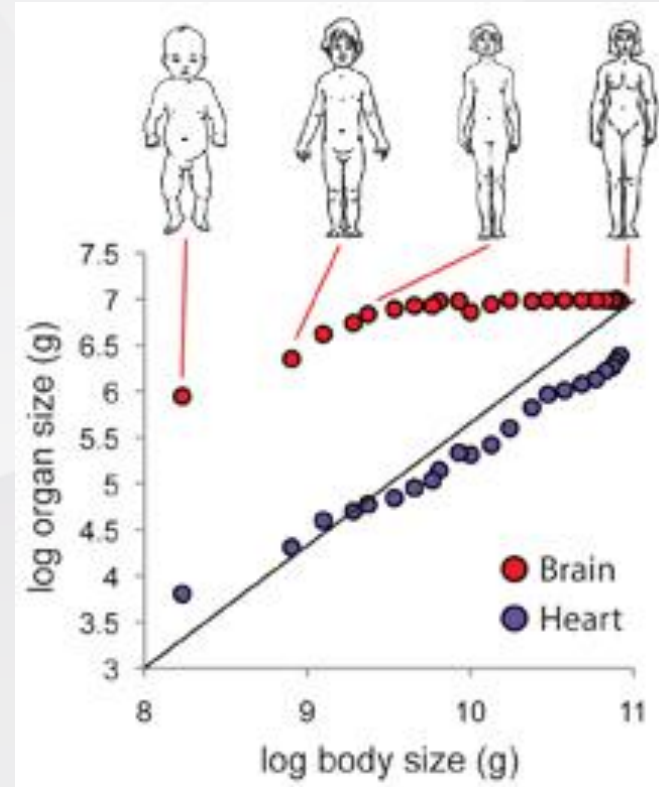
Static Allometry (across individuals) (Shingleton 2010)

Power equation: $Y = \alpha X^\beta$, where α is the intercept and β is the slope

West et al. (1997, Science) proposed a fractal model to interpret the power law.



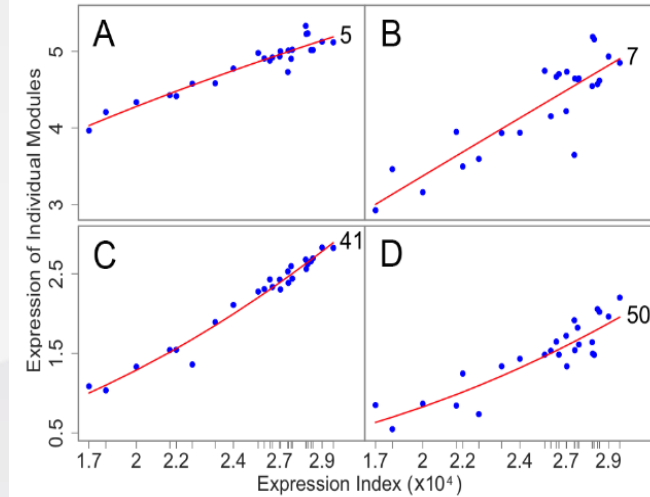
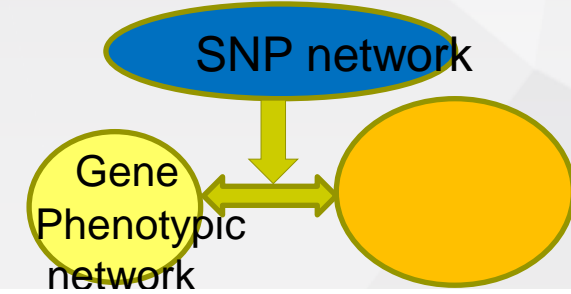
West, Brown, Enquist



Ontogenetic Allometry (across ages)

GTEX Data Structure

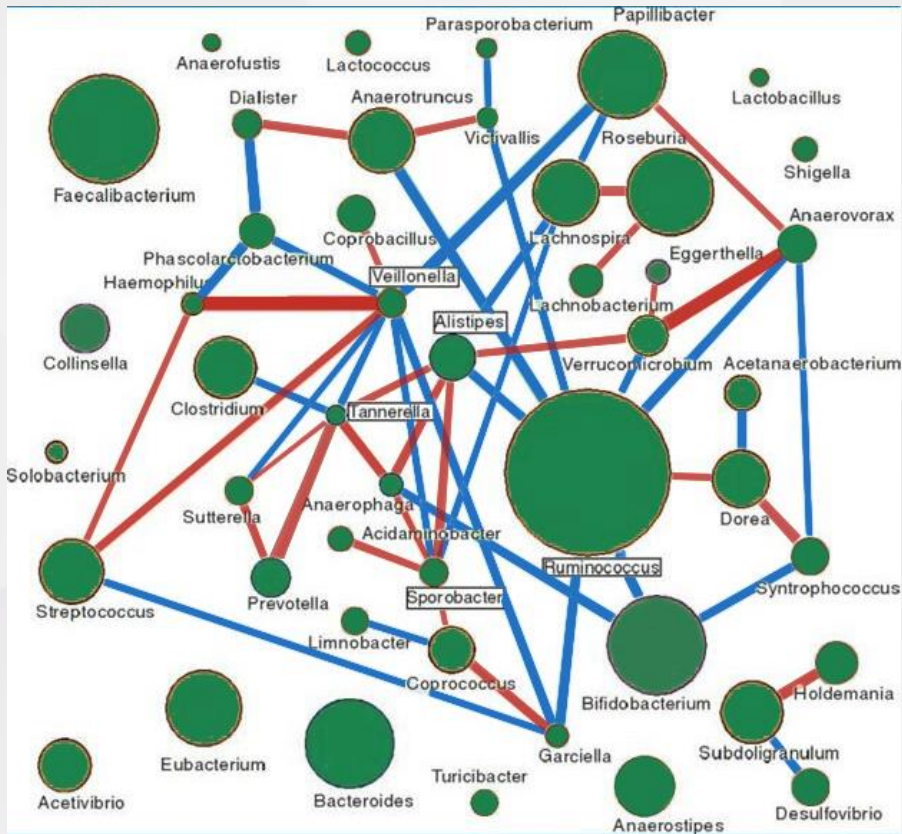
Individual	1				...	n			
Tissue	1	2	...	R_1		1	2	...	R_n
Gene expression									
1	$y_{11}(1)$	$y_{11}(2)$...	$y_{11}(R_1)$...	$y_{1n}(1)$	$y_{1n}(2)$...	$y_{1n}(R_n)$
2	$y_{21}(1)$	$y_{21}(2)$...	$y_{21}(R_1)$...	$y_{2n}(1)$	$y_{2n}(2)$...	$y_{2n}(R_n)$
...									
m	$y_{m1}(1)$	$y_{m1}(2)$...	$y_{m1}(R_1)$...	$y_{mn}(1)$	$y_{mn}(2)$...	$y_{mn}(R_n)$
	T_{11}	T_{12}	...	T_{1R_1}	...	T_{n1}	T_{n2}	...	T_{nR_n}
Histological and clinical									
1	$z_{11}(1)$	$z_{11}(2)$...	$z_{11}(R_1)$...	$z_{1n}(1)$	$z_{1n}(2)$...	$z_{1n}(R_n)$
2	$z_{21}(1)$	$z_{21}(2)$...	$z_{21}(R_1)$...	$z_{2n}(1)$	$z_{2n}(2)$...	$z_{2n}(R_n)$
...									
p	$z_{m1}(1)$	$z_{m1}(2)$...	$z_{m1}(R_1)$...	$z_{mn}(1)$	$z_{mn}(2)$...	$z_{mn}(R_n)$
SNP									
1		AA			...				aa
2		AA			...				Aa
...									
q		aa			...				AA



- Expression index (EI) is defined as the sum of expression of all genes
- Scaling individual genes vs. EI

Data Modeling of Hypernetworks: A Preliminary Result

The Gut Microbiota



Davenport et al.'s (2015) data

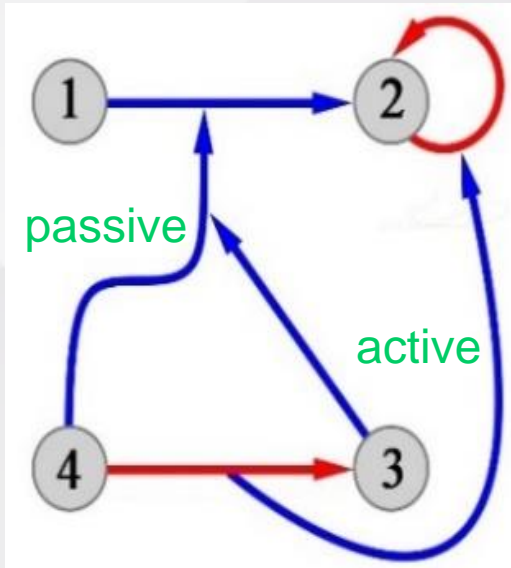
- Include 184 Amish samples from a founder, the Hutterites
- Measured at phylum, class, order, family, genus, and species levels in the winter and the coming summer.



Eight phyla: Hypergraph-based qdODEs

$$\dot{y}_j(T_i) = Q_j(y_j(T_i): \Theta_j) + \sum_{j'=1}^8 Q_{jj'}(y_{j'}(T_i): \Theta_{jj'}) + \sum_{j_1=1}^8 \sum_{j_2=1}^8 Q_{j \leftarrow j_1 j_2}(z_{j_1 j_2}(T_i): \Theta_{j \leftarrow j_1 j_2})$$

$$\dot{z}_{j_1 j_2}(T_i) = Q_{j_1 j_2}(z_{j_1 j_2}(T_i): \Theta_{j_1 j_2}) + \sum_{j=1}^8 Q_{j_1 j_2 \leftarrow j}(y_j(T_i): \Theta_{j_1 j_2 \leftarrow j})$$



High-order networks include

- (1) how a pairwise interaction **actively** affects a node (phylum)
- (2) how a dyadic interaction is **passively** affected by a node

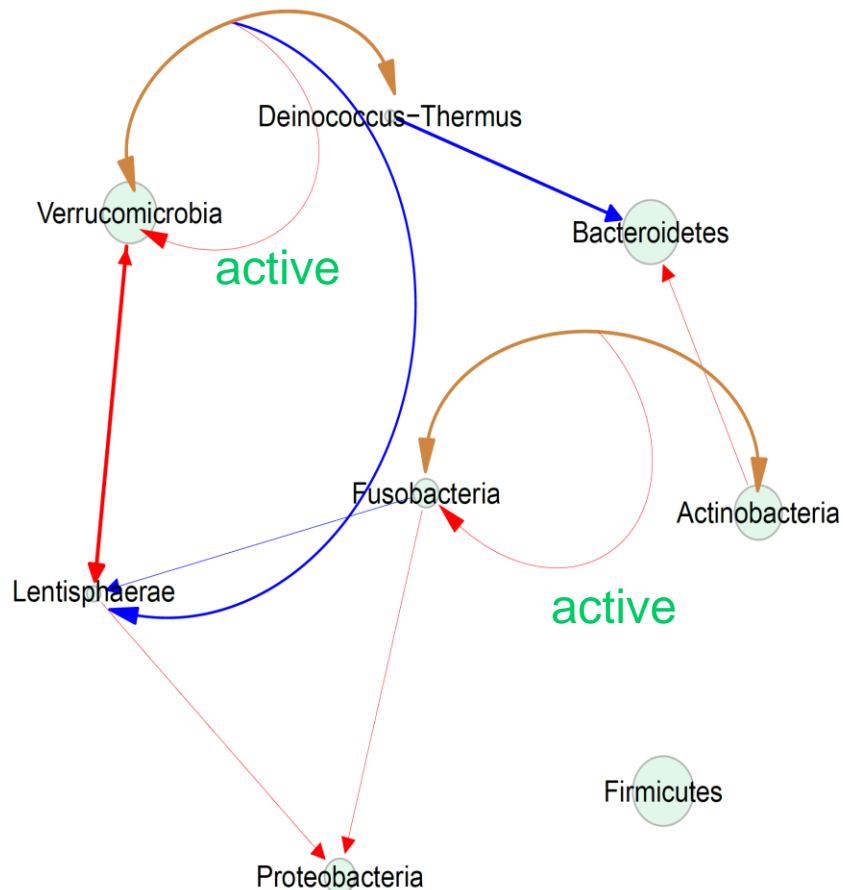
Pairwise interaction is defined by $z_{j_1 j_2}$

Active Hypernetworks (mutualism)

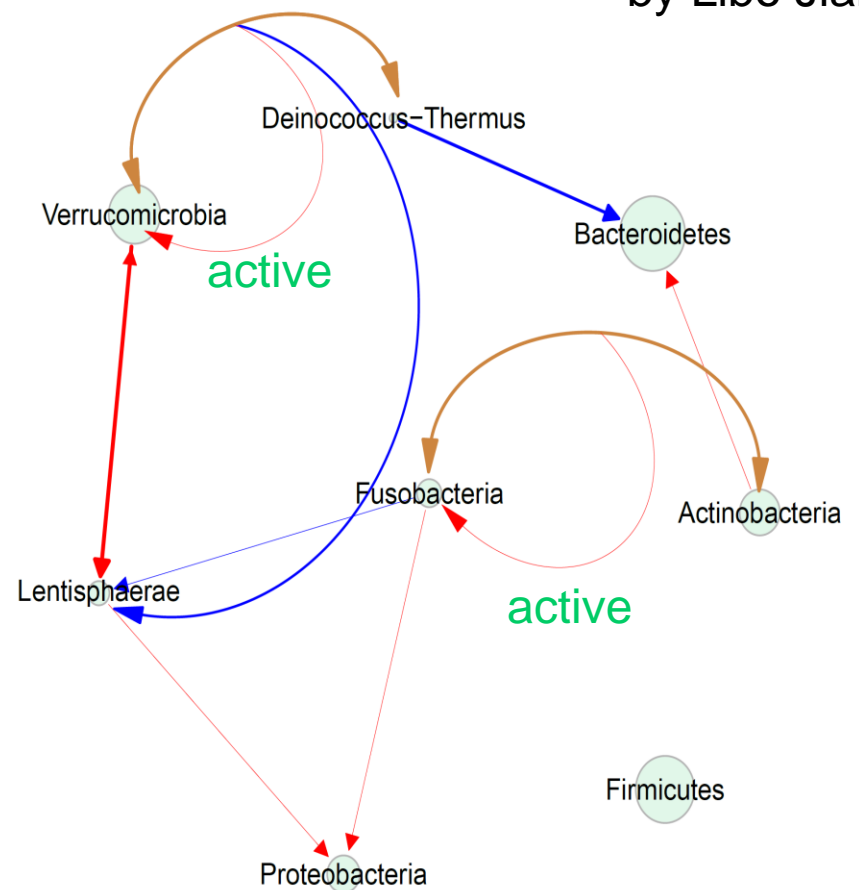


by Libo Jiang

Winter



Summer



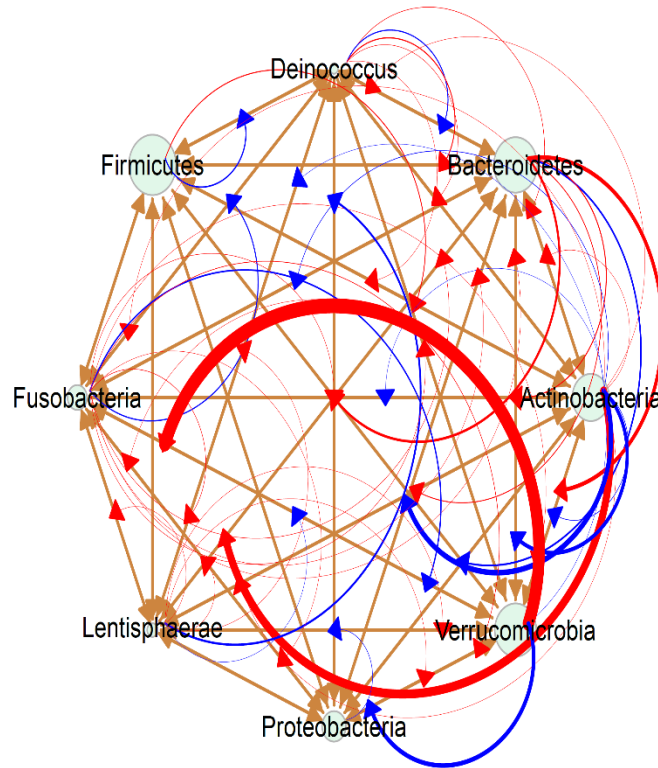
Passive Hypernetworks (mutualism)



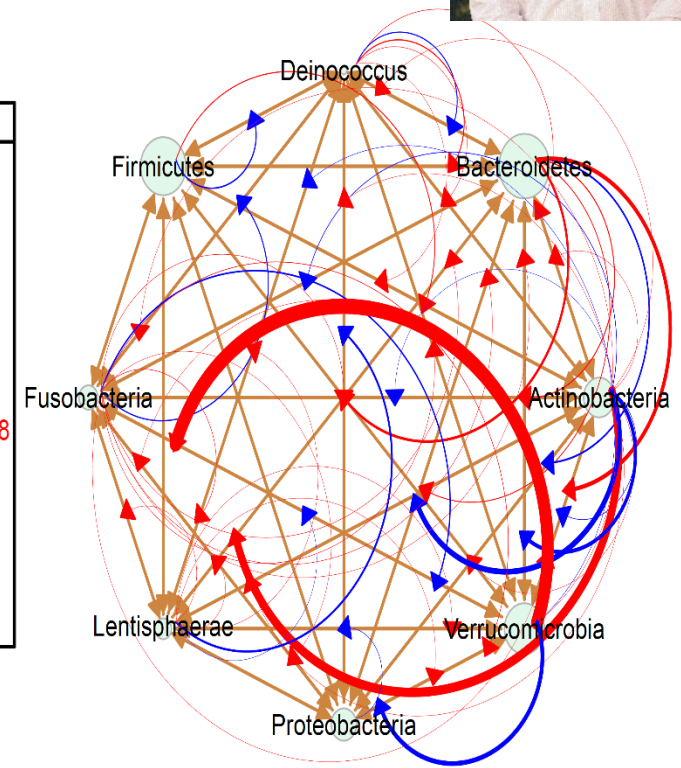
Winter

Summer

Highedge



Phylum	Mutualism
1 Actinobacteria	12 15 16 18 23 25 26 27 28 35
2 Bacteroidetes	16 17 18 26 28
3 Deinococcus -Thermus	12 13 14 23 24
4 Firmicutes	14 34
5 Fusobacteria	18 26 27 28 35 36 37 38 57 58 78
6 Lentisphaerae	36 37 56 57 58 68 78
7 Proteobacteria	68
8 Verrucomicrobia	36 57 58 78



passive

passive

Hypernetwork Implications

Four categories of hypernetworks:

- Mutualism-based hypernetworks
- Antagonism-based hypernetworks
- Altruism-based hypernetworks
- Aggression-based hypernetworks

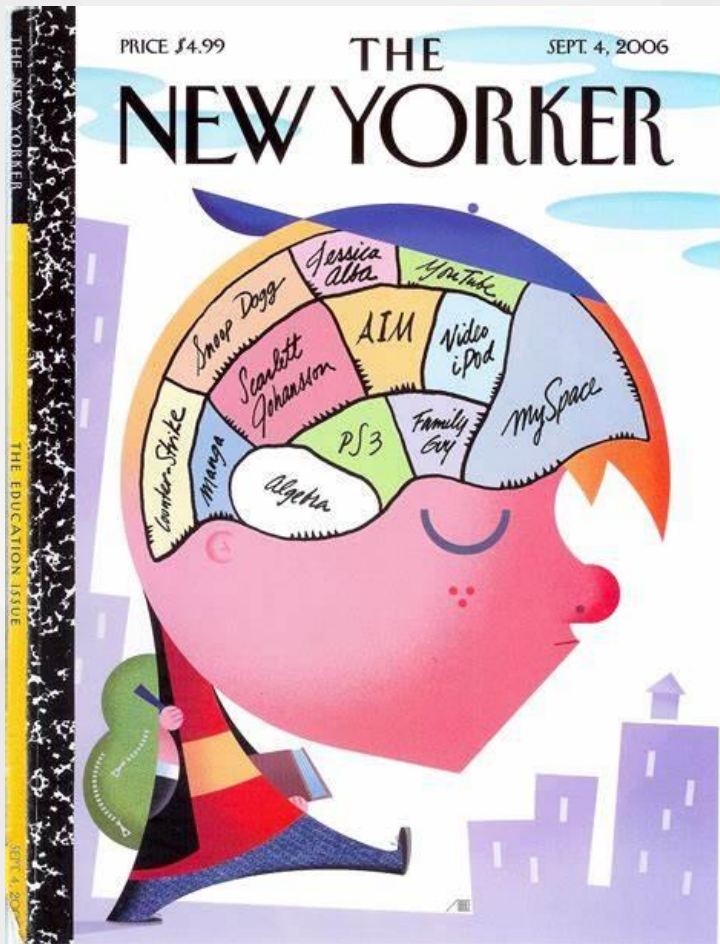
Cancer control as an example

- If a cell promotes the cooperation of two cancer cells, then a drug is developed to dismiss the function of this cell.
- If the cooperation of two cells activates the growth of a cancer cell, then a drug is designed to decouple their cooperation.

How to reconstruct networks from big data?

**We integrate developmental modularity
theory**

Modularity of Mind

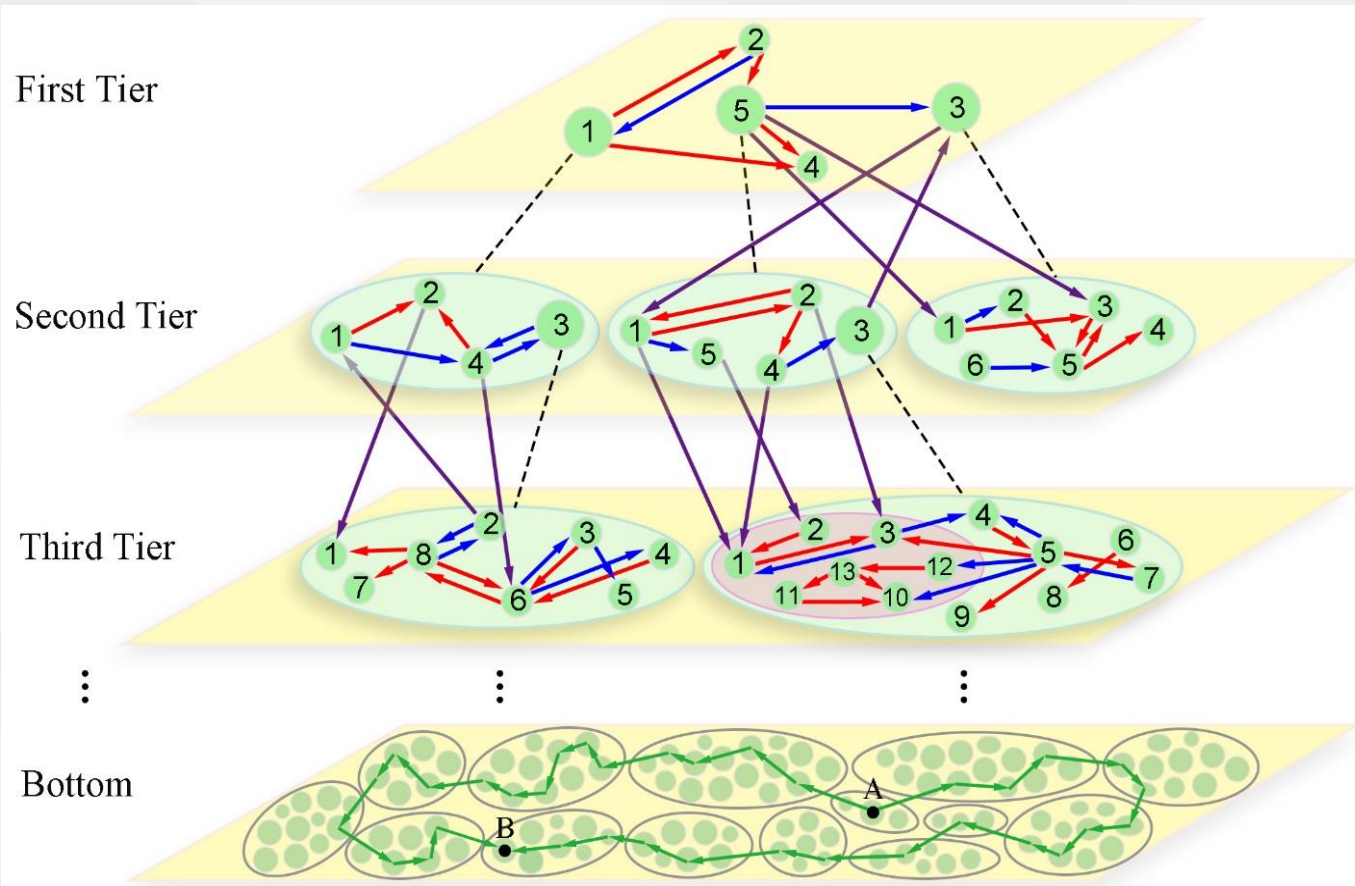


A human brain has distinct regions that think and function differently

- Divide all elements into distinct modules based on their similarity of dynamic change pattern by **functional clustering** (Kim et al. 2008, Genetics; Wang et al. 2012, Briefings in Bioinformatics).
- Divide each module into submodule
- Divide each submodule into sub-submodule.

- This process stops until the number of elements reaches Dunbar's number.

A multilayer, multiplex, and multifunctional network from any number of elements



Informative
- Causal
- Stable
- Sparse

Dynamic

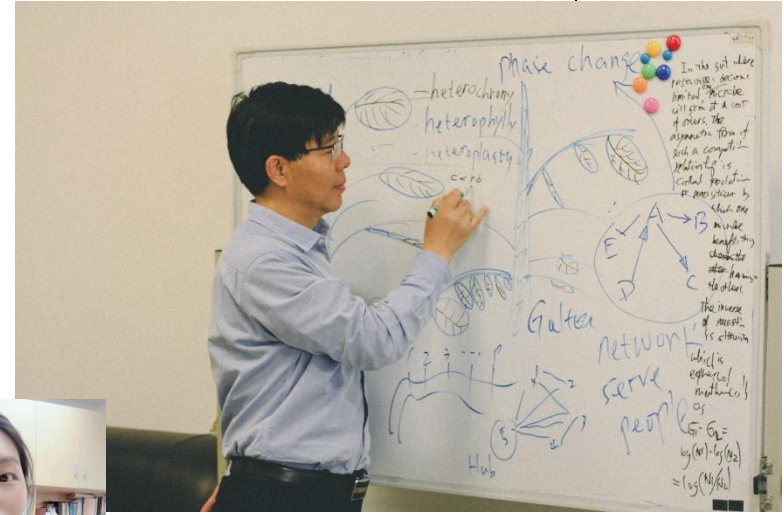
Omnidirectional

Personalized

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- Beijing Institute of Mathematical Sciences and Applications
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- Beijing Forestry University
- Pennsylvania State University



Christopher Griffin/Shawn Rice/
Claudia Gragnoli/Chandra Belani



董昂



冯莉



吴双



杨登程



王静



龚慧莹



车金灿



姜立波



桑蒙蒙



An interdisciplinary team



Thank You!

