

بِسْمِ اللَّهِ
الرَّحْمَنِ
الرَّحِيمِ

Balance Theory of Signed Genetic Interactions Reveals Differences in Cancerous and Healthy Cells

Abbas K. Rizi

Jan 13, 2020

CX Group, CS Dept.,
Aalto University

abbas.sitpor.org



Aalto University
School of Science





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"Social media has begun to reform the face of human life. We need to understand how our new societies work."

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We do research on statistical physics, social media, internet, and finance. CCNSD aims to bridge the gap between the research on social sciences and physics via its data-oriented projects. At our center, we try to create a popular work environment that fosters curiosity, creativity and imagination.



I am going to talk about:

- ◎ Balance Theory
- ◎ Molecular Biology
- ◎ Inference Problems (Inverse Stat. Phys.)
- ◎ Results!



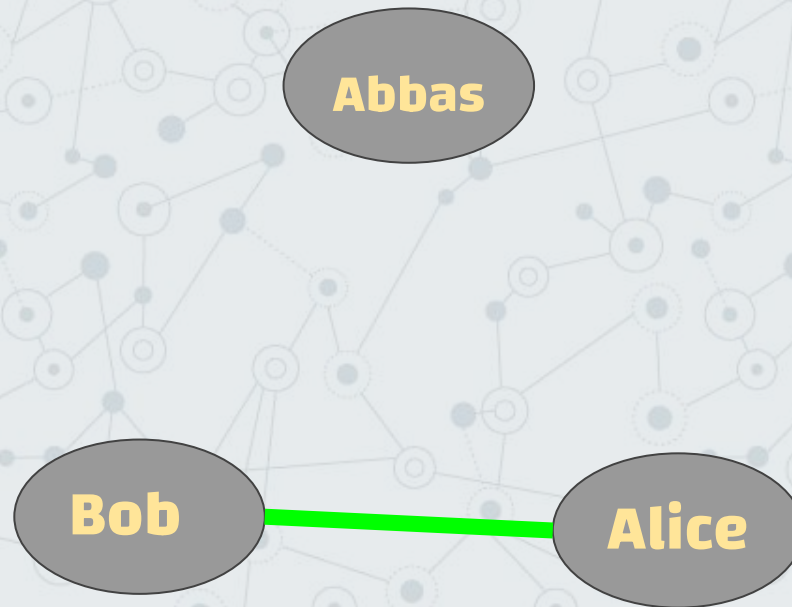
Balance Theory

Heider, Fritz (1958). *The Psychology of Interpersonal Relations*. John Wiley & Sons.



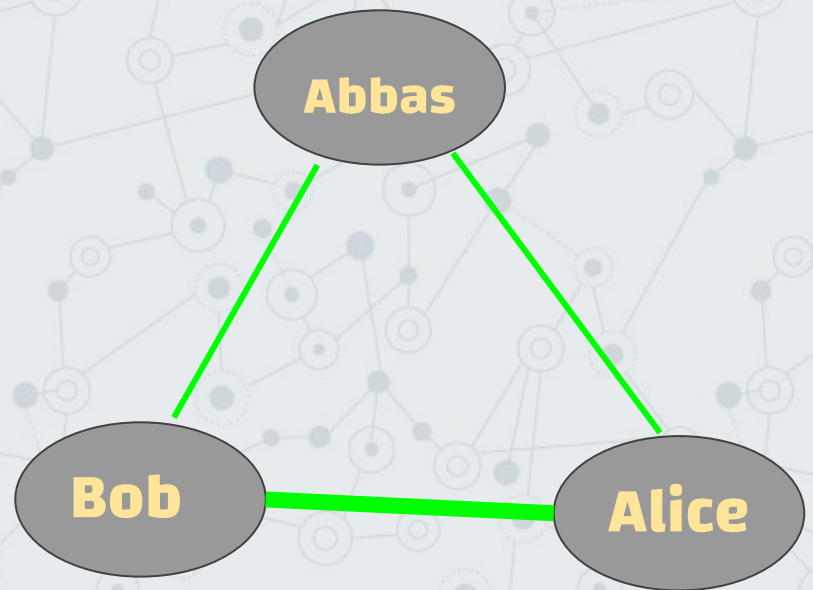
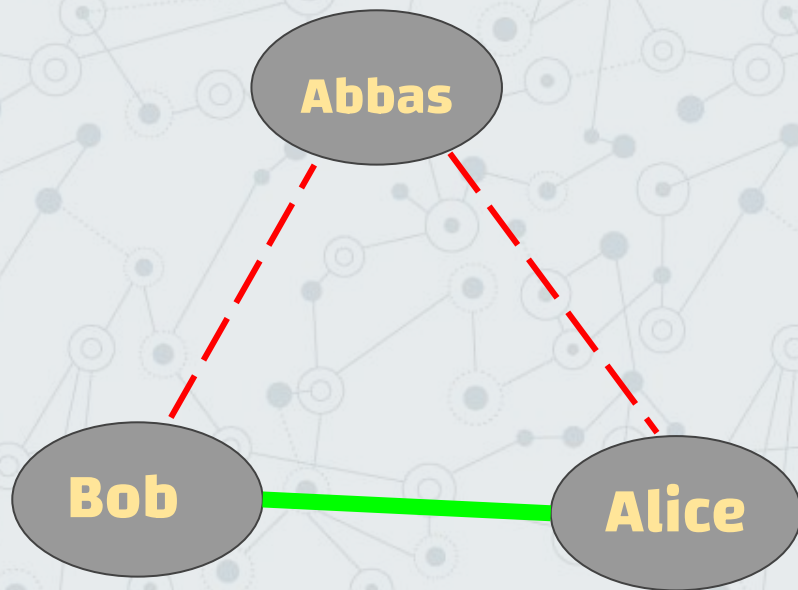
Balance Theory

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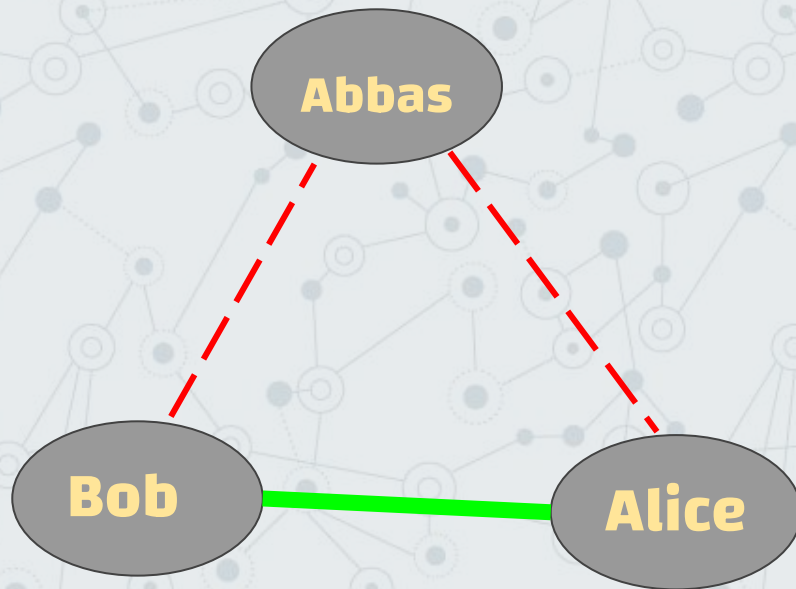
Balance Theory

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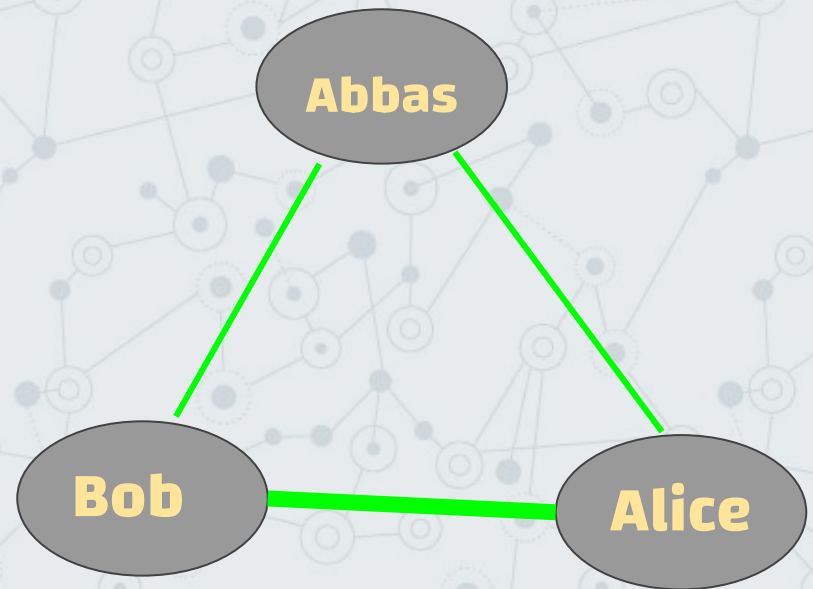


Balance Theory

Heider, Fritz (1958). *The Psychology of Interpersonal Relations*. John Wiley & Sons.



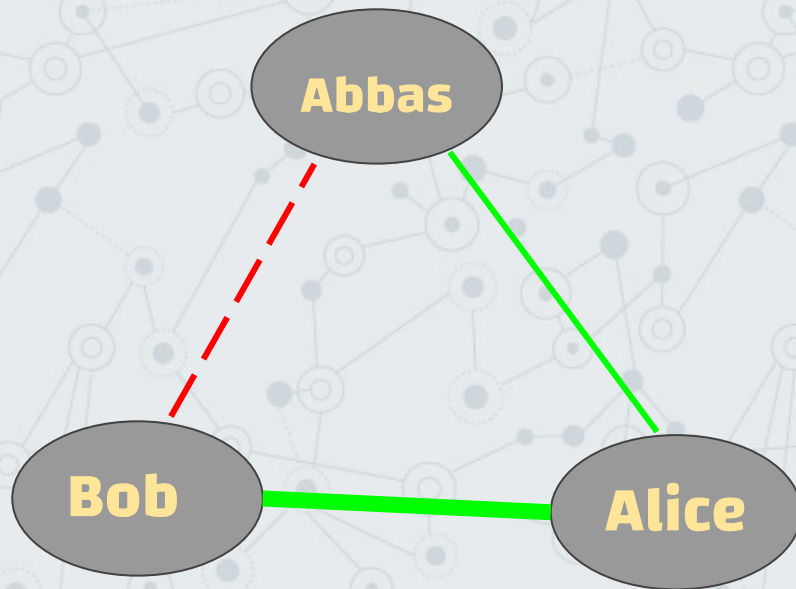
balanced



balanced

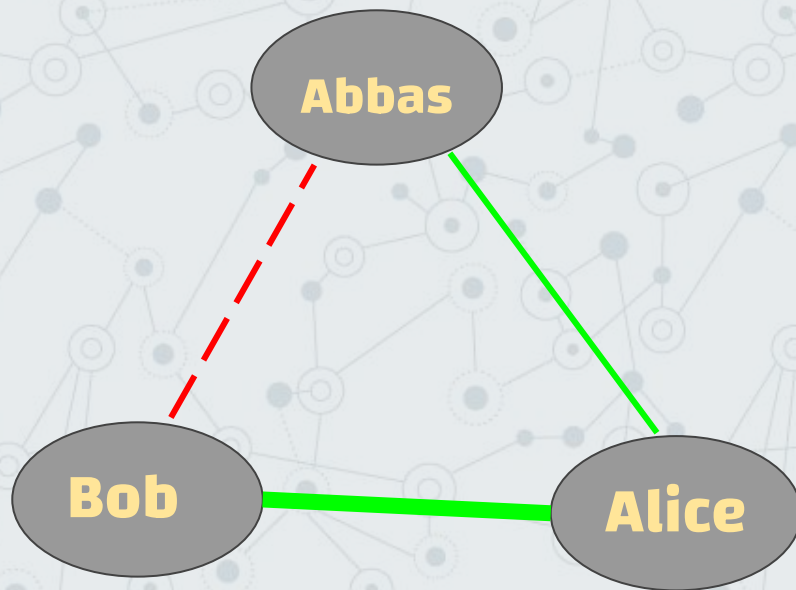
Balance Theory

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Balance Theory

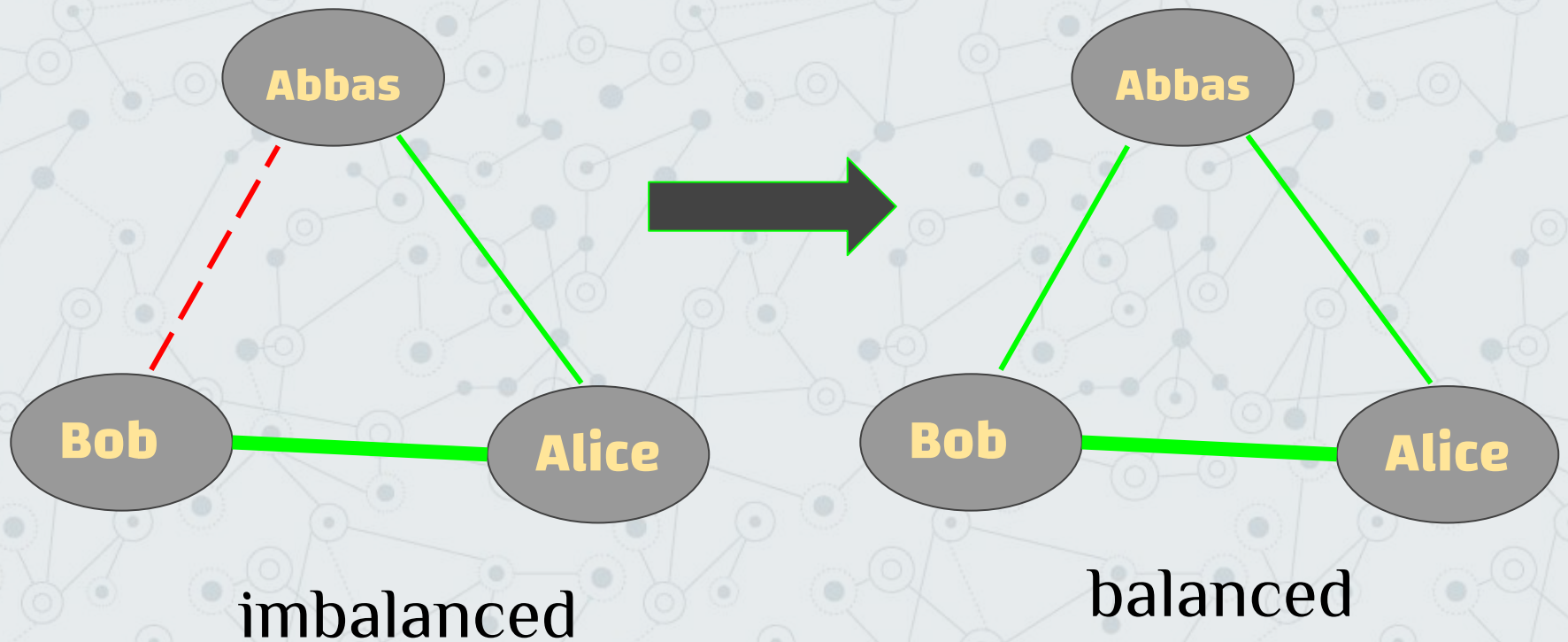
Heider, Fritz (1958). *The Psychology of Interpersonal Relations*. John Wiley & Sons.



imbalanced

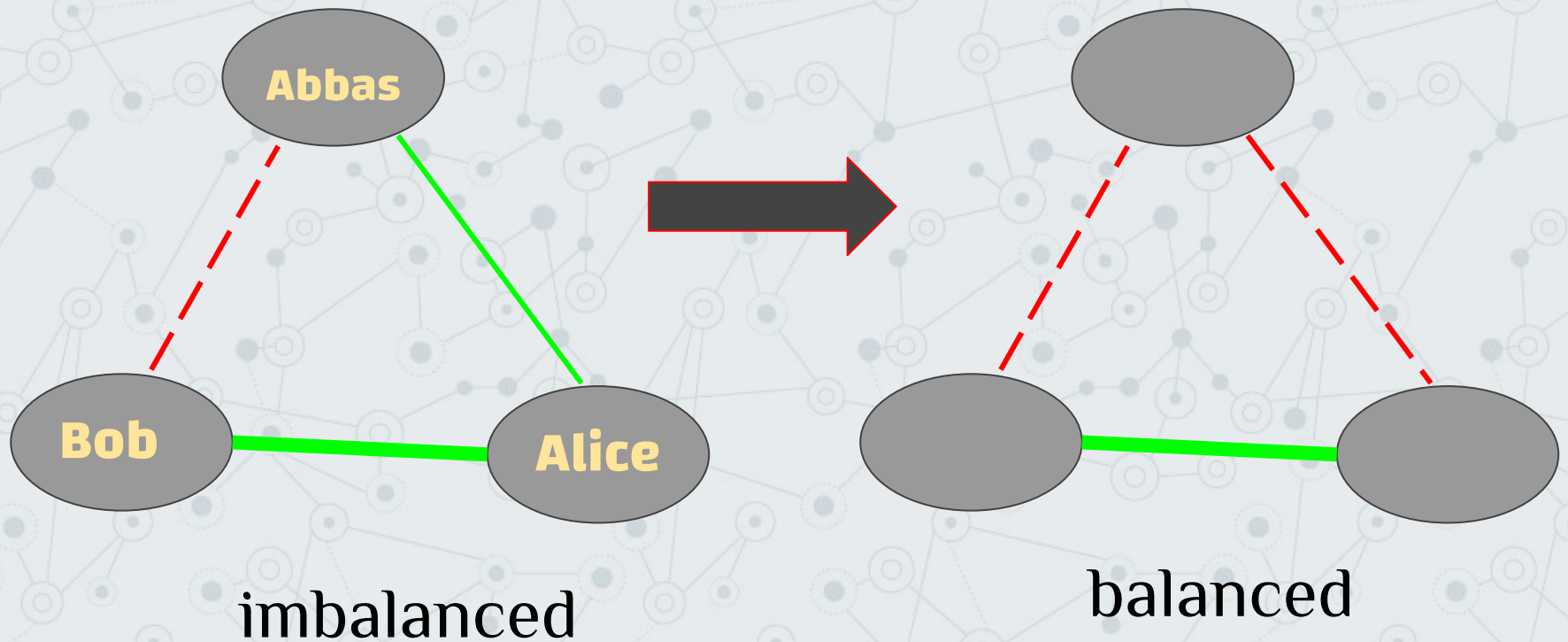
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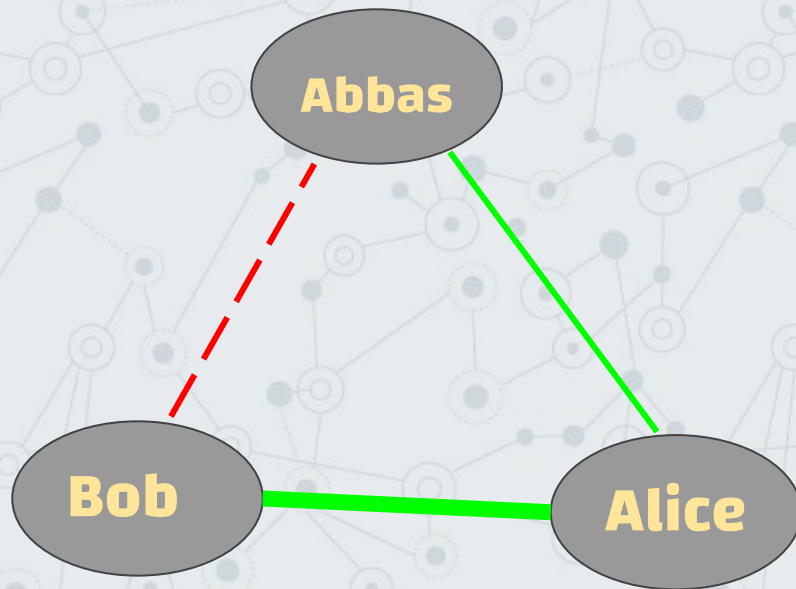
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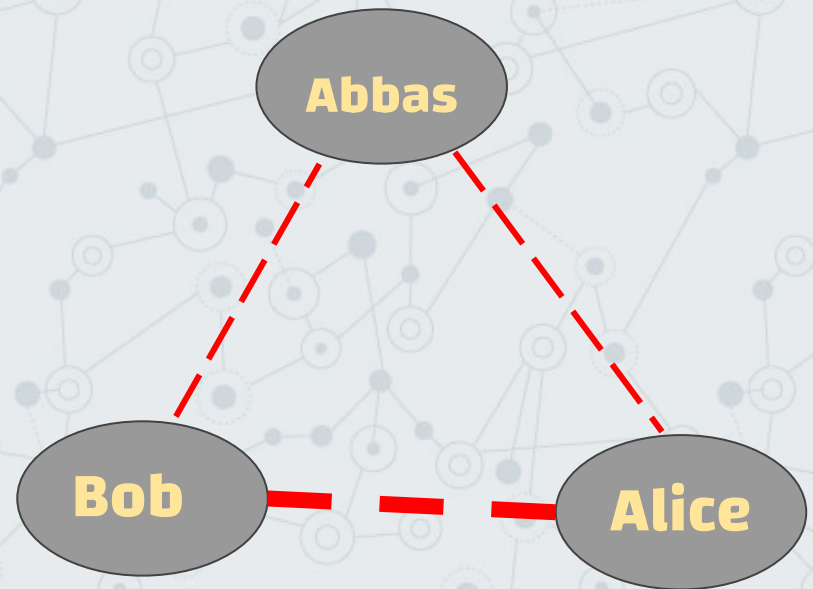


Balance Theory

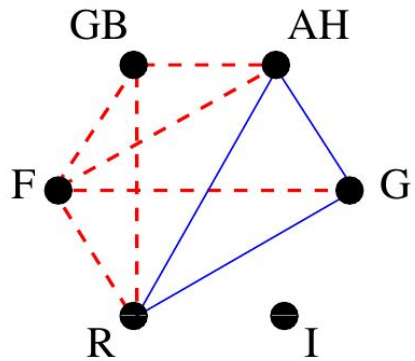
Heider, Fritz (1958). *The Psychology of Interpersonal Relations*. John Wiley & Sons.



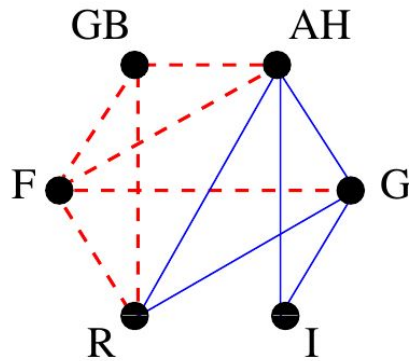
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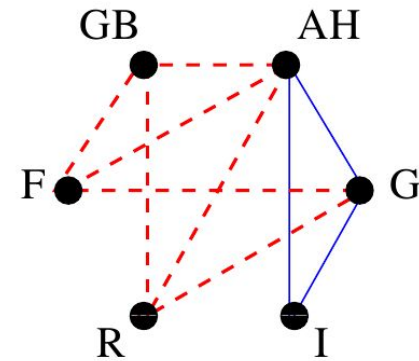
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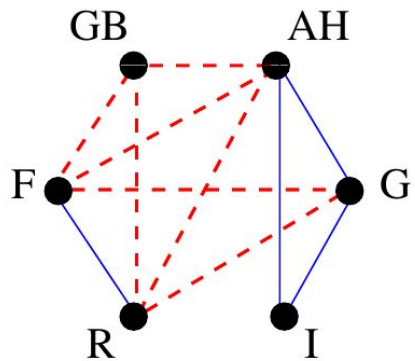
3 Emperor's league 1872–81



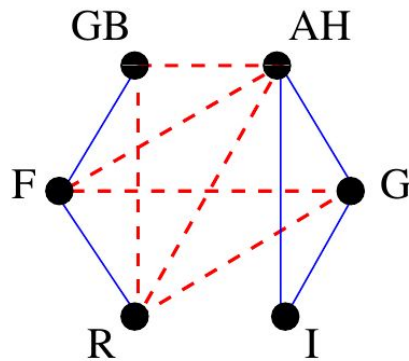
Triple Alliance 1882



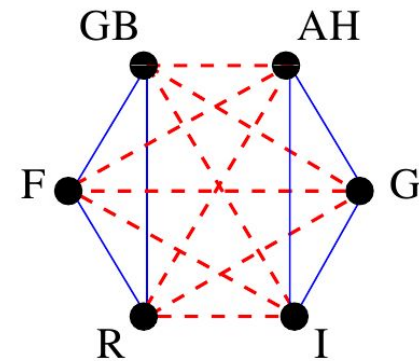
German–Russian Lapse 1890



French–Russian Alliance 1891–94



Entente Cordiale 1904



British–Russian Alliance 1907

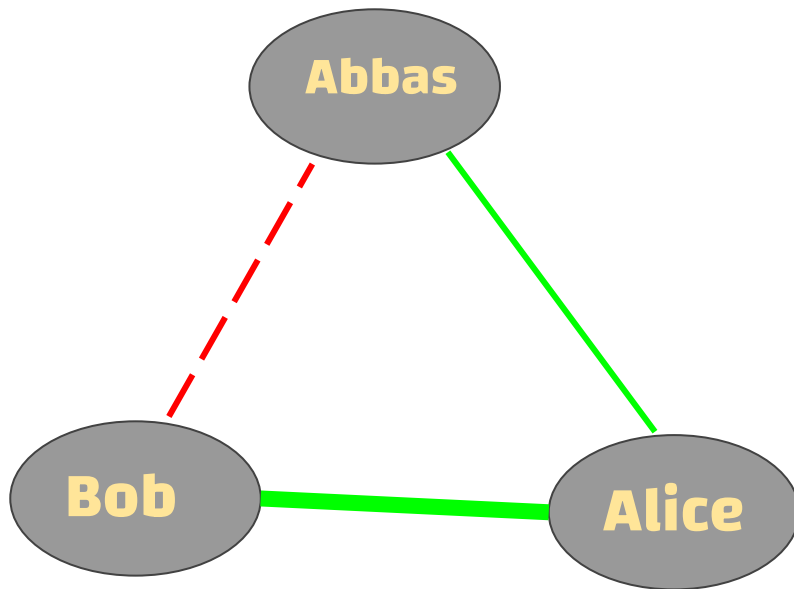
Evolution of the major relationship changes between the protagonists of World War I from 1872–1907.

GB = Great Britain, AH = Austria-Hungary, G = Germany, I = Italy, R = Russia, and F = France.

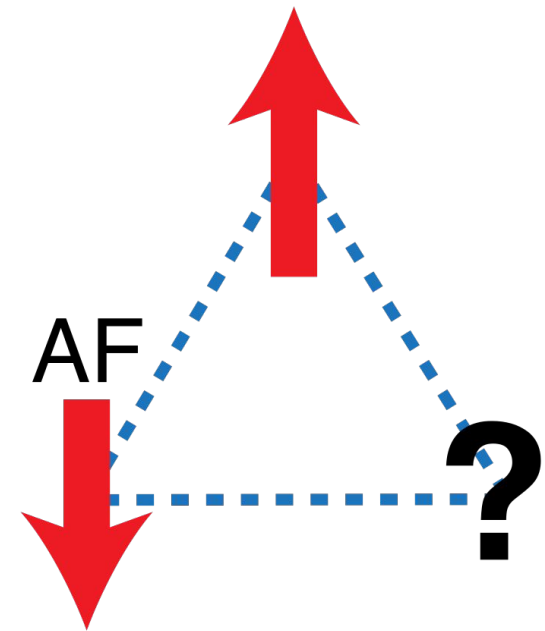
T. Antal, P. L. Krapivsky, and S. Redner,
 “Social Balance on Networks: The Dynamics of Friendship and Enmity,”

Balance Theory

Heider, Fritz (1958). *The Psychology of Interpersonal Relations*. John Wiley & Sons.



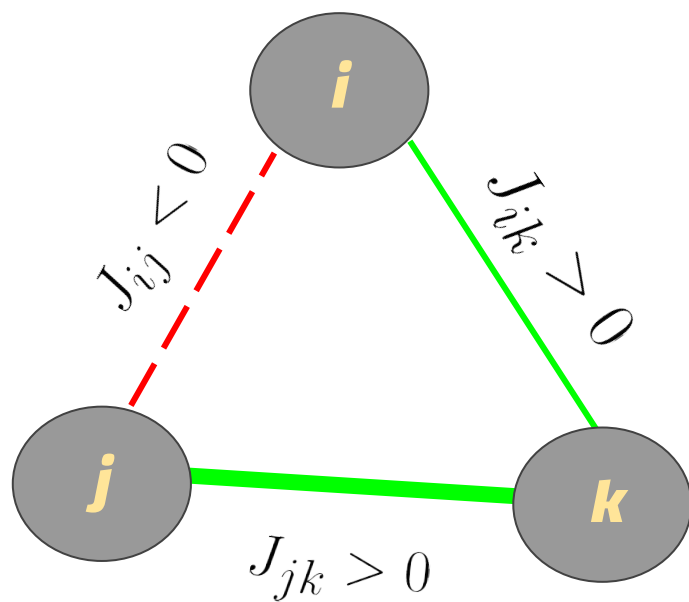
imbalanced



frustration

The Energy Landscape of Social Balance

Seth A. Marvel, Steven H. Strogatz, and Jon M. Kleinberg

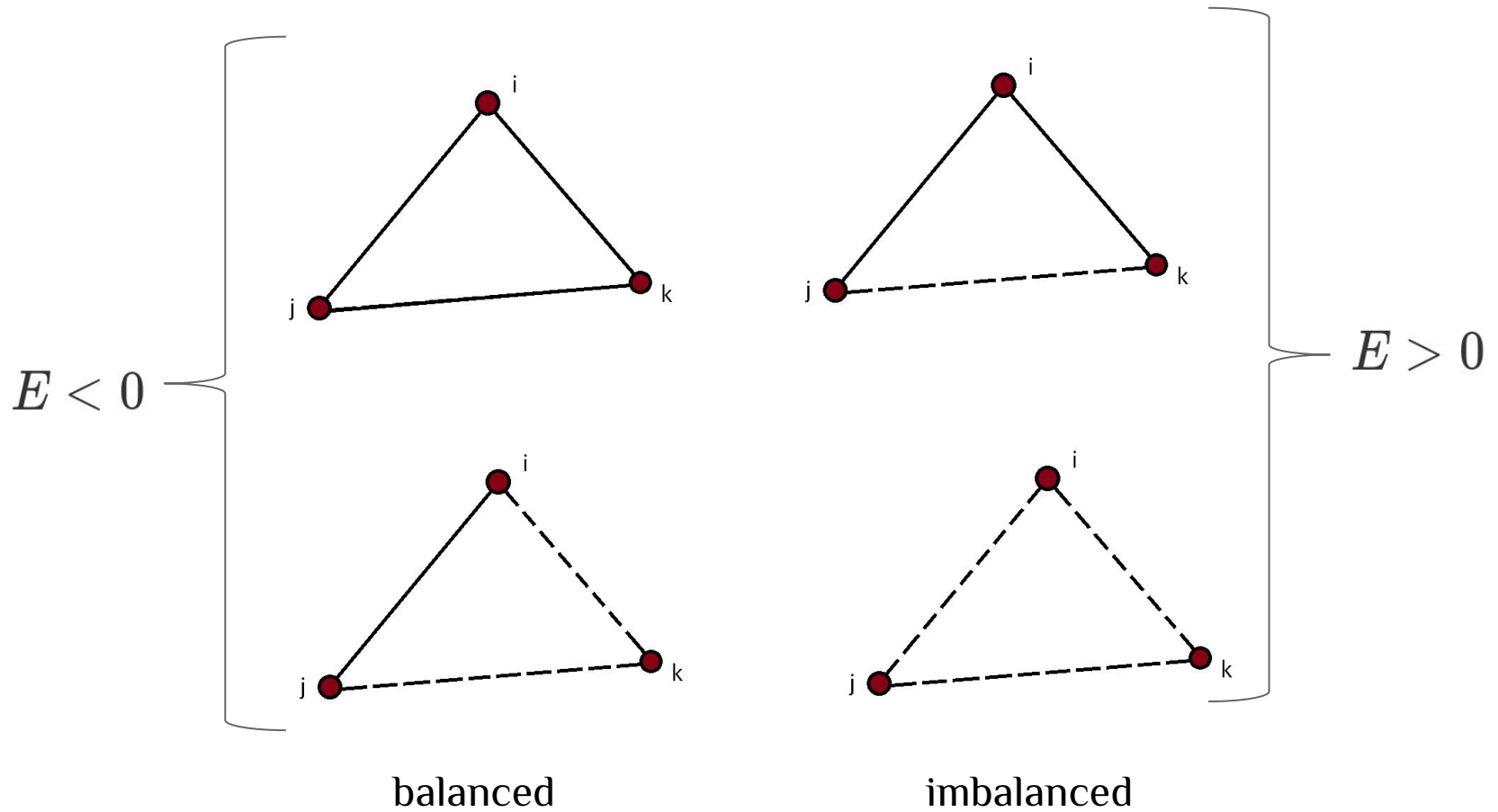


$$E_{\Delta} = -J_{ij}J_{jk}J_{ki}$$

$$J_{ij} \in R$$

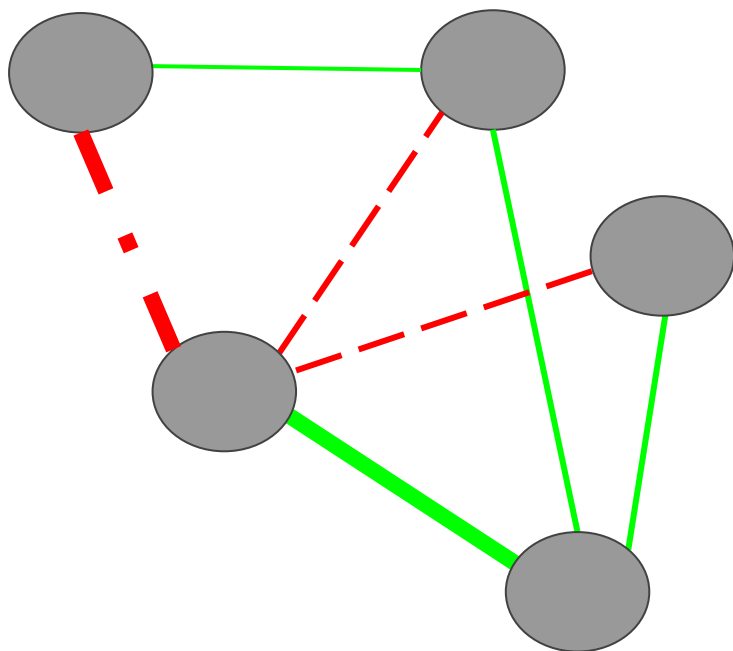
The Energy Landscape of Social Balance

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The Energy Landscape of Social Balance

Seth A. Marvel, Steven H. Strogatz, and Jon M. Kleinberg



$$E_{Net} = \sum_{\Delta} E_{\Delta}$$

$$E_{\Delta} = -J_{ij}J_{jk}J_{ki}$$

$$J_{ij} \in R$$

International Networks

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Continues-time Dynamics

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What About Biological Networks? *e.g. The GRN*

Central Dogma of Molecular Biology

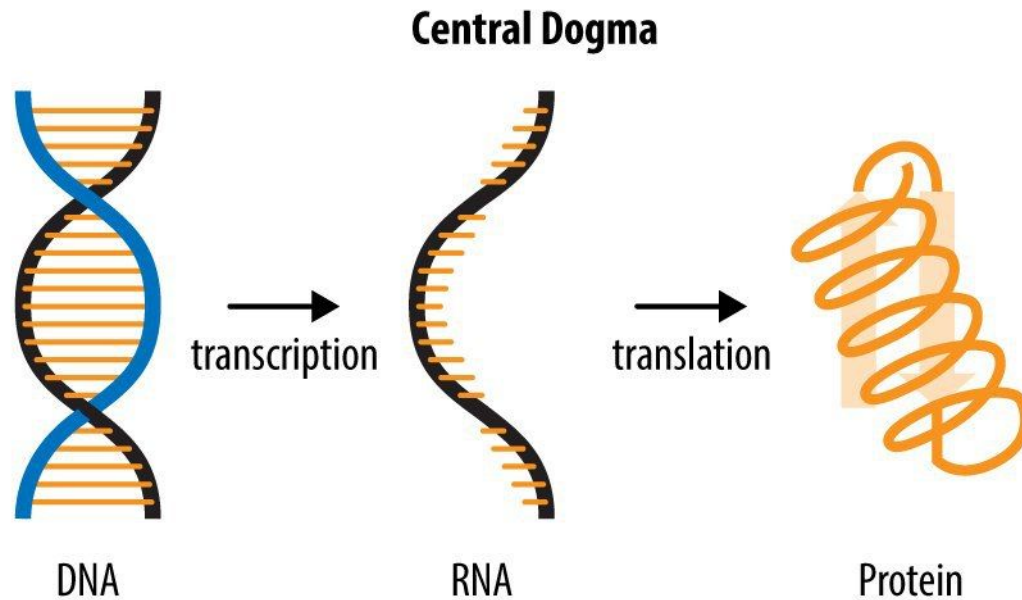
All the information needed to run a cell is stored in a molecule called **DNA**.

A section of DNA that encodes the information to make a protein (via mRNA) is called a **gene**.

The amount of mRNA produced from a given gene is called the **expression** of that gene.



Central Dogma of Molecular Biology



Gene Regulatory Network (GRN)



Regulatory effects

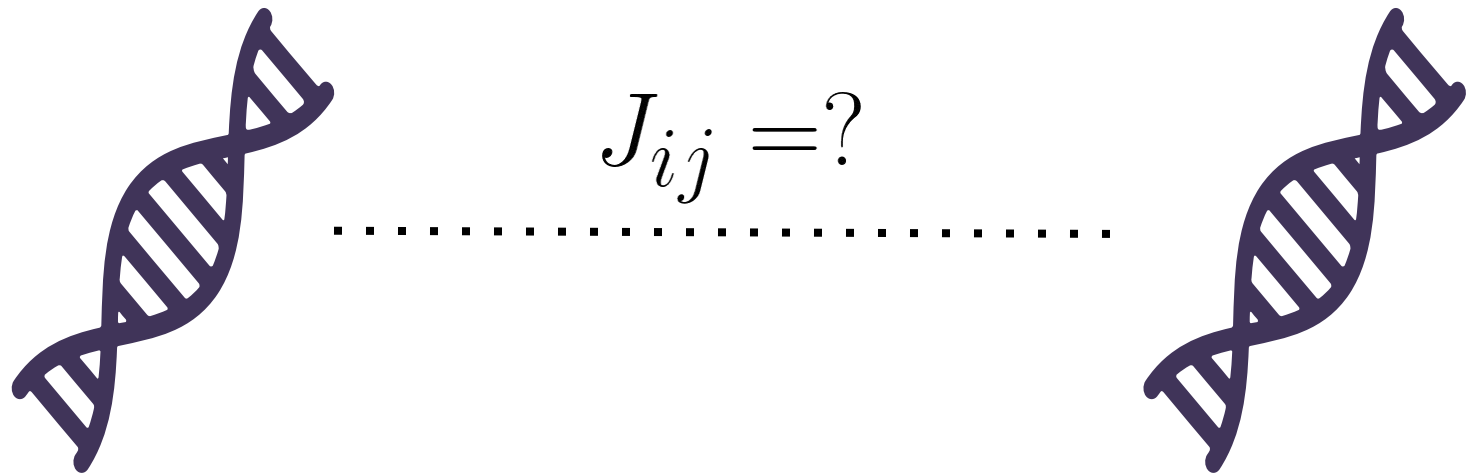


▲ up-regulation

▼ down-regulation

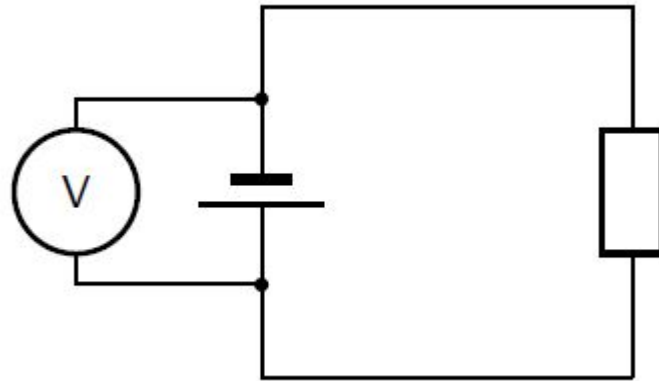


Gene Regulatory Network (GRN)





$$J_{ij} = ?$$



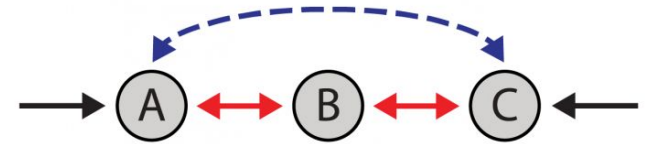
Data Set

- ◎ BRCA: Breast Invasive Carcinoma
- ◎ The Cancer Genome Atlas (TCGA)
- ◎ RNA Sequencing
(The state-of-the-art technology to measure mRNA)
- ◎ Expression Levels for 20532 Genes
- ◎ 114 Normal and 763 Cancerous Samples

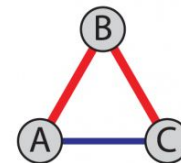
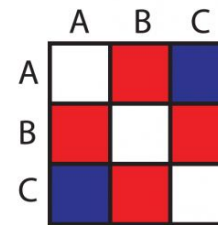


GRN Reconstruction

Reaction system reconstruction



Correlation



Partial correlation

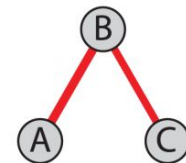
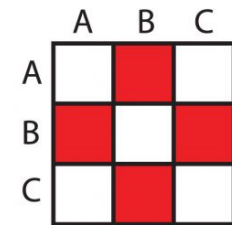
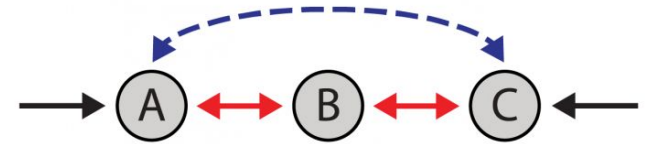


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<http://journals.plos.org/ploscompbiol/article?id=10.1371/journal.pcbi.1004182>

Inferring the Interactions

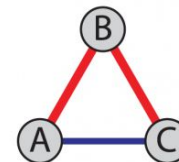
1. Fixing a Good Model!
2. Inferring the Parameters

Reaction system reconstruction



Correlation

	A	B	C
A		Red	Blue
B	Red		Red
C	Blue	Red	



Partial correlation

	A	B	C
A		Red	
B	Red		Red
C		Red	

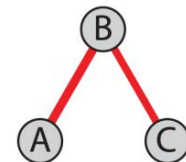


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<http://journals.plos.org/ploscompbiol/article?id=10.1371/journal.pcbi.1004182>

GRN Reconstruction

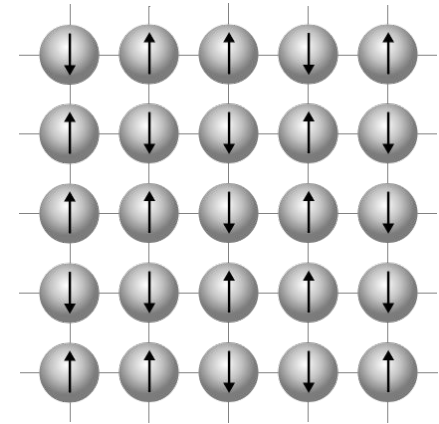
Statistical Physics

Fine-Grained
Microscopic Rules

Coarse-Grained
Observables

Inverse Stat. Phys.

Ising Model in Statistical Physics



Fine-Grained
Microscopic Rules

$$P(\{X_i\}) = \frac{1}{Z} \exp\left[\sum_{i<j} J_{ij} X_i X_j\right]$$

Ising Model in Statistical Physics

Coarse-Grained
Observables

$$\begin{cases} M \equiv \langle X_i \rangle \\ \xi \equiv \langle X_i X_j \rangle \end{cases}$$



Fine-Grained
Microscopic Rules

$$P(\{X_i\}) = \frac{1}{Z} \exp\left[\sum_{i<j} J_{ij} X_i X_j\right]$$



Inverse Ising Problem

Coarse-Grained
Observables

$$\begin{cases} M \equiv \langle X_i \rangle \\ \xi \equiv \langle X_i X_j \rangle \end{cases}$$



Fine-Grained
Microscopic Rules

to be inferred

$$P(\{X_i\}) = \frac{1}{Z} \exp\left[\sum_{i<j} J_{ij} X_i X_j\right]$$

GRN Reconstruction

1. Fixing a Good Model!

Undirected Pairwise Maximum-Entropy Probability Model

$$P(\{X_i\}) = \frac{1}{Z} \exp\left[\sum_{i < j} J_{ij} X_i X_j\right]$$

2. Inferring the Parameters,

$J = ?$



GRN Reconstruction

$$\text{maximize } S = - \int_{\mathbf{x}} P(\mathbf{x}) \ln P(\mathbf{x}) \, d\mathbf{x}$$

$$\left\{ \begin{array}{l} \int_{\mathbf{x}} P(\mathbf{x}) \, d\mathbf{x} = 1, \\ \langle x_i \rangle = \int_{\mathbf{x}} P(\mathbf{x}) x_i \, d\mathbf{x} = \frac{1}{M} \sum_{m=1}^M x_i^m = \overline{x_i}, \\ \langle x_i x_j \rangle = \int_{\mathbf{x}} P(\mathbf{x}) x_i x_j \, d\mathbf{x} = \frac{1}{M} \sum_{m=1}^M x_i^m x_j^m = \overline{x_i x_j} \end{array} \right.$$

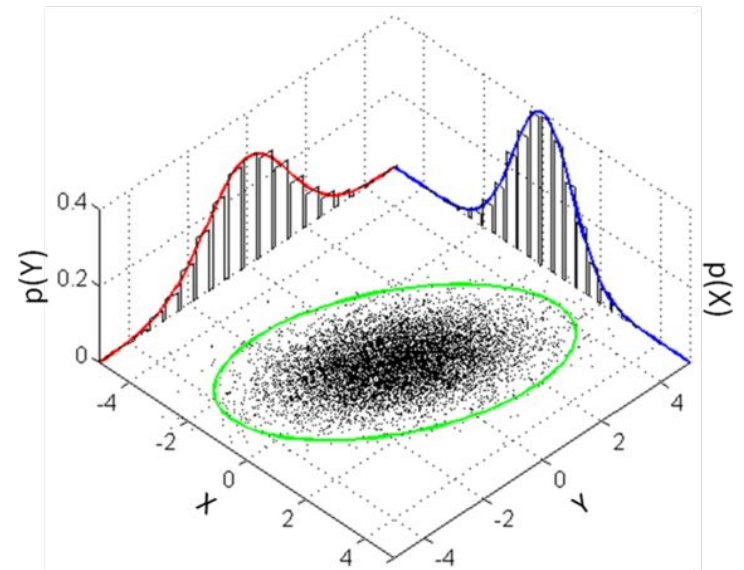
$$L = 20,532 \text{ and } M = 764 + 114$$

GRN Reconstruction

$$P(\mathbf{x}; \langle \mathbf{x} \rangle, C) = (2\pi)^{-L/2} \det(C)^{-1/2} \exp\left(-\frac{1}{2}(\mathbf{x} - \langle \mathbf{x} \rangle)^T C^{-1}(\mathbf{x} - \langle \mathbf{x} \rangle)\right)$$

Inferring Pairwise Interactions from Biological Data
Using Maximum-Entropy Probability Models

Richard R. Stein, Debora S. Marks, Chris Sander



GRN Reconstruction

$$P(\mathbf{x}; \langle \mathbf{x} \rangle, C) = (2\pi)^{-L/2} \det(C)^{-1/2} \exp\left(-\frac{1}{2}(\mathbf{x} - \langle \mathbf{x} \rangle)^T C^{-1}(\mathbf{x} - \langle \mathbf{x} \rangle)\right)$$

$$P(\{X_i\}) = \frac{1}{Z} \exp\left[\sum_{i < j} J_{ij} X_i X_j\right]$$

$$J = -C^{-1}$$

Inferring Pairwise Interactions from Biological Data
Using Maximum-Entropy Probability Models

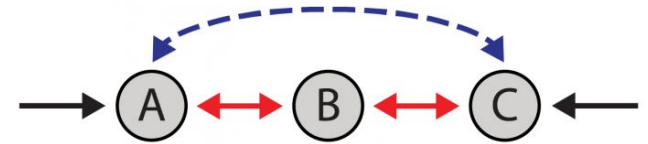
Richard R. Stein, Debora S. Marks, Chris Sander

GRN Reconstruction

$$J = -C^{-1}$$

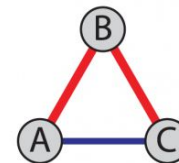
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Reaction system reconstruction



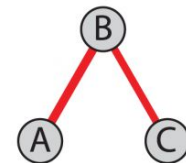
Correlation

	A	B	C
A		Red	Blue
B	Red		Red
C	Blue	Red	

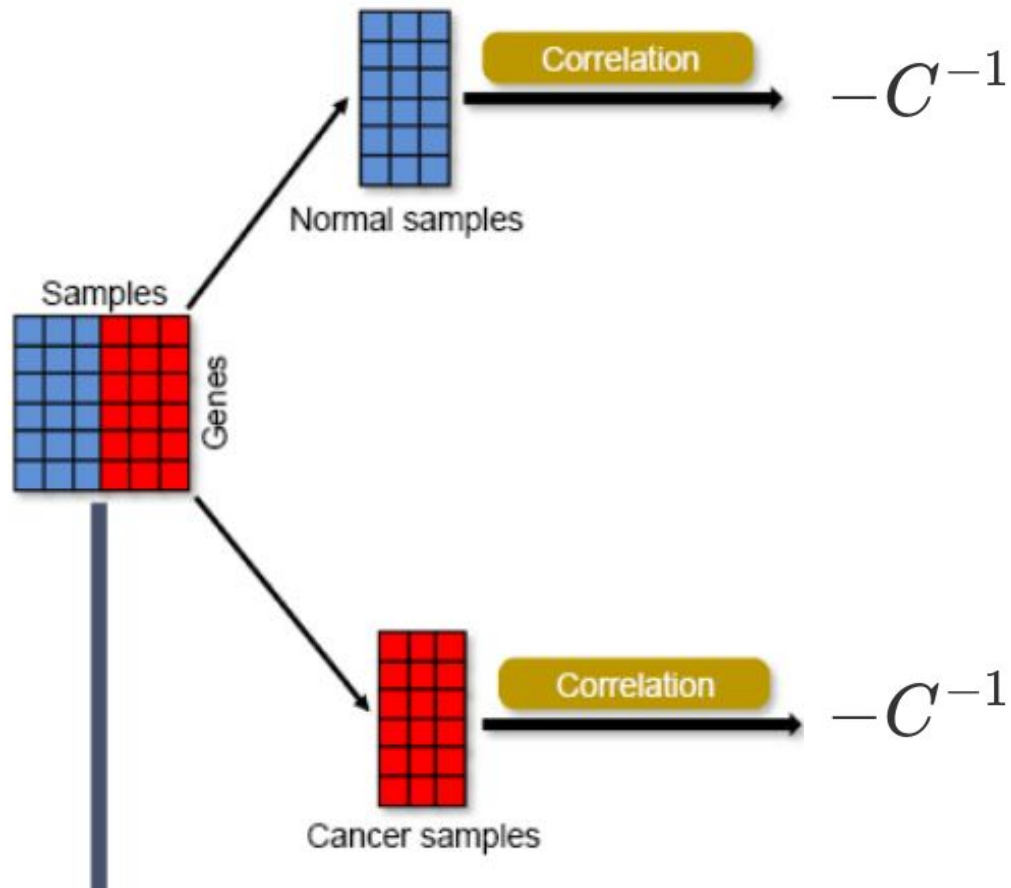


Partial correlation

	A	B	C
A		Red	
B	Red		Red
C		Red	



GRN Reconstruction



$$J = -C^{-1}$$

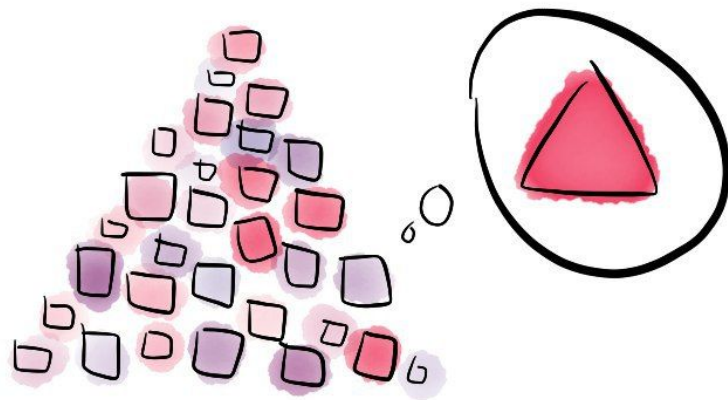
GLasso (Graphical Least Absolute Shrinkage and Selection Operator)

$$\left\{ \begin{array}{l} X \sim N(0, \Sigma) \\ X_1, X_2, \dots, X_n \end{array} \right. \quad \Theta = \Sigma^{-1}$$

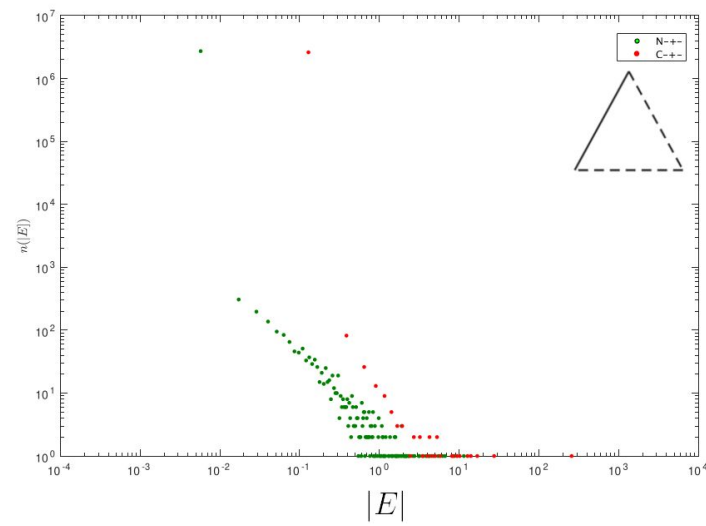
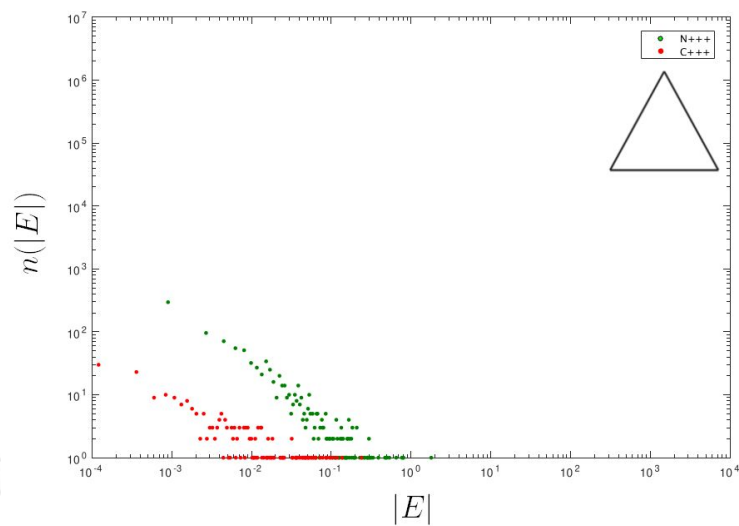
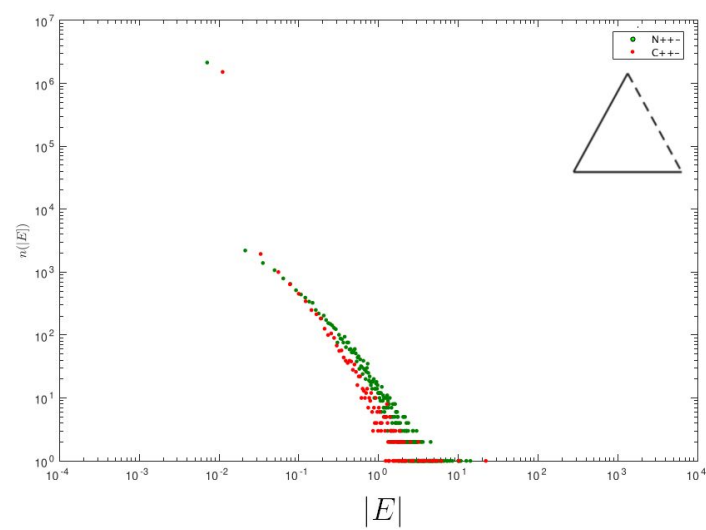
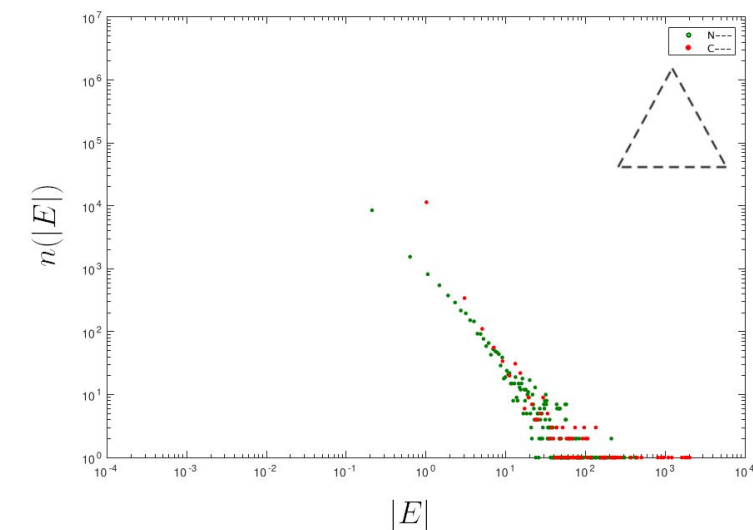
$$\hat{\Theta} = * \operatorname{argmin}_{\Theta \geq 0} \left(\operatorname{tr}(S\Theta) - \log \det(\Theta) + \lambda \sum_{j \neq k} |\Theta_{jk}| \right)$$

S: sample covariance

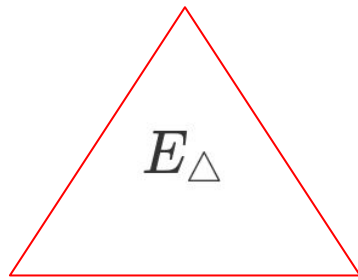
λ : penalizing parameter



Results



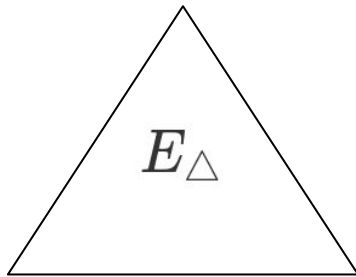
Coarse-graining the Network



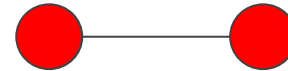
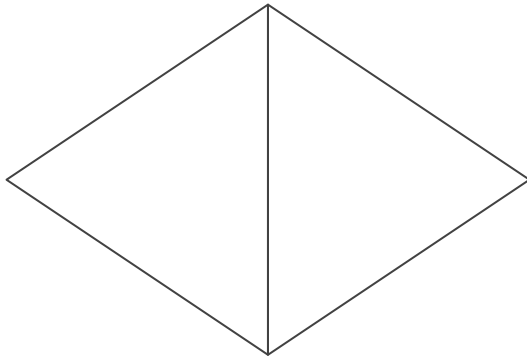
A node with energy E_{Δ}



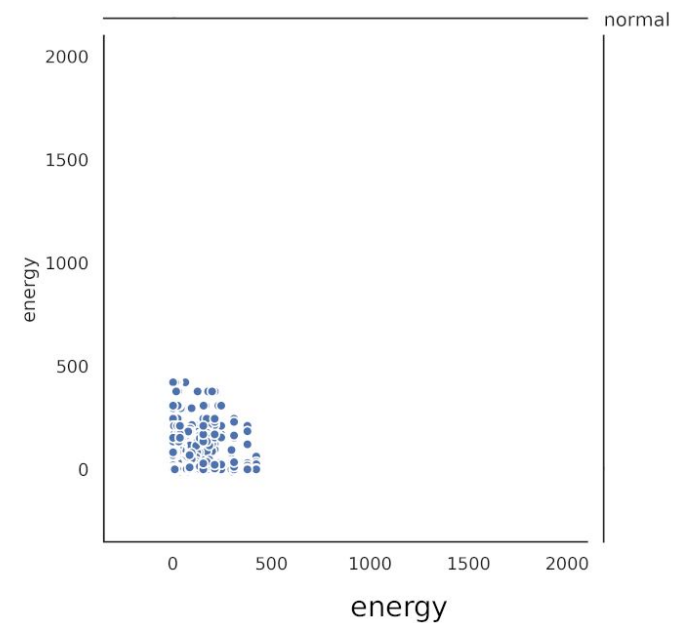
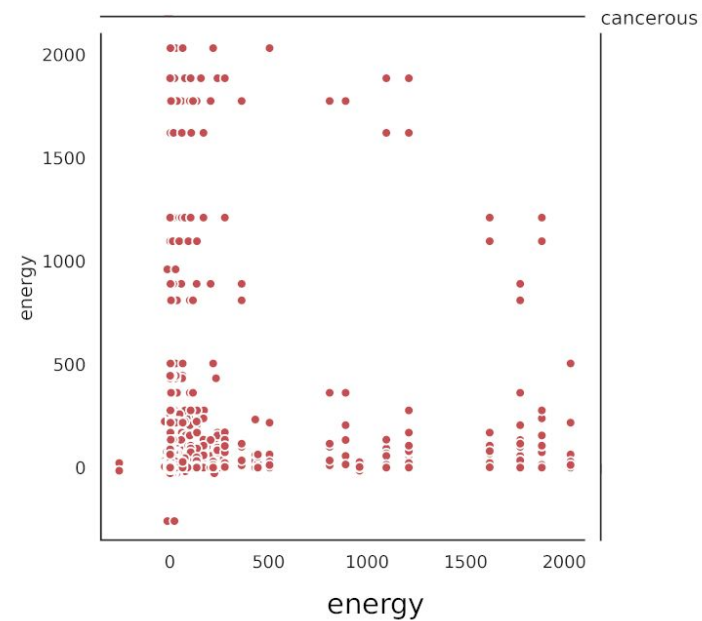
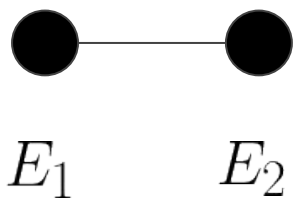
Coarse-graining the Network



 A node with energy E_Δ



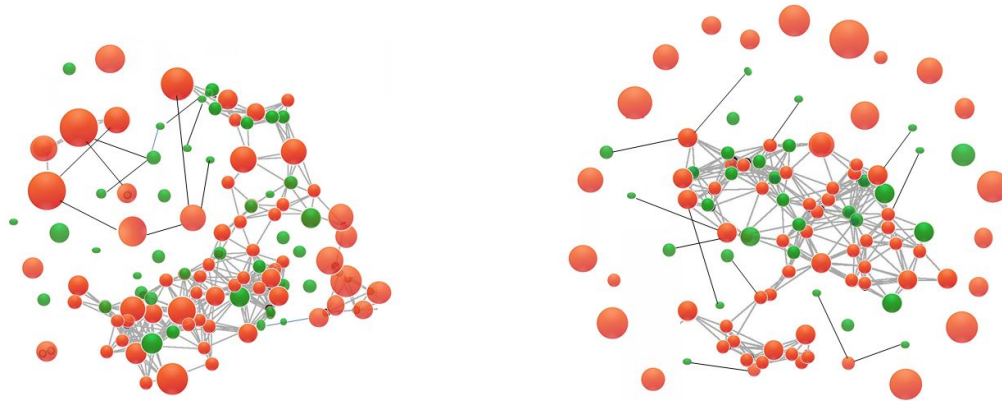
Results



Results

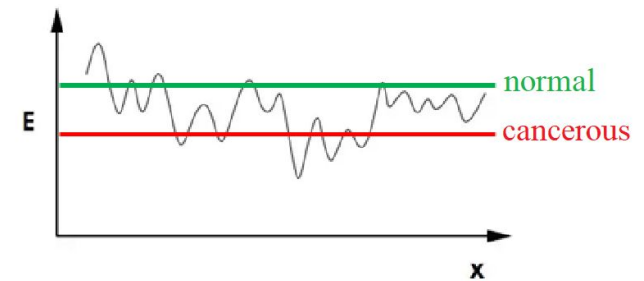
35,984 units
27,239 units

E Gap ~ 10,000 units



(a) cancerous coarse-grained network

(b) normal coarse-grained network



(c) energy levels and local minima

The background of the slide features a complex, light gray network pattern. It consists of numerous small circles, some of which are solid gray and others are hollow with a gray outline. These circles are interconnected by thin, light gray lines, creating a web-like structure that fills the entire background.

Thank You

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