

# A max-margin framework for supervised graph completion

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# Graph Completion

- given

- features for each potential edge

$$\mathbf{X} = \{\mathbf{x}_{j,k}\}$$

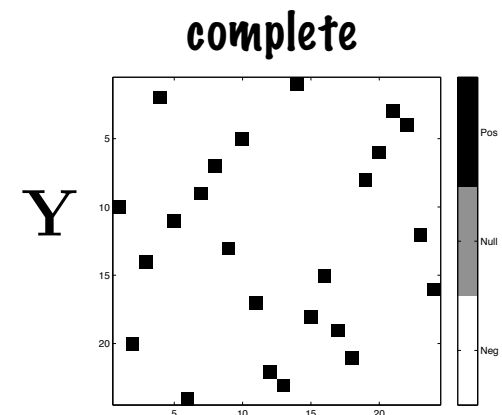
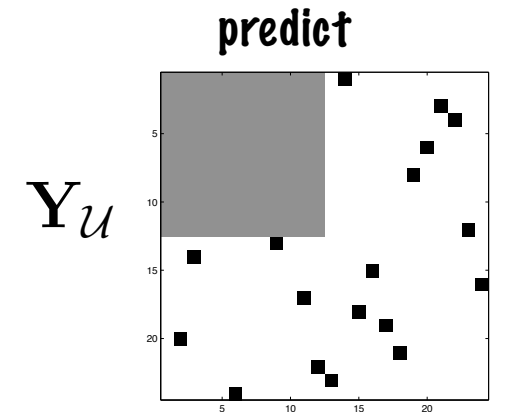
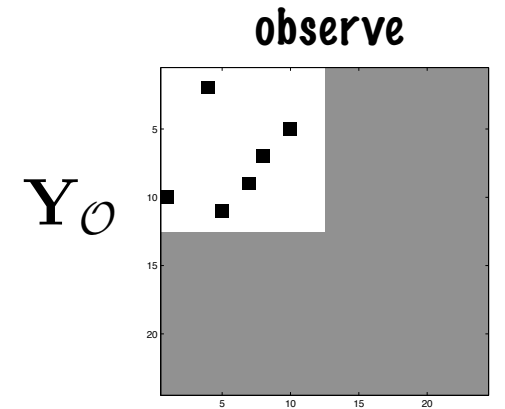
- partially observed connectivity

$$\mathbf{Y}_O \quad \text{from} \quad \mathbf{Y} = \{y_{j,k}\} \quad y_{j,k} \in \{0, 1\}$$

- task

- complete connectivity

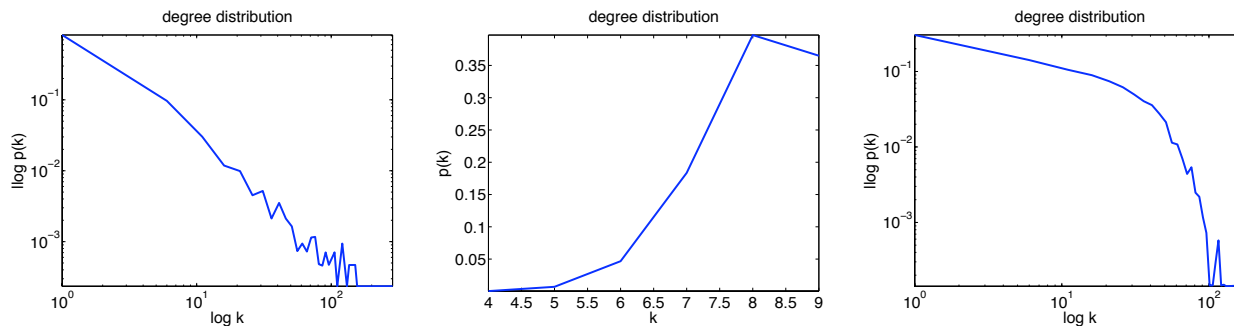
$$\mathbf{Y} = \mathbf{Y}_O \cup \mathbf{Y}_U$$



# Graph Completion

- approach 1: topology

- topology of complete connectivity is similar to observed connectivity
- e.g. Liben-Nowell and Kleinberg, 2003



- approach 2: attribute-value learning

- "on" / "off" property of edges is related to their features
- e.g. Yamanishi, Vert, and Kanehisa, 2003

- idea: a marriage of structured-output models and degree-constrained subgraphs

# Structured Output Models

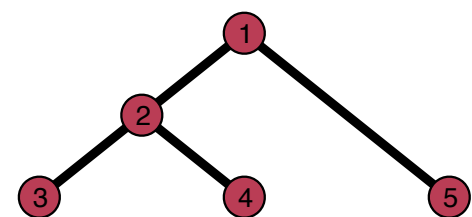
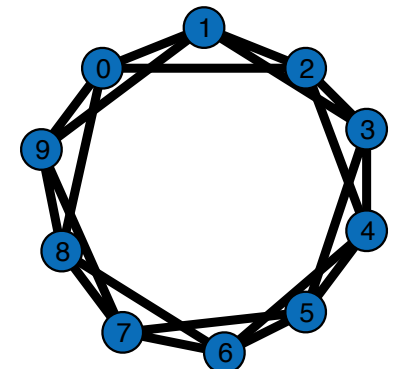
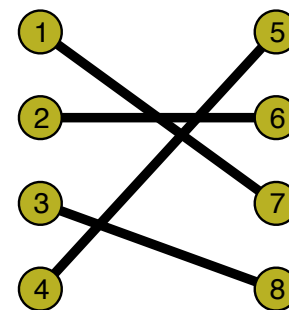
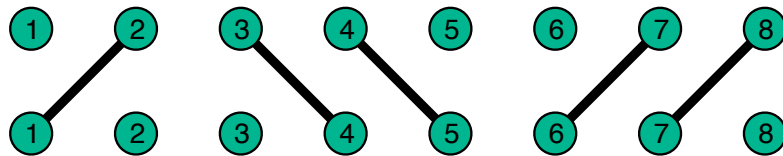
- multivariate interdependent outputs

- chains, trees, alignments, matchings, DCS

- parts  $\mathbf{X} = \{\mathbf{x}_i\}$

- labels  $\mathbf{Y} = \{y_i\} \in \mathbb{Y}$

$$\mathbb{Y} \subsetneq \{0, 1\}^N$$



# Structured Predictions

- part scores

$$s_i = \mathbf{w}^T \mathbf{x}_i$$

- structure scores

$$\sum_{i \in \text{parts}} \mathbf{w}^T \mathbf{x}_i z_i \quad \mathbf{Z} = \{z_i\} \quad \mathbf{Z} \in \mathbb{Y}$$

- predictions

$$f(\mathbf{X}) = \operatorname{argmax}_{\mathbf{Z} \in \mathbb{Y}} \sum_{i \in \text{parts}} \mathbf{w}^T \mathbf{x}_i z_i$$

- dynamic programming, combinatorial optimization, ...

# Degree-Constrained Subgraphs

- “DCS” have in-degree and out-degree constraints on each node

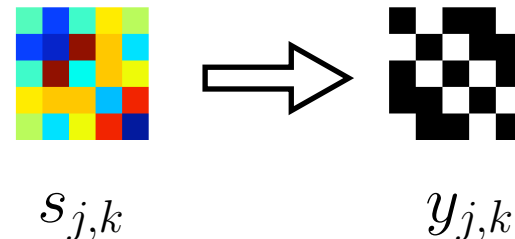
$$\left. \begin{array}{l}
 \bullet \text{ in-degree} \\
 \bullet \text{ out-degree}
 \end{array} \right\} \begin{array}{l}
 y_{j,k} \in \{0, 1\} \\
 \sum_j y_{j,k} \leq \delta_k^{in} \\
 \sum_k y_{j,k} \leq \delta_j^{out}
 \end{array} \quad \mathbf{Y} \in \mathbb{Y}^{\text{DCS}}$$

- degrees  $\delta_i^{in}$   $\delta_i^{out}$  often available, or reliably estimated

- maximum-weight DCS

- Given scores  $s_{j,k}$  finding the maximum-weight DCS is easy  $\mathcal{O}(n^3)$

$$\mathbf{Y} = \operatorname{argmax}_{\mathbf{Z} \in \mathbb{Y}^{\text{DCS}}} \sum_{j,k} s_{j,k} z_{j,k}$$



# Learning to Predict Graphs

- max-margin

- margin 
$$\gamma(\mathbf{X}, \mathbf{Y}) = \sum_{j,k} \mathbf{w}^T \mathbf{x}_{j,k} y_{j,k} - \max_{\substack{\mathbf{Z} \in \mathbb{Y} \\ \mathbf{Z} \neq \mathbf{Y}}} \sum_{j,k} \mathbf{w}^T \mathbf{x}_{j,k} z_{j,k}$$

- omitting slack variables for simplicity

$$\begin{aligned} & \min_{\mathbf{w}, \xi} \frac{1}{2} \|\mathbf{w}\|^2 && (\mathbf{X}^i, \mathbf{Y}^i) \sim P(\mathbf{X}, \mathbf{Y}) \\ \text{s.t. } & \sum_{j,k} \mathbf{w}^T \mathbf{x}_{j,k}^i y_{j,k}^i \geq \underbrace{\max_{\mathbf{Z}^i \in \mathbb{Y}} \sum_{j,k} \mathbf{w}^T \mathbf{x}_{j,k}^i z_{j,k}^i + \Delta^H(\mathbf{Y}^i, \mathbf{Z}^i)}_{\text{loss-augmented score of highest-scoring structure}}, \forall i \end{aligned}$$

- LHS: score of true structure
- RHS: loss-augmented score of highest-scoring structure (all incorrect structures are ranked “much” lower than the true structure, if and only if, the highest-scoring incorrect structure is ranked “much” lower)

- algorithms

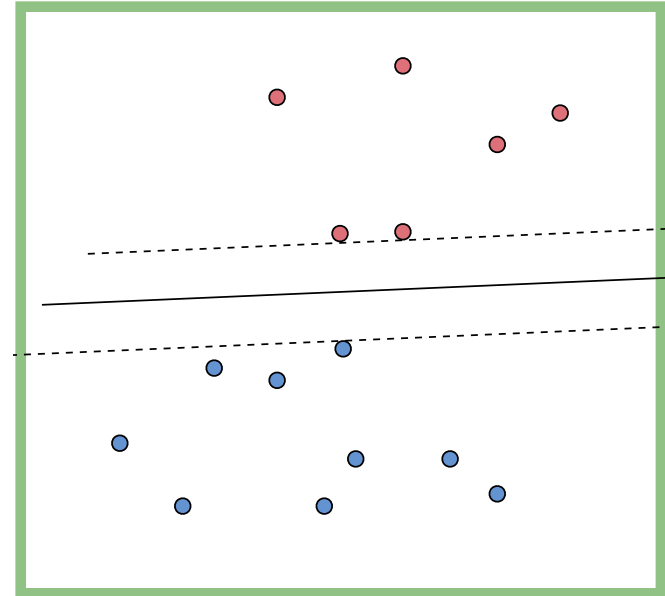
- perceptron, exponentiated gradient, SMO, cutting-planes, extra & dual extragradient, search-based optimization

# Inductive vs. Transductive

- inductive learning

- maximize margin  $\gamma(\mathbf{x}, y) = y(\mathbf{w}^T \mathbf{x} + b)$  over labeled examples

$$(\mathbf{x}^i, y^i) \sim P(\mathbf{x}, y) \longrightarrow f(\mathbf{x})$$



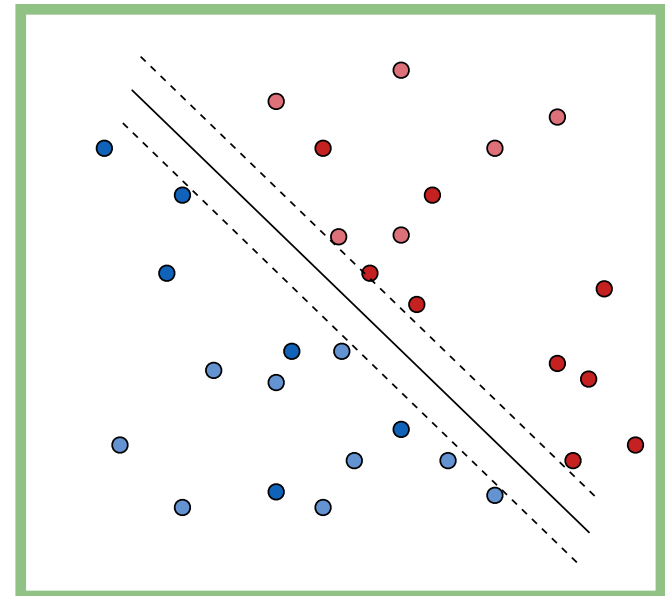
- transductive inference

- define self-reinforcing margin for unlabeled examples

$$\gamma(\mathbf{x}, *) = \max_{y=\pm 1} y(\mathbf{w}^T \mathbf{x} + b)$$

- maximize margin over ALL data

$$\begin{array}{ccc} (\mathbf{x}^i, y^i) & & (\mathbf{x}^i, y^i) \\ & \longrightarrow & \\ (\mathbf{x}^j, *) & & (\mathbf{x}^j, \hat{y}^j) \end{array}$$





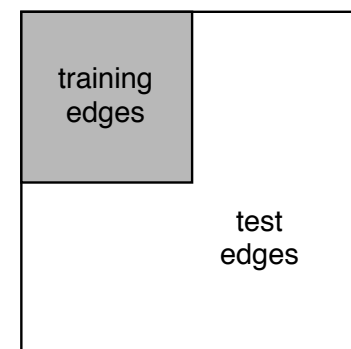
# Learning to Complete Graphs

- max-margin for partially-observed graphs

- similar to transductive inference

$$(\mathbf{X}, \mathbf{Y}_O, *) \mapsto (\mathbf{X}, \mathbf{Y}_O, \hat{\mathbf{Y}}_U)$$

$$\mathbf{Y} = \mathbf{Y}_O \cup \mathbf{Y}_U$$



- define self-reinforcing margin

$$\gamma(\mathbf{X}, \mathbf{Y}_O) = \max_{\substack{\mathbf{V} \in \mathbb{Y}^{\text{DCS}} \\ \mathbf{V}_O = \mathbf{Y}_O}} \sum_{j,k} \mathbf{w}^T \mathbf{x}_{j,k} u_{j,k} - \max_{\substack{\mathbf{Z} \in \mathbb{Y}^{\text{DCS}} \\ \mathbf{Z}_O \neq \mathbf{Y}_O}} \sum_{j,k} \mathbf{w}^T \mathbf{x}_{j,k} z_{j,k}$$

- maximize margin over ALL examples

$$\min_{\mathbf{w}, \xi} \frac{1}{2} \|\mathbf{w}\|^2$$

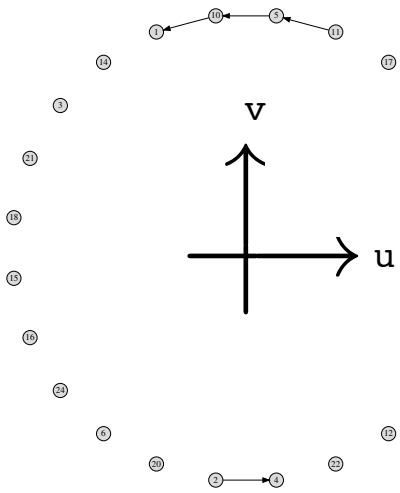
$$\text{s.t. } 0 \geq \min_{\substack{\mathbf{V} \in \mathbb{Y}^{\text{DCS}} \\ \mathbf{V}_O = \mathbf{Y}_O}} \max_{\mathbf{Z} \in \mathbb{Y}^{\text{DCS}}} \sum_{j,k} \mathbf{w}^T \mathbf{x}_{j,k} z_{j,k} + \Delta^H(\mathbf{V}, \mathbf{Z}) - \sum_{j,k} \mathbf{w}^T \mathbf{x}_{j,k} u_{j,k}$$

- algorithms

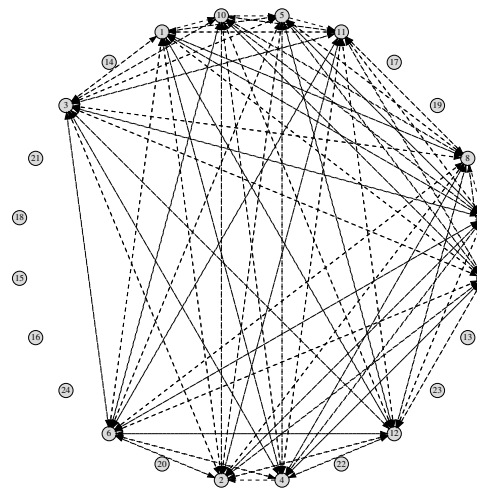
- perceptron, cutting-planes, dual extragradient (DXG)

# Circle

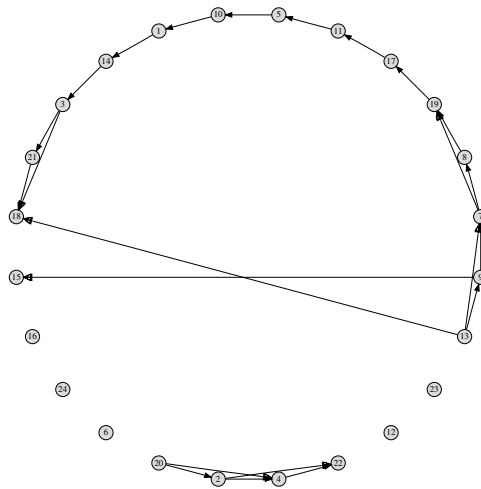
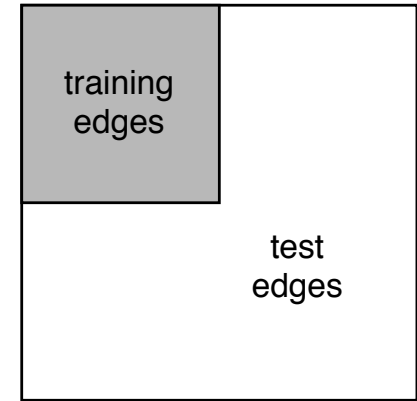
[nodes: 24, features: nodes  $\mathbf{x}_j = (u_j, v_j)$ , edges  $\mathbf{x}_{j,k} = \mathbf{x}_j \mathbf{x}_k^T$ ]



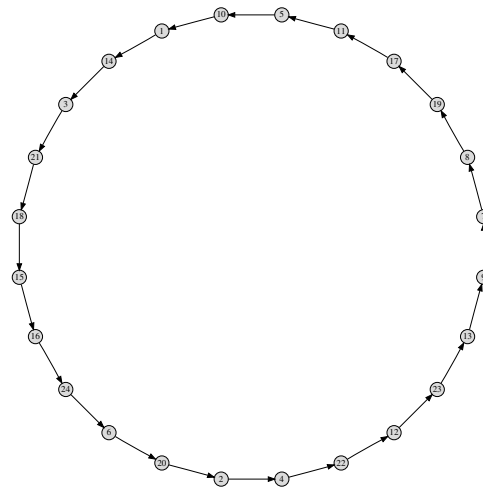
training edges



training non-edges



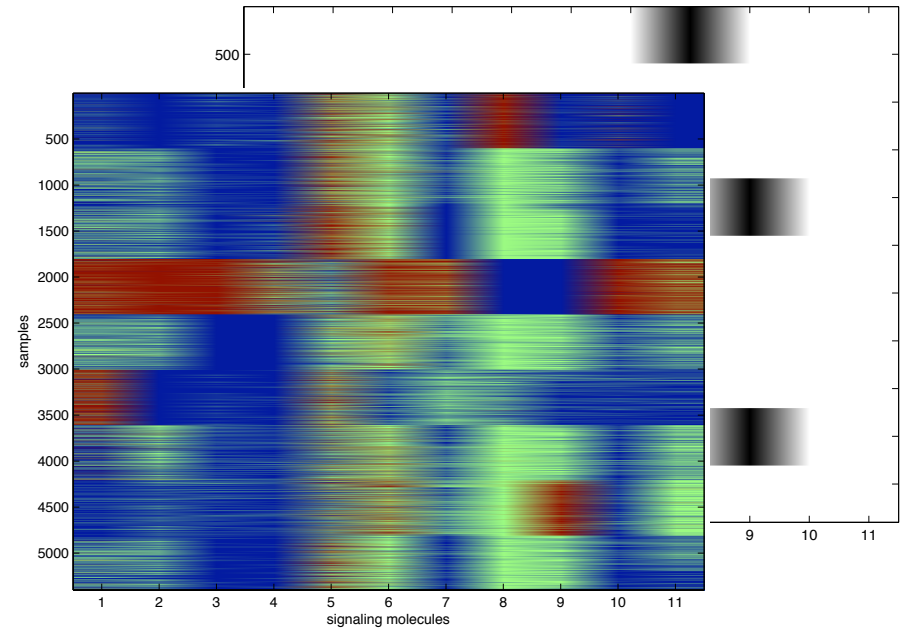
i.i.d. SVM



DXG  
w/ transduction

# Signal Transduction

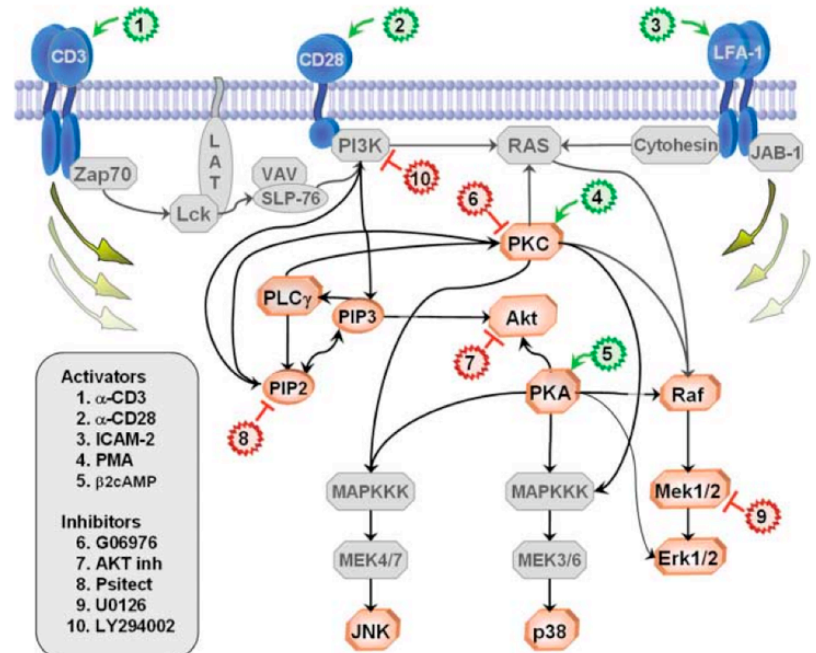
- flow cytometry expression levels  $\mathbf{x}_j$  and intervention states  $\bar{\mathbf{x}}_j$



- consensus network

$$Y_{j,k}$$

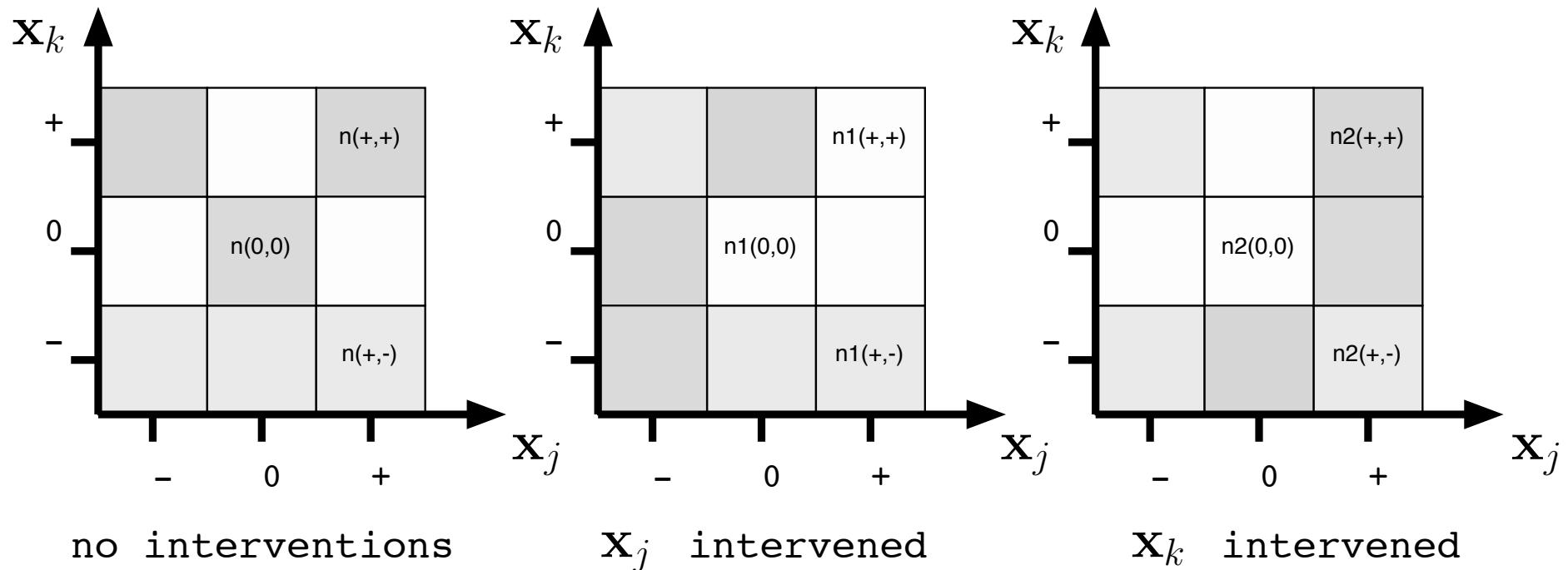
- published by Karen Sachs et al. Science 308, 523 (2005)  
[formatted by Eaton and Murphy, AISTAT, 2007]



# Signal Transduction

- edge features

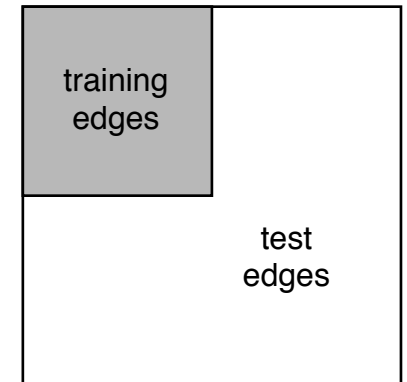
- based on expression level contingency tables



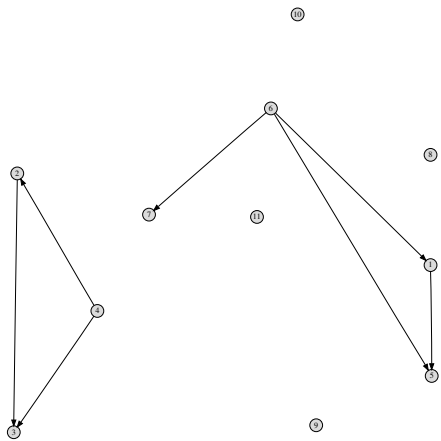
$$\mathbf{x}_{j,k} = (n_0(-, -), n_0(-, 0), \dots, n_2(+, 0), n_2(+, +))$$

# Signal Transduction

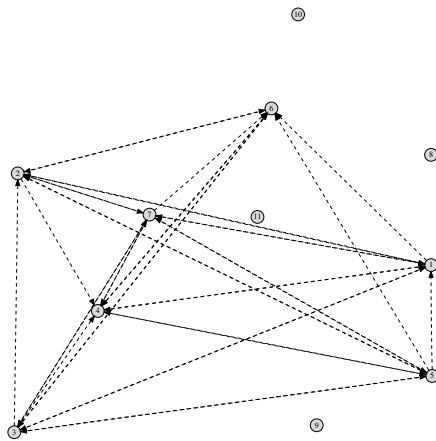
- graph completion protocol
  - blockwise partition of adjacency matrix
  - training edges labeled "on"/"off"
  - transductive max-margin training on all edges
  - predict "on"/"off" for test edges



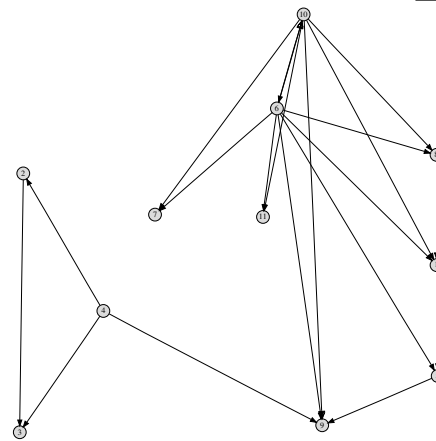
AUC 93%  
Recall 80%



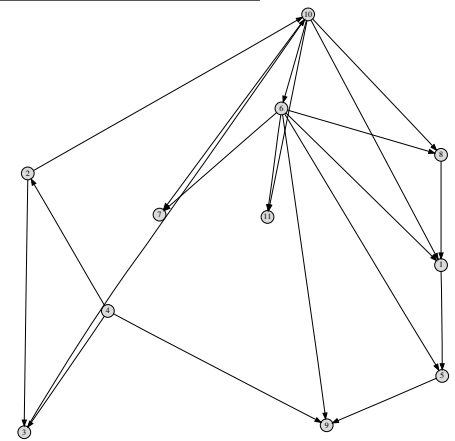
training edges



training non-edges



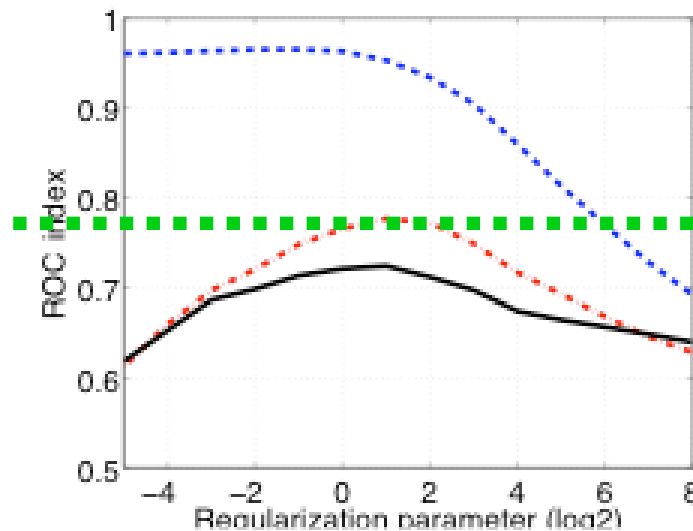
our completion



true completion

# Protein-Protein Interaction

- interactions derived from several sources
  - Yamanishi, Vert and Kanehisa, 2004
  - features: expression, Y2H, localization, phylogenetic  $x_{j,k}$
  - interactions: metabolic, physical and regulatory  $y_{j,k}$
  - 769 nodes
  - state-of-the-art performance w/o optimizing parameters



Our result  
AUC 76.2%  
Recall 54.4%

(d) Integrated kernel

Vert and Yamanishi

# Summary

- a marriage of structured-output models and degree-constrained subgraphs
  - topology and attribute-value learning
- evaluation on synthetic and real-world networks
  - task 1: complete circle network using x,y node positions
  - task 2: complete cell signaling network using flow cytometry expressions
  - task 3: complete protein-protein interactions using expression, Y2H, localization and phylogenetic features