

# Data Mining: Data

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Lecture Notes for Chapter 2---Data

# Similarity and Dissimilarity

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## ● Similarity

- Numerical measure of how alike two data objects are.
- Is higher when objects are more alike.
- Often falls in the range  $[0, 1]$

## ● Dissimilarity

- Numerical measure of how different are two data objects
- Lower when objects are more alike
- Minimum dissimilarity is often 0
- Upper limit varies

## ● Proximity refers to a similarity or dissimilarity

# Transformation (变换)

- Similarity [0, 1] scale

- 1, 2, ..., 10  $s' = (s - 1)/9$

相似度:  $s' = (s - \min_s) / (\max_s - \min_s)$

相异度:  $d' = (d - \min_d) / (\max_d - \min_d)$

- $[0, \infty]$   $d' = d / (1 + d)$

- 0, 0.5, 2, 10, 100, 1000
  - 0, 0.33, 0.67, 0.90, 0.99, 0.999

# Transformation (变换)

## ● Similarity to dissimilarity

- 0, 1, 10, 100 (dissimilarity)

- 1, 0.5, 0.09, 0.01

$$s = 1/(1 + d)$$

- 1.00, 0.37, 0.00, 0.00

$$s = e^{-d}$$

- 1.00, 0.99, 0.00, 0.00

$$s = 1 - \frac{(d - \min_d)}{(\max_d - \min_d)}$$

# Similarity/Dissimilarity for Simple Attributes

具有单个序数属性的对象（数据点）

- {poor, fair, OK, good, wonderful}

- {poor=0, fair=1, OK=2, good=3, wonderful=4}

$$d = (x - y) / (n - 1)$$

# Euclidean Distance

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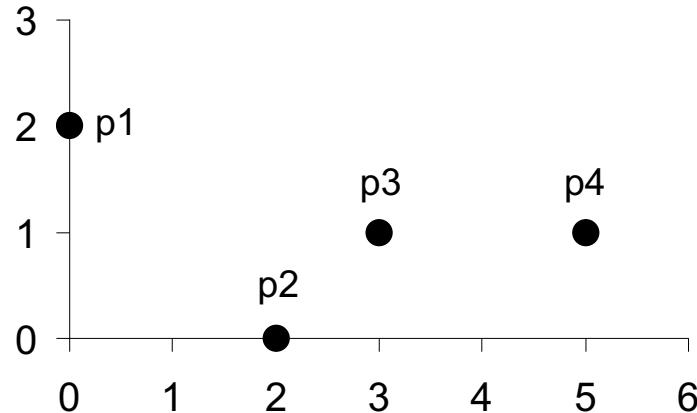
- Euclidean Distance

$$dist = \sqrt{\sum_{k=1}^n (p_k - q_k)^2}$$

Where  $n$  is the number of dimensions (attributes) and  $p_k$  and  $q_k$  are, respectively, the  $k^{\text{th}}$  attributes (components) of data objects  $p$  and  $q$ .

- Standardization is necessary, if scales differ.

# Euclidean Distance



point	x	y
p1	0	2
p2	2	0
p3	3	1
p4	5	1

	p1	p2	p3	p4
p1	0	2.828	3.162	5.099
p2	2.828	0	1.414	3.162
p3	3.162	1.414	0	2
p4	5.099	3.162	2	0

## Distance Matrix

# Minkowski Distance 闵可夫斯基距离

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- Minkowski Distance is a generalization of Euclidean Distance

$$dist = \left( \sum_{k=1}^n |p_k - q_k|^r \right)^{\frac{1}{r}}$$

Where  $r$  is a parameter,  $n$  is the number of dimensions (attributes) and  $p_k$  and  $q_k$  are, respectively, the  $k$ th attributes (components) of data objects  $p$  and  $q$ .

# Minkowski Distance: Examples

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- $r = 1$ . City block (Manhattan, taxicab,  $L_1$  norm) distance
- $r = 2$ . Euclidean distance
- $r \rightarrow \infty$ . “supremum(上确界)” ( $L_{\max}$  norm,  $L_\infty$  norm) distance
  - This is the maximum difference between any component of the vectors
- Do not confuse  $r$  with  $n$ , i.e., all these distances are defined for all numbers of dimensions.

$$dist = \lim_{r \rightarrow \infty} \left( \sum_{k=1}^n |p_k - q_k|^r \right)^{\frac{1}{r}} = \max |p_k - q_k|$$

# Minkowski Distance

point	x	y
p1	0	2
p2	2	0
p3	3	1
p4	5	1

L1	p1	p2	p3	p4
p1	0	4	4	6
p2	4	0	2	4
p3	4	2	0	2
p4	6	4	2	0

L2	p1	p2	p3	p4
p1	0	2.828	3.162	5.099
p2	2.828	0	1.414	3.162
p3	3.162	1.414	0	2
p4	5.099	3.162	2	0

$L_\infty$	p1	p2	p3	p4
p1	0	2	3	5
p2	2	0	1	3
p3	3	1	0	2
p4	5	3	2	0

## Distance Matrix

# Common Properties of a Distance

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- Distances, such as the Euclidean distance, have some well known properties.
  1.  $d(p, q) \geq 0$  for all  $p$  and  $q$  and  $d(p, q) = 0$  only if  $p = q$ . (Positive definiteness)
  2.  $d(p, q) = d(q, p)$  for all  $p$  and  $q$ . (Symmetry)
  3.  $d(p, r) \leq d(p, q) + d(q, r)$  for all points  $p$ ,  $q$ , and  $r$ . (Triangle Inequality)where  $d(p, q)$  is the distance (dissimilarity) between points (data objects),  $p$  and  $q$ .
- A distance that satisfies these properties is a **metric**

# Common Properties of a Similarity

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- Similarities, also have some well known properties.
  1.  $s(p, q) = 1$  (or maximum similarity) only if  $p = q$ .
  2.  $s(p, q) = s(q, p)$  for all  $p$  and  $q$ . (Symmetry)

where  $s(p, q)$  is the similarity between points (data objects),  $p$  and  $q$ .

# Similarity Between Binary Vectors

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- Common situation is that objects,  $p$  and  $q$ , have only binary attributes
- Compute similarities using the following quantities

$M_{01}$  = the number of attributes where  $p$  was 0 and  $q$  was 1

$M_{10}$  = the number of attributes where  $p$  was 1 and  $q$  was 0

$M_{00}$  = the number of attributes where  $p$  was 0 and  $q$  was 0

$M_{11}$  = the number of attributes where  $p$  was 1 and  $q$  was 1

- **Simple Matching and Jaccard Coefficients**

SMC = number of matches / number of attributes

$$= (M_{11} + M_{00}) / (M_{01} + M_{10} + M_{11} + M_{00})$$

$J$  = number of “11” matches / number of not-both-zero attributes values

$$= (M_{11}) / (M_{01} + M_{10} + M_{11})$$

# SMC versus Jaccard: Example

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$p = 1000000000$

$q = 0000001001$

$M_{01} = 2$  (the number of attributes where p was 0 and q was 1)

$M_{10} = 1$  (the number of attributes where p was 1 and q was 0)

$M_{00} = 7$  (the number of attributes where p was 0 and q was 0)

$M_{11} = 0$  (the number of attributes where p was 1 and q was 1)

$$SMC = (M_{11} + M_{00}) / (M_{01} + M_{10} + M_{11} + M_{00}) = (0+7) / (2+1+0+7) = 0.7$$

$$J = (M_{11}) / (M_{01} + M_{10} + M_{11}) = 0 / (2 + 1 + 0) = 0$$

# Cosine Similarity

- If  $d_1$  and  $d_2$  are two document vectors, then

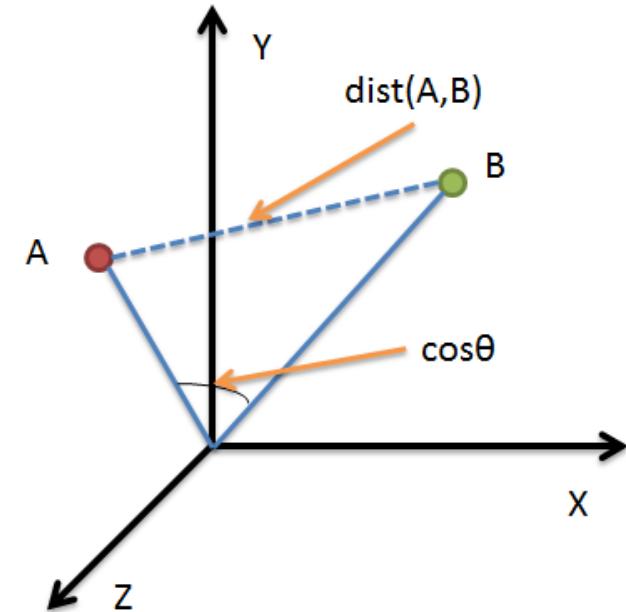
$$\cos(d_1, d_2) = (d_1 \bullet d_2) / \|d_1\| \|d_2\| ,$$

where  $\bullet$  indicates vector dot product  
and  $\|d\|$  is the length of vector  $d$ .

- Example:

$$d_1 = 3 \ 2 \ 0 \ 5 \ 0 \ 0 \ 0 \ 2 \ 0 \ 0$$

$$d_2 = 1 \ 0 \ 0 \ 0 \ 0 \ 0 \ 0 \ 1 \ 0 \ 2$$



$$d_1 \bullet d_2 = 3*1 + 2*0 + 0*0 + 5*0 + 0*0 + 0*0 + 0*0 + 2*1 + 0*0 + 0*2 = 5$$

$$\|d_1\| = (3^2 + 2^2 + 0^2 + 5^2 + 0^2 + 0^2 + 0^2 + 2^2 + 0^2 + 0^2)^{0.5} = (42)^{0.5} = 6.481$$

$$\|d_2\| = (1^2 + 0^2 + 0^2 + 0^2 + 0^2 + 0^2 + 0^2 + 1^2 + 0^2 + 2^2)^{0.5} = (6)^{0.5} = 2.245$$

$$\cos(d_1, d_2) = 0.3150$$

# Correlation

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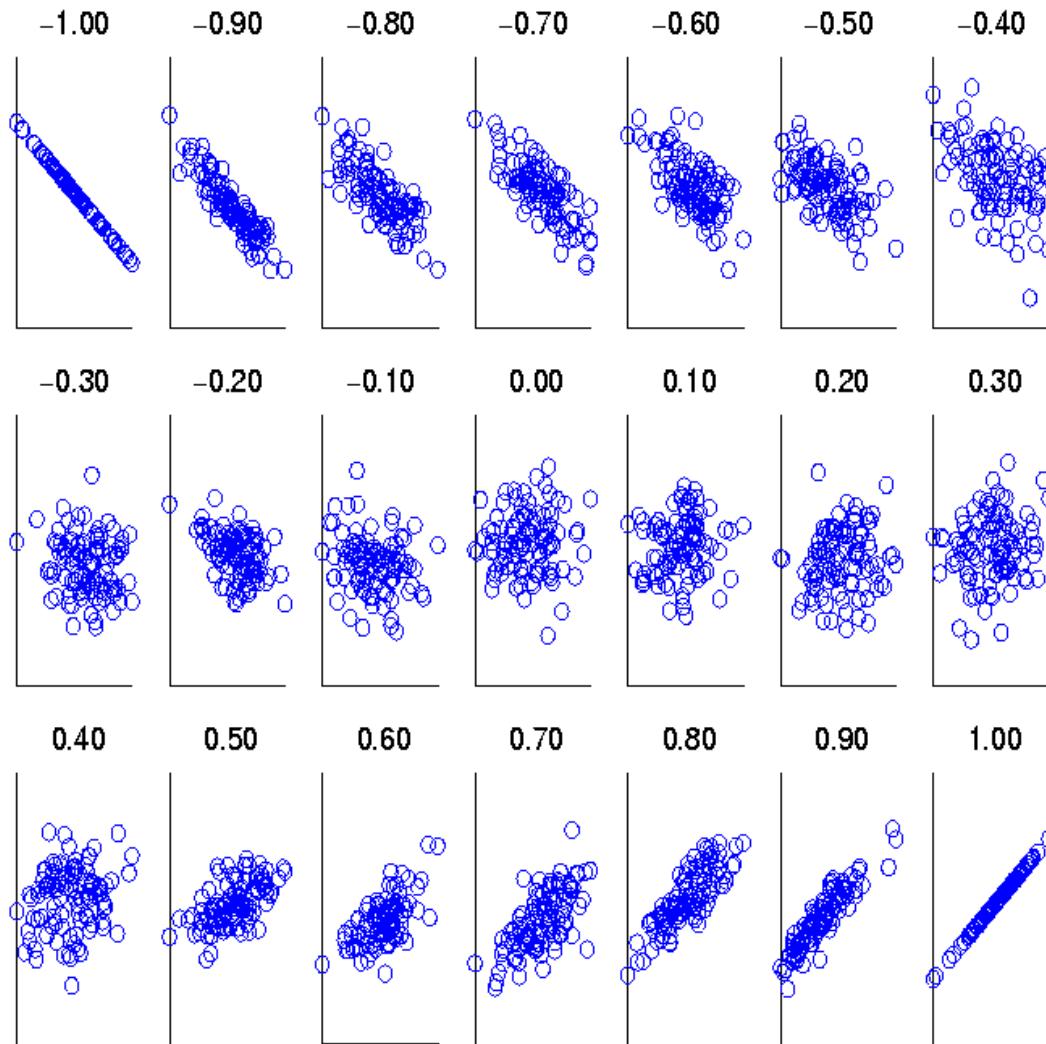
- Correlation measures the linear relationship between objects
- To compute correlation, we standardize data objects,  $p$  and  $q$ , and then take their dot product

$$p'_k = (p_k - \mathbf{mean}(p)) / \mathbf{std}(p)$$

$$q'_k = (q_k - \mathbf{mean}(q)) / \mathbf{std}(q)$$

$$\mathbf{correlation}(p, q) = p' \bullet q'$$

# Visually Evaluating Correlation



**Scatter plots  
showing the  
similarity from  
-1 to 1.**

# General Approach for Combining Similarities

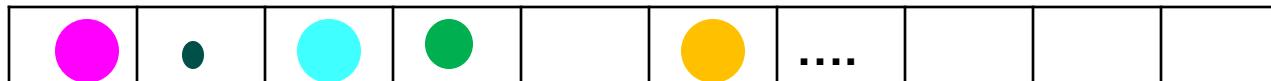
- Sometimes attributes are of many different types, but an overall similarity is needed.

1. For the  $k^{th}$  attribute, compute a similarity,  $s_k$ , in the range  $[0, 1]$ .
2. Define an indicator variable,  $\delta_k$ , for the  $k_{th}$  attribute as follows:

$$\delta_k = \begin{cases} 0 & \text{if the } k^{th} \text{ attribute is a binary asymmetric attribute and both objects have} \\ & \text{a value of 0, or if one of the objects has a missing values for the } k^{th} \text{ attribute} \\ 1 & \text{otherwise} \end{cases}$$

3. Compute the overall similarity between the two objects using the following formula:

$$\text{similarity}(p, q) = \frac{\sum_{k=1}^n \delta_k s_k}{\sum_{k=1}^n \delta_k}$$



# Using Weights to Combine Similarities

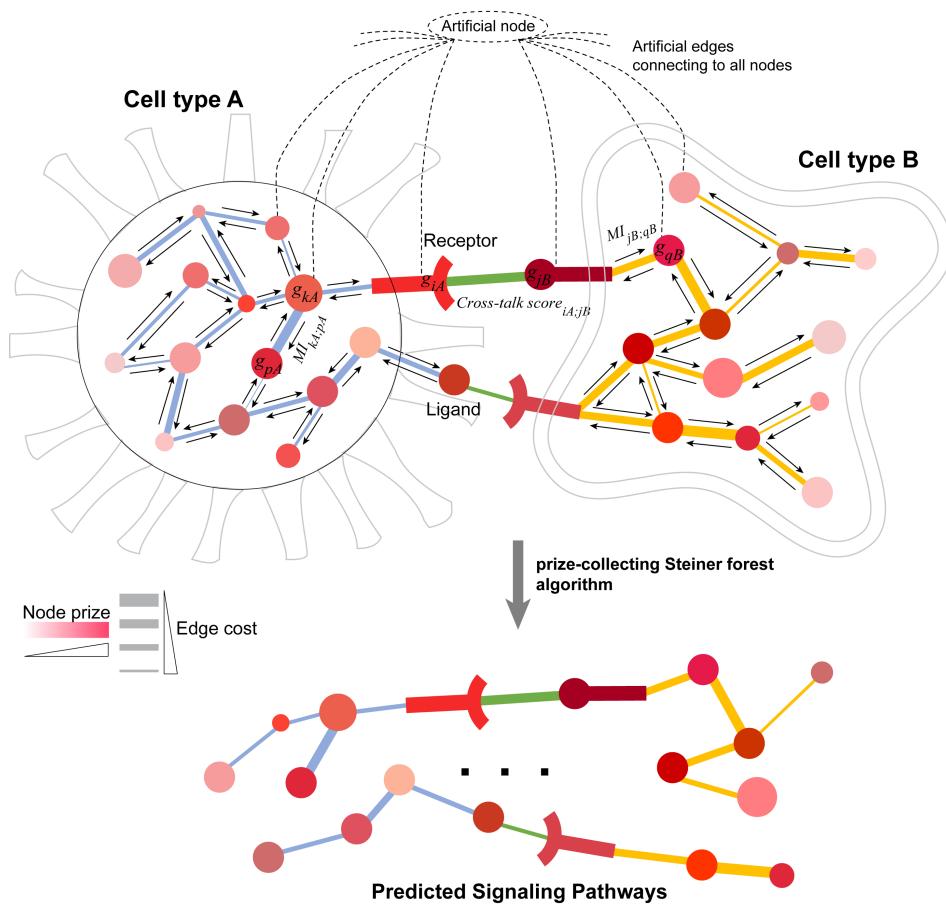
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- May not want to treat all attributes the same.
  - Use weights  $w_k$  which are between 0 and 1 and sum to 1.

$$similarity(p, q) = \frac{\sum_{k=1}^n w_k \delta_k s_k}{\sum_{k=1}^n \delta_k}$$

$$distance(p, q) = \left( \sum_{k=1}^n w_k |p_k - q_k|^r \right)^{1/r}$$

# 相似性应用: CytoTalk算法解析细胞通讯



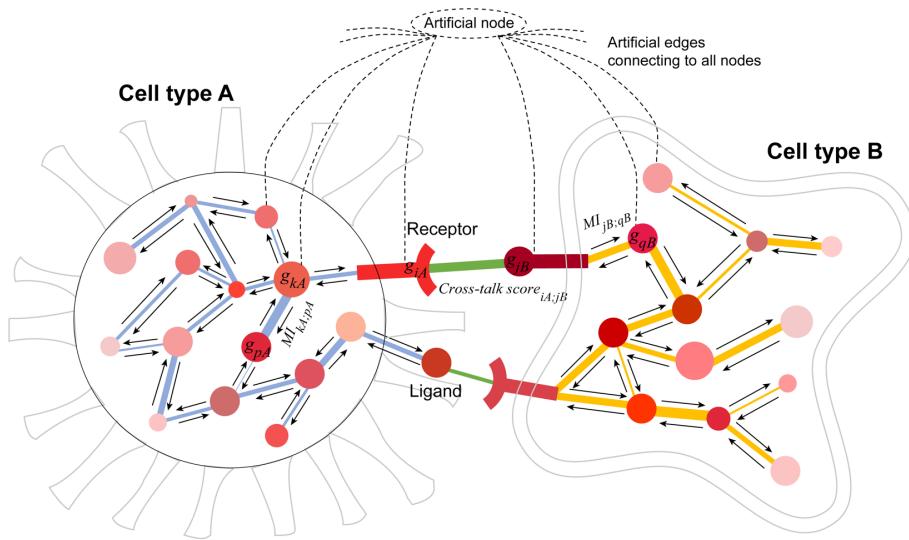
**Mathematical modeling:**  
Formulate the prediction of cell type-specific signaling network as a prize-collecting Steiner forest problem.

**Key question:**  
How to define node prize and edge cost in the input cross-cell-type gene network?

Hu Y, Gao L, et al. *Science Advances*, 2021

# 相似性应用: CytoTalk算法解析细胞通讯

Node prize:



$$Prize_{iA} = Relevance_{iA} \times PEM_{iA}$$

(1) Cell-type-specificity (preferential expression measure)

$$PEM_{iA} = \log_{10}(Expr_{iA} / e_{iA})$$

$$e_{iA} = \sum_{m=1}^M Expr_{im} \cdot \frac{S_{*A}}{\sum_{m=1}^M S_{*m}}$$

(2) Closeness to the ligand or receptor genes in the network

$$\mathbf{Relevance}_A^t = \alpha \mathbf{W}' \mathbf{Relevance}^{t-1} + (1 - \alpha) \mathbf{Relevance}^0$$

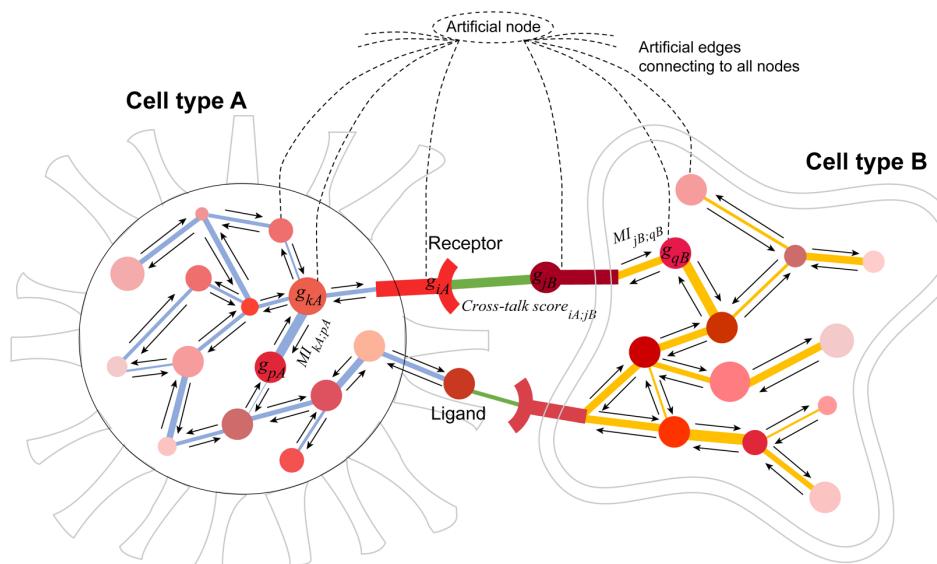
# 相似性应用：CytoTalk算法解析细胞通讯

## Edge cost:

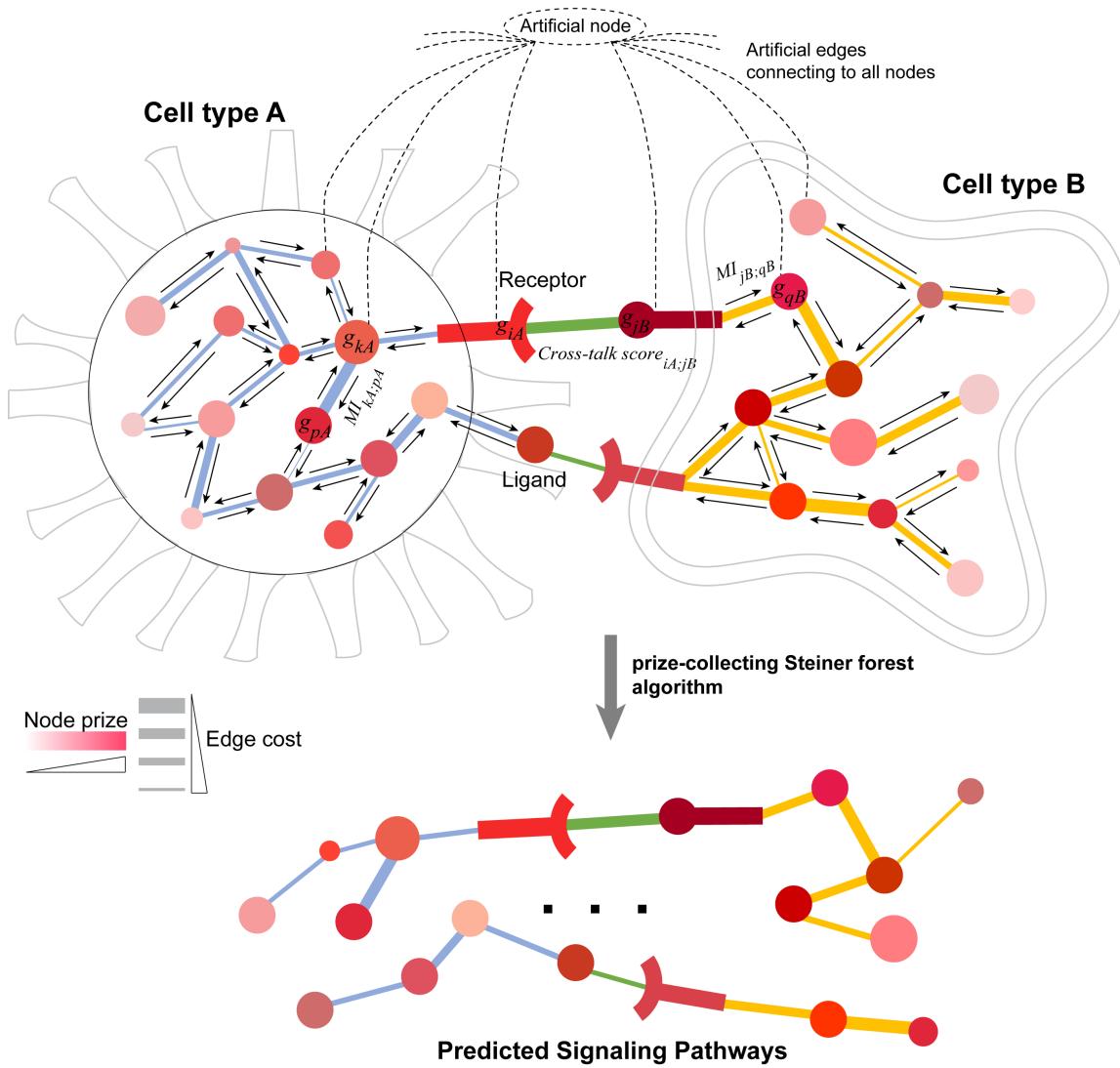
$$\text{Crosstalk score}_{iA,jB} = \text{Norm}(\text{Expression score}_{iA,jB}) \times \text{Norm}(\text{Non-self-talk score}_{iA,jB})$$

$$\text{Expression score}_{iA,jB} = (PEM_{iA} + PEM_{jB})/2$$

$$\text{Non-self-talk score}_{iA,jB} = [(-\log_{10} \frac{MI_{iA;jA}}{\min\{H_{iA}, H_{jA}\}}) + (-\log_{10} \frac{MI_{iB;jB}}{\min\{H_{iB}, H_{jB}\}})]/2$$



# 相似性应用：CytoTalk算法解析细胞通讯



Objective function:

$$\min_F c(F) + \beta \times p(\bar{F}) + \omega \times k$$



感谢各位同学！  
下次课再见！