

# Subgraph Matching via Optimal Transport

**Theory, Algorithms, Applications**

Wenxin Pan

*2023 Master Thesis*

# Outline

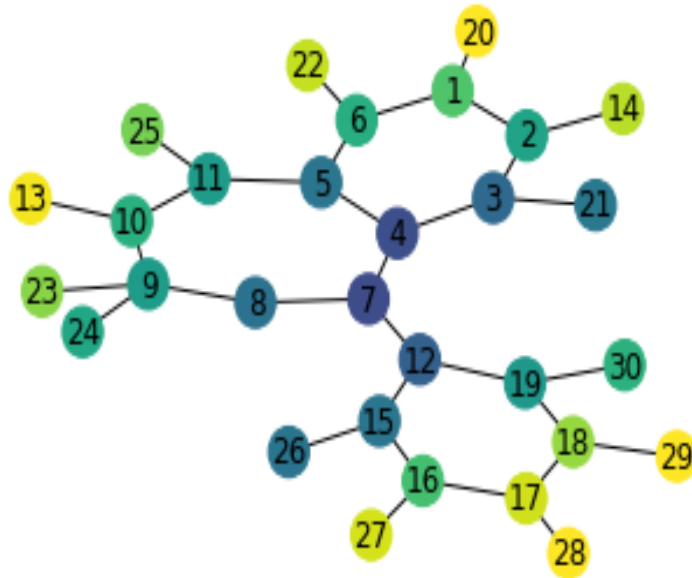
- **Introduction** – Problem formulation & motivation
- **Theory** – Optimal Transport on graphs
- **Algorithms** – Subgraph matching algorithms
- **Applications** – for biological datasets

# Outline

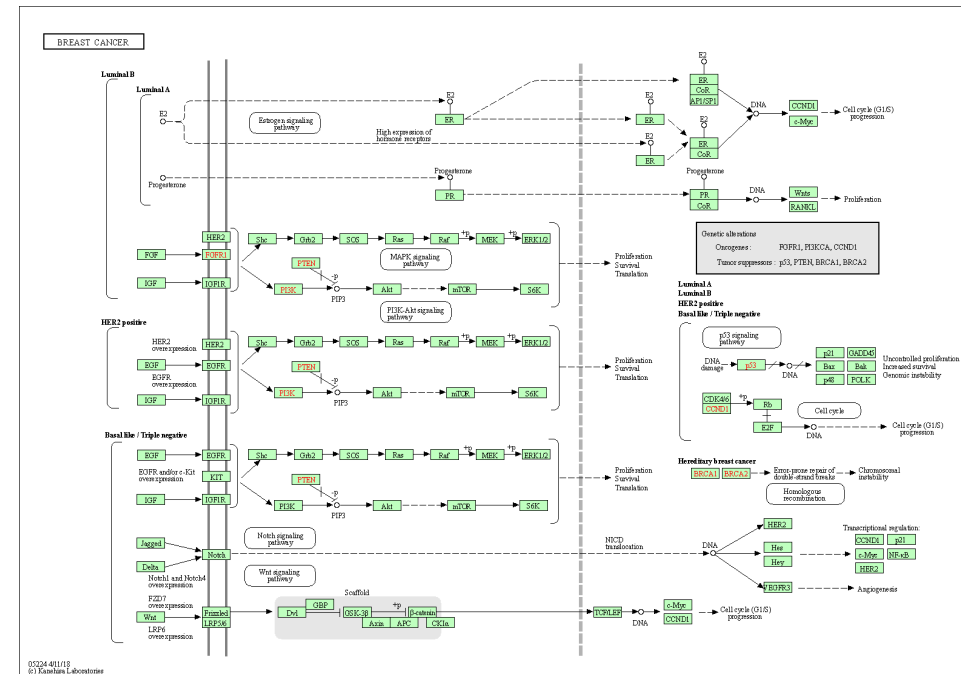
- **Introduction** – Problem formulation & motivation
- **Theory** – Optimal Transport on graphs
- **Algorithms** – Subgraph matching algorithms
- **Applications** – for biological datasets

# Introduction

- Find predefined patterns within an object



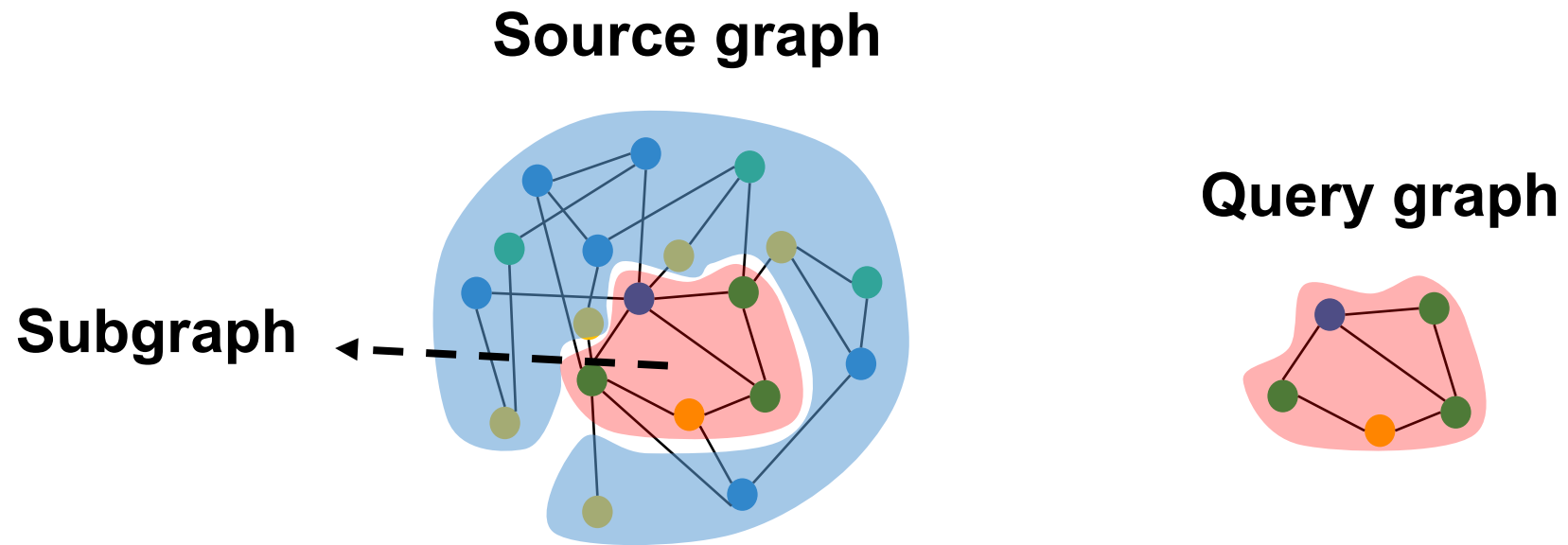
Find **rings** in a chemical compound



Find **signaling pathways** in a biomedical pathway

# Introduction

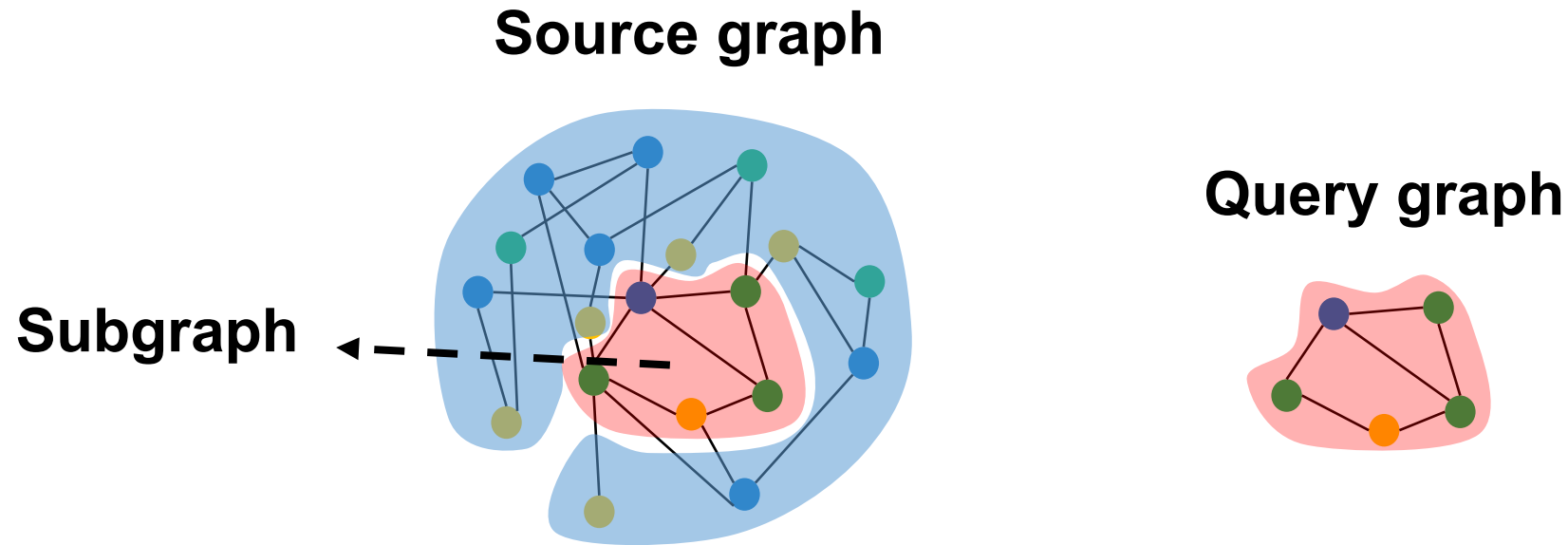
- **Problem formulation**
  - Subgraph matching



# Introduction

- **Goals**

- Is there any *subgraph* same/similar to the *query graph* in the *source graph* ?
- Where is the *subgraph* ?
- How similar are the *subgraph* and *query graph* ?





# Classic Formulation: Quadratic Assignment Problems (QAPs)

With adjacency matrices  $A^s$  and  $A^q$ ,  
find a *binary* mapping matrix between source and query

- For two graphs of *equal size*:  
that each column and row has exactly one “1”

$$\mathbf{X} \in \{0, 1\}^{n \times n}, \mathbf{X}\mathbf{1}_n = \mathbf{1}_n, \mathbf{X}^\top \mathbf{1}_n = \mathbf{1}_n$$

- For two graphs of *different sizes*:  
that each column (or row) has exactly one “1”

$$\mathbf{X} \in \{0, 1\}^{n \times m}, \mathbf{X}\mathbf{1}_m \leq \mathbf{1}_n, \mathbf{X}^\top \mathbf{1}_n = \mathbf{1}_m$$

■				
	■			
		■		
			■	
				■

## Koopmans-Beckmann's QAP

$$\min_{\mathbf{X} \in \mathcal{X}} -\langle \mathbf{A}^s, \mathbf{X} \mathbf{A}^q \mathbf{X}^\top \rangle_F$$

## Lawler's QAP

$$\min_{\mathbf{X} \in \mathcal{X}} \text{vec}(\mathbf{X})^\top \mathbf{Q} \text{vec}(\mathbf{X}),$$
$$\mathbf{Q}_{n(j-1)+i, n(j'-1)+i'} = d^S(\mathbf{A}_{i,i'}^s, \mathbf{A}_{j,j'}^q)$$

## Remarks

### The Quadratic Assignment Problems (QAPs)

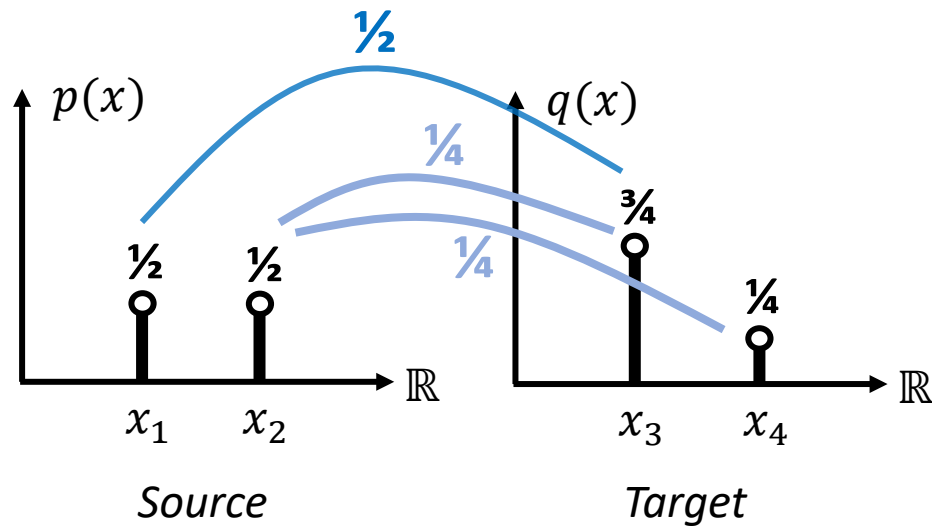
- are complex *combinatorial* optimization problems
- do not consider *node features*

# Outline

- **Introduction** – Problem formulation & motivation
- **Theory** – Optimal Transport on graphs
- **Algorithms** – Subgraph matching algorithms
- **Applications** – for biological datasets

# Optimal Transport

- How to transport the *mass* from one probability distribution to another with minimum cost ?



Cost Matrix  $M$ :

$ x_3 - x_1 $	$ x_4 - x_1 $
$ x_3 - x_2 $	$ x_4 - x_2 $

A possible Transport Matrix  $T$ :

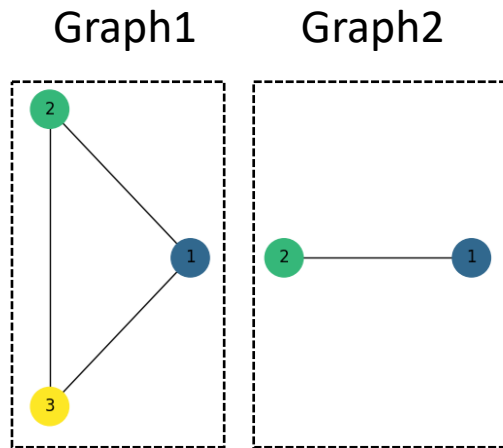
$\frac{1}{2}$	$0$
$\frac{1}{4}$	$\frac{1}{4}$

- The total minimum cost is defined as the **Wasserstein distance**

$$\mathcal{W}(p(x), q(x)) = \min_T \left\{ \langle T, M \rangle_F \stackrel{\text{def.}}{=} \sum_{i,j} T_{i,j} M_{i,j} \right\}$$

# Optimal Transport on Graphs

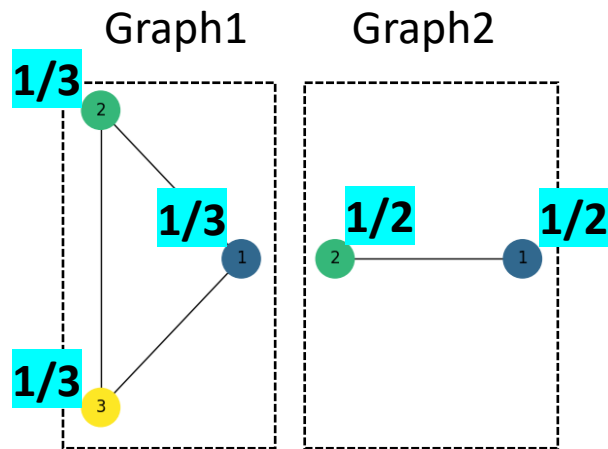
- How to transport the *information* from one graph to another with minimum cost ?
- *Information: Feature & Structure*



$$T = \begin{bmatrix} ? \\ ? \\ ? \end{bmatrix}_{3 \times 2}$$

# Optimal Transport on Graphs

- How to transport the *information* from one graph to another with minimum cost ?
- *Information: Feature & Structure*
- *Feature*



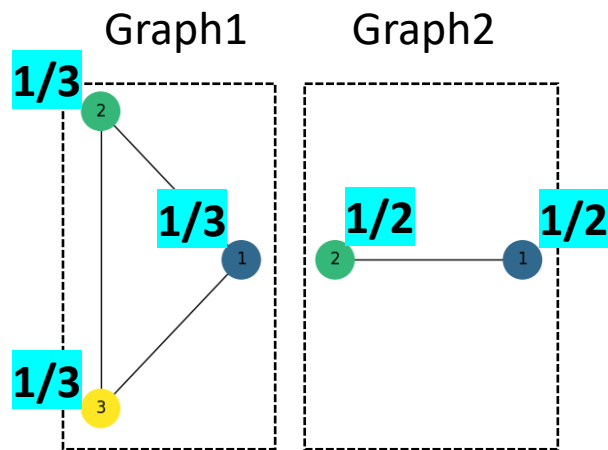
$$M = \begin{matrix} & \begin{matrix} g \\ b \\ \gamma \end{matrix} & \begin{bmatrix} 0 & 1 \\ 1 & 0 \\ 1 & 1 \end{bmatrix} \end{matrix}$$

$$T^* = \begin{bmatrix} 1/3 & 0 \\ 0 & 1/3 \\ 1/6 & 1/6 \end{bmatrix}$$

**Wasserstein distance:**  $\mathcal{W}(p, q, M) = \min_T \langle T, M \rangle_F$

# Optimal Transport on Graphs

- How to transport the *information* from one graph to another with minimum cost ?
- *Information: Feature & Structure*
- *Structure*



$$C^s = \begin{bmatrix} 0 & 1 & 1 \\ 1 & 0 & 1 \\ 1 & 1 & 0 \end{bmatrix} \quad C^t = \begin{bmatrix} 0 & 1 \\ 1 & 0 \end{bmatrix} \quad L_{i,i',j,j'} = \mathcal{L}(C_{i,i'}^s, C_{j,j'}^t)$$

6 optimal transport matrices:  $T^* = \begin{bmatrix} 1/3 & 0 \\ 0 & 1/3 \\ 1/6 & 1/6 \end{bmatrix} \quad T^* = \begin{bmatrix} 0 & 1/3 \\ 1/3 & 0 \\ 1/6 & 1/6 \end{bmatrix} \dots$

**Gromov-Wasserstein distance:**  $\mathcal{GW}(p, q, C^s, C^t) = \min_T \sum_{i,i',j,j'} \mathcal{L}(C_{i,i'}^s, C_{j,j'}^t) T_{i,j} T_{i',j'}$

## Wasserstein Distance

$$\mathcal{W}(\mathbf{p}, \mathbf{q}, \mathbf{M}) = \min_{\mathbf{T}} \left\{ \langle \mathbf{T}, \mathbf{M} \rangle_F \stackrel{\text{def.}}{=} \sum_{i,j} \mathbf{T}_{i,j} \mathbf{M}_{i,j} \right\}$$

## Gromov-Wasserstein Distance

$$\mathcal{GW}(\mathbf{p}, \mathbf{q}, \mathbf{C}^s, \mathbf{C}^t) = \min_{\mathbf{T}} \left\{ \langle \mathbf{L} \otimes \mathbf{T}, \mathbf{T} \rangle_F \stackrel{\text{def.}}{=} \sum_{i,i',j,j'} \mathcal{L}(\mathbf{C}_{i,i'}^s, \mathbf{C}_{j,j'}^t) \mathbf{T}_{i,j} \mathbf{T}_{i',j'} \right\}$$

with  $(\mathbf{L} \otimes \mathbf{T})_{i,j} \stackrel{\text{def.}}{=} \sum_{i',j'} \mathbf{L}_{i,i',j,j'} \mathbf{T}_{i',j'}$

## Fused Gromov-Wasserstein Distance

$$\mathcal{FGW}(\mathbf{p}, \mathbf{q}, \mathbf{M}, \mathbf{C}^s, \mathbf{C}^t) = \min_{\mathbf{T}} (1 - \alpha) \langle \mathbf{T}, \mathbf{M} \rangle_F + \alpha \langle \mathbf{L} \otimes \mathbf{T}, \mathbf{T} \rangle_F$$

## FGW distance

### Fused Gromov-Wasserstein Distance

$$\mathcal{FGW}(\mathbf{p}, \mathbf{q}, M, \mathbf{C}^s, \mathbf{C}^t) = \min_{\mathbf{T}} (1 - \alpha) \langle \mathbf{T}, M \rangle_F + \alpha \langle \mathbf{L} \otimes \mathbf{T}, \mathbf{T} \rangle_F$$

$$\text{with } (\mathbf{L} \otimes \mathbf{T})_{i,j} \stackrel{\text{def.}}{=} \sum_{i',j'} L_{i,i',j,j'} \mathbf{T}_{i',j'}$$

$$\text{Constraints} \quad \text{s.t. } \mathbf{T} \mathbf{1}_m = \mathbf{p}; \mathbf{T}^\top \mathbf{1}_n = \mathbf{q}; \mathbf{T} \in \mathbb{R}_+^{n \times m}$$

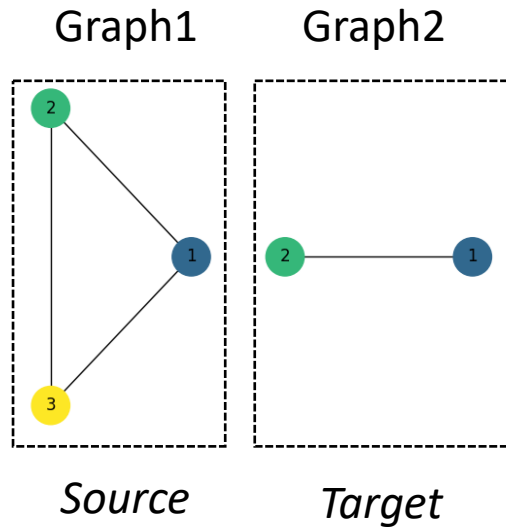
- Probability vectors  $\mathbf{p}$  and  $\mathbf{q}$  indicate *weights/importance* on nodes

# Remarks

## The FGW distance

- defines a *distance/metric* between two labeled graphs
- is a *generalization* of classic Quadratic Assignment Problems (QAPs)
- is naturally a *continuous* problem
- naturally allows graphs of *different sizes*

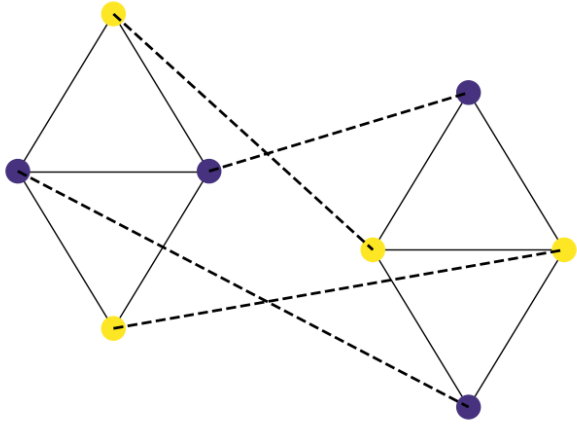
# Examples



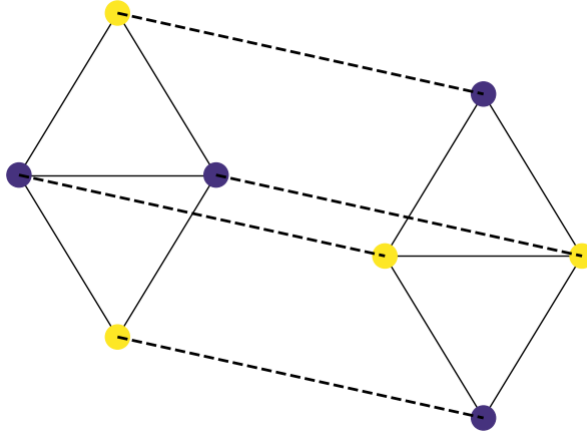
$$\mathbf{T}^* = \begin{bmatrix} 1/3 & 0 \\ 0 & 1/3 \\ 1/6 & 1/6 \end{bmatrix}$$

$$\mathcal{FGW}(\mathbf{p}, \mathbf{q}, M, \mathbf{C}^s, \mathbf{C}^t) = 0.306$$

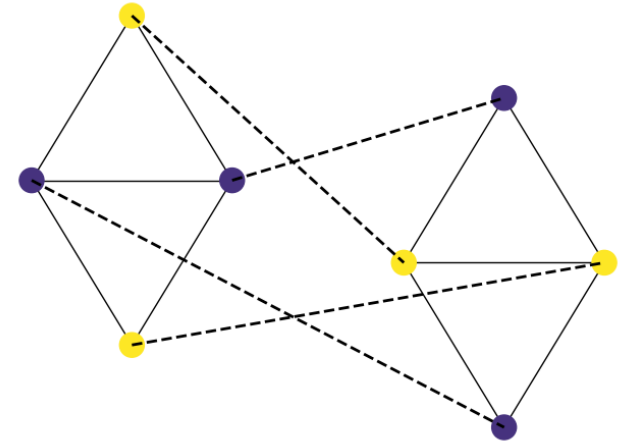
# Examples



Wasserstein distance = 0



GW distance = 0

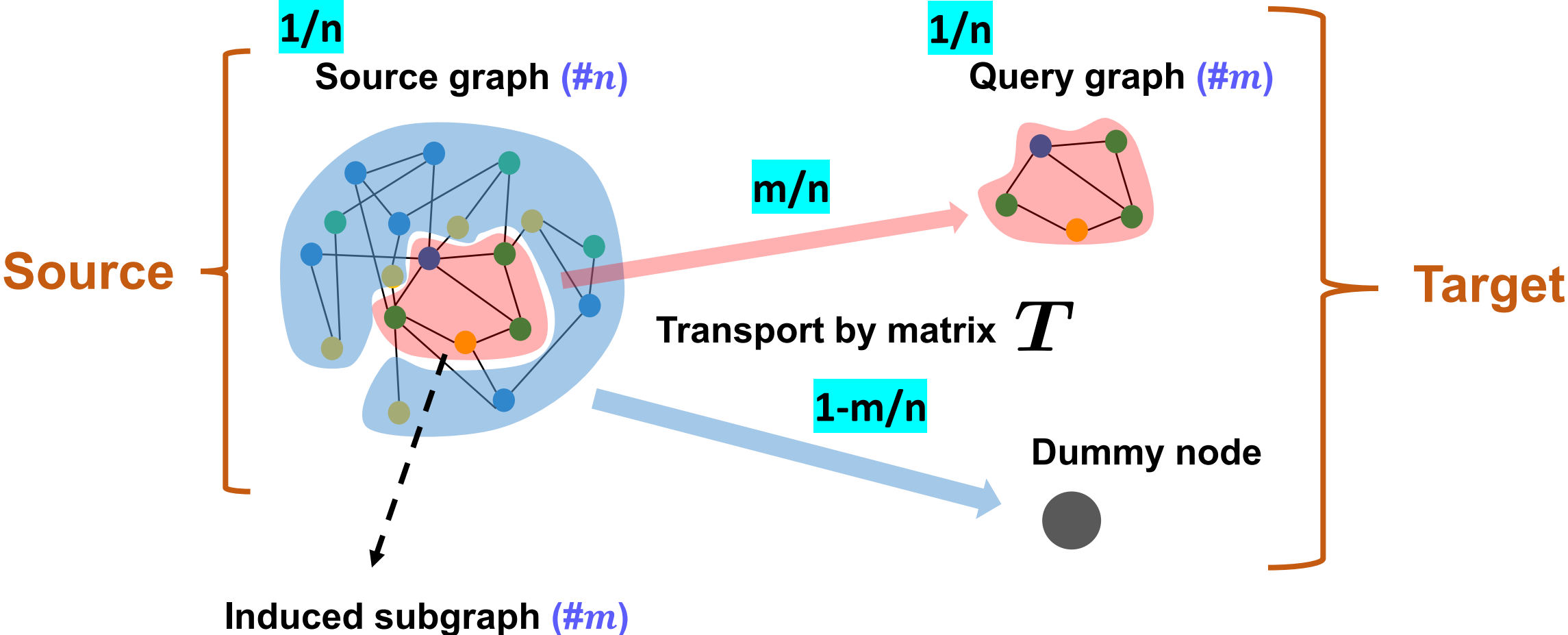


FGW distance = 0.125 ( $\alpha = 0.5$ )

# Outline

- **Introduction** – Problem formulation & motivation
- **Theory** – Optimal Transport on graphs
- **Algorithms** – Subgraph matching algorithms
- **Applications** – for biological datasets

# Subgraph Matching – Framework 1



# Subgraph Matching – Framework 1

$$\mathcal{FGW}(\mathbf{p}, \mathbf{q}, \mathbf{M}, \mathbf{C}^s, \mathbf{C}^t) = \min_{\mathbf{T}} (1 - \alpha) \langle \mathbf{T}, \mathbf{M} \rangle_F + \alpha \langle \mathbf{L} \otimes \mathbf{T}, \mathbf{T} \rangle_F$$

- mass of *source graph* are transported to
  - query graph with **original costs**
  - dummy node **"for free"**

$$M_{i,j} = \begin{cases} d(a_i, b_j), & \text{for } j = 1, \dots, m \\ 0, & \text{for } j = m + 1. \end{cases}$$

$$L_{i,i',j,j'} = \begin{cases} \mathcal{L}(\mathbf{C}_{i,i'}^s, \mathbf{C}_{j,j'}^t), & \text{for } j = 1, \dots, m \text{ and } j' = 1, \dots, m \\ 0, & \text{for } j = m + 1 \text{ or } j' = m + 1 \end{cases}$$

- Minimum value of FGW distance is still 0.

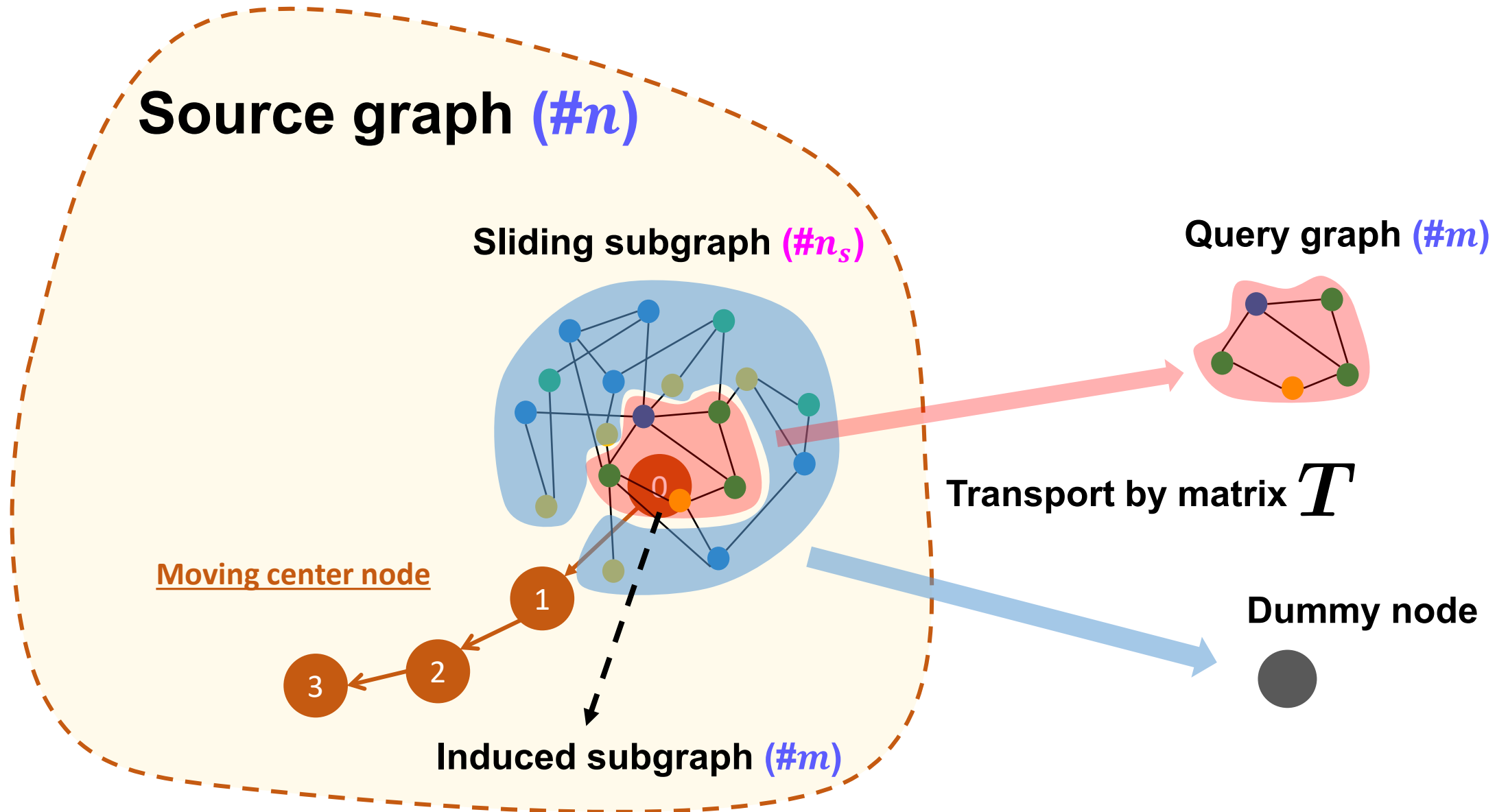
## Issues raised in large graphs

$$\mathcal{FGW}(\mathbf{p}, \mathbf{q}, \mathbf{M}, \mathbf{C}^s, \mathbf{C}^t) = \min_{\mathbf{T}} (1 - \alpha) \langle \mathbf{T}, \mathbf{M} \rangle_F + \alpha \langle \mathbf{L} \otimes \mathbf{T}, \mathbf{T} \rangle_F$$

- **Large Computation** of the tensor-matrix product  $\mathbf{L} \otimes \mathbf{T}$
- **Non-convexity** due to the product  $\mathbf{T}_{i,j} \mathbf{T}_{i',j'}$ 
  - larger graphs create more local minima

Potential solution: “shrink” the graph that is directly used in the FGW distance

# Subgraph Matching – Framework 2



## Subgraph Matching – Framework 2

**For node in Source graph:**

**Create a sliding subgraph**

Framework 1

**End**

# Outline

- **Introduction** – Problem formulation & motivation
- **Theory** – Optimal Transport on graphs
- **Algorithms** – Subgraph matching algorithms
- **Applications** – for biological datasets

## Application - example

### **Frequent subgraph matching**

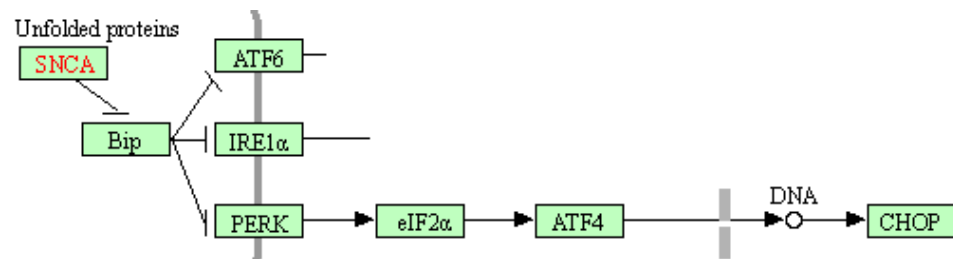
- Detect the similarities within a set of graphs

## Application - example

Detect signaling pathways within Neurodegenerative disease pathways

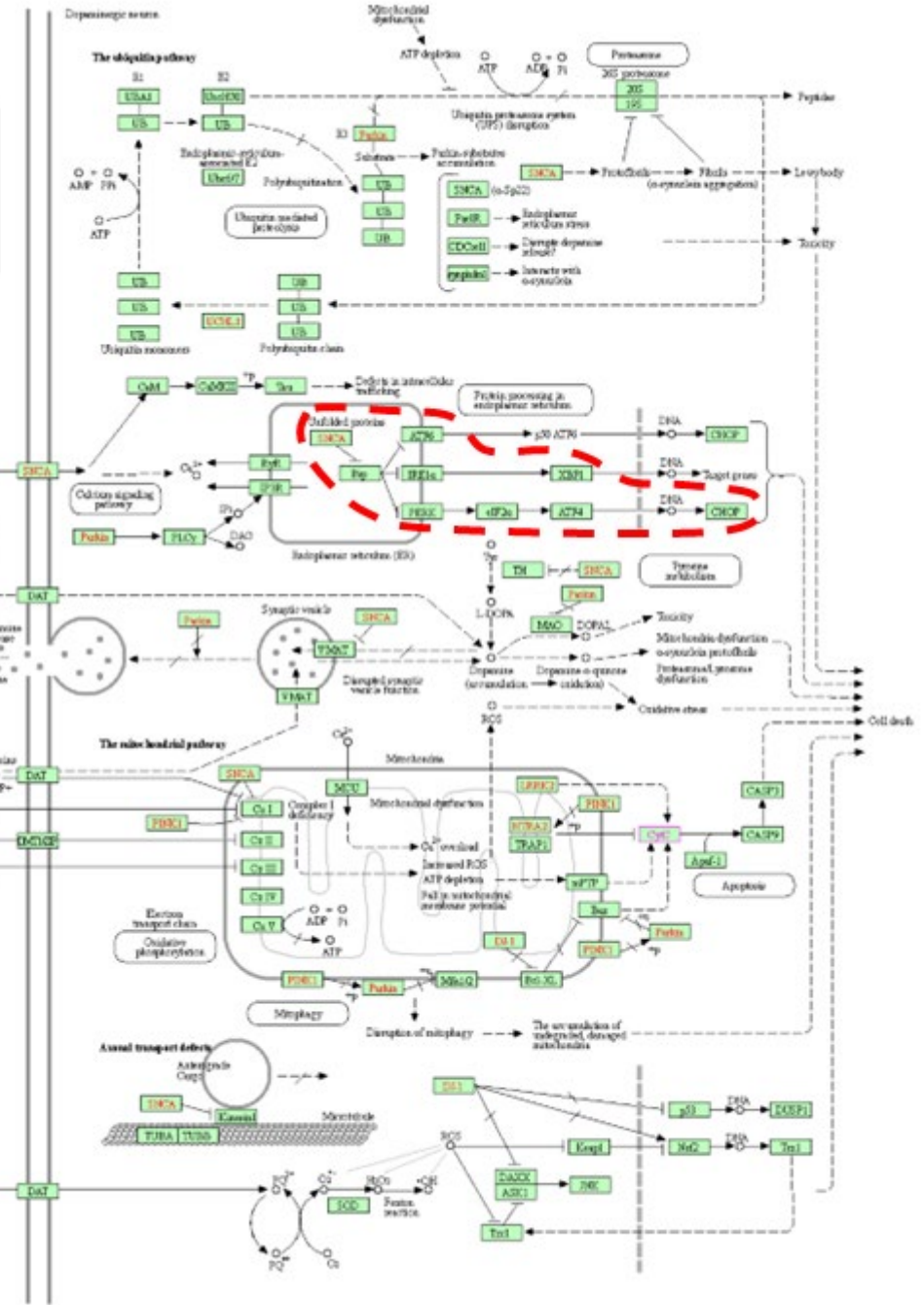
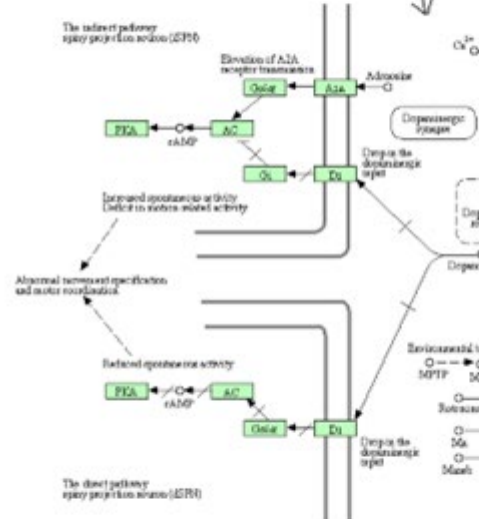
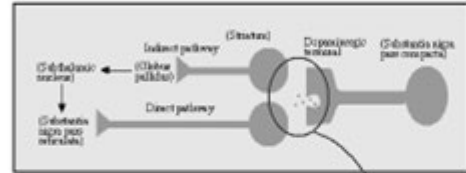
- Parkinson
- Alzheimer

Query graph: unfolded protein response (UPR)



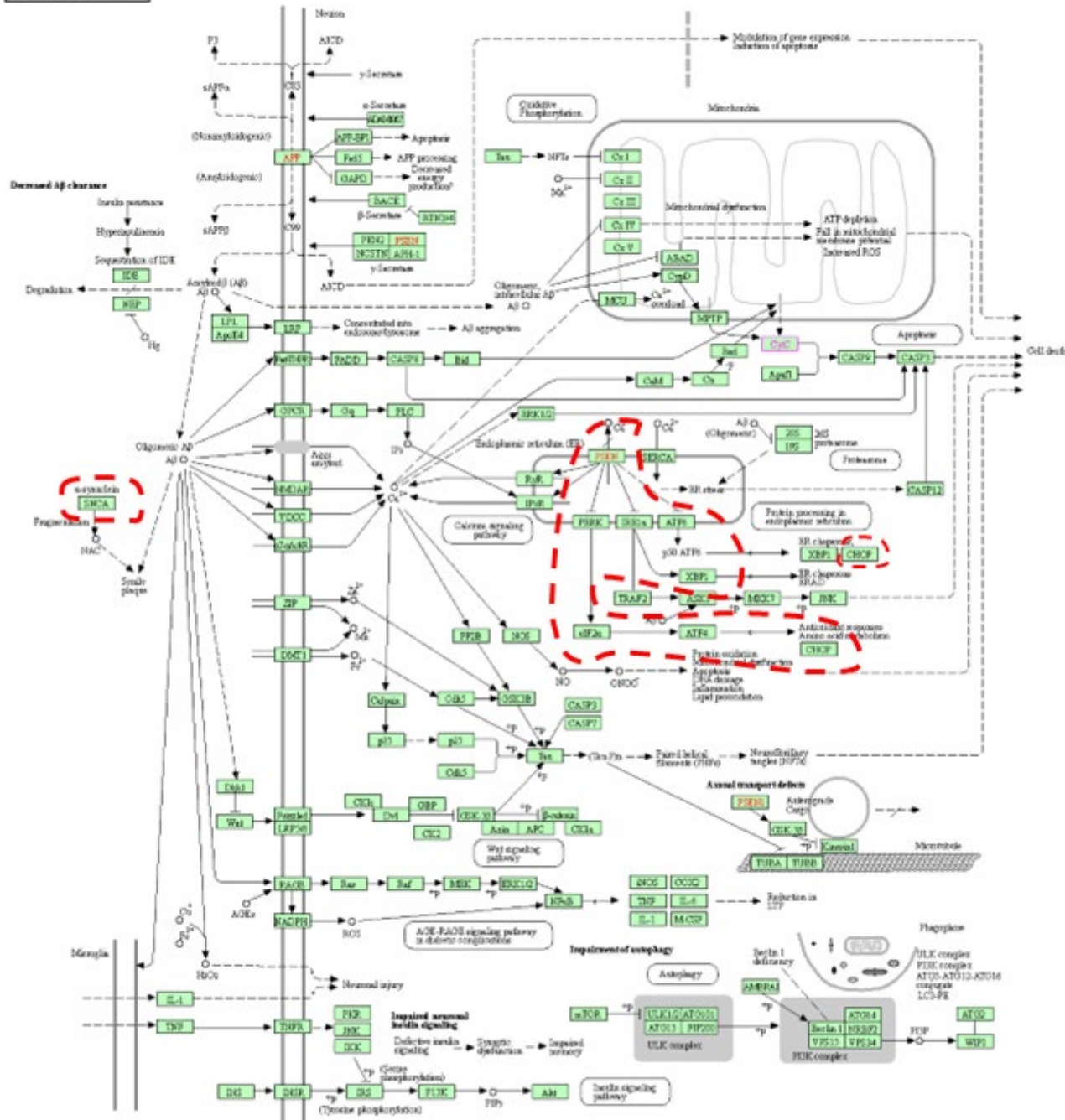
# Parkinson

PARKINSON DISEASE



# Alzheimer

ALZHEIMER DISEASE



# Conclusion

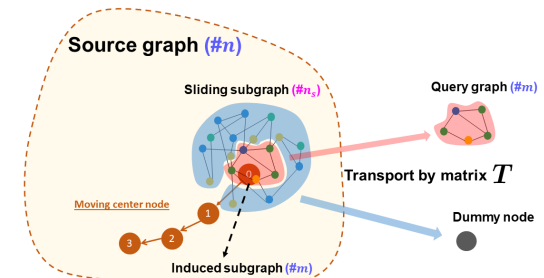
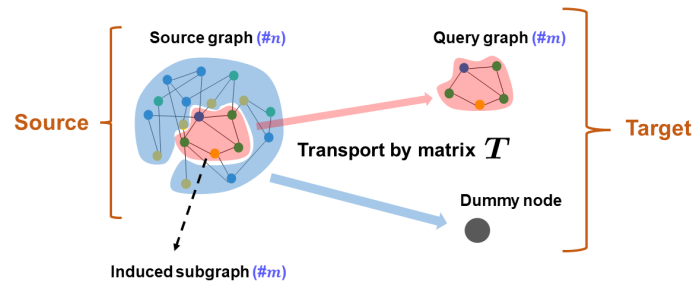
Theory

Algorithms

Applications

## Fused Gromov-Wasserstein Distance

$$\mathcal{FGW}(p, q, M, C^s, C^t) = \min_{\mathbf{T}} (1 - \alpha) \langle \mathbf{T}, \mathbf{M} \rangle_F + \alpha \langle \mathbf{L} \otimes \mathbf{T}, \mathbf{T} \rangle_F$$



Frequent subgraph matching

**Thank you!**