

Course

# Introduction to Graph Theory



# Motivation

## Graph theory

- Study mathematical structure and relationship

## Applications

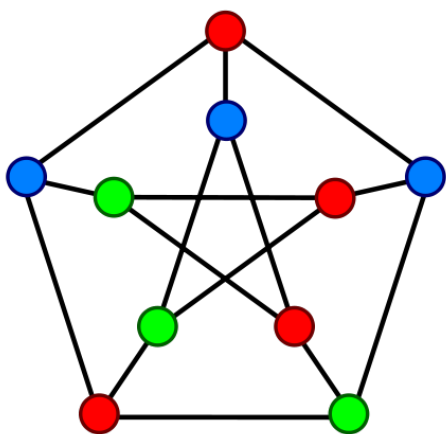
- Linguistic, computer, chemistry, biology, network

Identify the problem  
Re-use existing algorithms

NP-hard problems

*nondeterministic polynomial time*

- Traveling salesman
- Color labelling



P problems

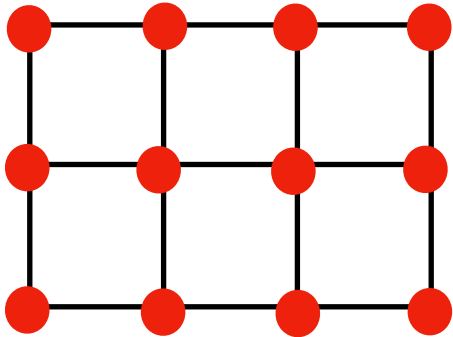
- Voronoi
- Minimum span tree
- Shortest path
- Assignment
- Max-flow / min-cut

Graph Neural Network

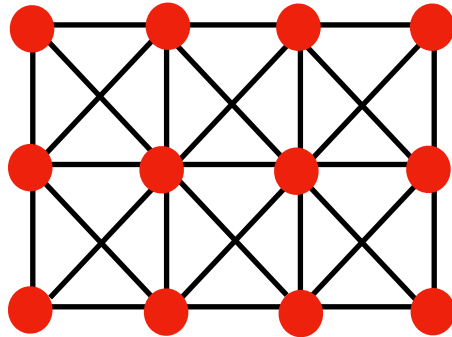
## Graphs in Bioimage Analysis

### Data modeling

- Representation of image at pixel level

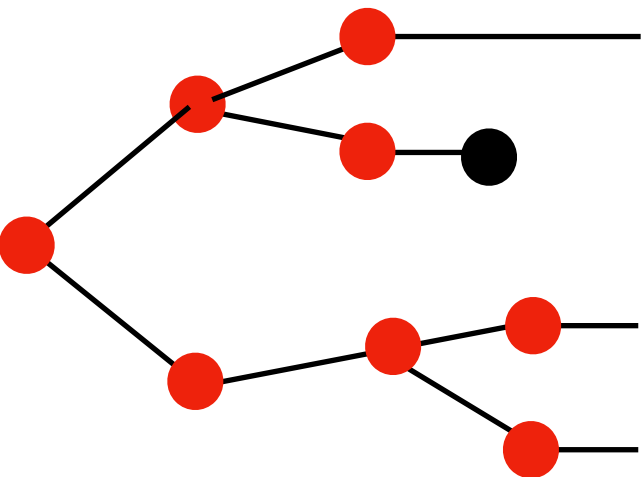


2D: 4-connected  
3D: 6-connected

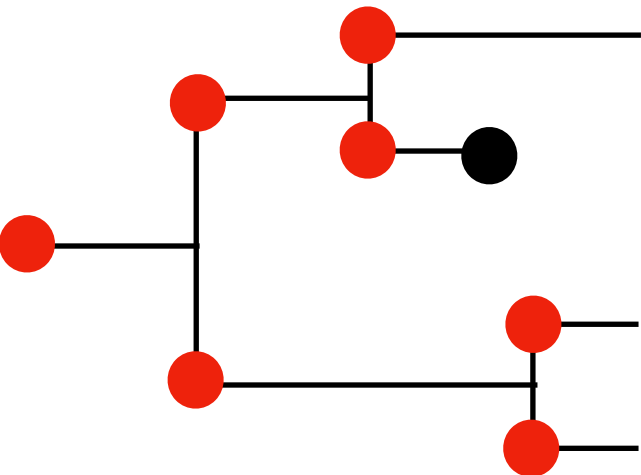


2D: 8-connected  
3D: 26-connected

- Dimension: scale, space, time
- Object: particles, cell
- Spatial organization:
- Vascular structure: tree
- Relation: cell lineage, tracking, ...



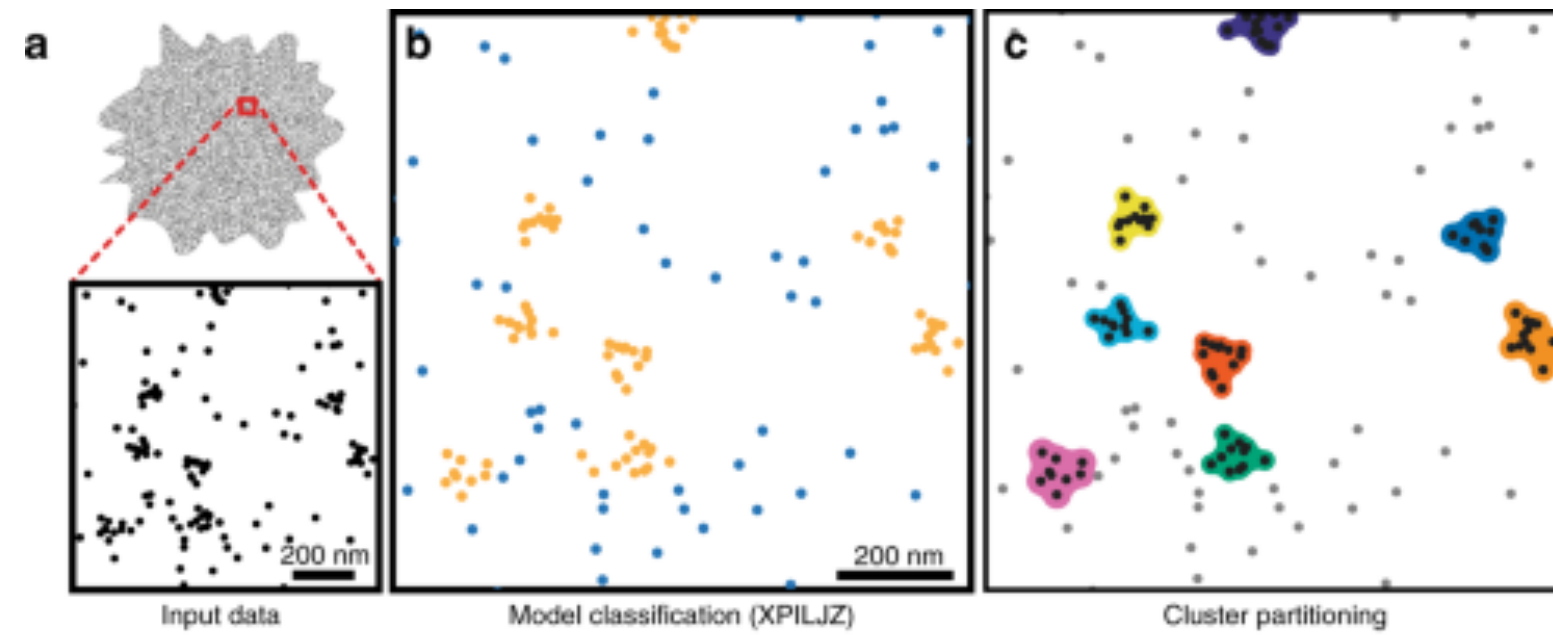
- Mosaic of images



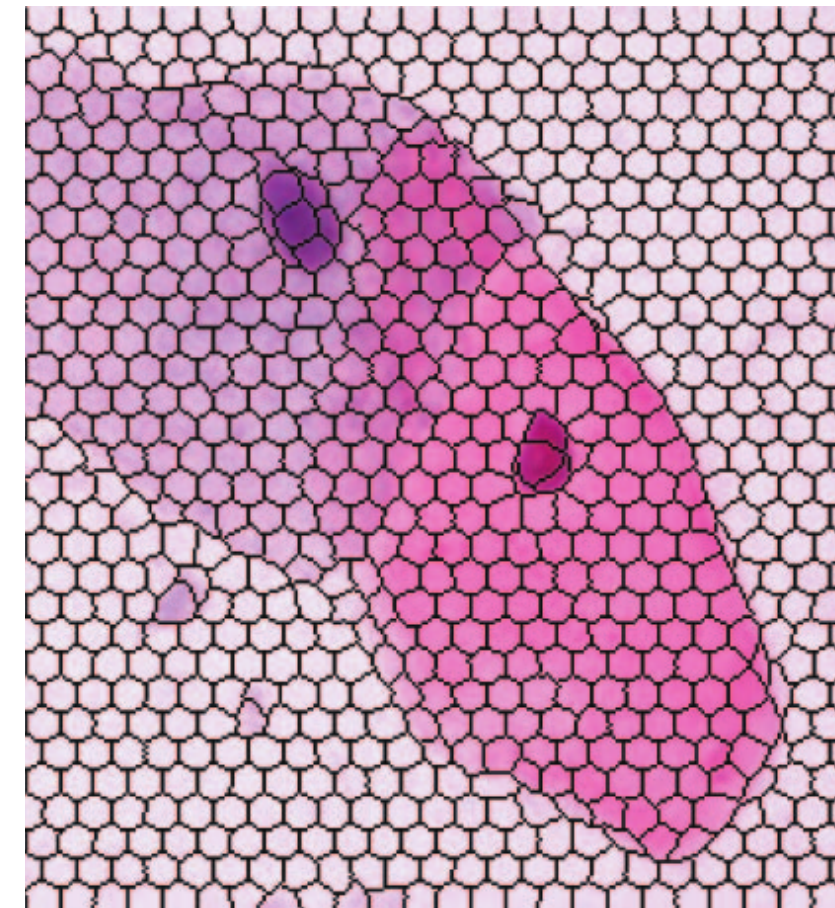




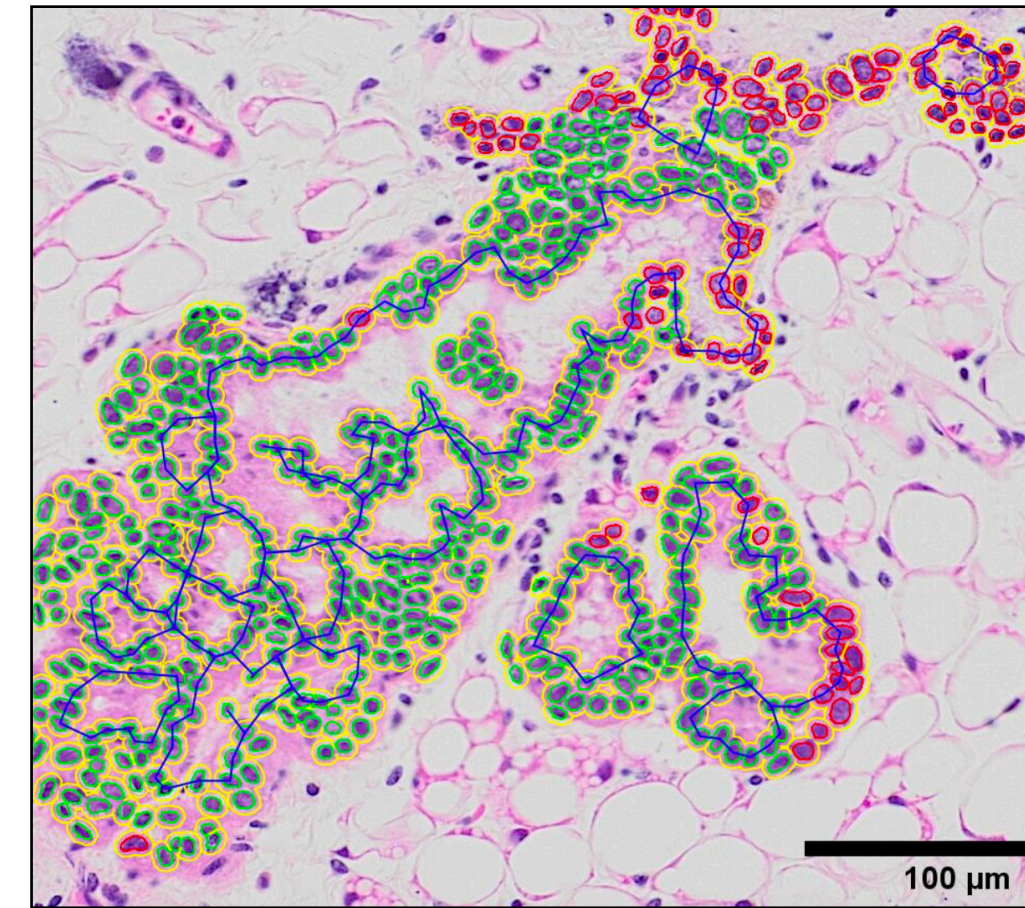
# Graph Modelling



Molecules ➡ Objects



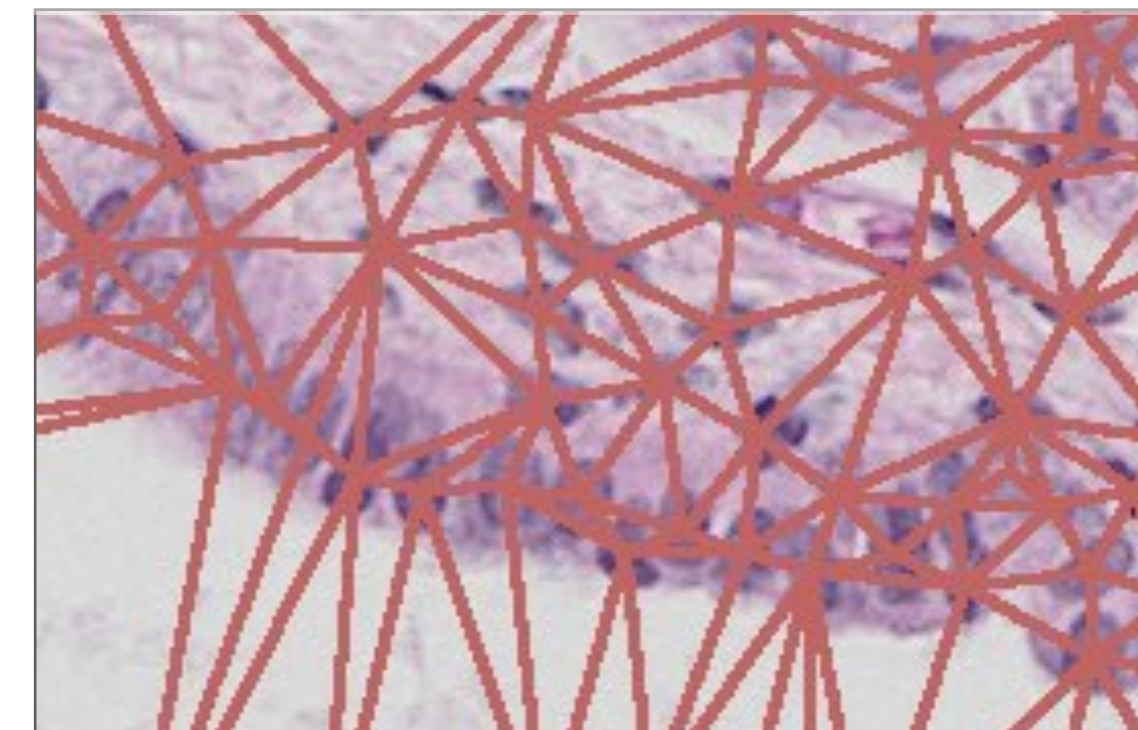
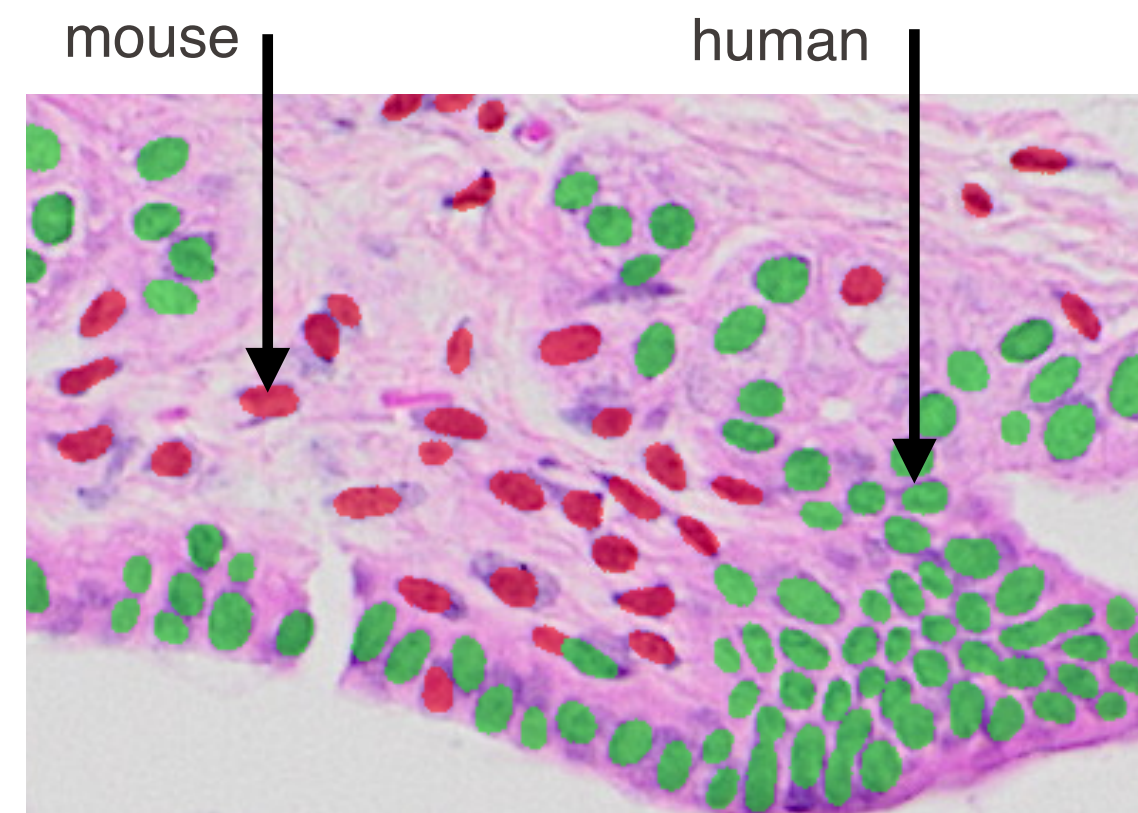
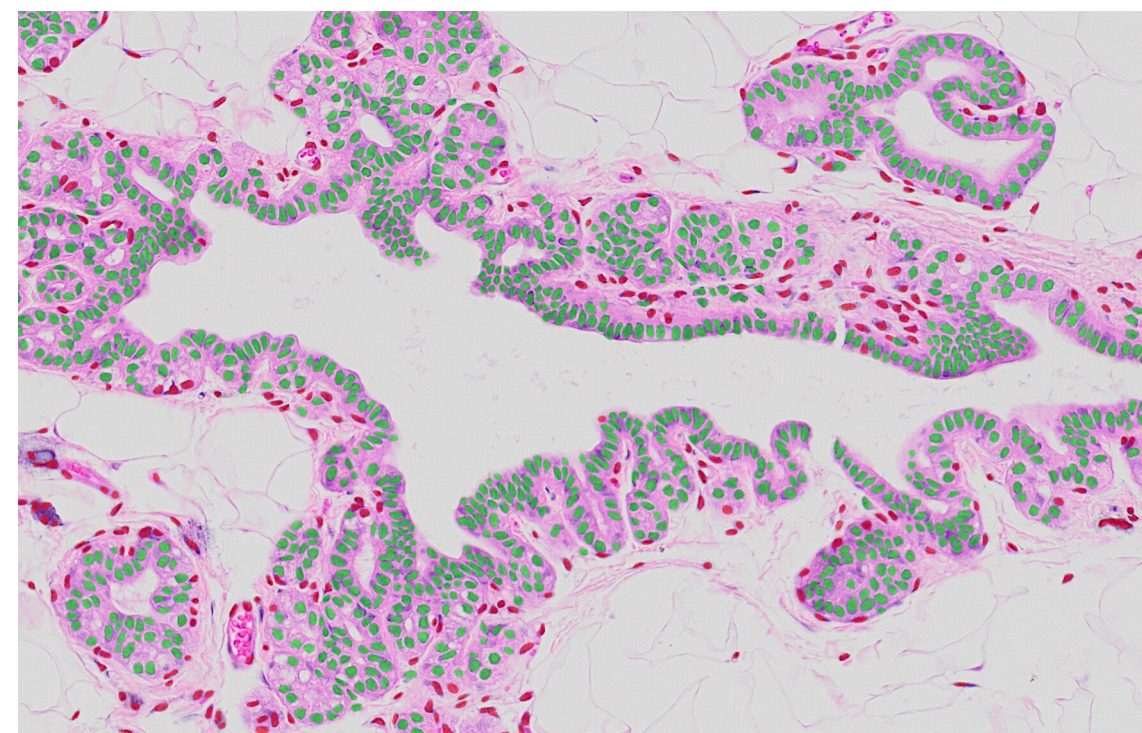
Superpixel ➡ Objects



Cell ➡ Objects



Image ➡ Objects



Morphometric features: 68.8 %  
Textural information: 85.7%  
Contextual, connectivity: 96.4%

Q. Juppert, et al., Journal of  
Mammary Gland Biology and  
Neoplasia, 2021





# Graph Structure

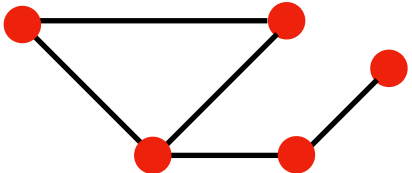
**G**raph

**V**ertex (nodes, points)

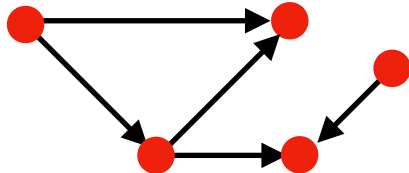
**E**edges (arc, line)

$$G = (V, E)$$

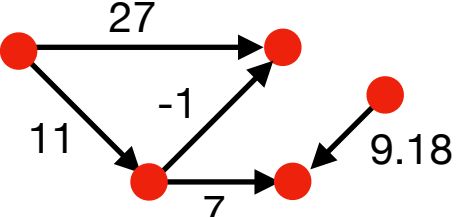
Type



undirected

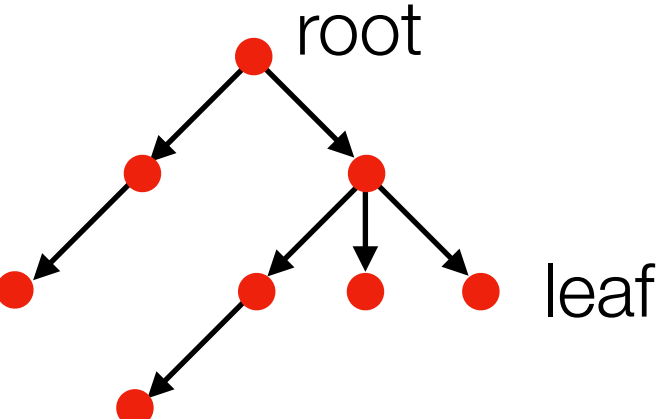


directed

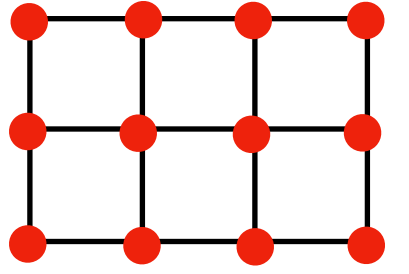


weighted

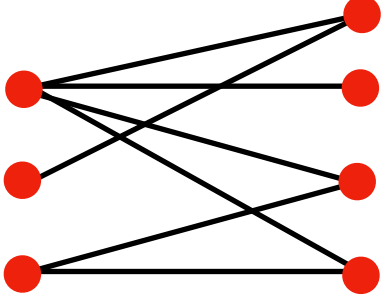
Structure



tree

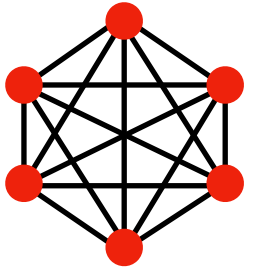


adjacency graph

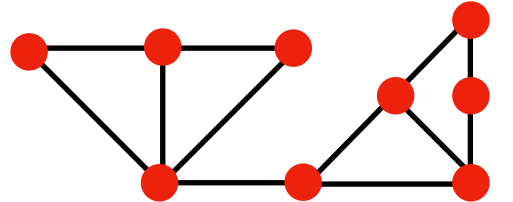


bipartite graph

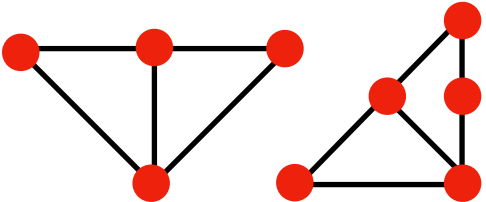
Connectivity



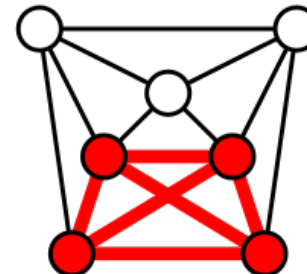
full



weak

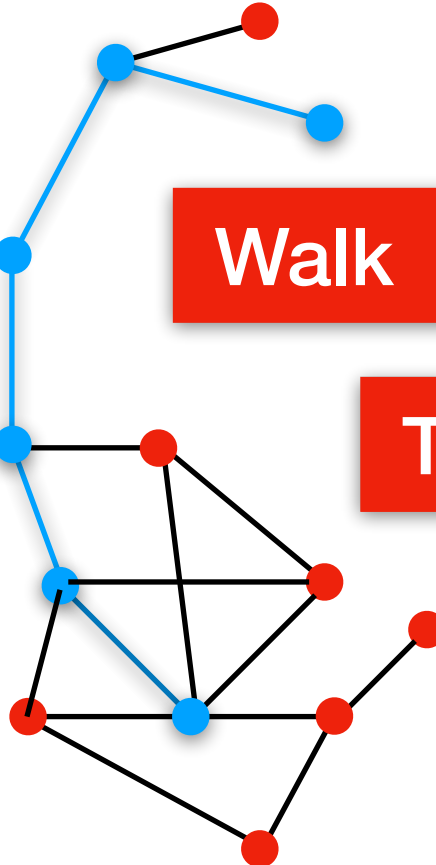


disconnected



clique

subgraph fully connected  
Find cliques is a  
NP-hard problem



**Walk**

sequence of alternating vertices and edges

**Trial**

walk with no repeated edges

**Path**

open trail with no repeated vertices

**Cycle**

closed trail with no repeated vertices





# Graph Representation in Java

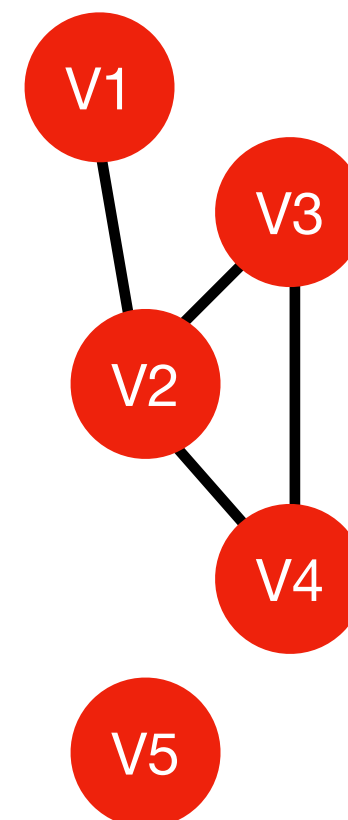
## Data Structure

- e.g. Object, ArrayList, Linked List, Map

```
public class Vertex {
    public String name;
    public ArrayList<Vertex> vertices;
    public Vertex(String name) {
        vertices = new ArrayList<Vertex>();
        this.name = name;
    }
}

public GraphExample() {
    List<Vertex> graph =
        new ArrayList<Vertex>();
    for(int i=1; i<5; i++)
        graph.add(new Vertex("V" + i));
    addEdge(graph, 1, 2);
    addEdge(graph, 2, 3);
    addEdge(graph, 3, 4);
    addEdge(graph, 2, 4);
}

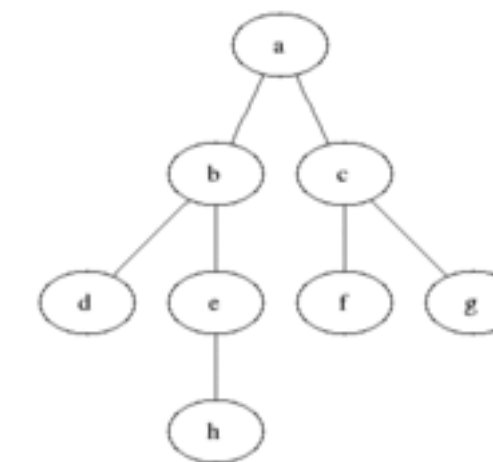
void addEdge(List<Vertex> g, int a, int b) {
    g.get(a).vertices.add(g.get(b));
    g.get(b).vertices.add(g.get(a));
}
```



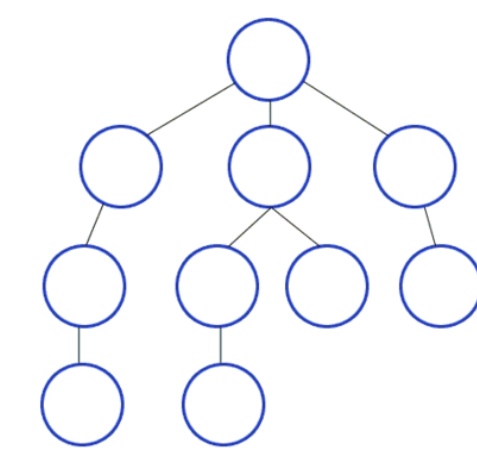
## Recursive algorithm

- e.g. graph traversal

```
void dfs(int i, int[][] mat, boolean[] visited) {
    if(!visited[i]) {
        visited[i] = true;
        System.out.print( (i+1) + " ");
        for (int j = 0; j < mat[i].length; j++)
            if (mat[i][j] == 1 && !visited[j])
                dfs(j, mat, visited); // Visit
    }
}
```



Breath First Search



Depth First Search

## Third-party Java libraries: JGraphT

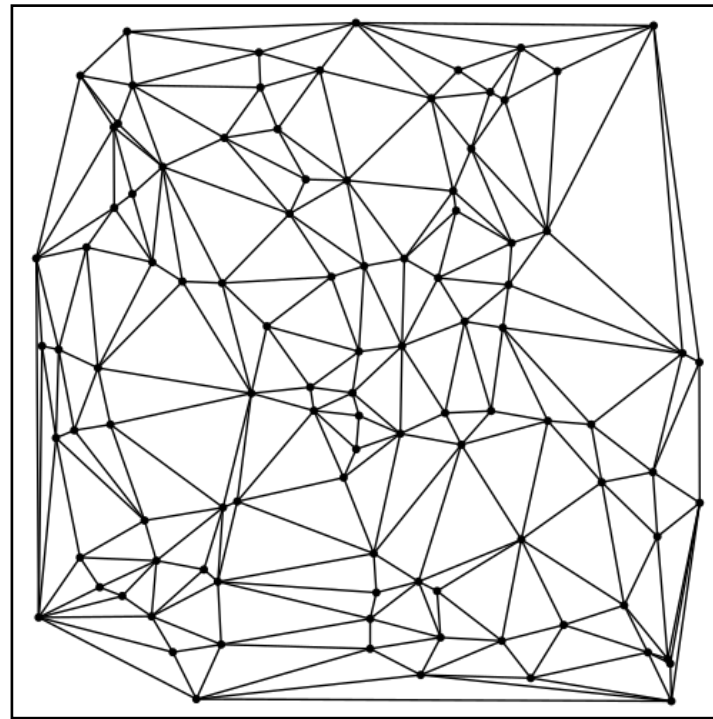
- e.g. Voronoi, Delaunay, Hungarian



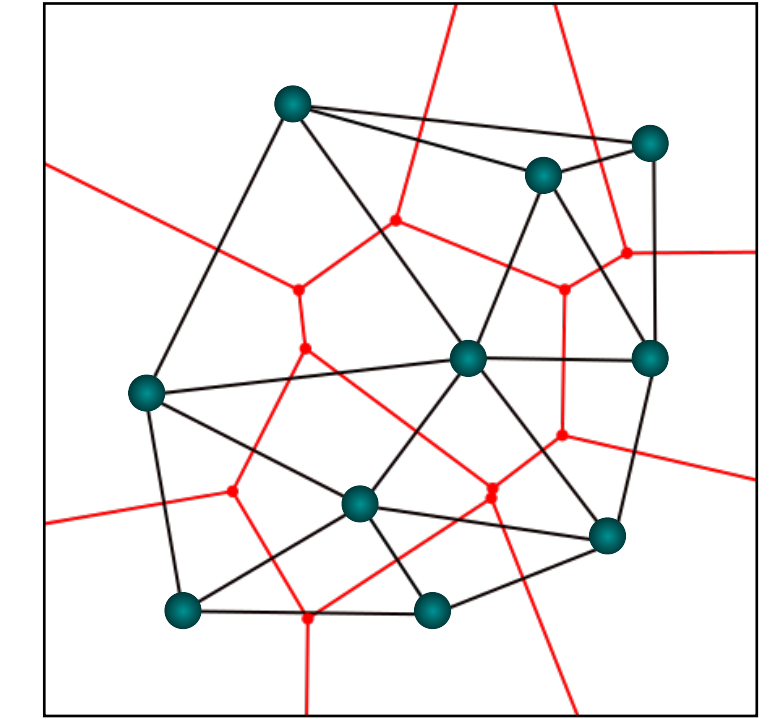
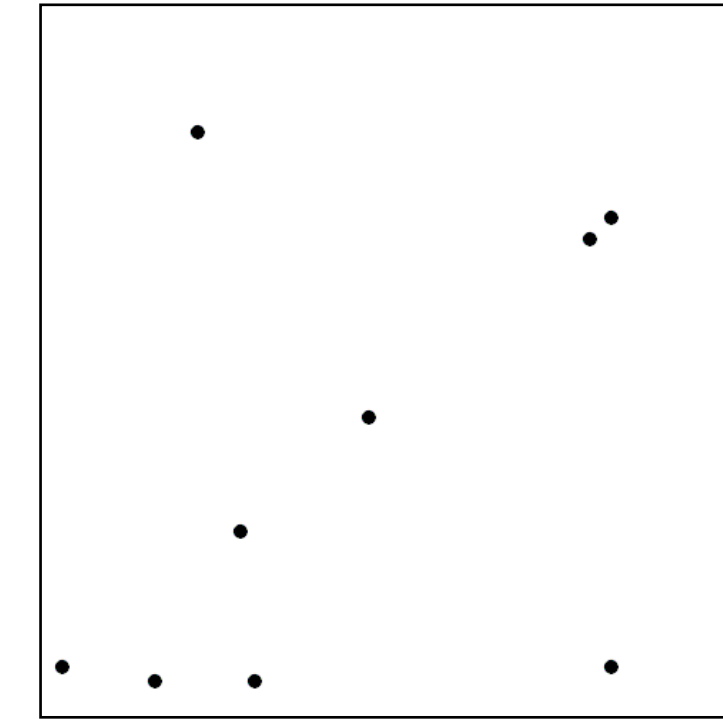
# Planar Graph

A **planar graph** is drawn without edges crossing → the edges are defining regions (faces)

Delaunay triangulation  
built from points

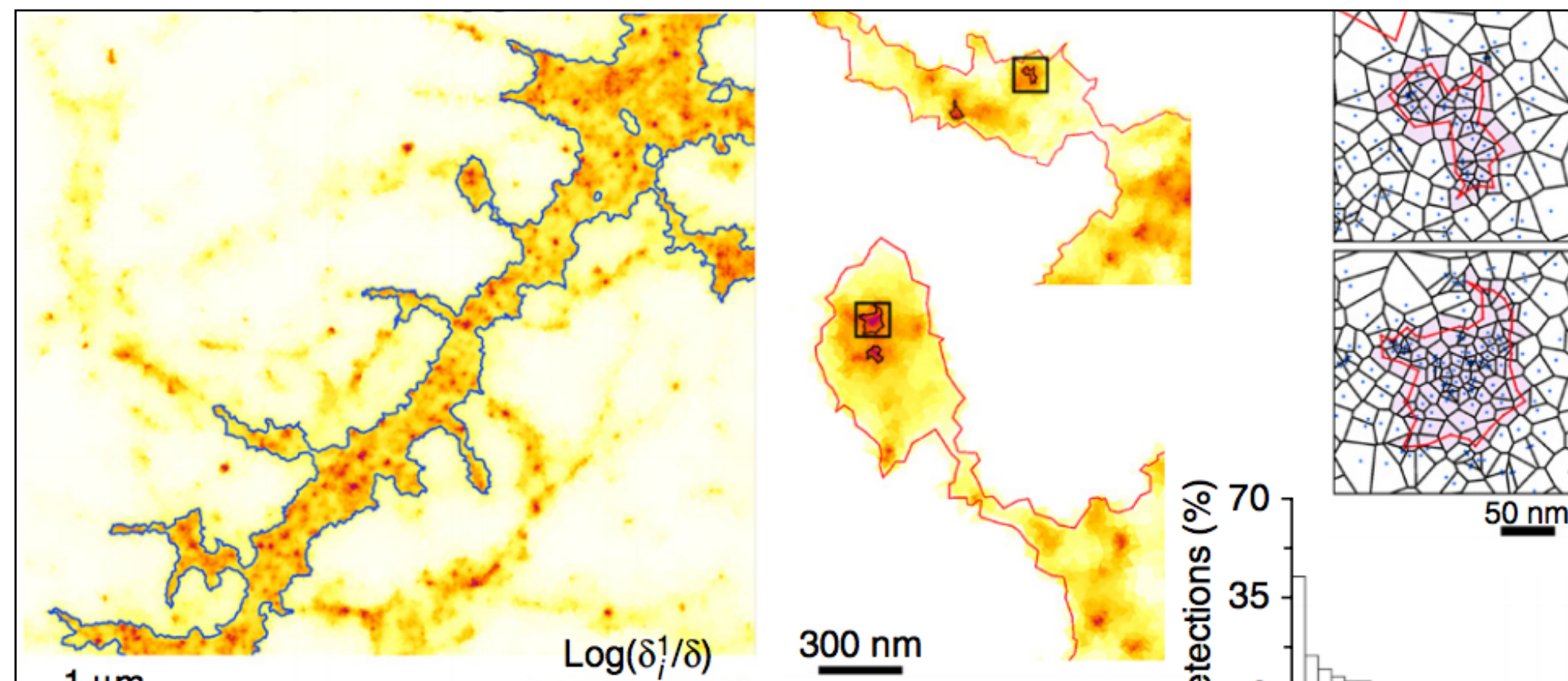


Voronoi  
tessellation  
from seed  
points

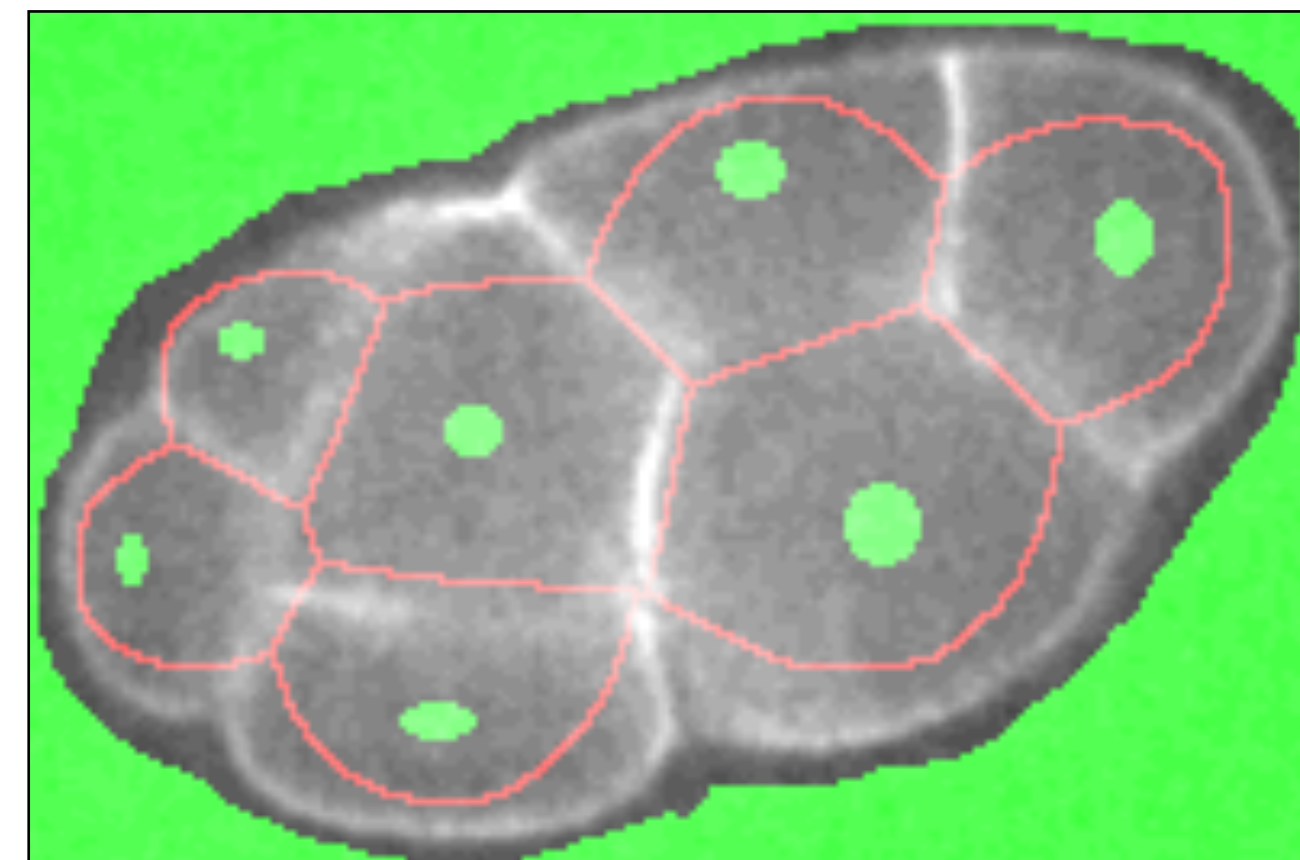


Dual graph

- Voronoi diagram
- Delaunay triangulation



Florian Levet. SR-Tesseler: segment and quantify localization-based super-resolution microscopy data, *Nature Method* 2015



*C. elegans* embryo, cell membrane and nucleus. Courtesy of Radek Janeke, EPFL





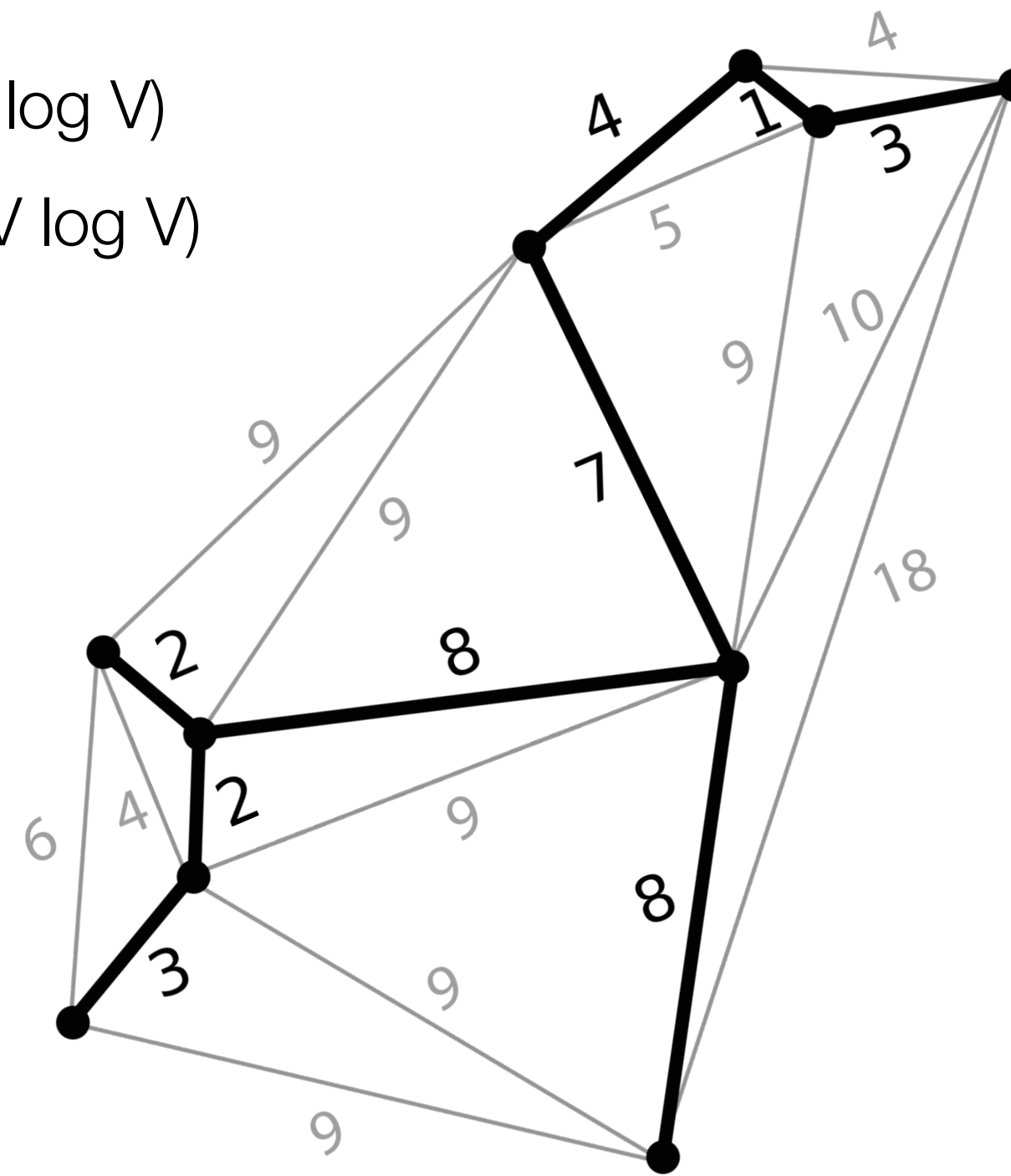
# Minimum Spanning Tree

## Tree-like structure

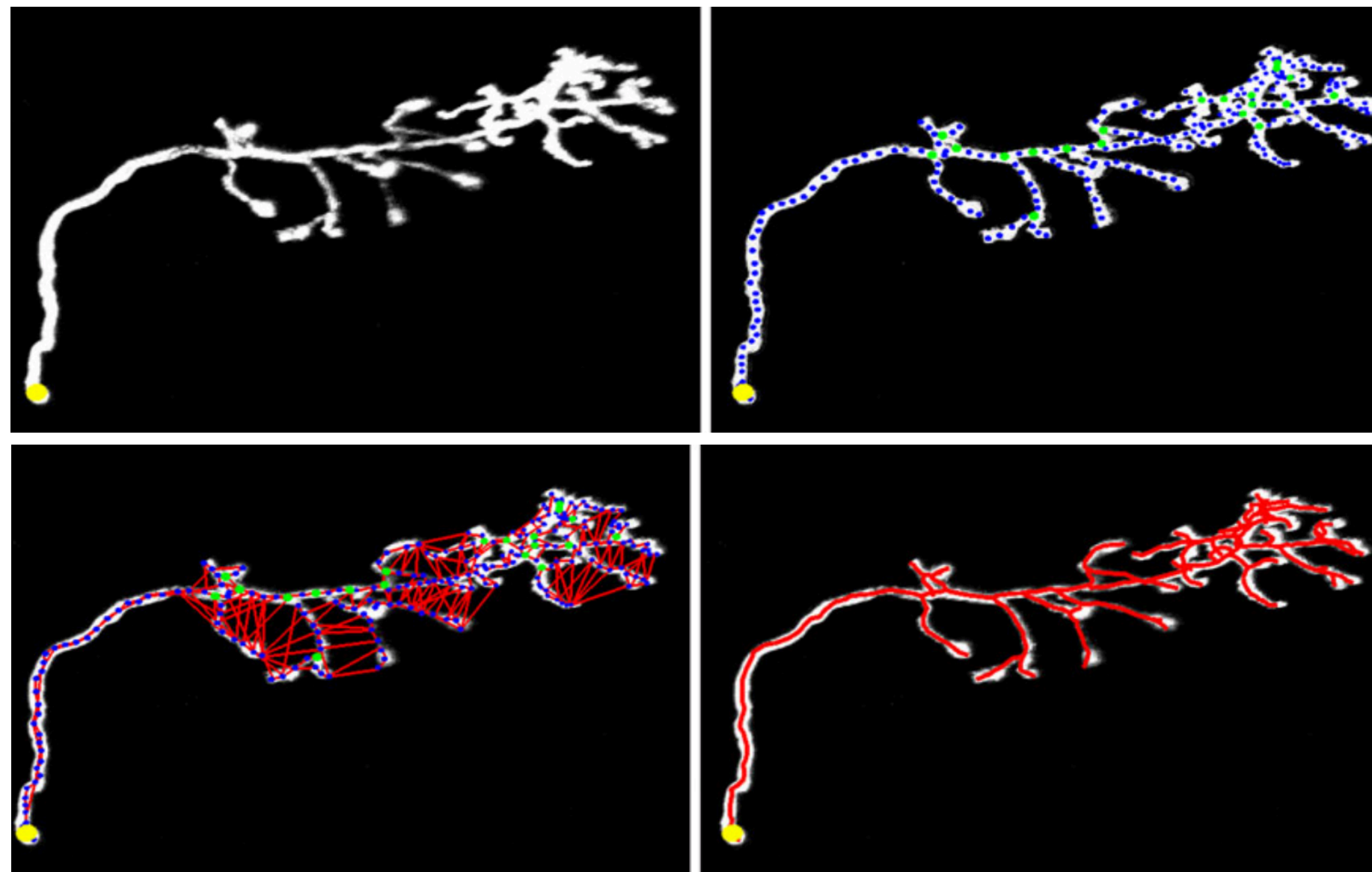
- Vascular network, dendrites, ...
- Space 2D or 3D
- Time: Cell lineage

## Solver

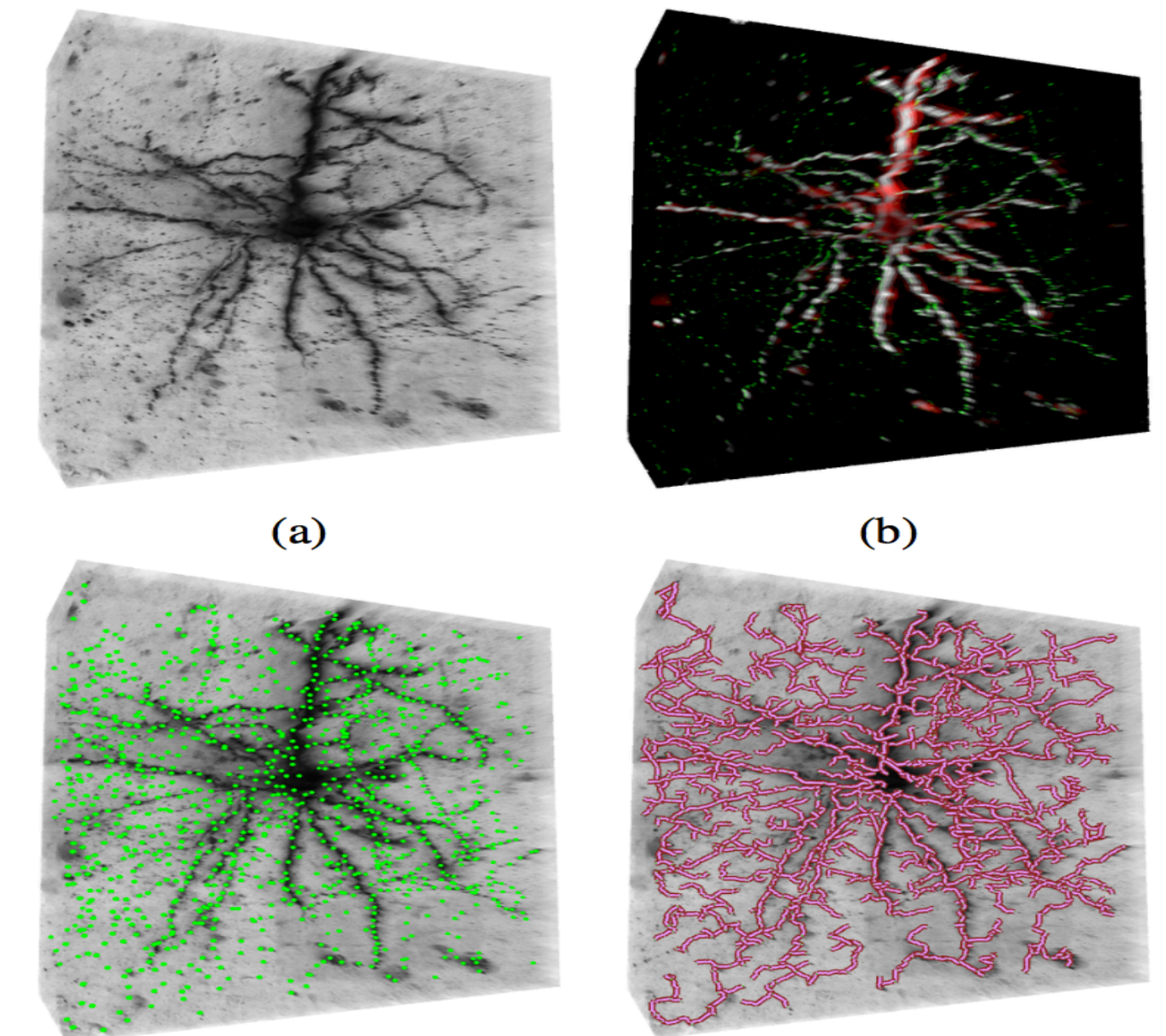
- Kruskal  $\mathcal{O}(E \log V)$
- Prim  $\mathcal{O}(E + V \log V)$



<https://visualgo.net/en/mst>



Olfactory projection fibers, E. Türetken, Neuroinform, 2011



Dendrites. German Gonzalez, ECCV 2008





# Bipartite Graph Matching

$$G = (V, E)$$

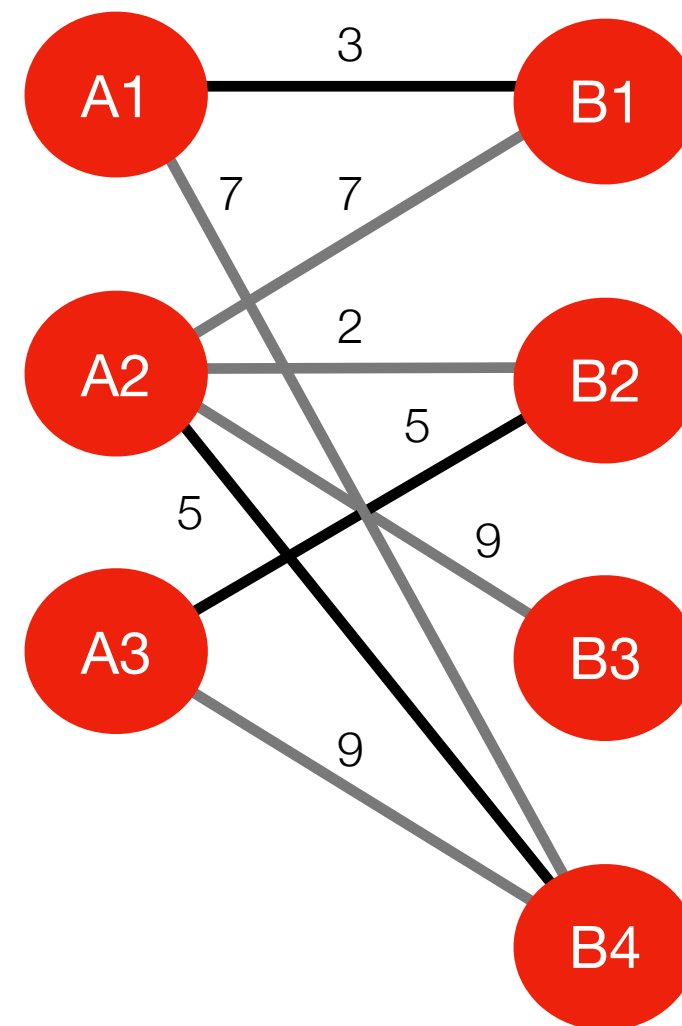
$$V = V_A \cup V_B$$

$$V_A \cap V_B = \emptyset$$

- Directed
- Weight (cost  $\xi_{ij}$ )
- Balanced

## Assignment Problem

- Minimum weight perfect matching
- Hungarian algorithm  $O(E V^2)$



	A1	A2	A3	A4	A5
B1	5	9			6
B2	4	6			
B3	8	5	1		
B4			9	3	2
B5			6	5	3

?

Is it the best match?

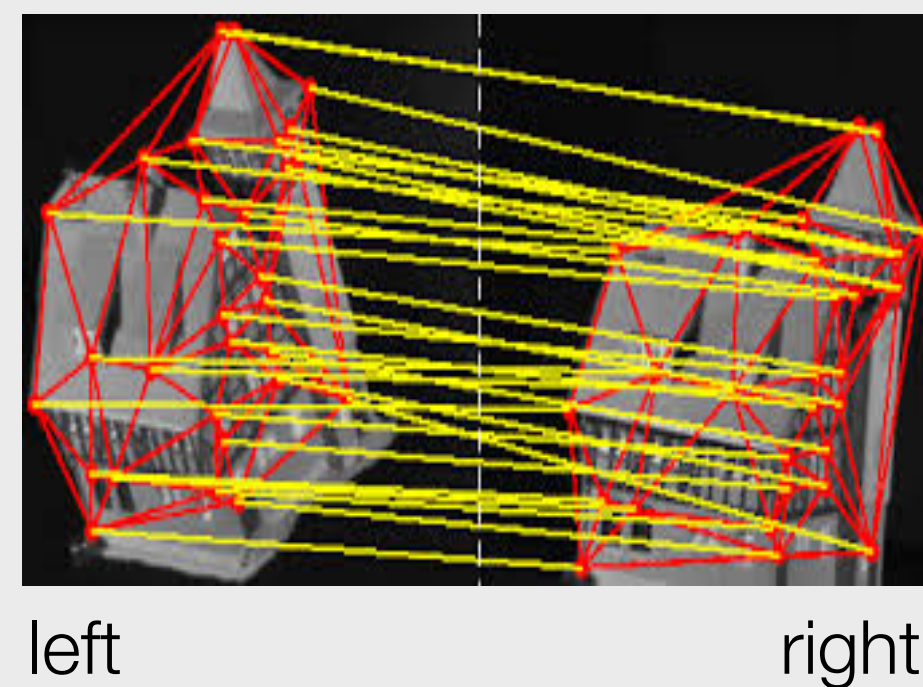
	A1	A2	A3	A4	A5
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B3	8	5	1		
B4			9	3	2
B5			6	5	3

First Minimum  
4+5+6+3+6=24

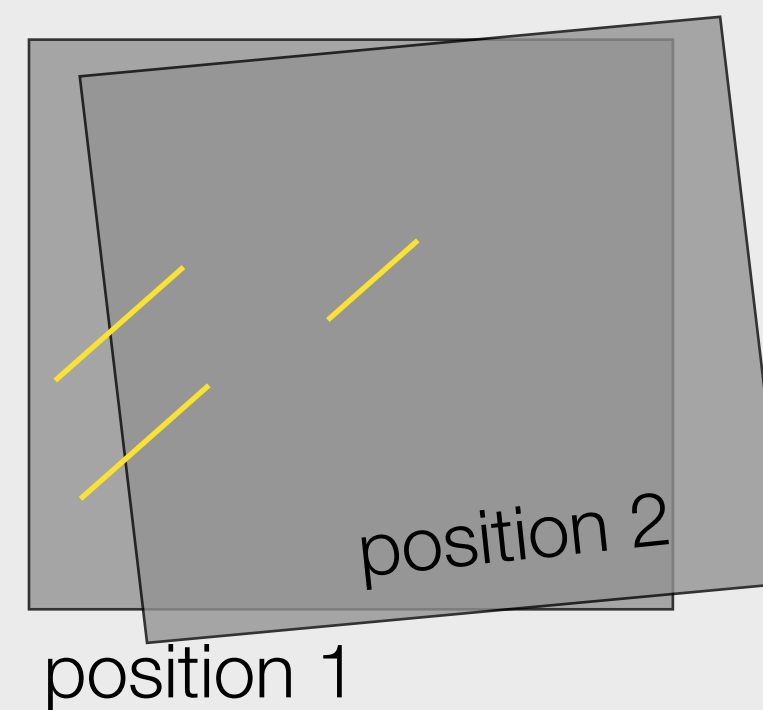
	A1	A2	A3	A4	A5
B1	5	9			6
B2	4	6			
B3	8	5	1		
B4			9	3	2
B5			6	5	3

Greedy  
1+2+4+5+9=21

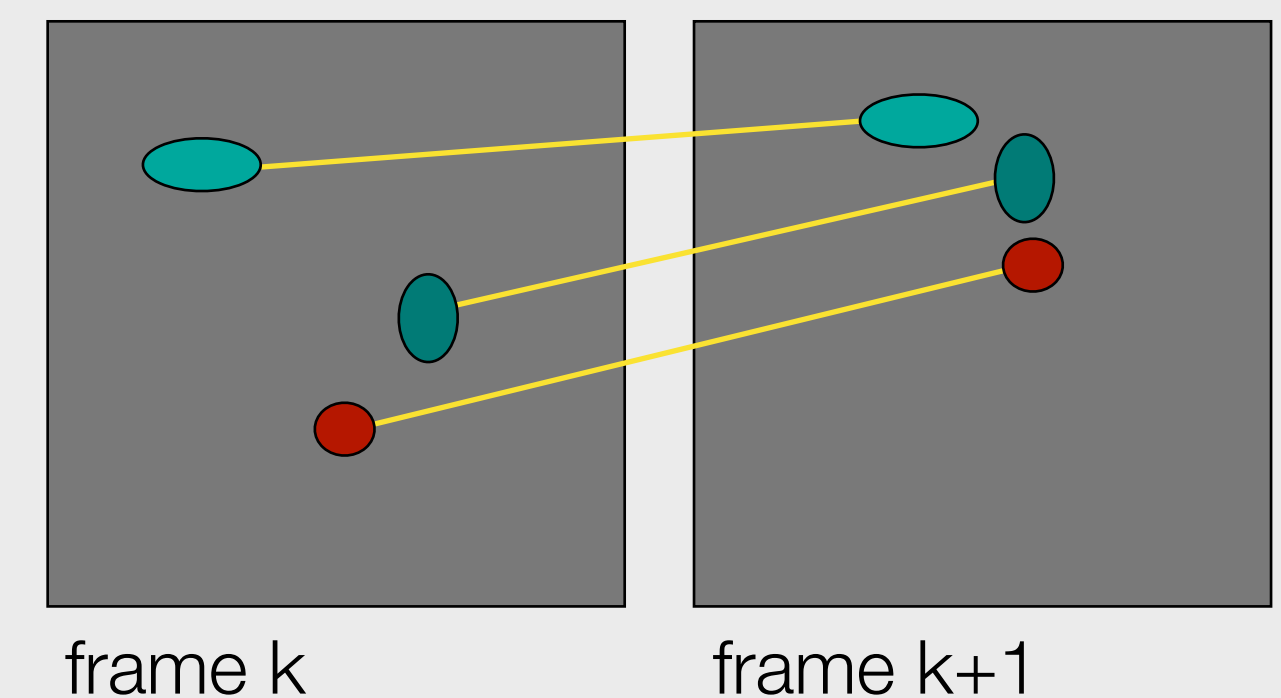
## Stereo matching



## Registration



## Tracking



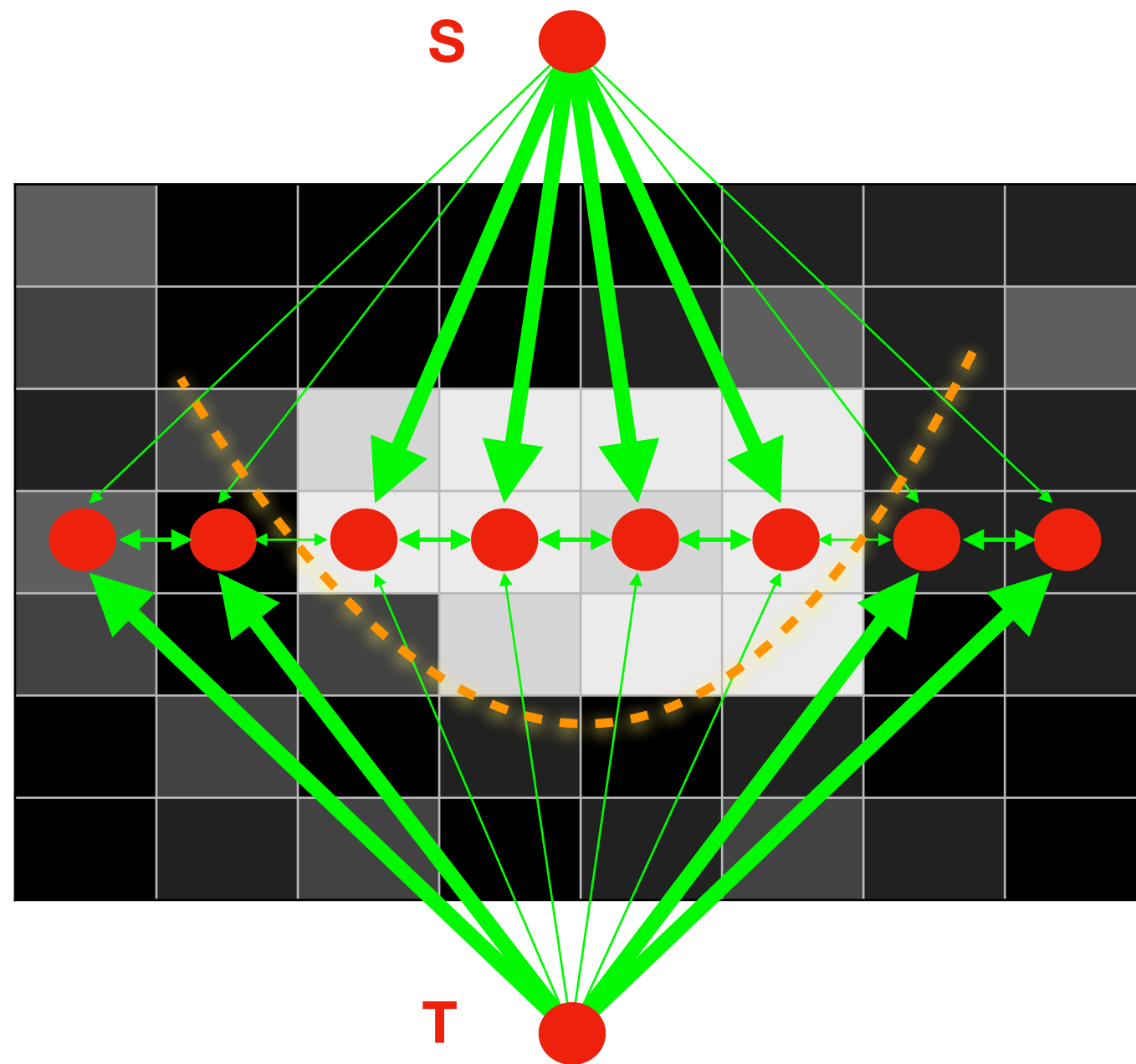




# Graph Cut

## Minimal cut

- Build a graph with a source **S** and a sink **T**
- Based on min-cut max-flow theorem
- Efficient way to compute (Boykov-Kolmogorov)
- Interactivity: user guide partition

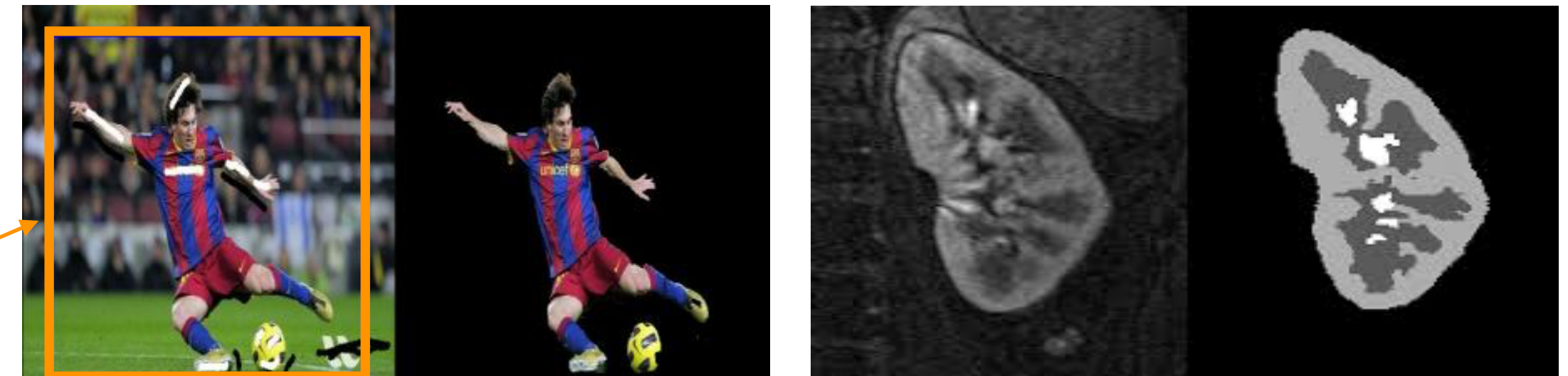


## Weights for image segmentation

- **Source** with probability to belong to the foreground
- **Sink** with probability to belong to the foreground
- **Neighborhood** with a measure of similarity of adjacent pixels

## Application to image segmentation with complex background

background



Interactive Graph Cuts [www.csd.uwo.ca](http://www.csd.uwo.ca)

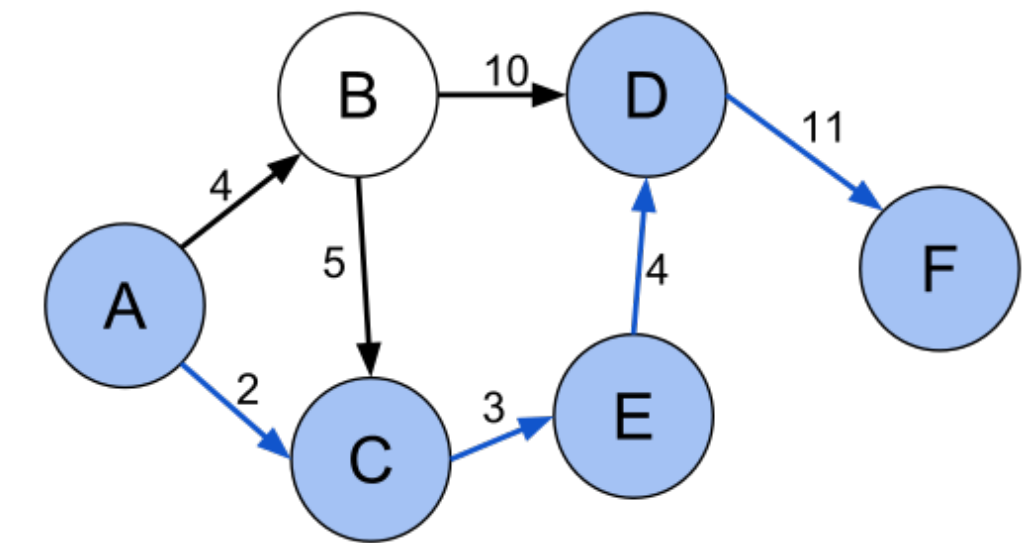




# Shortest Path

## Shortest path problem in weighted graph

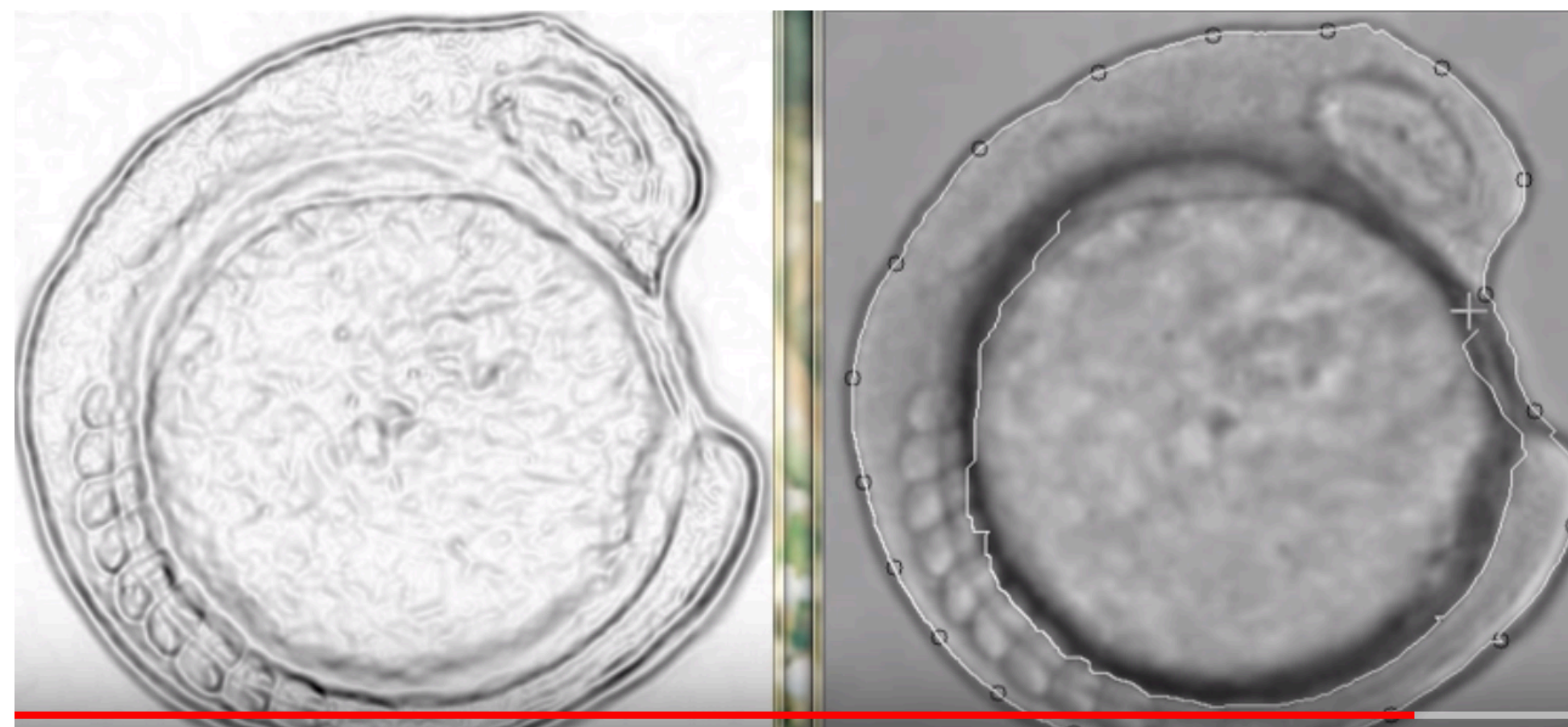
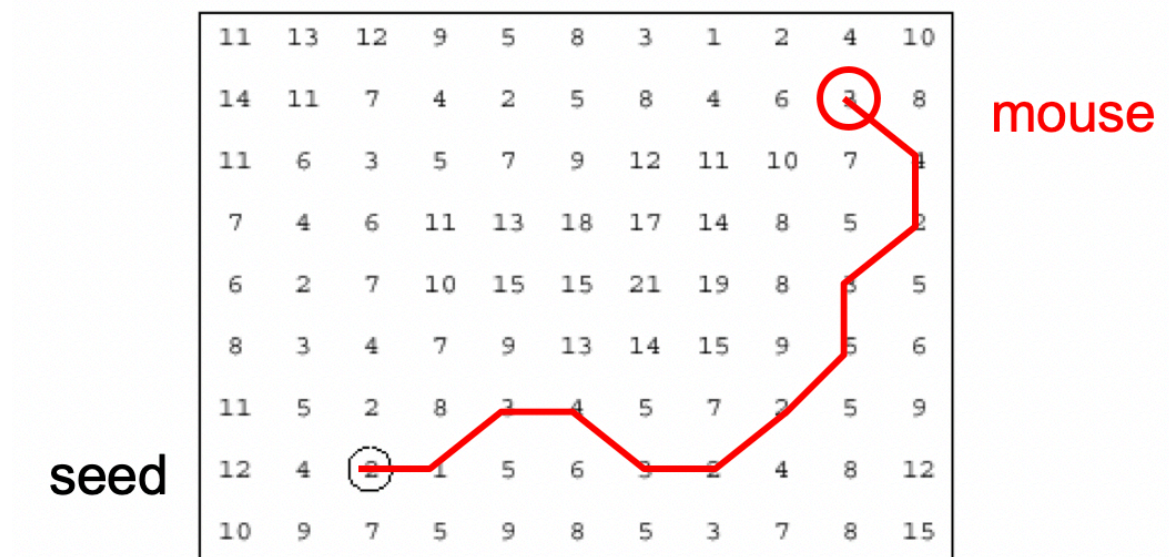
- Find a path between a starting vertex and an ending vertex
- Minimise the sum of weights of edges
- Applications: GPS route, intelligent scissors in imaging, clipping path in photo...



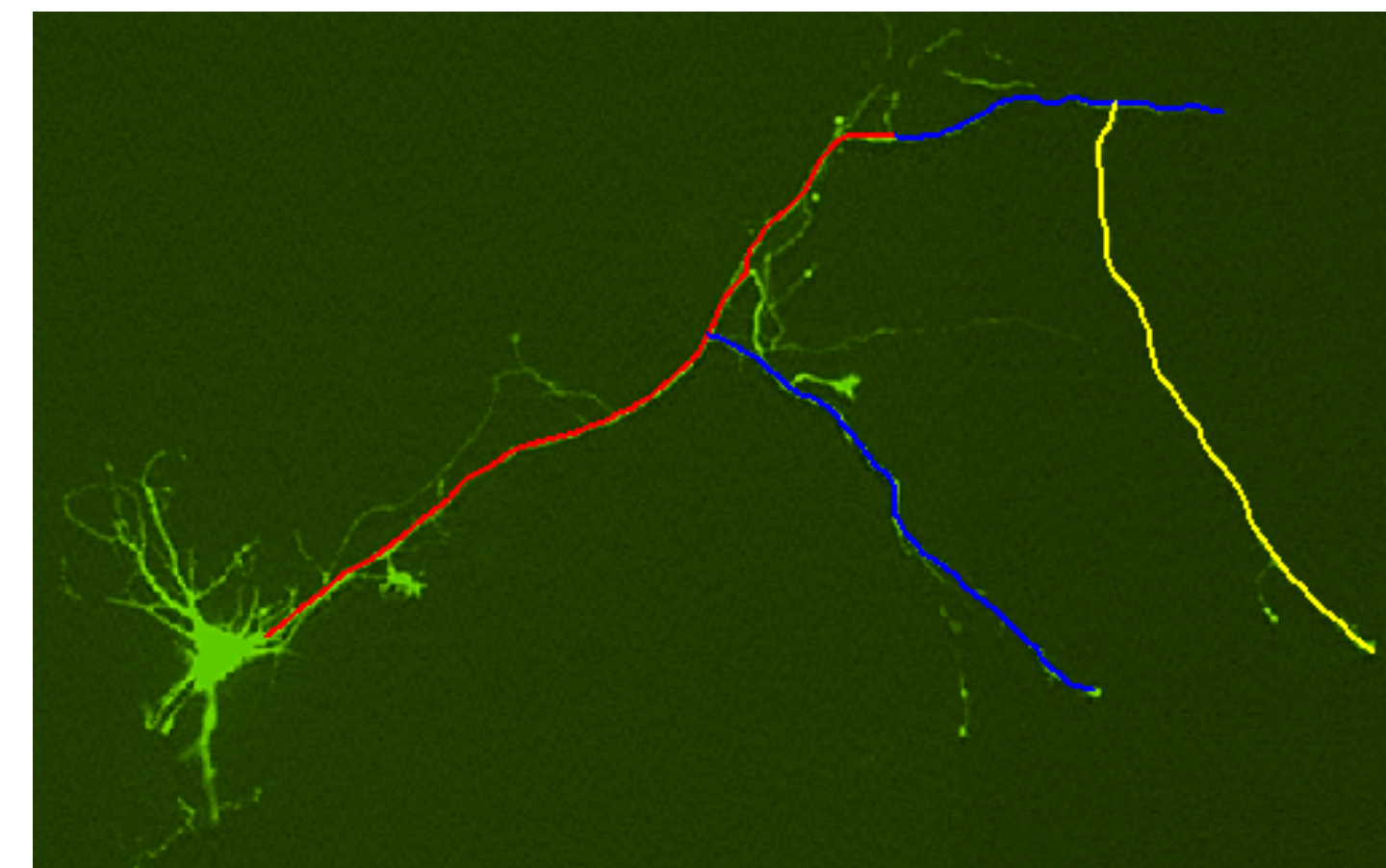
## Dijkstra Algorithm

### Explore to the end point and backtrack to start point

- Picks the unvisited vertex with the lowest distance
- Distance through it to each unvisited neighbor
- Direct implementation  $O(V)$  → Fast  $O(E+V\log(V))$



[https://www.youtube.com/watch?v=X\\_dZ\\_7xAclM](https://www.youtube.com/watch?v=X_dZ_7xAclM)  
 Franklin Fang, Intelligent Scissor



**Livewire**  
 Erik Meijering, Neuronj



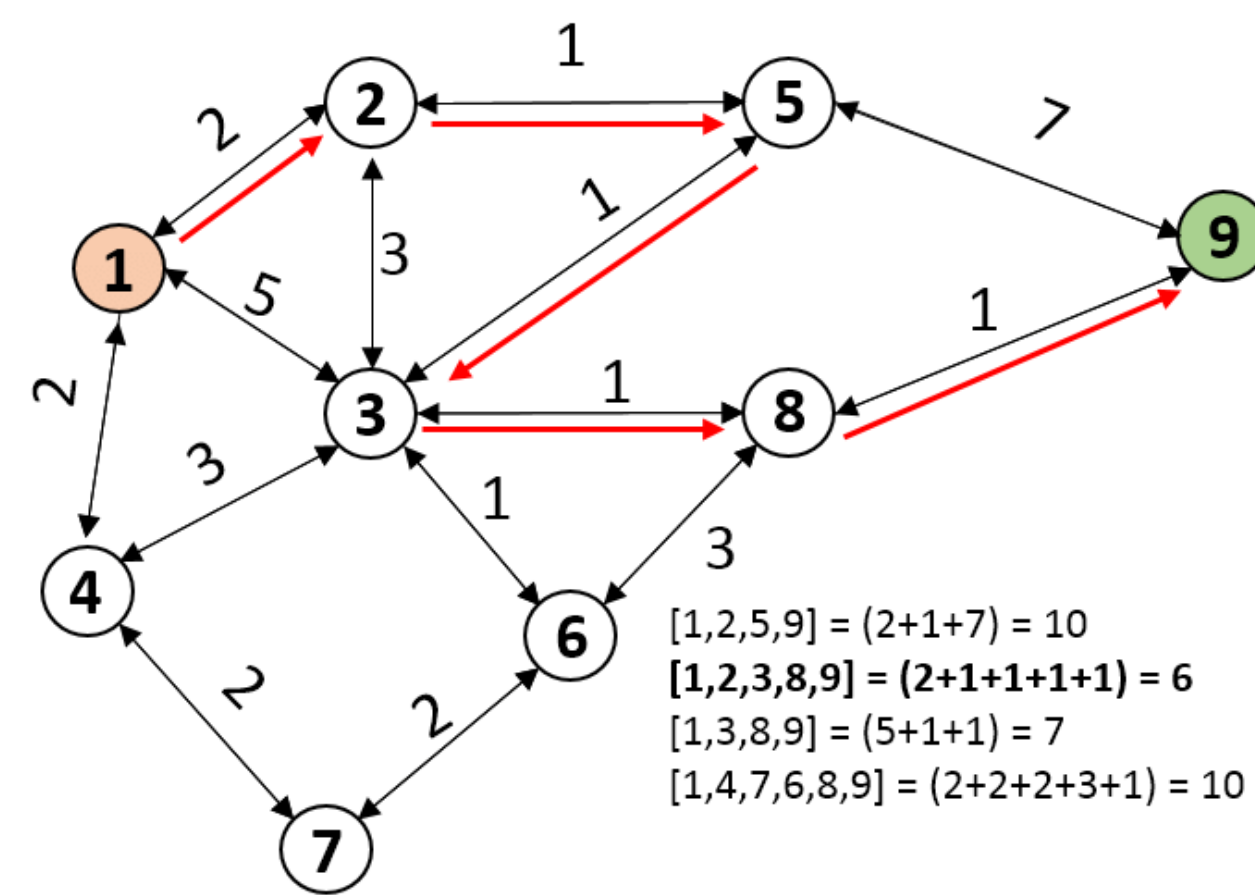


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## Shortest Path

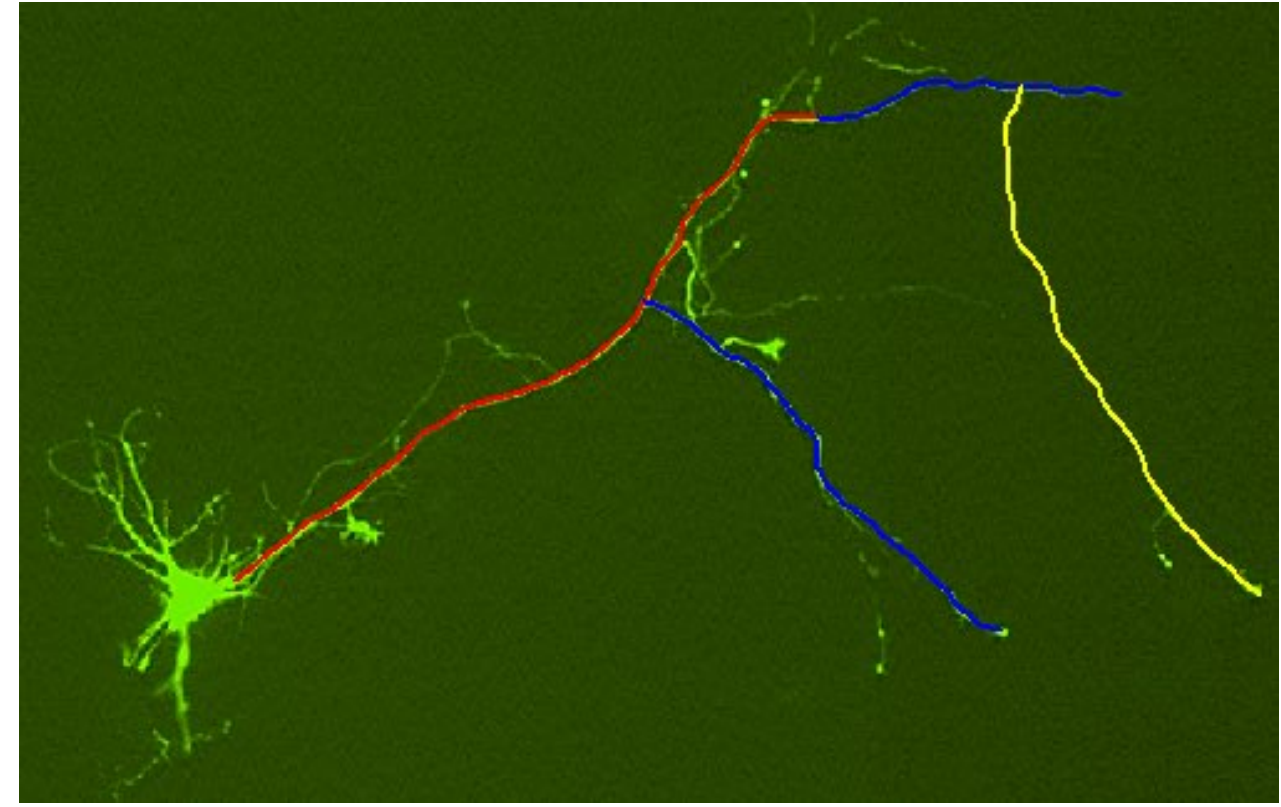


## Dijkstra Algorithm

### Find the shortest route



### NeuronJ



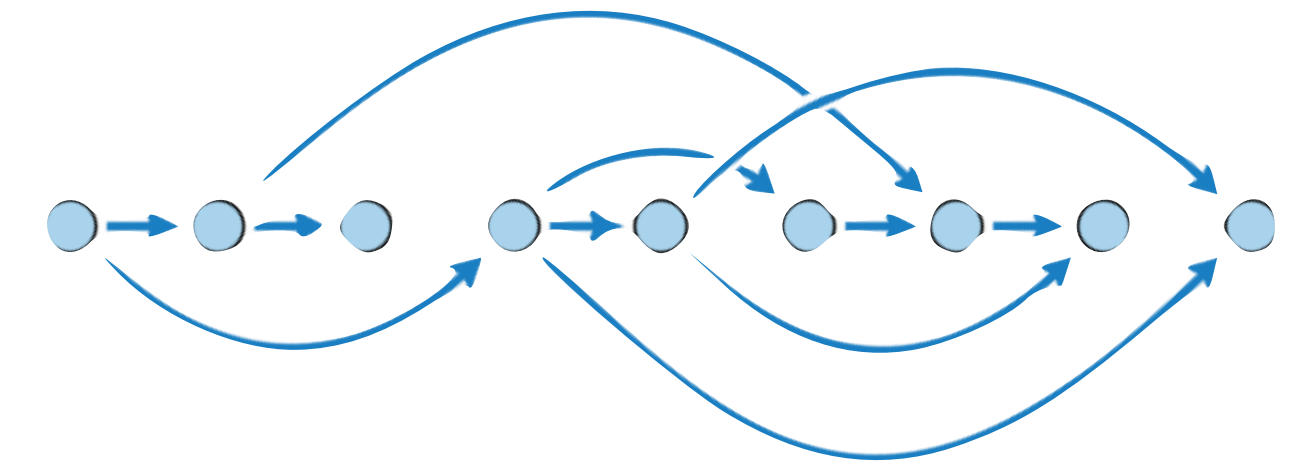
E. Meijering et al. Cytometry 2004



# Dynamic Programming

A **directed acyclic graph (DAG)** is a directed graph with no directed cycles

→ A DAG has a topological ordering of its vertices into a sequence



## Dynamic Programming

### Viterbi algorithm

**Bellman's principle:** break this decision problem into smaller subproblems

#### Building the graph on the pixel image (connectivity)

- Possible paths from start point to end point
- **Smoothness** aperture  $\Delta$  controls the possible jump
- **Confinement area** limitation the search area

#### DP engine

- Explore the graph in sequence
- Backtrack
- Minimising the sum of costs

#### Cost a the edge

- from a vertex A to a vertex B
- customise to a specific problem
- cost = weight of edge



# DP Optimal Detection of Curves

## Powerful and fast optimiser for curve detector in images

- Discretization along the initial line
- Cost: weighted sum of penalisation / attraction
- Easy to add constraint point

$$\xi_{\mathbf{u}\mathbf{v}} = \lambda_1 \frac{f(\mathbf{u})}{N_{int}} + \lambda_2 \frac{|\mathbf{u} - \mathbf{v}|}{N_s}$$

$\uparrow$  data term       $\uparrow$  regularisation term

