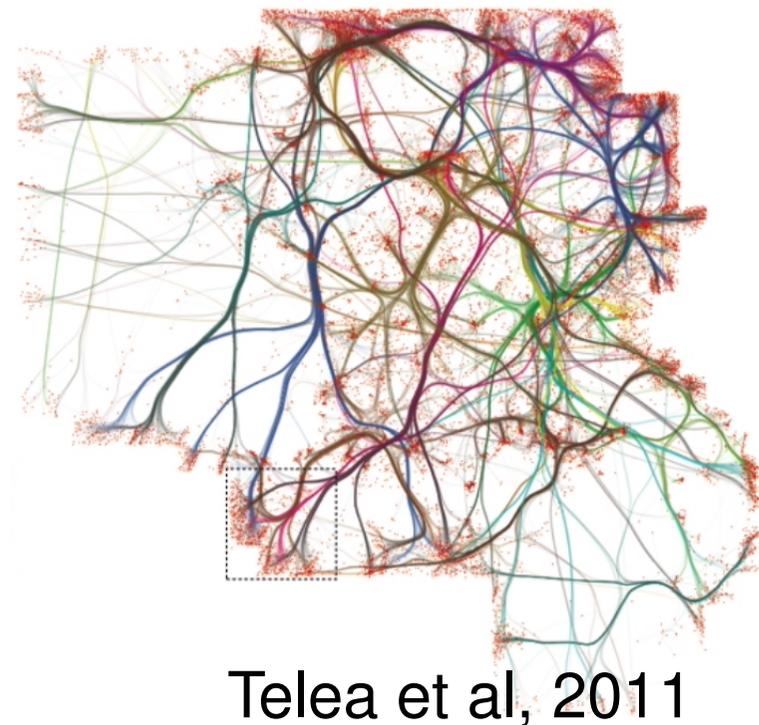
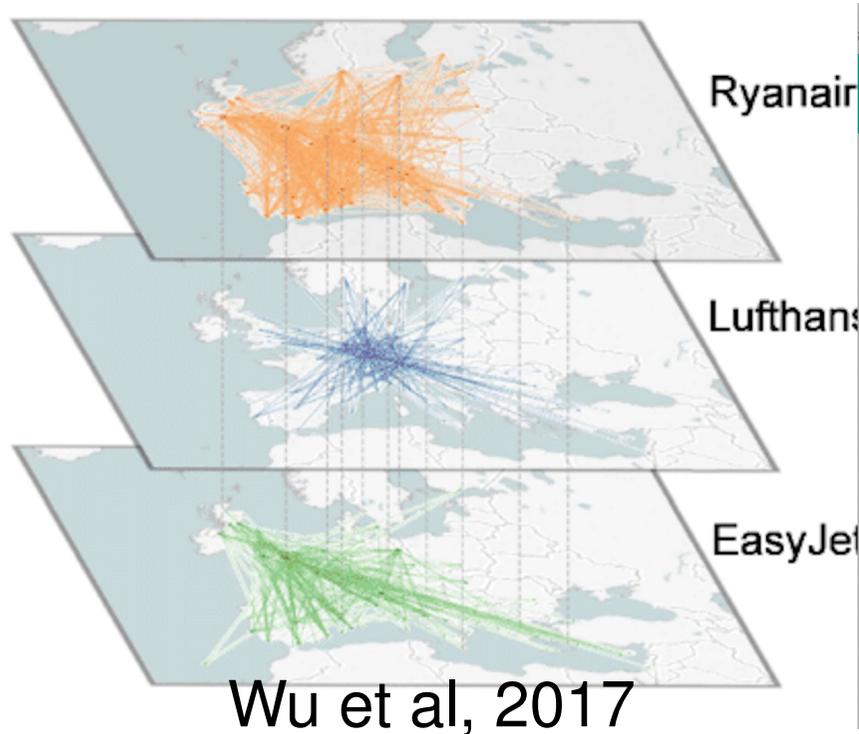


# Multilayer Network Visualization

**Course :** Data Visualization

**Lecturer :** Tamara Mchedlidze

Utrecht University, Dept. of Information and Computing Sciences



# Lecture Overview

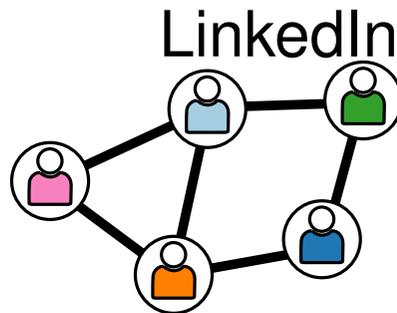
- **Multilayer network**
- **Visualization types for multilayer networks**
- **Algorithm for visualization in 2.5D**
- **Edge simplification - bundling**
- **An algorithm for edge bundling**
- **Proposed technique for the implementation**

# Adding complexity

- definition of network/graph we used till now (nodes, edges and perhaps labels) is a simplification of reality, where the network are often way more complex

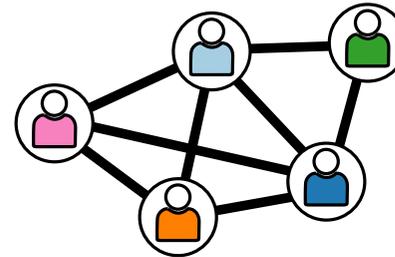
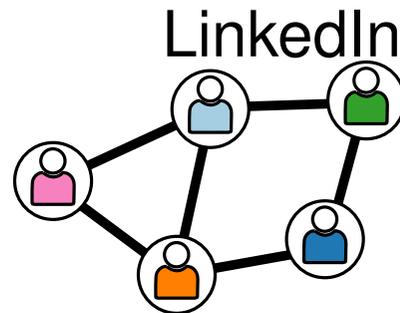
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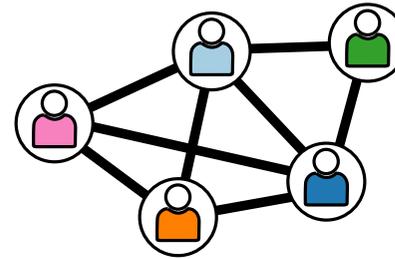
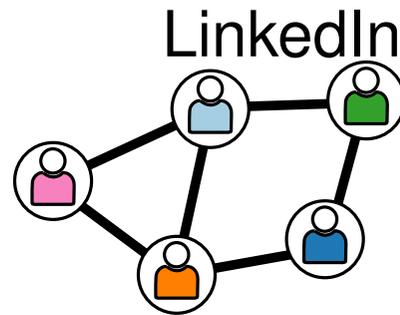
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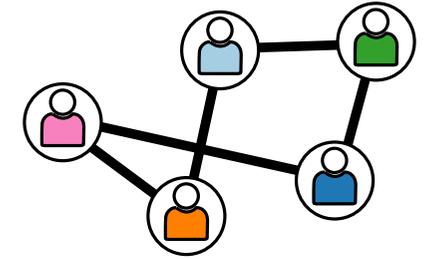
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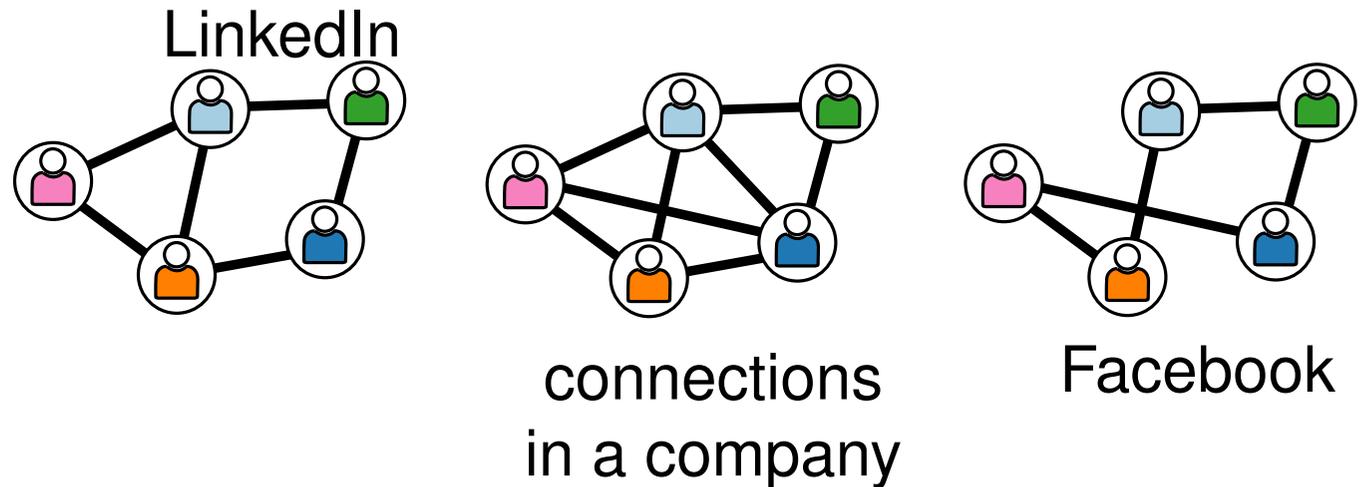
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Facebook

# Adding complexity

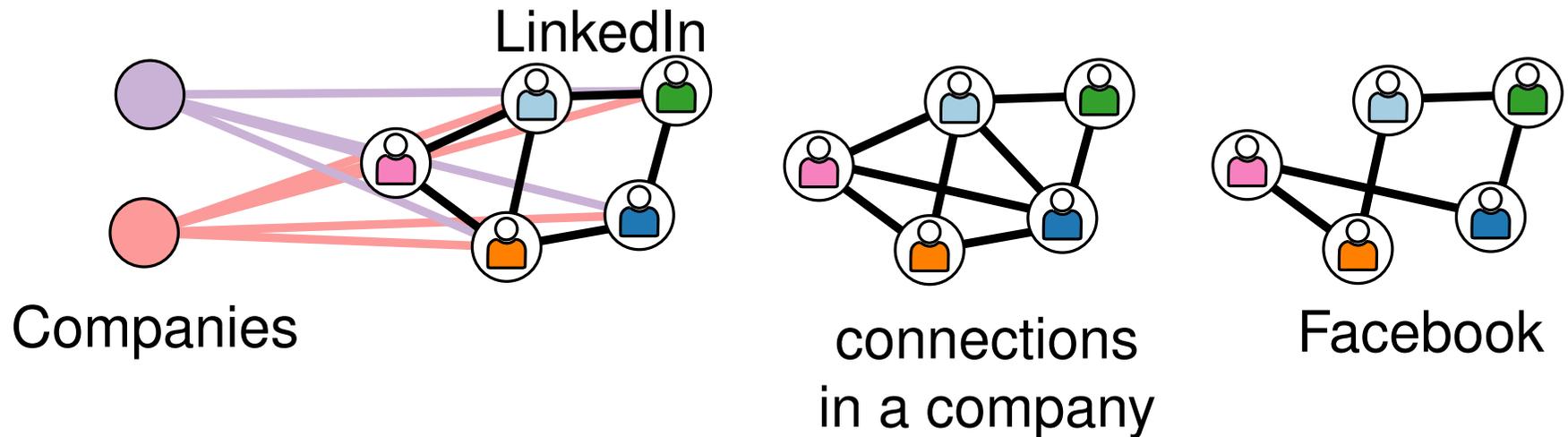
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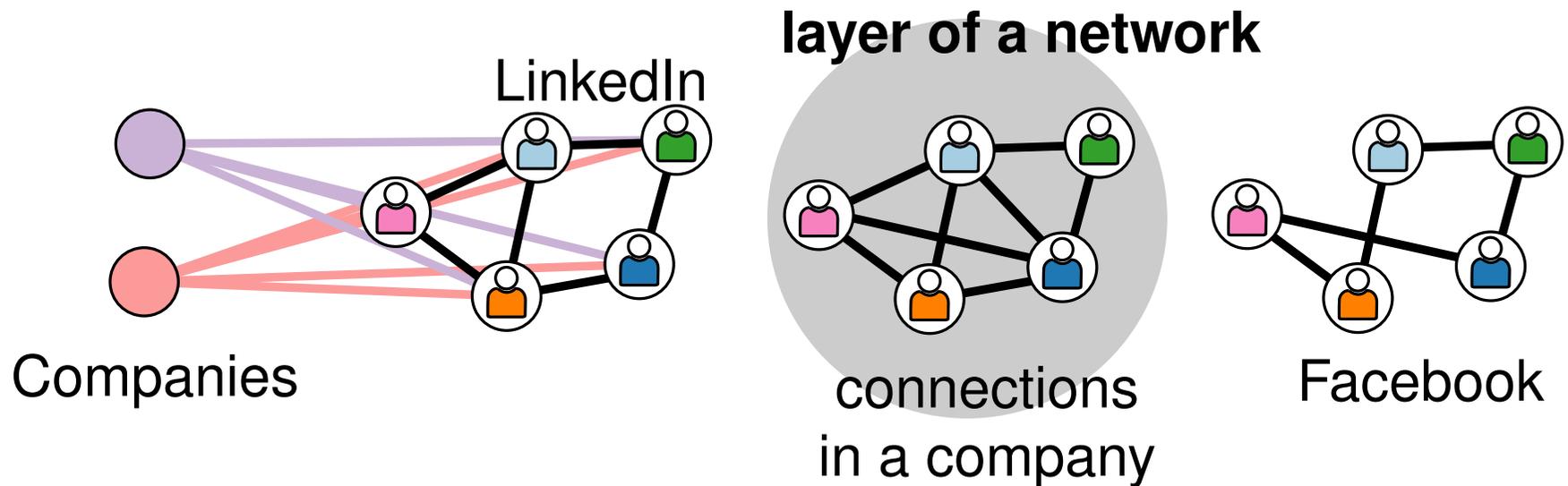
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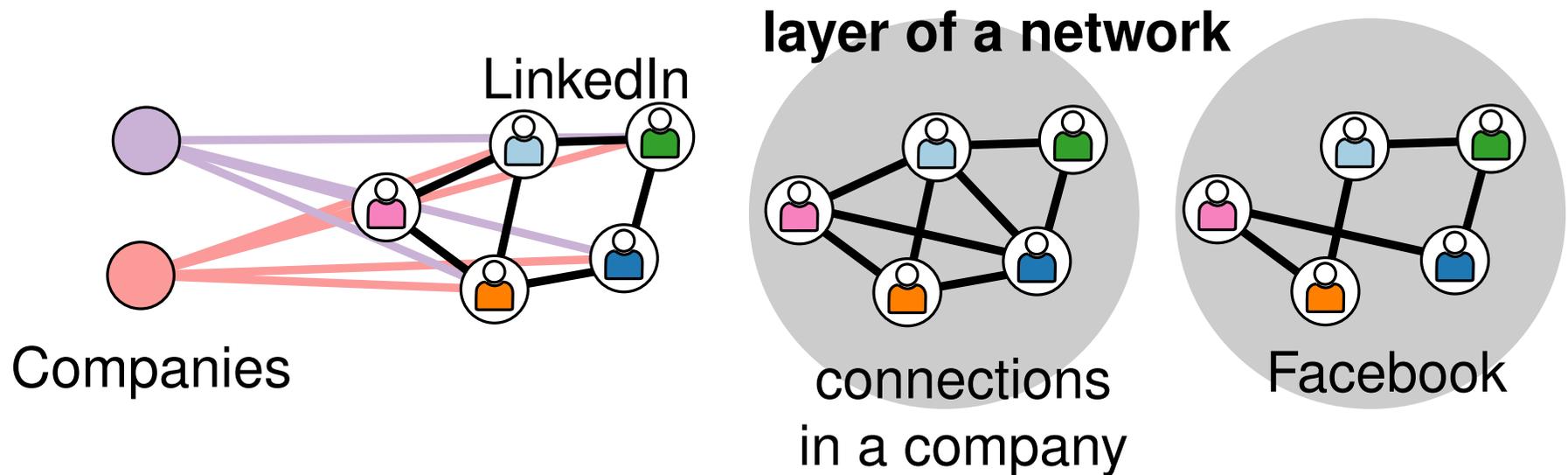
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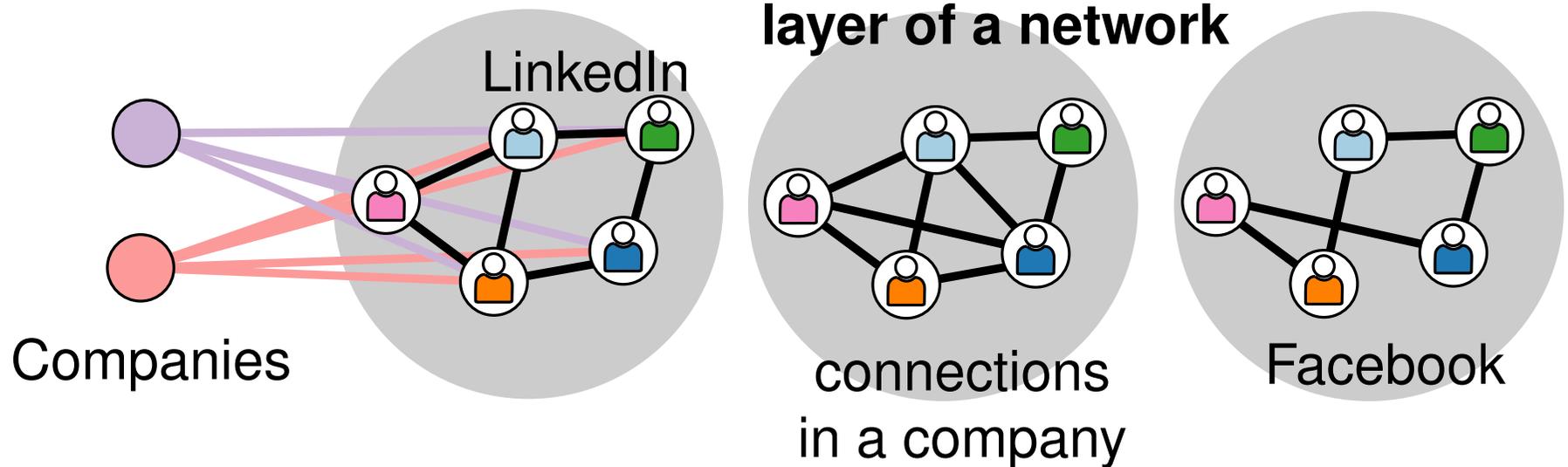
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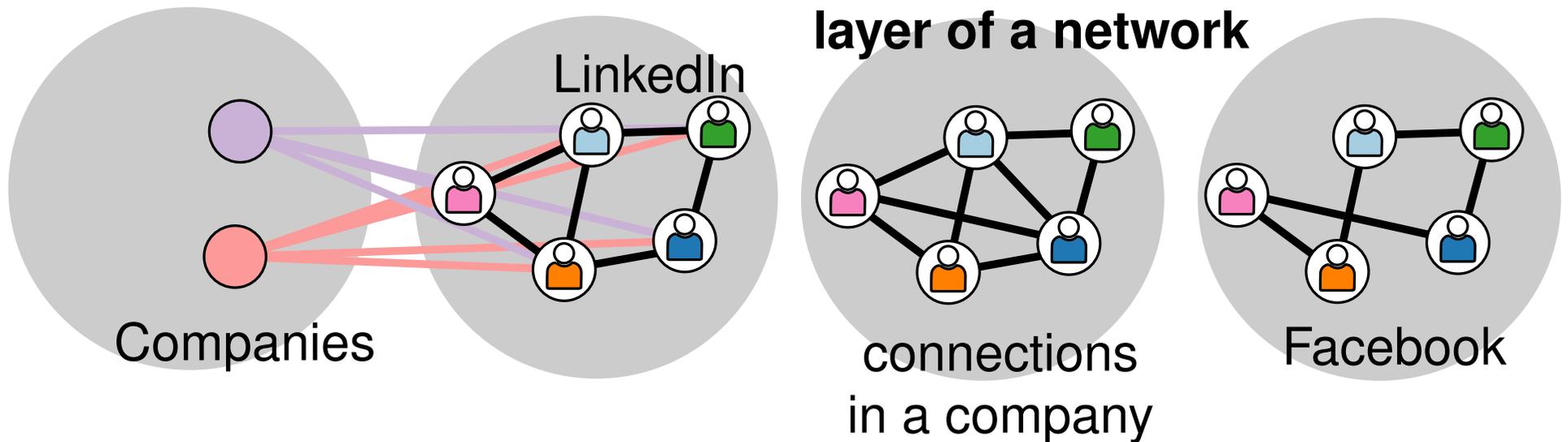
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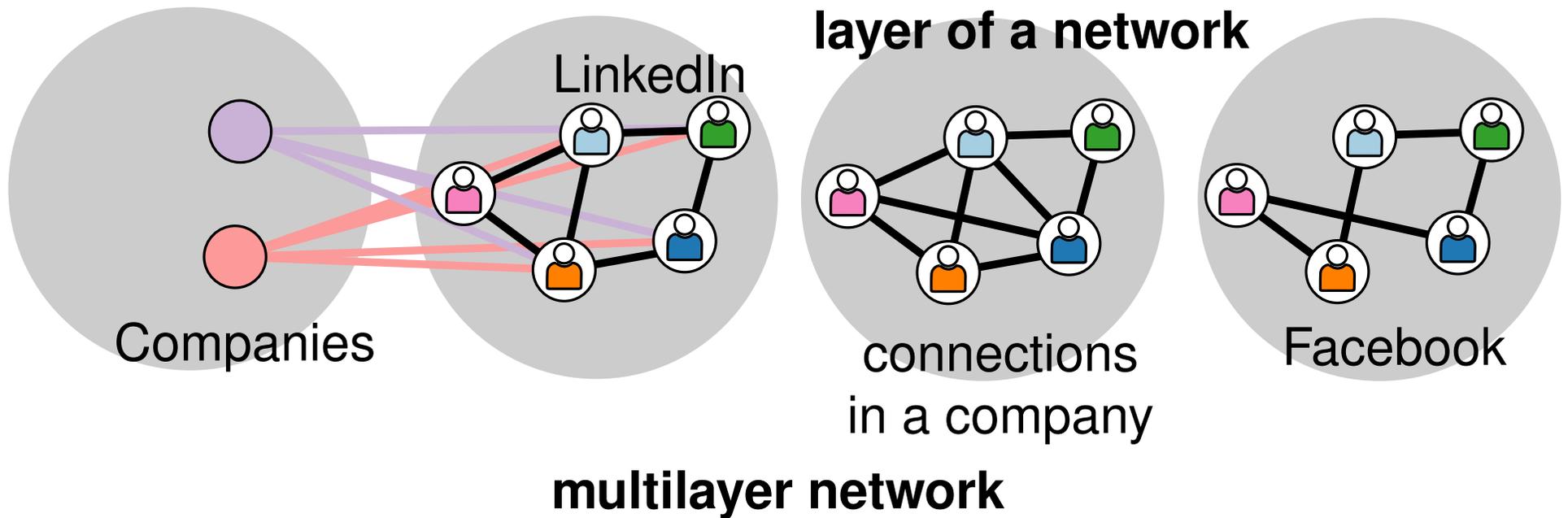
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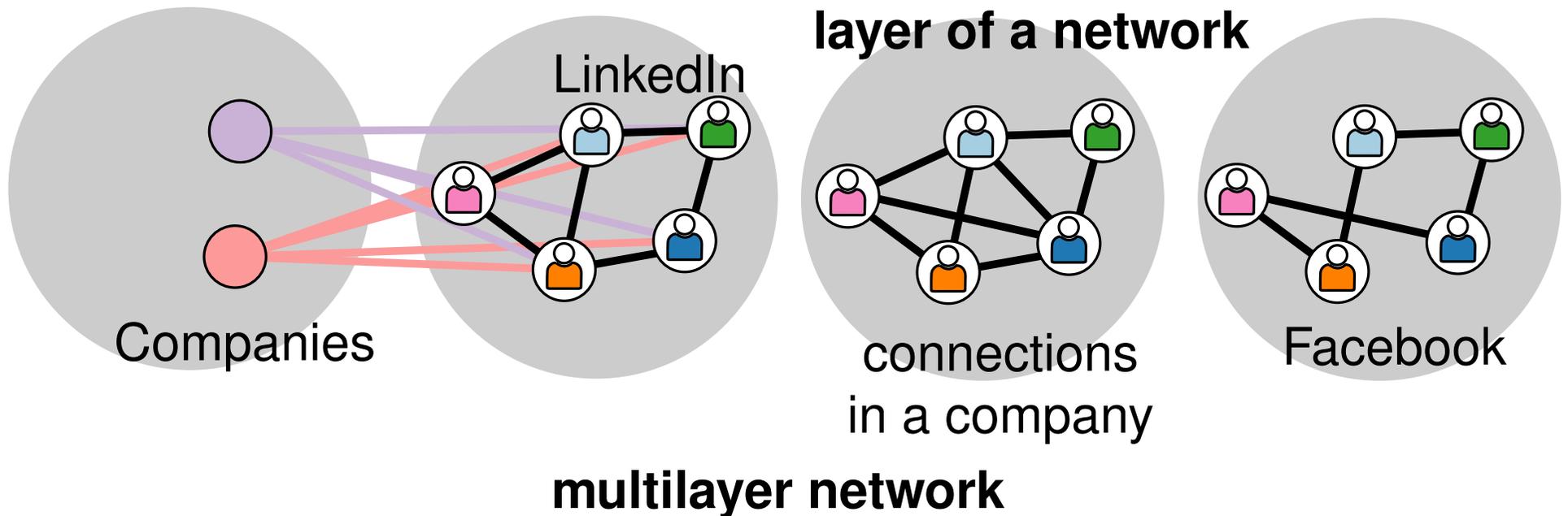
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- changes in one network effect changes in the other
- nodes of other types
- analysis of graph patterns across layers reveal complex facts about the data

# Multilayer Network

- Standard graph definition  $G = (V, E)$ ,  $E \subseteq V \times V$

# Multilayer Network

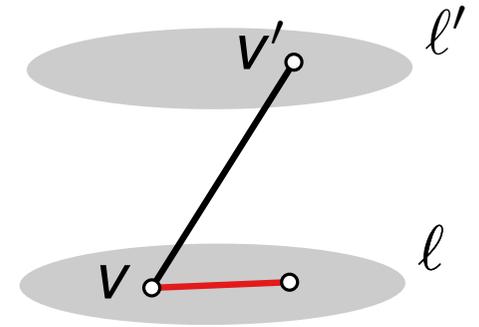
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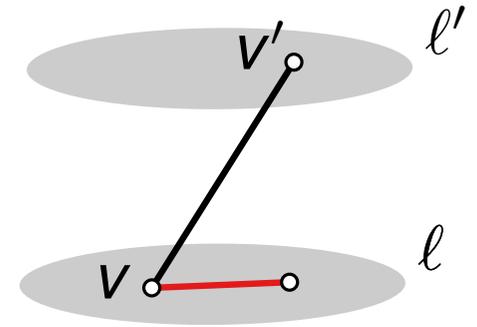
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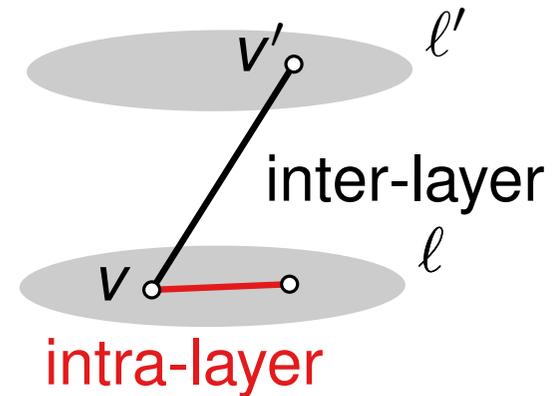
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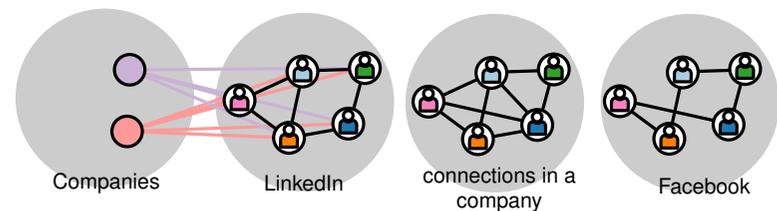
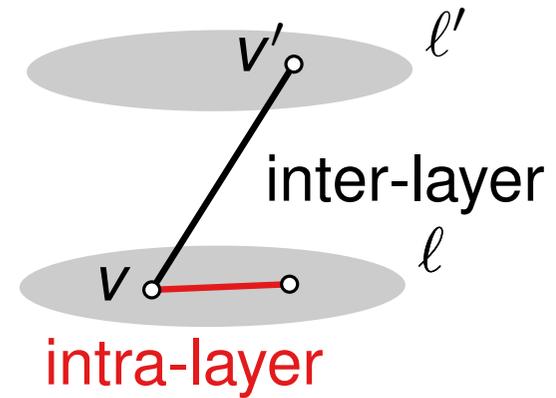
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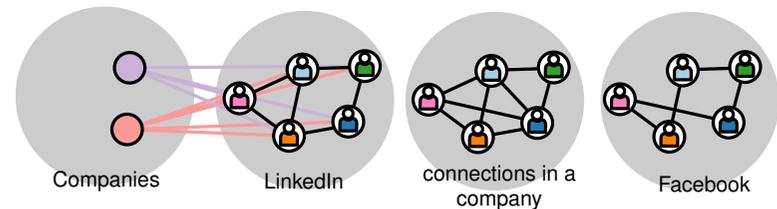
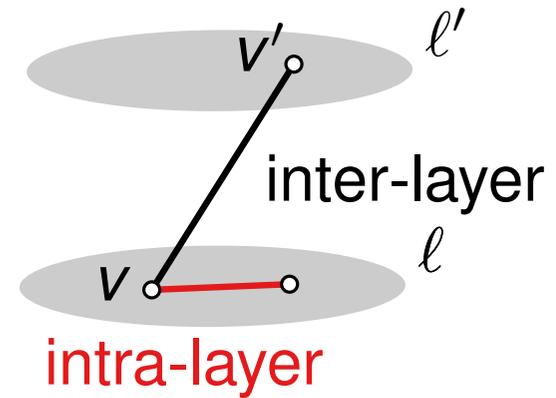
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- here layers are  $\{l_1, l_2, l_3, l_4\}$ ,  
 $l_1$  =LinkedIn,  
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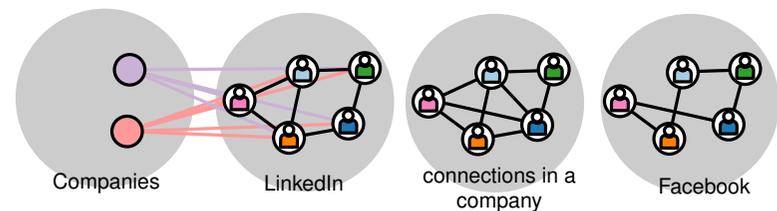
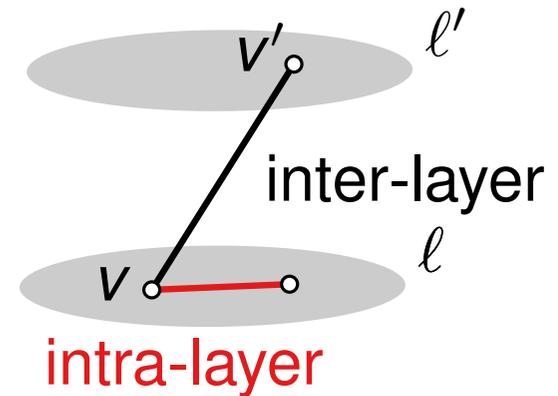
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- node  $v = \text{👤}$  appears as  $(v, \ell_1)$ ,  $(v, \ell_2)$ ,  $(v, \ell_3)$  in  $V_m$

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multilayer networks appear as models in

- biology: genomic, proteomic and metabolomic data to model intricate biological processes

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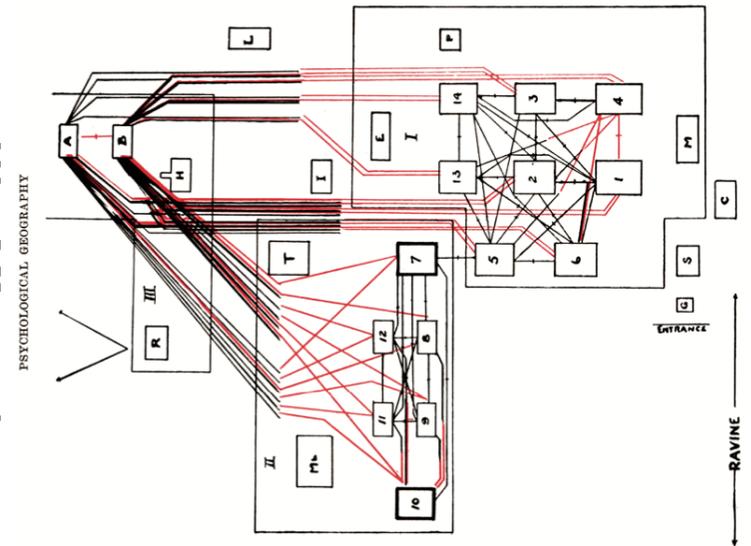
- biology: genomic, proteomic and metabolomic data to model intricate biological processes
- civil infrastructure: urban planning taking into account the interplay between multiple networks such as transportation networks, energy networks, telecommunication networks and water/wastewater networks
- epidemiology, sociology (including criminology), digital humanities

# Multilayer Network

multilayer networks appear as mode

- biology: genomic, proteomic and model intricate biological processes
- civil infrastructure: urban planning, the interplay between multiple networks, transportation networks, energy and telecommunication networks and networks

date back to 50s – notion of many relationships between individuals in the sociograms introduced by Moreno

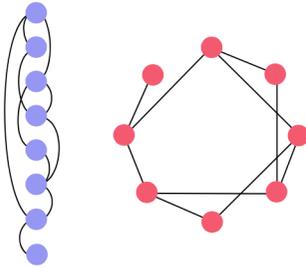






# Multilayer Network Visualizations

Types of visualizations of multilayer networks

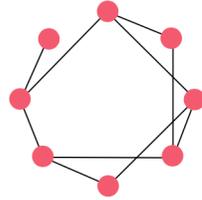
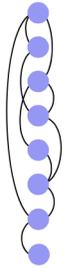


1-dimensional: circular,  
linear

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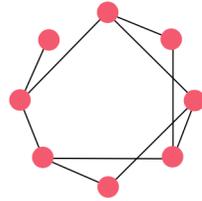
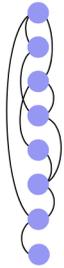


Gestalt principle:  
continuation, continuity

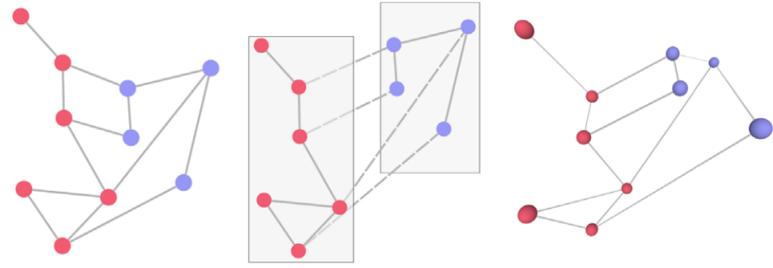
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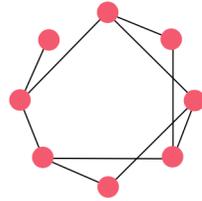
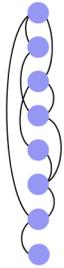


2D, 2.5D, 3D  
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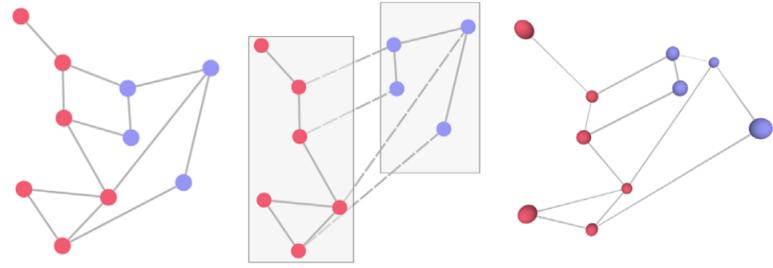
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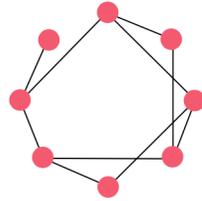
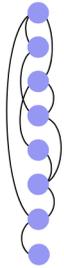


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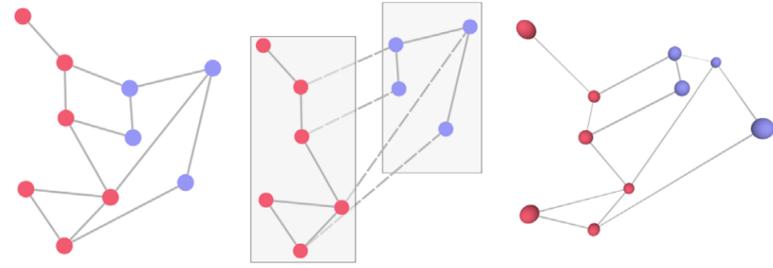
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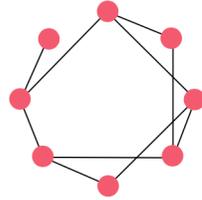
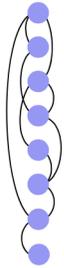


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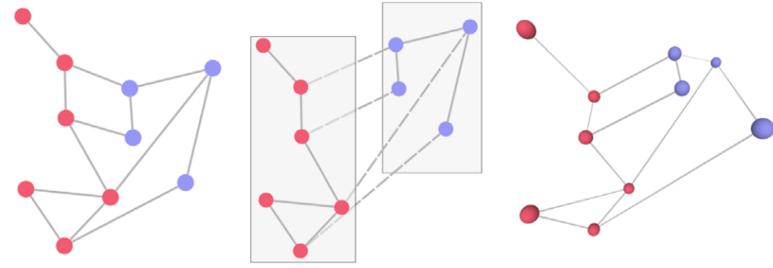
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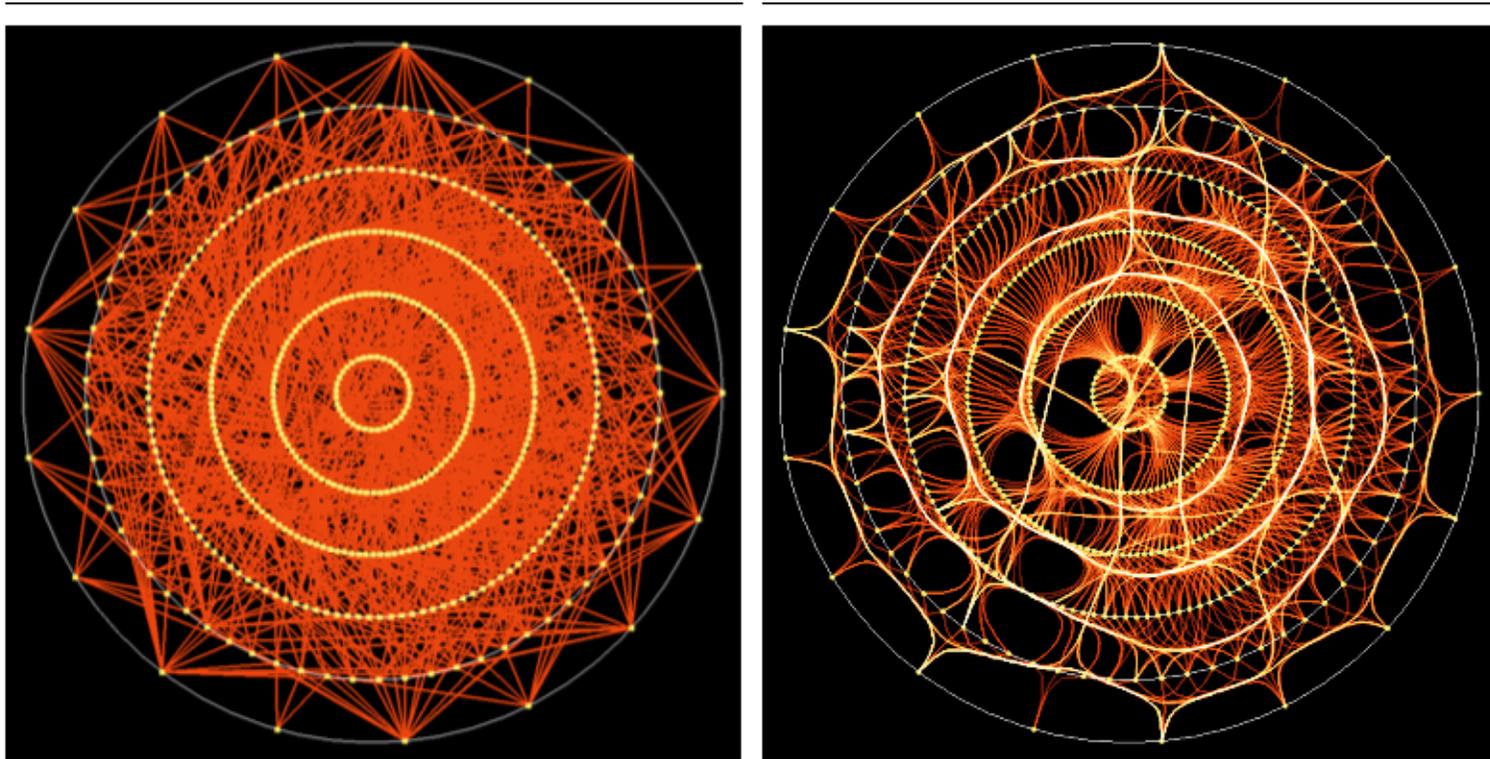


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- 2D - layers indicated by separation (proximity Gestalt principle), color (similarity)
- 2.5D - layers are are different planes stacked next to each other
- 3D – depth is indicating the layer, camera movement is necessary

# 1-dimensional representation: circular

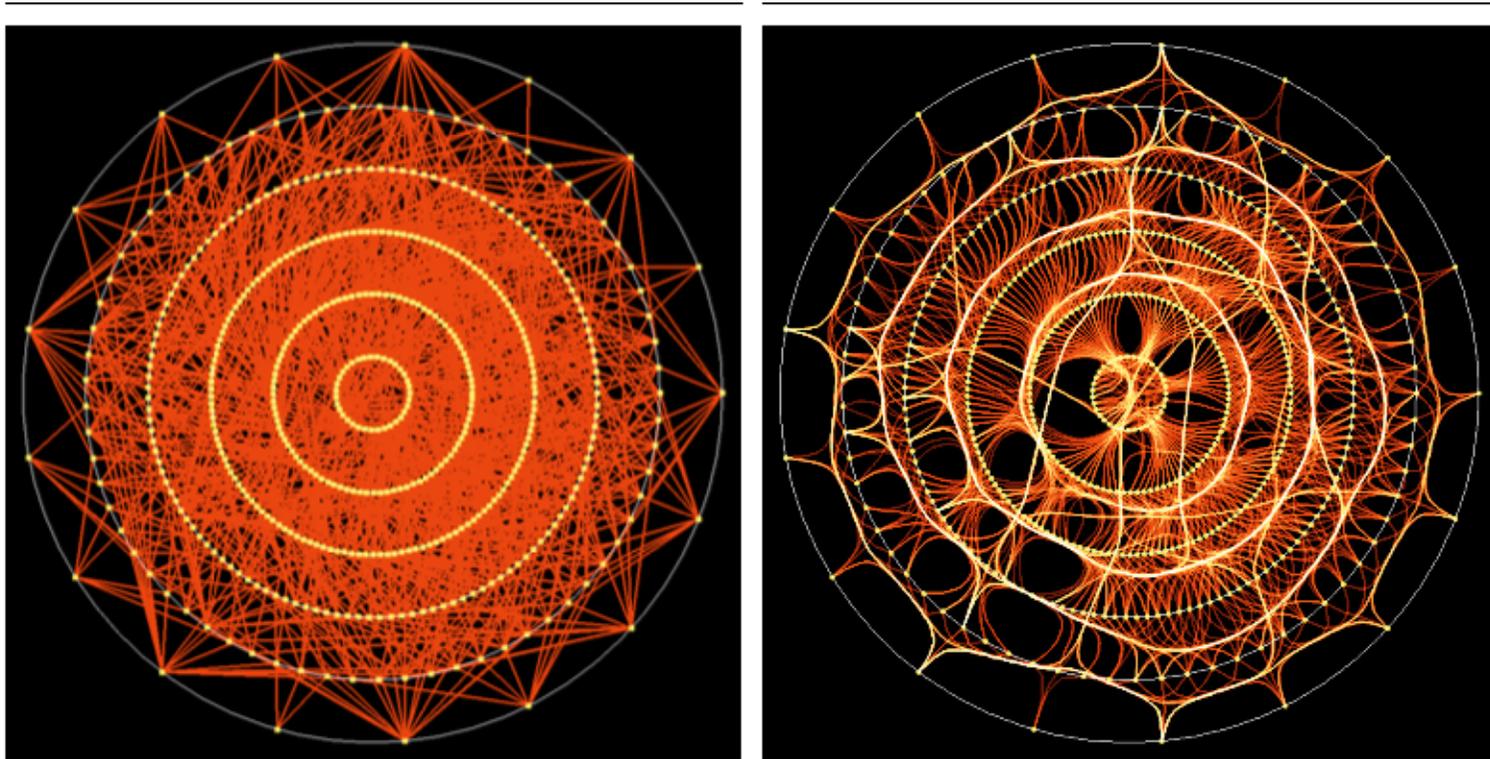
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Visualization of Frequent Itemsets with Nested Circular Layout and Bundling Algorithm, Bothorel et al 2013

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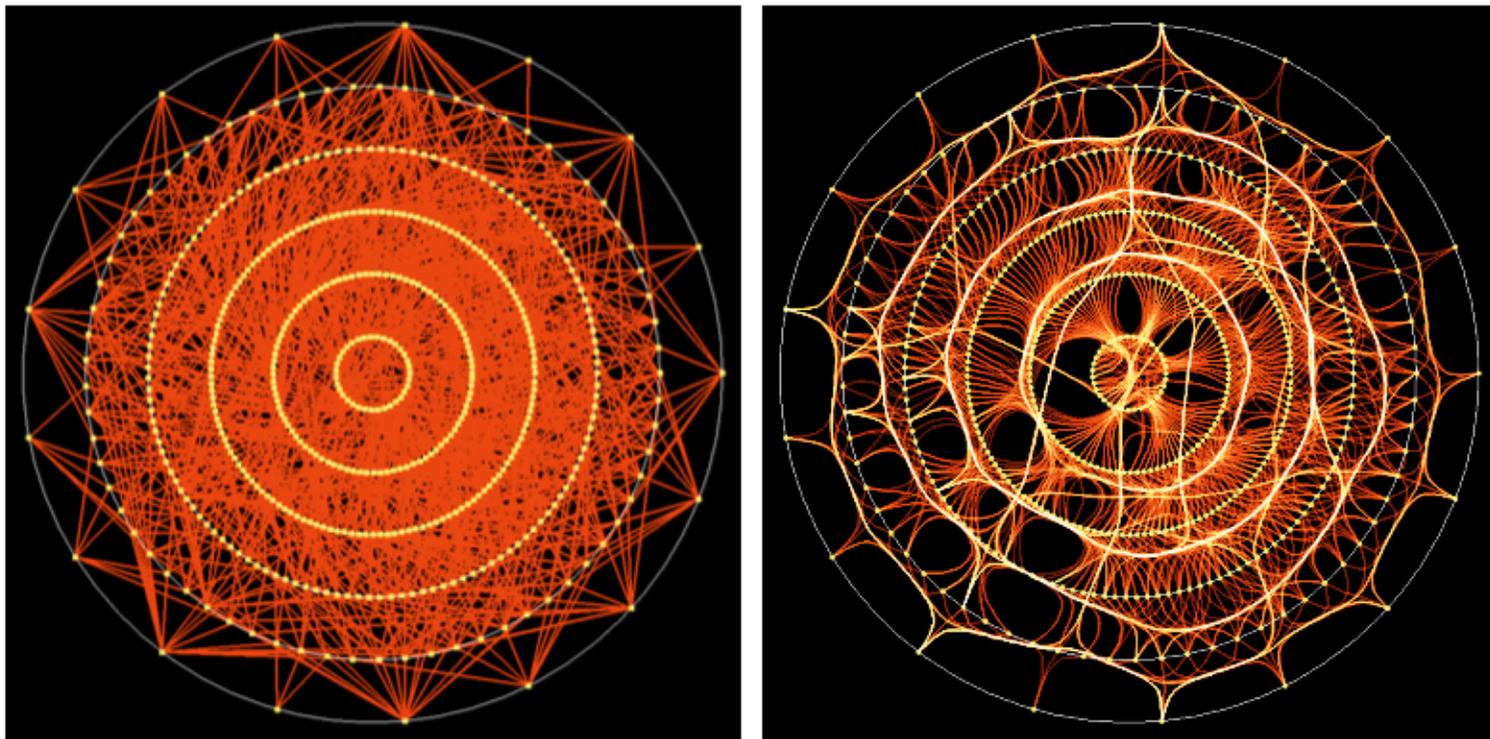
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- nodes – mushrooms
- layers – attributes: the edibility, the cap shape, the odor, the ring, etc.



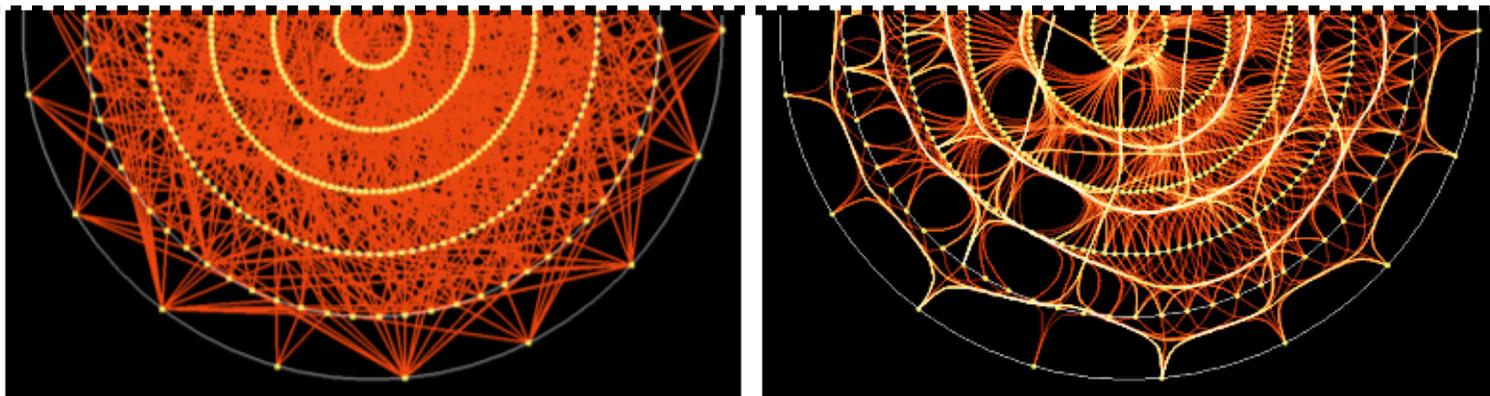
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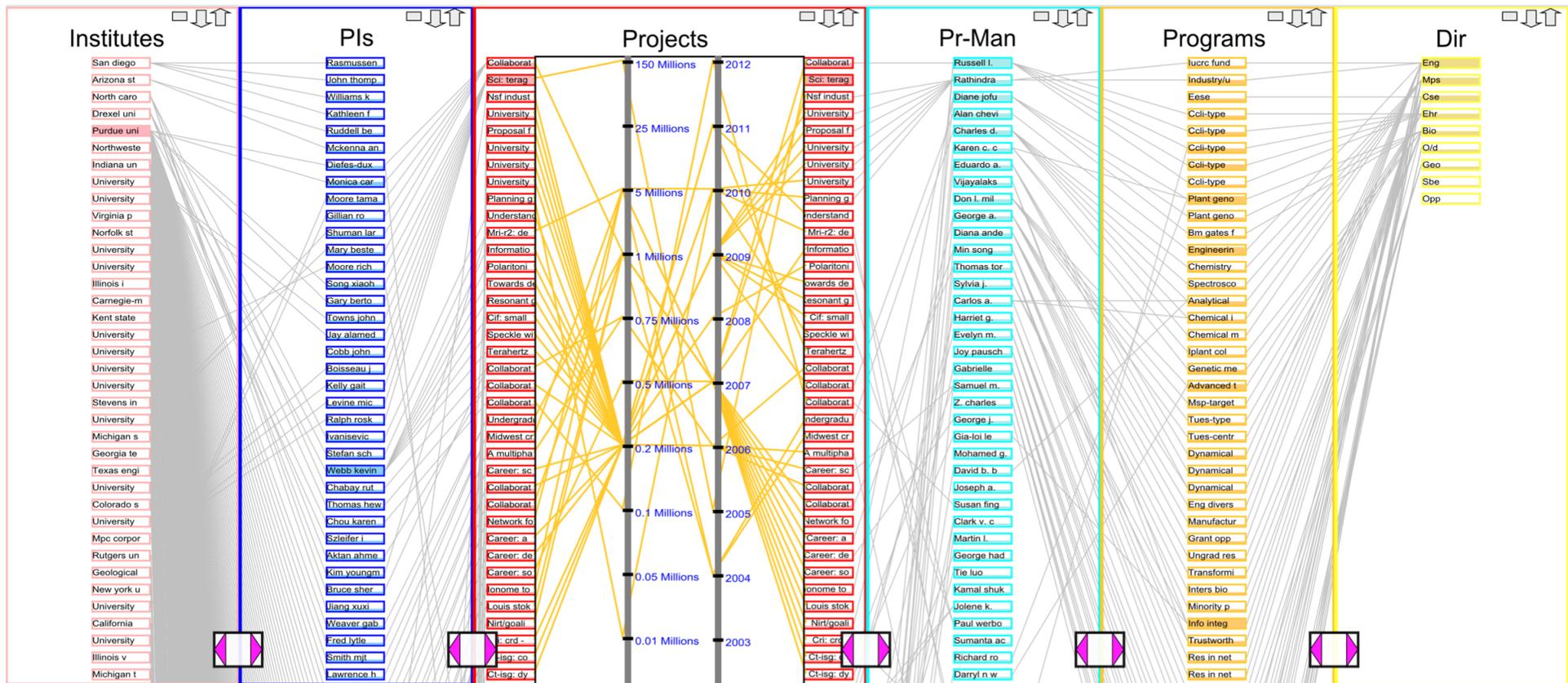
Which of the techniques you know can you use to construct this layout?



Visualization of Frequent Itemsets with Nested Circular Layout and Bundling Algorithm, Bothorel et al 2013

# 1-dimensional representation: linear

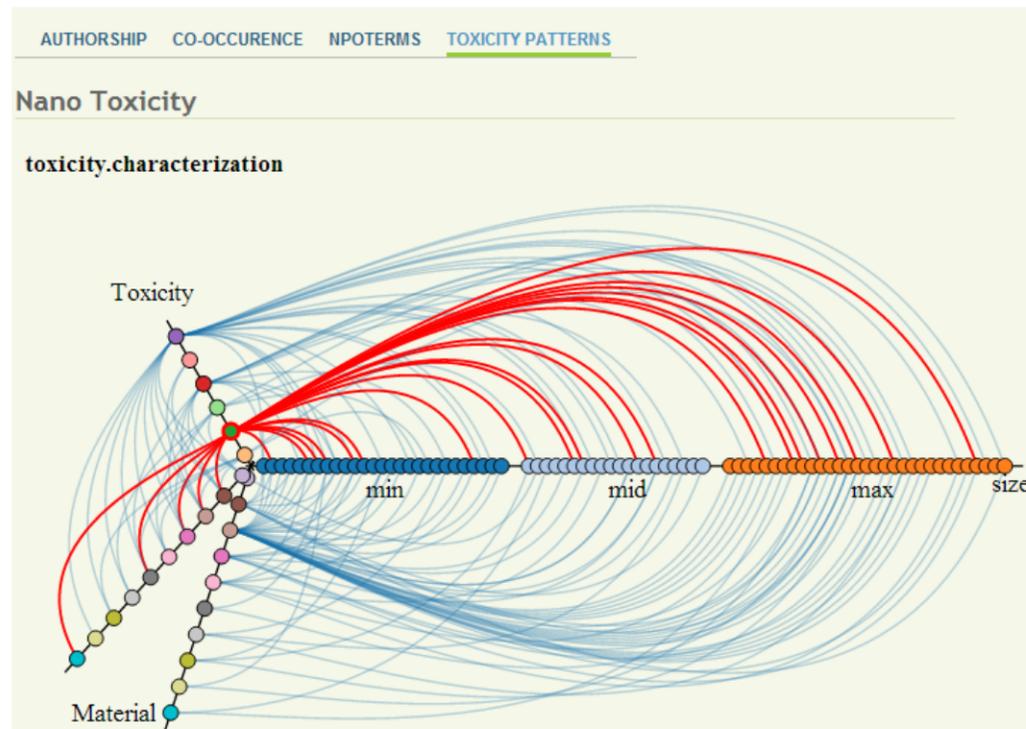
- multimodal NSF funding data consisting of Institutions, PIs (and Co-PIs), Projects, program managers (Pr-Man), NSF programs (Programs), and NSF directorates (Dir)
- remind parallel coordinate plots



Visual Analytics for Multimodal Social Network Analysis: A Design Study with Social Scientists, Ghani et al, 2013

# 1-dimensional representation: linear

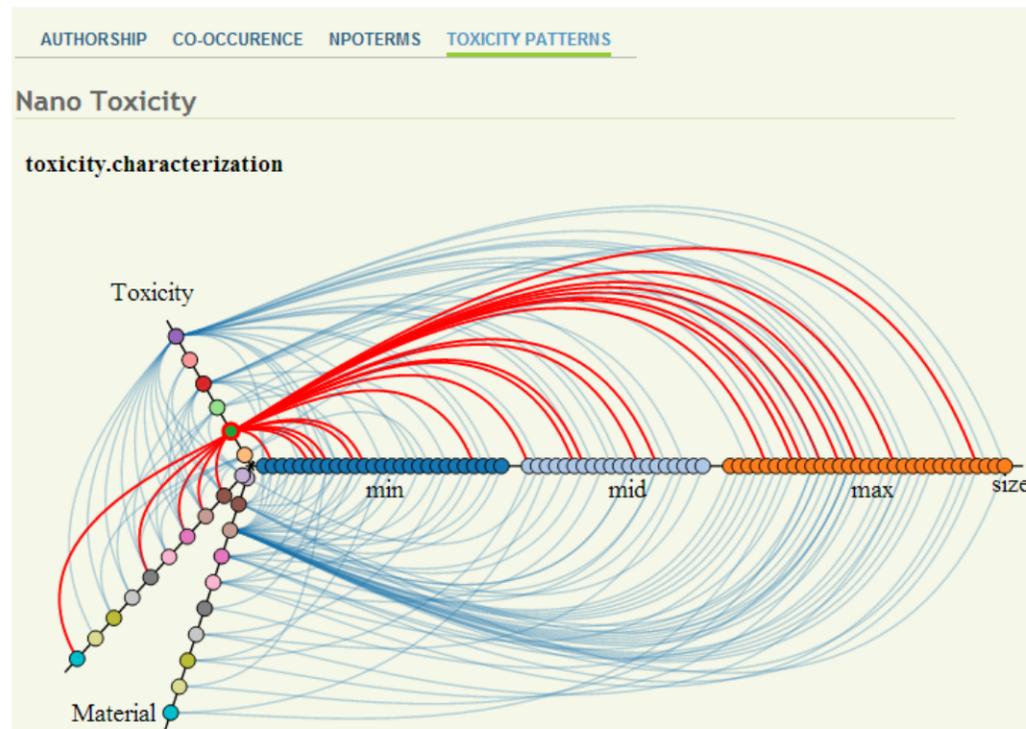
- Hive plot: axes are arranged radially
- investigation among nano-toxicity type, nanomaterial and particle size



A user-centred approach to information visualisation in nano-health,  
Yang et al, 2016

# 1-dimensional representation: linear

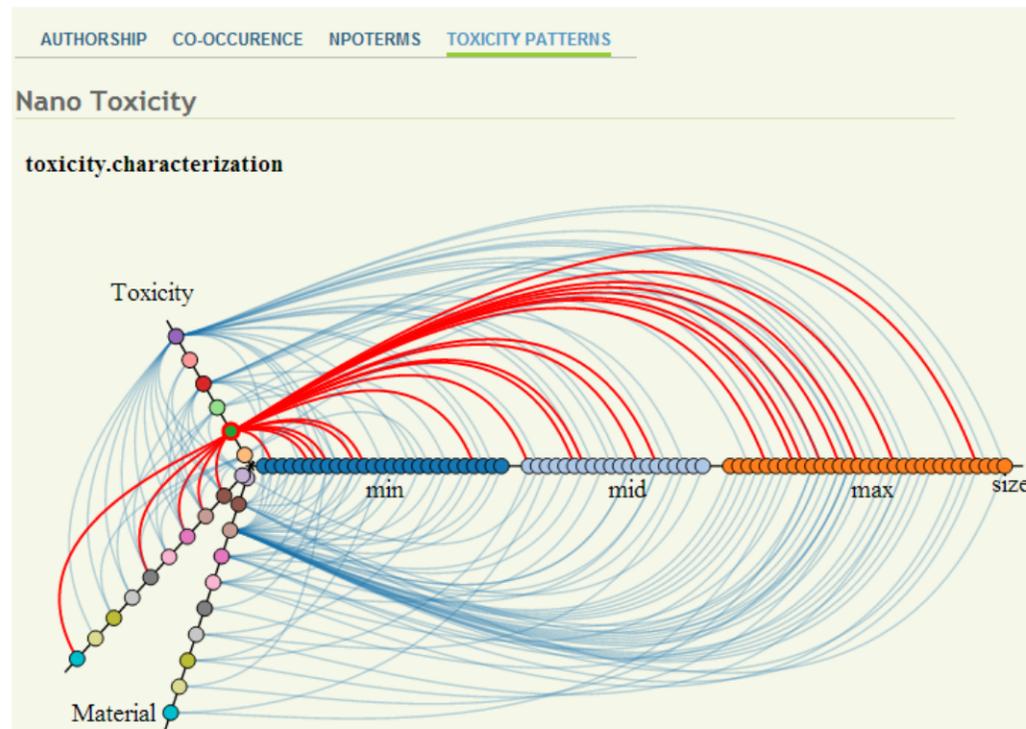
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- investigation among nano-toxicity type, nanomaterial and particle size
- edges are displayed only between adjacent layers



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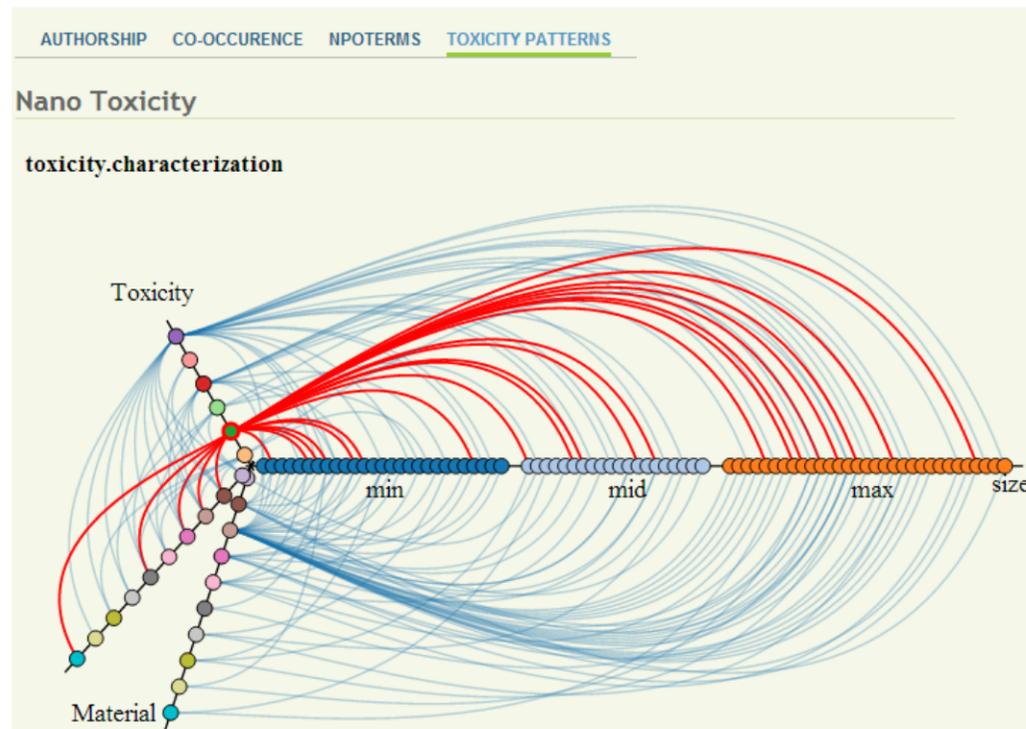
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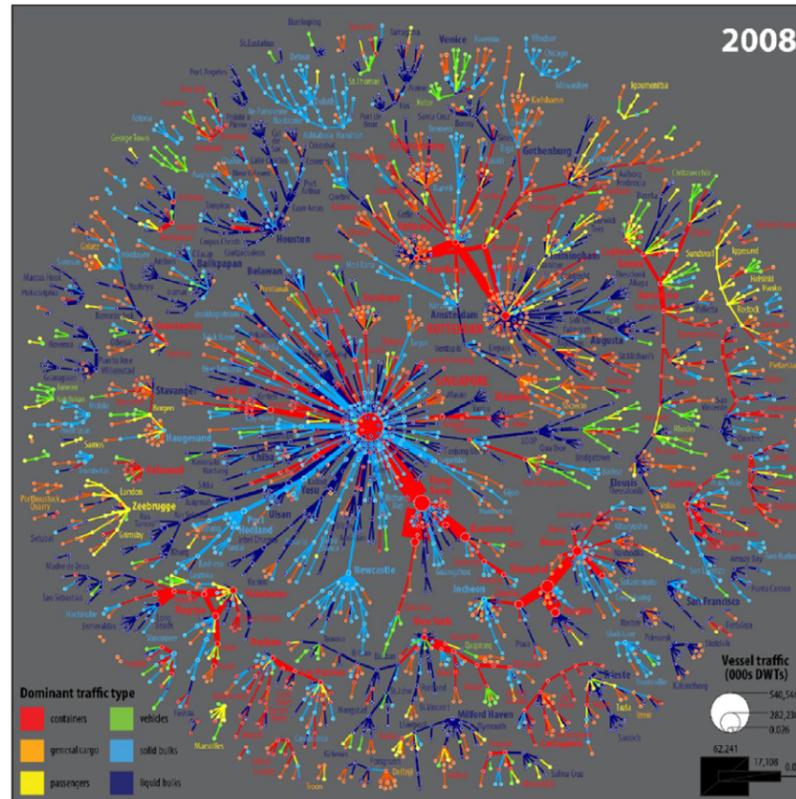
- Hive plot: axes are arranged radially
- investigation among nano-toxicity type, nanomaterial and particle size
- edges are displayed only between adjacent layers
- nodes are arranged based on graph metric (e.g. degree)
- reduce clutter using layer duplication



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Yang et al, 2016

# 2-dimensional representaiton: color

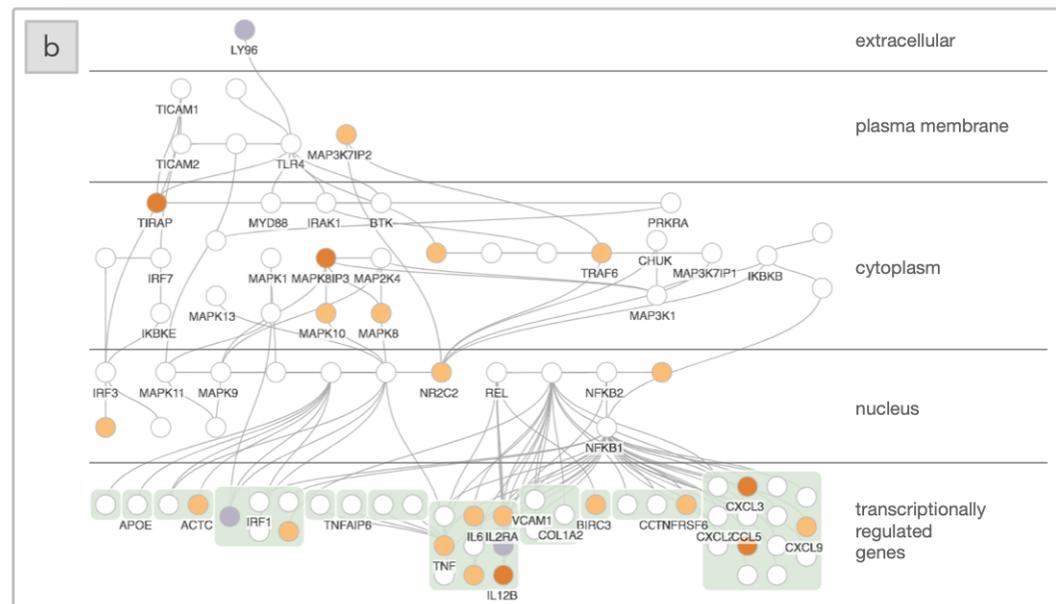
- flow of maritime traffic: nodes represent ports and different edge colours represent different modes of shipping



Multilayer dynamics of complex spatial networks: The case of global maritime flows, Ducuet, 2017

# 2-dimensional representation: separation

- use constrained layouts to separate the nodes of different layers spatially

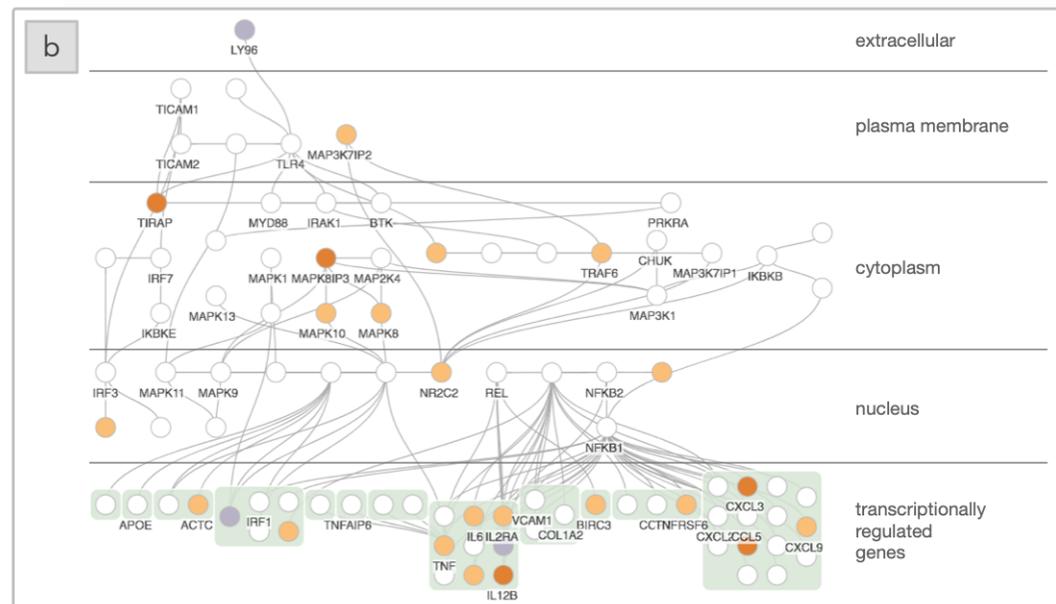


nodes – physical compounds in a cell; that are separated by physical membranes, creating compartments defining their subcellular location  
– layered edge; edges interactions among nodes

SetCoLa: High-Level Constraints for Graph Layout, Hoffswell et al, 2018

# 2-dimensional representation: separation

- use constrained layouts to separate the nodes of different layers spatially
- not intended for multilayer networks but can be used for them

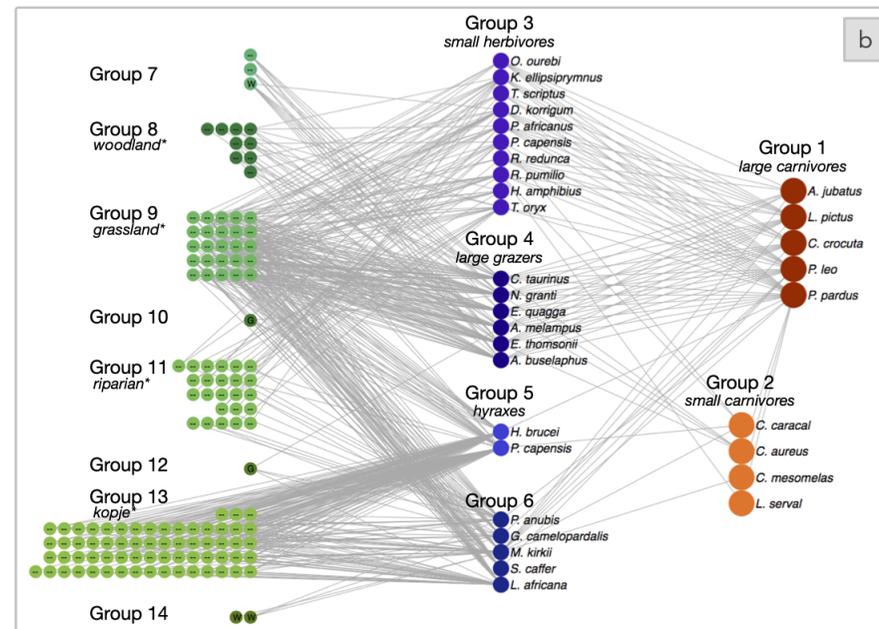


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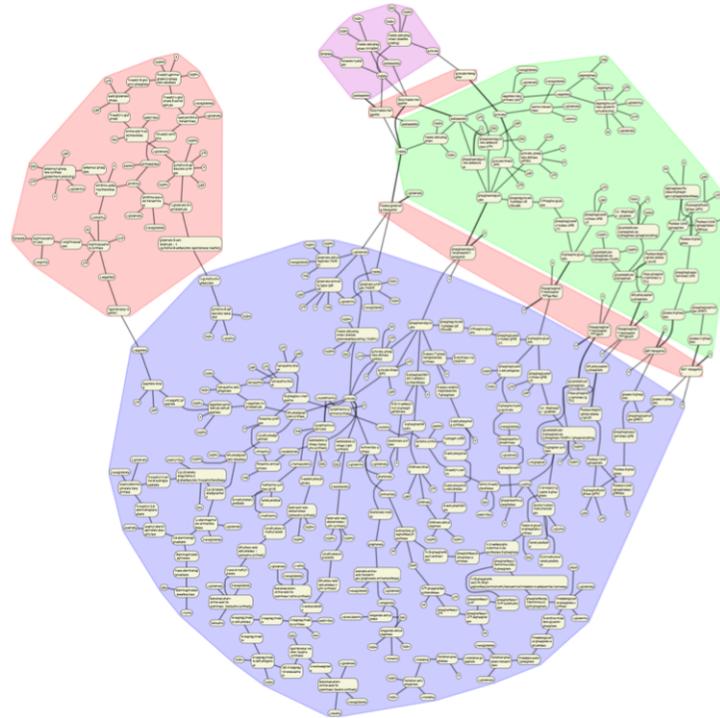


161 plants, herbivores, and carnivores with 592 links between entities  
– feeding links, groups – clustering, layers – trophic hierarchy

SetCoLa: High-Level Constraints for Graph Layout, Hoffswell et al, 2018

# 2-dimensional representation: separation

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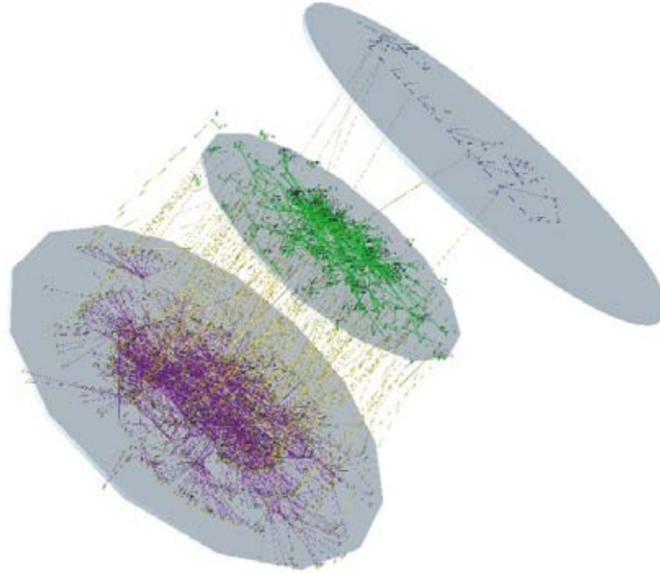


Biological pathways: nodes – proteins, edges–interactions. Rather visualization of clusters, but can be used to show layers too.

Scalable, Versatile and Simple Constrained Graph Layout, Dwyer 2009

## 2.5-dimensional representation

- each layer is drawn on a plane and planes are stacked in 3D parallel to each other
- use 2D layout algorithms for a single layer
- same node can appear on many layers – similar positions are desired. Same for reducing edge clutter.

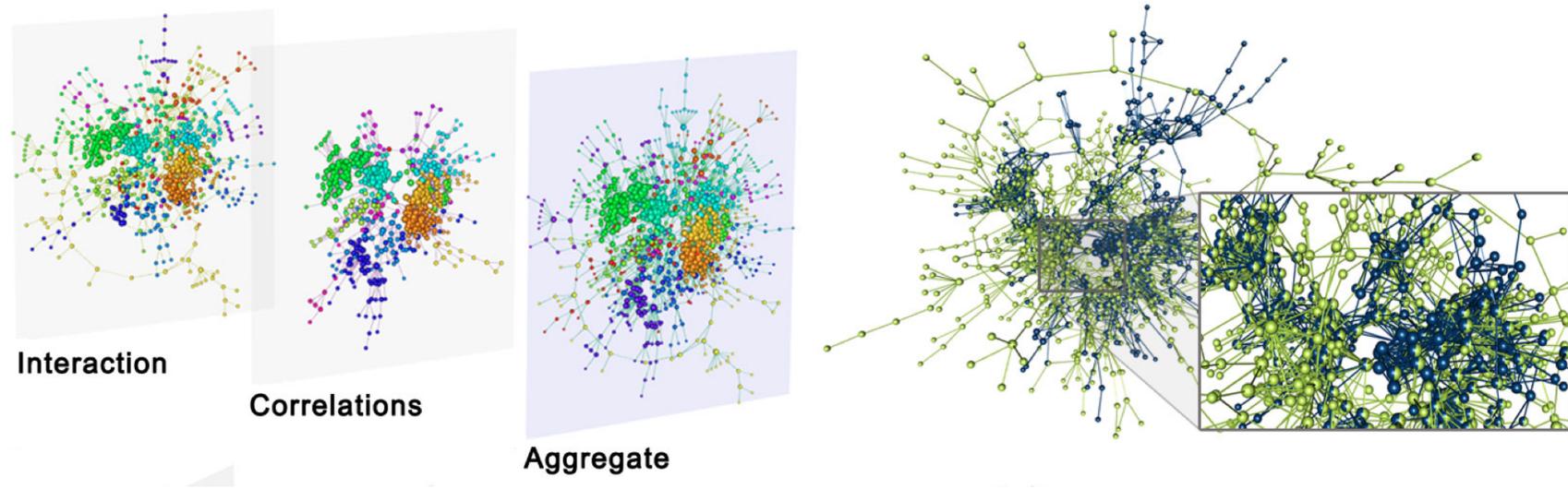


metabolic network, protein interaction networks and gene regulatory network; inter-layer edges: proteins are the result of gene expression, special proteins known as enzymes help transforming metabolites to another.

Visual Analysis of Overlapping Biological Networks, Fung et al, 2009

# 2.5-dimensional representation

- same node can appear across layers and lie at the same position
- aggregated layer is possible

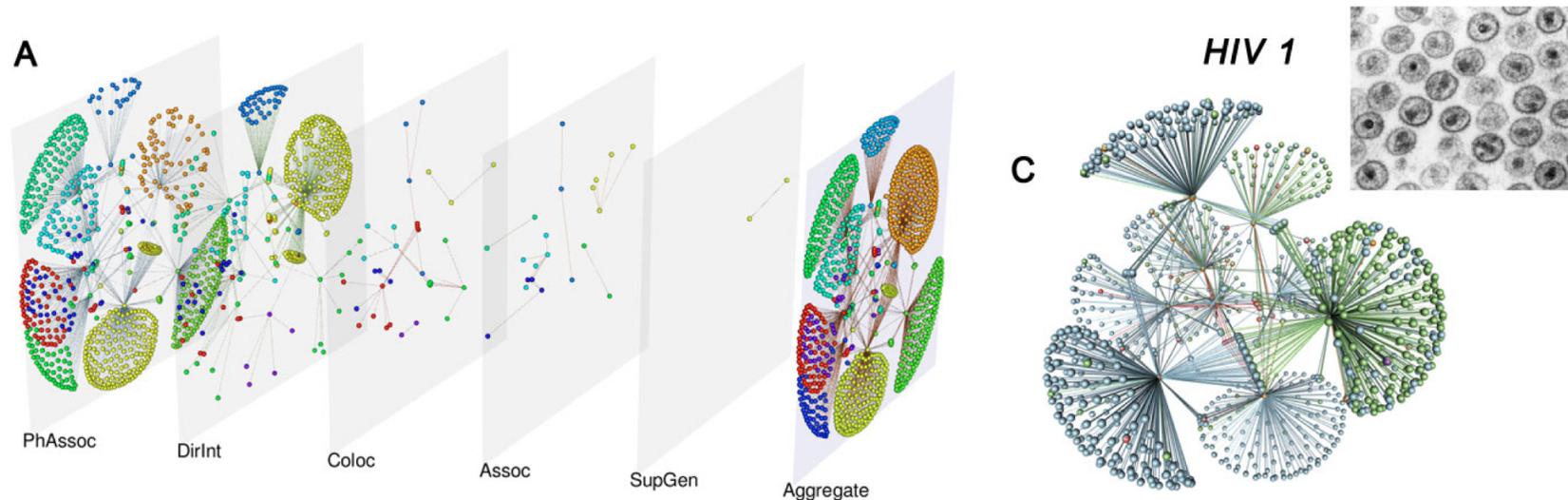


- first layer: interaction of genes in *Saccharomyces cerevisiae*;
- second layer: genes with similar interaction profiles are connected to each other;
- third layer: aggregated network
- right - edge colors represent layers

MuxViz: A Tool for Multilayer Analysis and Visualization of Networks,  
De Domenico et al, 2015.

# 2.5-dimensional representation

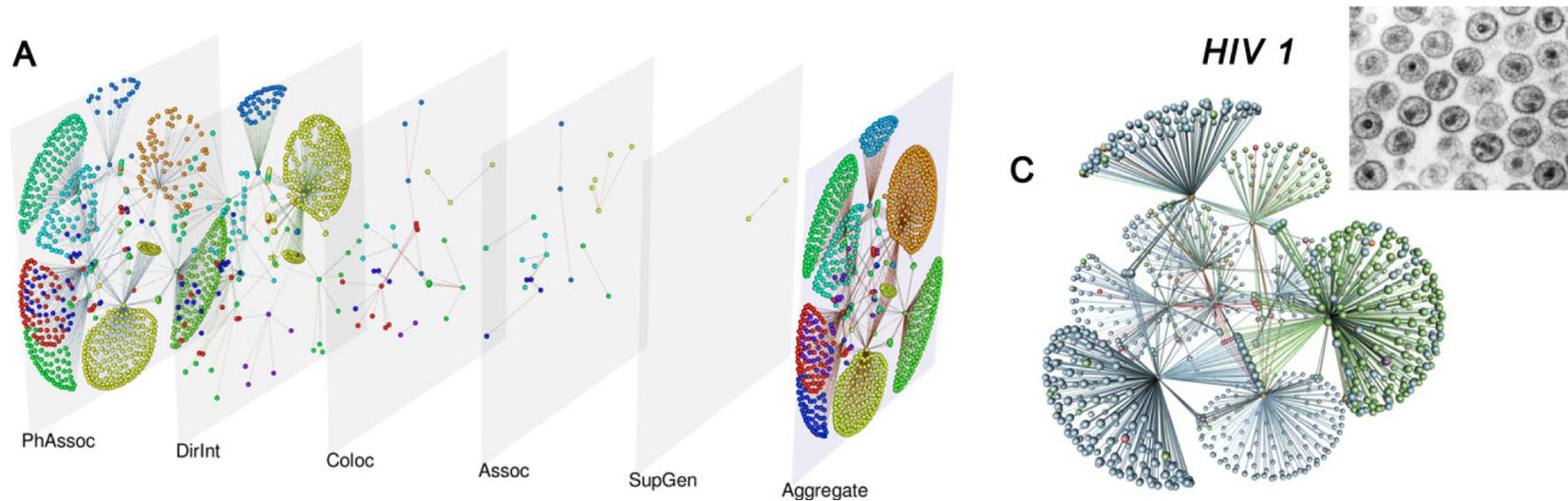
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- Multilayer analysis of HIV-1 genetic interaction network

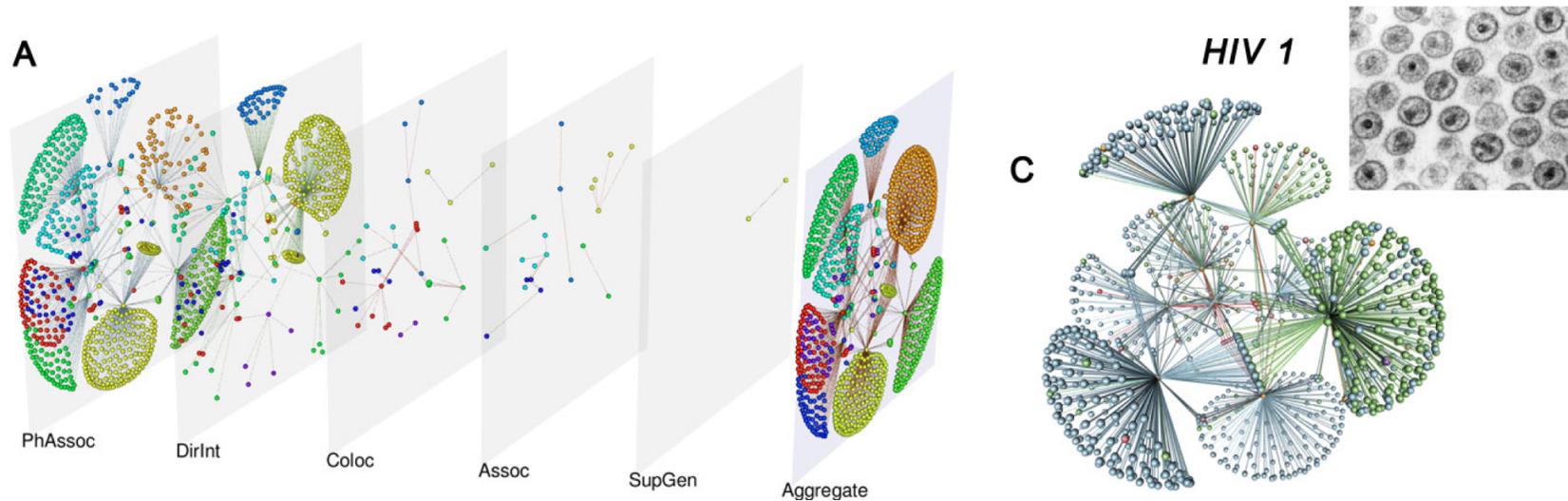
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# Generating 2.5D representations



- If it is essential that same node has exactly the same position over the layers

# Generating 2.5D representations

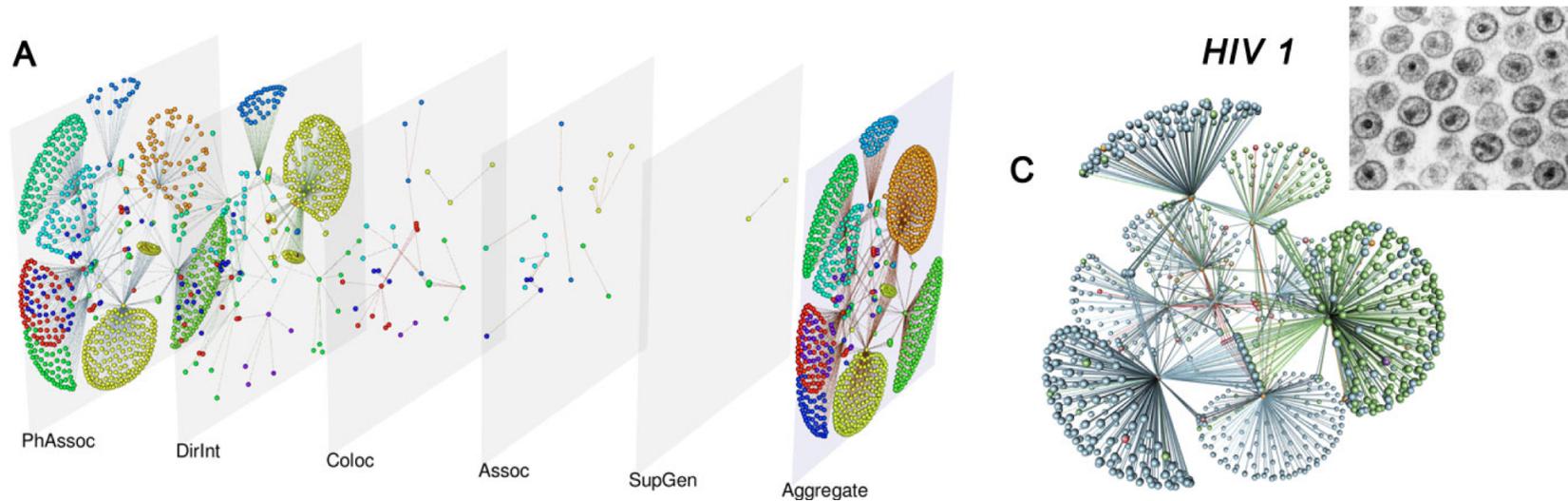


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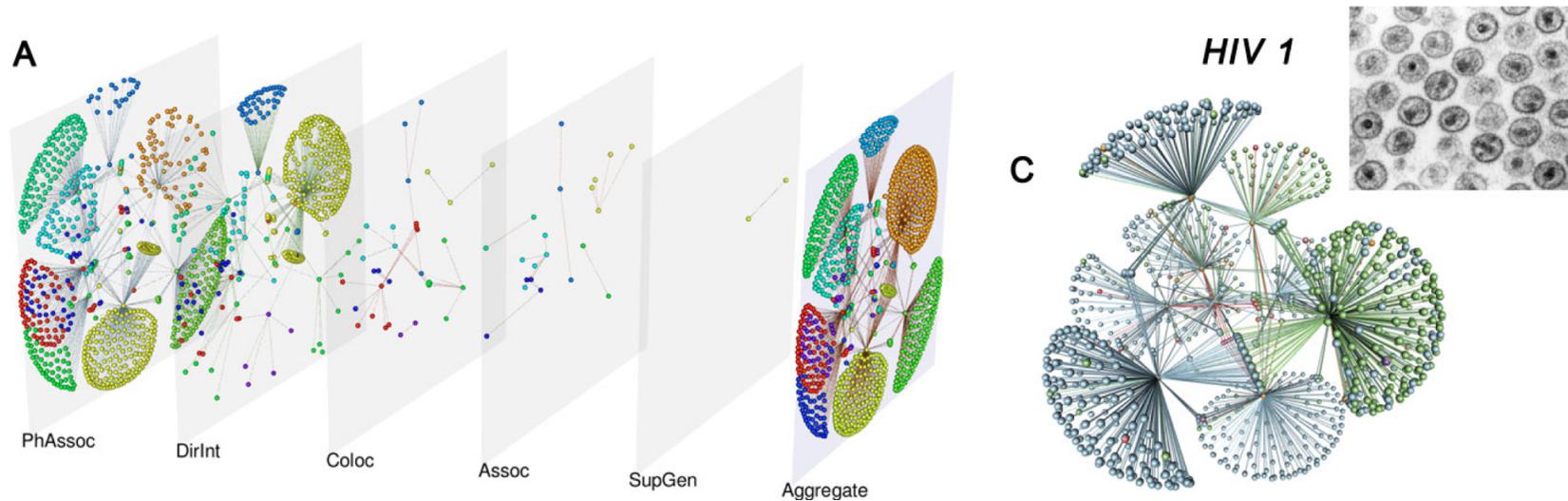
How to construct this representation?

# Generating 2.5D representations



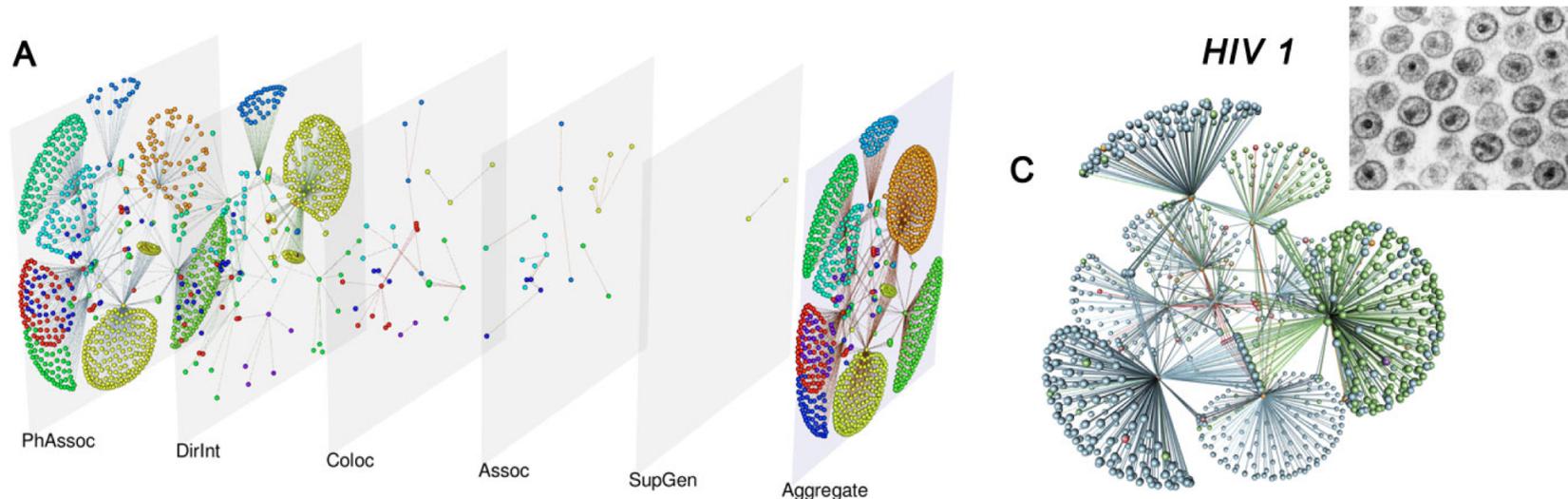
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# Generating 2.5D representations



- If it is essential that same node has exactly the same position over the layers
- Aggregate the graphs over the layers into a single graph  $G = (V, E)$
- Layout  $G$  with a favorite layout method – aggregated layer

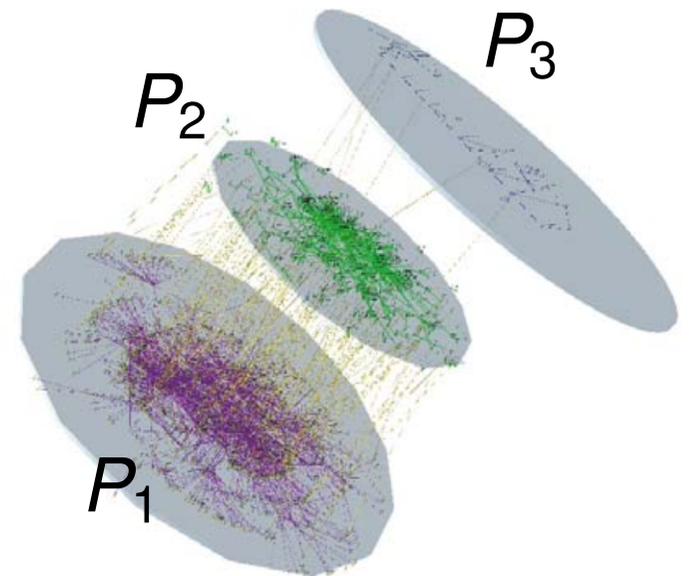
# Generating 2.5D representations



- If it is essential that same node has exactly the same position over the layers
- Aggregate the graphs over the layers into a single graph  $G = (V, E)$
- Layout  $G$  with a favorite layout method – aggregated layer
- Use coordinates of the nodes of  $G$  to construct the layouts of the rest layers

# Generating 2.5D representations Fung et al, 2009

- If it is not essential that same node has exactly the same position over the layers, or there are not many identical nodes over the layers

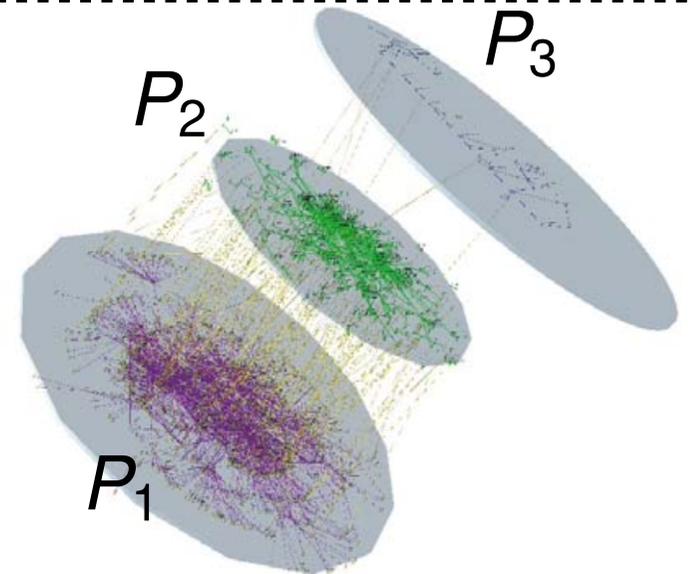


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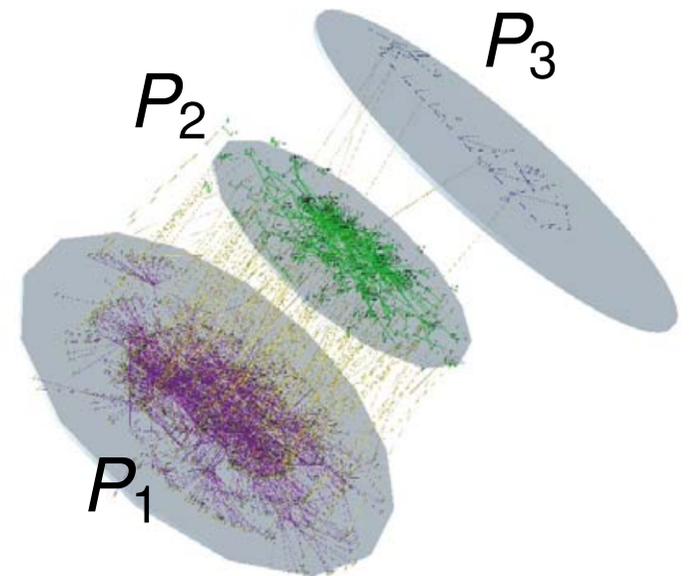


Same method as before is possible, but how to use the flexibility in node position in order to construct better layout?



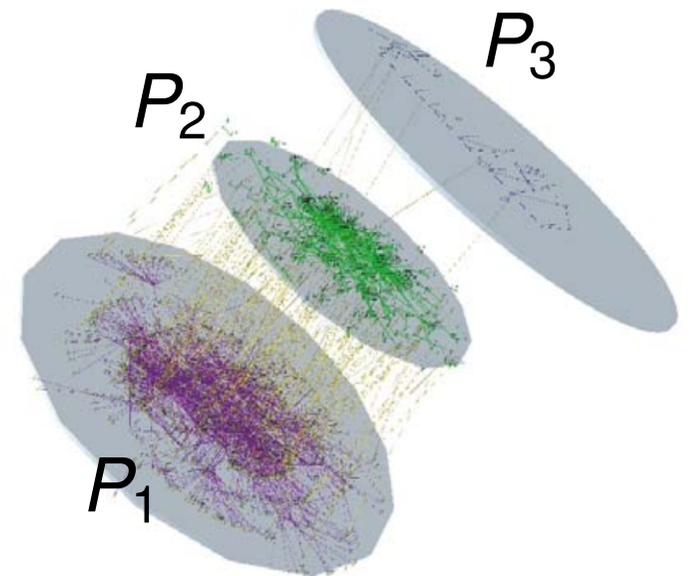
# Generating 2.5D representations Fung et al, 2009

- If it is not essential that same node has exactly the same position over the layers, or there are not many identical nodes over the layers
- Assume we have 3 layers  $\ell_1, \ell_2, \ell_3$ , let  $G_i$  be graph induced by  $\{(v, \ell_i) \in V_m : v \in V\}$ ,  $i = 1, 2, 3$



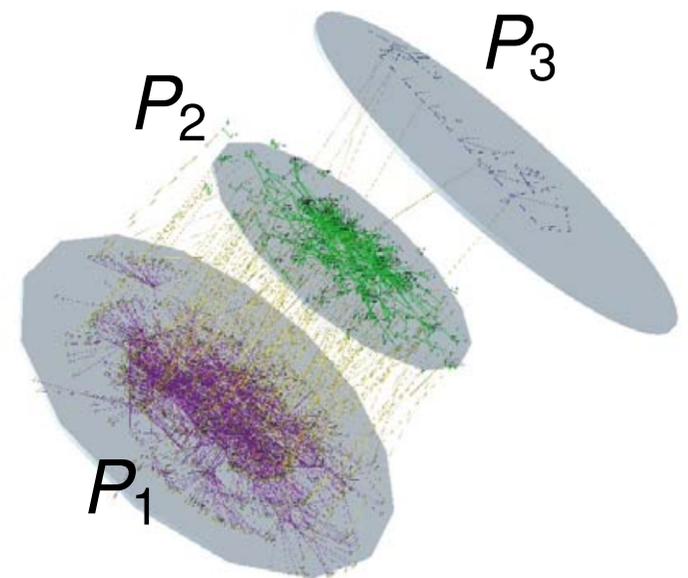
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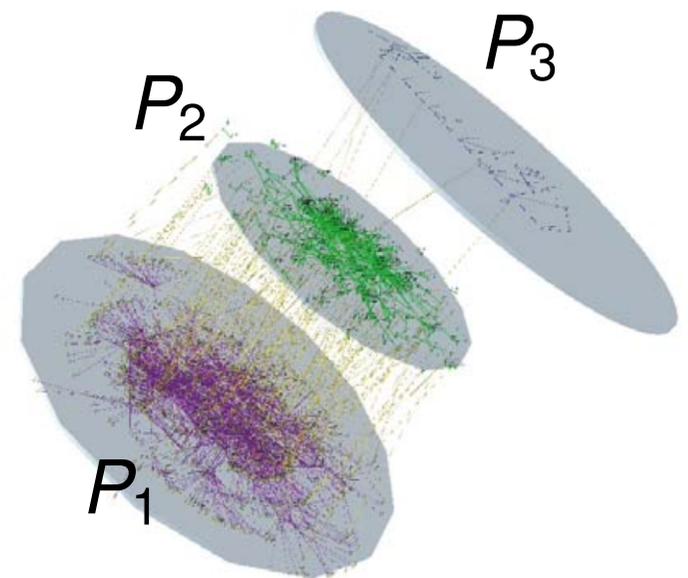
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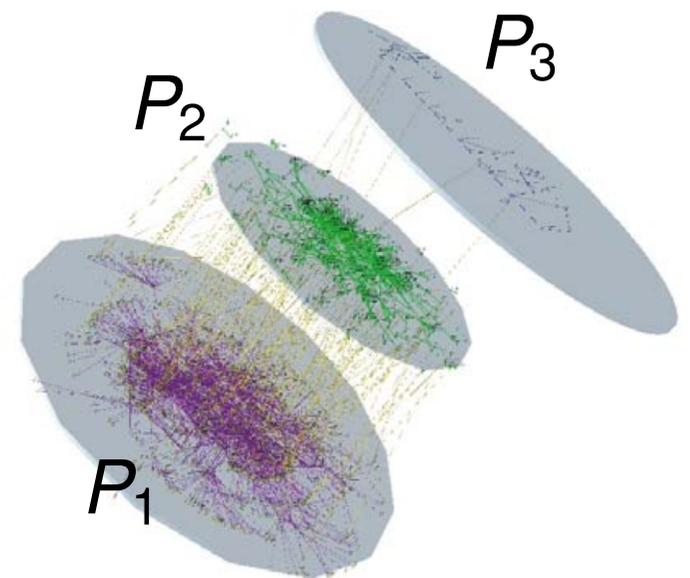
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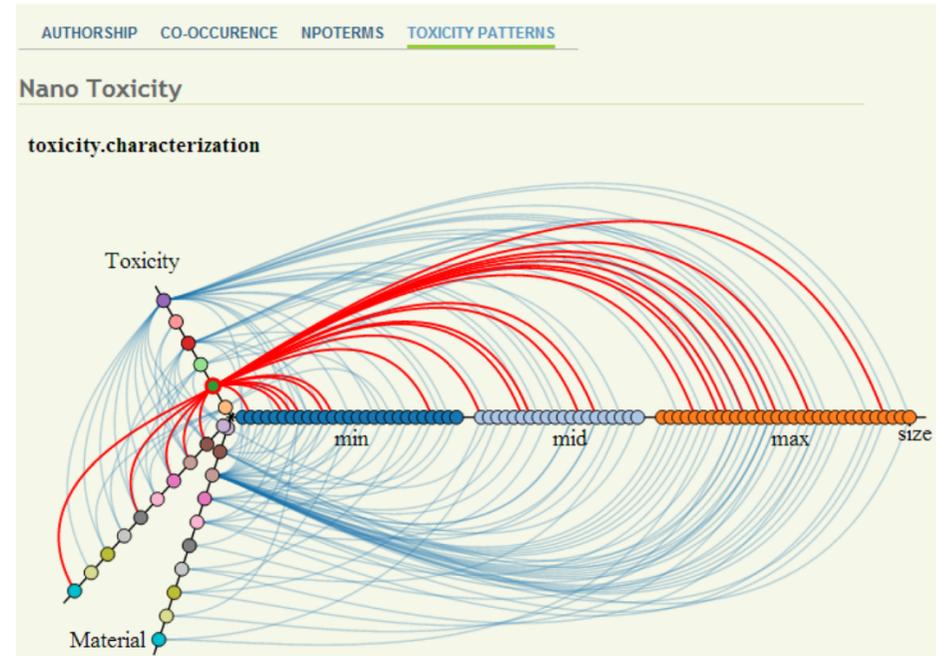
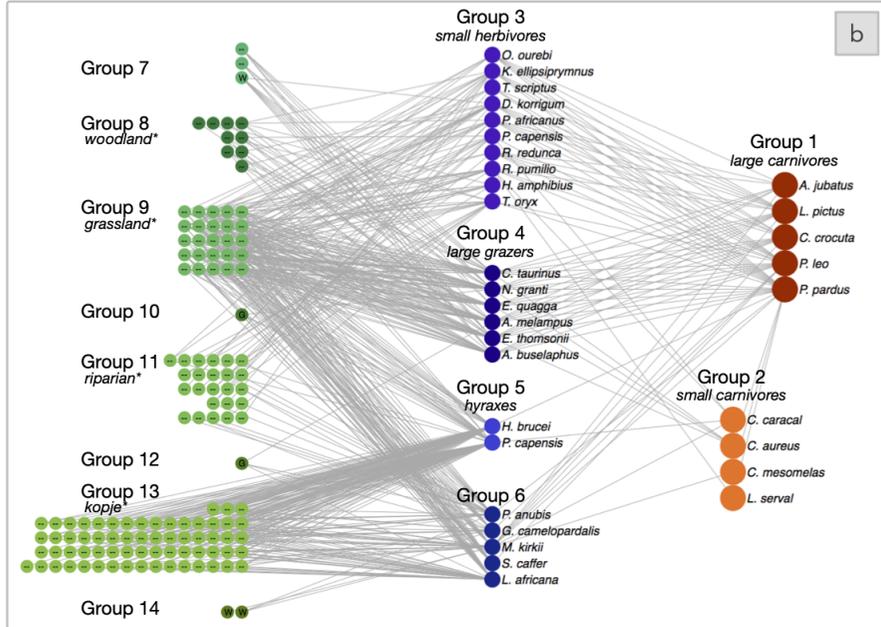


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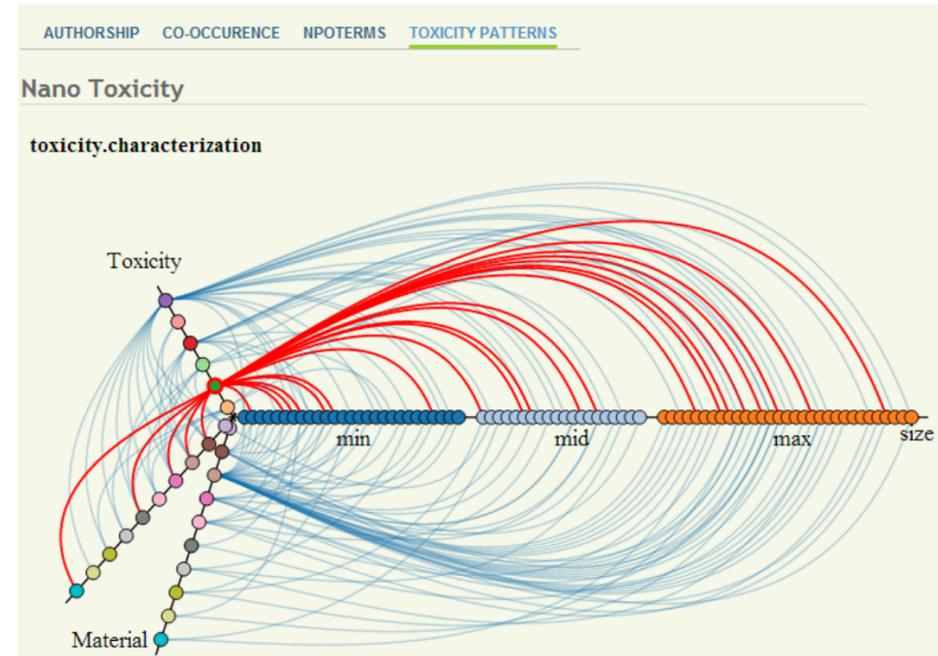
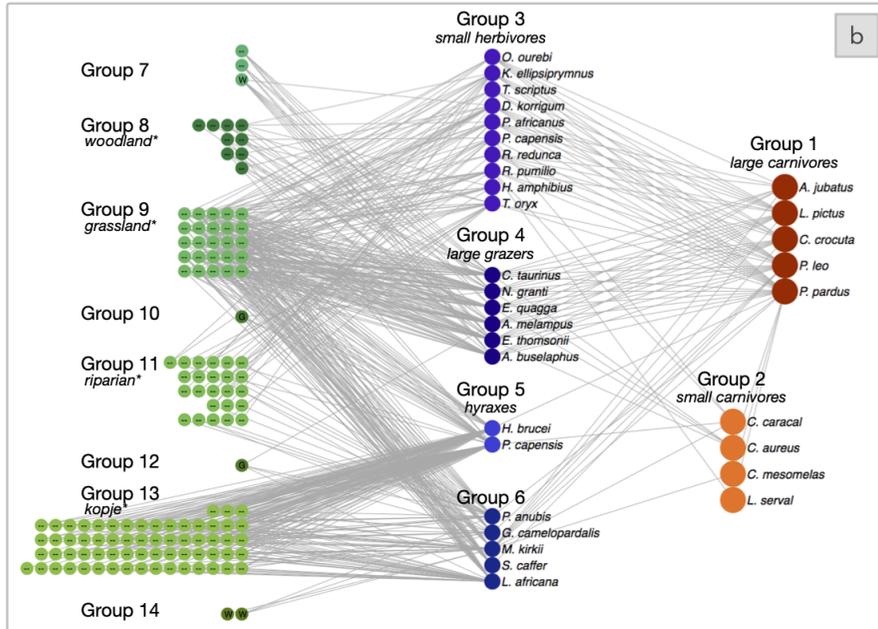
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- Model inter-layer edges as zero-length spring (attraction only)
- Draw  $G_2$  and the inter-layer edges using a force directed layout



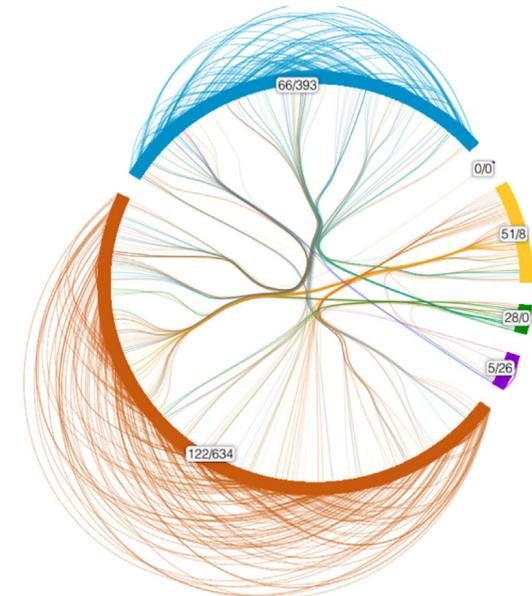
# Edge clutter in multilayer visualizations



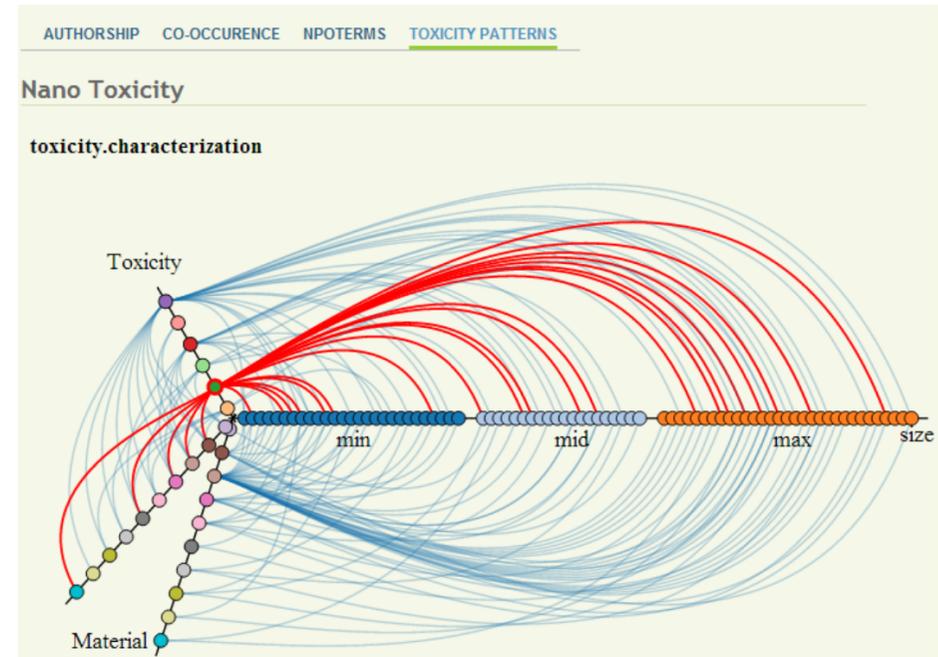
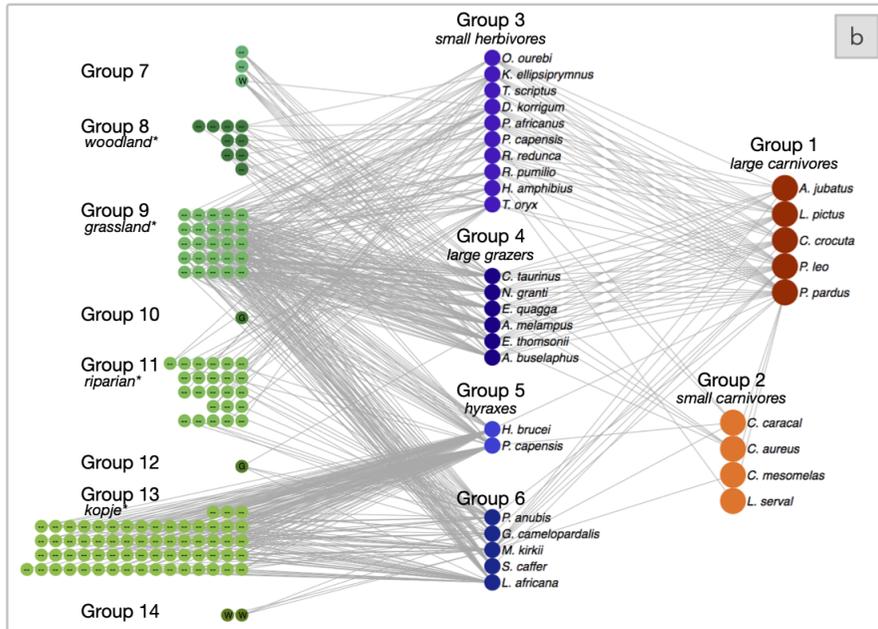
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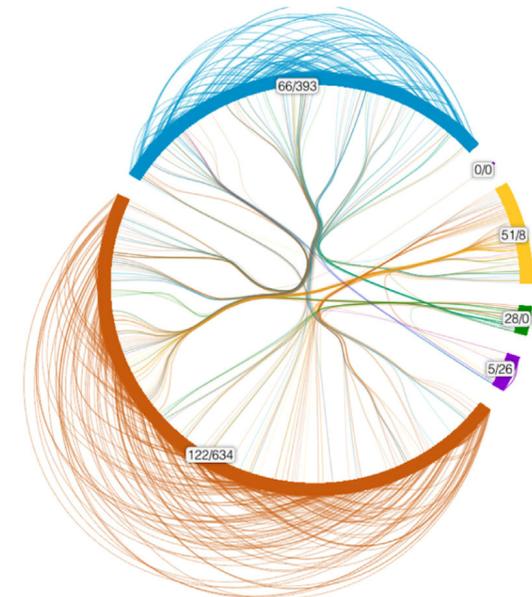
- edge bundling as a method to layout edges in multilayer network visualizations



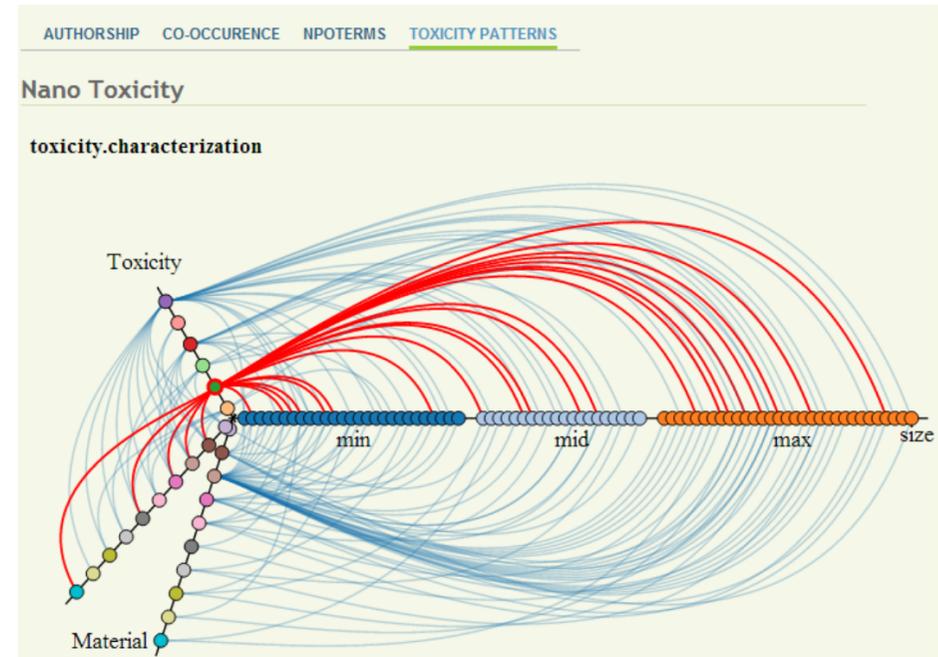
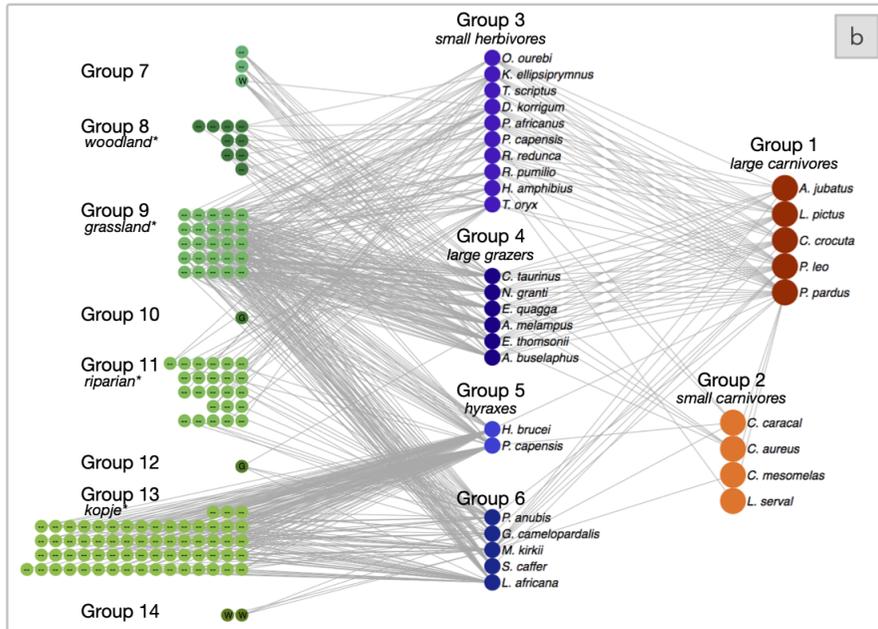
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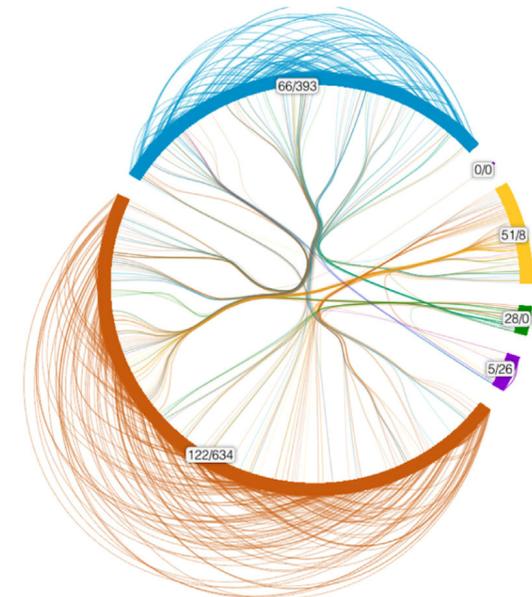
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# Edge clutter in multilayer visualizations



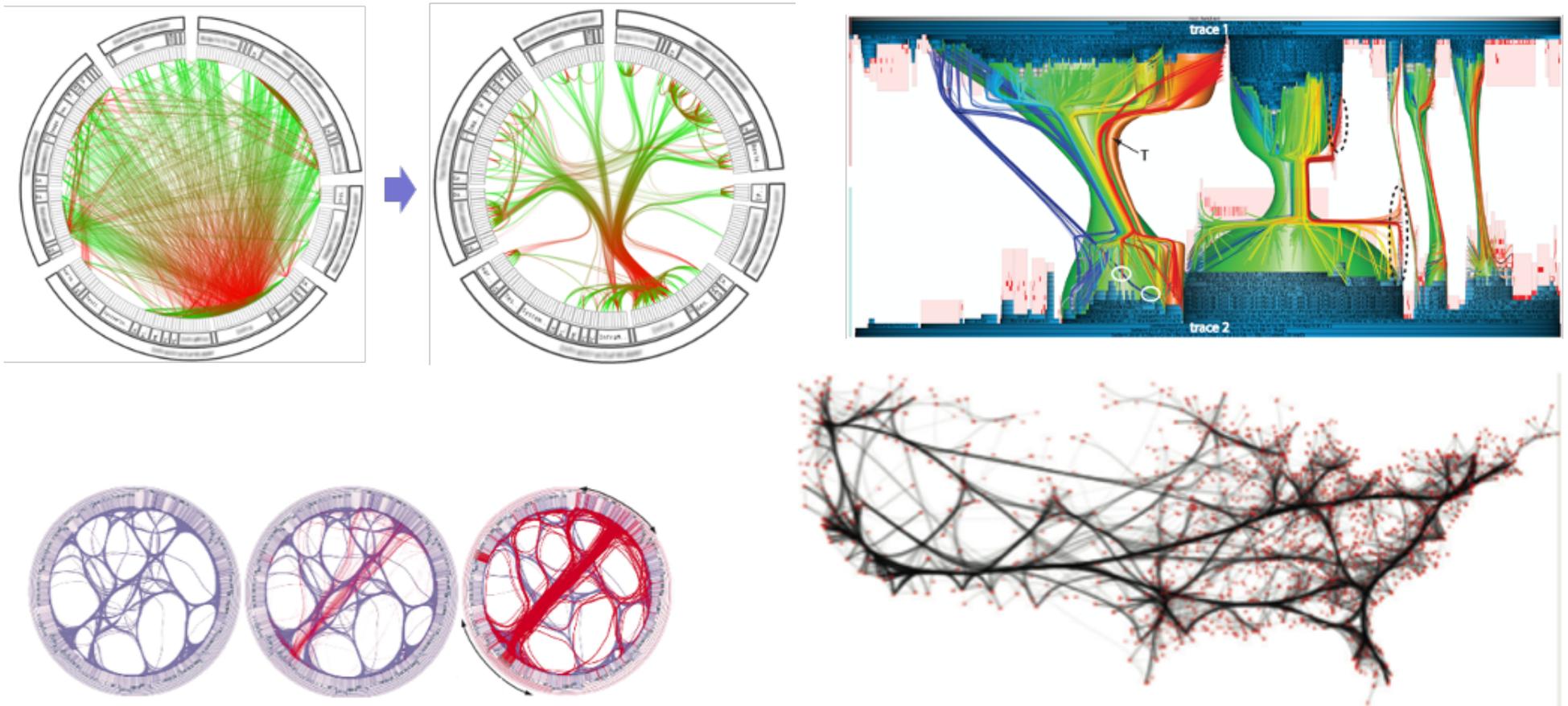
- edge bundling as a method to layout edges in multilayer network visualizations
- bundle only the inter-layer (or intra-layer) edges
- edge bundling is not specific for multilayer network visualizations



# Edge bundling

Method for reduction of clutter in a graph layout

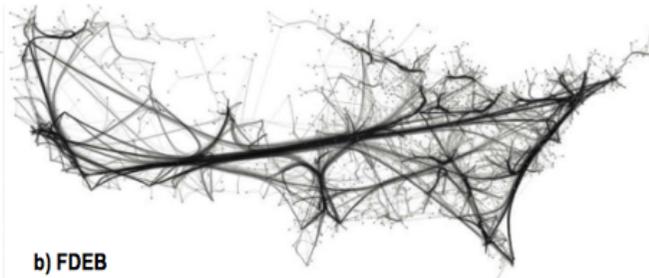
“Change the shape of edges by visually bundling them together analogous to the way electrical wires and network cables are merged into bundles...” [Holten, van Wijk, 09]



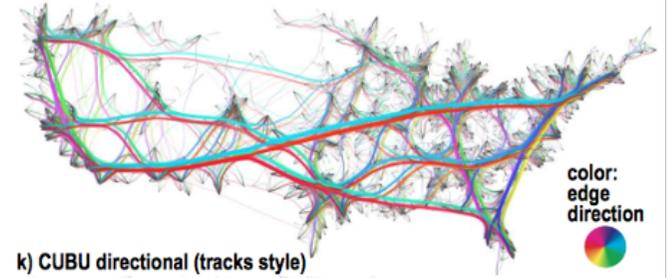
# Many methods



a) MINGLE



b) FDEB



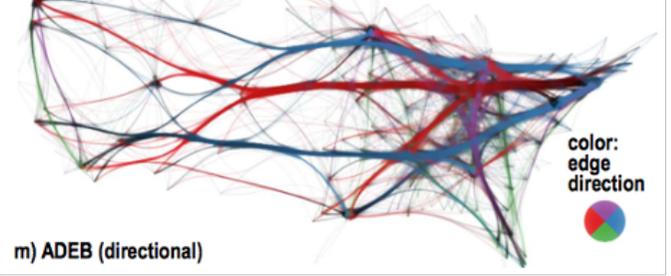
k) CUBU directional (tracks style)



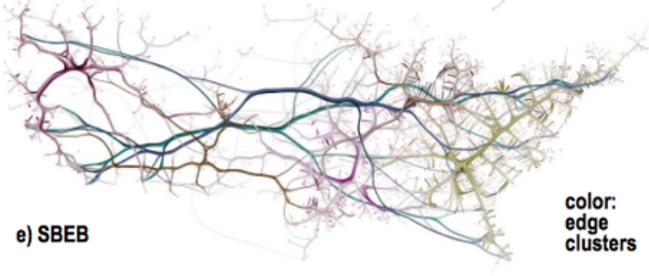
c) WR



d) KDEEB

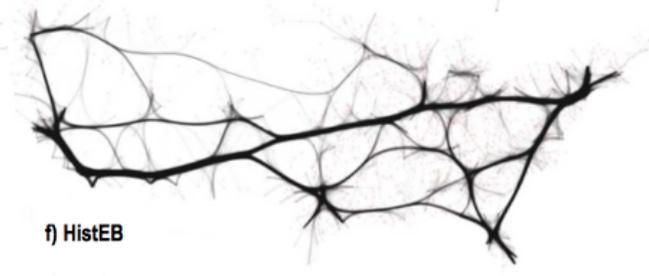


m) ADEB (directional)



e) SBEB

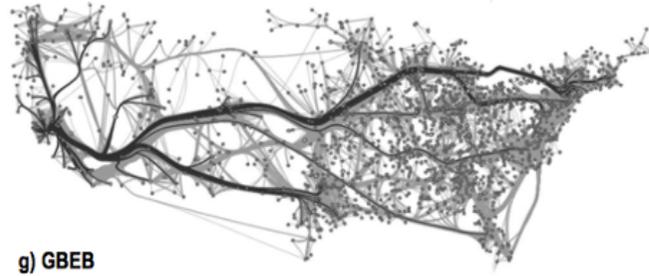
color:  
edge  
clusters



f) HistEB



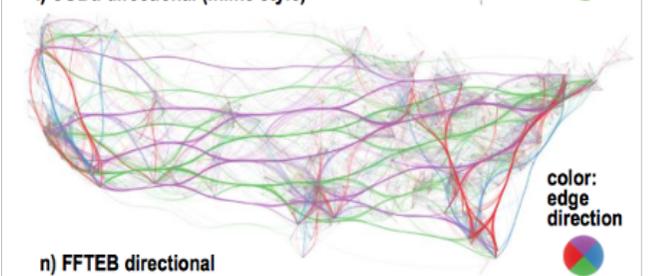
l) CUBu directional (inline style)



g) GBEB



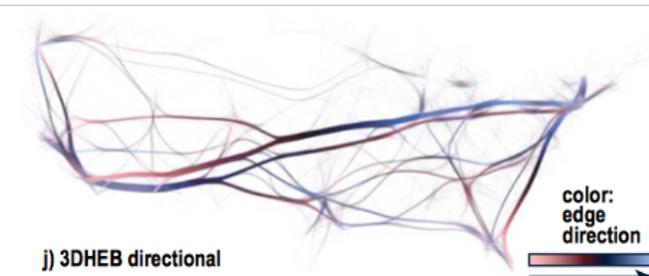
h) CUBu



n) FFTEB directional



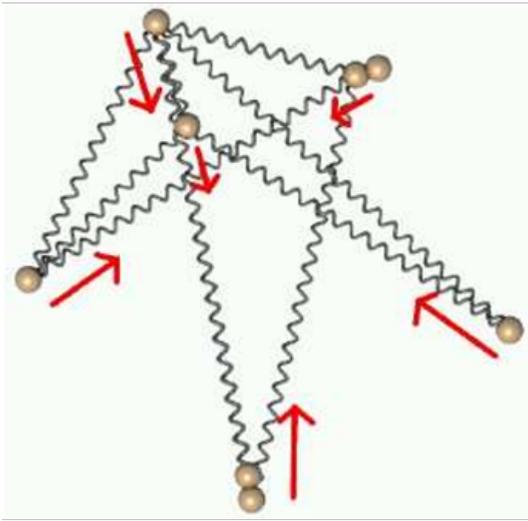
i) FFTEB



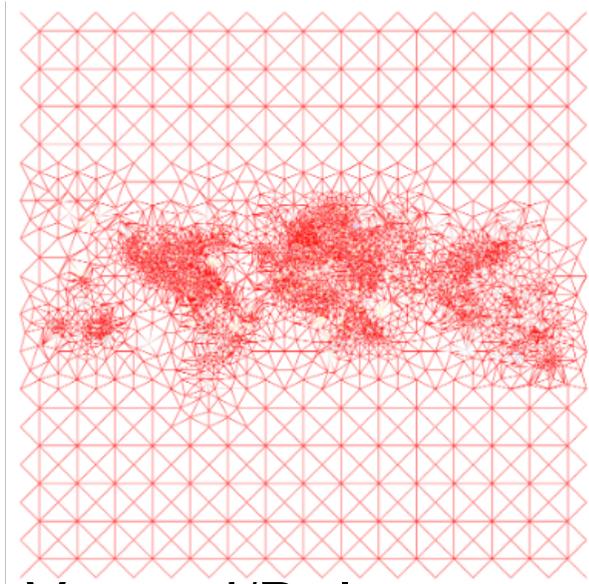
j) 3DHEB directional

color:  
edge  
direction

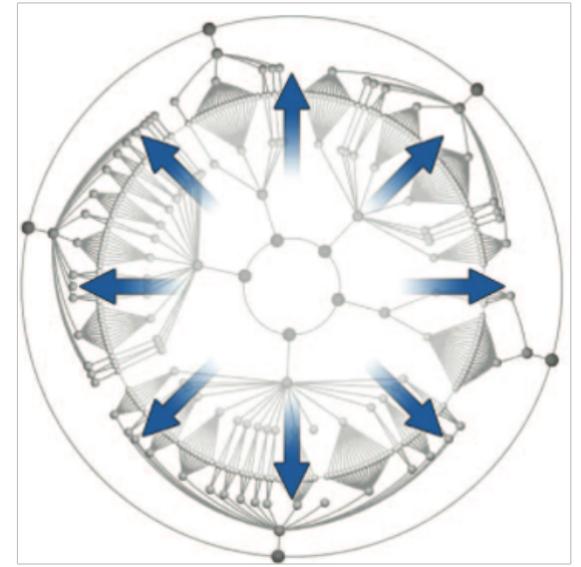
# Multiple techniques



spring embedders  
(FDEB)



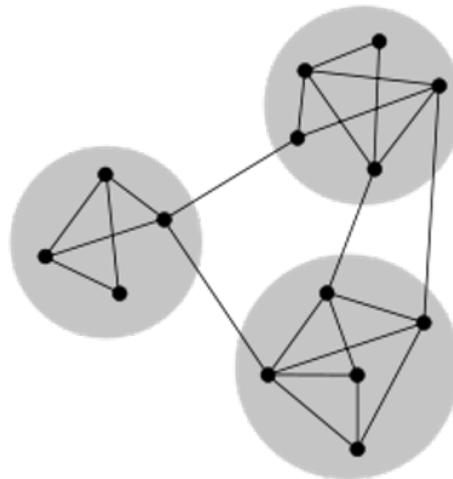
Voronoi/Delaunay  
diagrams



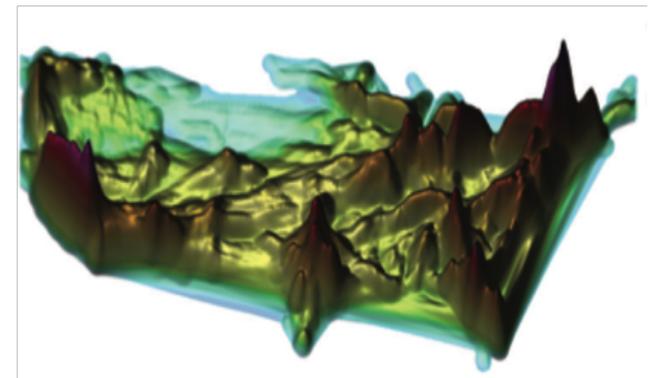
tree layouts & splines



medial axes

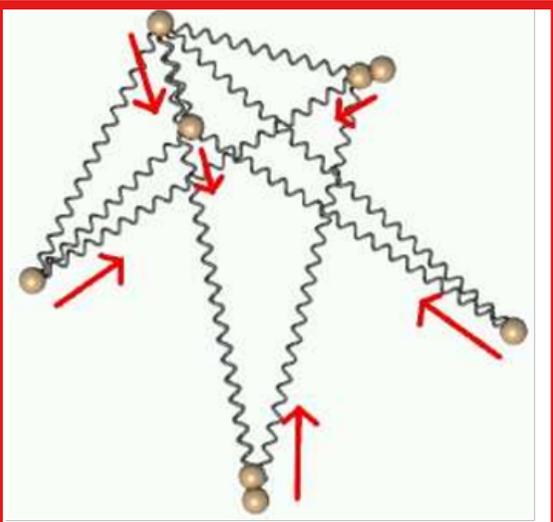


graph clustering

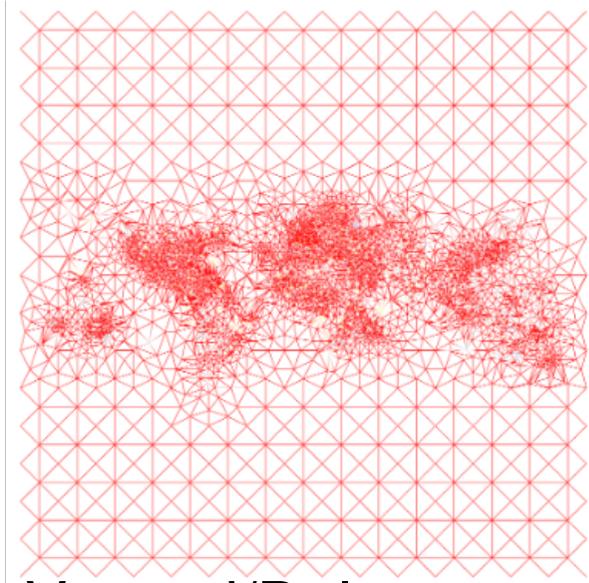


kernel density estimation

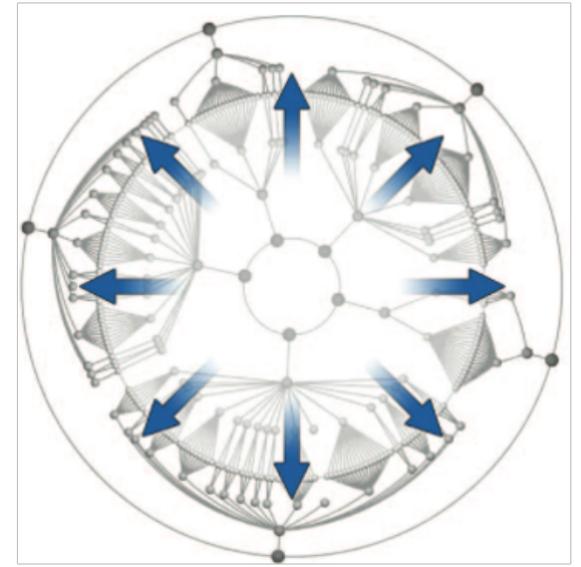
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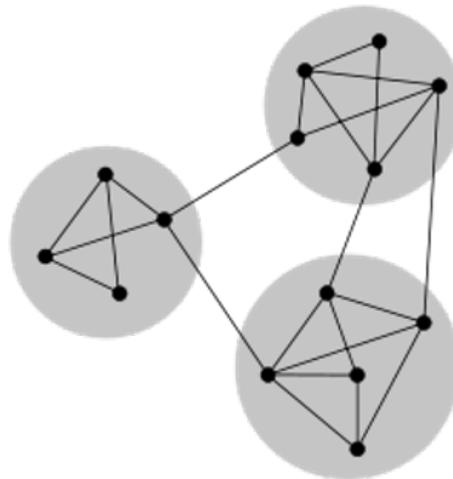
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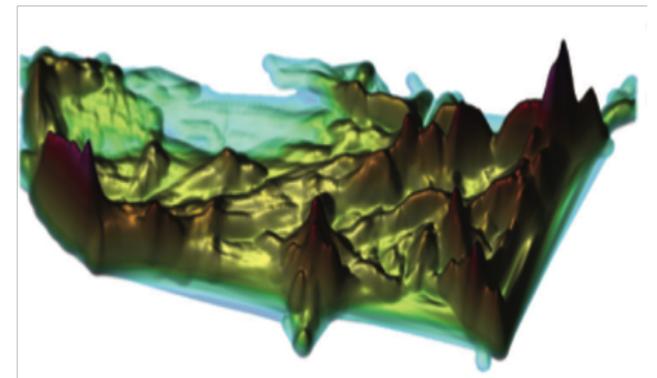
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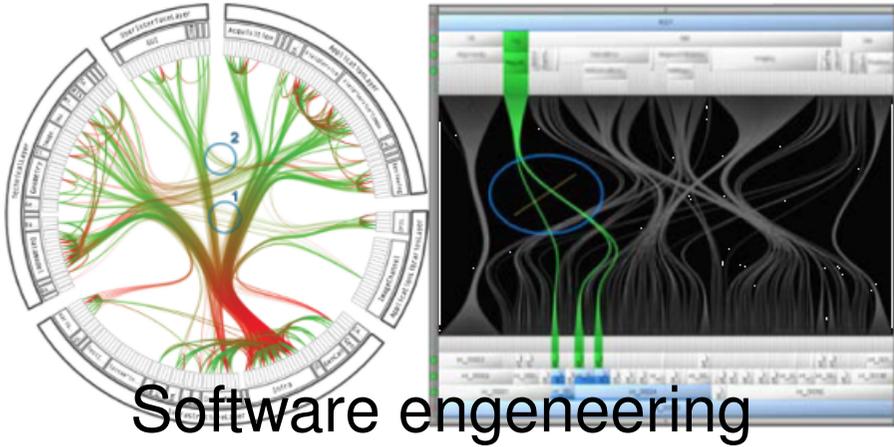


graph clustering



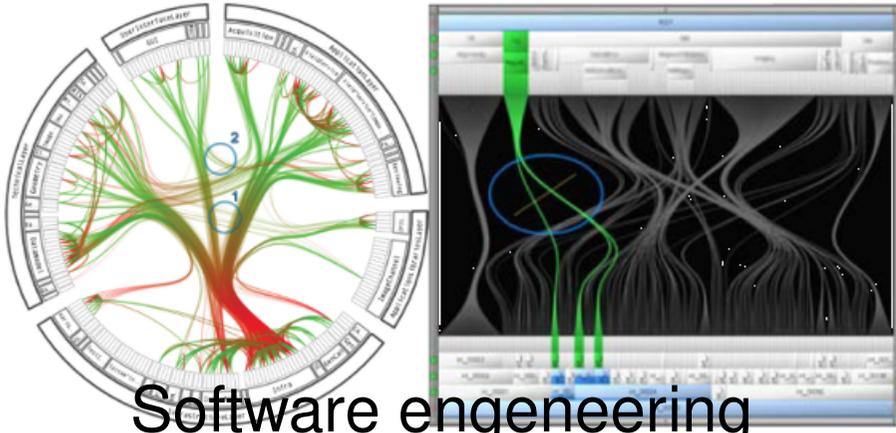
kernel density estimation

# Edge bundling: applications

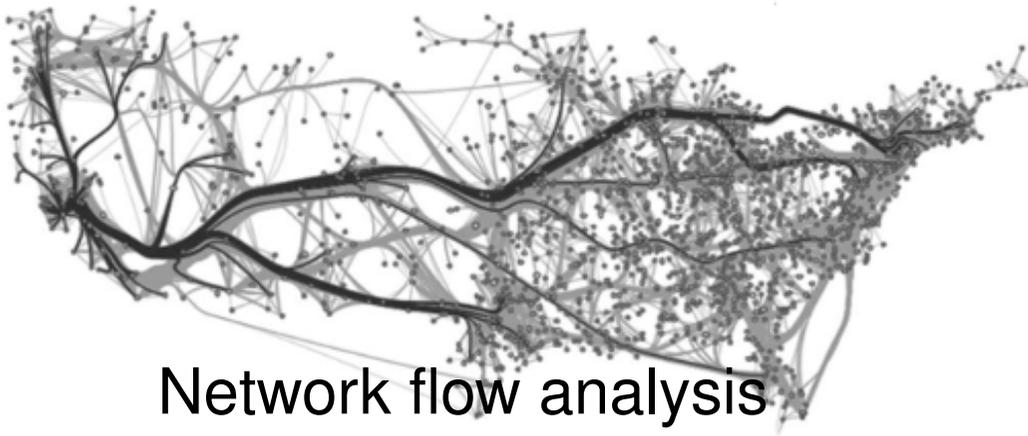


Software engineering

# Edge bundling: applications

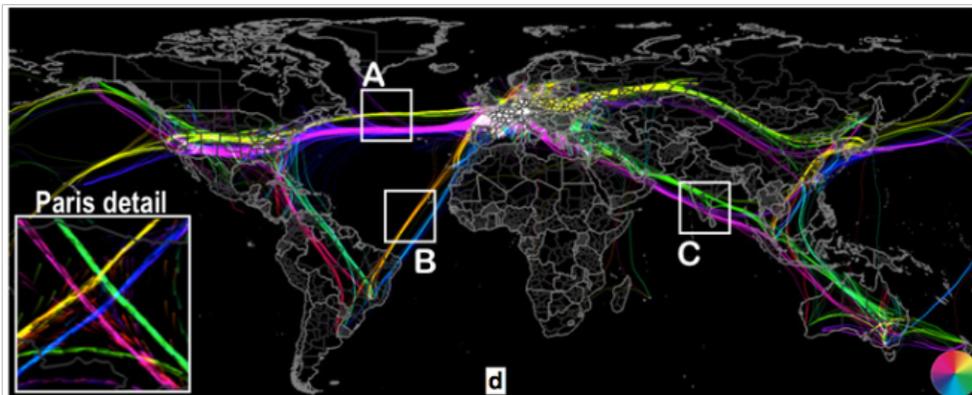
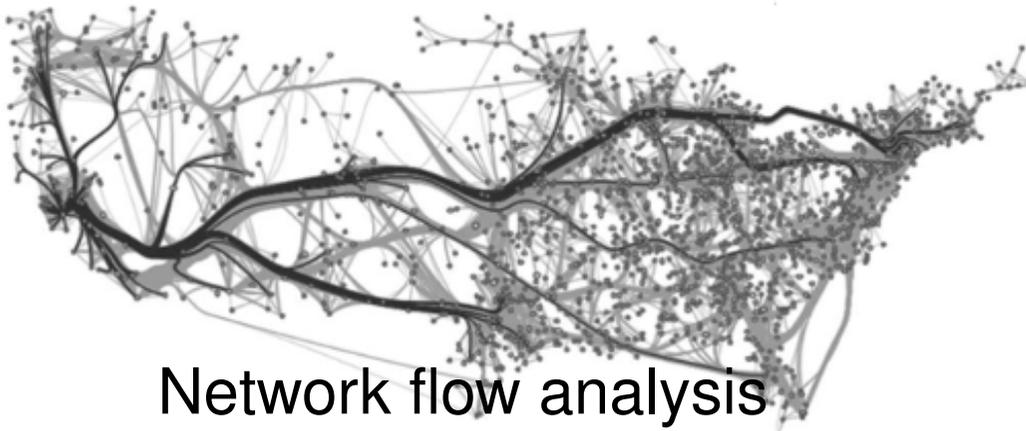
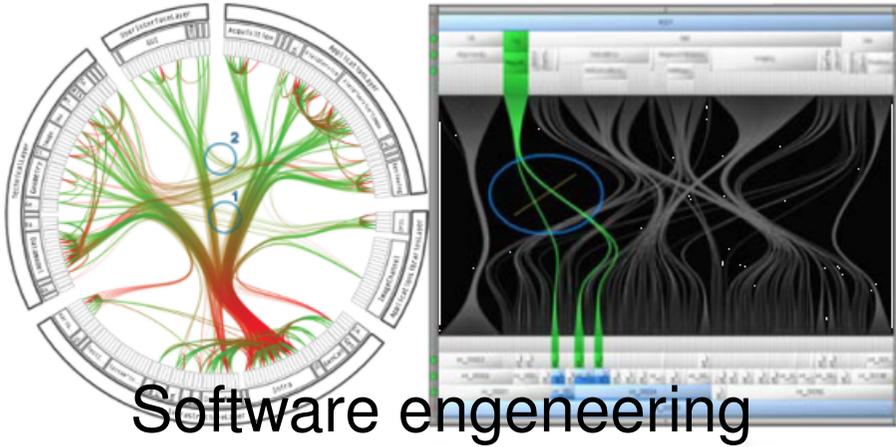


Software engineering



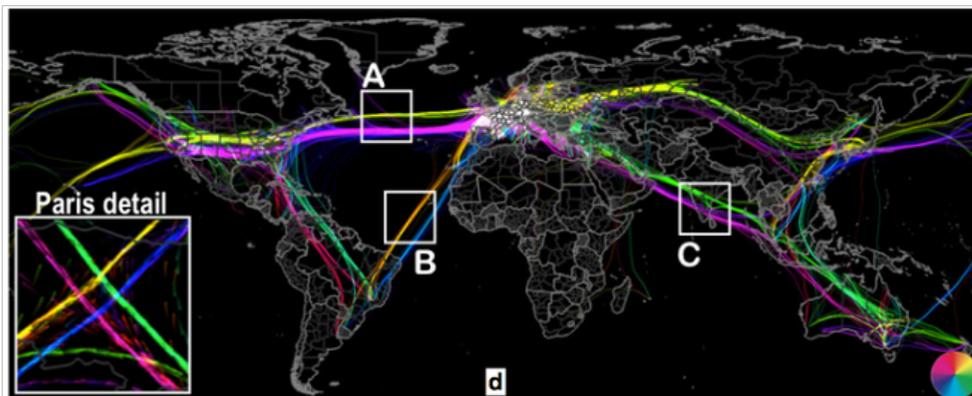
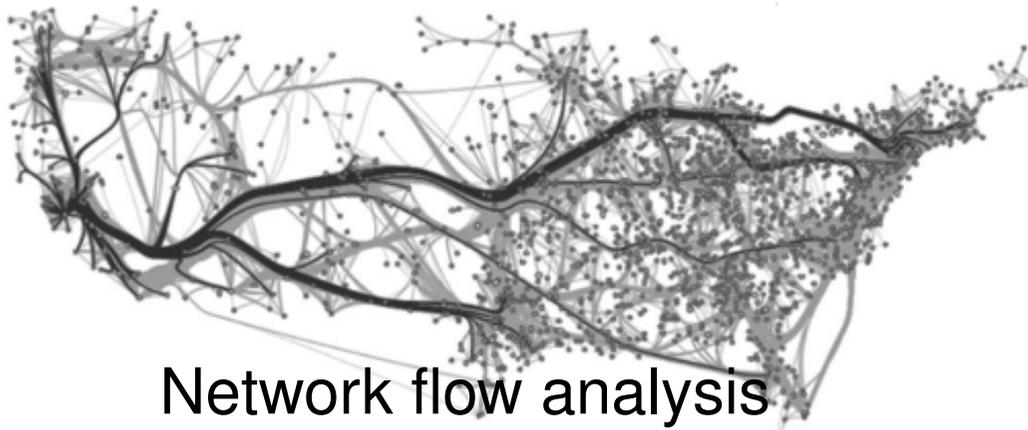
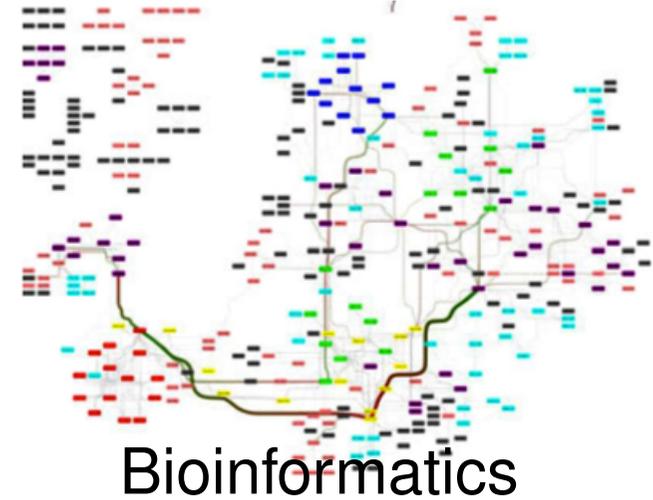
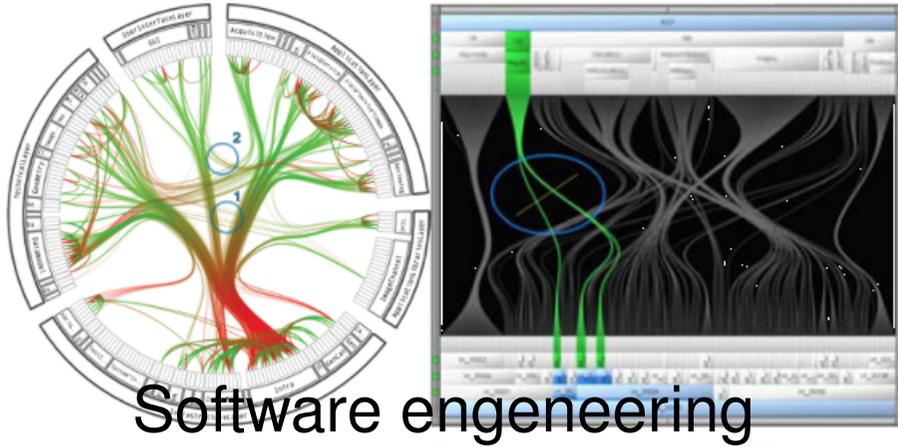
Network flow analysis

# Edge bundling: applications



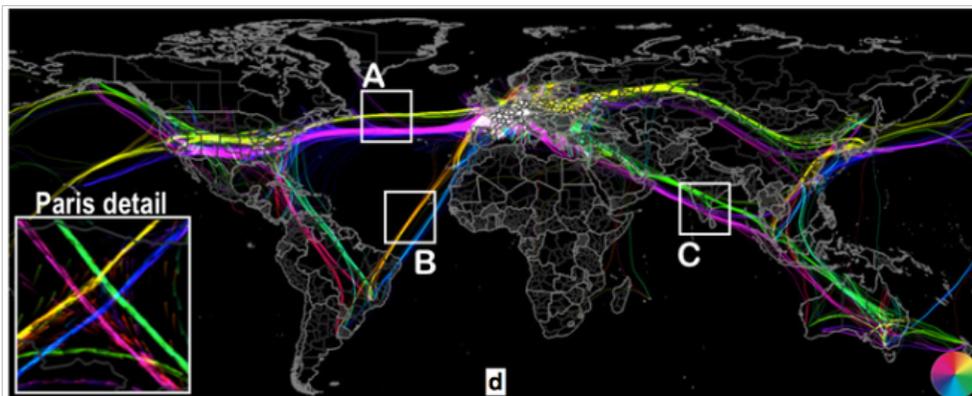
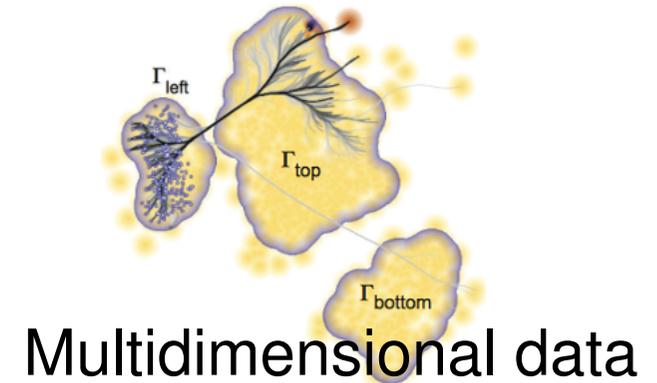
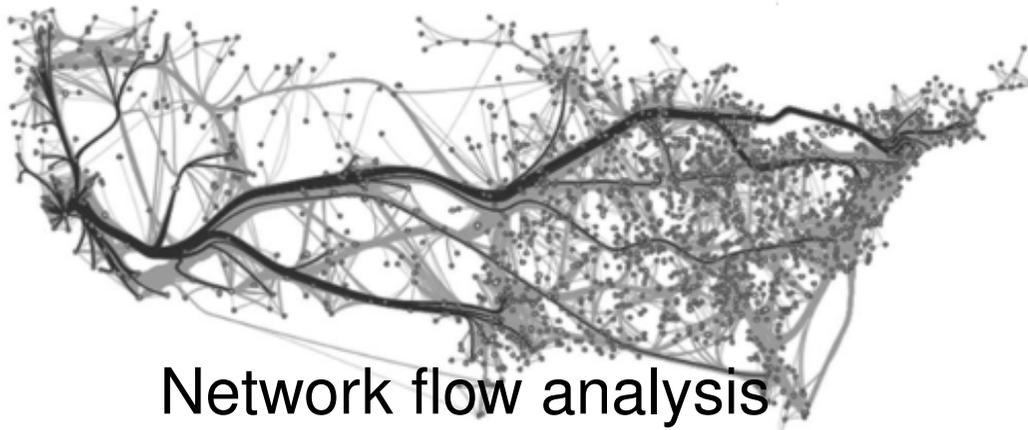
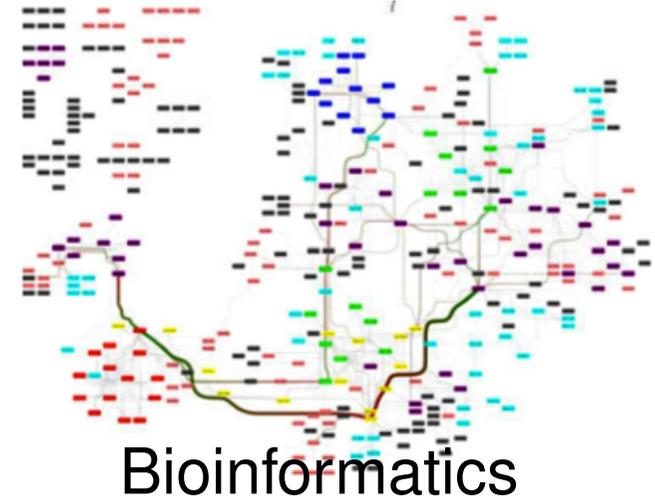
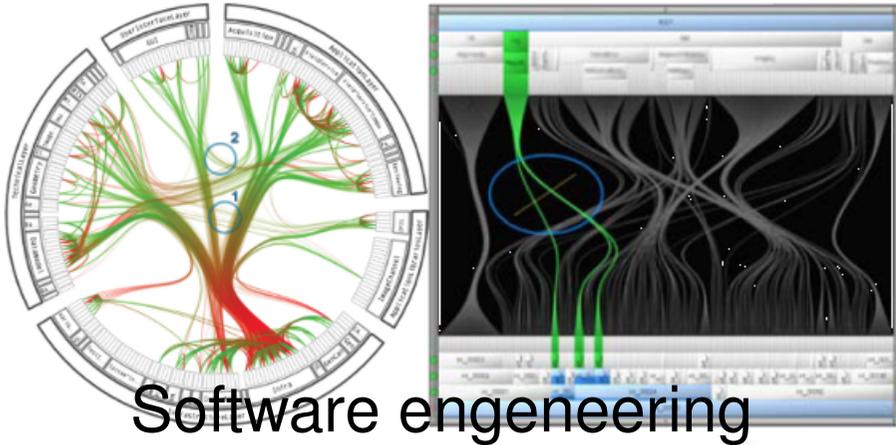
Air traffic control

# Edge bundling: applications



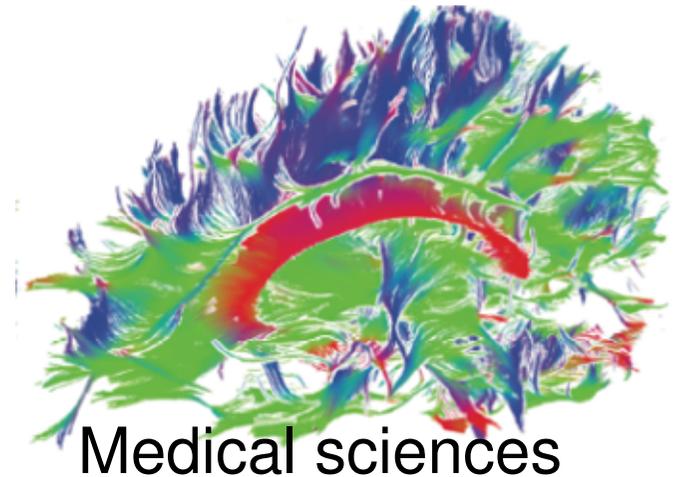
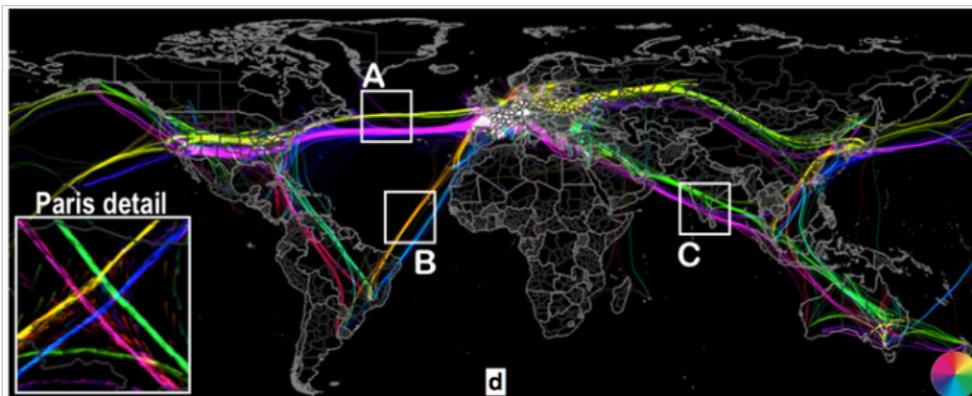
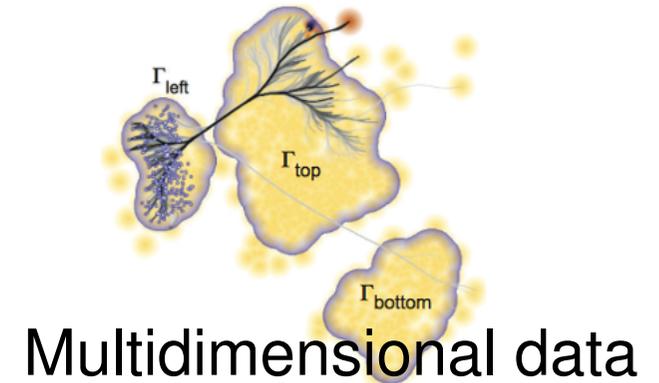
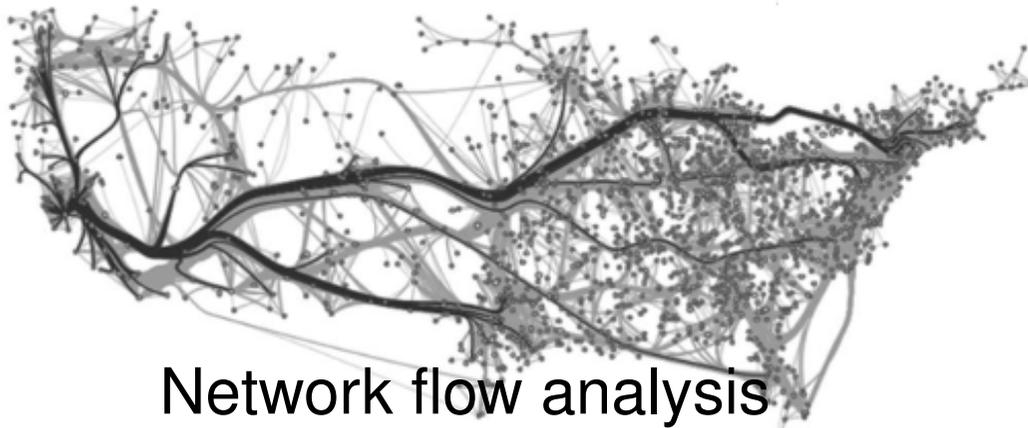
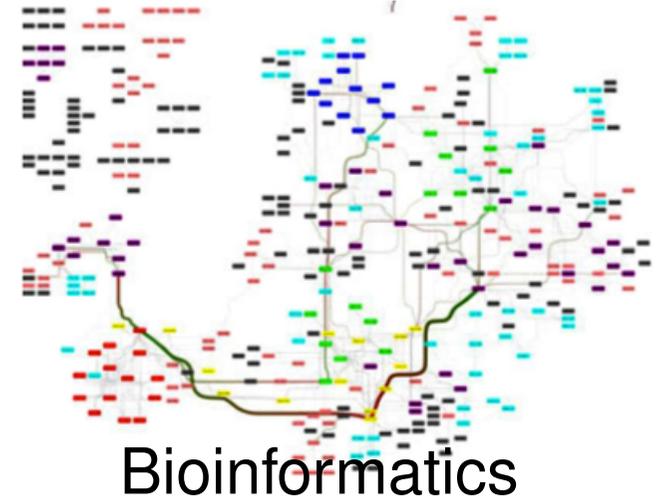
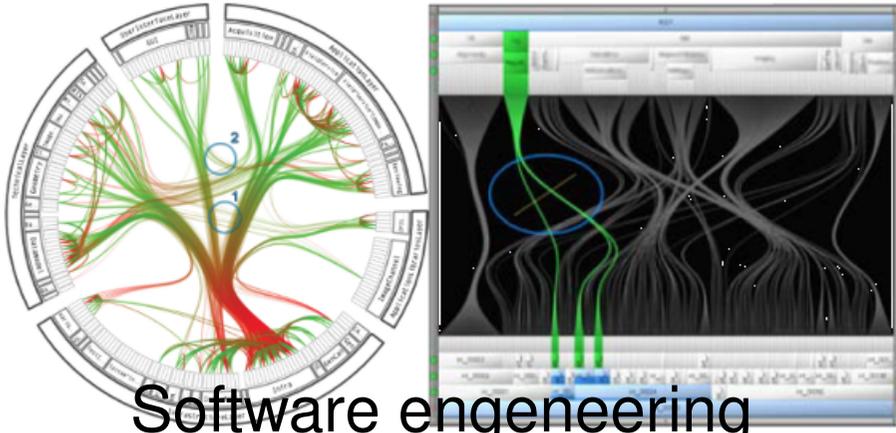
Air traffic control

# Edge bundling: applications

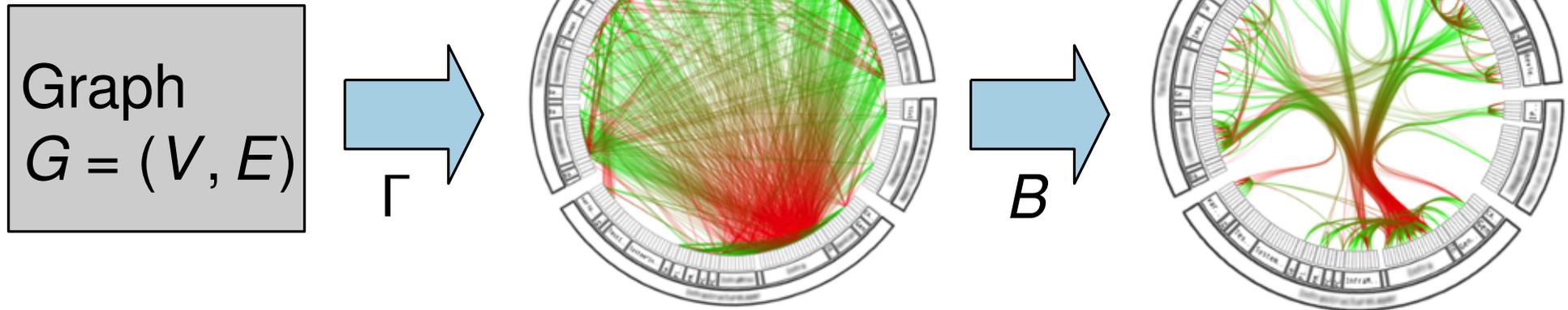


Air traffic control

# Edge bundling: applications



# Edge bundling: definition



$\Gamma$  – drawing/layout function

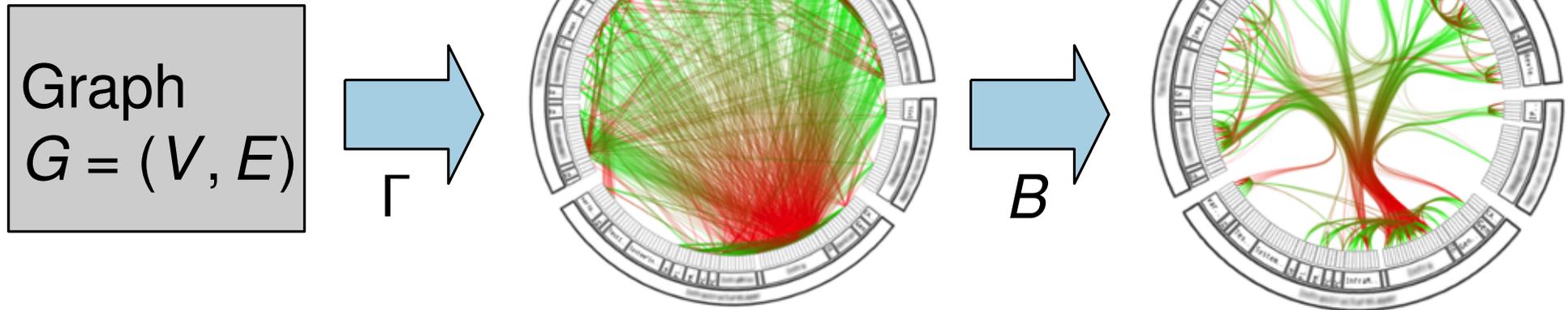
$B$  – bundling function

$$\forall (e_i, e_j) \in E \times E \text{ such that } e_i \neq e_j \wedge k(e_i, e_j) < k_{\max} \rightarrow \\ \delta(B(\Gamma(e_i)), B(\Gamma(e_j))) \ll \delta(\Gamma(e_i), \Gamma(e_j))$$

$k_{\max}$  – maximum similarity of the edges that still need to be bundled

$k$ –similarity of two edges;  $\delta$  – similarity of two curves

# Edge bundling: definition



$\Gamma$  – drawing/layout function

$B$  – bundling function

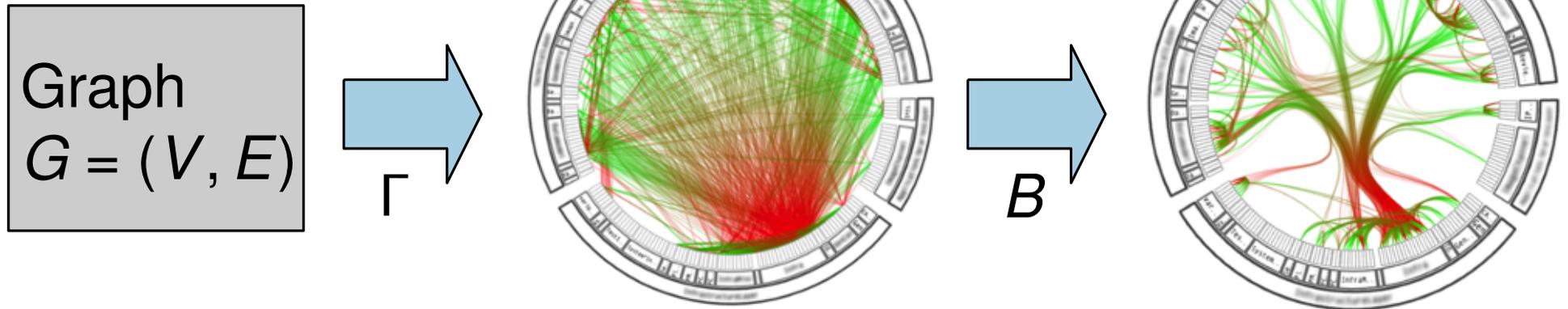
$\forall (e_i, e_j) \in E \times E$  such that  $e_i \neq e_j \wedge$  **two edges are similar**  $k(e_i, e_j) < k_{\max} \rightarrow$

$$\delta(B(\Gamma(e_i)), B(\Gamma(e_j))) \ll \delta(\Gamma(e_i), \Gamma(e_j))$$

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the distance between curves after bundling is small

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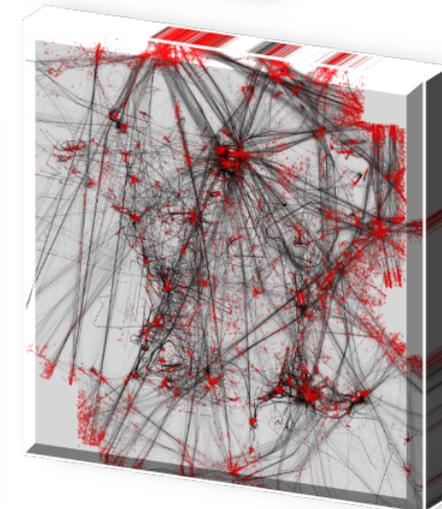
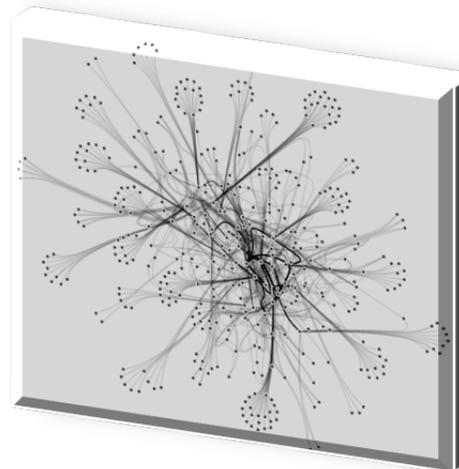
## Data-based similarities

- Structured-based
- Attribute-based

	100 h	0 h	100 h	200 h	300 h	400 h	500 h	600 h	700 h	800 h
50 h	5.1	5.6	5.1	4.9	5.0	4.8	4.9	4.8	4.8	4.7
100 h	1.4	1.4	1.4	1.4	1.3	1.2	1.2	1.2	1.1	1.1
150 h	6.7	6.5	6.4	6.4	6.3	6.3	6.2	6.2	6.1	6.1
200 h	1.8	1.8	1.7	1.7	1.6	1.6	1.5	1.5	1.4	1.4
250 h	8.4	8.3	8.1	8.1	7.9	7.8	7.8	7.7	7.5	7.4
300 h	2.2	2.2	2.1	2.1	2.0	2.0	1.9	1.9	1.8	1.7
350 h	10.5	10.3	10.3	10.0	10.0	9.8	9.8	9.6	9.4	9.2
400 h	2.6	2.6	2.5	2.5	2.4	2.4	2.3	2.3	2.2	2.1
450 h	12.9	12.8	12.5	12.3	12.2	12.0	11.9	11.7	11.7	11.4
500 h	3.1	3.1	3.0	3.0	2.9	2.9	2.7	2.7	2.6	2.5
550 h	15.6	15.3	15.2	14.9	14.8	14.5	14.4	14.1	14.0	13.7
600 h	3.4	3.4	3.3	3.3	3.2	3.2	3.0	3.0	2.9	2.8
650 h	17.0	16.8	16.6	16.4	16.2	16.0	15.8	15.6	15.4	15.2
700 h	3.7	3.7	3.6	3.6	3.5	3.5	3.3	3.3	3.2	3.1
750 h	18.3	18.1	17.9	17.7	17.5	17.3	17.1	16.9	16.7	16.5
800 h	4.0	4.0	3.9	3.9	3.8	3.8	3.6	3.6	3.5	3.4

## Drawing-based similarities

- Geometric-based
- Image-based



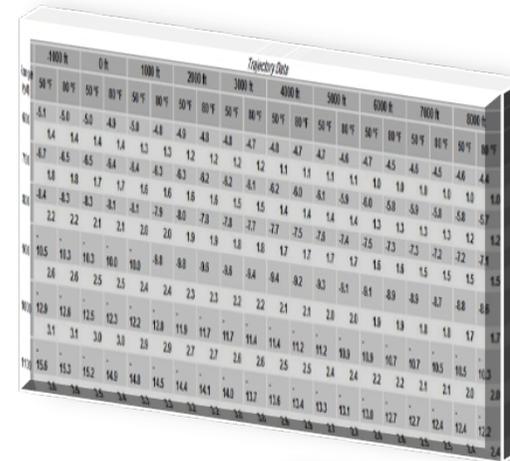
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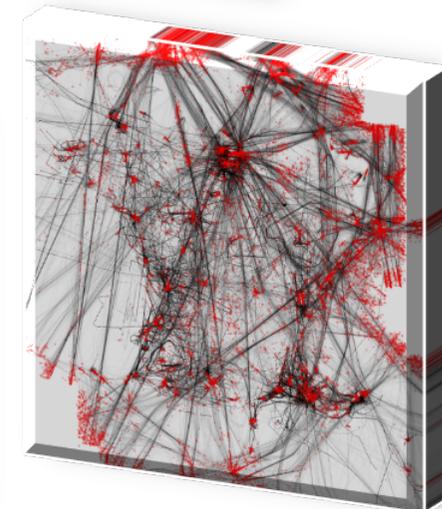
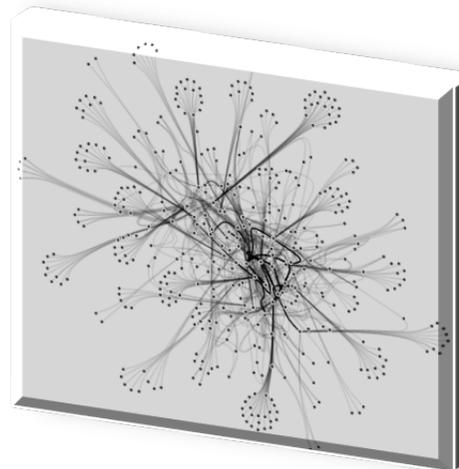
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	100 h	0 h	100 h	200 h	300 h	400 h	500 h	600 h	700 h	800 h										
50'	5.1	5.6	5.1	4.9	5.0	4.8	4.9	4.8	4.8	4.7	4.8	4.7	4.7	4.6	4.7	4.5	4.5	4.5	4.6	4.4
1 h	1.4	1.4	1.4	1.4	1.3	1.3	1.2	1.2	1.2	1.2	1.1	1.1	1.1	1.1	1.1	1.0	1.0	1.0	1.0	1.0
5 h	0.7	0.5	0.4	0.4	0.4	0.3	0.3	0.2	0.2	0.1	0.2	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0
10 h	1.8	1.8	1.7	1.7	1.6	1.6	1.6	1.5	1.5	1.4	1.4	1.4	1.4	1.3	1.3	1.3	1.2	1.2	1.2	1.2
30 h	0.4	0.3	0.3	0.3	0.3	0.2	0.2	0.2	0.2	0.1	0.2	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1
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5 h	10.5	10.3	10.3	10.0	10.0	9.8	9.8	9.6	9.6	9.4	9.4	9.2	9.3	9.1	9.1	8.9	8.9	8.7	8.8	8.8
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30 h	12.9	12.8	12.5	12.3	12.2	12.0	11.9	11.7	11.7	11.6	11.4	11.2	11.2	11.0	10.9	10.7	10.7	10.5	10.5	10.3
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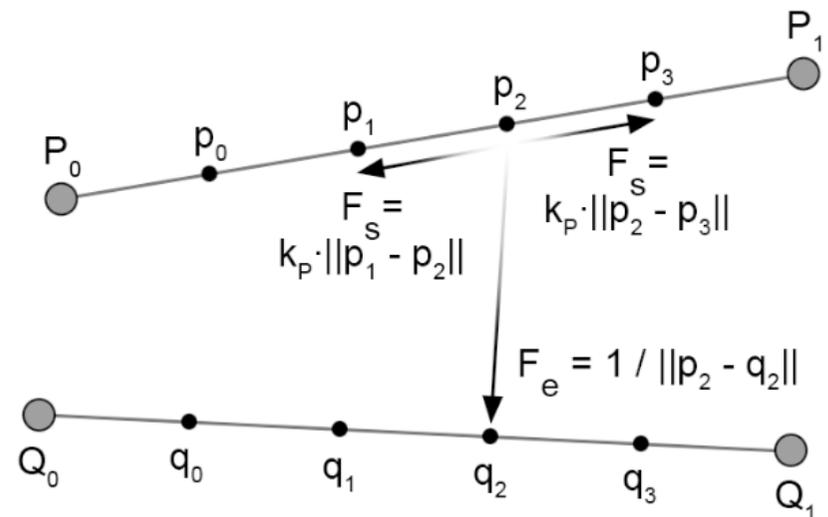
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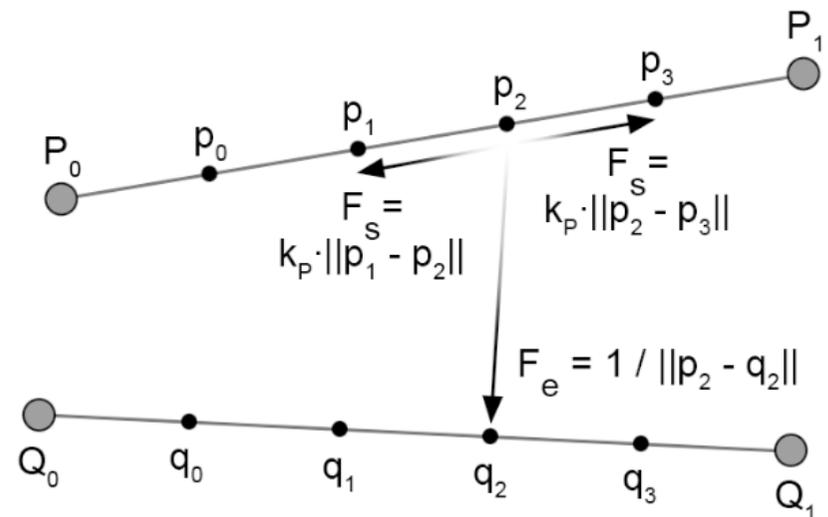
# Edge bundling Holten and van Wijk, 2009

- Assume two edges  $P$  and  $Q$  need to be bundled (which – later). We say they are *interacting*.



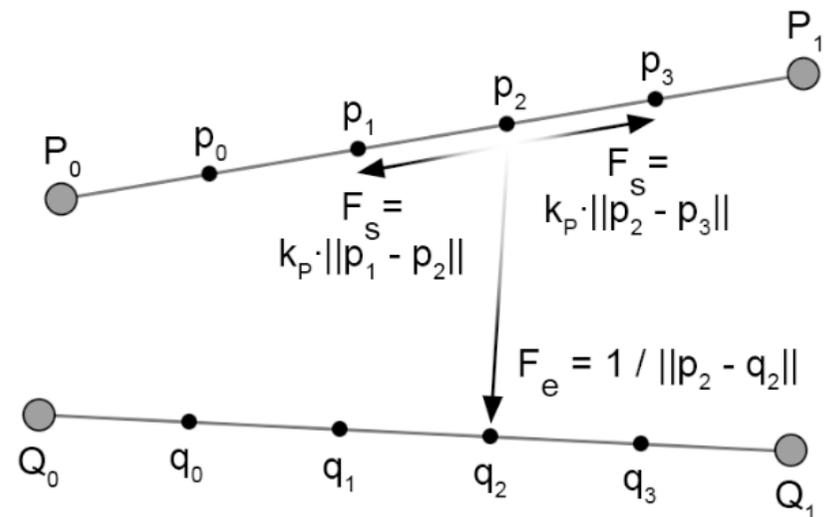
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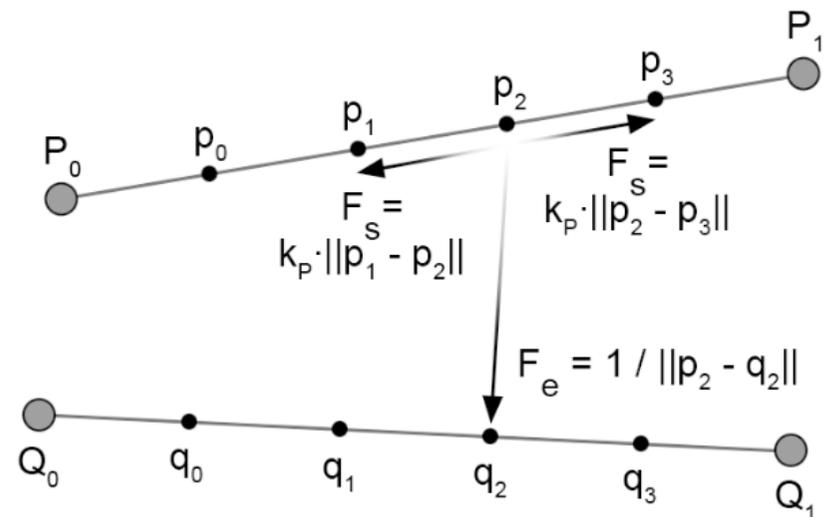
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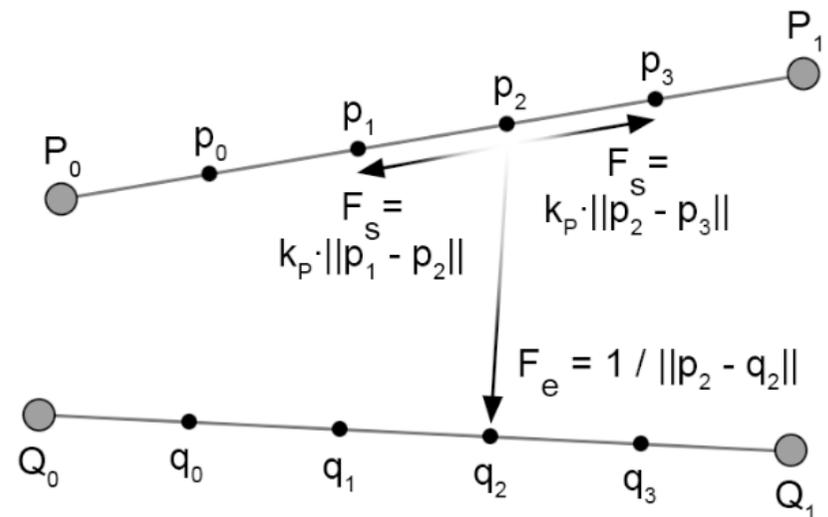


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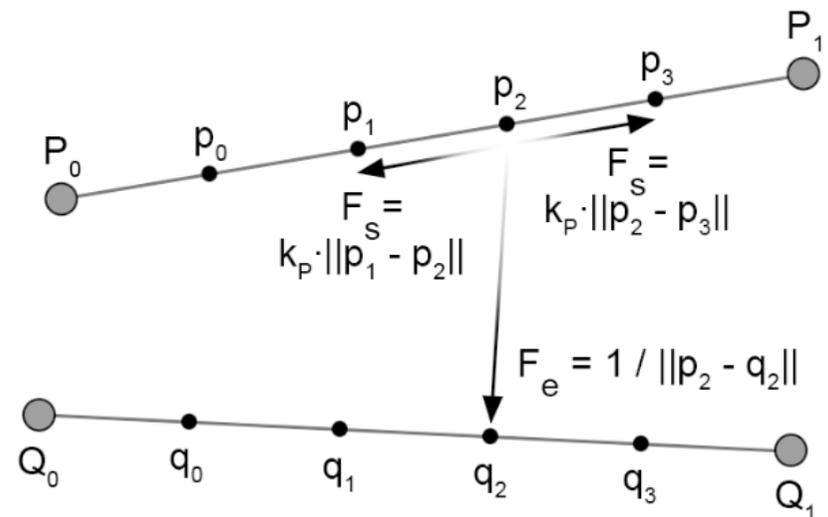
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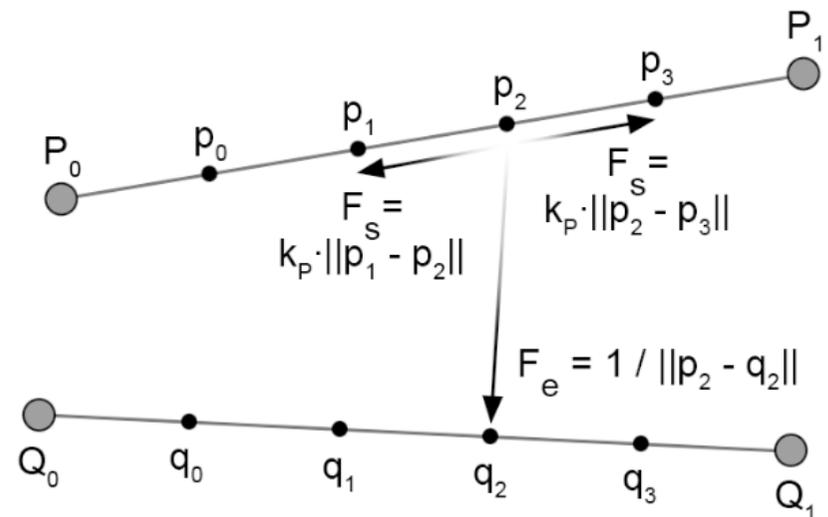
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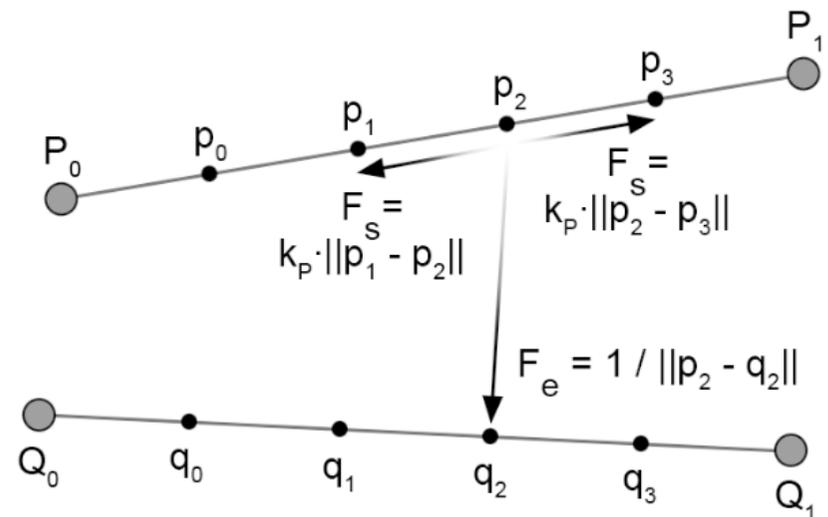
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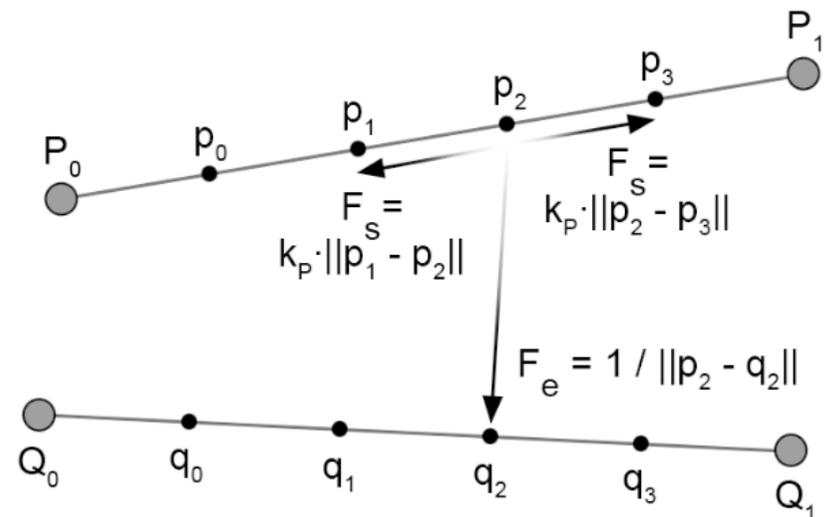
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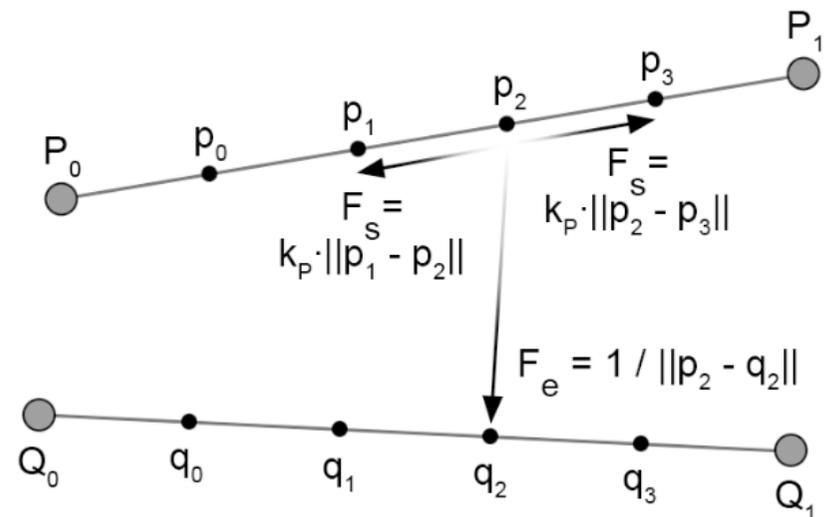
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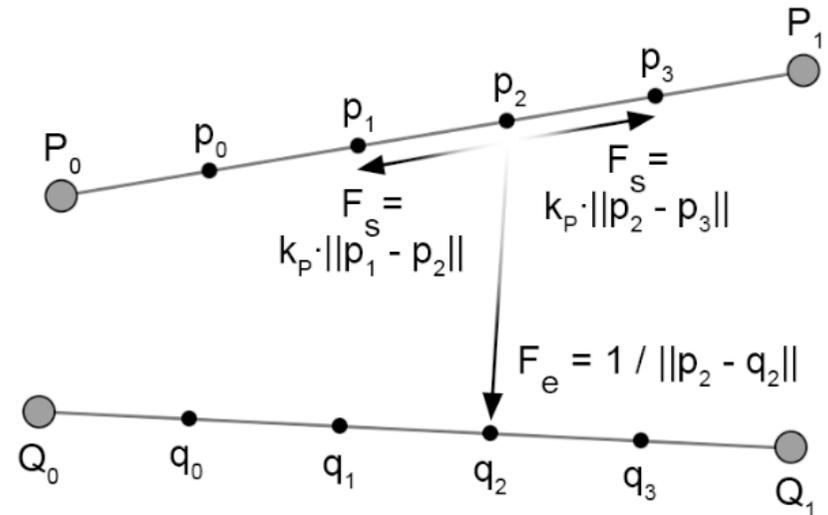
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Large values of  $K$  make system very stiff



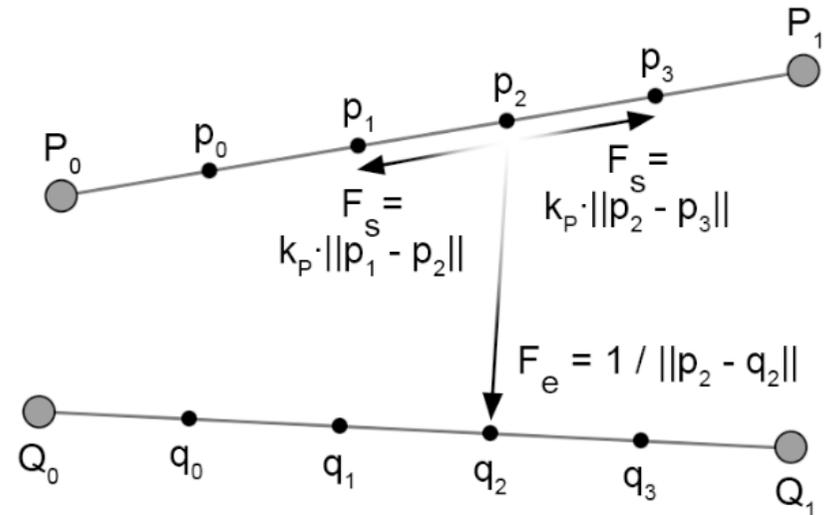
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- An attraction electrostatic force  $F_e(\{p_i, q_i\}) = \frac{1}{\|p_i - q_i\|}$  is used between each pair of corresponding subdivision points of  $P$  and  $Q$ , thus between  $p_0$  and  $q_0$ ,  $p_1$  and  $q_1$ , ...



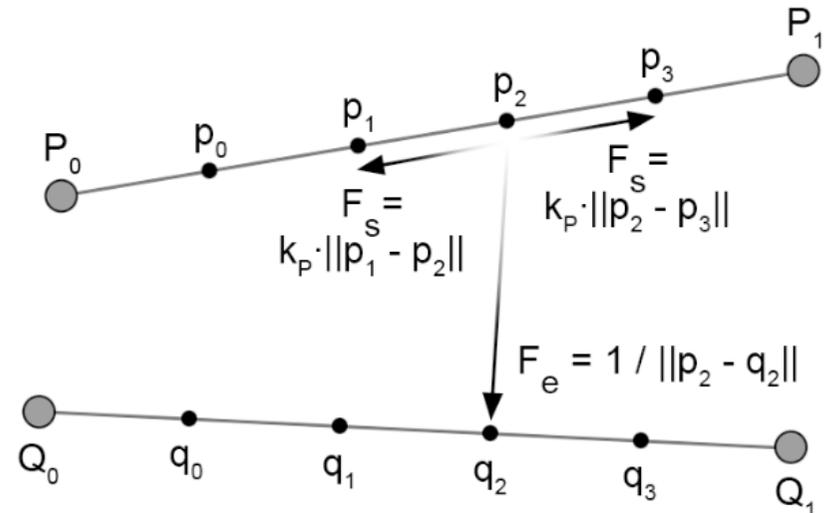
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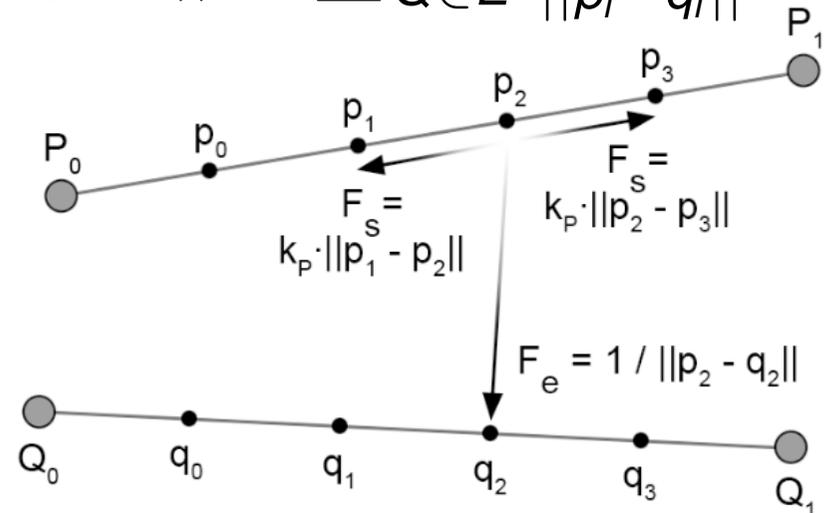
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$$F_{p_i} = k_P \cdot (\|p_{i-1} - p_i\| + \|p_i - p_{i+1}\|) + \sum_{Q \in E} \frac{1}{\|p_i - q_i\|}$$



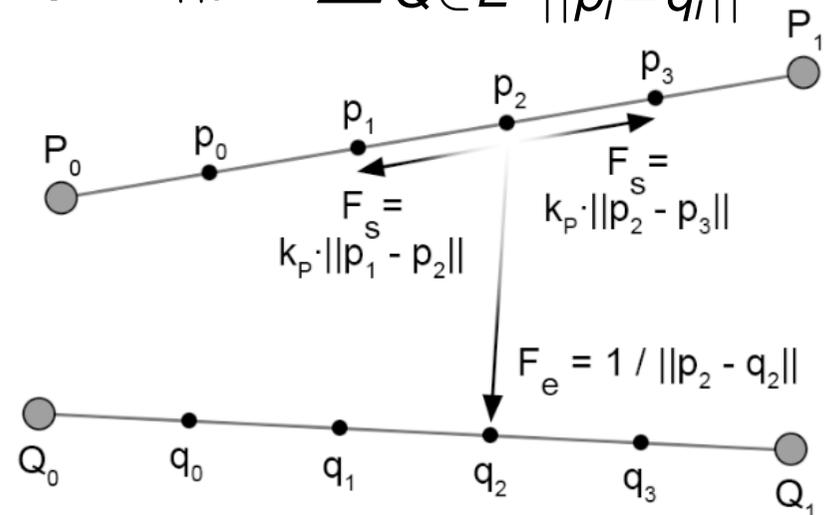
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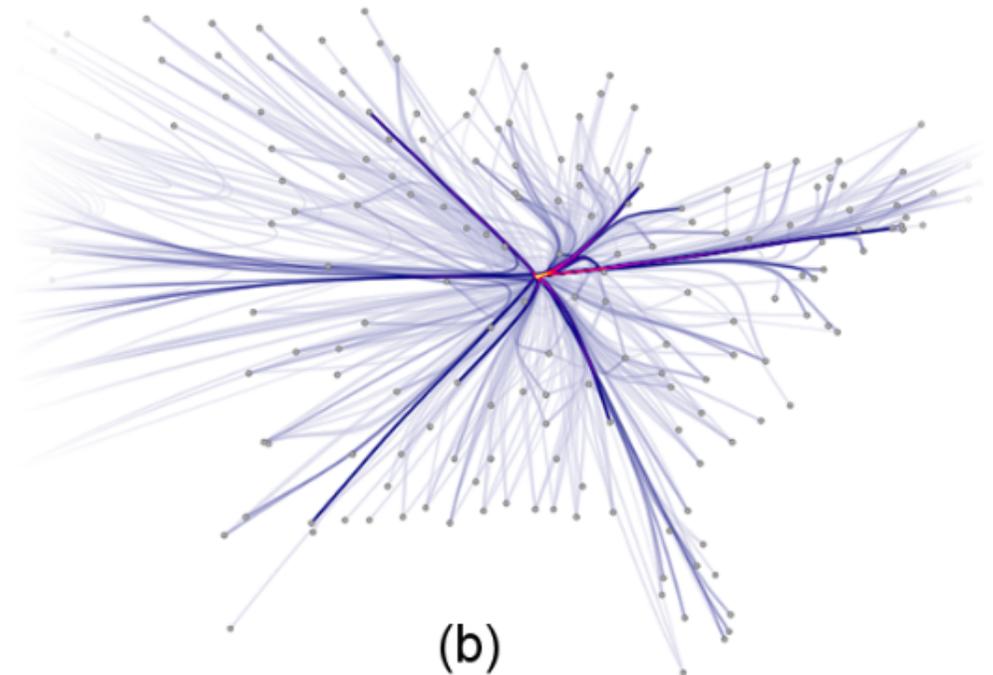
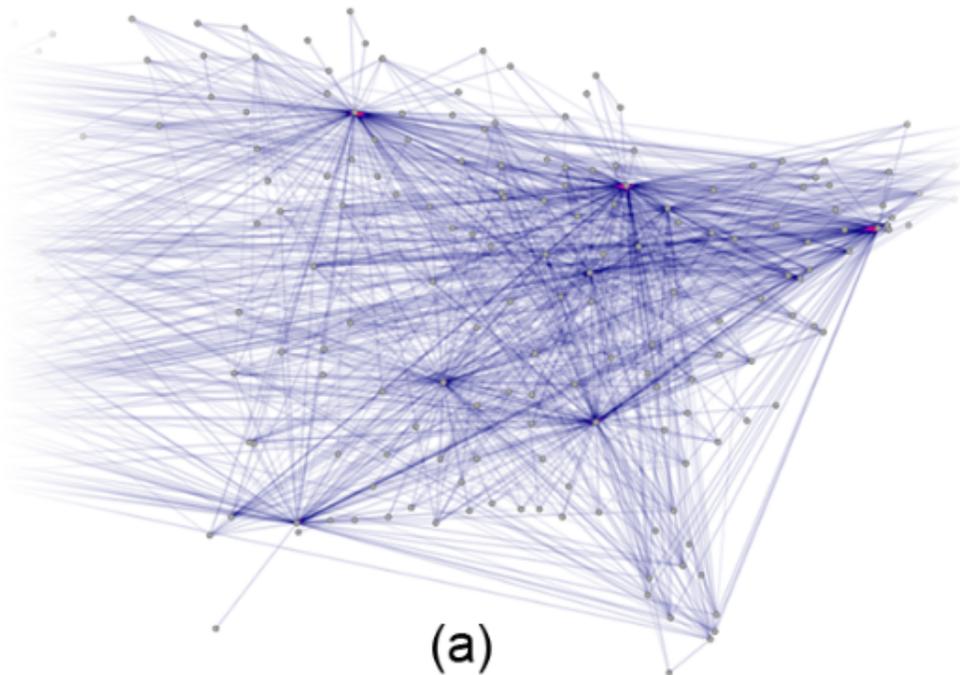
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$k_P$  – constant for edge  $P$



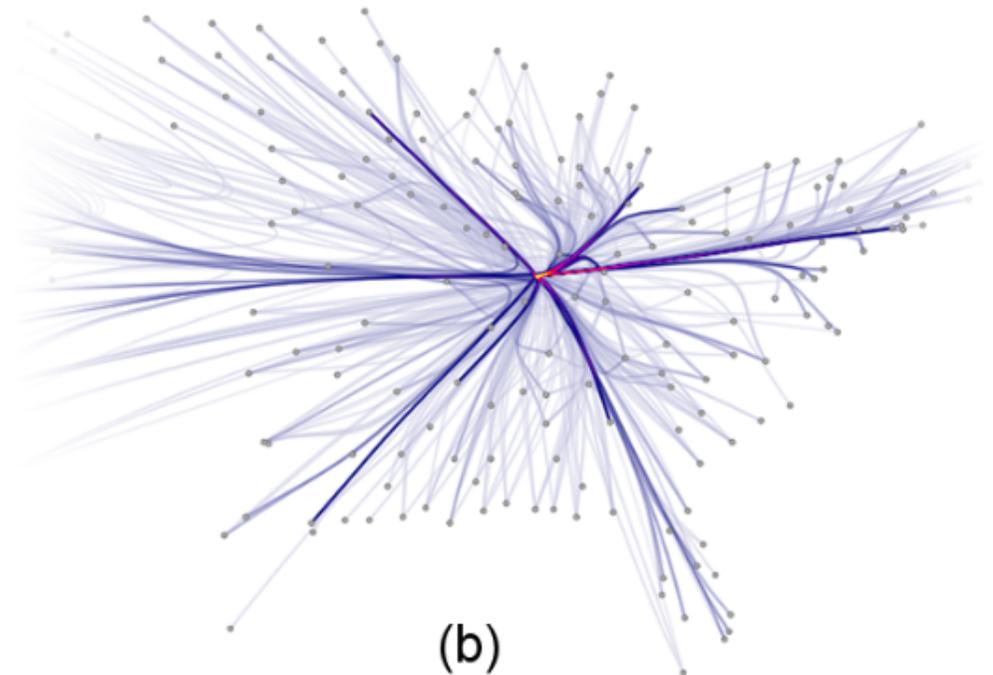
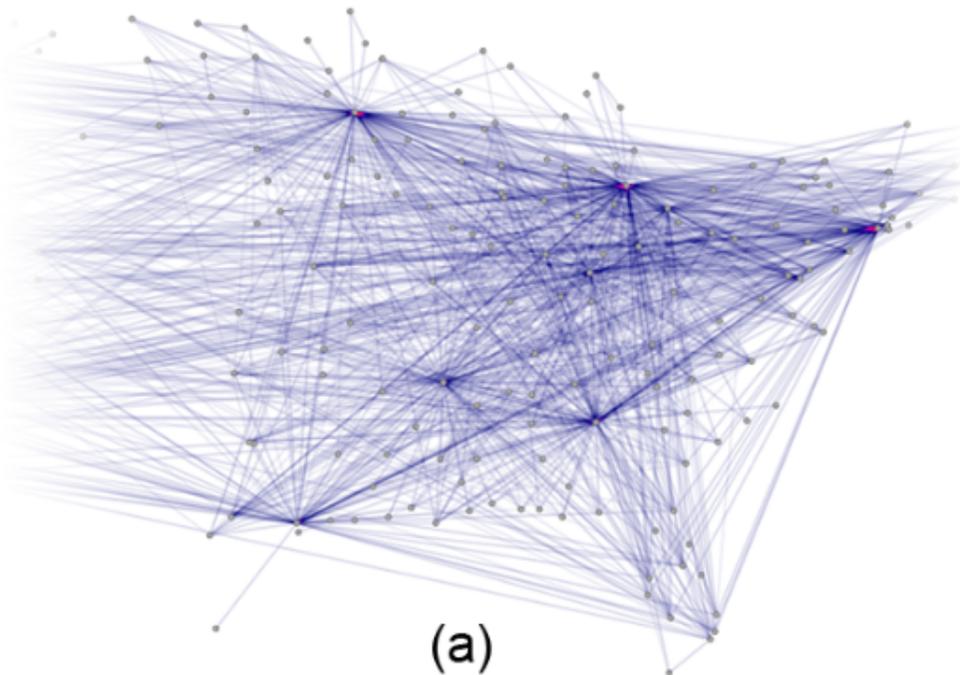
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- Fig.b – performance of the model given up to now. Here all edges interact with all.



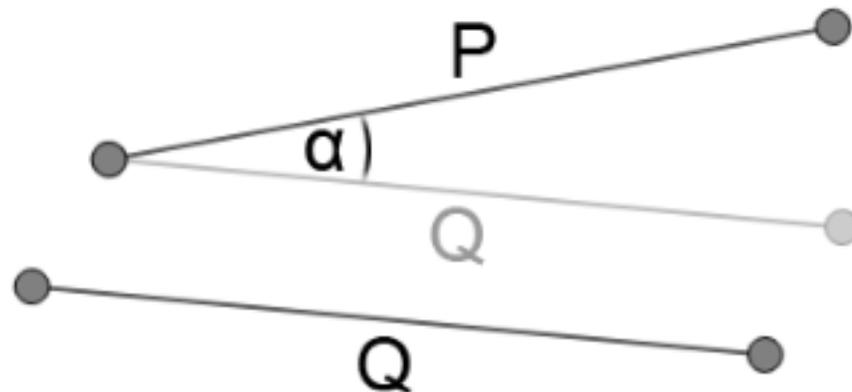
# Edge bundling: performance

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- Increasing the value of  $K$  gives less bundling overall and therefore in parts of the graph where a high amount of bundling is still desirable



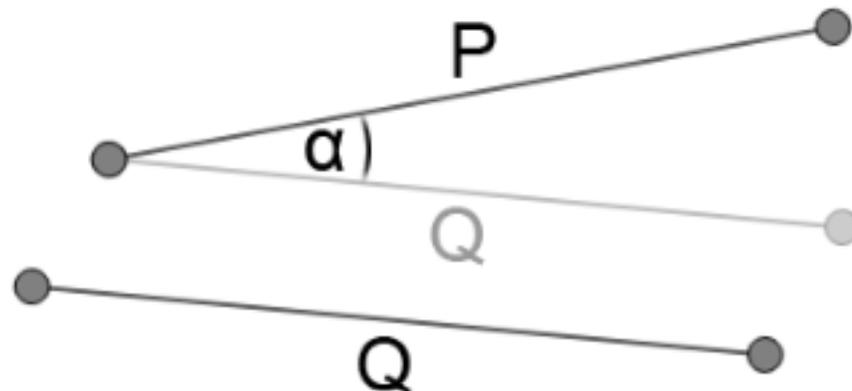
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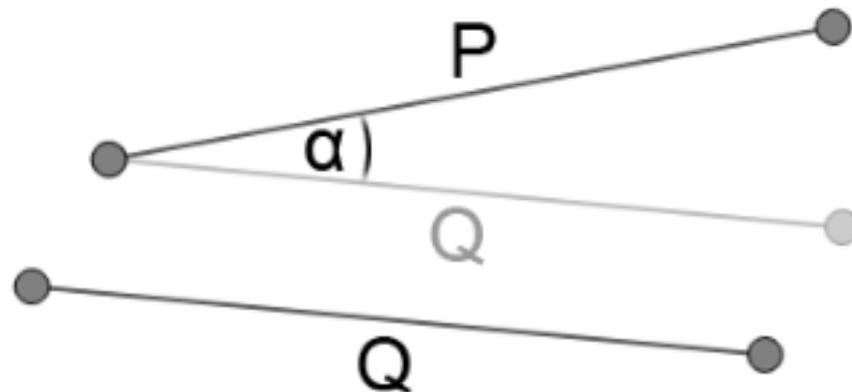
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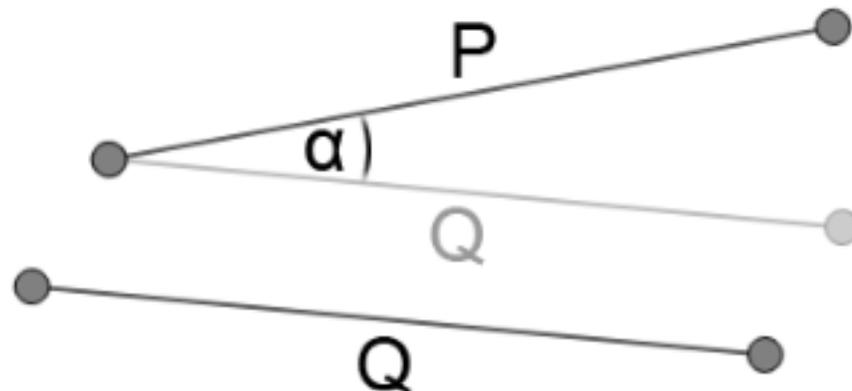
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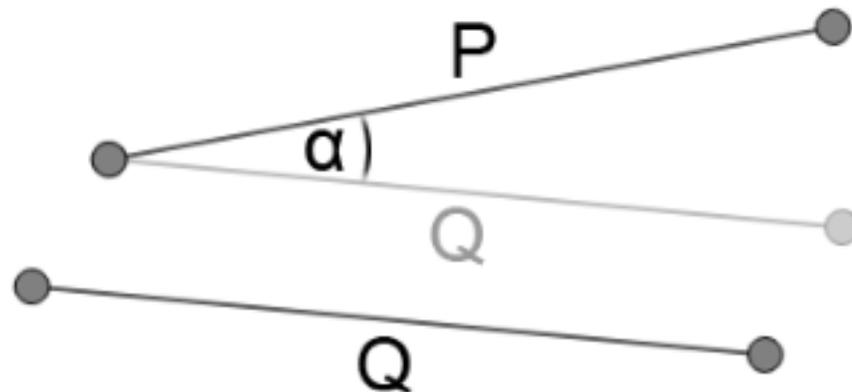
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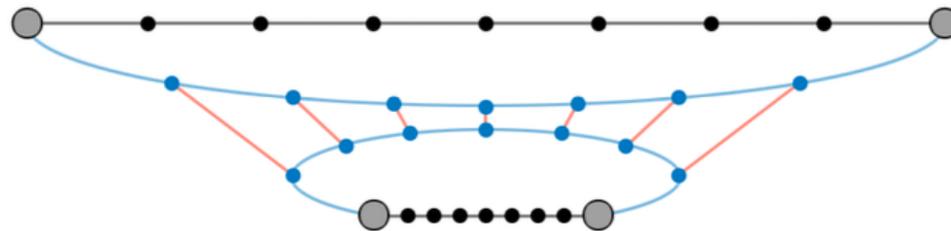
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- the larger  $\alpha$ , the smaller  $C_a(P, Q)$
- $C_a(P, Q) = 0$  if  $\alpha = 90^\circ$  and  $C_a(P, Q) = 1$  if  $\alpha = 0$ , i.e.  $P$  and  $Q$  are parallel



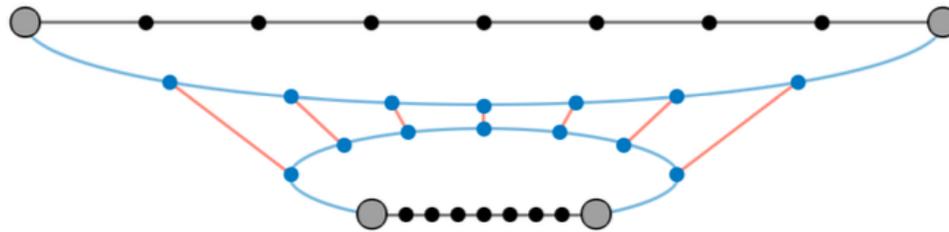
# Edge compatibility measures: scale

- edges that differ a lot in length should not be bundled together



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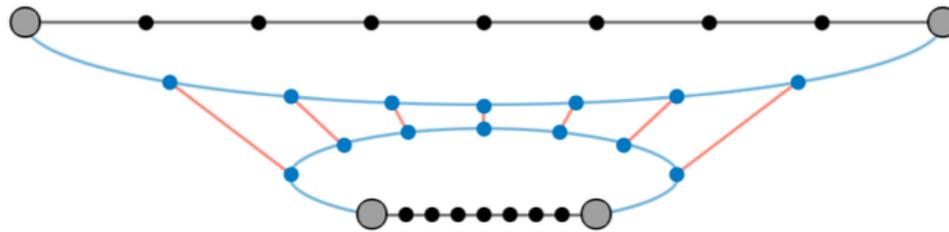
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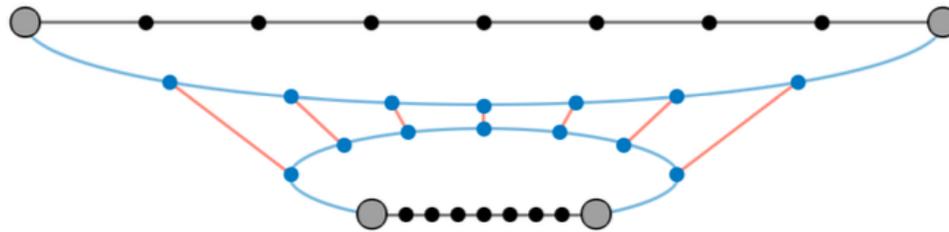


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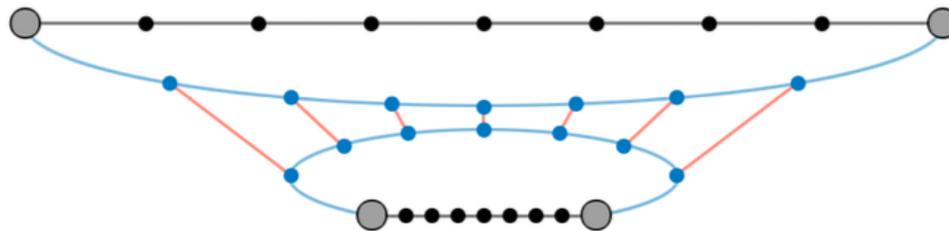


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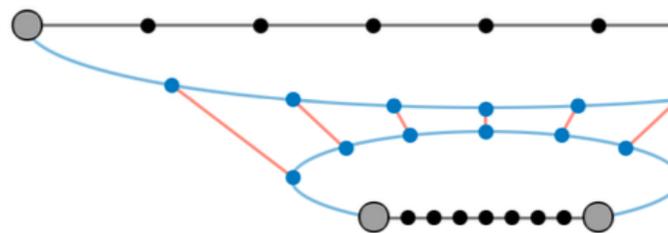


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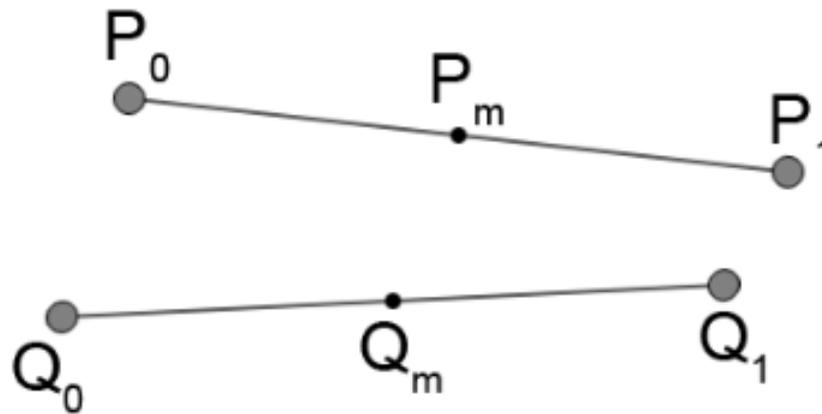


## !!!Correction:

Set  $|P| = |P| / \min(|P|, |Q|)$   
and  $|Q| = |Q| / \min(|P|, |Q|)$  -  
normalize so that shortest  
has length one, otherwise  
 $C_s(P, Q) > 1$

# Edge compatibility measures: distance

- edges that are far apart should not be bundled together

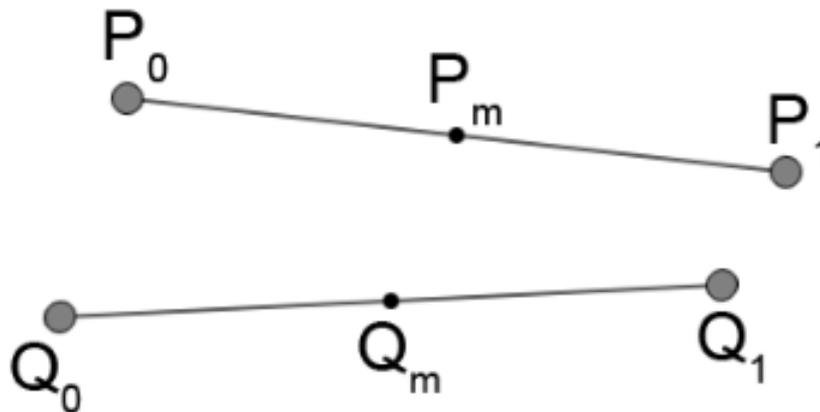


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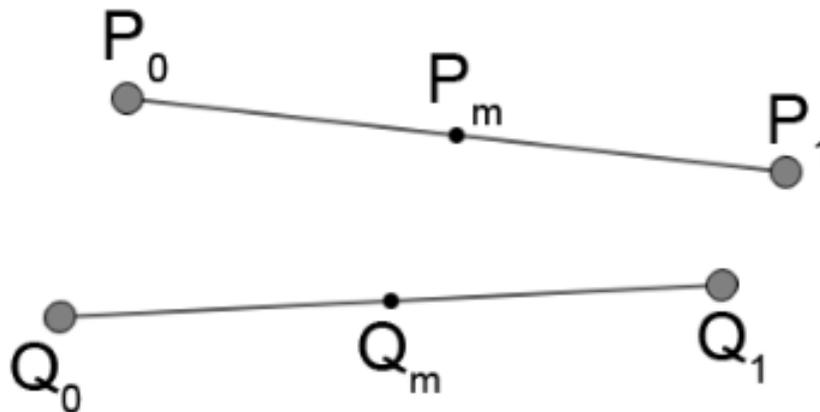
$$C_p(P, Q) = \ell_{\text{avg}} / (\ell_{\text{avg}} + \|P_m - Q_m\|),$$

with  $P_m$  and  $Q_m$  – midpoints of  $P$  and  $Q$ , respectively



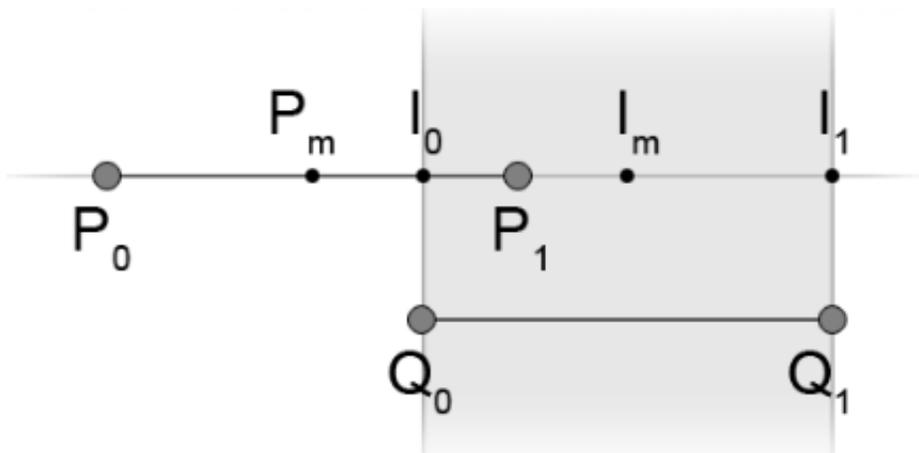
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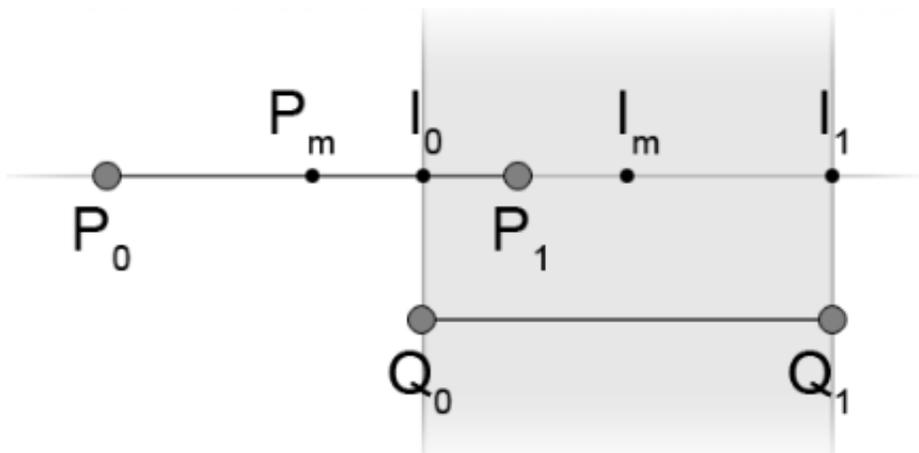
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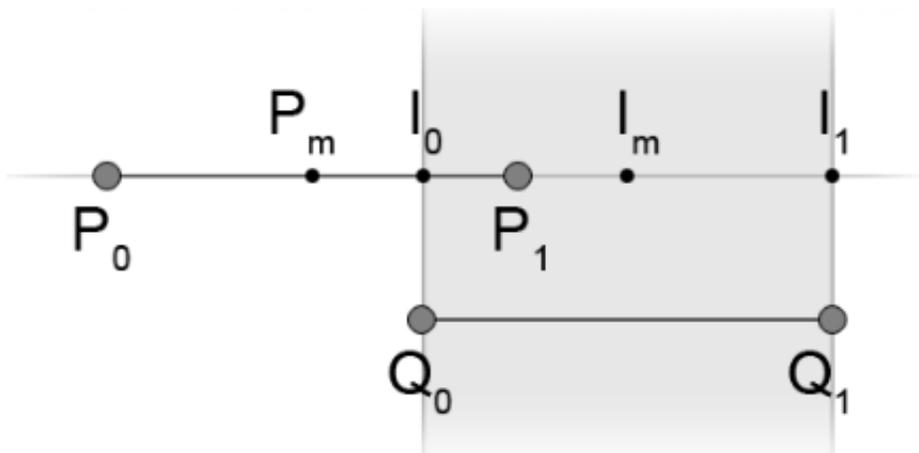
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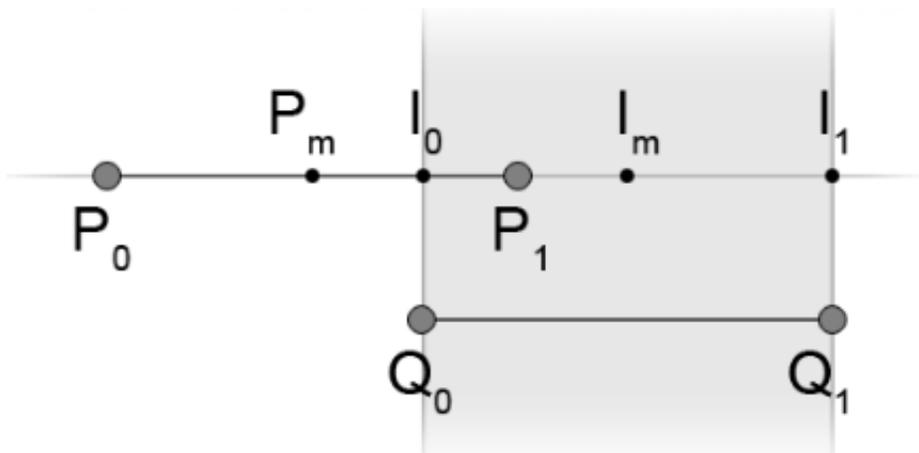


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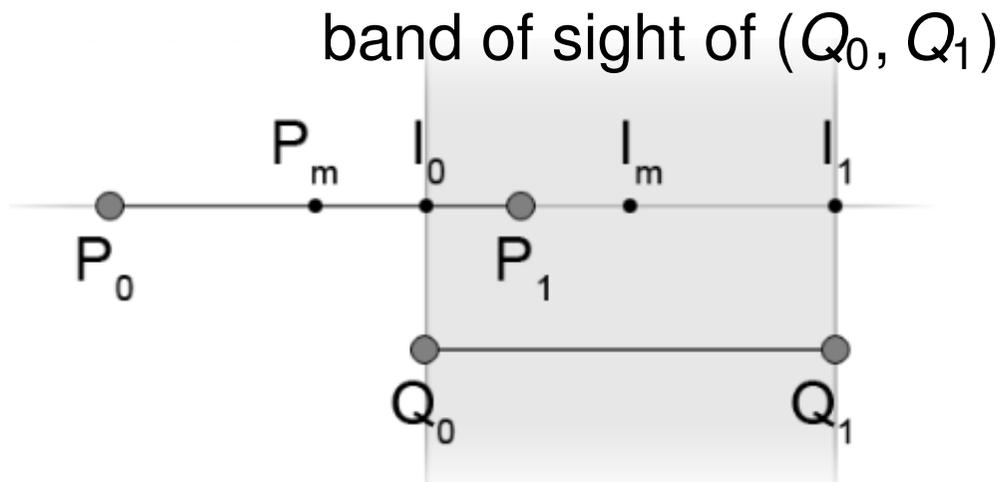


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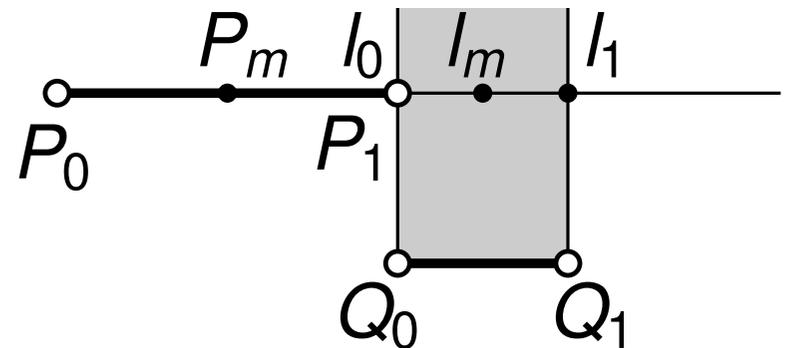
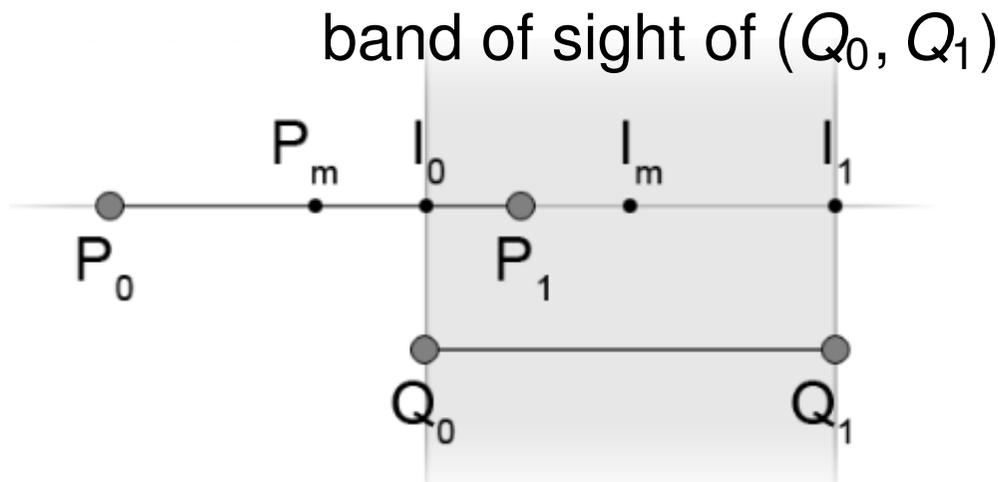


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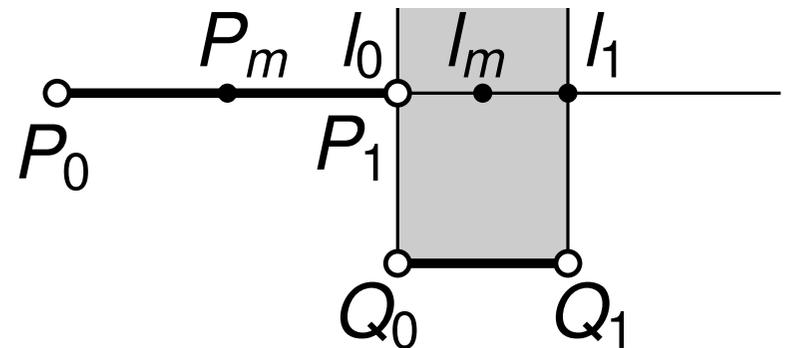
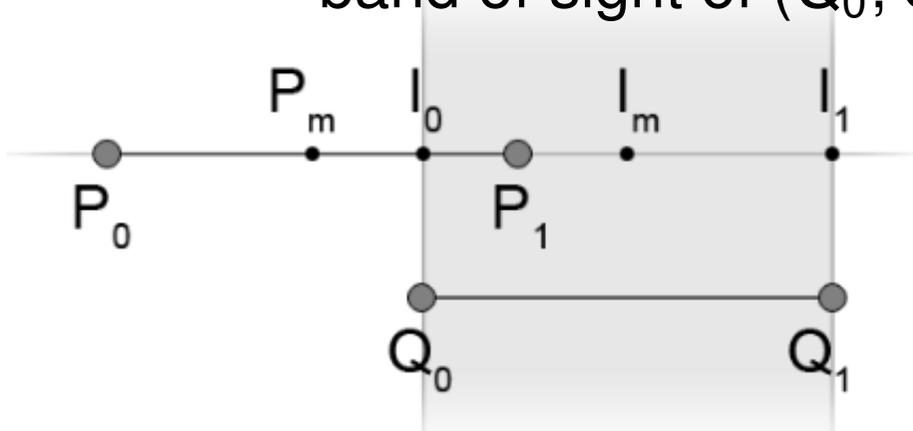
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band of sight of  $(Q_0, Q_1)$



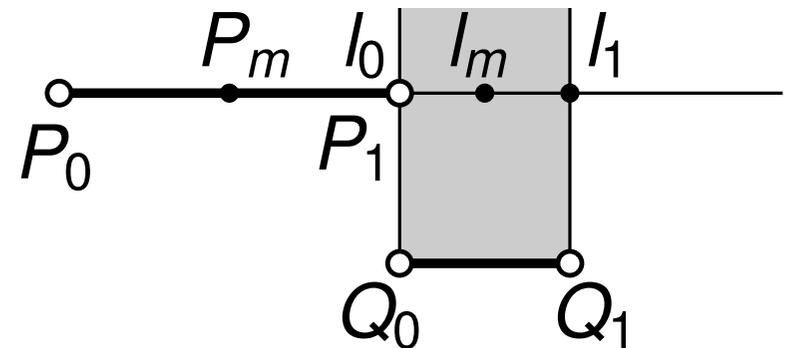
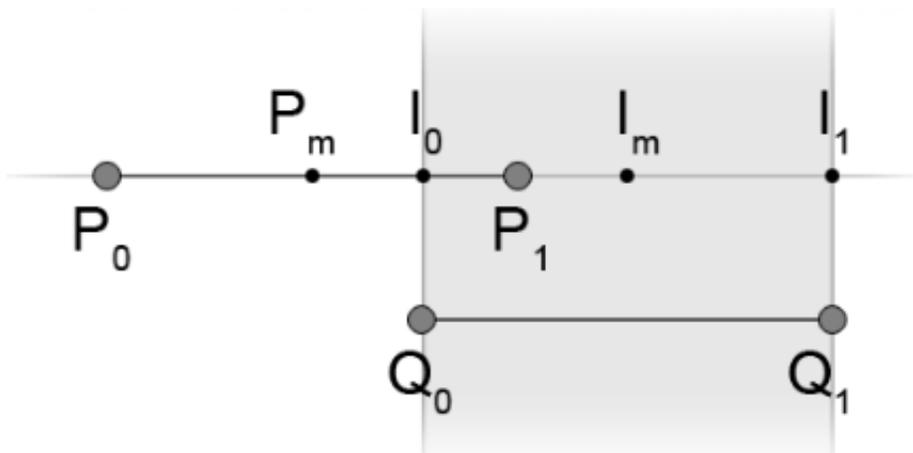
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# Edge compatibility measures: combined

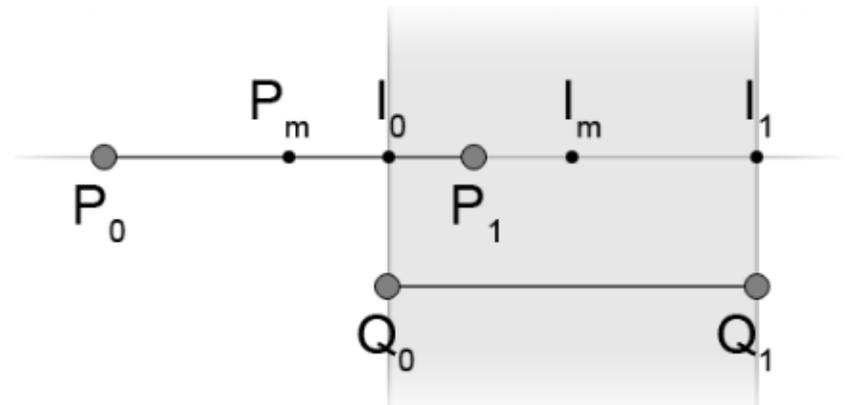
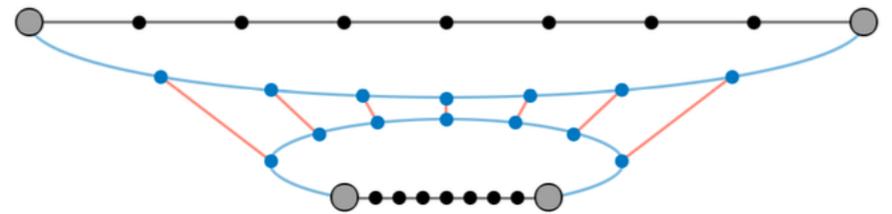
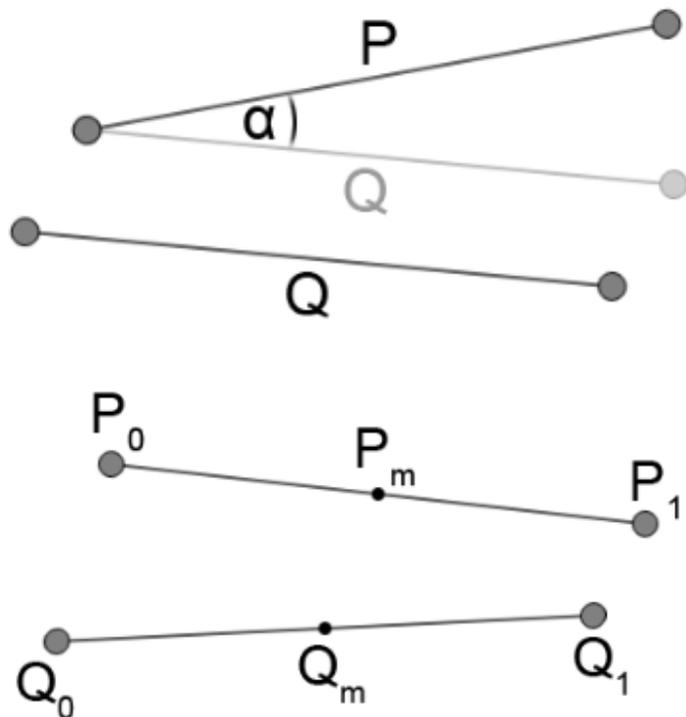
- The overall compatibility is defined as

$$C_e(P, Q) = C_a(P, Q) \cdot C_s(P, Q) \cdot C_p(P, Q) \cdot C_v(P, Q)$$

- The overall force on point  $p_i$  is then redefined as

$$F_{p_i} = k_P \cdot (\|p_{i-1} - p_i\| + \|p_i - p_{i+1}\|) + \sum_{Q \in E} \frac{C_e(P, Q)}{\|p_i - q_i\|}$$

$k_P$  – constant for edge  $P$



# Edge bundling summary

**Input:**  $G = (V, E)$  undirected graph with vertex placement,

number of cycles  $C \in \mathbb{N}$ ,

number of iterations in the first cycle  $l_0 \in \mathbb{N}$ ,

step size  $s_0 \in \mathbb{N}$ ,

number of subdivision points in the first cycle  $n_0$

interaction function  $C_e : E \times E \rightarrow \mathbb{R}$

**Output:** Layout with bundled edges

$n \leftarrow n_0$  initial number of subdivisions

$t \leftarrow 1$  iteration counter

$l \leftarrow l_0$  number of iterations in the first cycle

$c \leftarrow 1$  cycle counter

$s \leftarrow s_0$  step size

# Edge bundling summary

**while**  $c < C$  **do**

**foreach**  $P \in E$  **do**

    subdivide  $P$  by  $n$  points  $P_1 \dots P_n$ ;  $B \leftarrow B \cup \bigcup_{P \in E} \{P_1 \dots P_n\}$

**foreach**  $P \in E$  **do**

**foreach**  $0 < i < n$  **do**

$$F_{P_i} = k_P \cdot (\|P_{i-1} - P_i\| + \|P_i - P_{i+1}\|)$$

**foreach**  $Q \neq P \in E$  **do**

**foreach**  $0 < j < n$  **do**

$$F_{P_i} = F_{P_i} + \frac{C_e(P, Q)}{\|P_i - Q_j\|};$$

**foreach**  $p \in B$  **do**

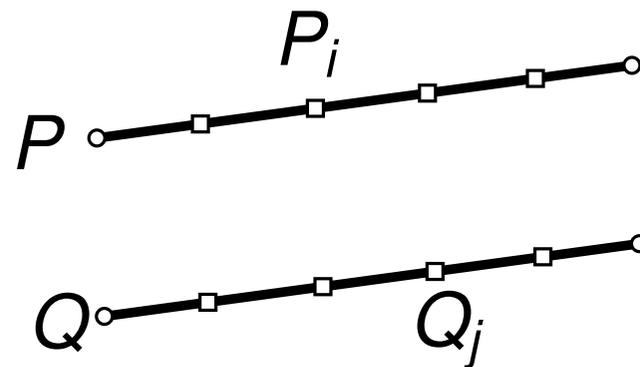
$p \leftarrow p + s \cdot F_p$

$t \leftarrow t + 1$

**if**  $t == l$  **then**

$t \leftarrow 1$ ;  $c \leftarrow c + 1$ ;  $n \leftarrow 2n$ ;

$s \leftarrow s/2$ ; **decrease**( $l$ );



# Edge bundling: experiments

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 $n_0 = 1$ ,  $s_0 = 0.04$ ,  $C = 6$ , and  $l_0 = 50$ .

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cycle	0	1	2	3	4	5
$n$	1	2	4	8	16	32
$S$	.04	.02	.01	.005	.0025	.00125
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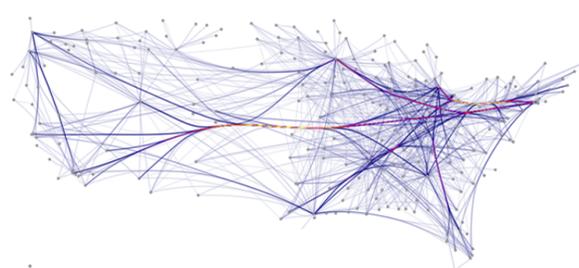
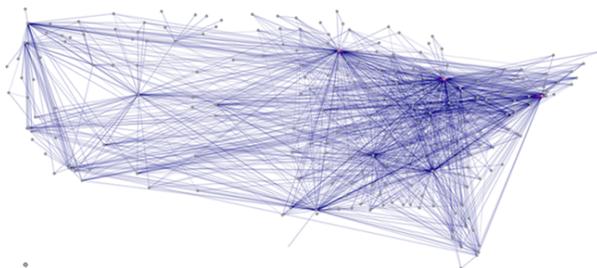
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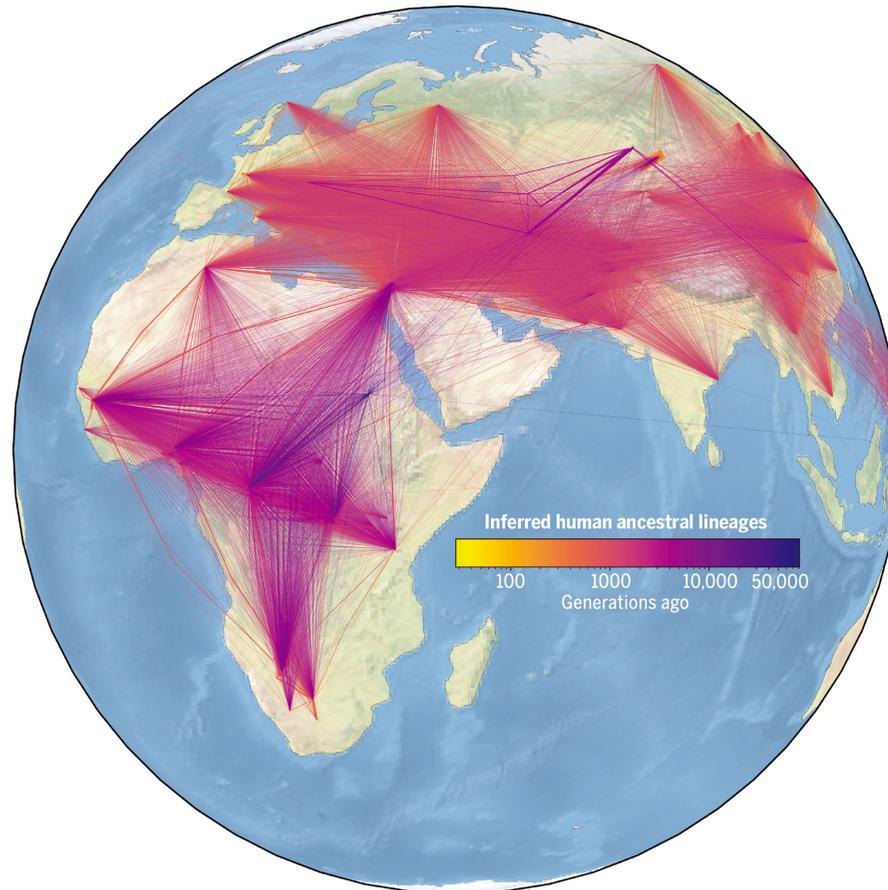
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- US airlines graph with inverse linear and inverse quadratic model



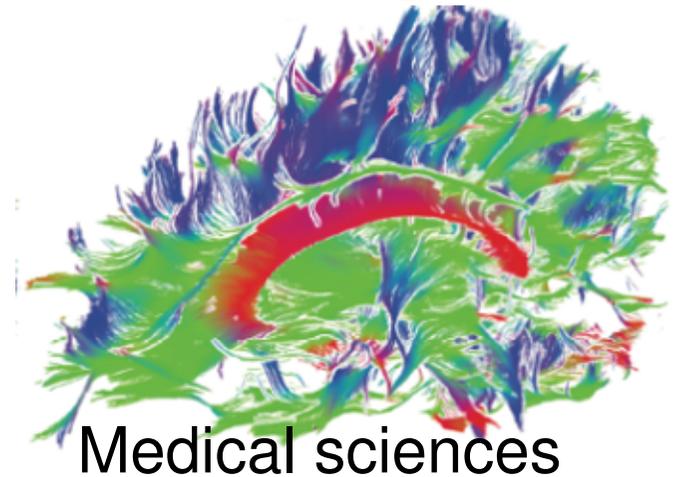
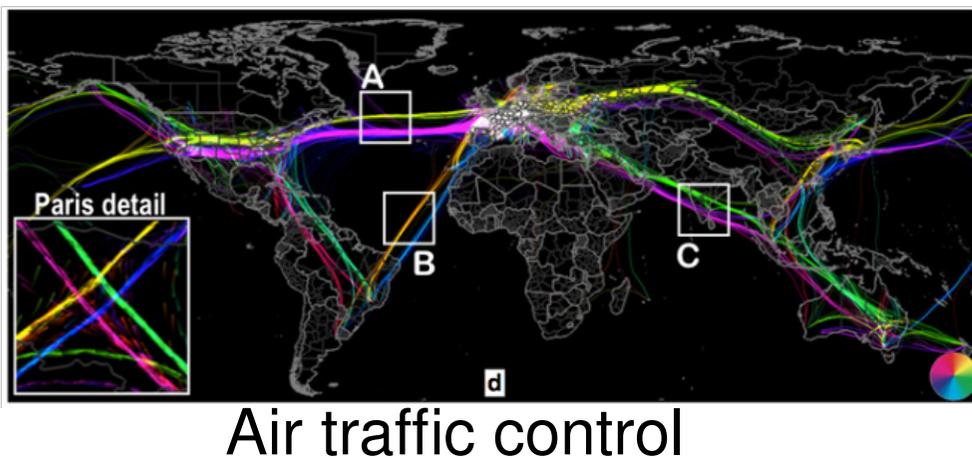
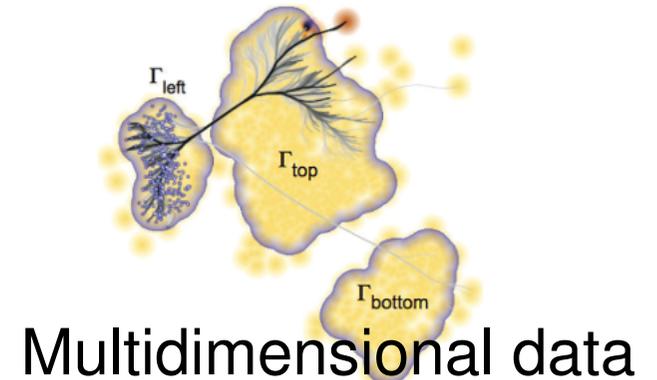
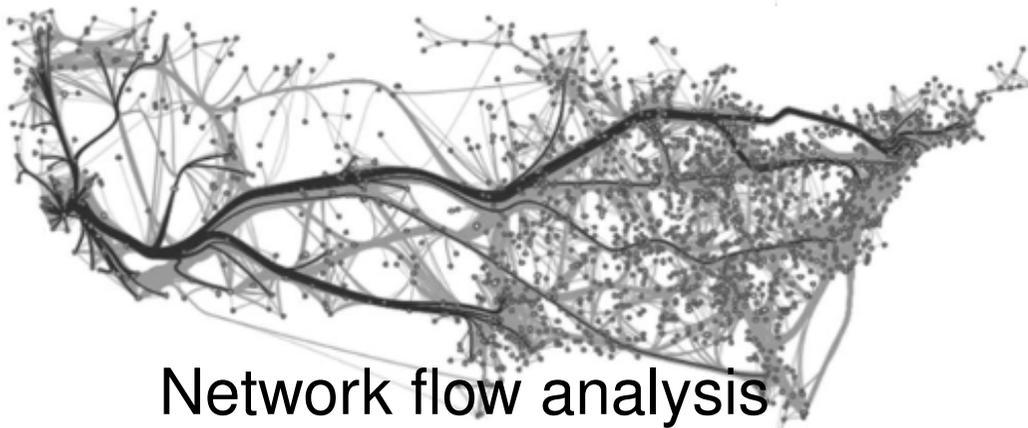
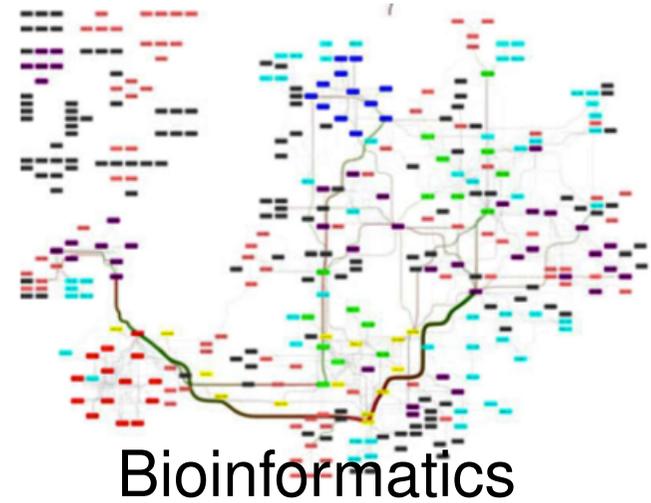
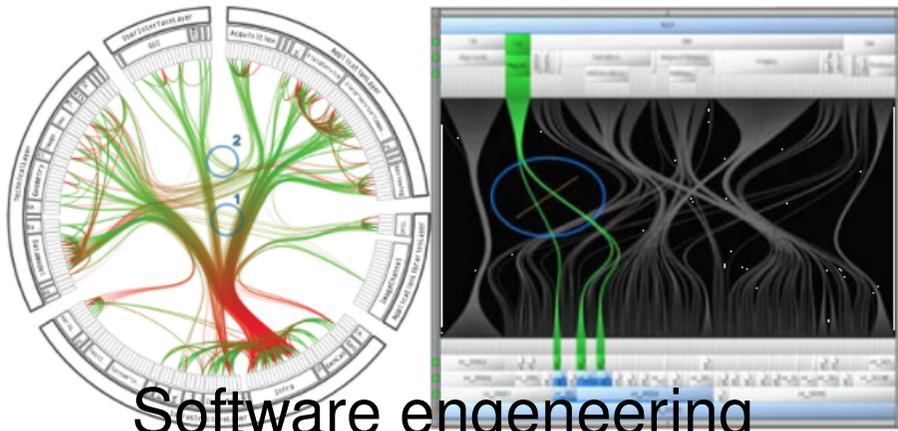
# Edge bundling: inspiration

Inspiration: edges are ancestor-descendant relationship in the genealogy of modern and ancient genomes. Edge width – how many times the relationship is observed, color – age of the ancestor

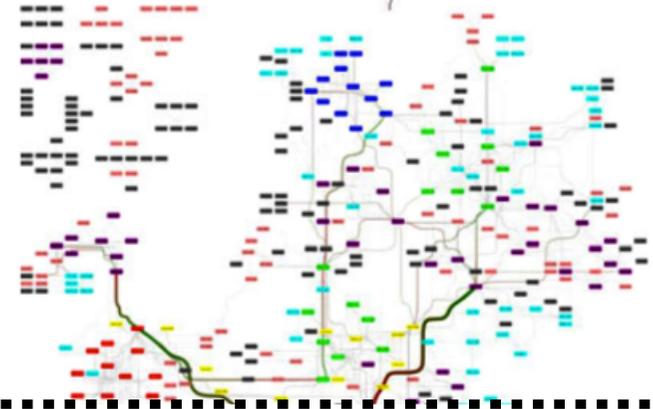
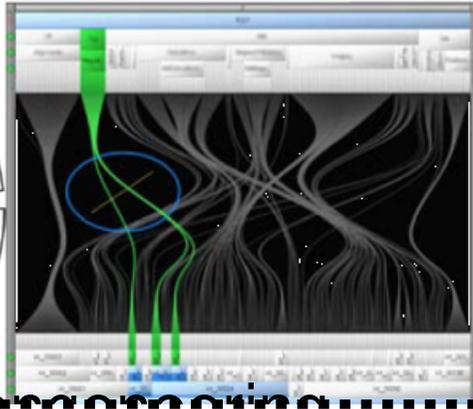
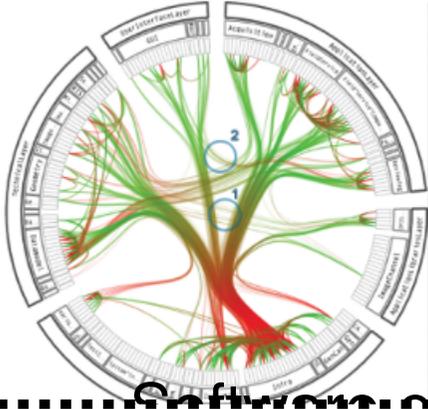


A unified genealogy of modern and ancient genomes, Wohns et al.  
Nature 2022

# Edge bundling: discussion



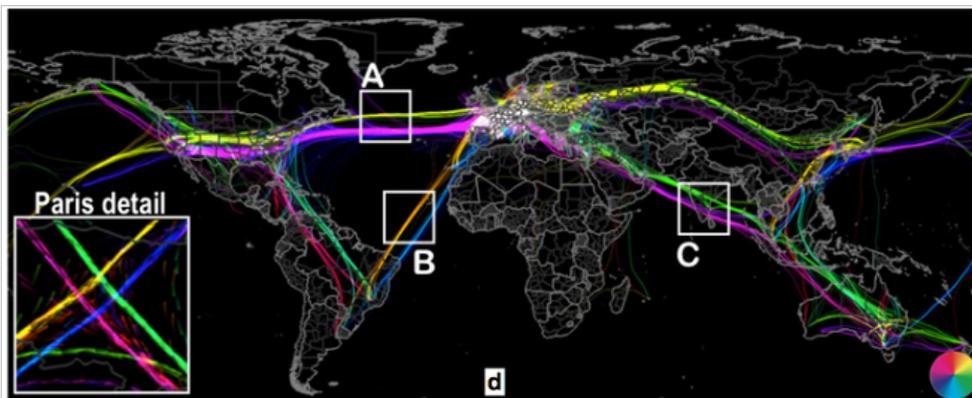
# Edge bundling: discussion



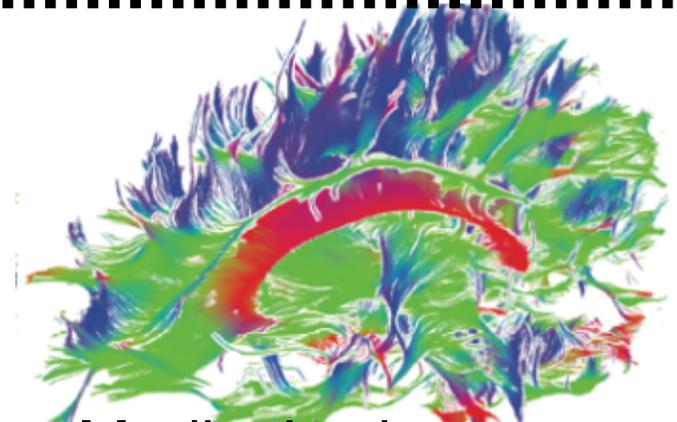
Software engineering



- What are the benefits and the drawbacks of the bundled layouts?
- When are the edge bundling techniques appropriate to use?



Air traffic control



Medical sciences

# Tutorial task

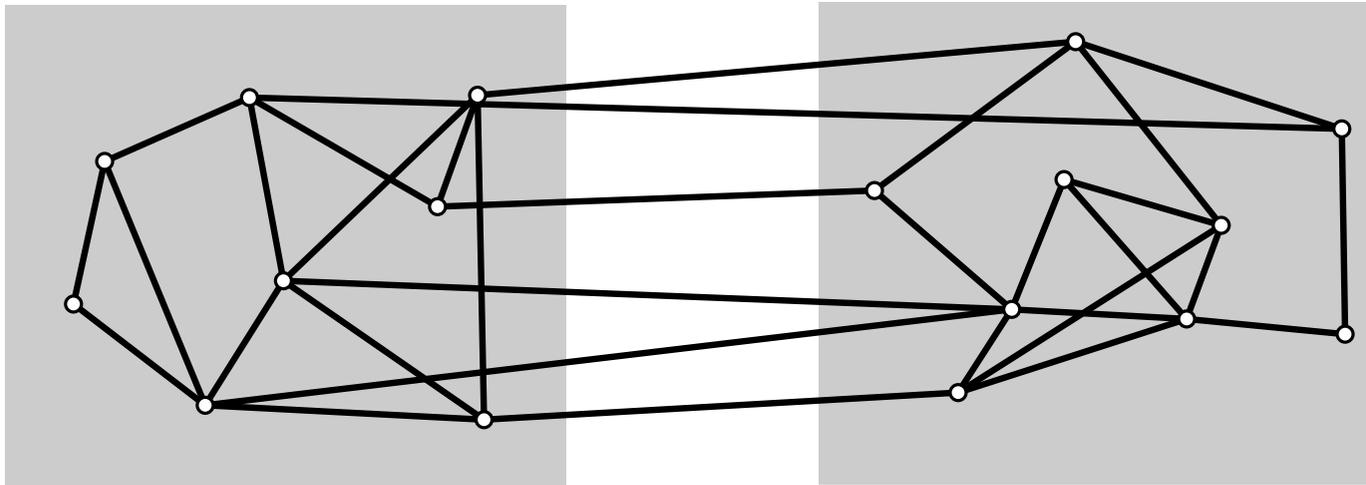
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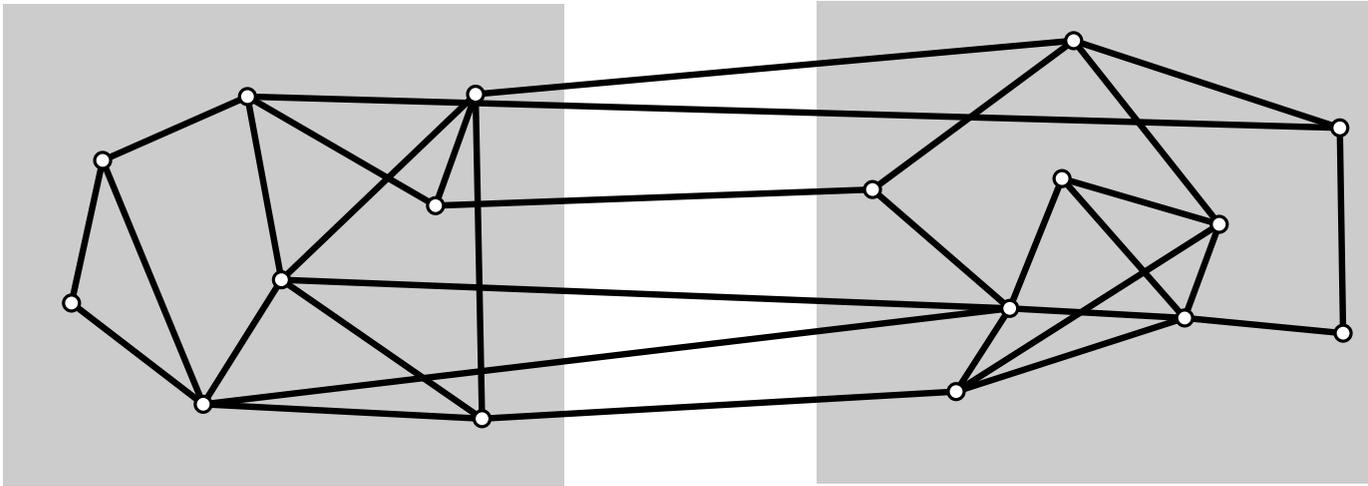
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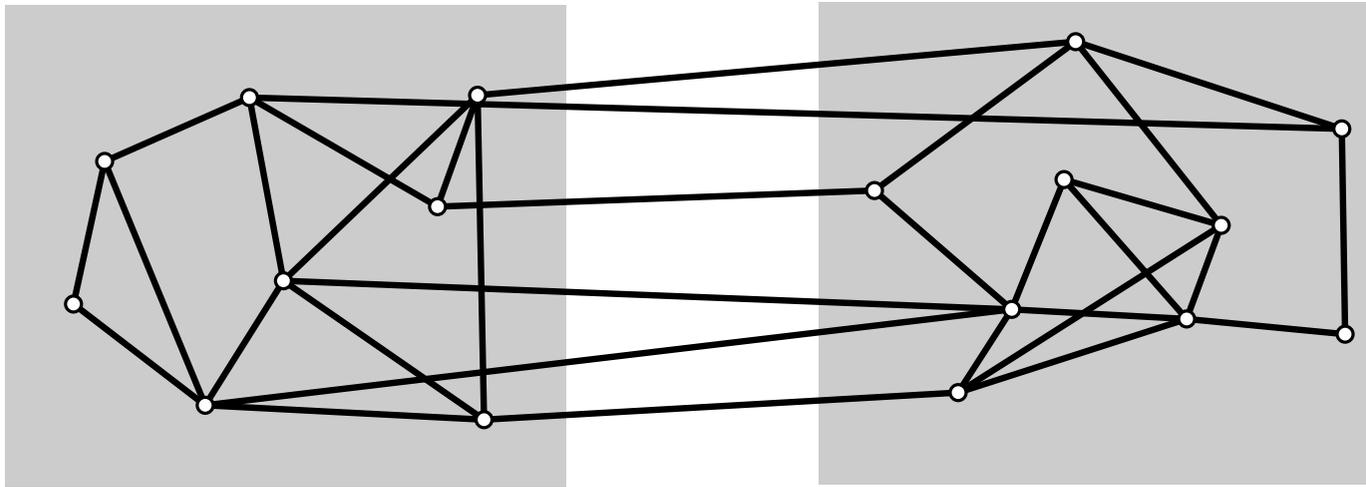
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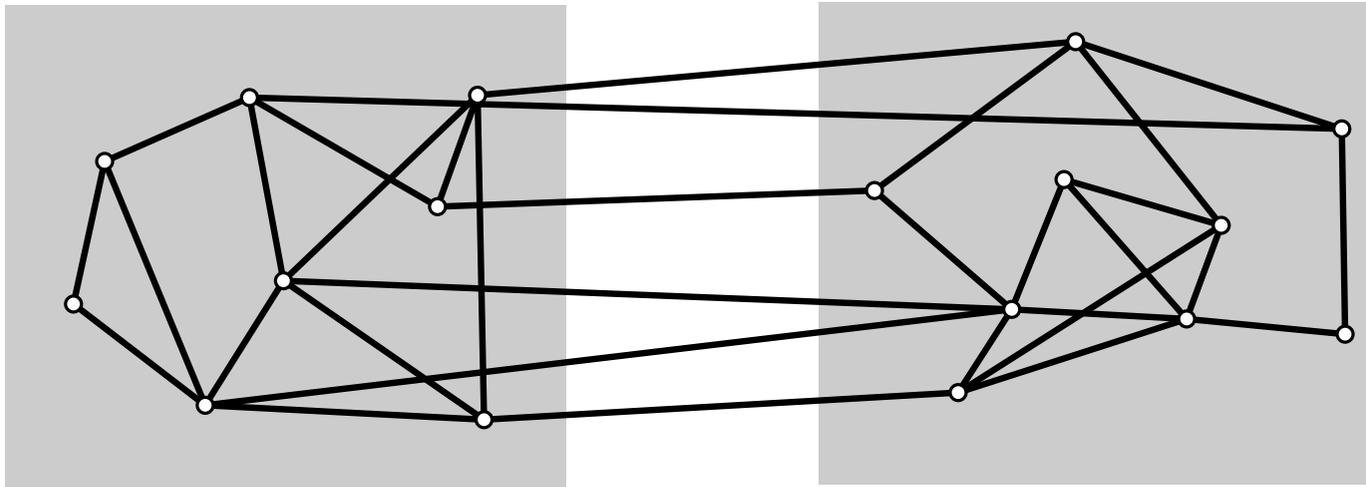
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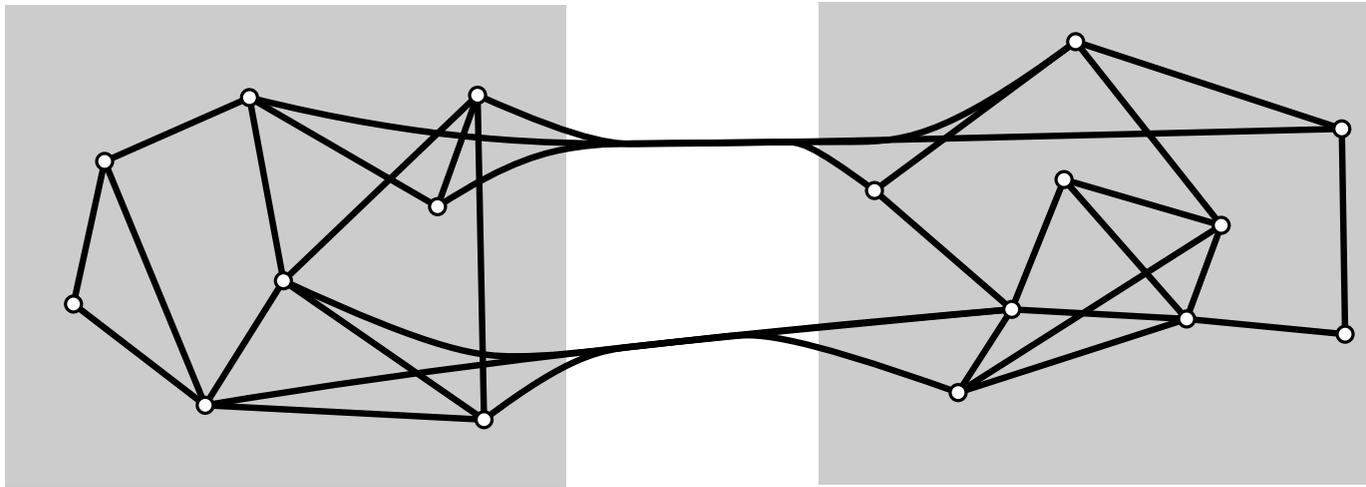
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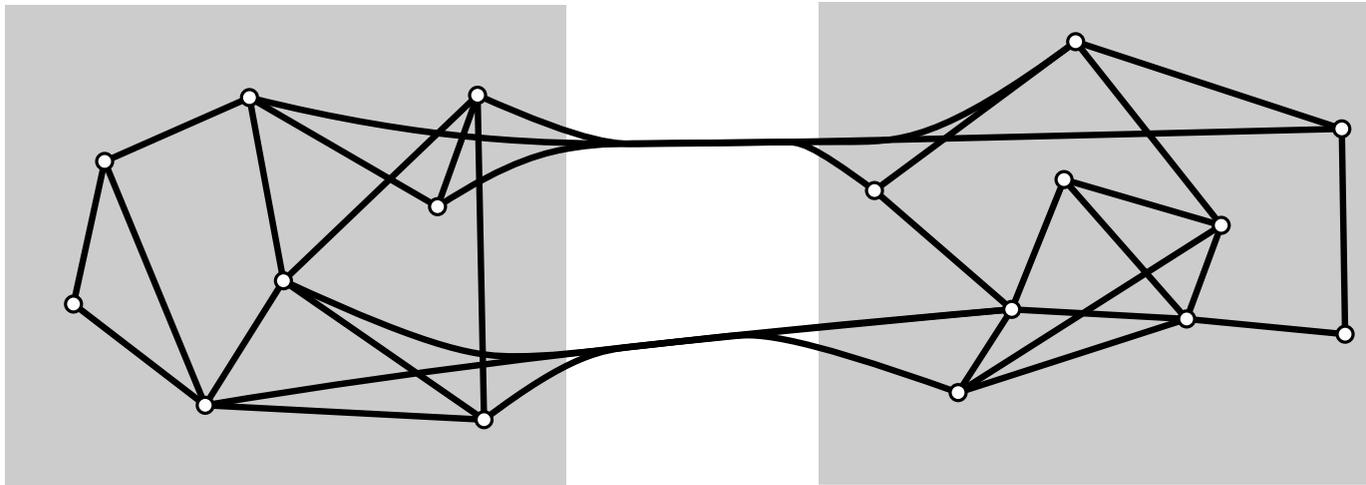
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- experiment with political blogosphere, argument network (besides the two clusters, nodes and edges have different types)



# Reading and Next



## Additional Reading

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## Next

	March 6	-	Visualization of multilevel networks	Tamara	
11	March 11	-	Tutorial: Step 5	Alister	Step 4
	March 13	-	High-dimensional data visualization	Alex	
12	March 18	-	Tutorial: Step 6	Alister	Step 5
	March 20	-	High-dimensional data visualization: advanced	Alex	
13	March 25	-	Tutorial: Step 7	Alister	Step 6
	March 27	-	—		
14	April 2	17:15-19:00	Final Presentations	Students	Step 7
	April 3	9:00-12:45	Final Presentations	Students	
	April 8				Final deliverables

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