



**Cody Dunne**  
Northeastern University

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TREES & NETWORKS

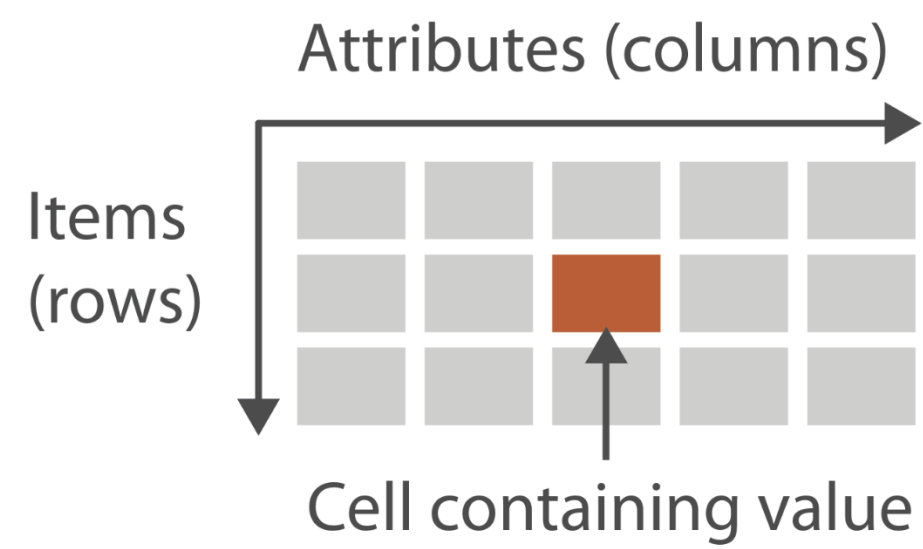
# GOALS FOR TODAY

- Learn the definition of a network (including node, edge)
- Learn the definition of a tree
- Learn common visual encoding techniques for network data (i.e., node-link diagram, adjacency matrix), and the advantages of each one.

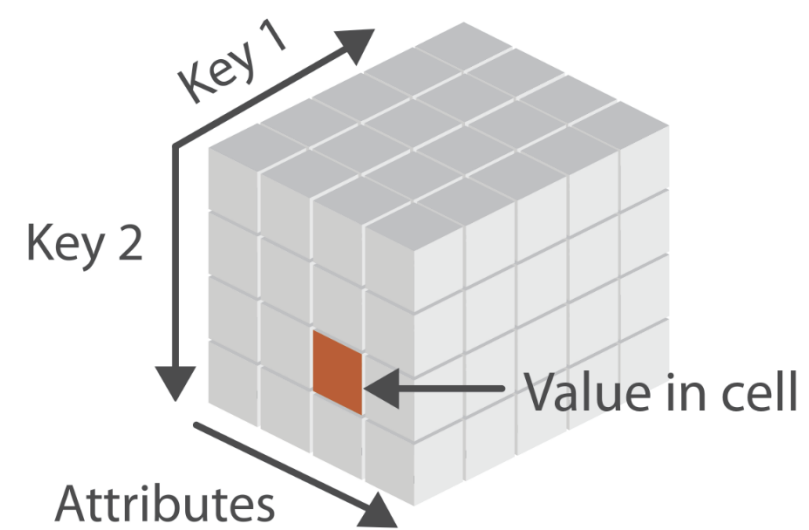
# Networks

# ➔ Dataset Types

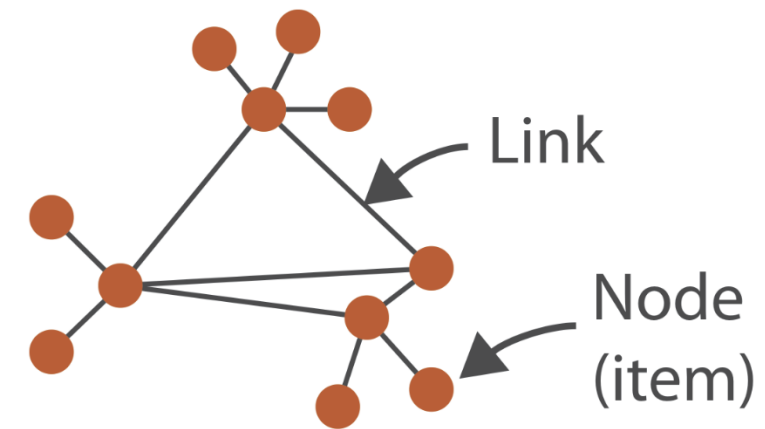
➔ Tables



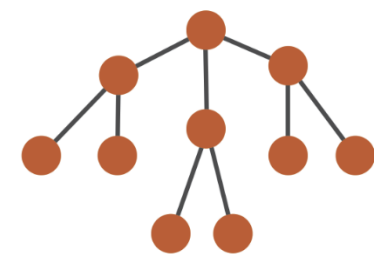
➔ *Multidimensional Table*



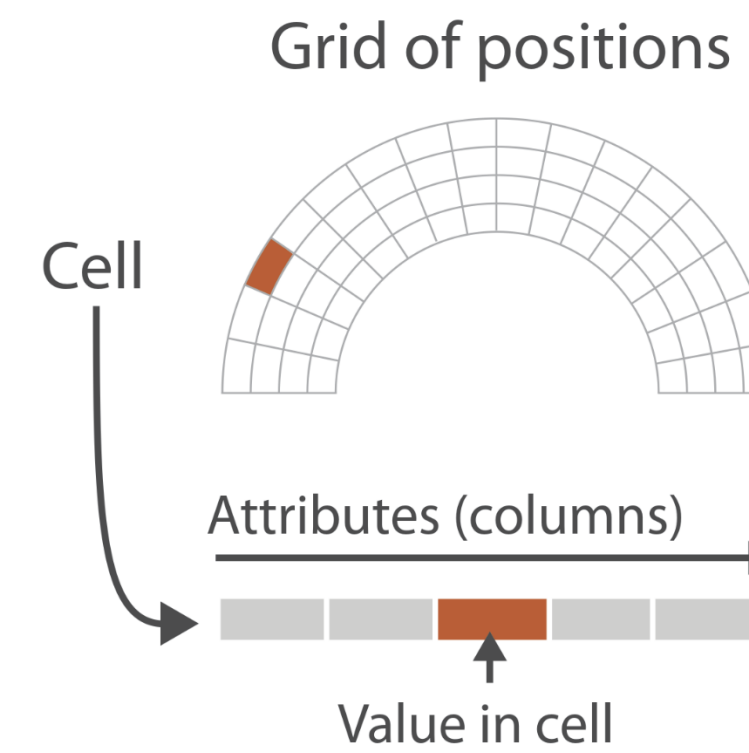
➔ Networks



➔ Trees



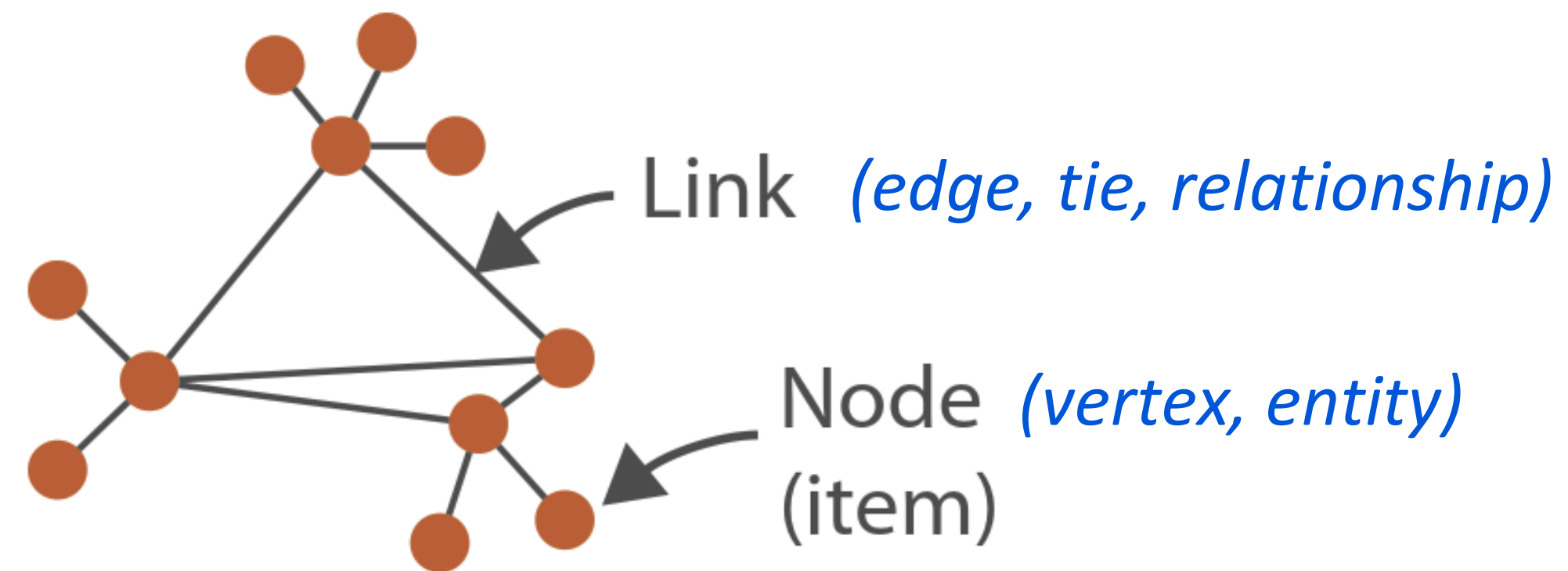
➔ Fields (Continuous)



➔ Geometry (Spatial)

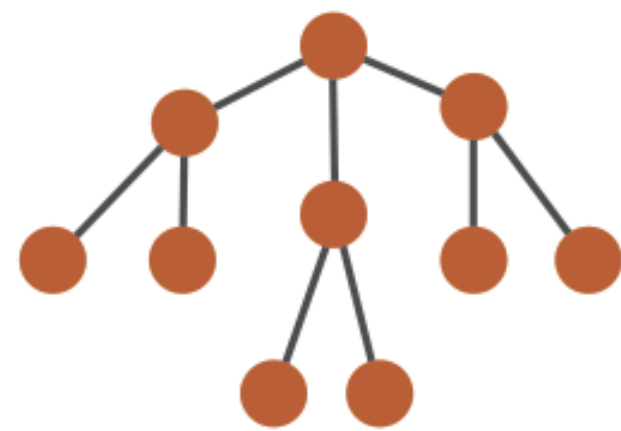


## → Networks *(graphs)*



Network = entities and relationships between them

## → Trees

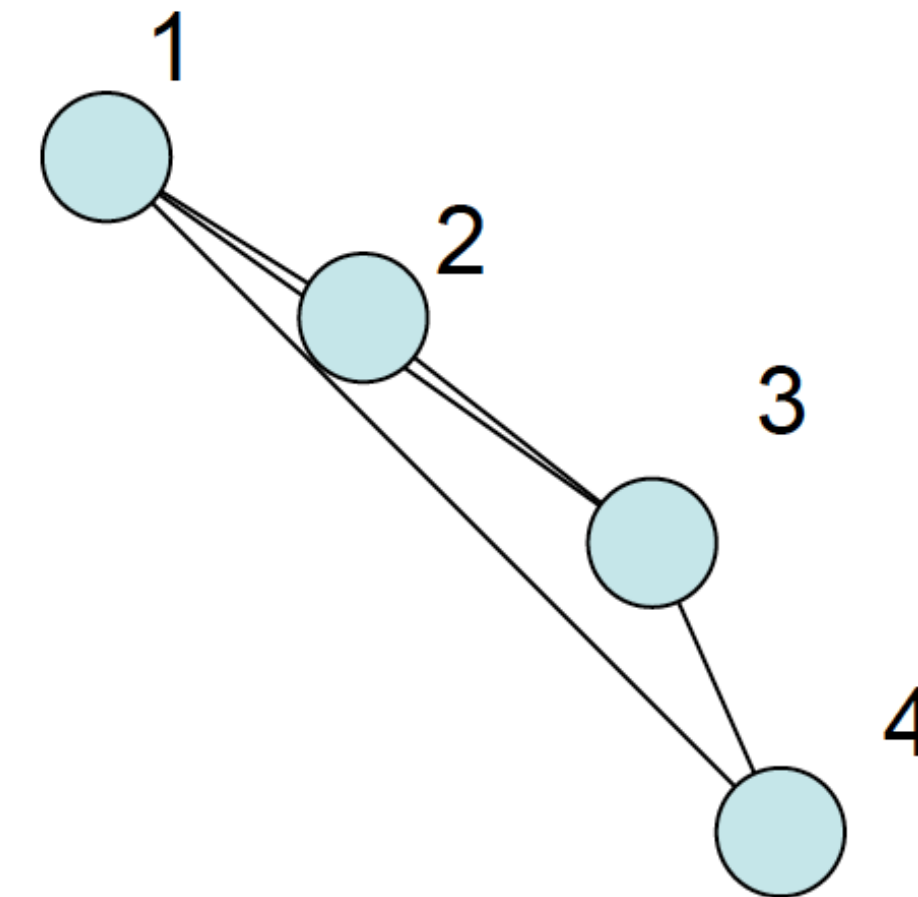
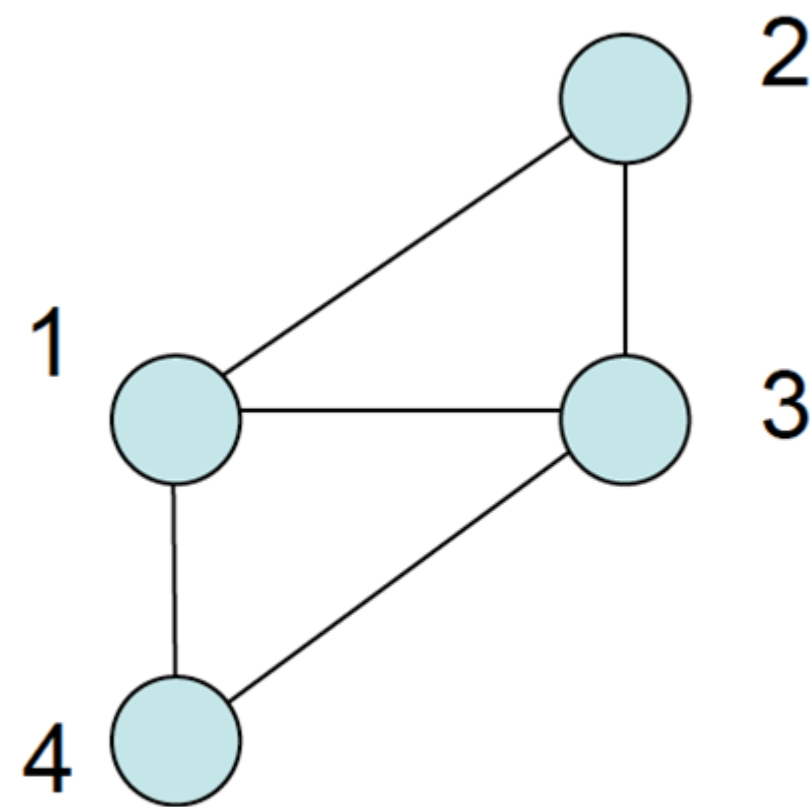
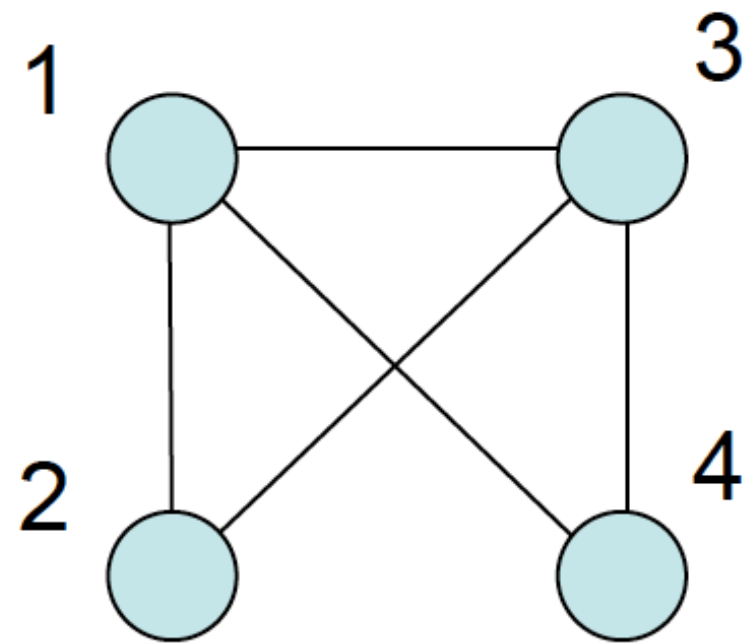


Tree = *undirected, connected, acyclic* network

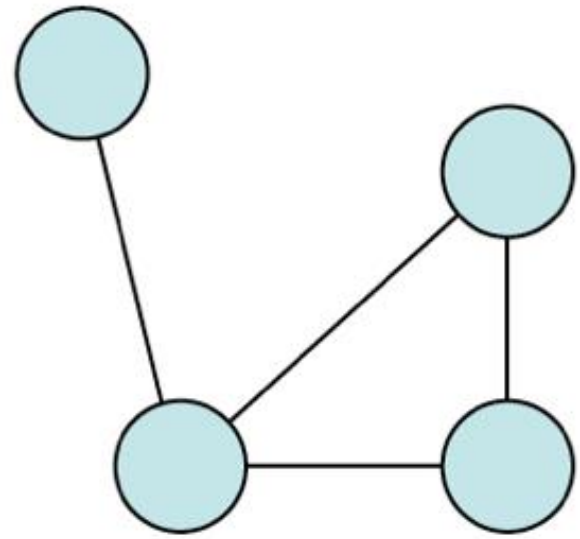
# Networks

- A network  $G$  consists of a set of nodes  $N$  and a set of edges  $E$
- An edge  $e_{n1,n2} \in E$  connects two nodes  $n1, n2 \in N$
- E.g.,  $G = \{1,2,3,4\}$ ,  $E = \{(1,2),(1,3), (2,3),(3,4),(4,1)\}$

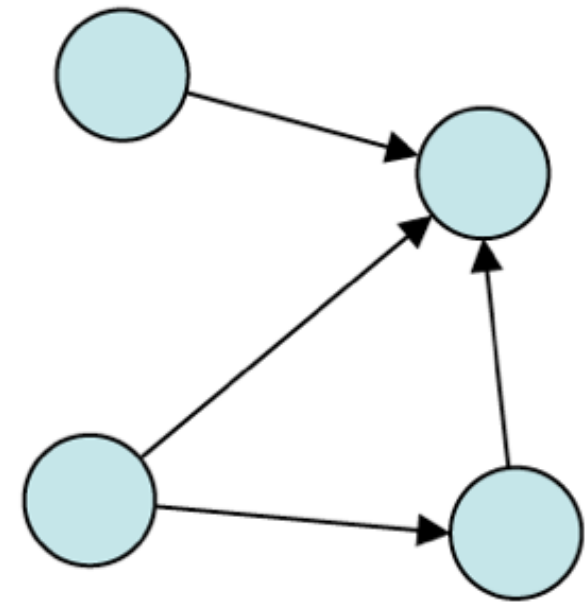
Note all the same network,  
just different layouts!



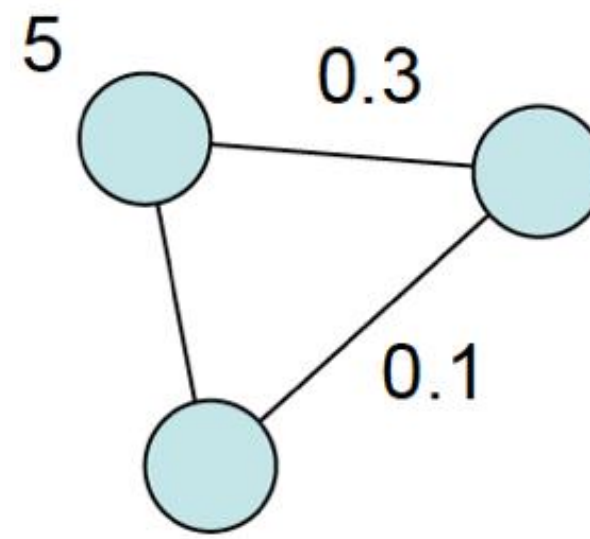
# A bunch of definitions



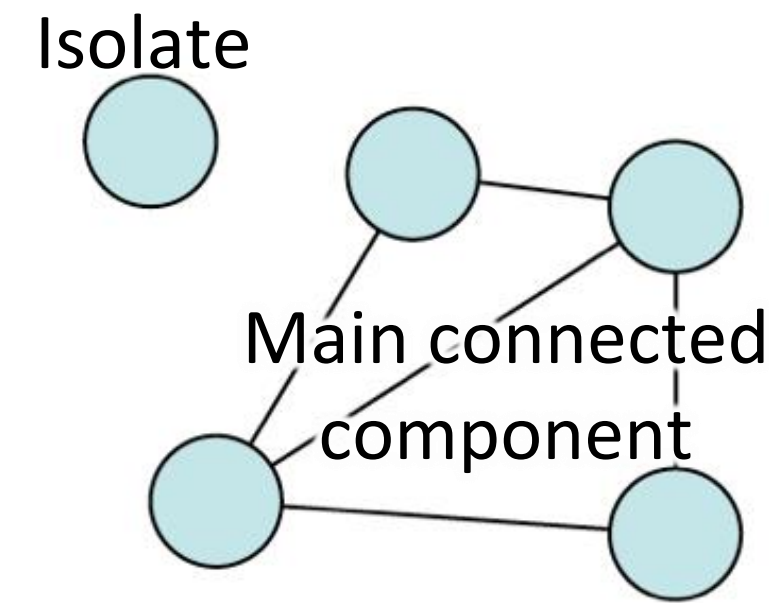
An undirected graph



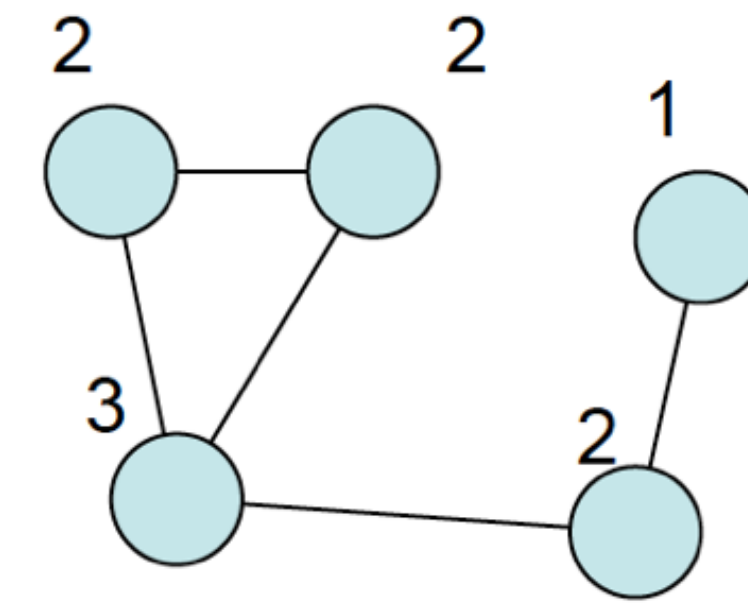
A directed graph



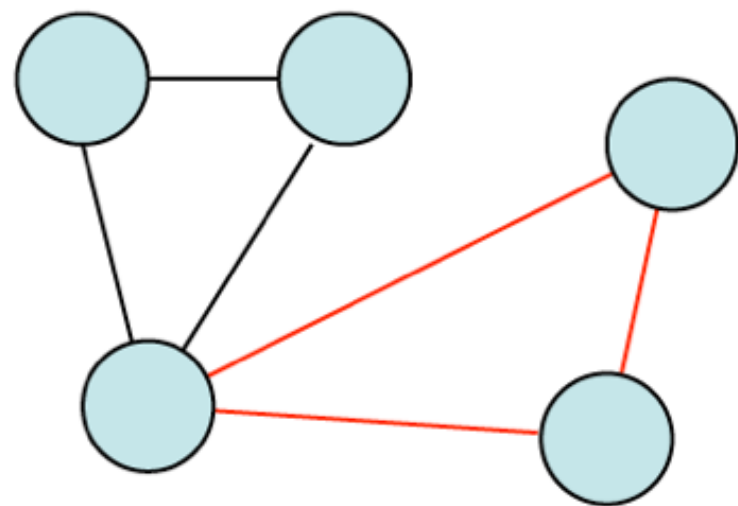
Weighted



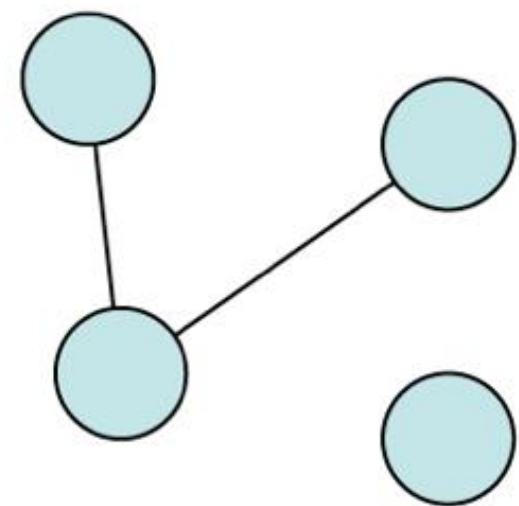
Unconnected



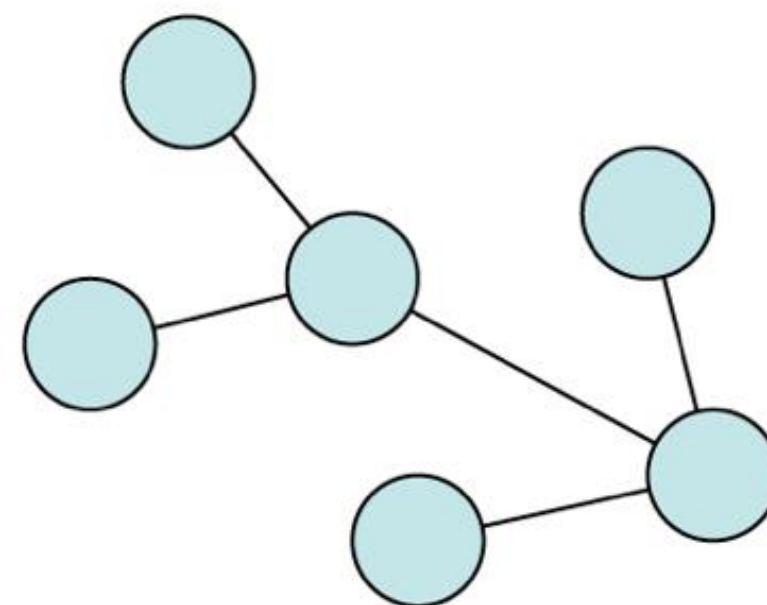
Node degrees



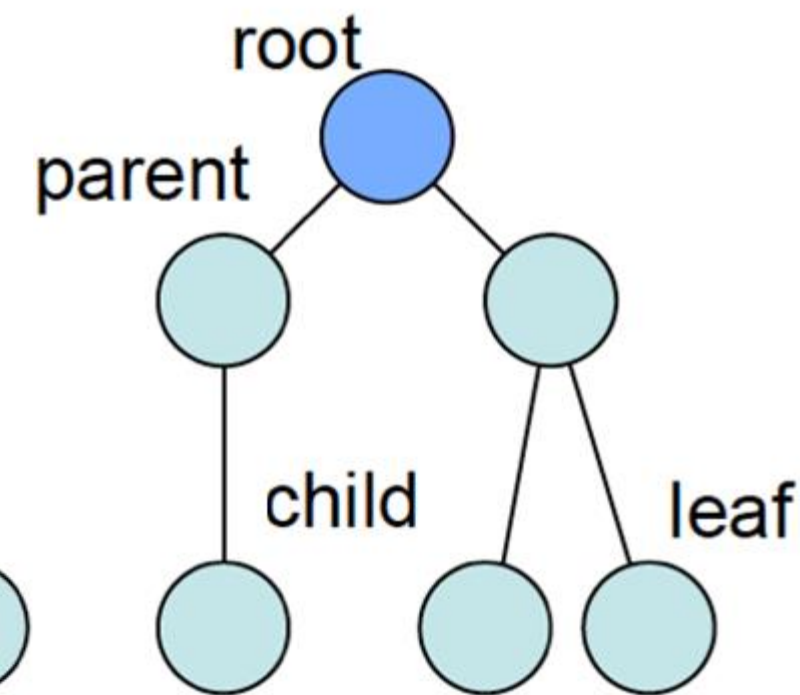
A **cycle**



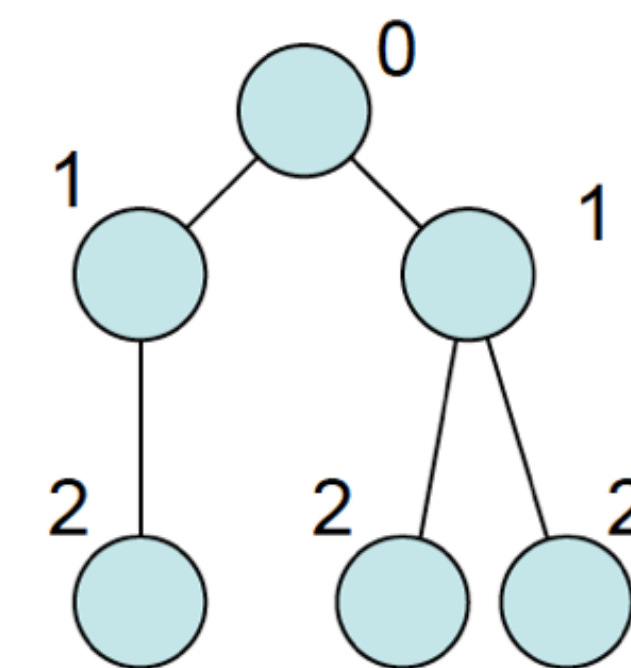
An acyclic graph



A connected acyclic graph, a.k.a. a **tree**



A rooted tree or hierarchy

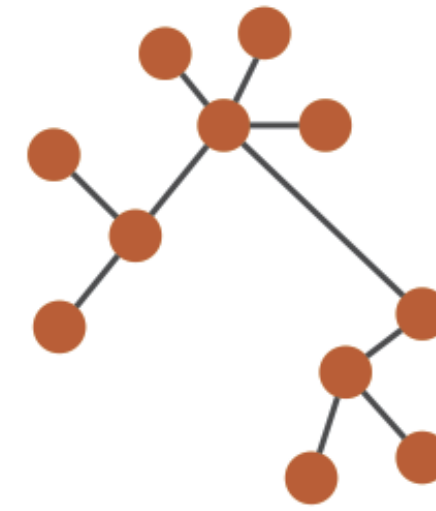


Node depths

# Arrange Networks and Trees

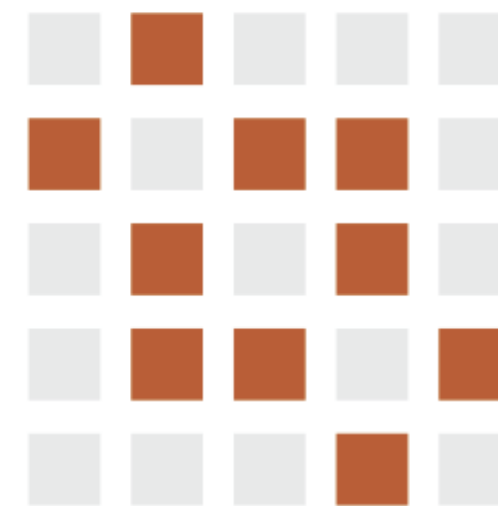
→ **Node-Link Diagrams**  
Connection Marks

✓ NETWORKS    ✓ TREES



→ **Adjacency Matrix**  
Derived Table

✓ NETWORKS    ✓ TREES



→ **Enclosure**  
Containment Marks

✗ NETWORKS    ✓ TREES



“Treemap”





# Node-Link Diagrams

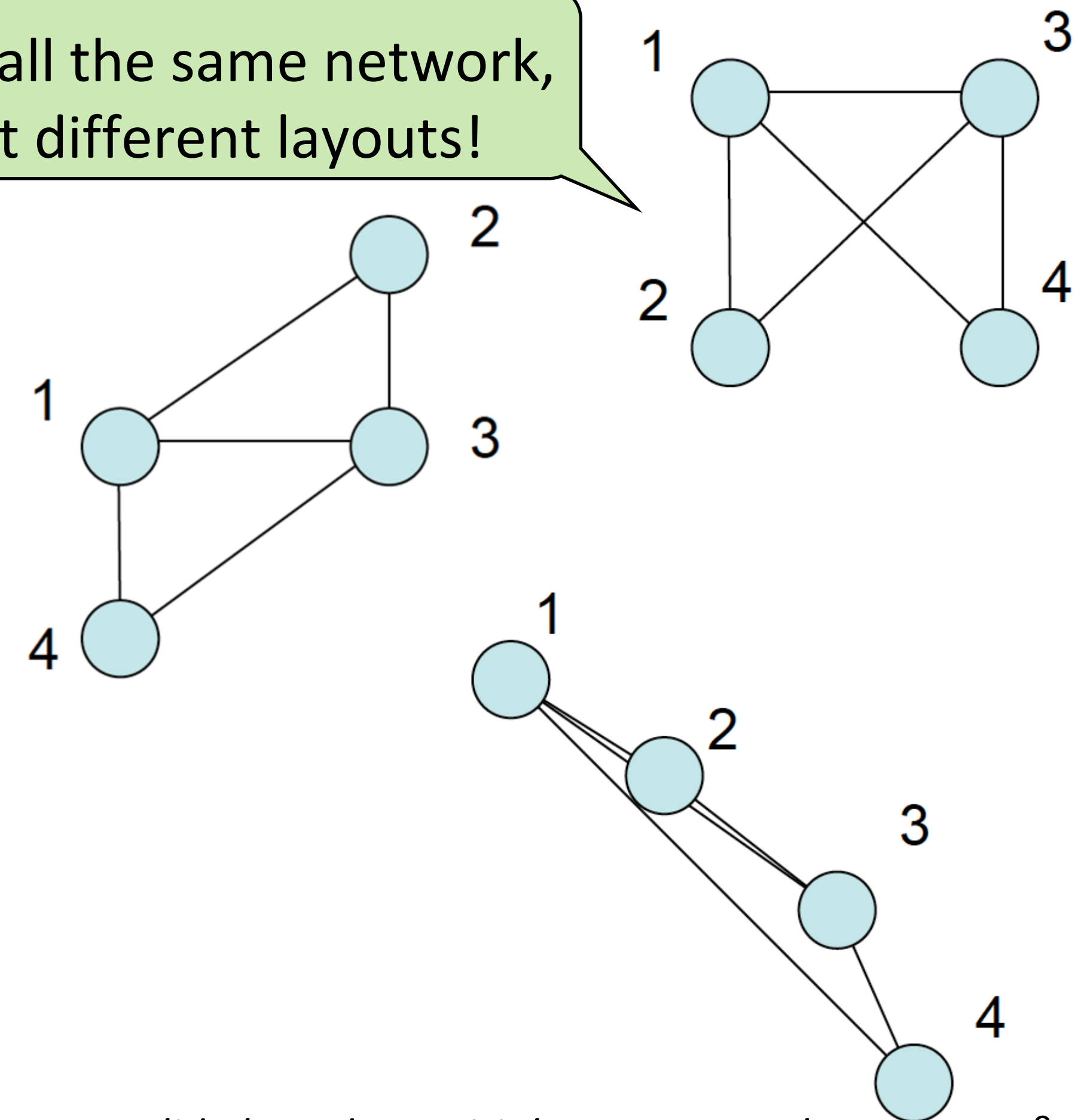
Connection Marks

- ✓ NETWORKS
- ✓ TREES

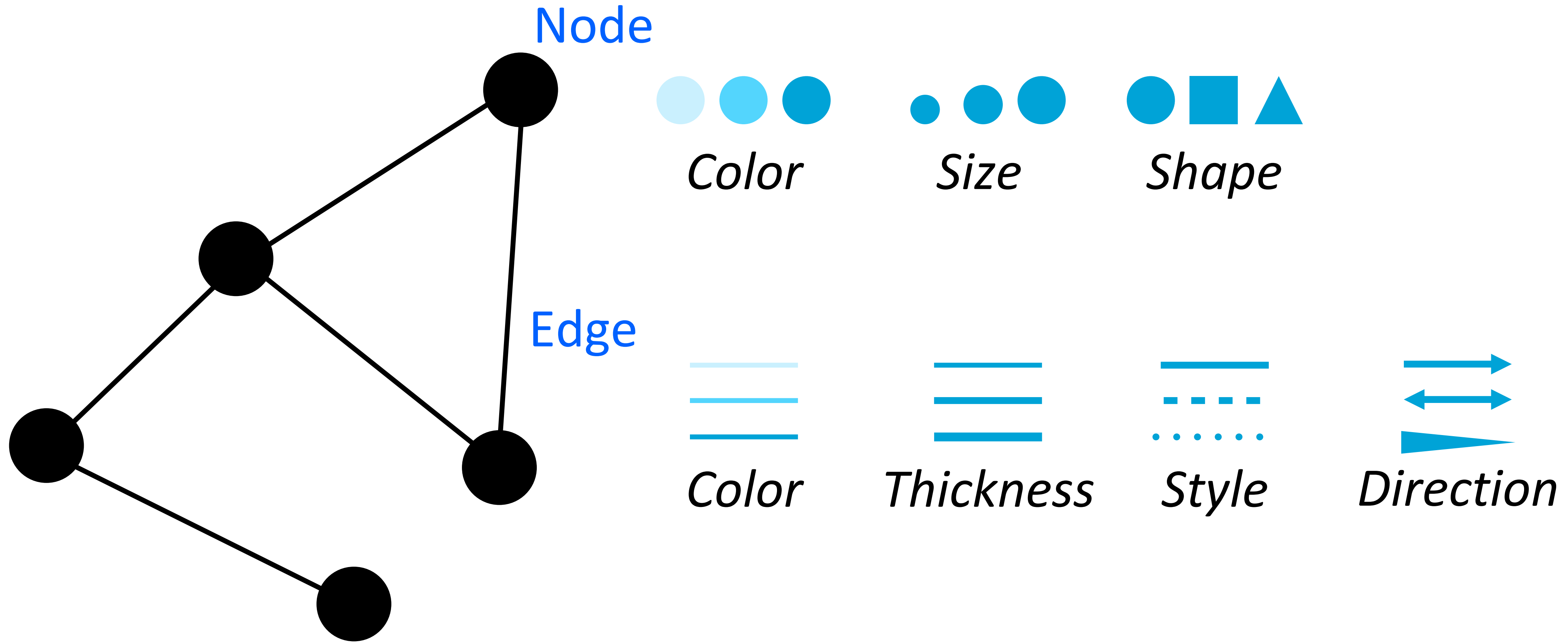


- Primary concern is the *spatial layout* of nodes and edges, a.k.a. *graph drawing*
- The goal is often to effectively depict the graph structure for *topology-based tasks*:
  - connectivity, path-following
  - network distance
  - clustering
  - ordering (e.g., hierarchy level)
- But not always topology-based tasks. E.g., understanding attributes, statistics, metrics

Note all the same network, just different layouts!



# Marks & Channels



# An Extended Evaluation of the Readability of Tapered, Animated, and Textured Directed-Edge Representations in Node-Link Graphs

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Eindhoven University of Technology

Petra Isenberg†

INRIA

Jarke J. van Wijk‡

Eindhoven University of Technology

Jean-Daniel Fekete§

INRIA

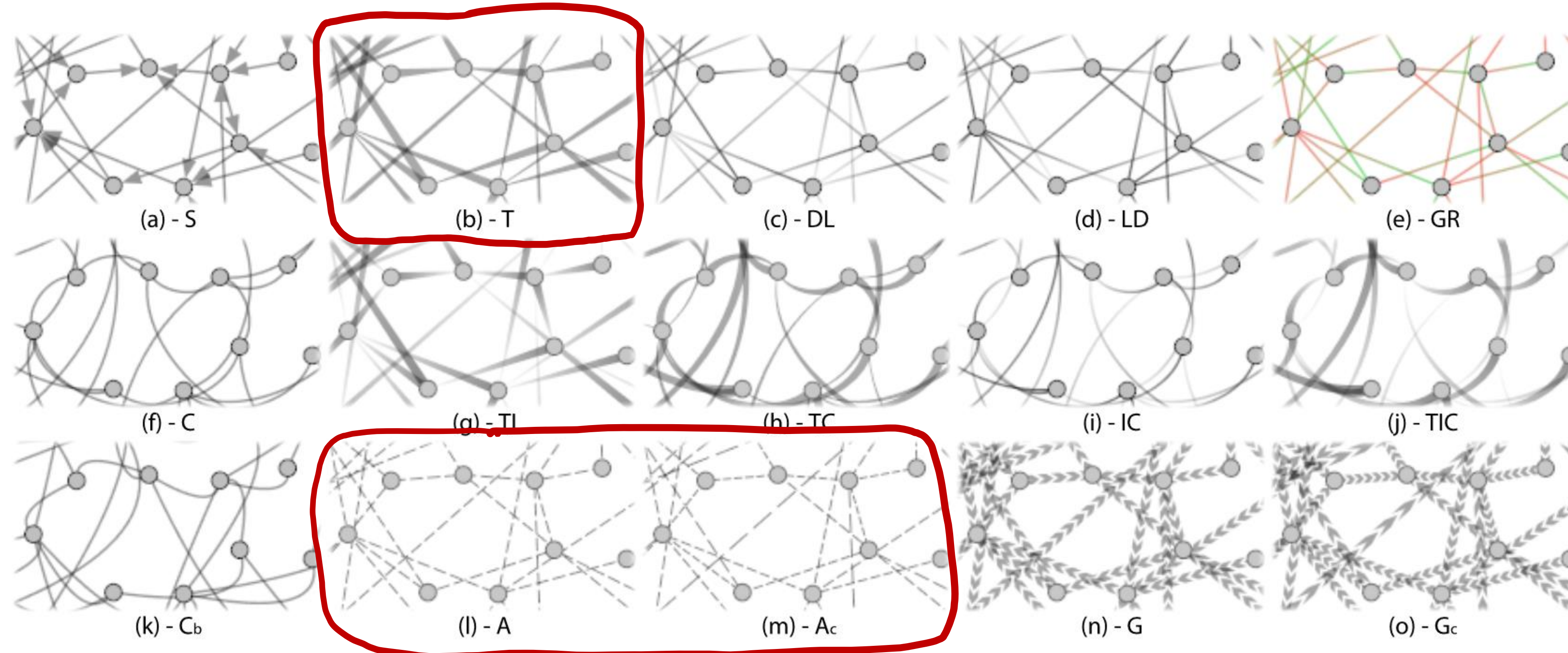


Figure 1: All directed-edge representations used in our initial (a to j), follow-up (b, k, l), and current study (b, l, m, n, o). (a) standard arrow – S, (b) tapered – T, (c) dark-to-light – DL (a.k.a intensity – I), (d) light-to-dark – LD, (e) green-to-red – GR, (f) curvature – C, (g) tapered-intensity – TI, (h) tapered-curvature – TC, (i) intensity-curvature – IC, (j) tapered-intensity-curvature – TIC, (k) biased curvature – C<sub>b</sub>, (l) animated – A, (m) animated compressed – A<sub>c</sub>, (n) glyph – G, and (o) glyph compressed – G<sub>c</sub>.

Hall of Fame?

or

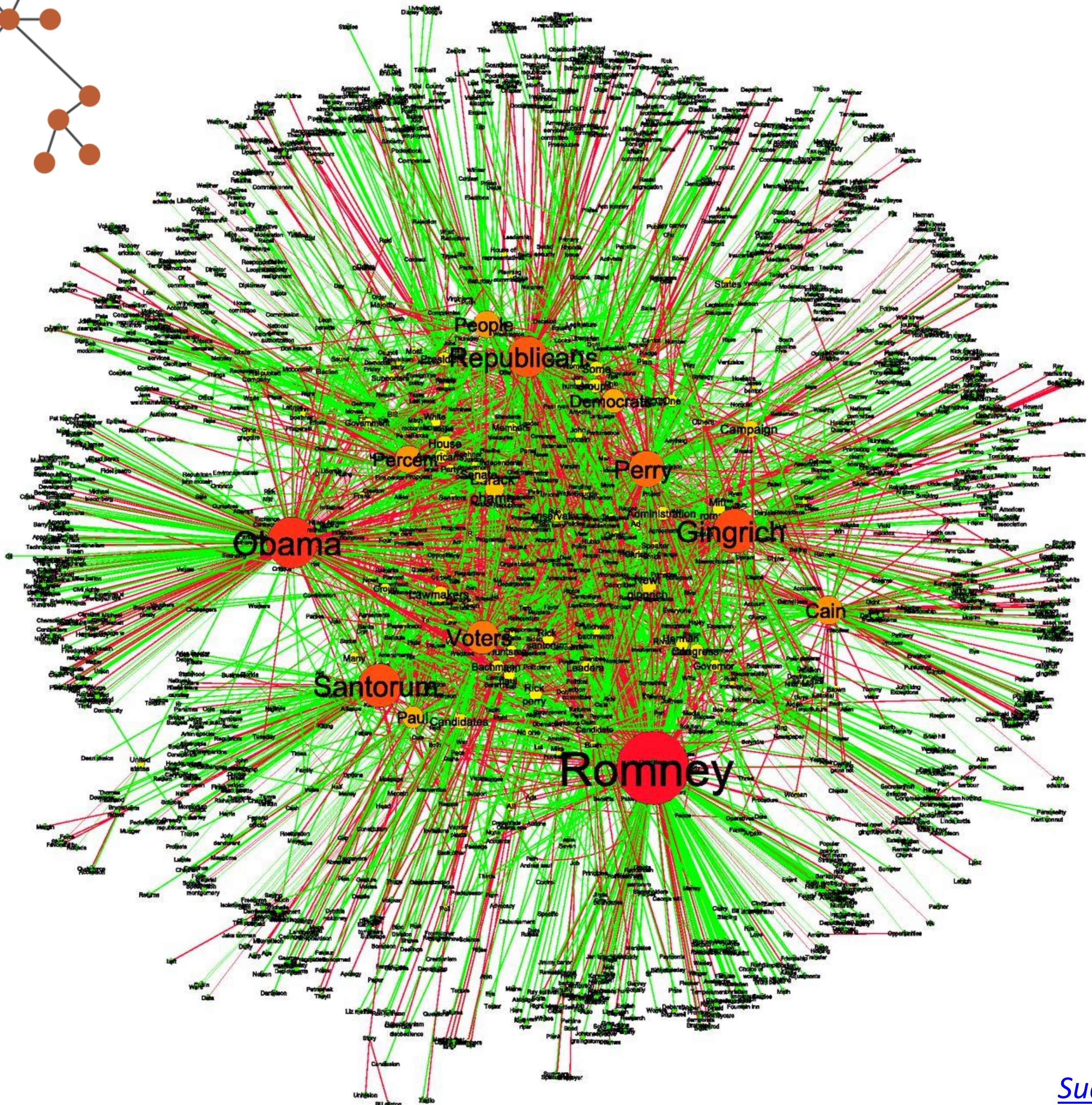
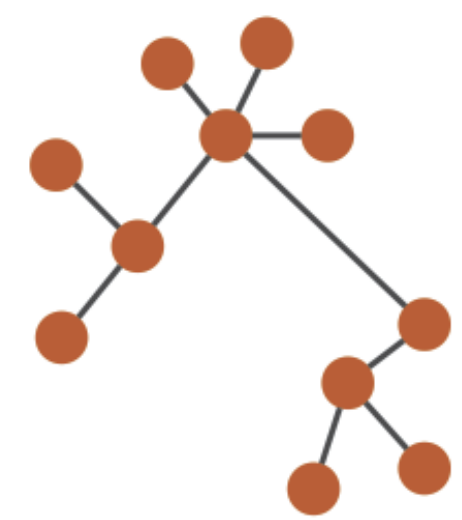
Hall of Shame?



# Node-Link Diagrams

Connection Marks

- ✓ NETWORKS
- ✓ TREES



US presidential election network for 2012 primaries. 81k articles from 490 news outlets

- Nodes: key entities from noun phrases. Sized by degree.
- Edges: relationships from verbs. Colored by positive (green) and negative (red) weights.

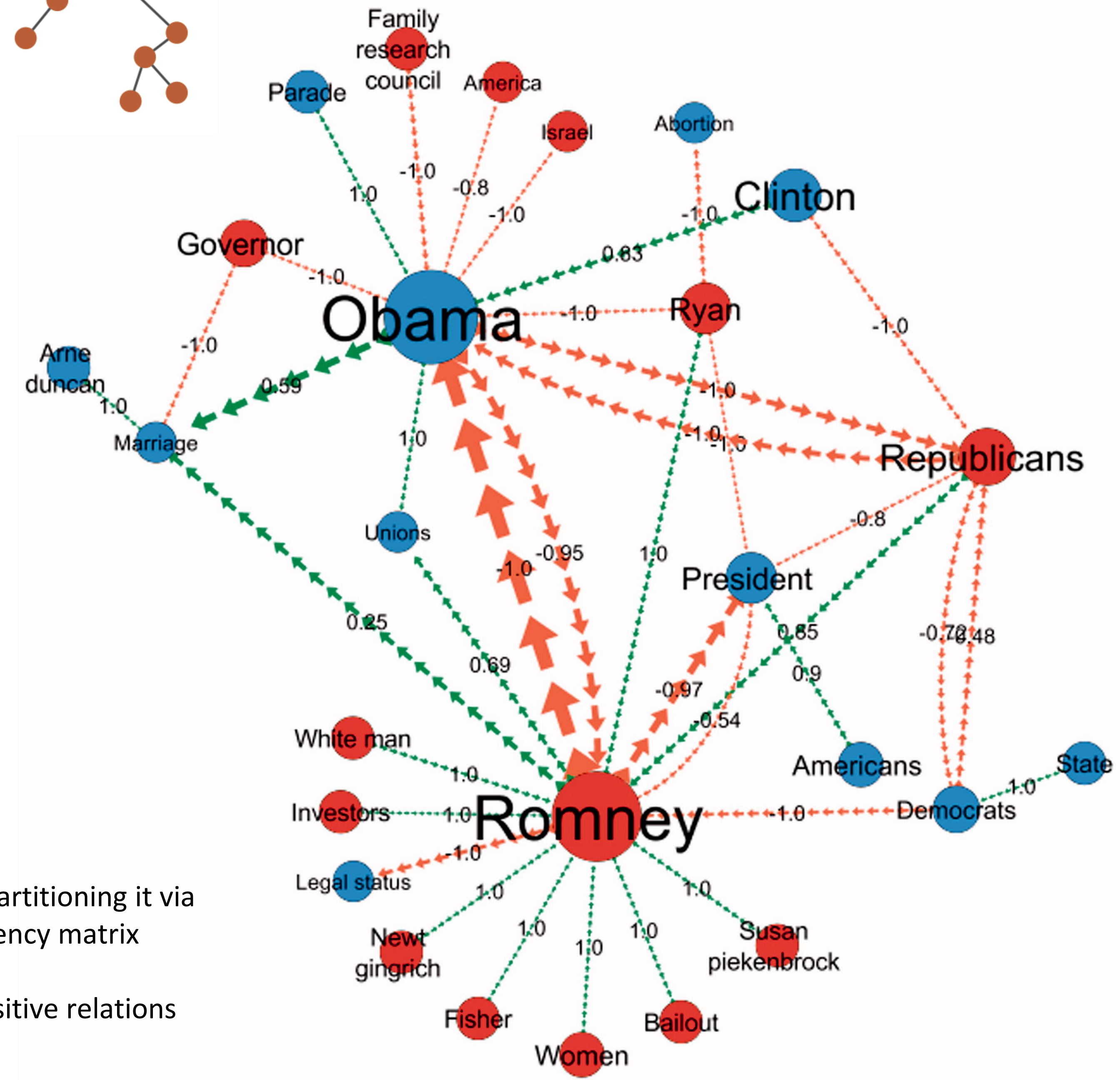
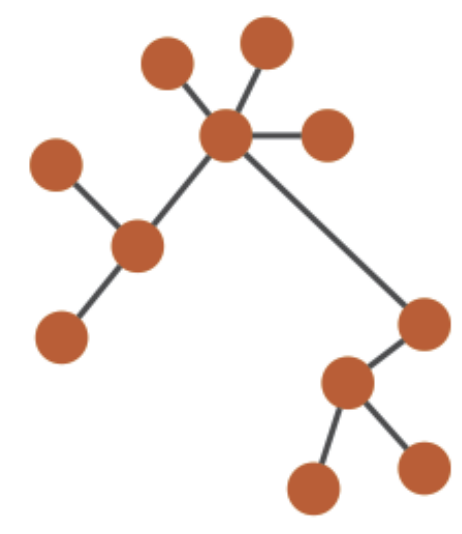


# Node-Link Diagrams

Connection Marks

✓ NETWORKS

✓ TREES

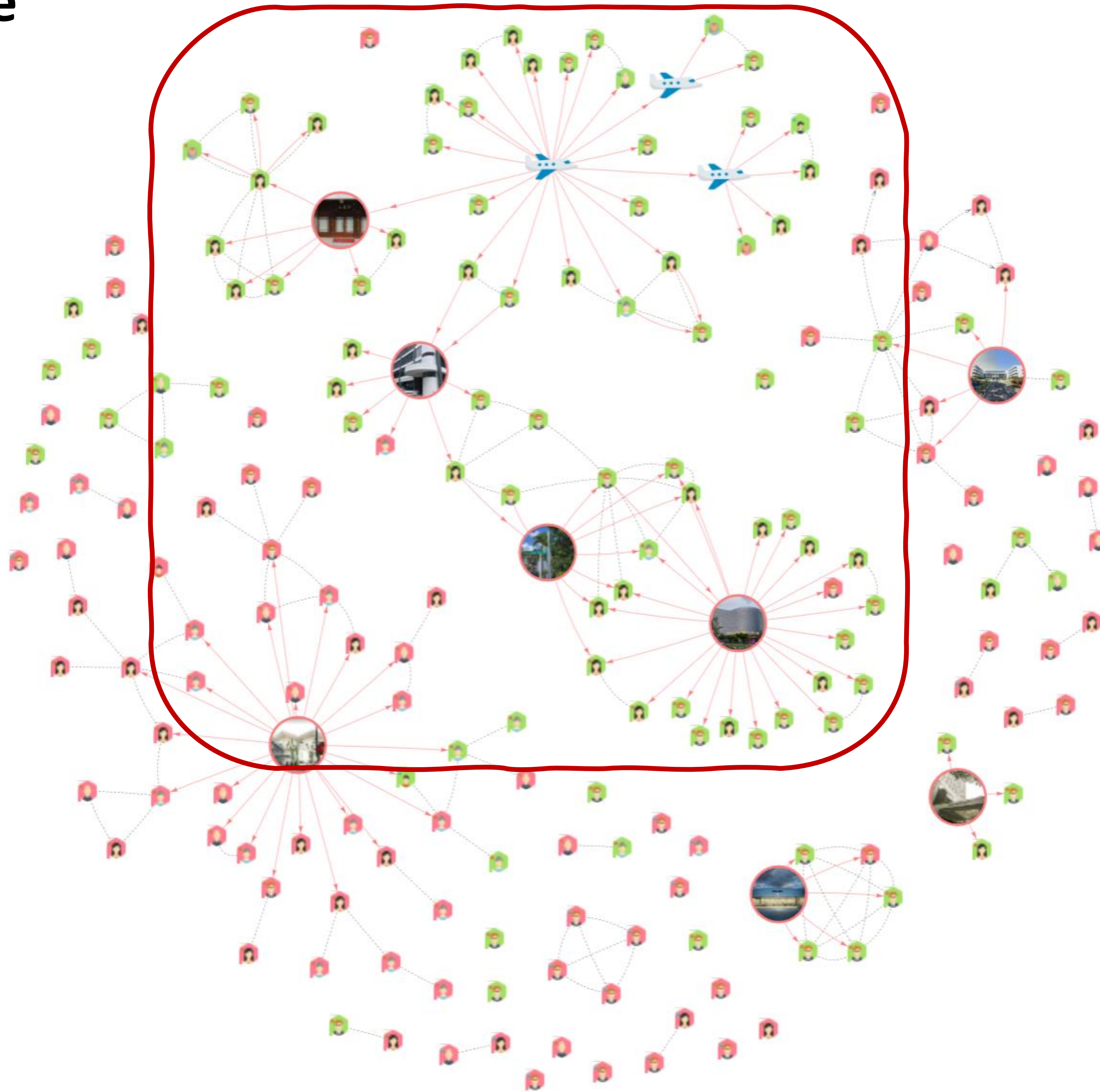


subset of the election network, coloured by partitioning it via the first eigenvalue of the symmetrised adjacency matrix

orange and green links show negative and positive relations between entities.

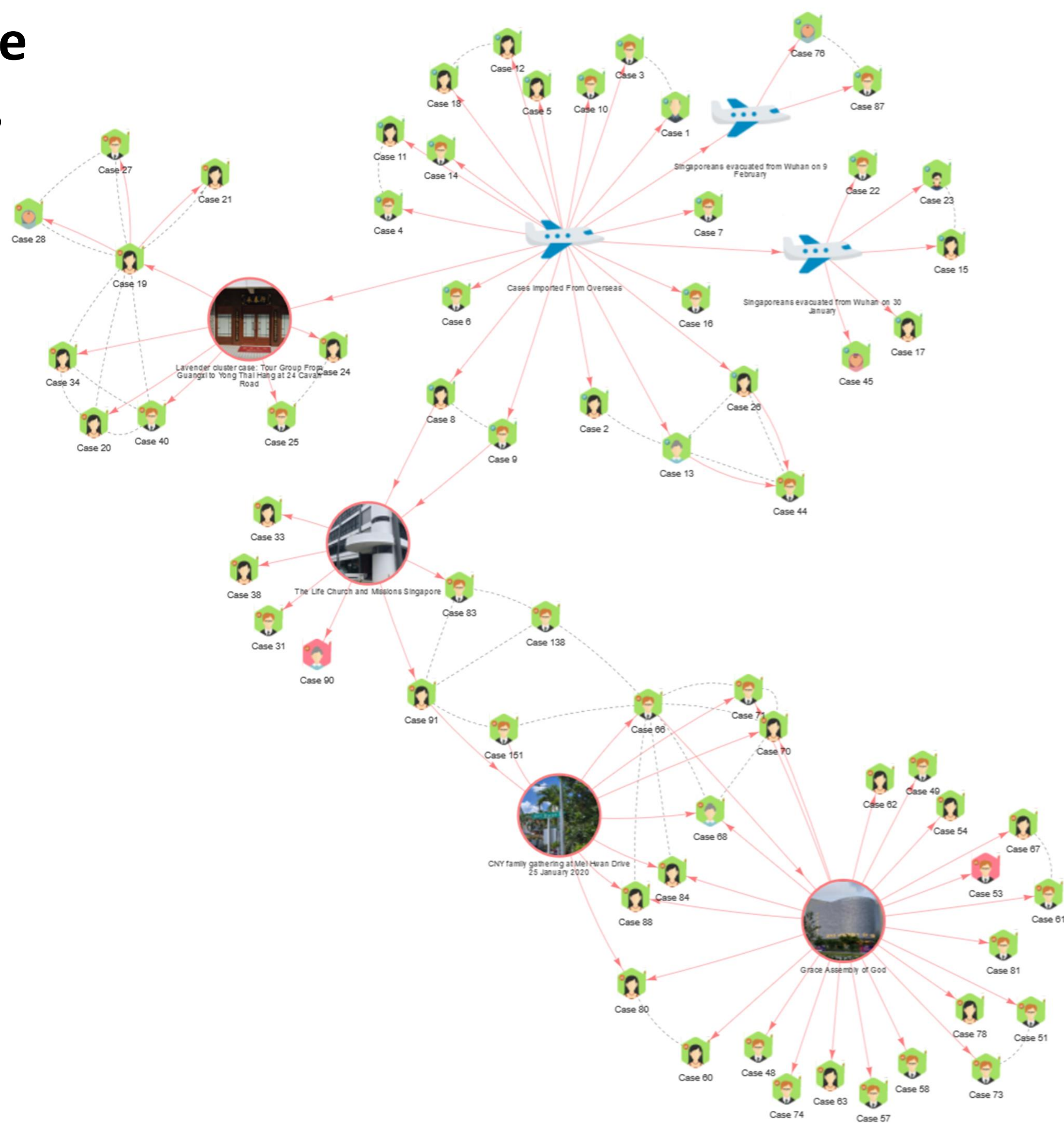
# Dashboard of the COVID-19 Virus Outbreak in Singapore

2020-01-21 – 03-12



# Dashboard of the COVID-19 Virus Outbreak in Singapore

2020-01-21 – 03-12





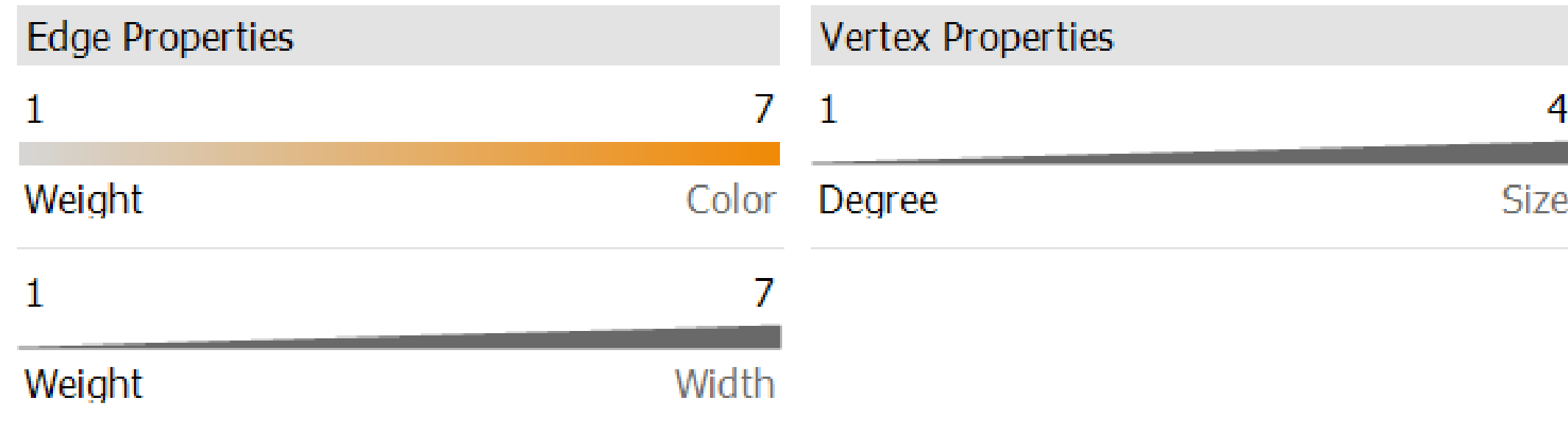
In-class exercise

# Drawing a Node-Link Visualization

| Nodes: | <u>ID</u> | <u>Type</u> |
|--------|-----------|-------------|
|        | 1         | A           |
|        | 2         | B           |
|        | 3         | A           |
|        | 4         | A           |
|        | 5         | B           |

| Edges: | <u>Source</u> | <u>Target</u> | <u>Weight</u> |
|--------|---------------|---------------|---------------|
|        | 1             | 2             | 1             |
|        | 1             | 3             | 7             |
|        | 2             | 3             | 4             |
|        | 3             | 4             | 2             |
|        | 4             | 1             | 2             |
|        | 5             | 1             | 1             |

# Drawing a Node-Link Visualization

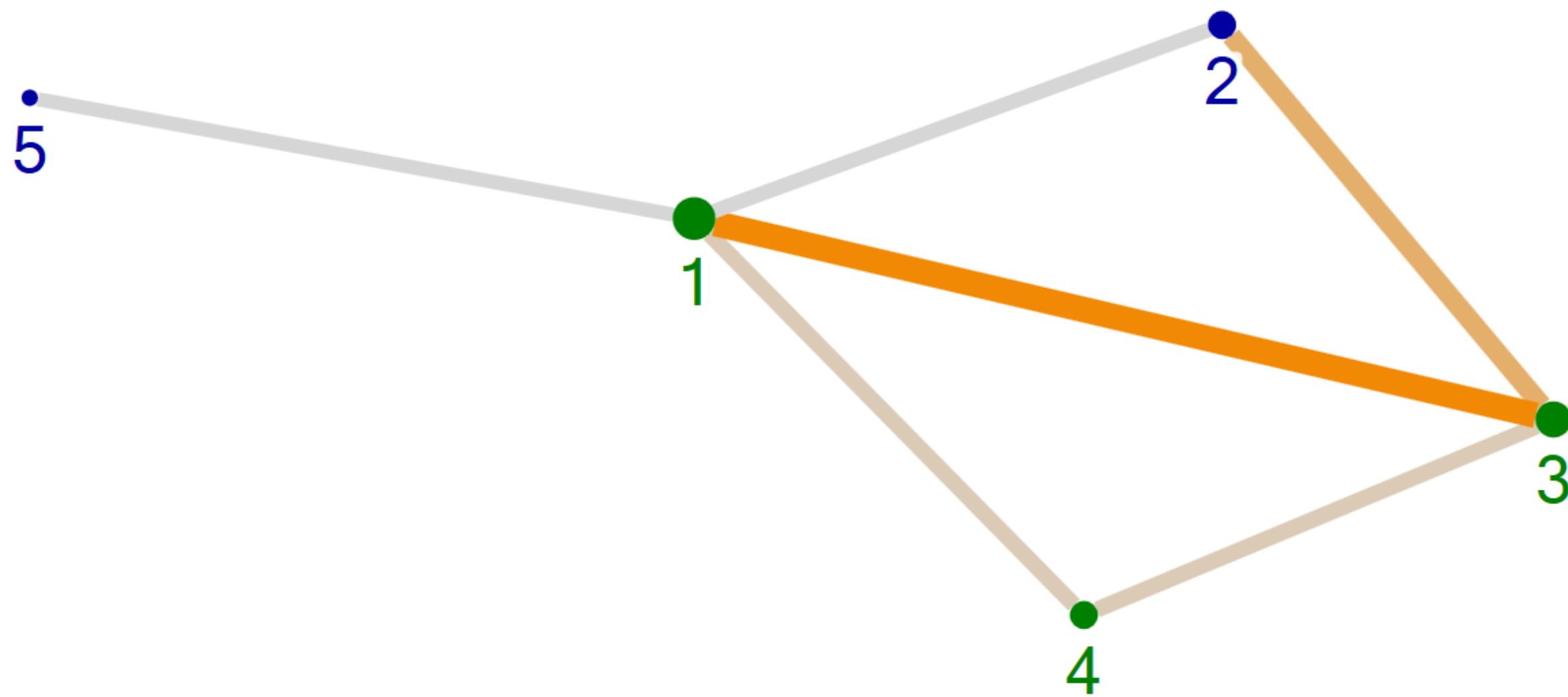


Nodes:

| ID | Type |
|----|------|
| 1  | A    |
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| 3  | A    |
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| 5  | B    |

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| 5      | 1      | 1      |



Let's re-create this in [NodeXL...](#)

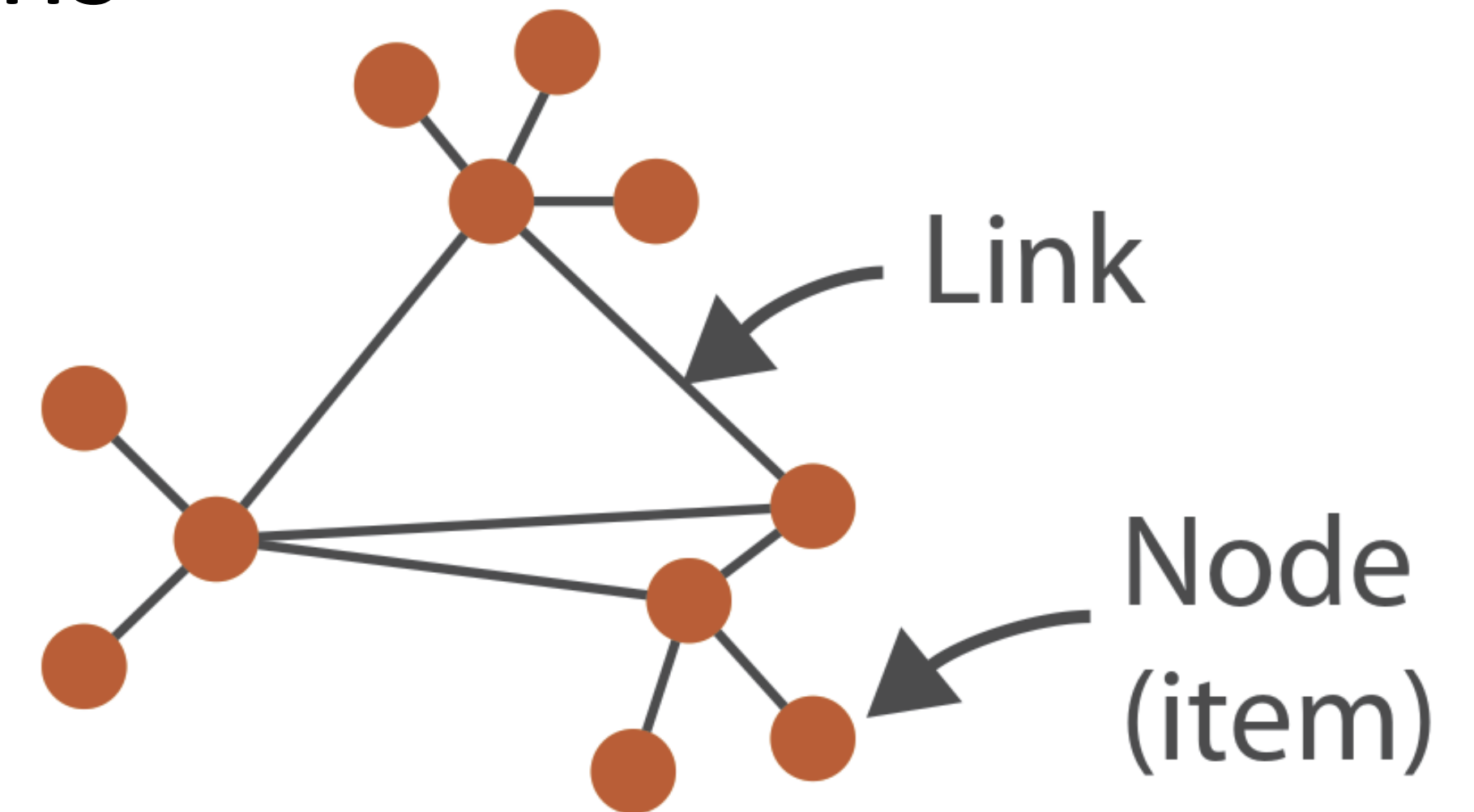
# Node-Link Visualizations

Pros:

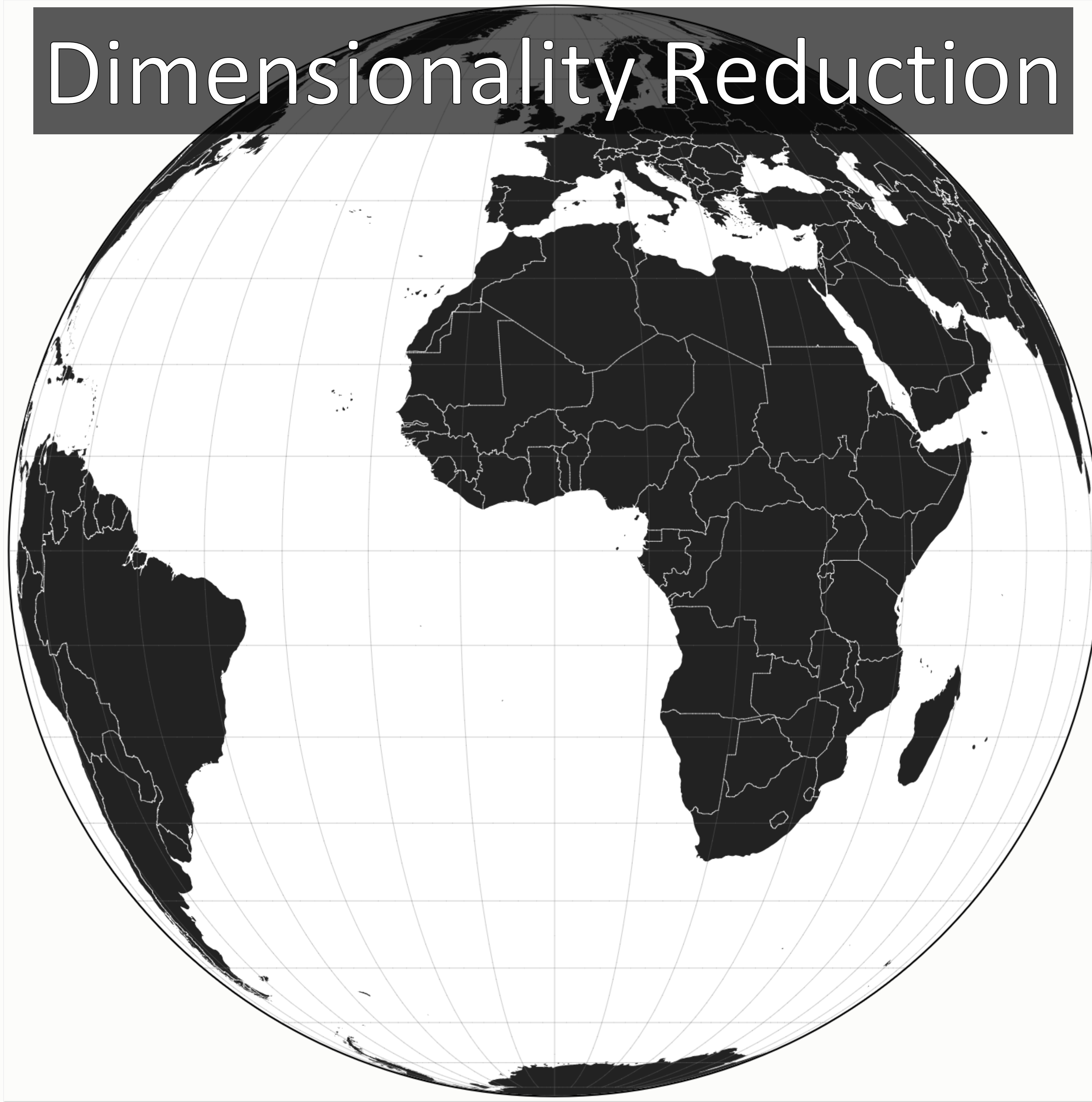
- understandable visual mapping
- can show overall structure, clusters, paths
- flexible, many variations

Cons:

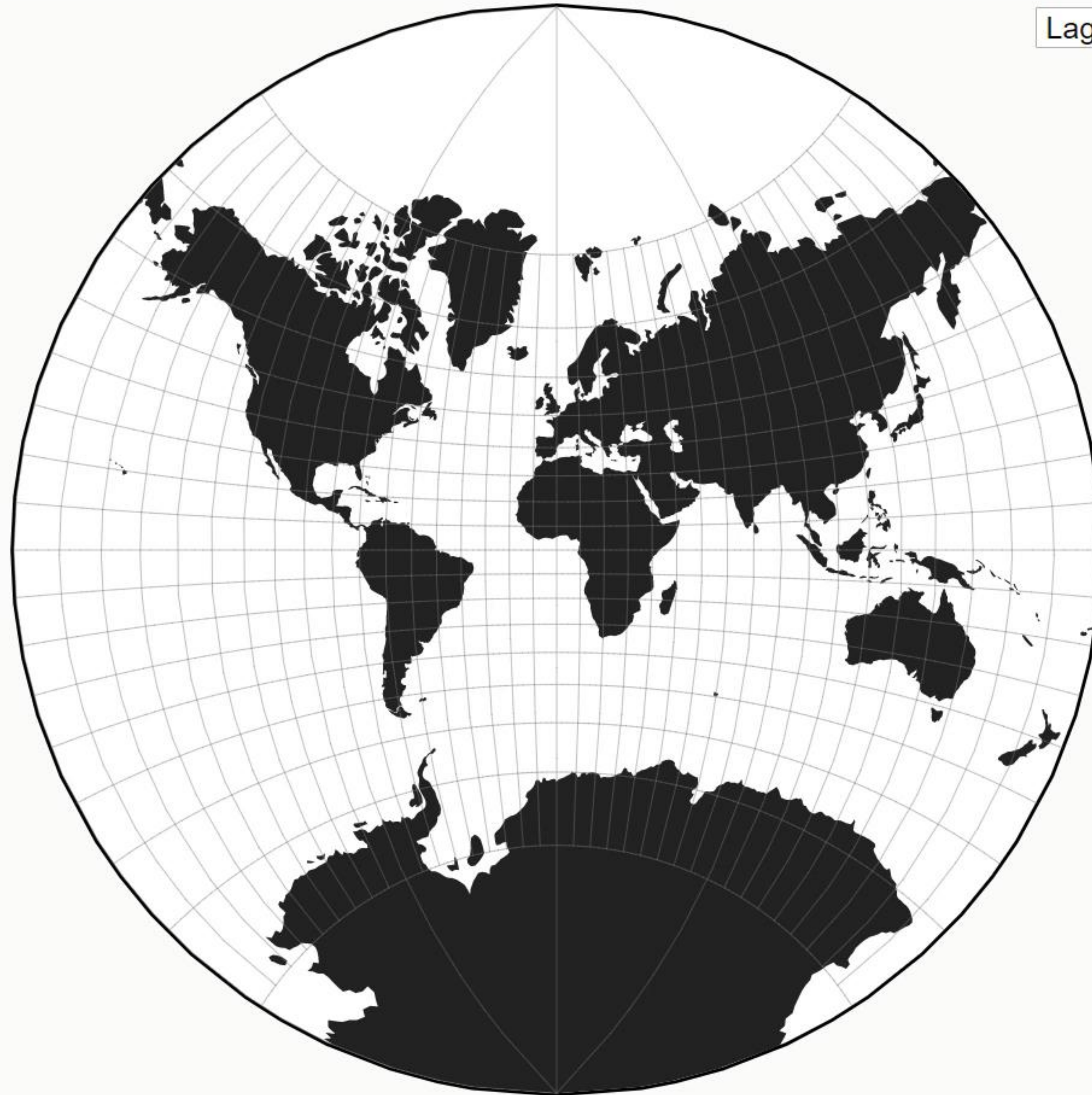
- automatic layout algorithm deficiencies
  - time consuming to run
  - non-deterministic results
  - heuristics with sometimes poor results
- not good for dense graphs - hairball problem!



# Dimensionality Reduction



# Projection Transitions

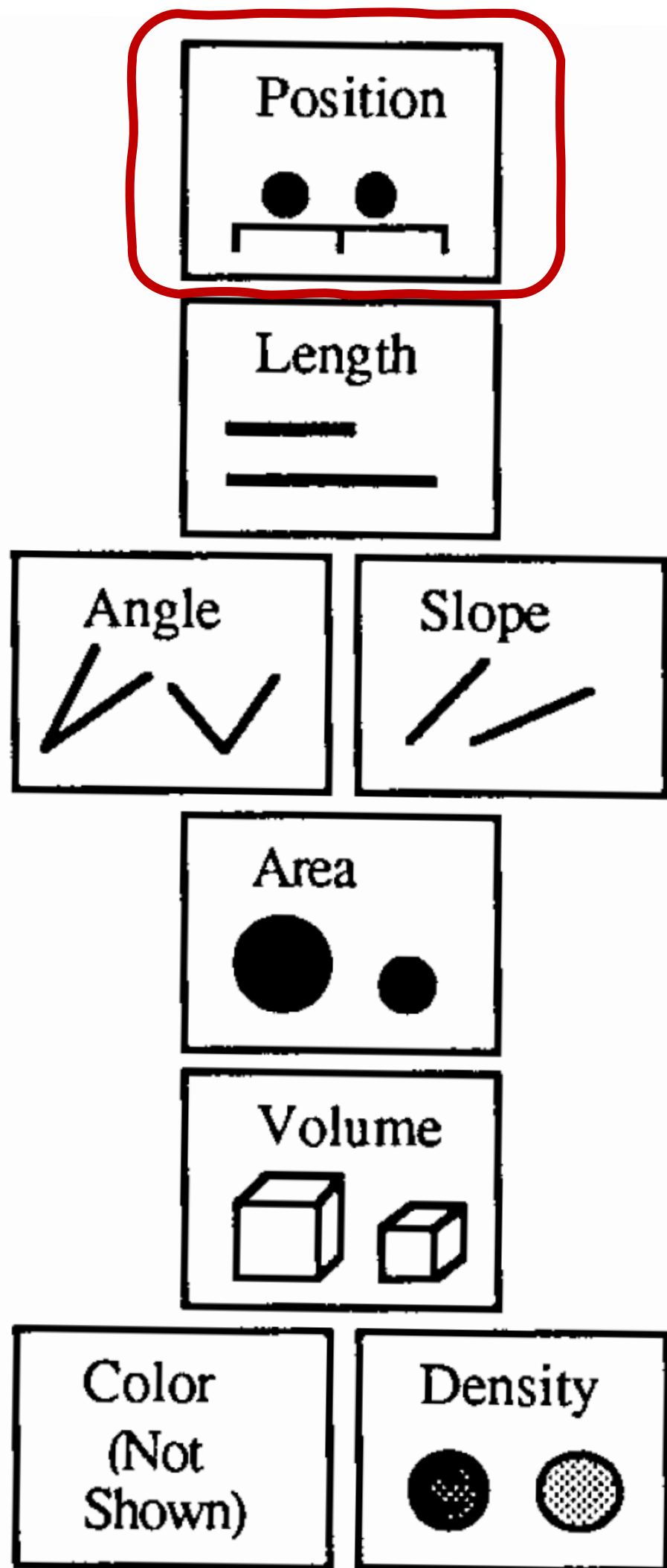


Lagrange ▼

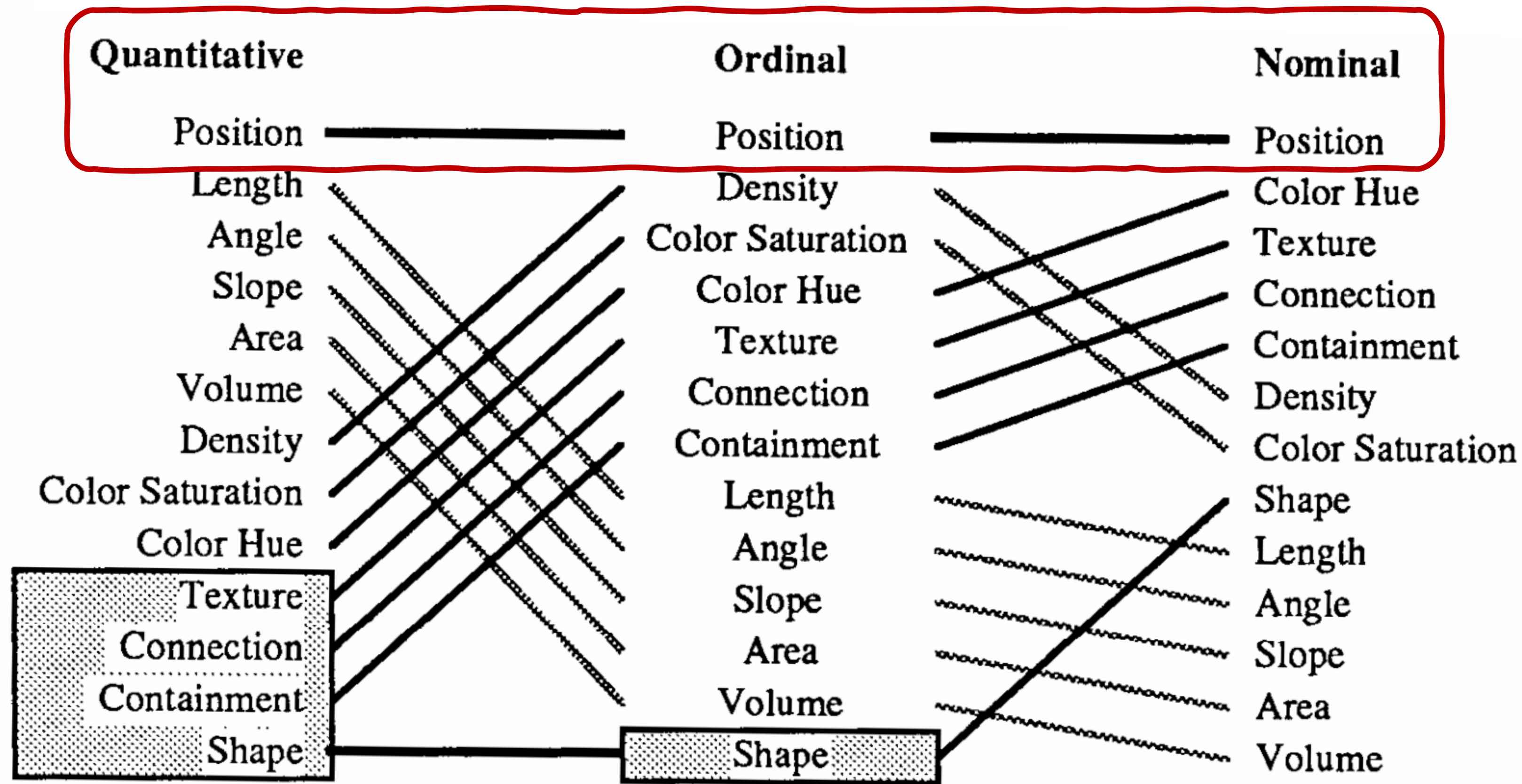
# Spatial Layout

## Quantitative Tasks

More accurate



Less accurate

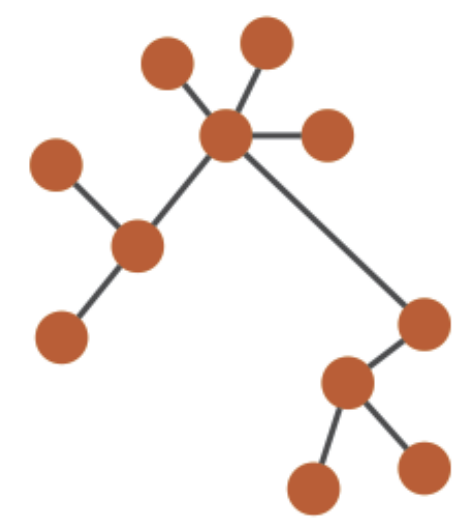




# Node-Link Diagrams

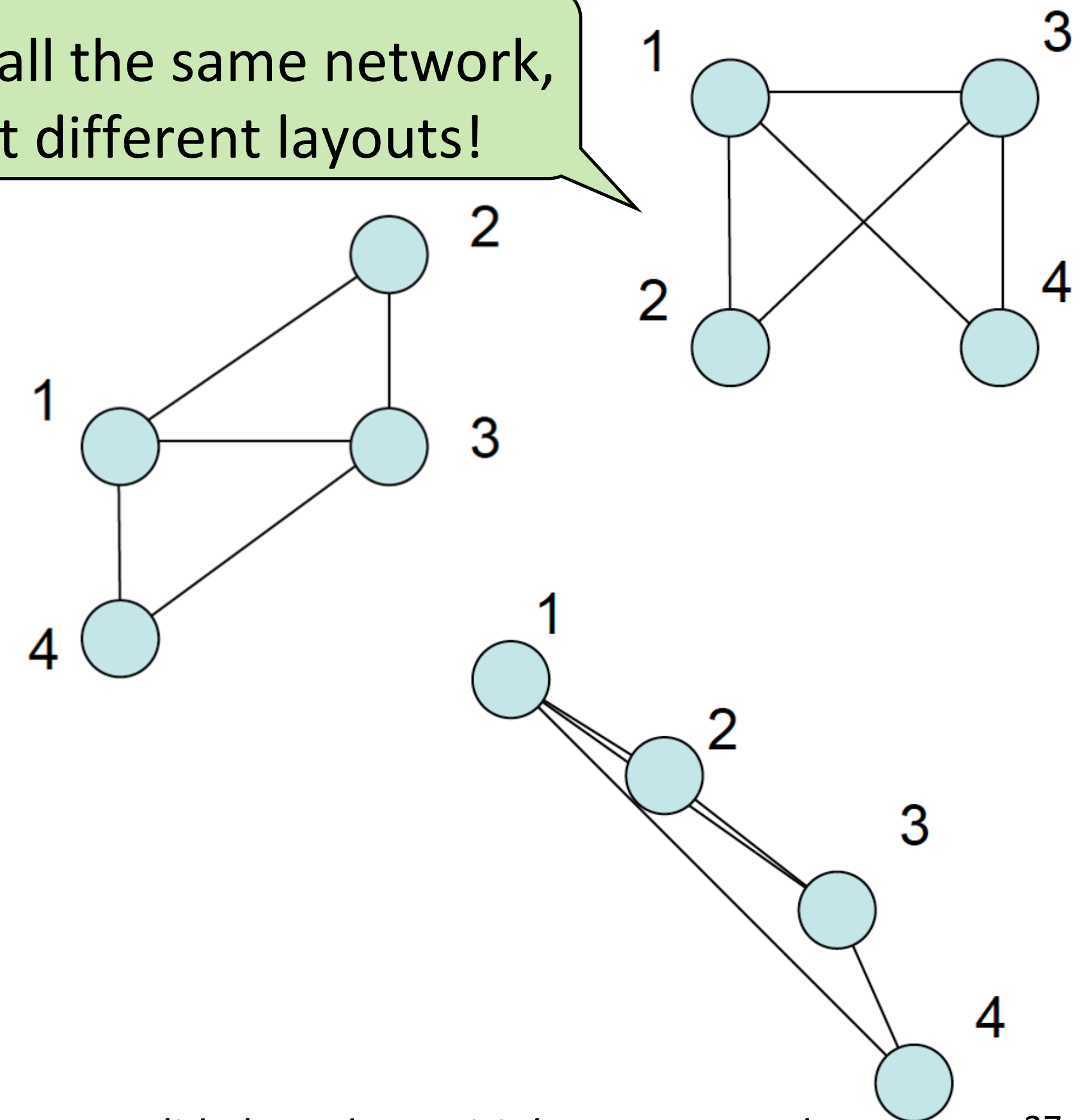
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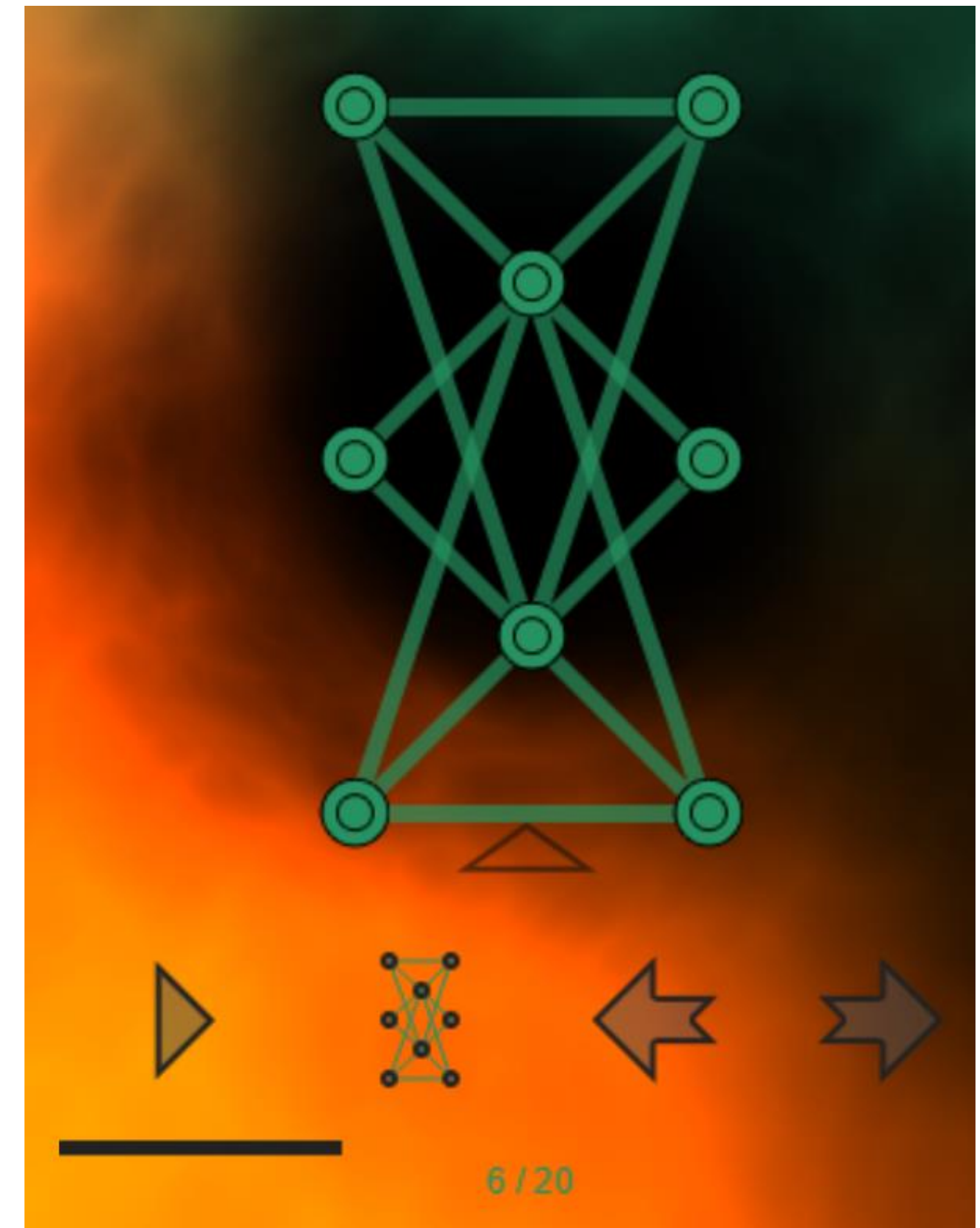




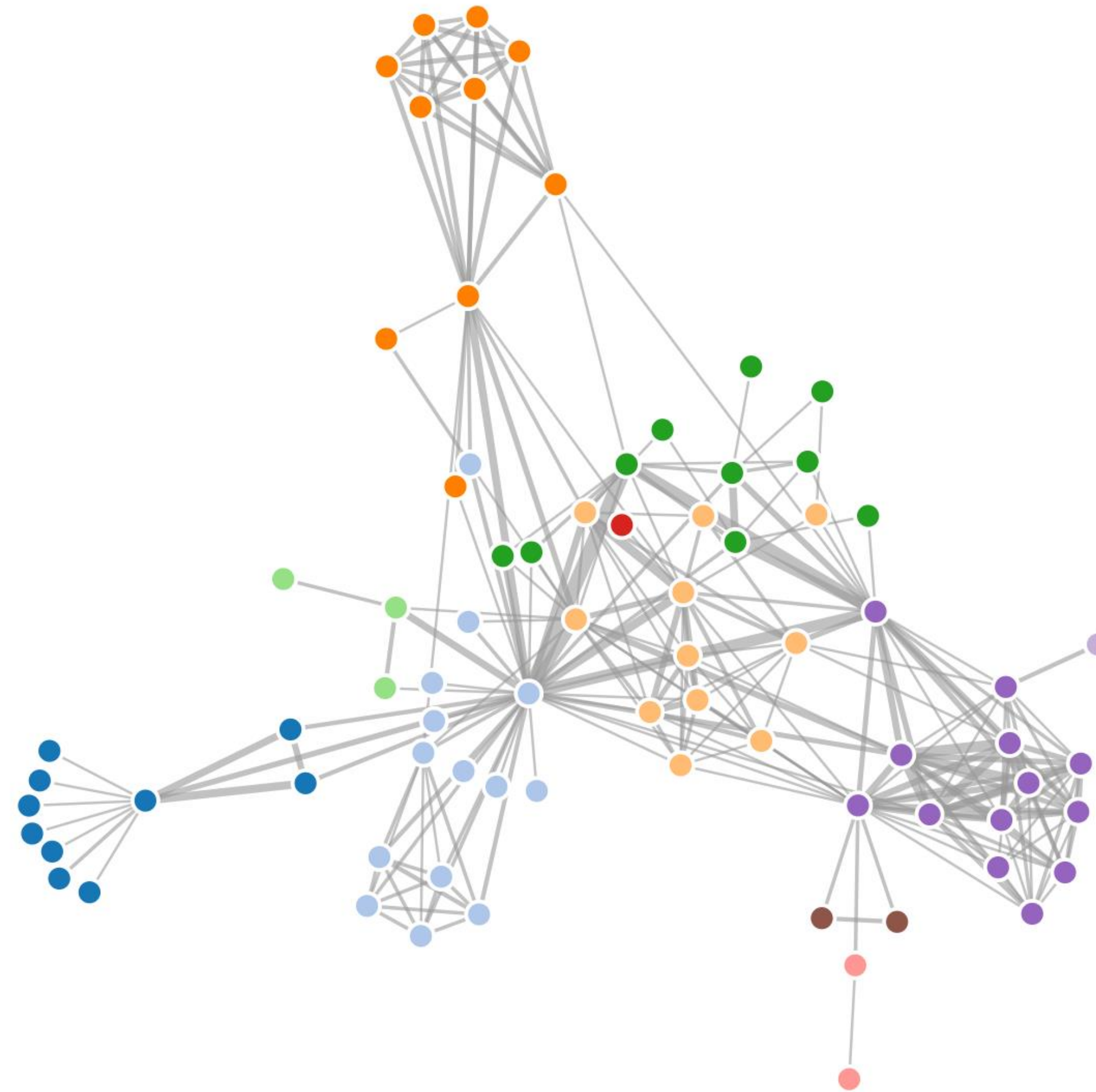
In-class activity:  
planarity

<https://treksit.com>

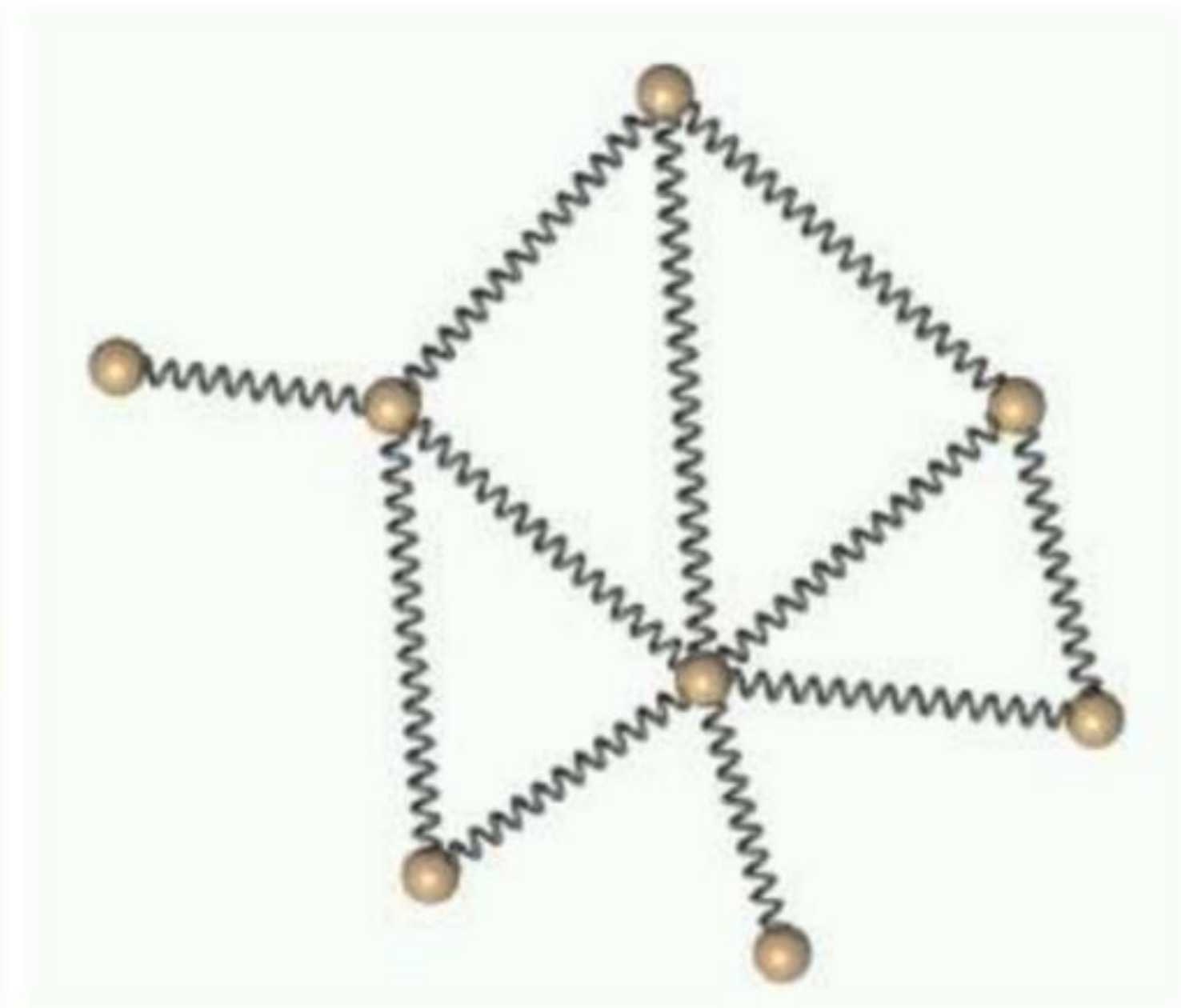
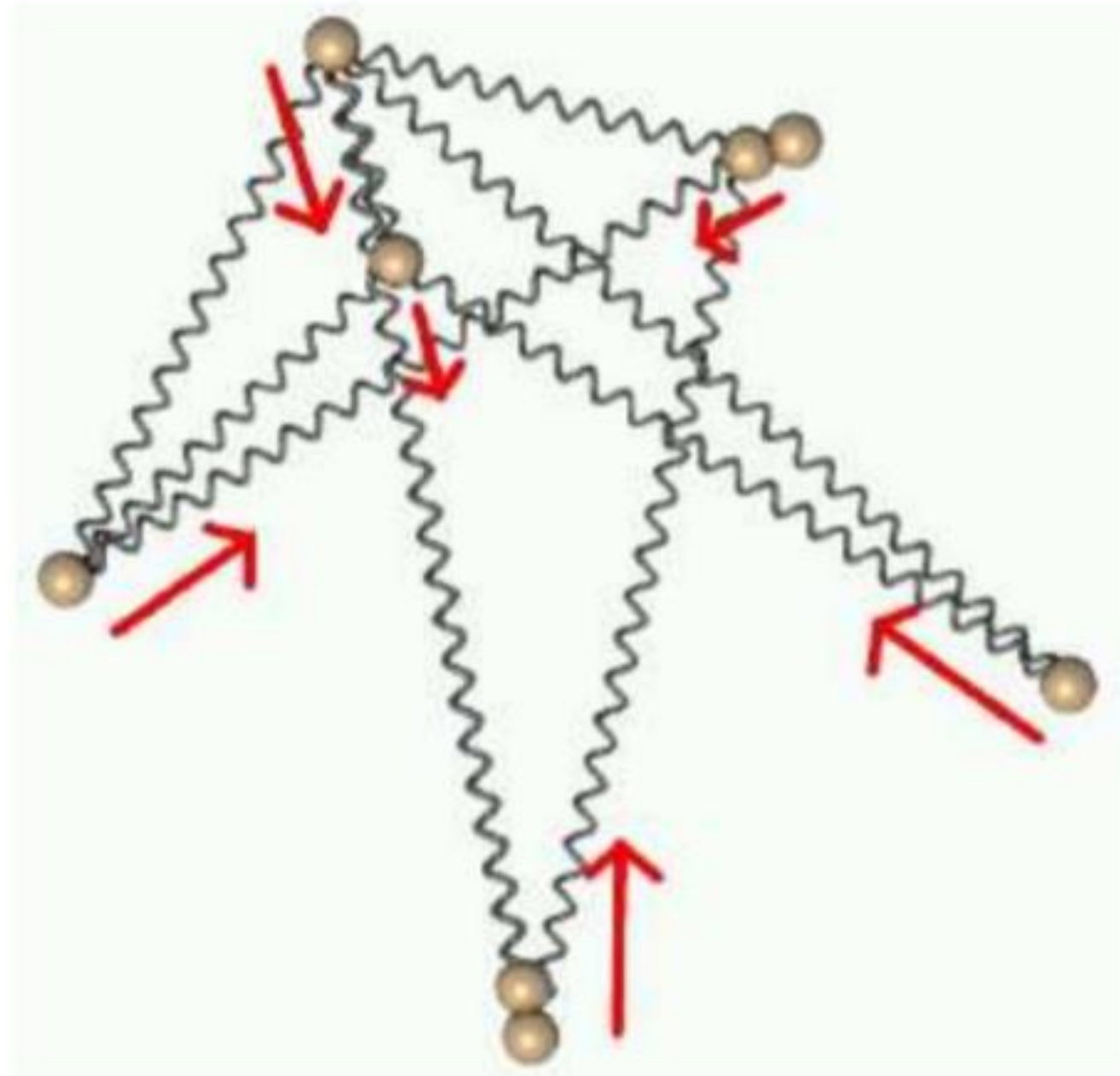
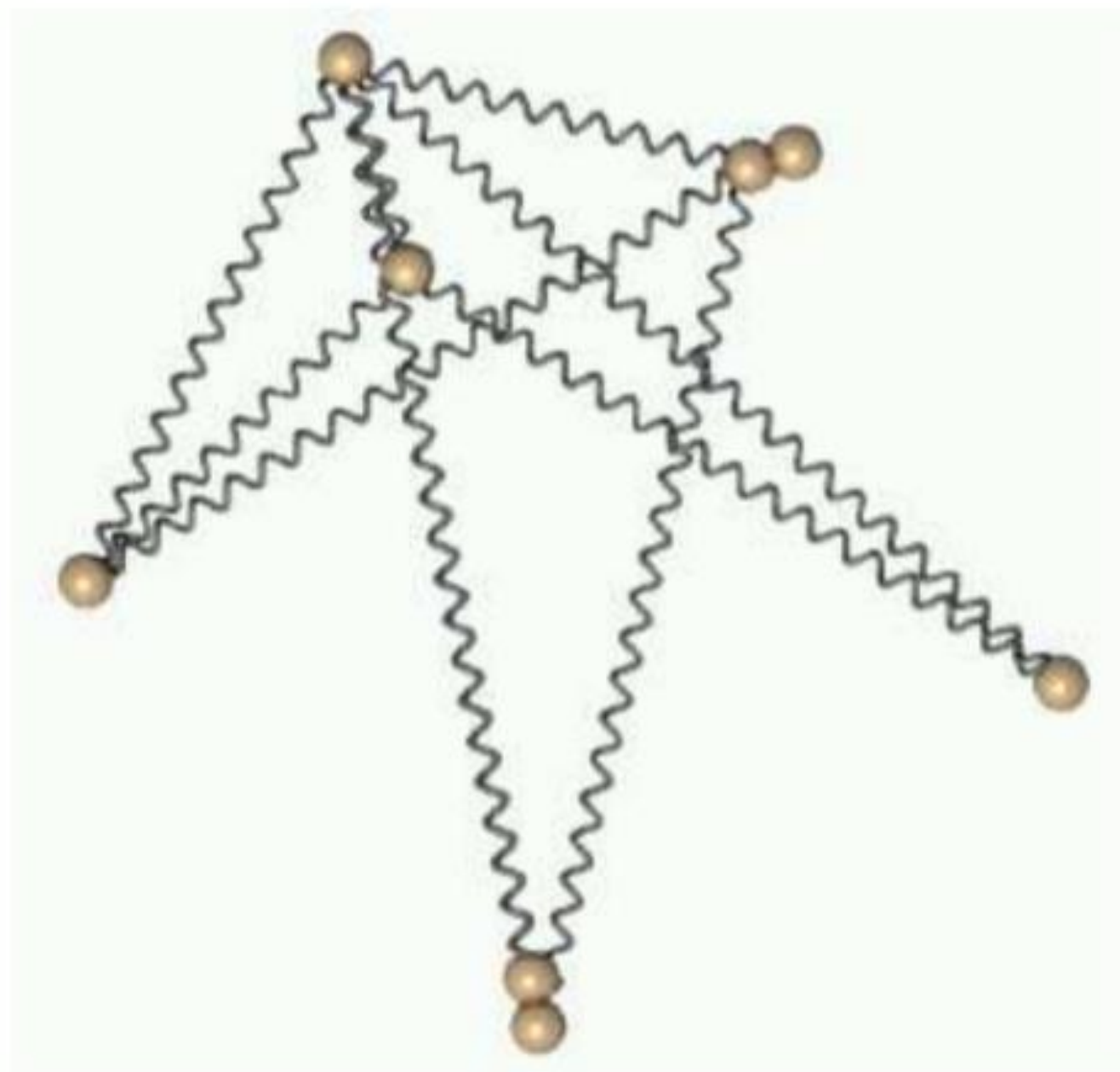
~15 min

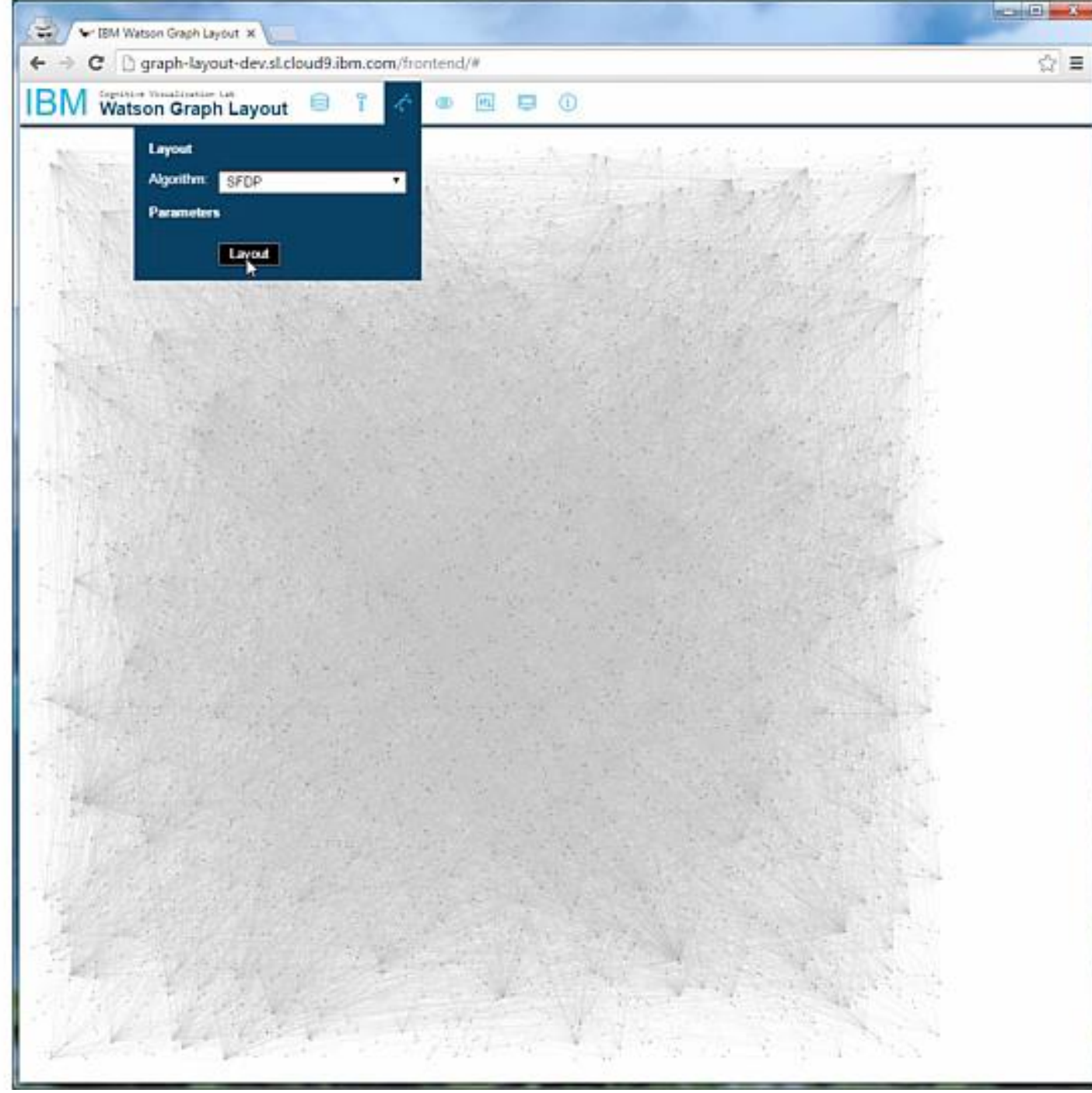
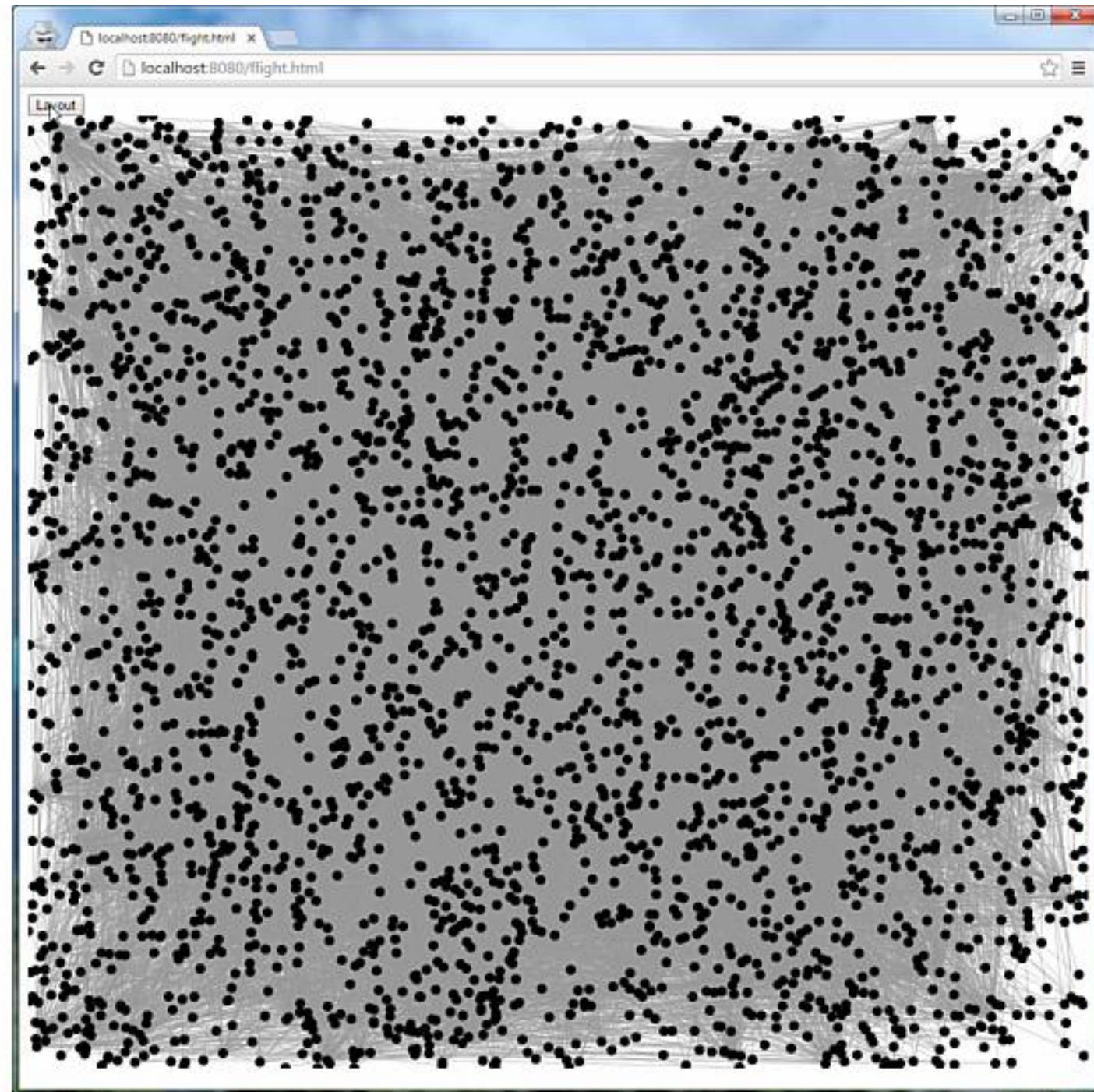
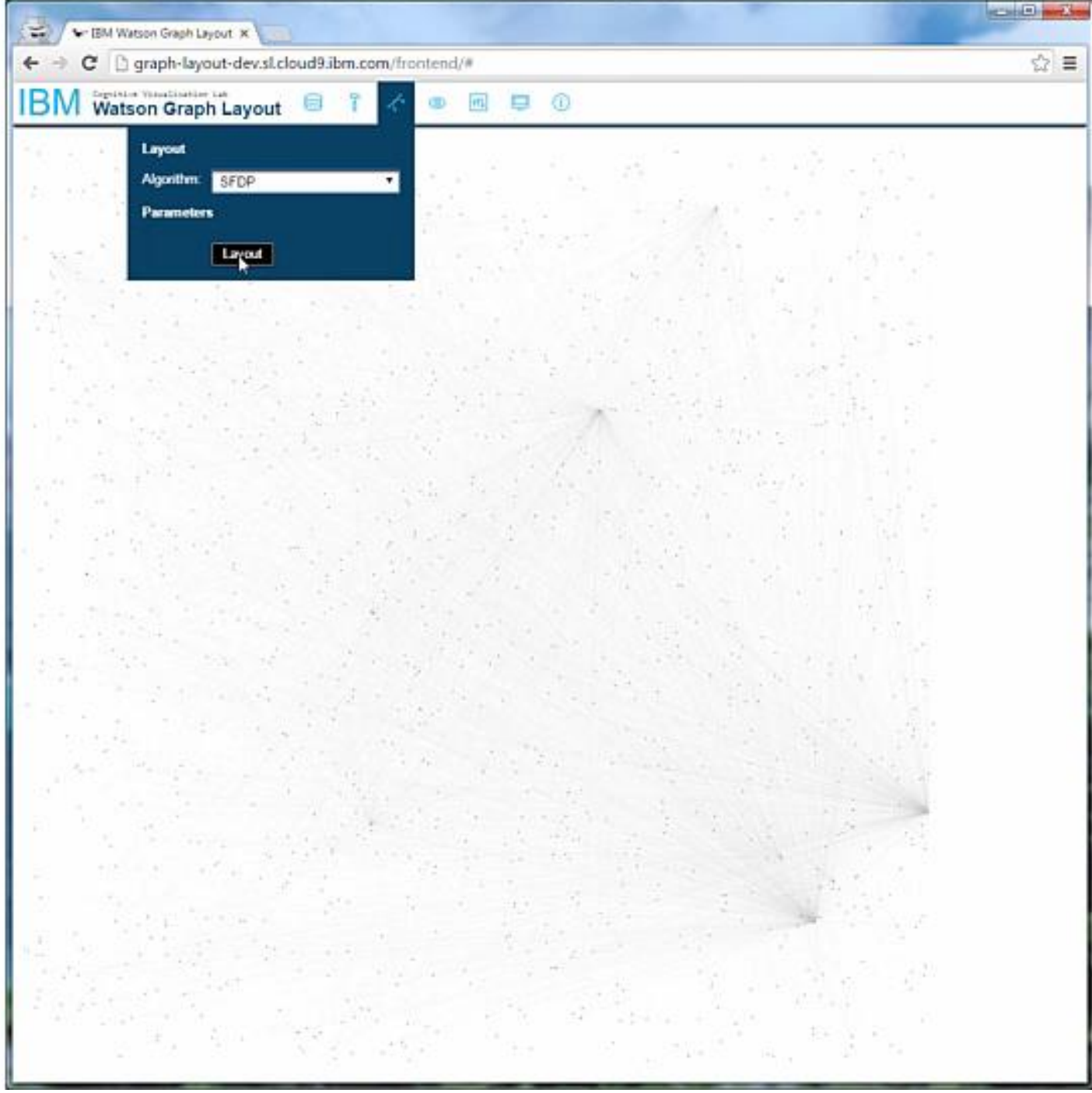
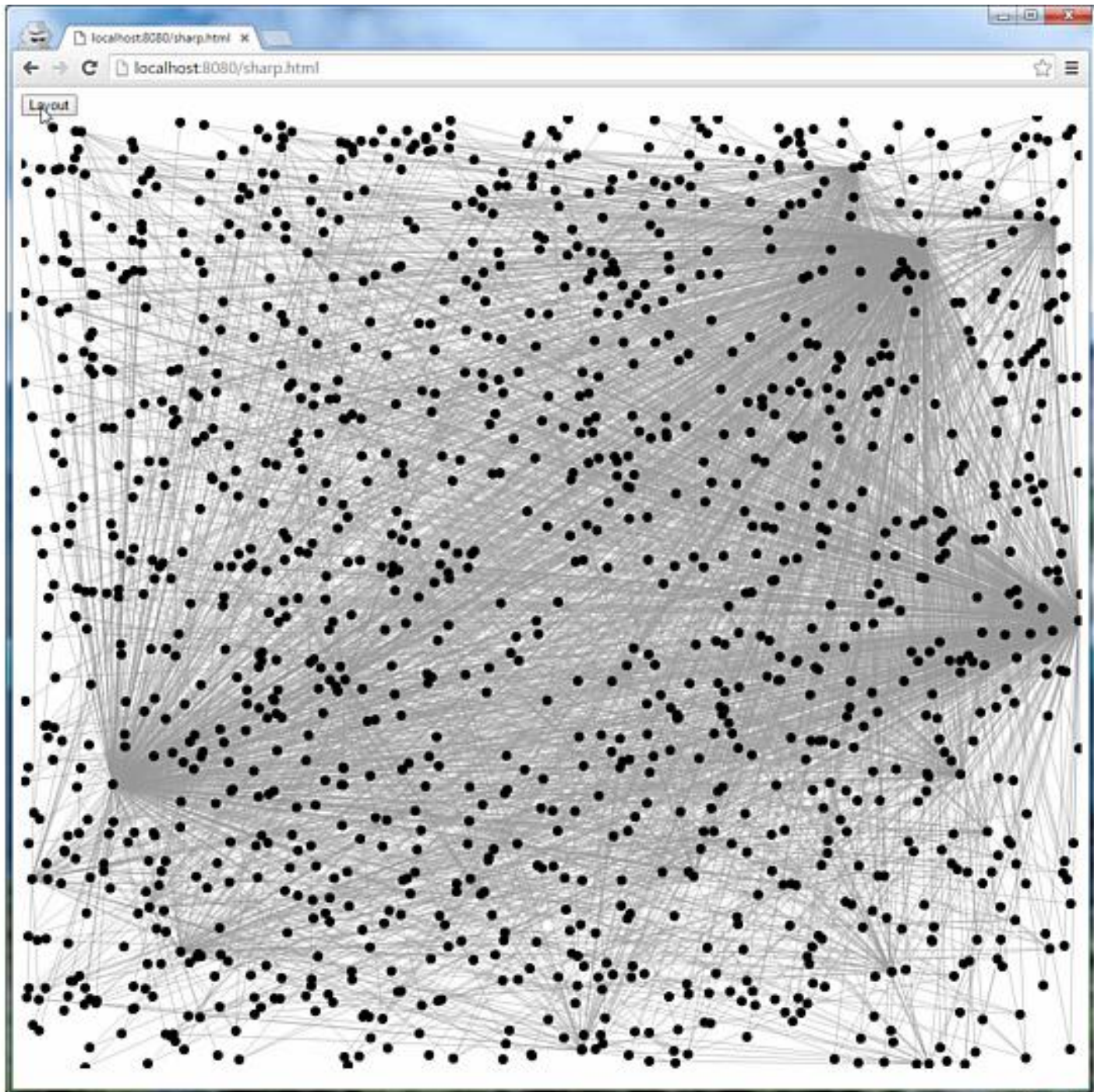
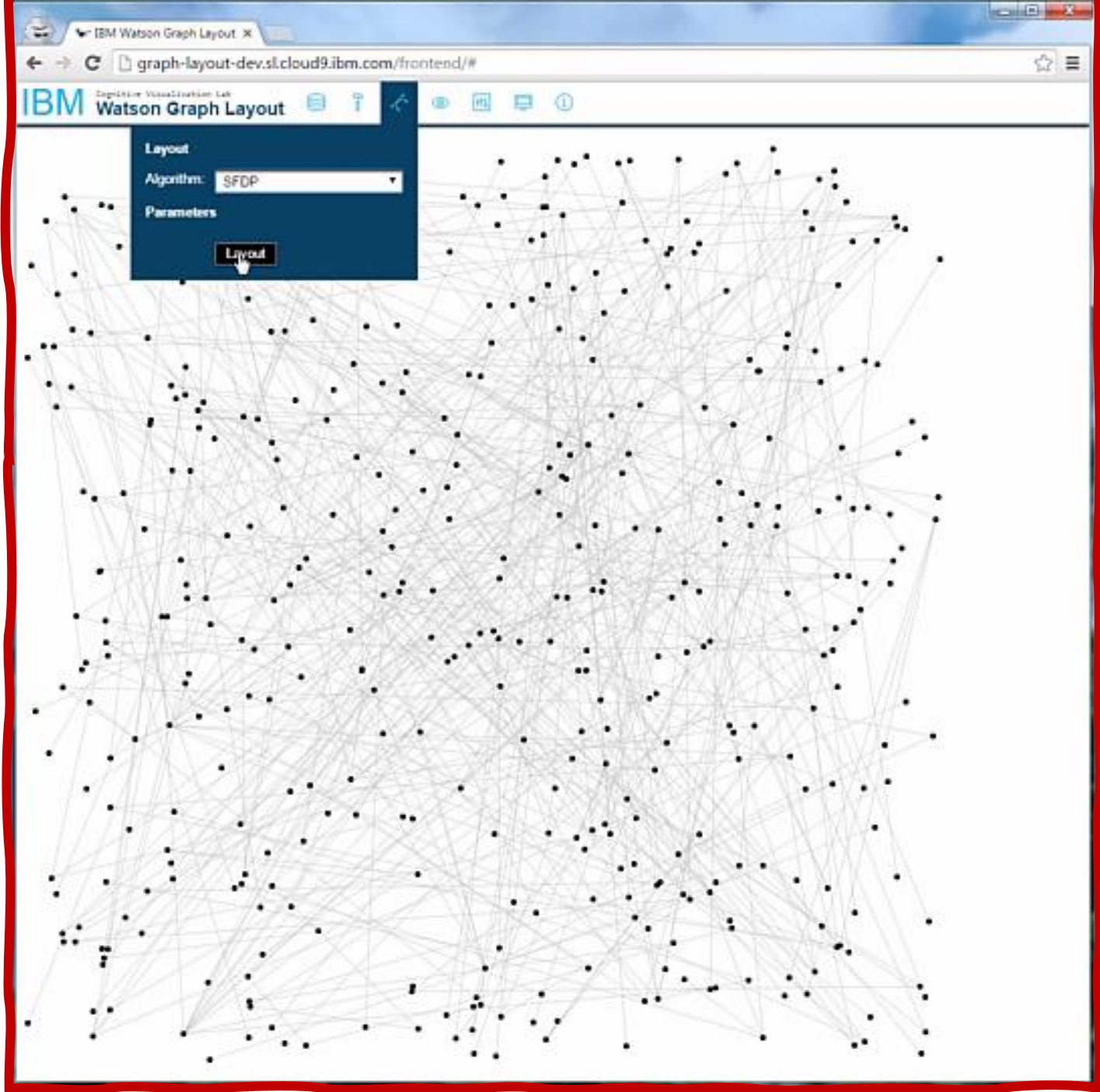
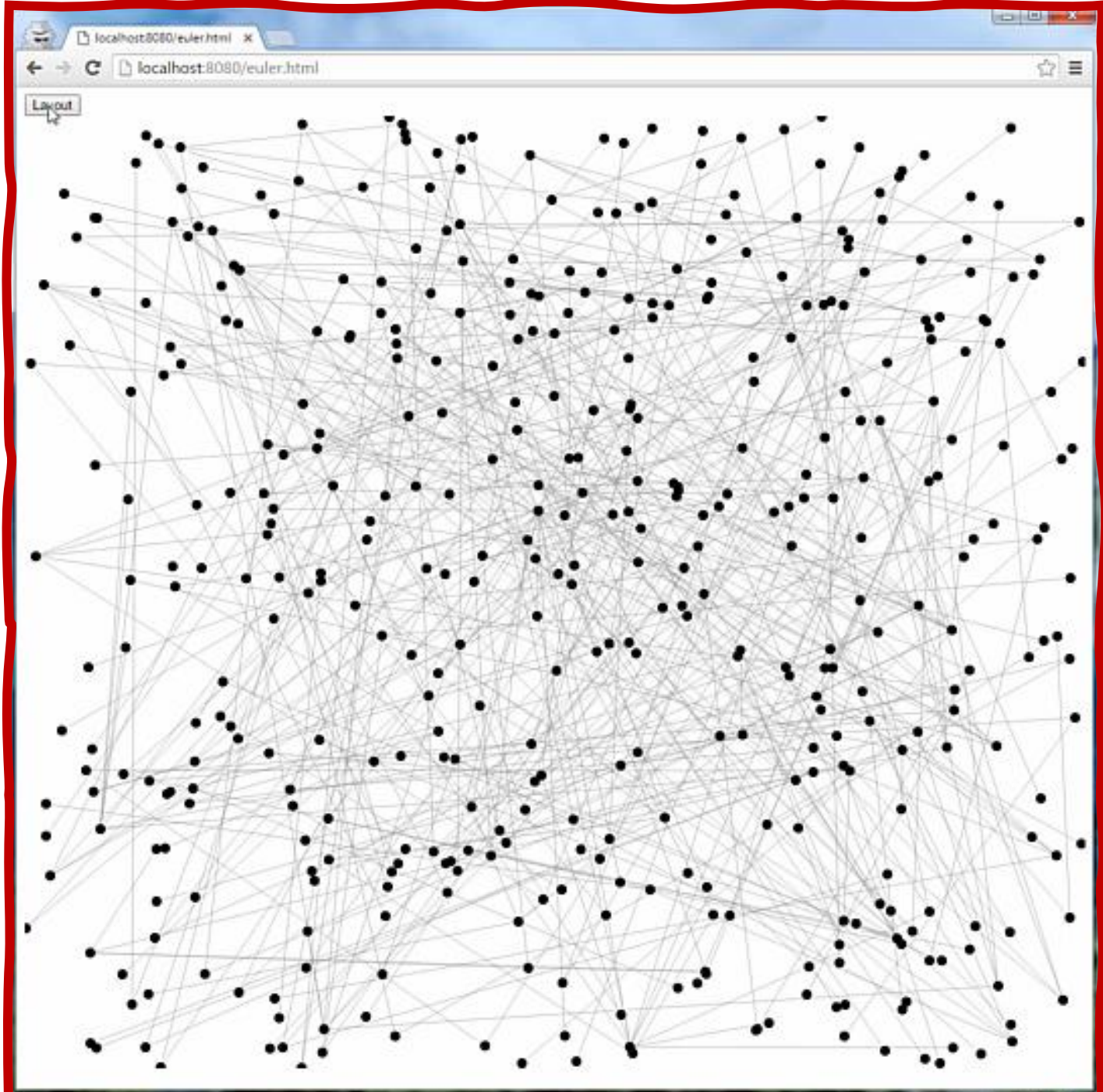


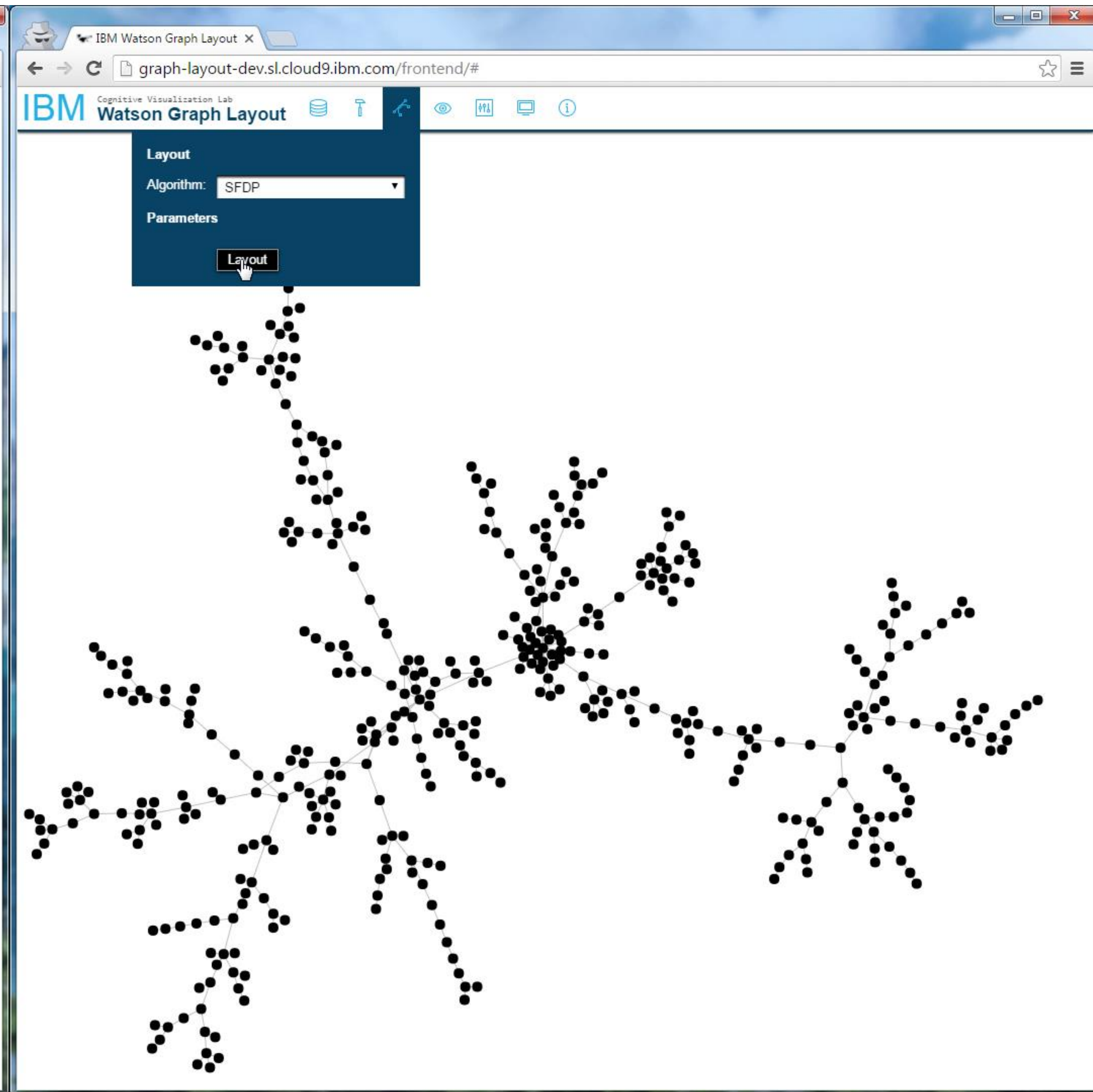
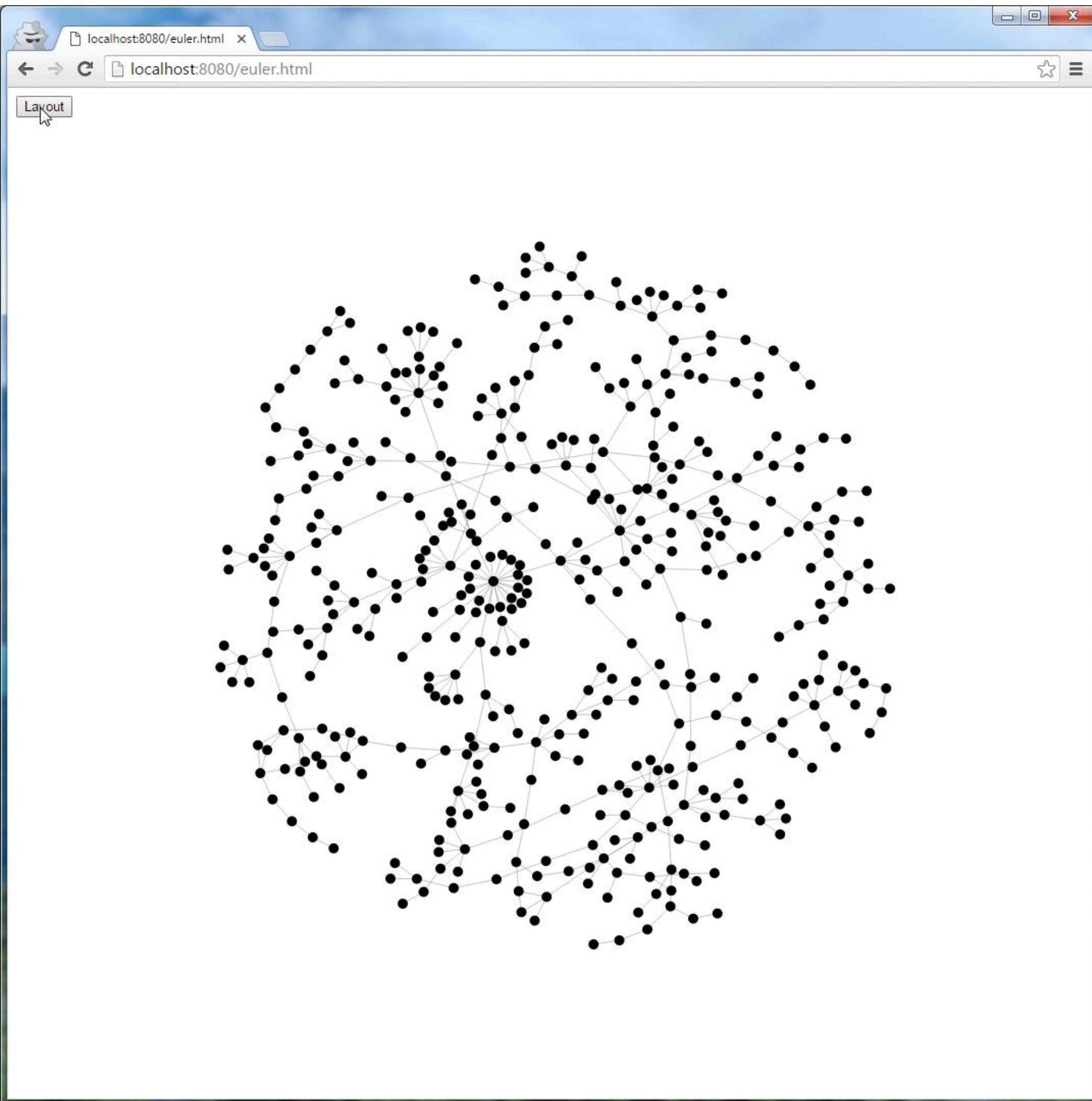
# D3 Force-Directed Layout



# Force-Directed Algorithms

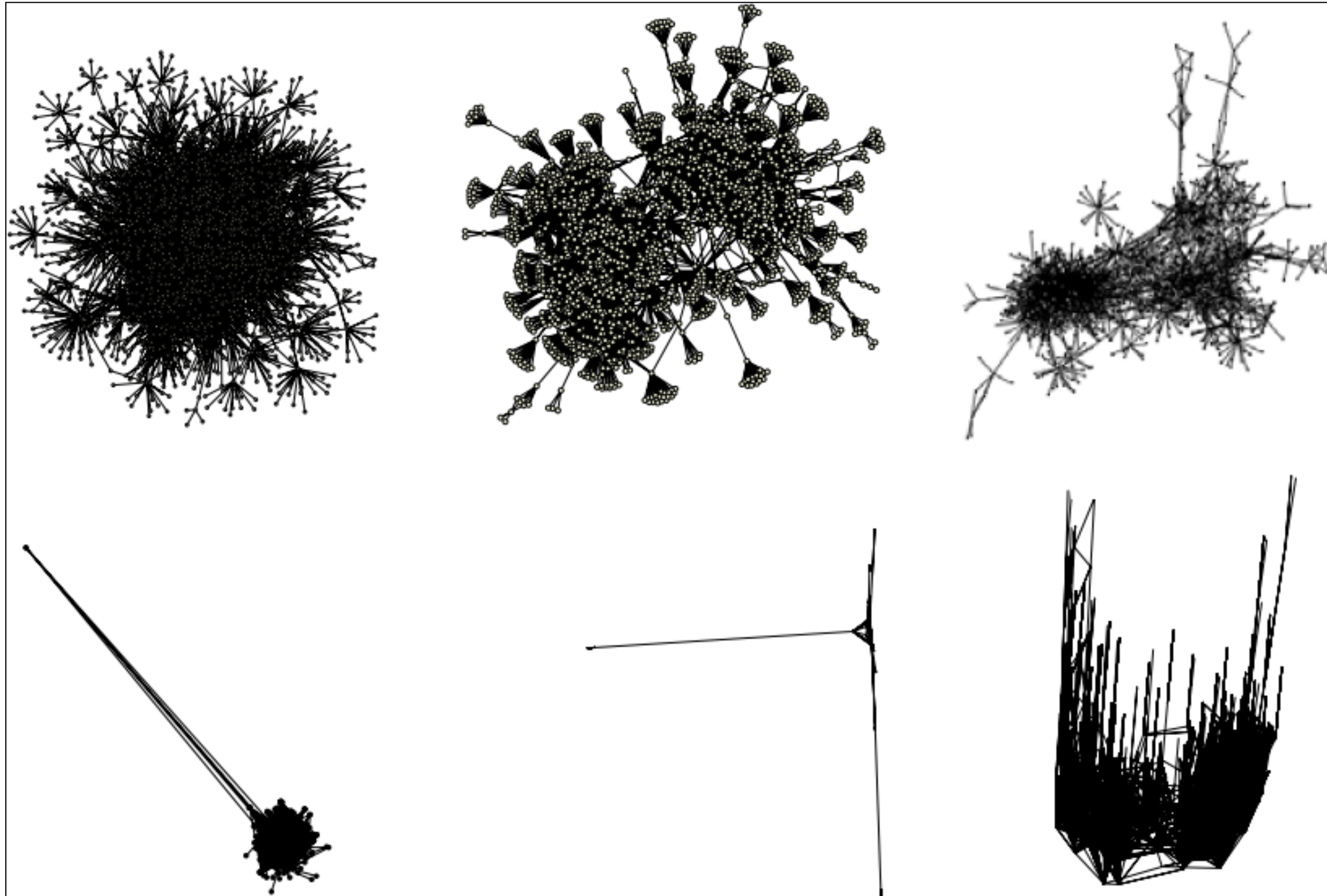




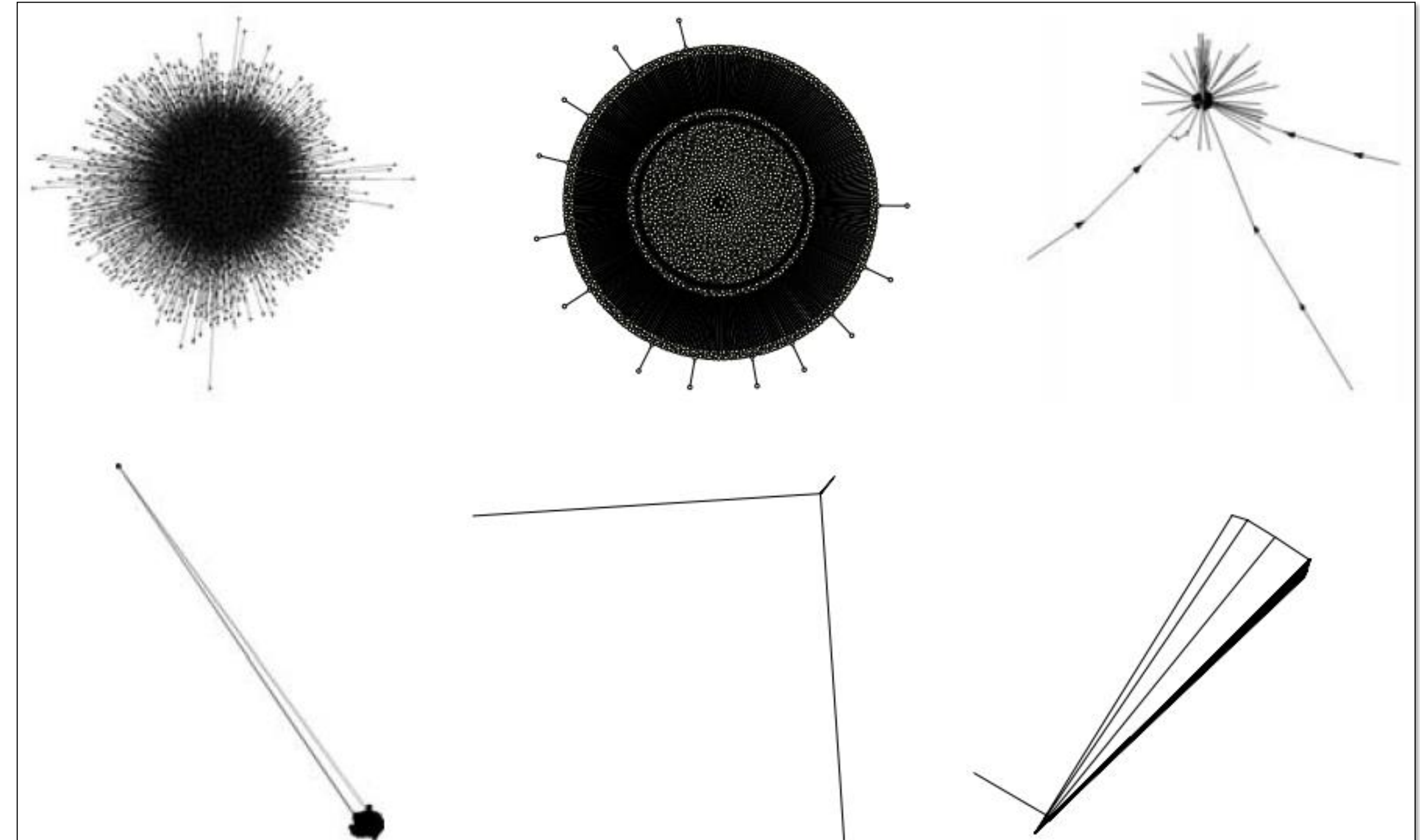


# Algorithm Comparisons

Graph A



Graph B



# How to compare?

User performance,  
controlled experiments

[Huang et al., 2007](#), etc.

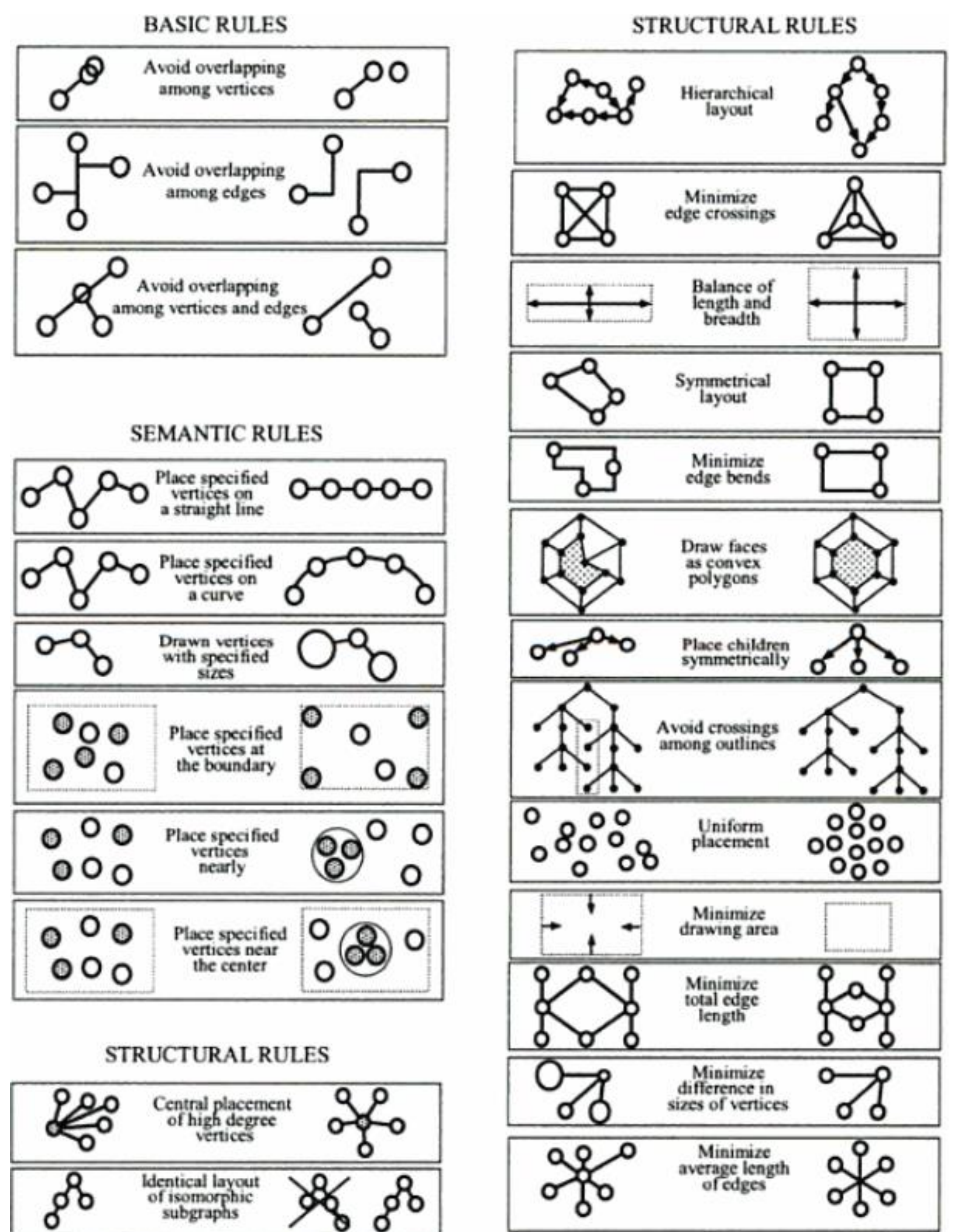
Simple rules or heuristics

[Davidson & Harel, 1996](#)

Computational evaluations,  
Global and local readability metrics

[Purchase et al., 2002](#)

[Dunne et al., 2015](#)



## Benchmark datasets

AT&T  
Biological Pathways (KEGG)  
Co-Phylogenetic Trees  
Complete Graphs  
World Maps  
Graphviz Examples  
KnownCR  
Militarized Interstate Disputes (MID)  
North DAGs  
RandDAG  
Rome-Lib  
Scotch Graph Collection  
SteinLib  
Storylines (Movie Plots)  
WebCompute

## Established Network Repositories

Matrix Market  
Network Repository  
Pajek  
SNAP (Stanford Network Analysis Platform)  
SuiteSparse Matrix Collection

## Aggregate collections

Airlines, Migrations, and Air Traffic  
Assorted Collaboration Network  
Blogposts, Tweets, and Forums  
Code Dependency Graphs

# Graph Layout Benchmark Datasets

The following is a list of benchmark datasets for testing graph layout algorithms. The list was collected at the [Northeastern University Visualization Lab](#), and is maintained by the same. Our collection methodology targeted layout algorithms specifically - we do acknowledge the existence of other repositories that target other network-related purposes more in detail. The collection and supplemental material is also accessible at <https://osf.io/j7ucv/>

Click on the names of the collections to expand them and access information about their contents and a list of papers using them.

If you find our work useful for your research, consider citing our paper as well as the linked "Origin Paper" for each dataset used:

```
@Misc{DiBartolomeo2023CollectionBenchmarkDatasets,  
  author      = {Di~Bartolomeo, Sara and Puerta, Eduardo and Wilson, Connor and Crnovrsanin, Tarik and Du  
  howpublished = {Under submission to Graph Drawing Posters},  
  title       = {A collection of benchmark datasets for evaluating graph layout algorithms},  
  year        = {2023},  
  url         = {https://visdunneright.github.io/gd_benchmark_sets/},  
}
```

**Contributing:** Please open an issue here: [https://github.com/VisDunneRight/gd\\_benchmark\\_sets](https://github.com/VisDunneRight/gd_benchmark_sets).  
Alternatively, reach out to [dibartolomeo.sara@gmail.com](mailto:dibartolomeo.sara@gmail.com)

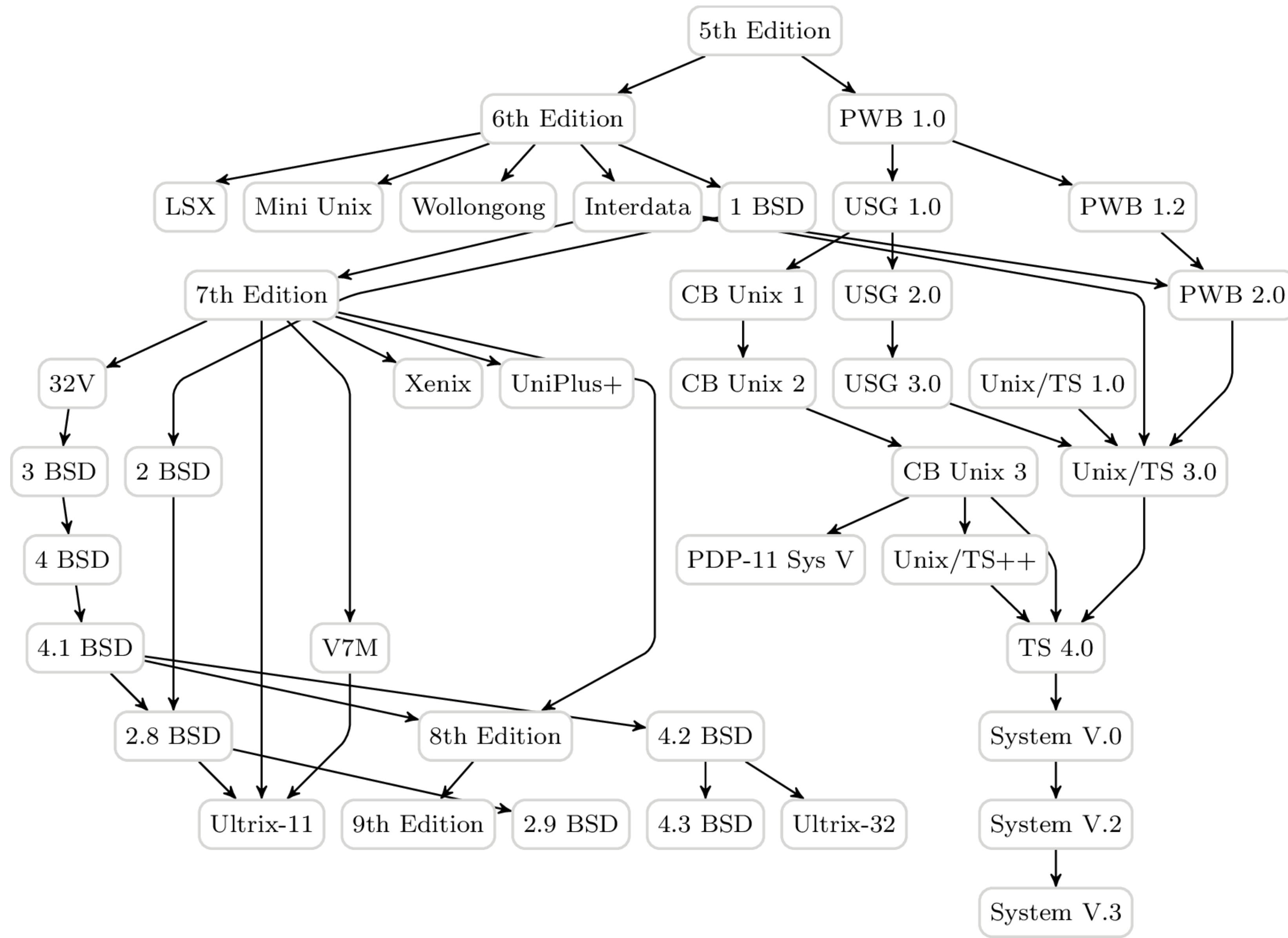
## Benchmark datasets

These are collections of graphs that have been frequently used in graph drawing papers. By clicking on each collection name, you can see additional information. We also provide an analysis of the contents of each collection, and a list of papers that use them, their sources, and various types of information.

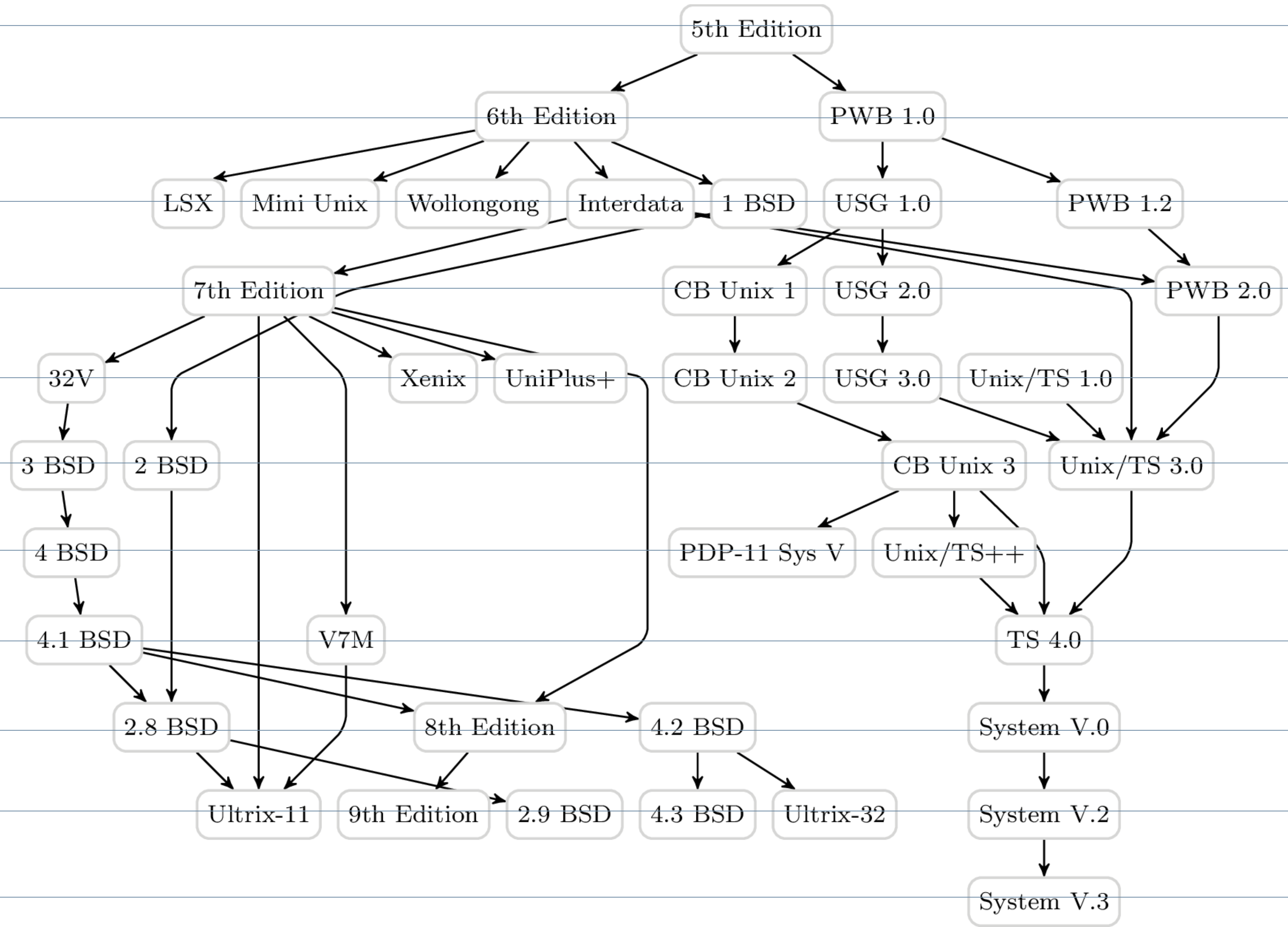
| Name                                       | Node Distr. | Min nodes | Max nodes | Features   |
|--|-------------|-----------|-----------|--|
| <a href="#">AT&amp;T</a>                   |             | 10        | 100       | acyclic, directed edges                            |
| <a href="#">Biological Pathways (KEGG)</a> |             | 47        | 292       | clusters, directed edges, large, partition         |
| <a href="#">Co-Phylogenetic Trees</a>      |             | 13        | 773       | trees  |
| <a href="#">Complete Graphs</a>            |             | 5         | 80        | bipartite, generic, known crossing number          |
| <a href="#">World Maps</a>                 |             | 48        | 514       | categorical nodes, dynamic, node weighted, spatial |



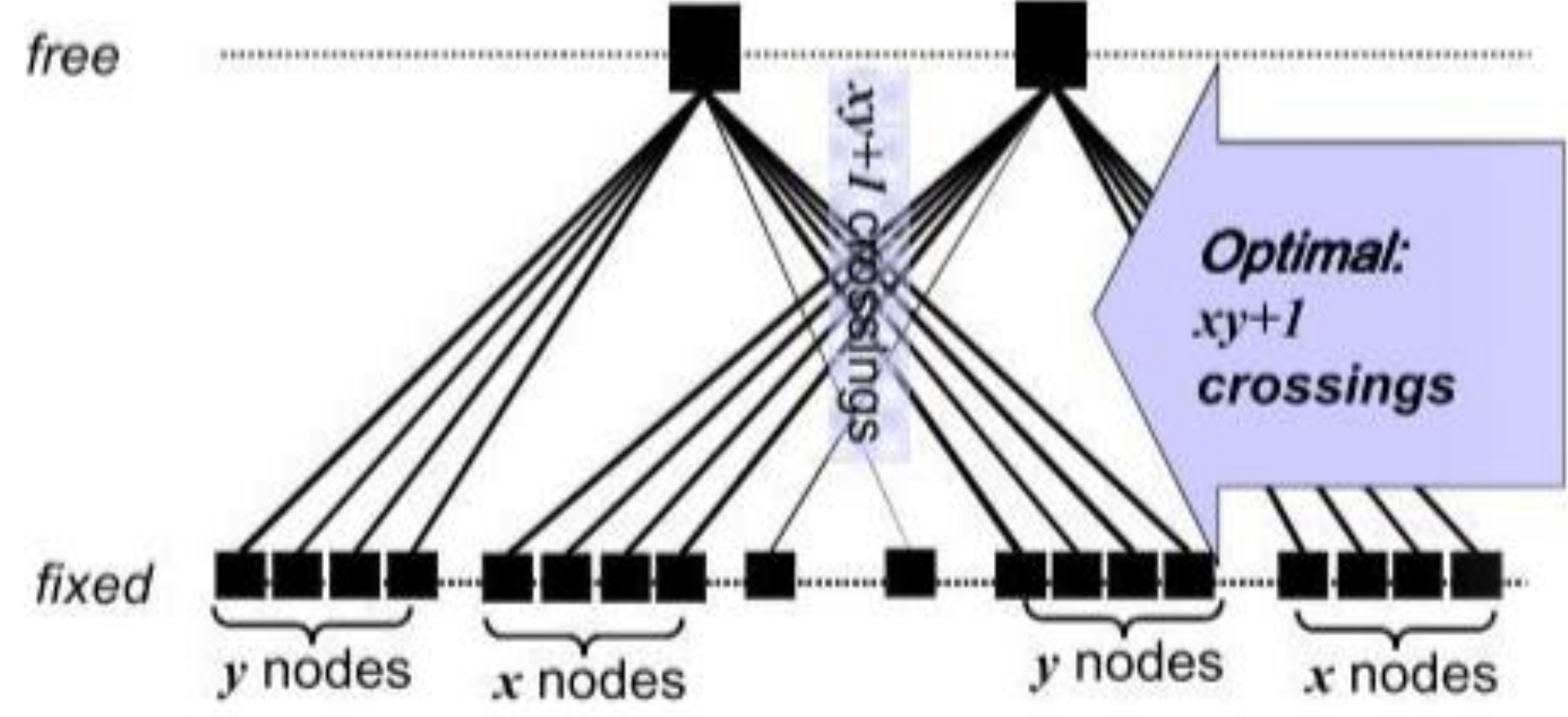
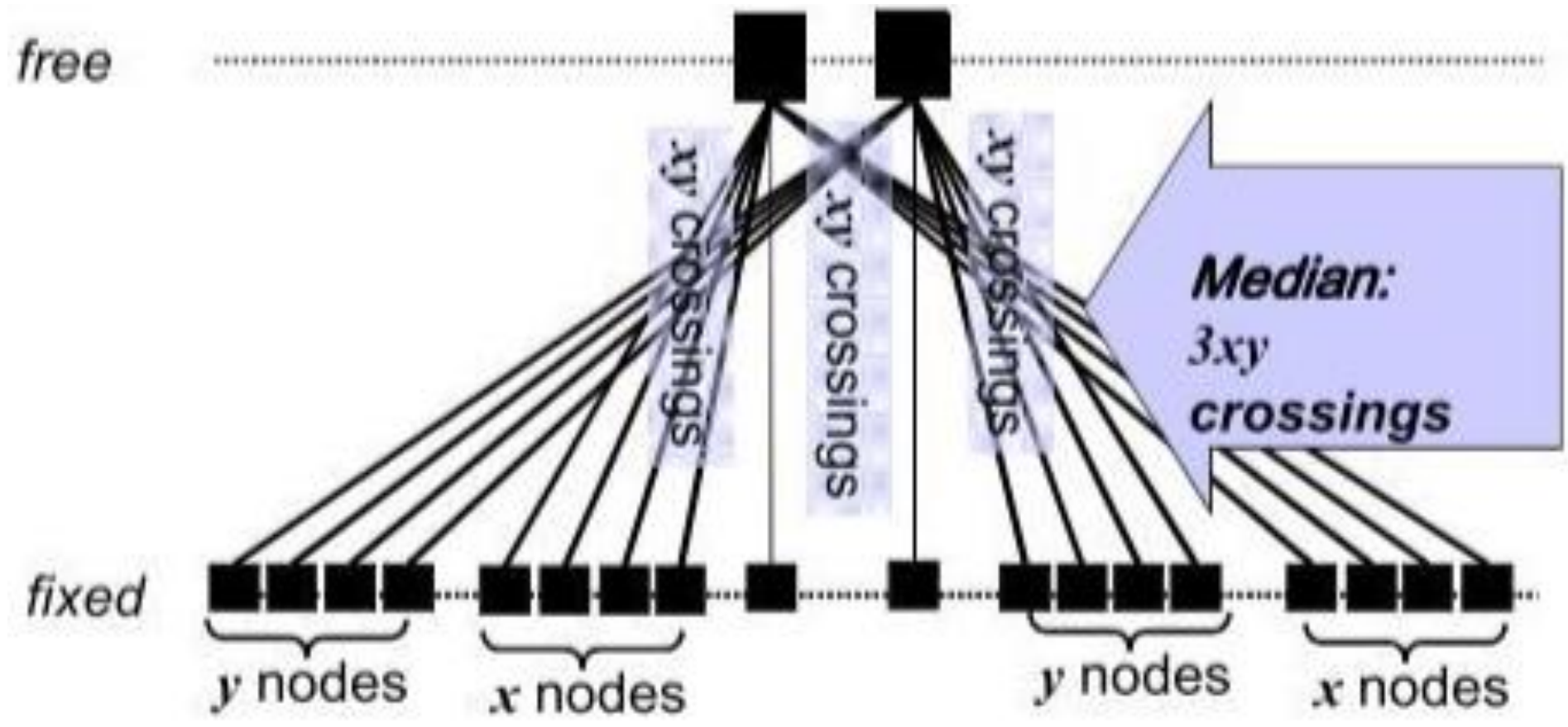
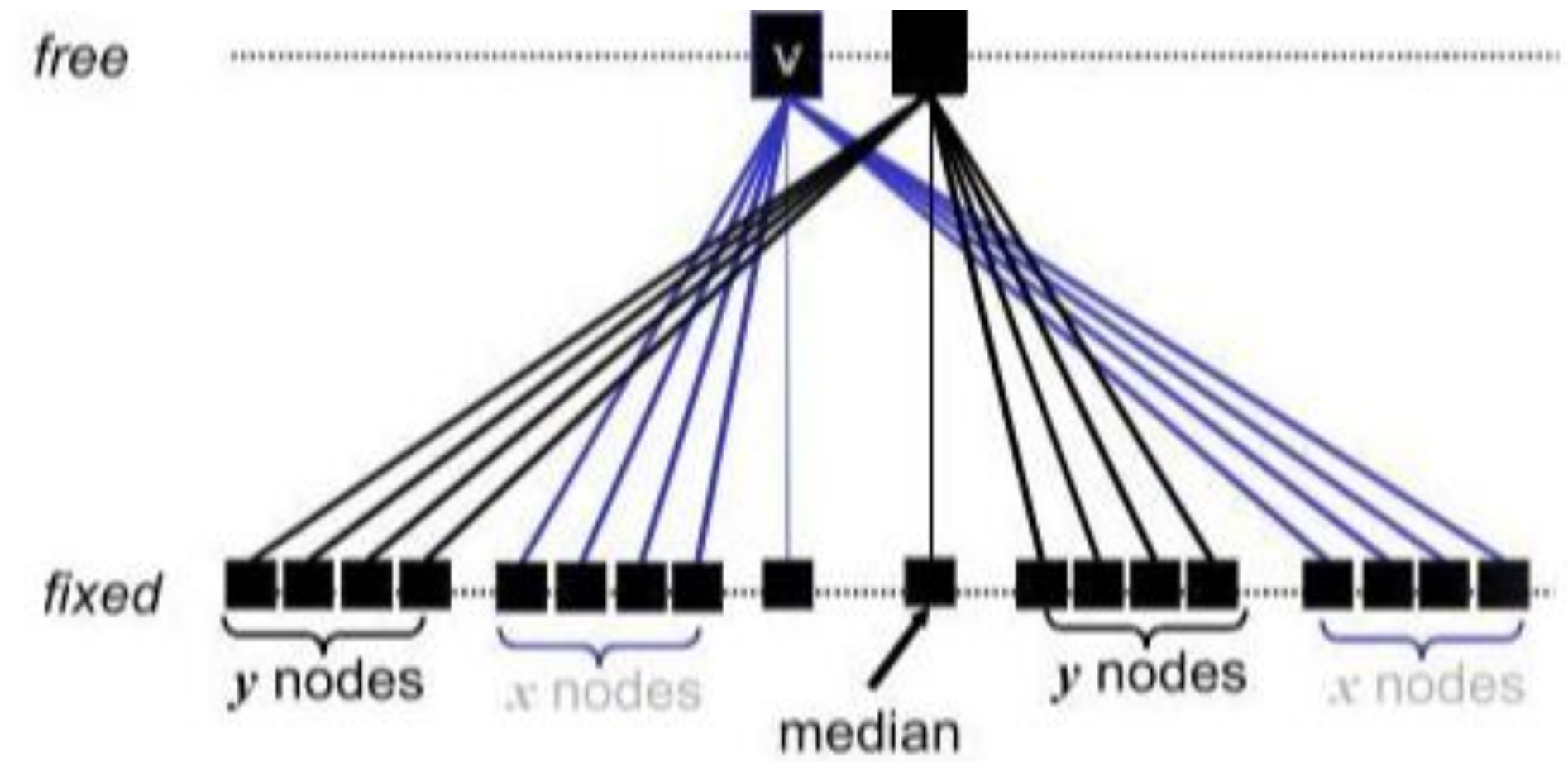
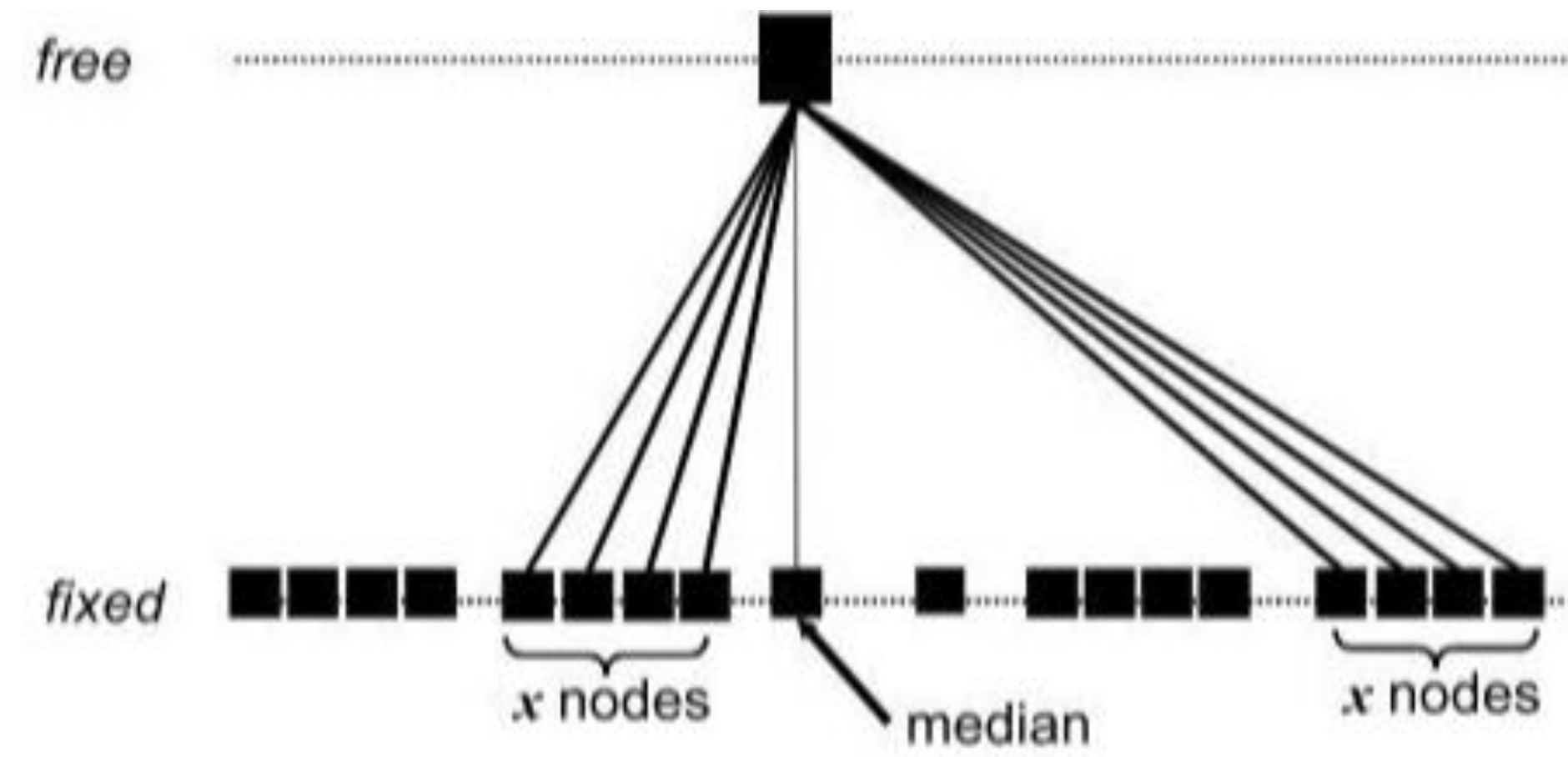
# Rooted trees / layered graph drawing



# Back-and-Forth Sweeps



# Median Heuristic



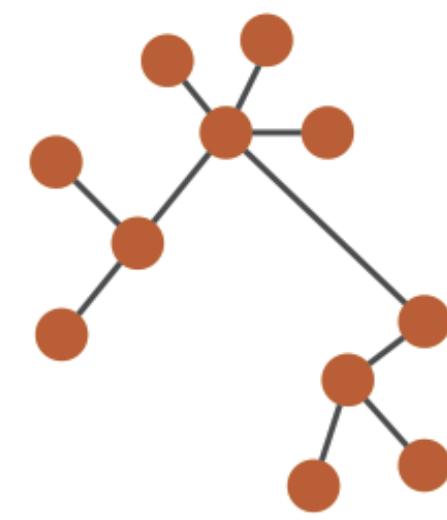


## Node-Link Diagrams

Connection Marks

✓ NETWORKS

✓ TREES



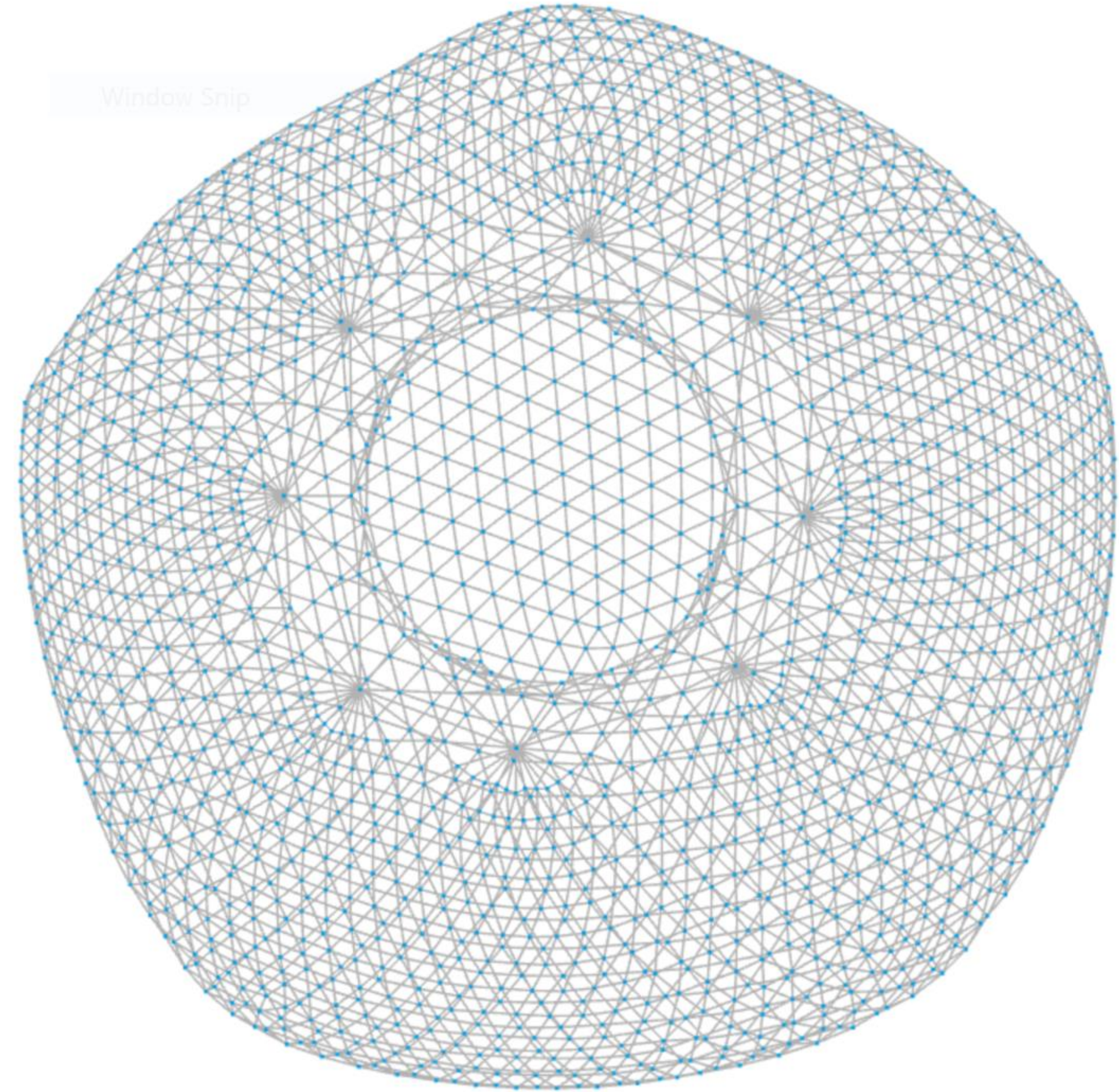
# Scale Problems...

- Quickly run out of space!
- Tree breadth often grows exponentially
- Layout algorithms are slow and heuristics
- Slow rendering
- Solutions:
  - scrolling or panning
  - filtering or zooming
  - aggregation & simplification
  - faster but trickier rendering approaches

### Choose Graph:

FAVORITE GRAPHS

- HB/blckhole**
- Bai/rw5151
- HB/bcsstm13
- HB/jagmesh6
- HB/watt\_1
- HB/lshp1882
- HB/plat1919
- HB/bcsstk26
- Bai/dw256A
- Bai/tols2000
- Bai/dw1024
- Bai/rdb2048
- Pajek/CSphd
- GHS\_indef/laser
- BAI
- bfwa398
- bfwa62
- bfwb398
- bfwb62
- bfwb782
- bwm200
- cdde1
- cdde2
- cdde3
- cdde4
- cdde5
- cdde6
- ck104
- ck400
- ck656



### Layout Settings

- Spring Coeff:
- Spring Length:
- Gravity Coeff:
- Drag Coeff:
- Theta Coeff:

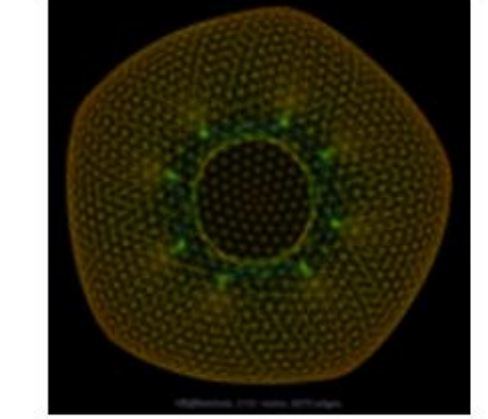
Reset to default

### HB/blckhole

Nodes: 2121

Edges: 6370

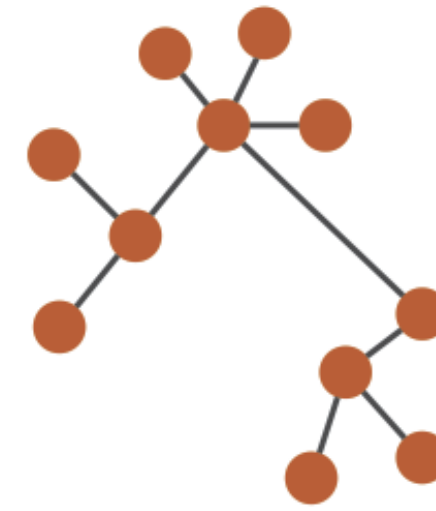
Image:



# Arrange Networks and Trees

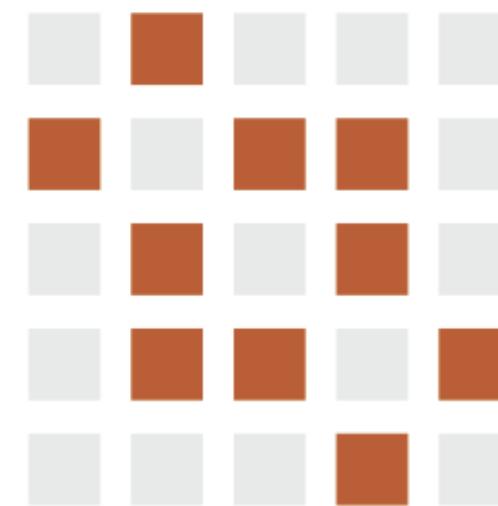
→ **Node-Link Diagrams**  
Connection Marks

✓ NETWORKS    ✓ TREES



→ **Adjacency Matrix**  
Derived Table

✓ NETWORKS    ✓ TREES



→ **Enclosure**  
Containment Marks

✗ NETWORKS    ✓ TREES



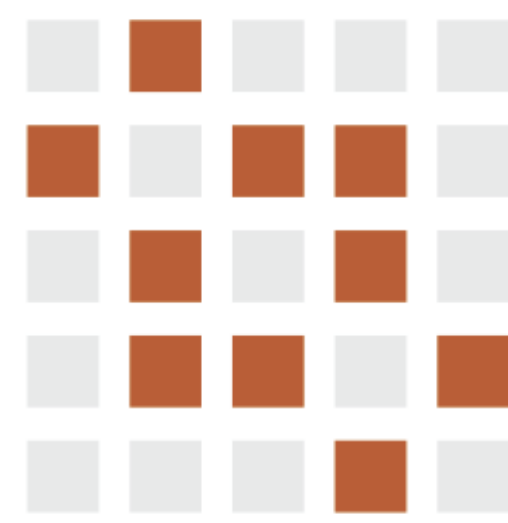
“Treemap”

# → Adjacency Matrix

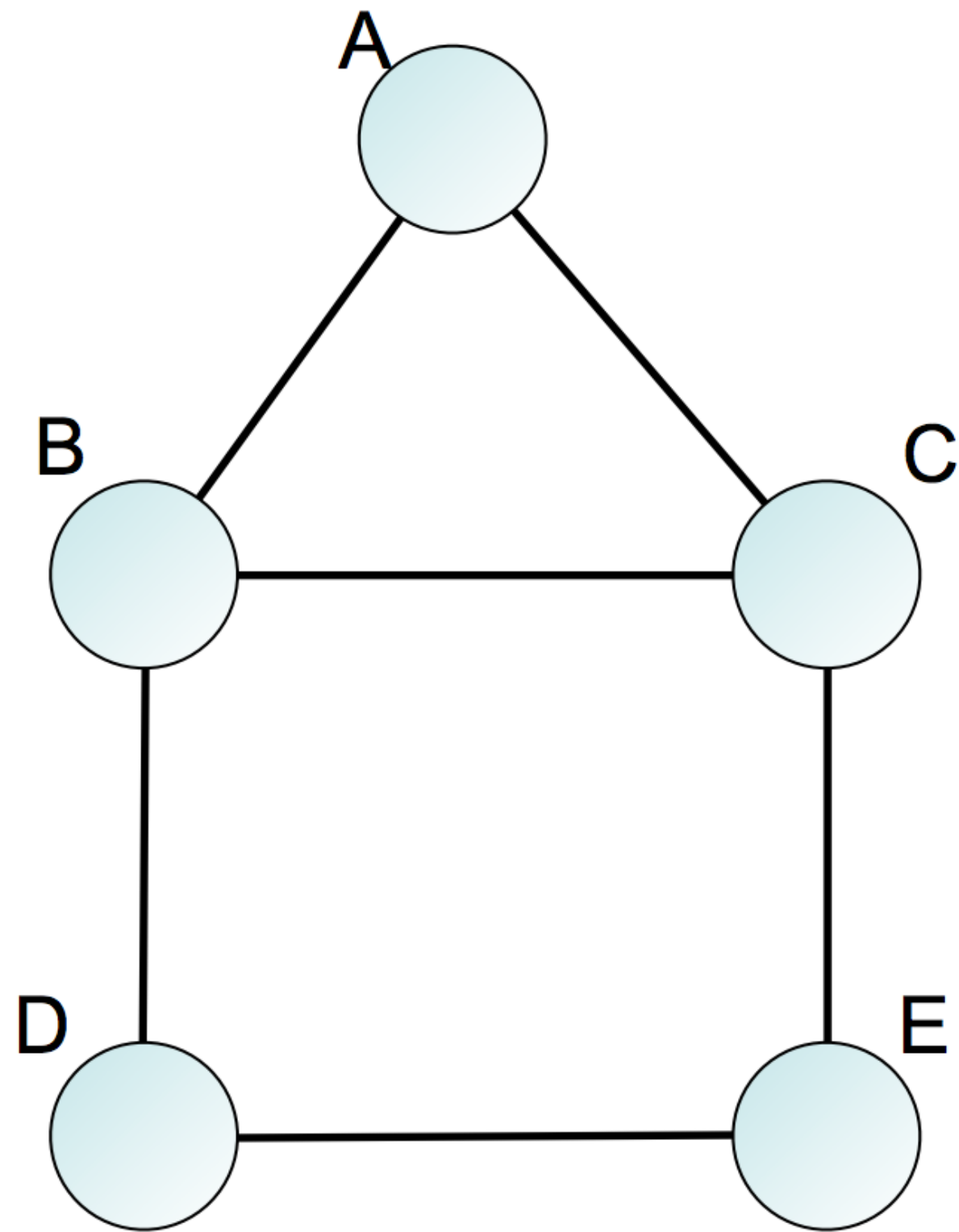
Derived Table

✓ NETWORKS

✓ TREES

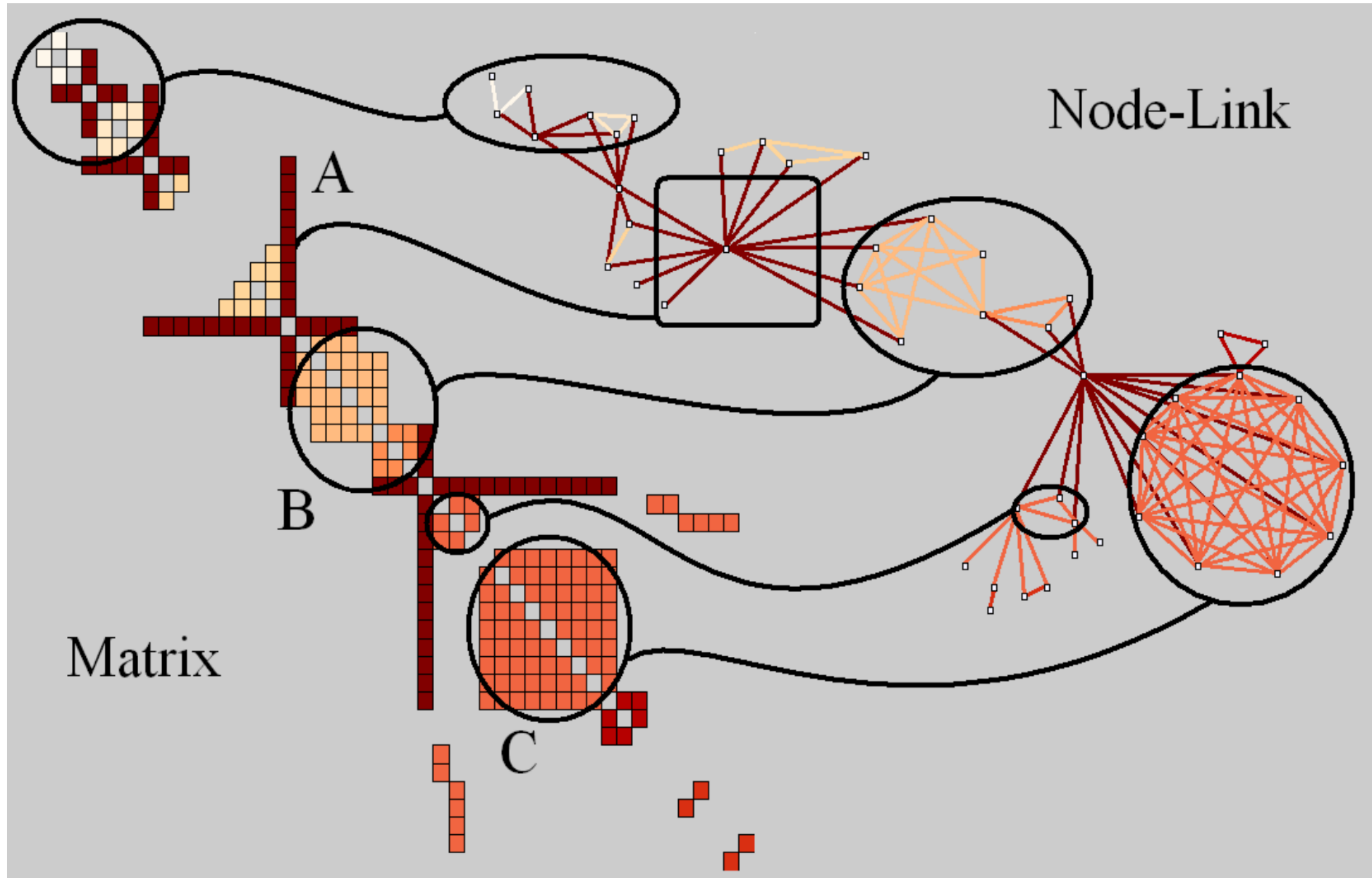


Alternate to node-link visualization for dense & weighted networks



|   | A | B | C | D | E |
|---|---|---|---|---|---|
| A |   | ■ | ■ |   |   |
| B | ■ |   | ■ | ■ |   |
| C | ■ | ■ |   |   | ■ |
| D |   | ■ |   |   | ■ |
| E |   |   | ■ | ■ |   |

# Adjacency Matrix



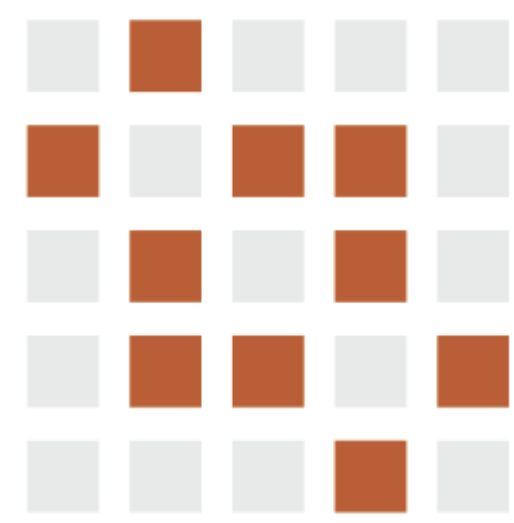


## → Adjacency Matrix

Derived Table

✓ NETWORKS

✓ TREES



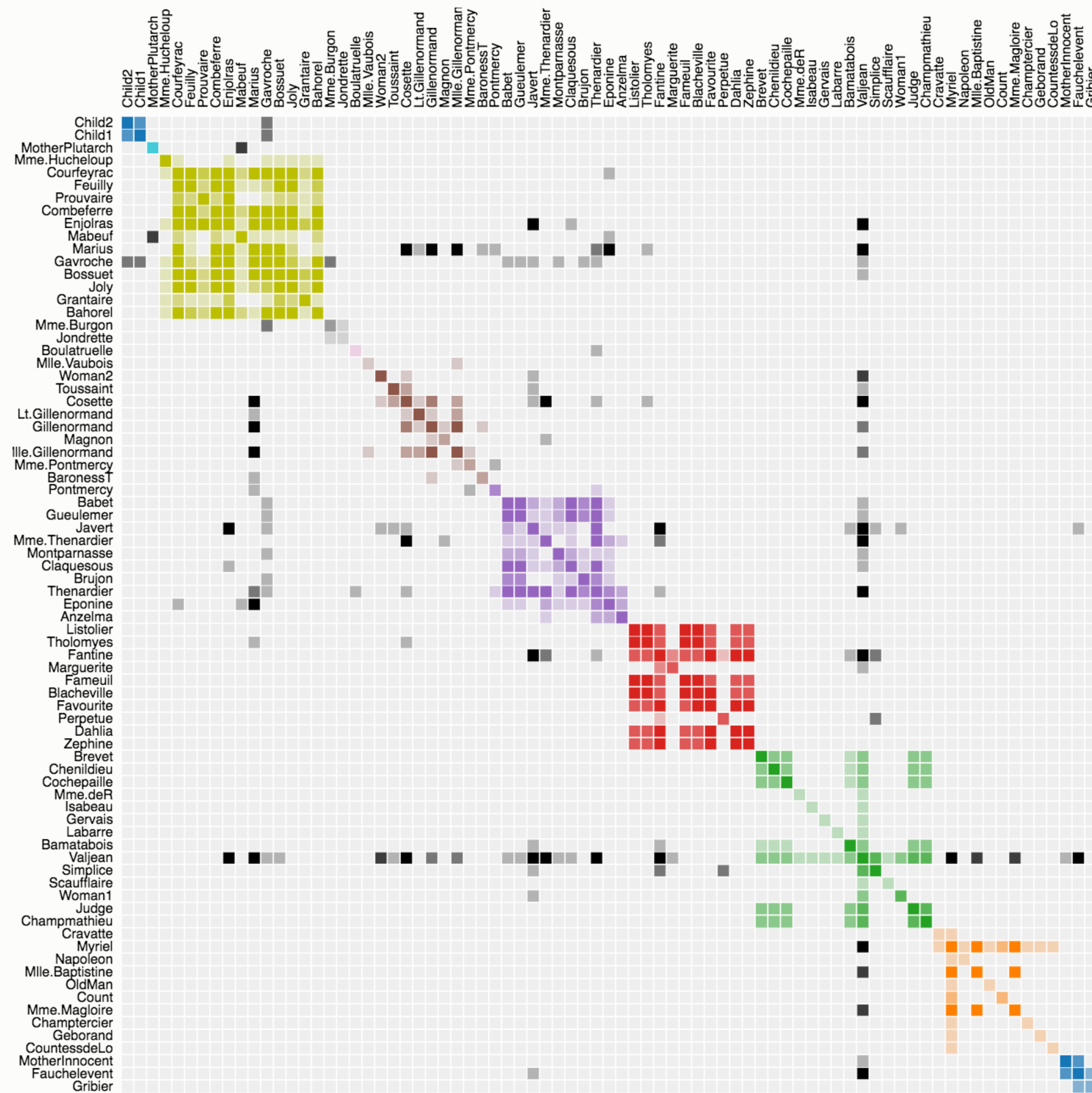
### Pros:

- great for dense graphs
- visually scalable
- can spot clusters

### Cons:

- **row order affects what you can see**
- abstract visualization
- hard to follow paths

# Les Misérables Co-occurrence



Source: [The Stanford GraphBase](#).

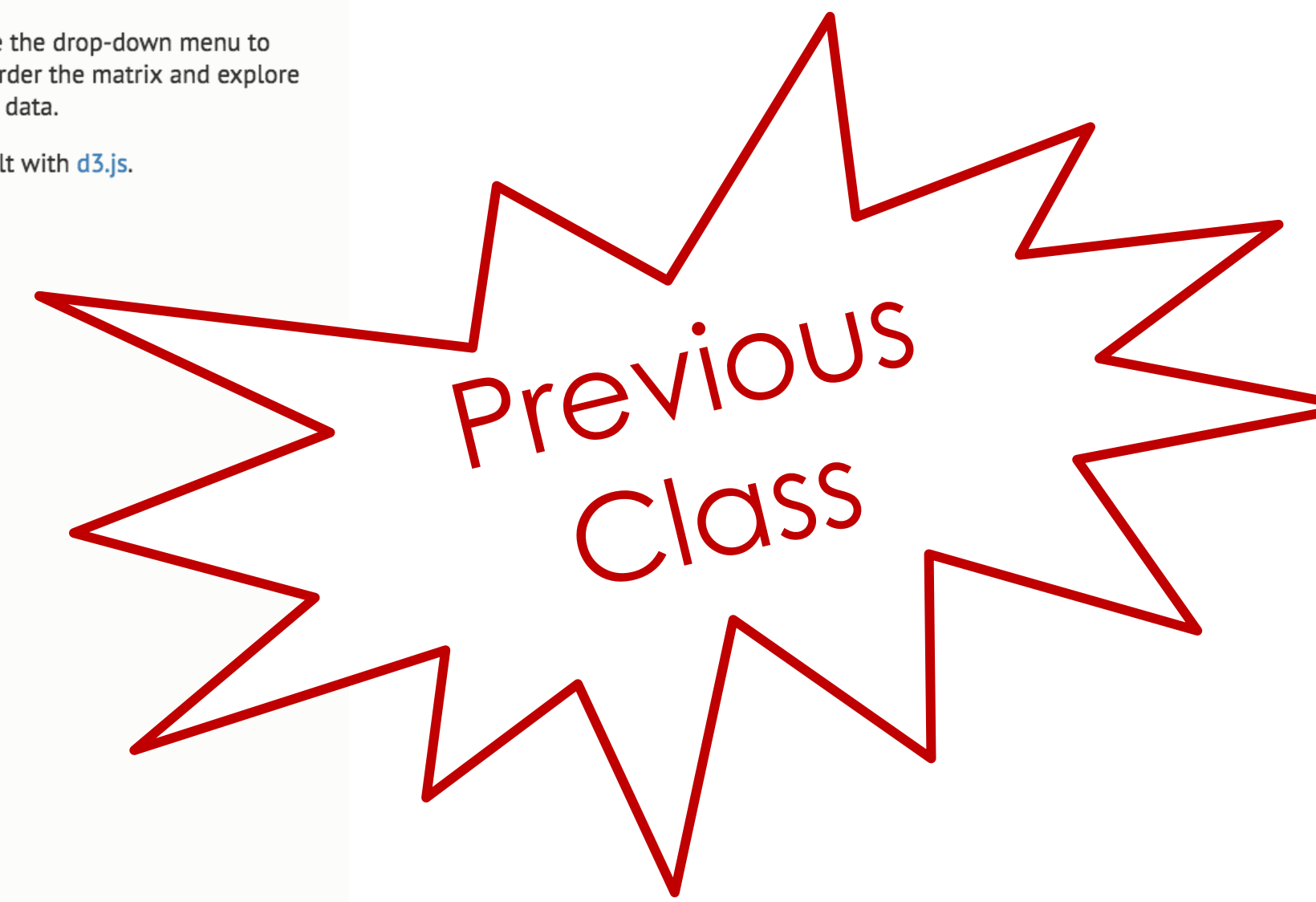
Order:

This matrix diagram visualizes character co-occurrences in Victor Hugo's *Les Misérables*.

Each colored cell represents two characters that appeared in the same chapter; darker cells indicate characters that co-occurred more frequently.

Use the drop-down menu to reorder the matrix and explore the data.

Built with [d3.js](#).



<https://bost.ocks.org/mike/miserables/>

# WDA-LS clustered co-occurrence

Use the drop-down menu to reorder the matrix and explore the data.

When ordered by cluster, rows and columns are clustered by affinity values using hierarchical agglomerative clustering.

Distance measure: Euclidean.

Linkage technique: Single.

Rows and columns are then arranged using leaf reordering using the algorithm from: Sakai, Ryo, et al. "Dendsort: modular leaf ordering methods for dendrogram representations in R." *F1000Research* 3 (2014).

Cell labels show count and color shows normalized affinity.

[Cody Dunne](#) and [Tim Stutts](#), IBM Watson Health [Cognitive Visualization Lab](#)

Dataset:

Order:

The query was for genes related to the genes *SOX9*, *TCF7L1*, *SMAD4*, *PIK3CA*, *KRAS* in Medline.

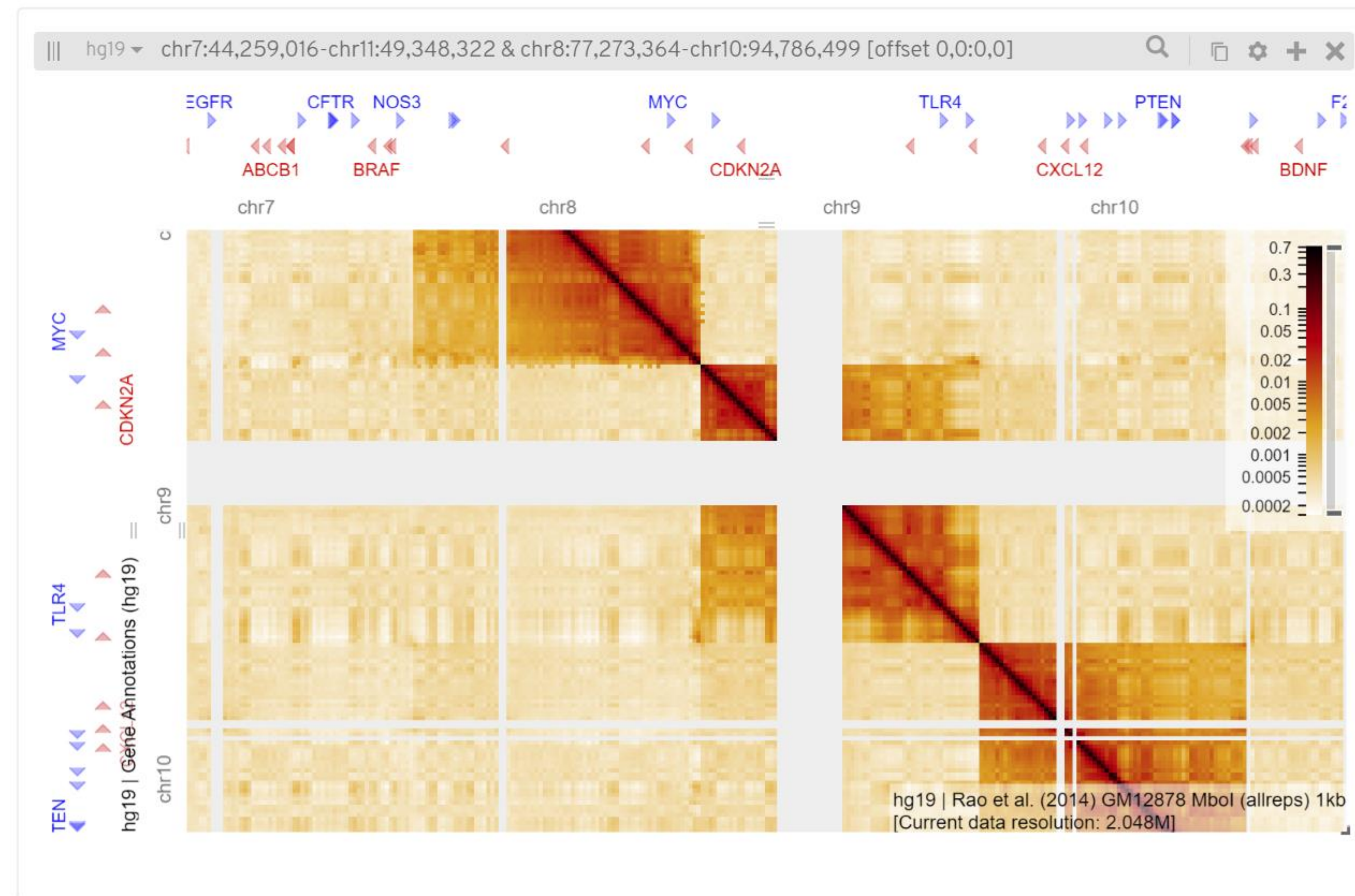
|        | SOX9 | TCF7L1 | SMAD4 | KRAS  | PIK3CA |
|--------|------|--------|-------|-------|--------|
| tp53   | 33   | 4      | 406   | 1295  | 726    |
| apc    | 10   | 1      | 106   | 255   | 91     |
| kras   | 10   | 1      | 166   | 11277 | 926    |
| nras   | 0    | 0      | 20    | 878   | 269    |
| hras   | 0    | 0      | 9     | 659   | 107    |
| f2     | 2    | 0      | 5     | 407   | 0      |
| raf1   | 3    | 1      | 12    | 760   | 266    |
| alk    | 0    | 0      | 11    | 339   | 126    |
| ns2    | 0    | 0      | 0     | 228   | 0      |
| sos1   | 0    | 0      | 0     | 286   | 8      |
| hspb3  | 0    | 0      | 4     | 279   | 9      |
| ptpn11 | 0    | 0      | 6     | 192   | 21     |
| cd8a   | 4    | 0      | 7     | 190   | 25     |
| cd4    | 0    | 0      | 11    | 152   | 34     |
| ifng   | 0    | 0      | 14    | 118   | 12     |
| myc    | 18   | 1      | 50    | 278   | 80     |
| mlh1   | 0    | 1      | 34    | 190   | 50     |
| smad4  | 13   | 1      | 3052  | 166   | 53     |
| smad2  | 21   | 1      | 828   | 12    | 12     |
| smad3  | 20   | 0      | 658   | 6     | 12     |
| smad7  | 5    | 0      | 281   | 0     | 0      |
| smad1  | 17   | 0      | 262   | 0     | 6      |
| tgfb1  | 23   | 0      | 230   | 16    | 7      |
| inhbe  | 12   | 0      | 164   | 0     | 0      |
| tgfb2  | 5    | 0      | 123   | 22    | 6      |
| crkn2a | 13   | 0      | 222   | 330   | 150    |



HiGlass is a tool for exploring genomic contact matrices and tracks. Please take a look at the [examples and documentation](#) for a description of the ways that it can be configured to explore and compare contact matrices. To load private data, HiGlass can be [run locally within a Docker container](#). The HiC data in the examples below is from Rao et al. (2014) [2].

A preprint of the paper describing HiGlass is [available on bioRxiv](#) [1].

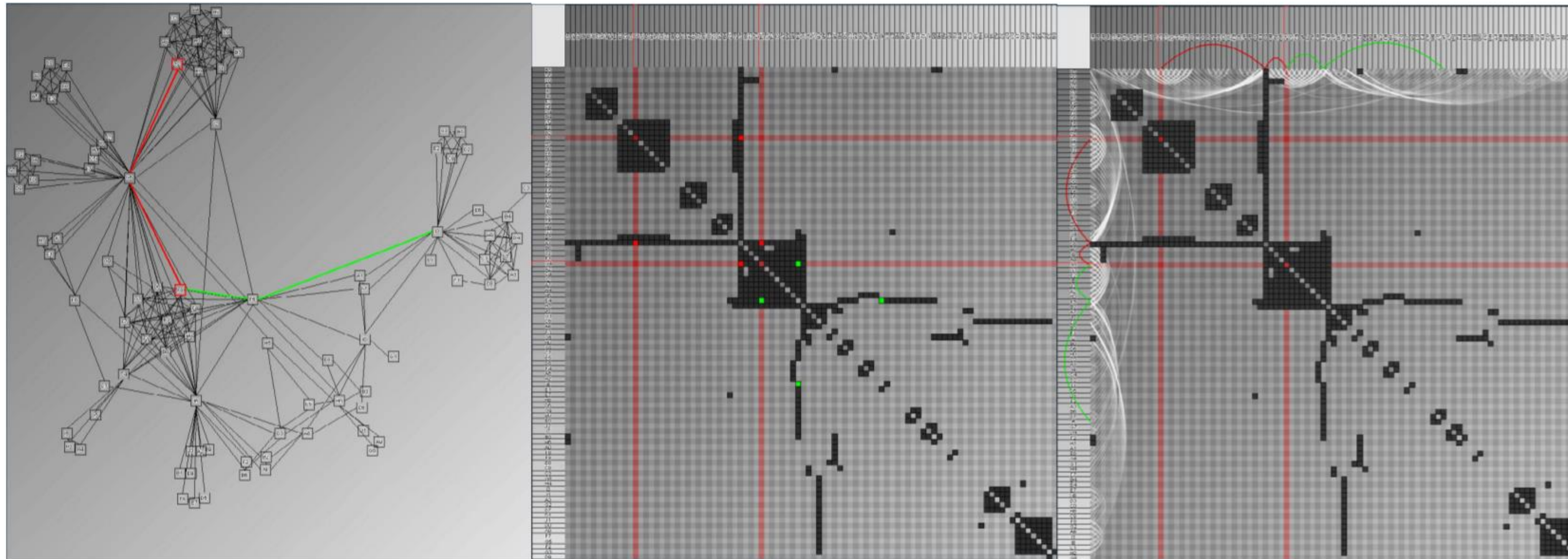
### Single View



<http://higlass.io/>



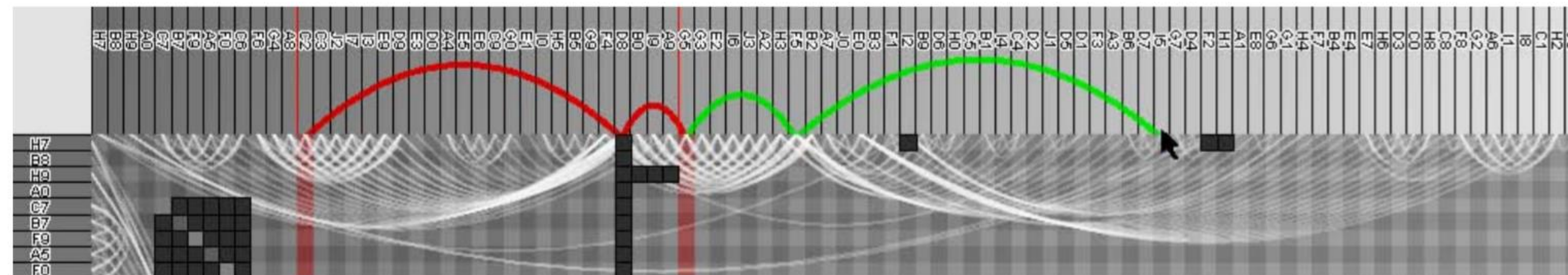
# MatLink



(a) Node-Link(NL)

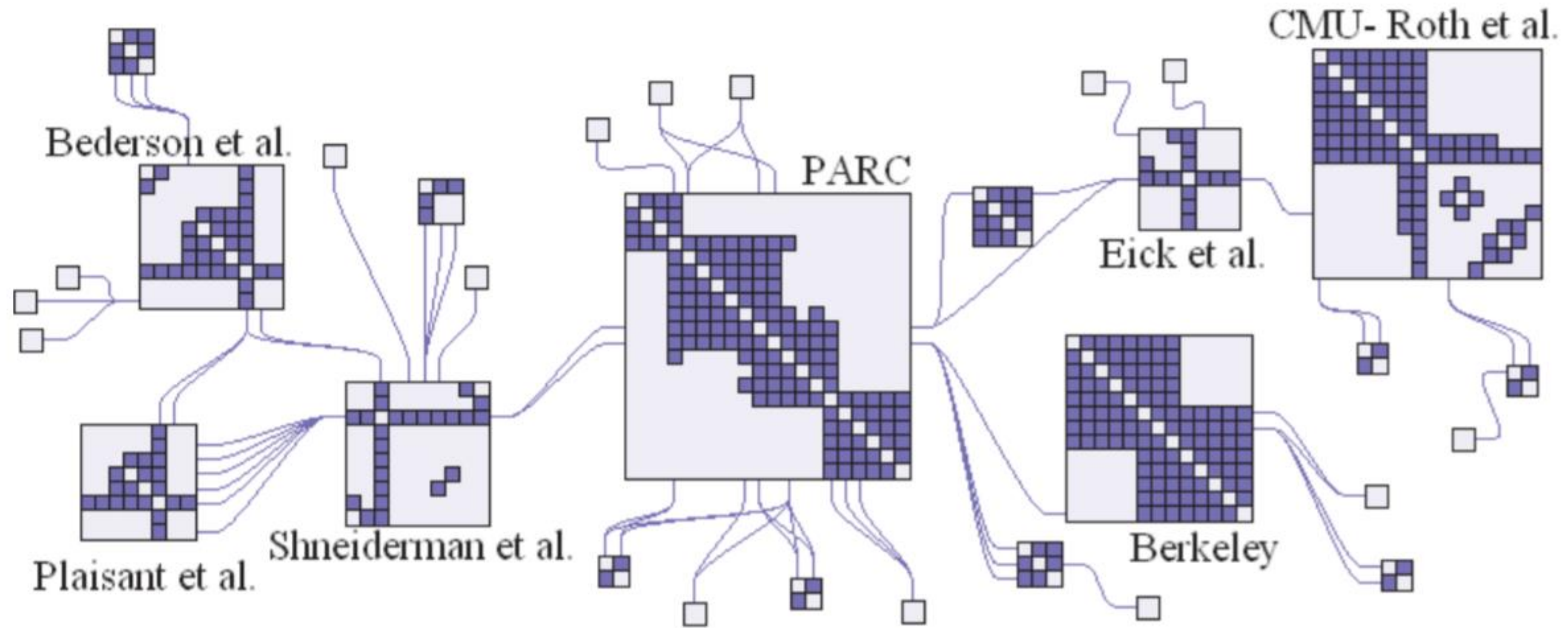
(b) Matrix(MAT)

(c) MatLink

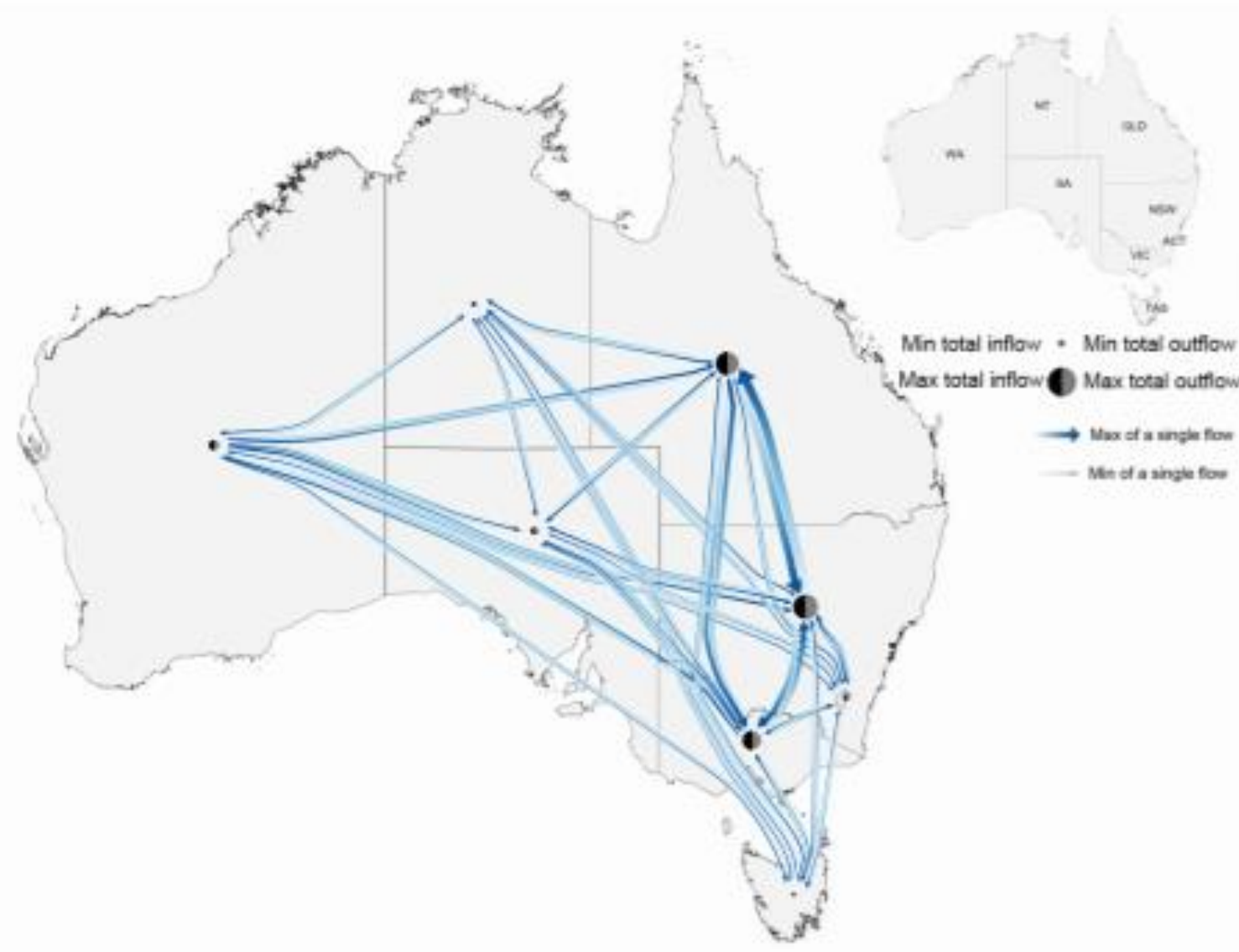


(d) Zoom on MatLink

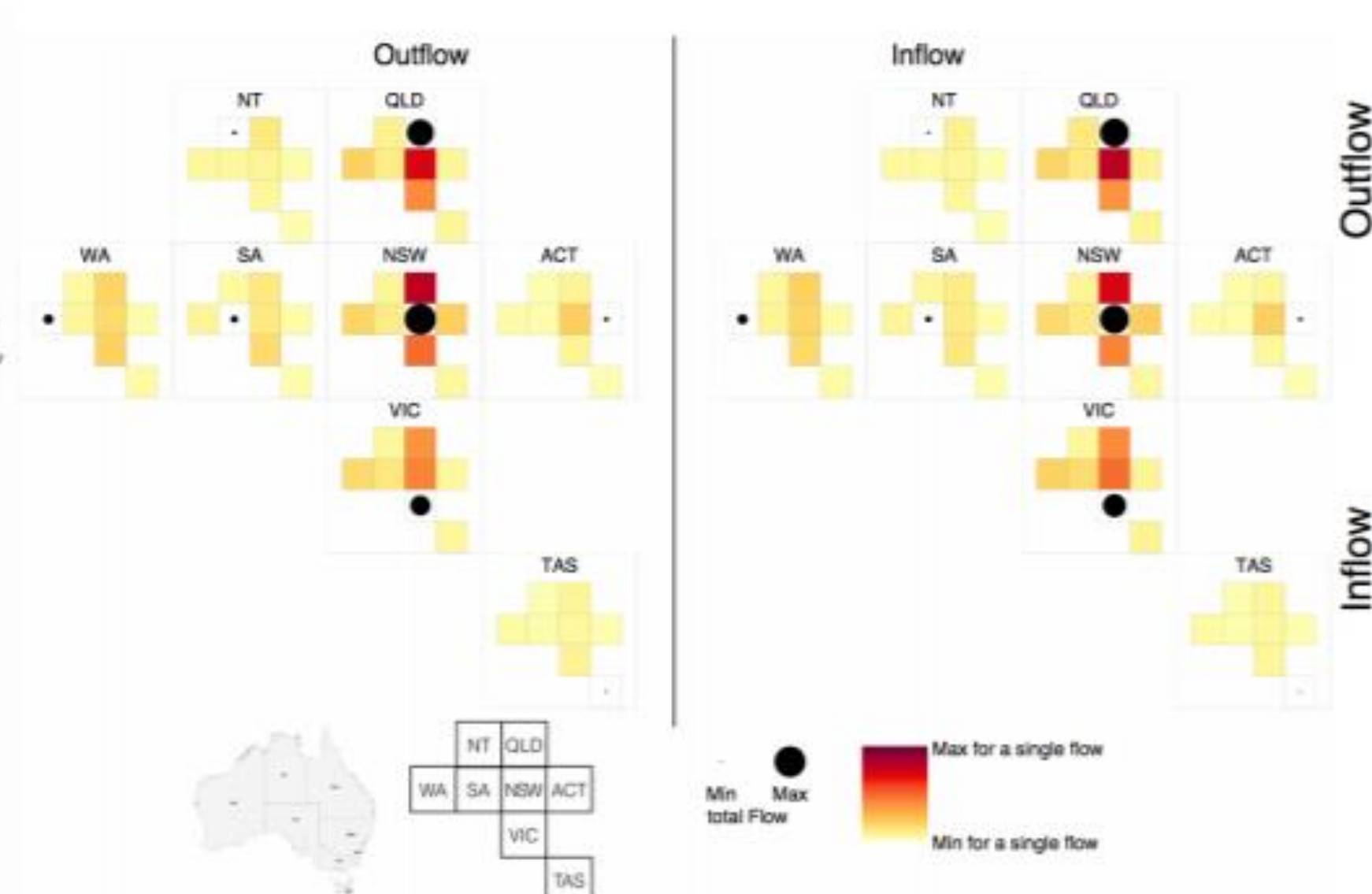
# NodeTrix



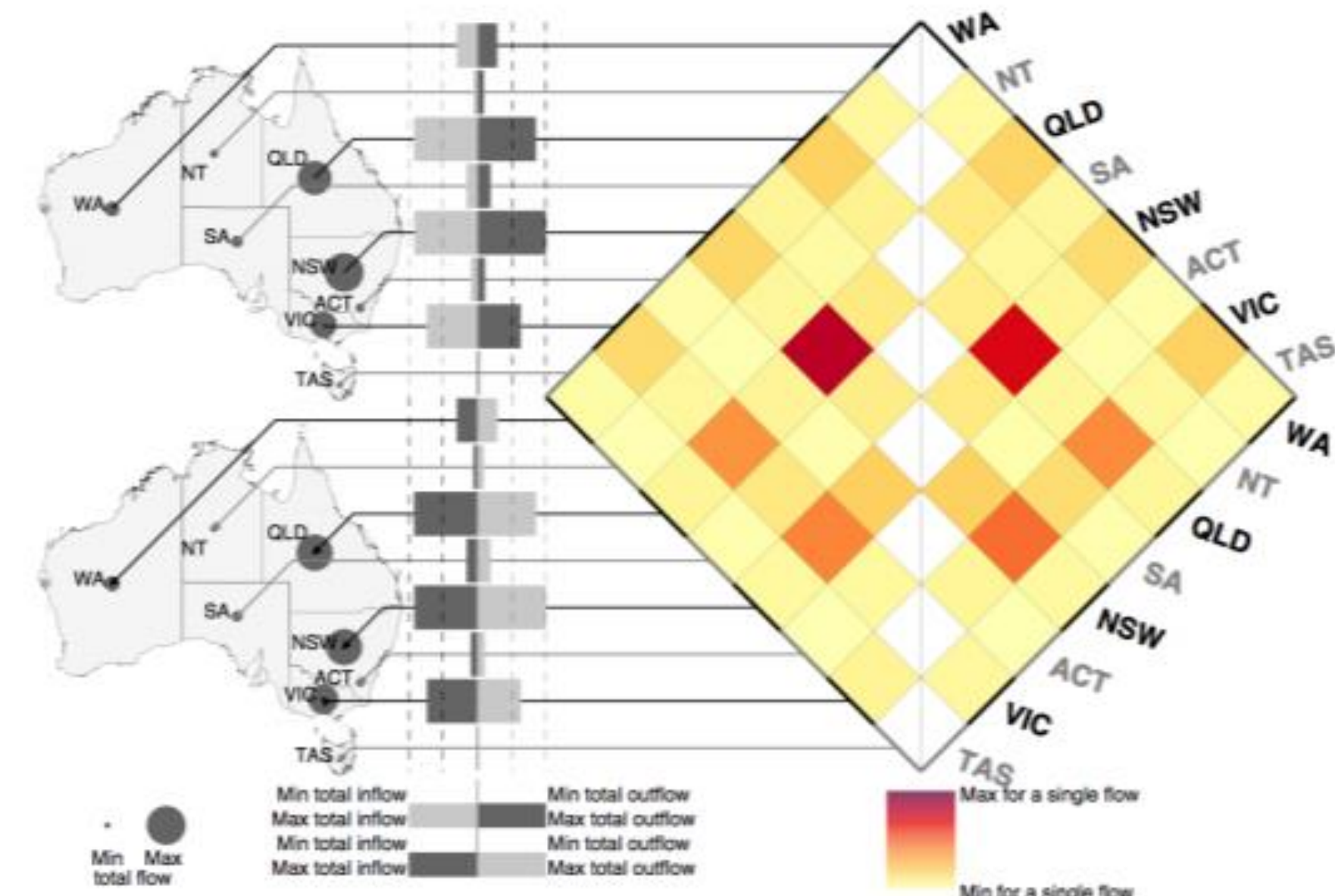
# MapTrix



(a) Bundled Flow Map



(b) OD Map

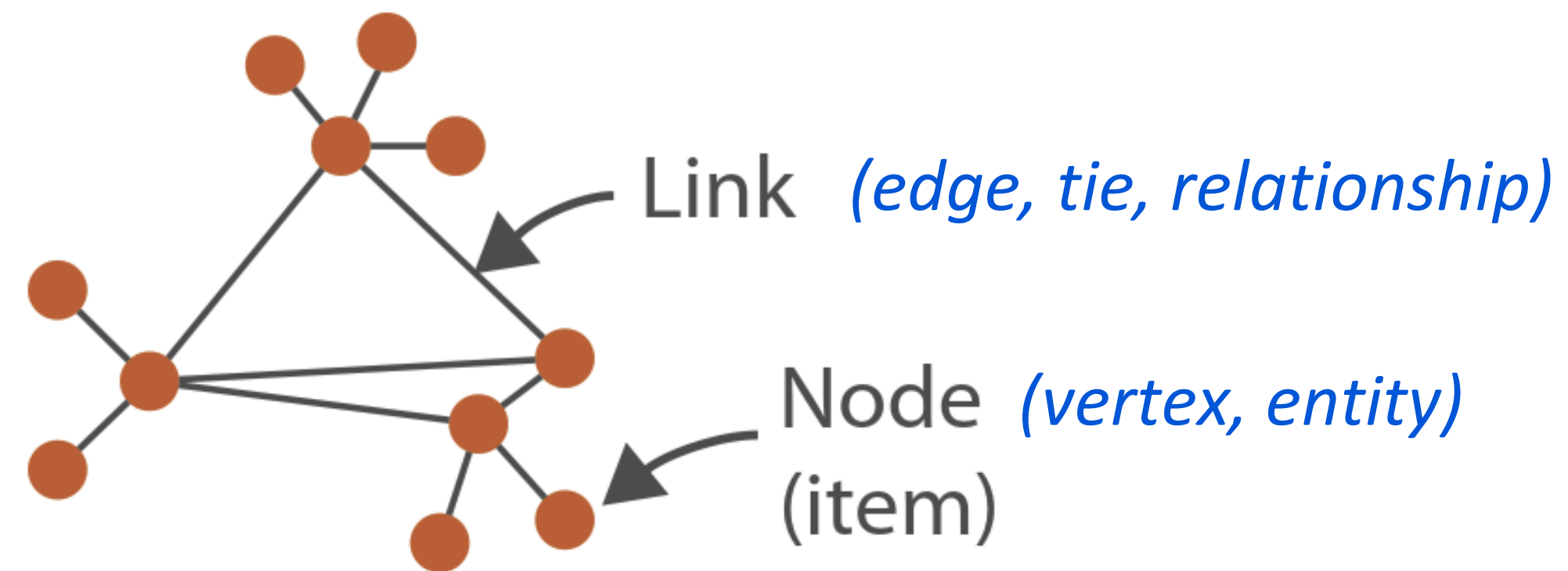


(c) MapTrix

<https://vimeo.com/182970812>  
<https://vimeo.com/278433529>

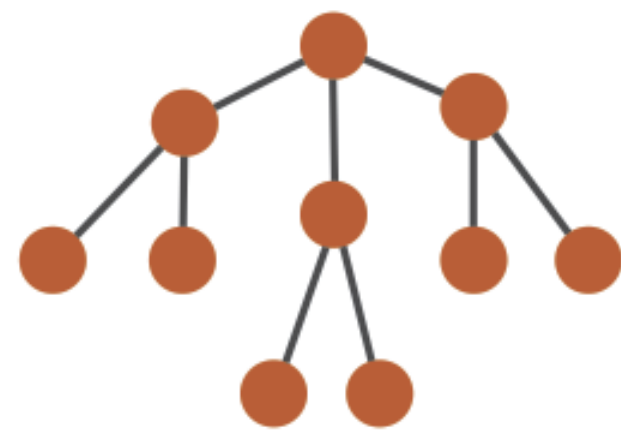
[Yang et al., 2016; Demo](#)

## → Networks *(graphs)*



Network = entities and relationships between them

### → *Trees*



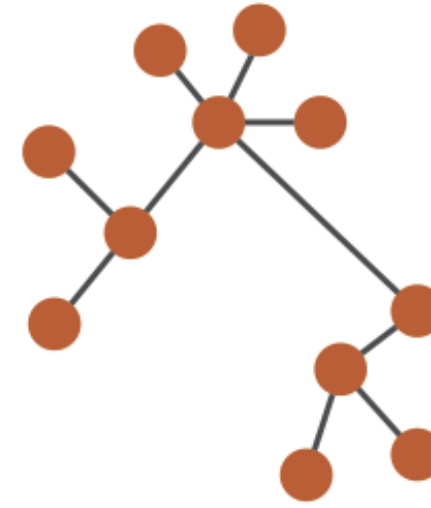
Tree = *undirected, connected, acyclic* network



# Arrange Networks and Trees

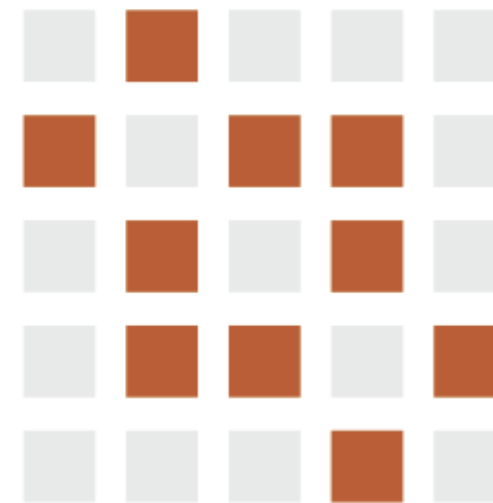
→ **Node-Link Diagrams**  
Connection Marks

✓ NETWORKS ✓ TREES



→ **Adjacency Matrix**  
Derived Table

✓ NETWORKS ✓ TREES



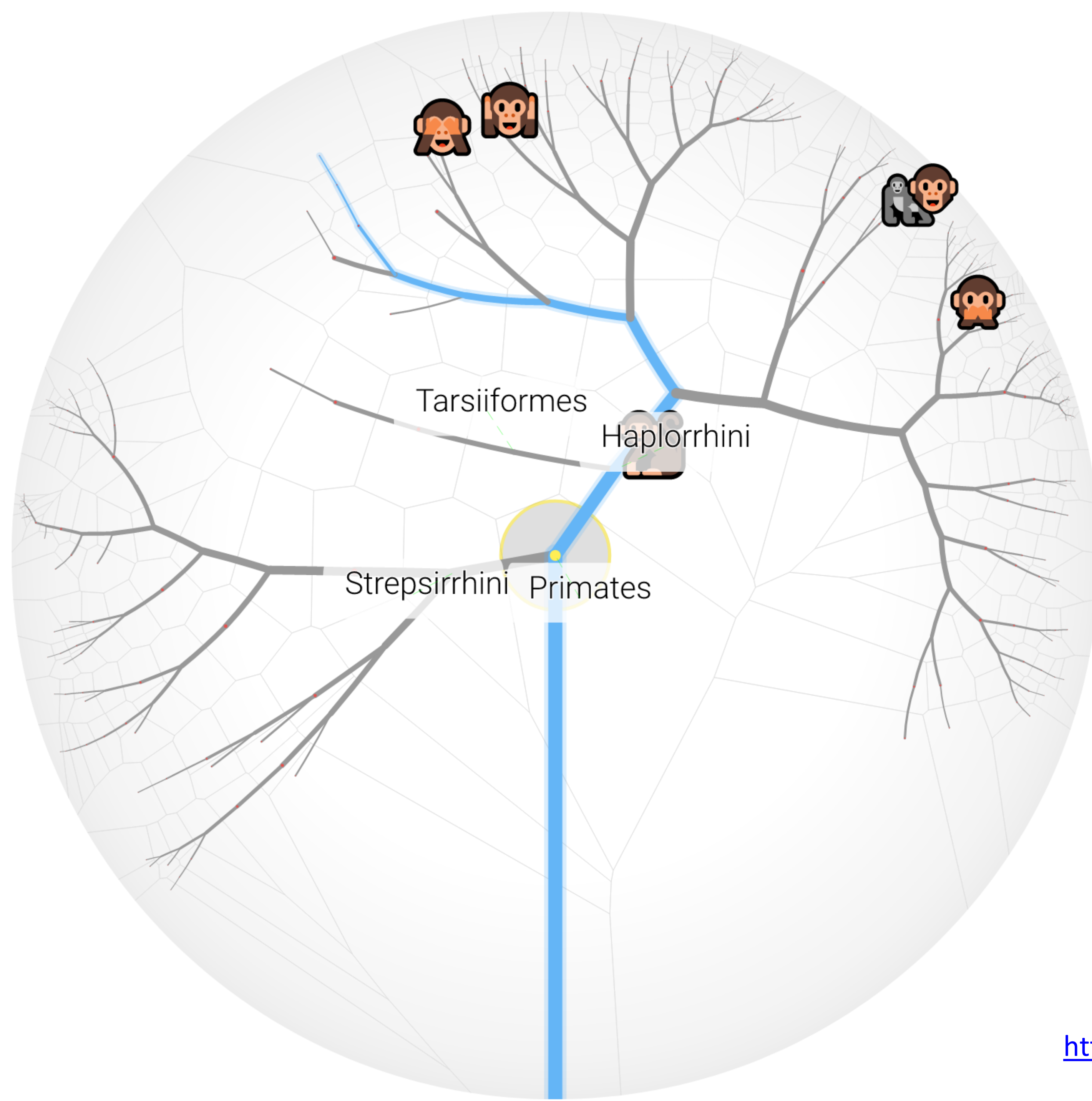
→ **Enclosure**  
Containment Marks

✗ NETWORKS ✓ TREES



“Treemap”

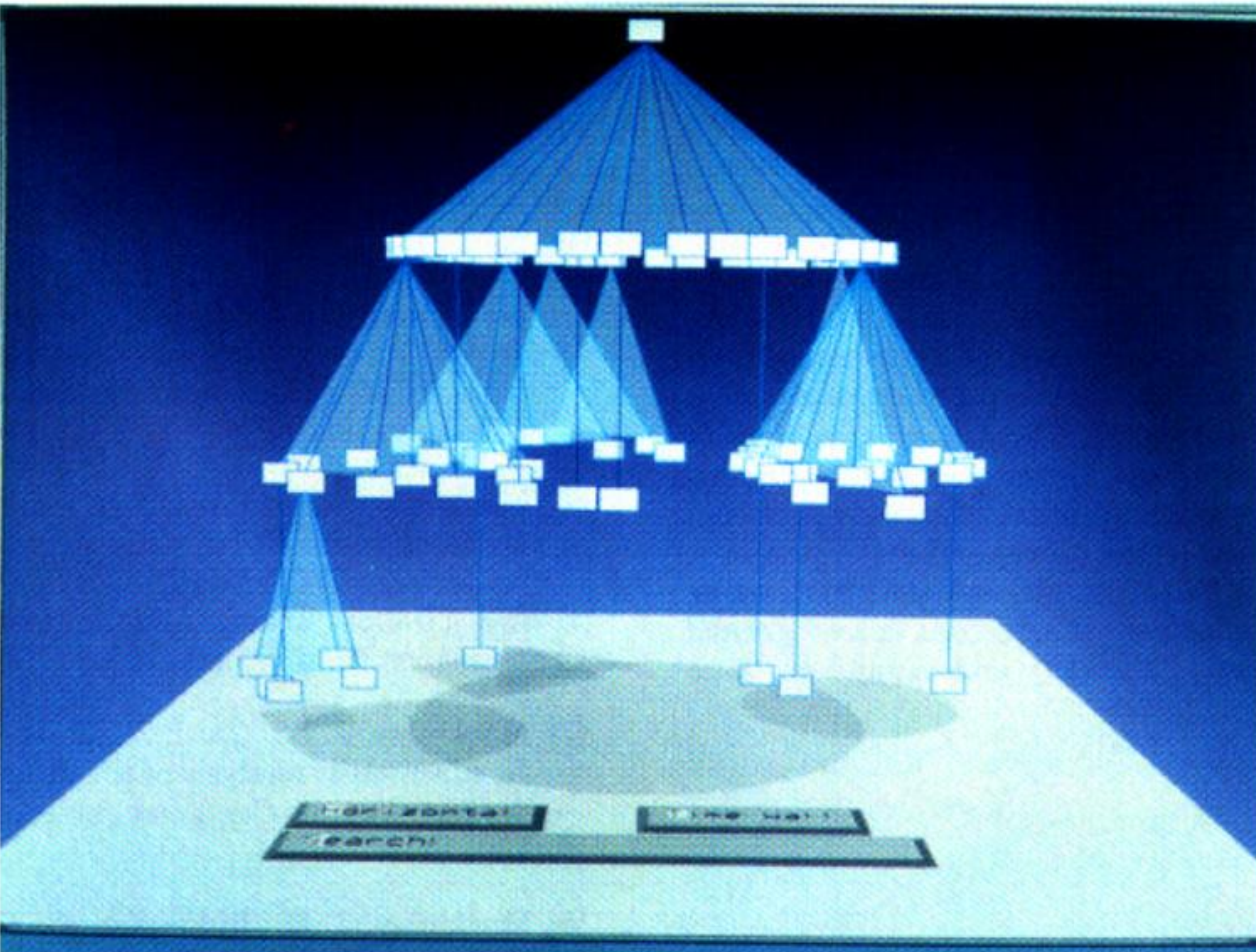
# Hyperbolic trees



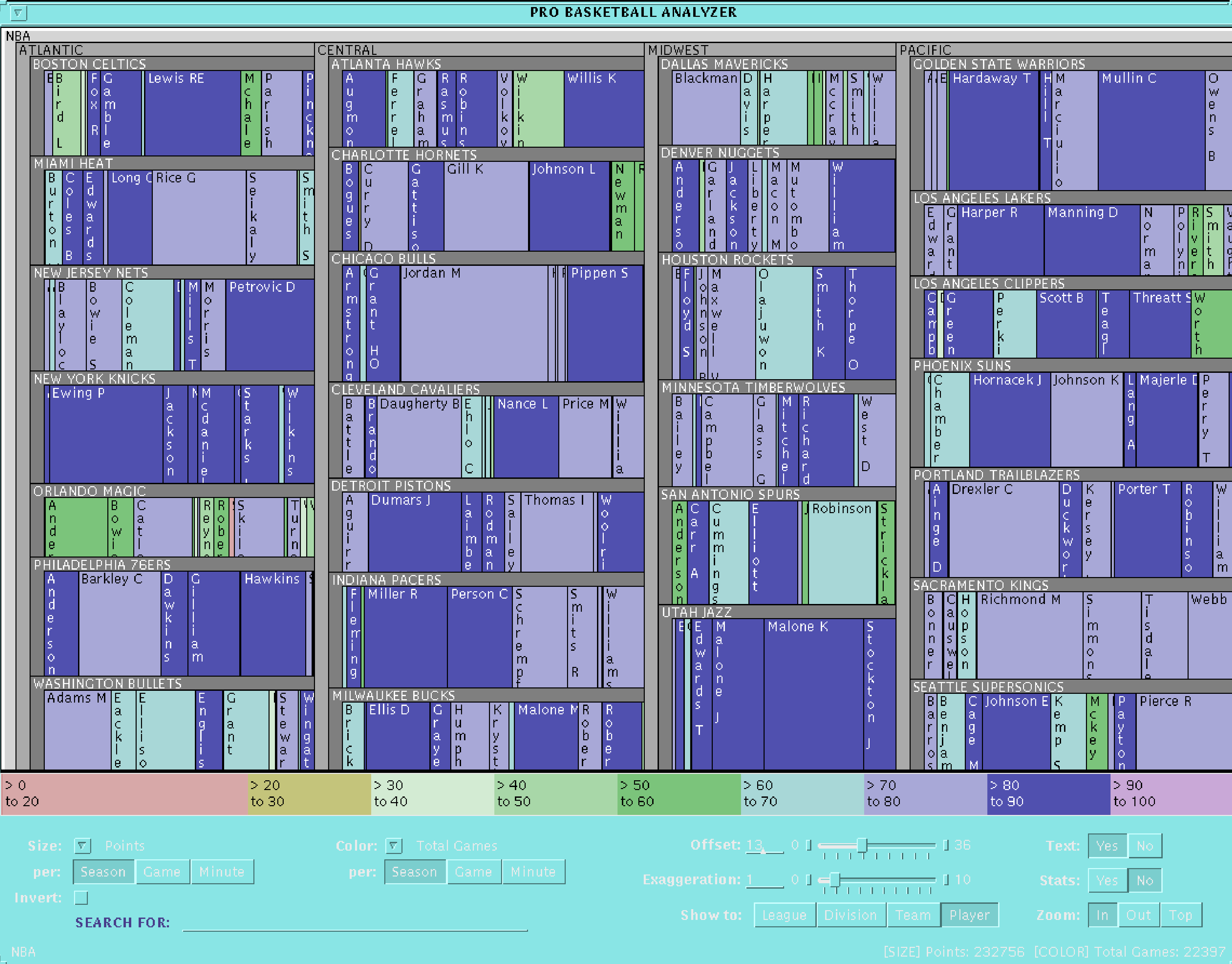
<https://glouwa.github.io/d3-hypertree-examples/demo/>



# Cone Trees



# Slice and Dice Treemaps



# Cluster / Squarified Treemaps

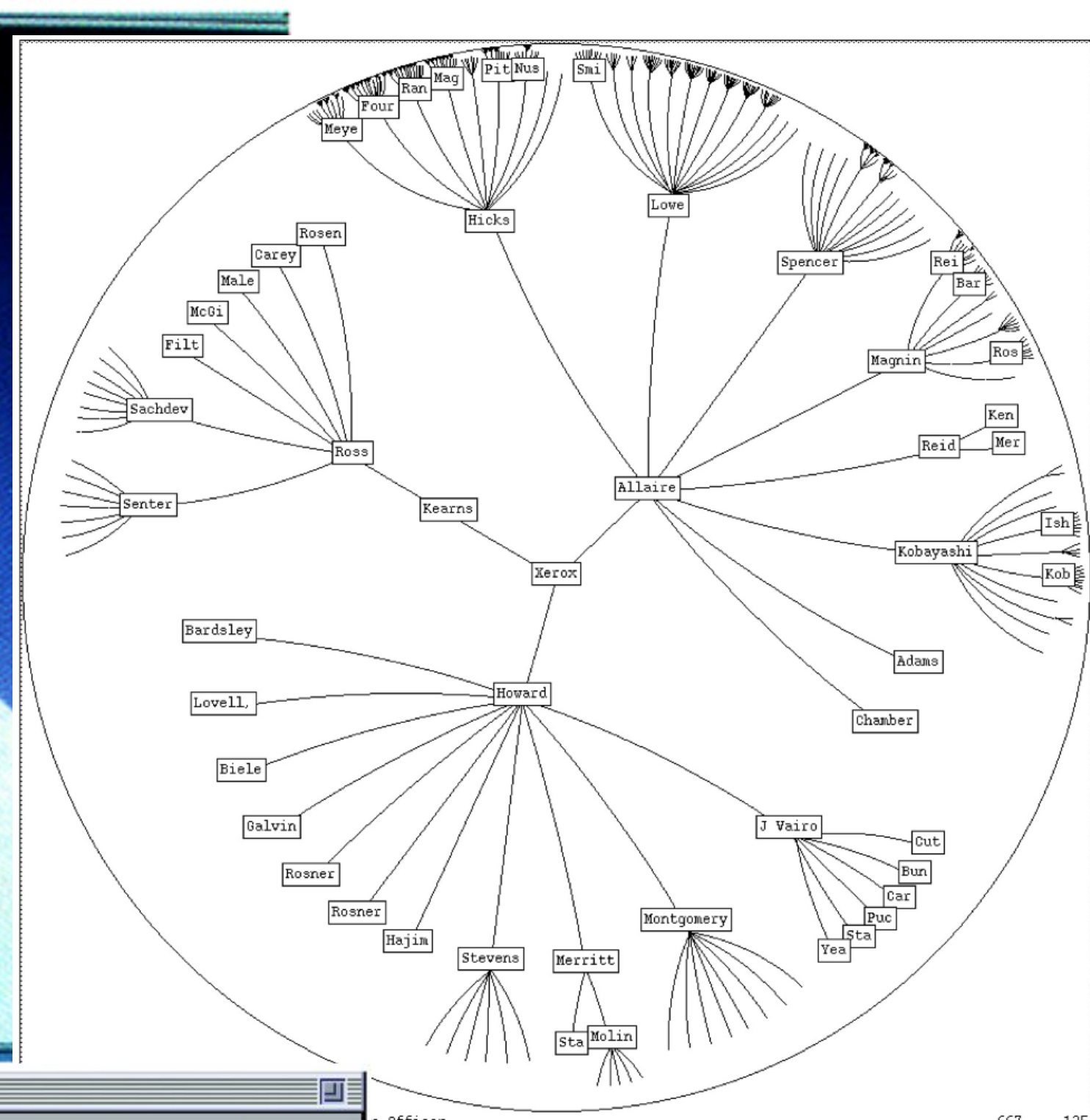
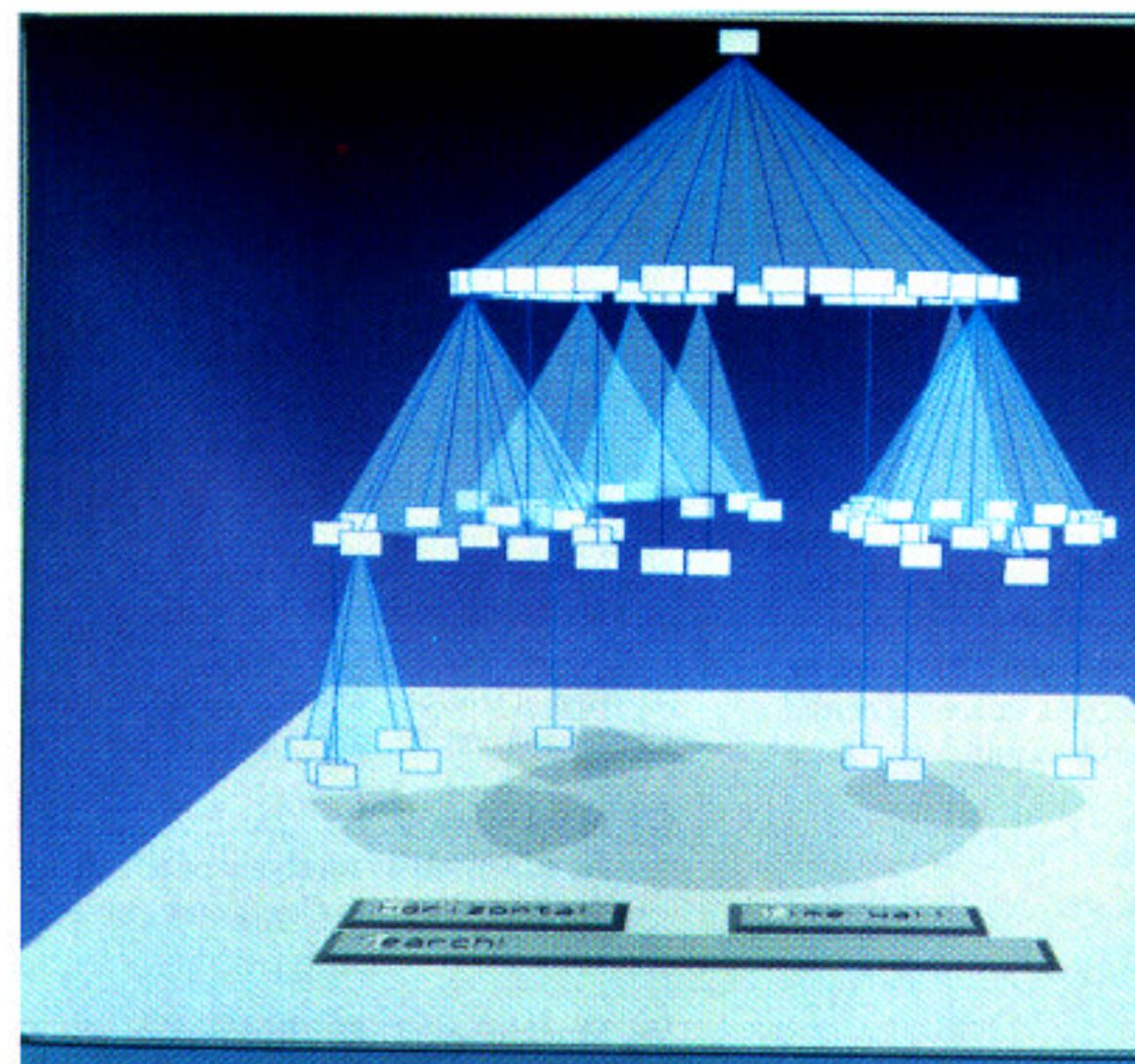
finviz

S&P 500 • 1 DAY PERFORMANCE • Thu MAR 19 2020 9:57 AM EST



[Wattenberg, 1999](#); [Bruls et al., 2000](#); [finviz live site](#); Snapshot: [finviz, 2020](#)





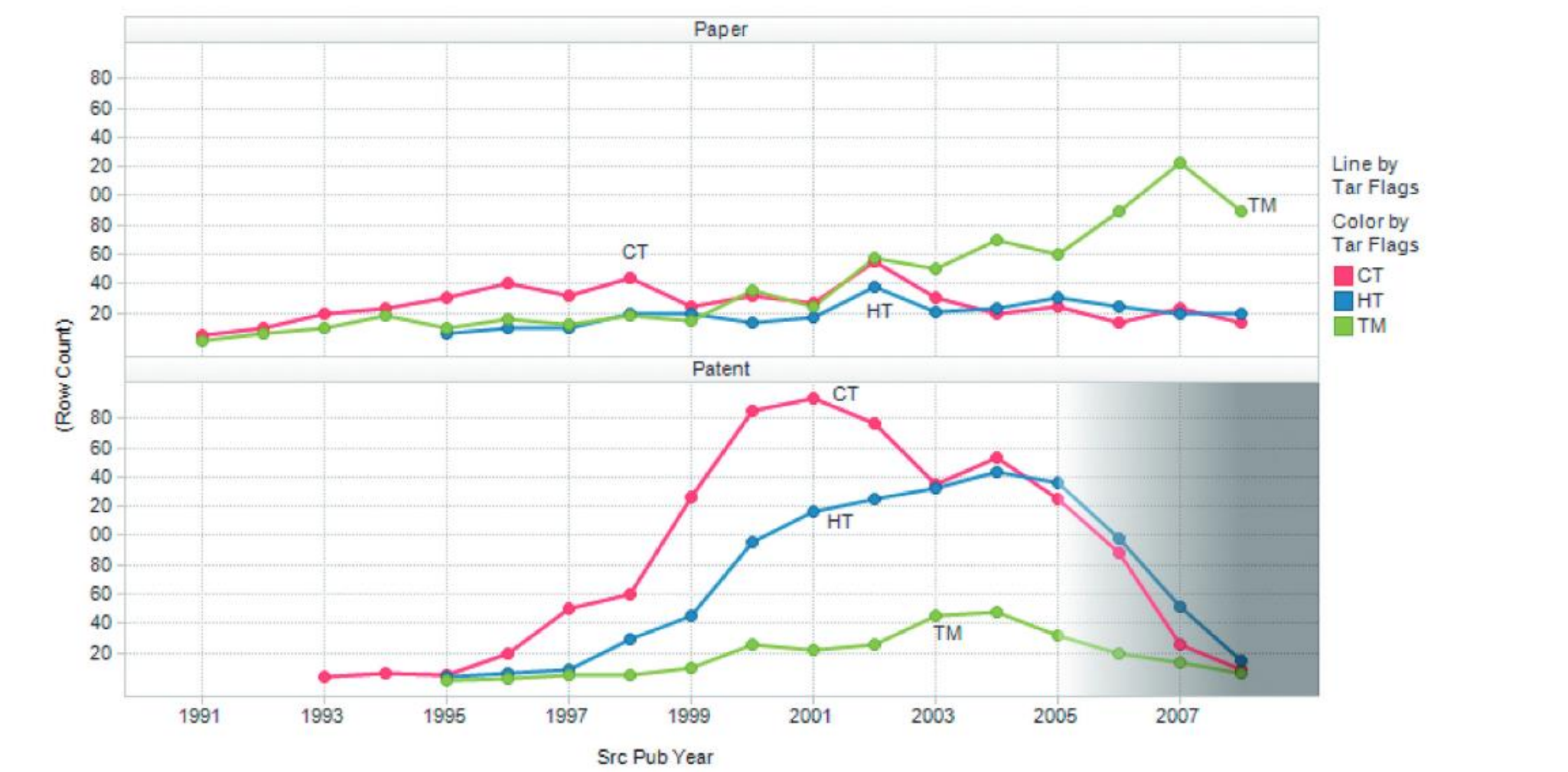
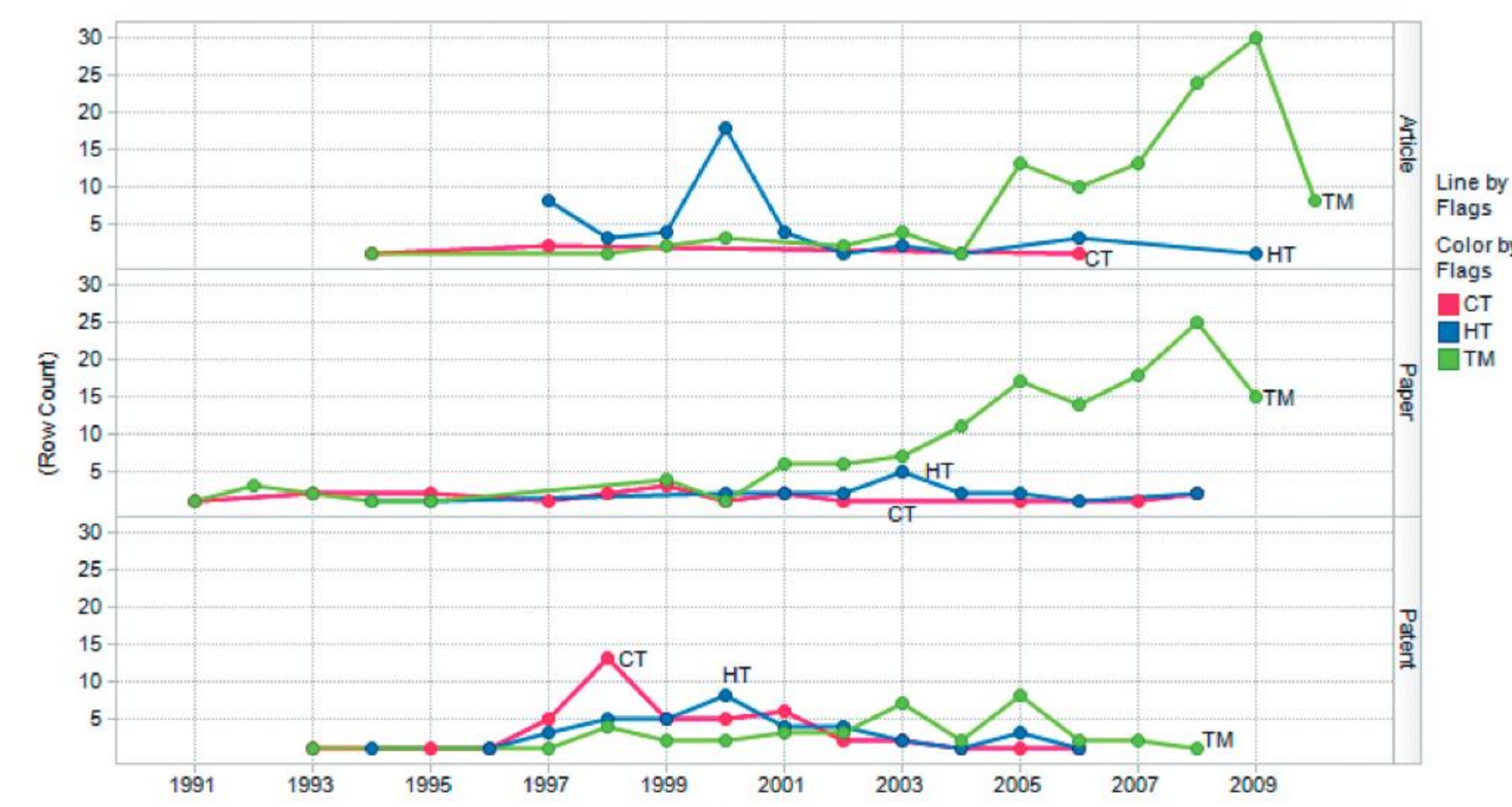
**-- History -- Treemap**

| 1991       | 1992                  | 1993                      | 1994                   |
|------------|-----------------------|---------------------------|------------------------|
| HCIL 90-91 | -cv 12/92             | -cv 2/93                  | AHP video2             |
| HCIL 90-91 | 91-92 ann rpt journa  | -UIS94-slides             | annual report letter   |
| Open house | annual report letter  | 92-93 ann rpt journalists | Bell Atlantic proposal |
| SWIFT rev  | CFAR Annual Report 9  | annual report letter      | King, UCI, ISR, Asahi  |
| tree-map 1 | HCIL 91-92 ann rpt le | NYU.slides                | Open house slides 94   |
| Treemaps 3 | Open house 92         | Open house 93             | VIS.chiformat          |
| vid-CHI'91 | Teittinen reference   | Open house slides 93      | DQtutorial             |
|            | letters typed by Lian | Letters typed by Lian     | Outline                |
|            |                       |                           | short summ             |

Outline 233 K 10% Mw2D MwII 7/20/94 7/22/94

-- History --:1994:DQtutorial:

Unknown Text Graphics Archives/Stacks Programming Applications System

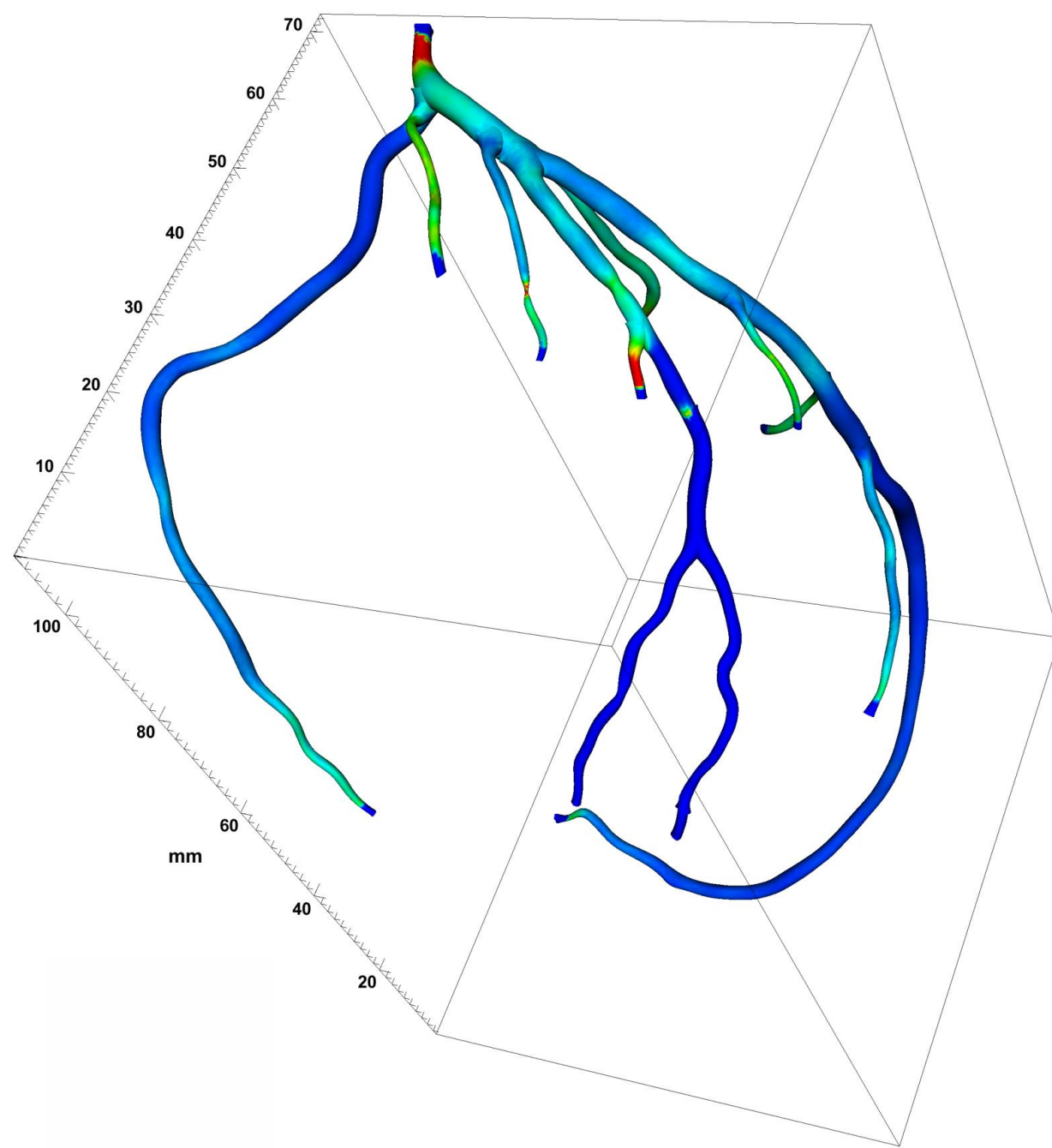


Previous  
Class

Rainbow:

3D: 39%

2D: 62%

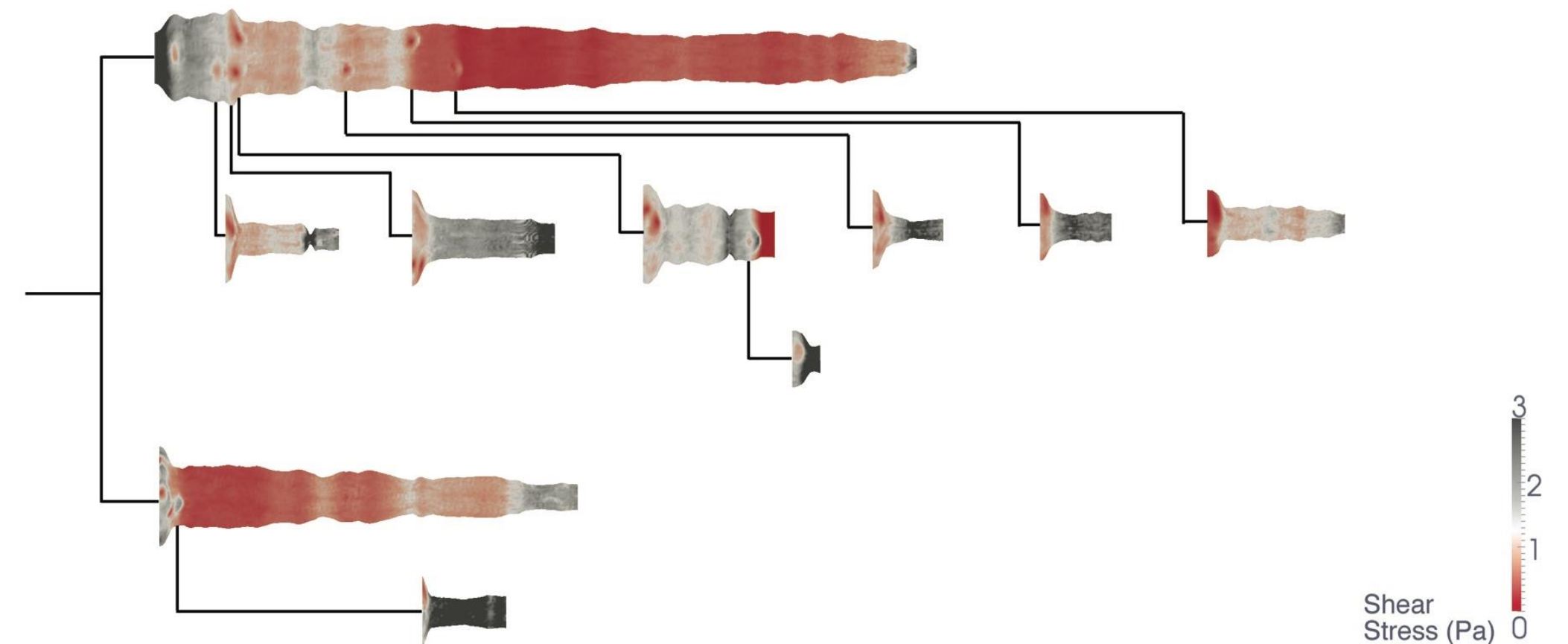


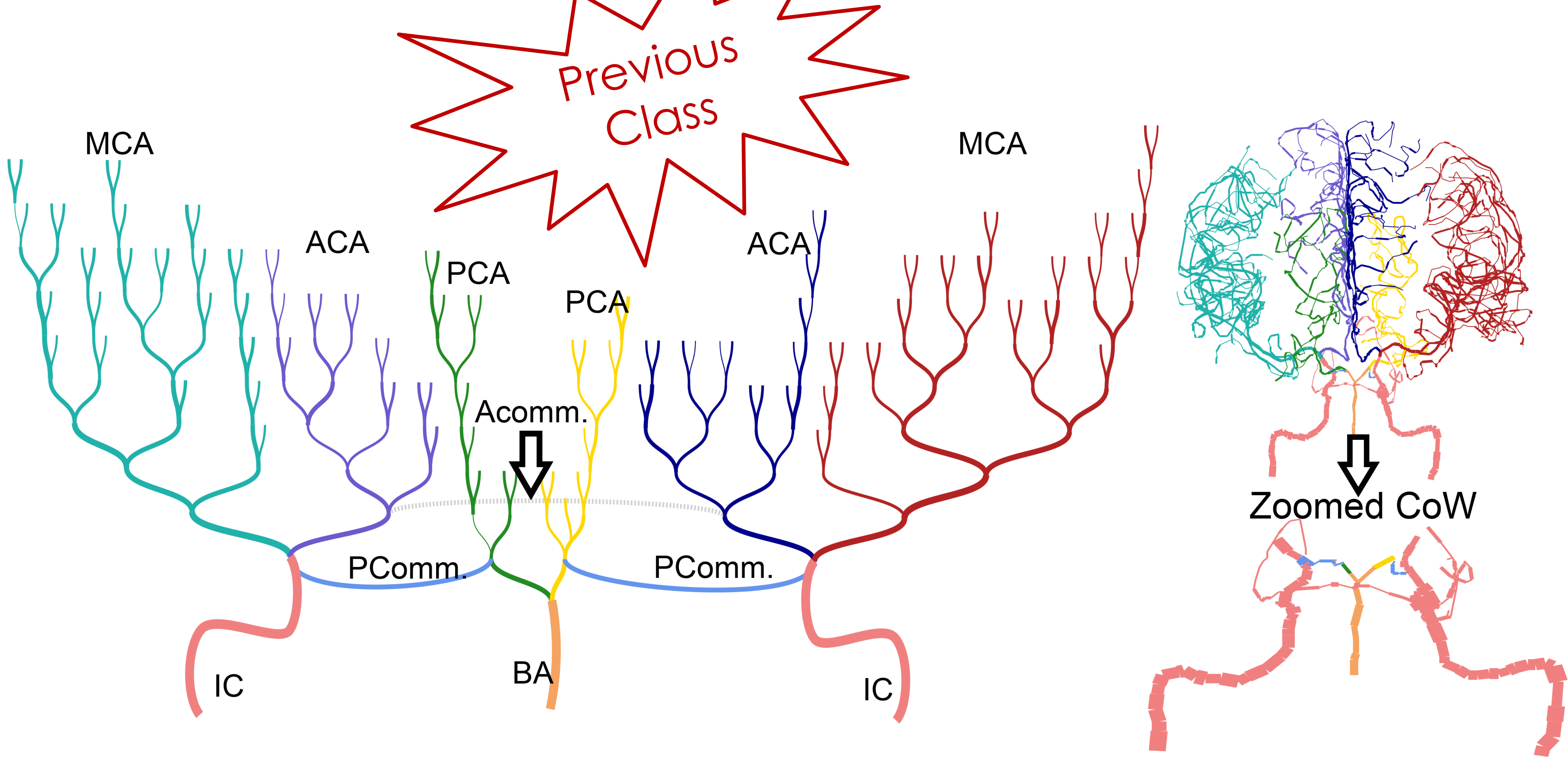
How many diseased  
regions found?

Diverging:

3D: 71% ( $\Delta$  +31%)

2D: 91% ( $\Delta$  +29%)



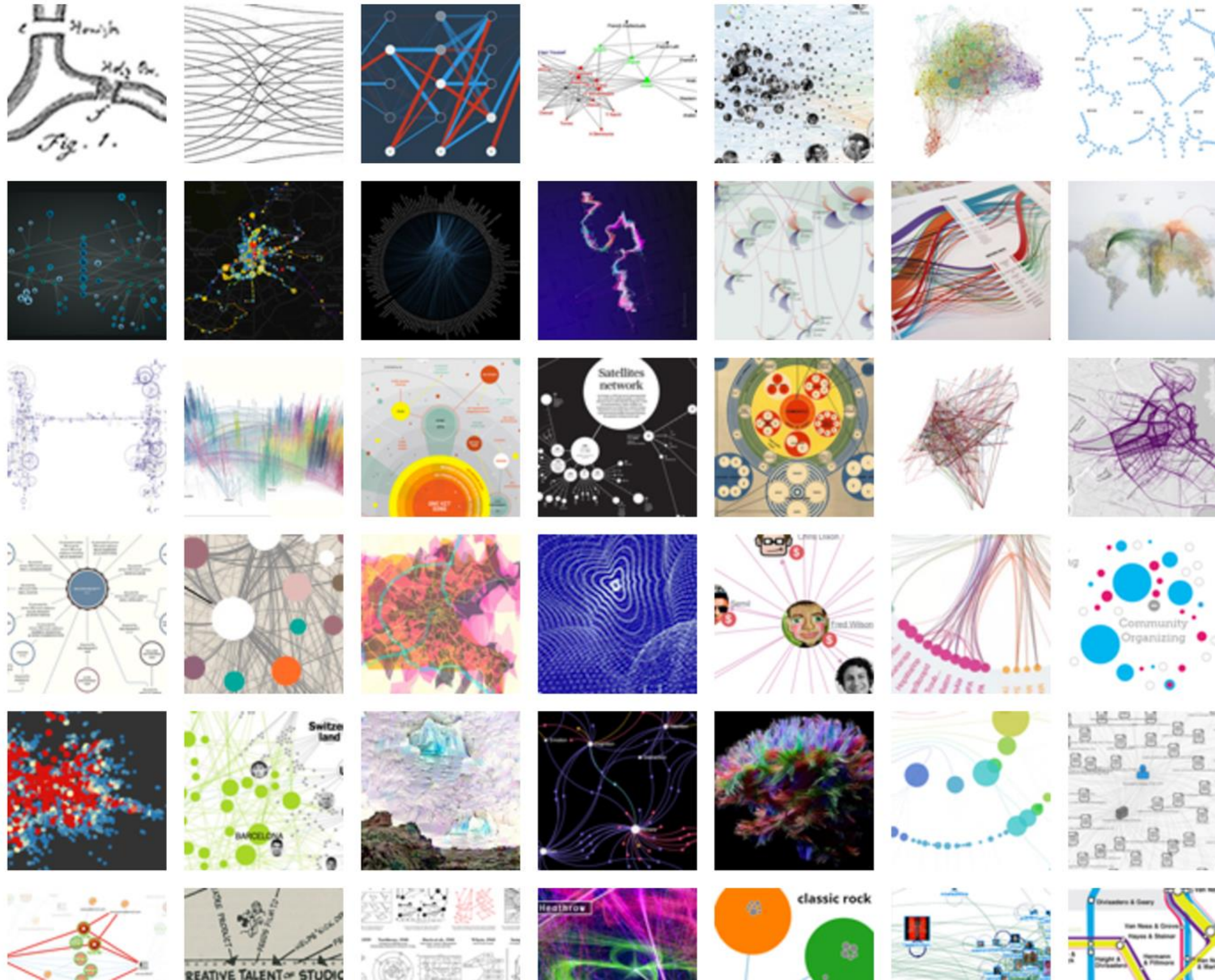




GO

### Latest Projects:

Indexing **1000** projects

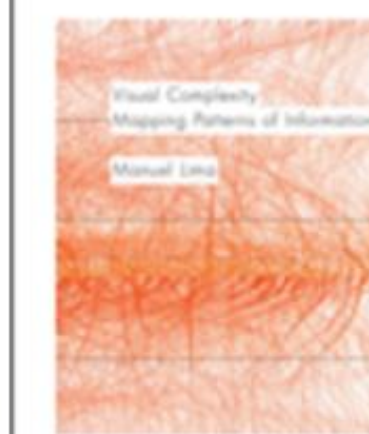


### Filter by:

SUBJECT

- Art (74)
- Biology (60)
- Business Networks (50)
- Computer Systems (39)
- Food Webs (16)
- Internet (35)
- Knowledge Networks (141)
- Multi-Domain Representation (70)
- Music (47)
- Others (77)
- Pattern Recognition (53)
- Political Networks (34)
- Semantic Networks (44)
- Social Networks (135)
- Transportation Networks (70)
- World Wide Web (55)

See All (1000)



visual complexity  
Mapping Patterns of Information

Buy now

Dimensionality



Representation



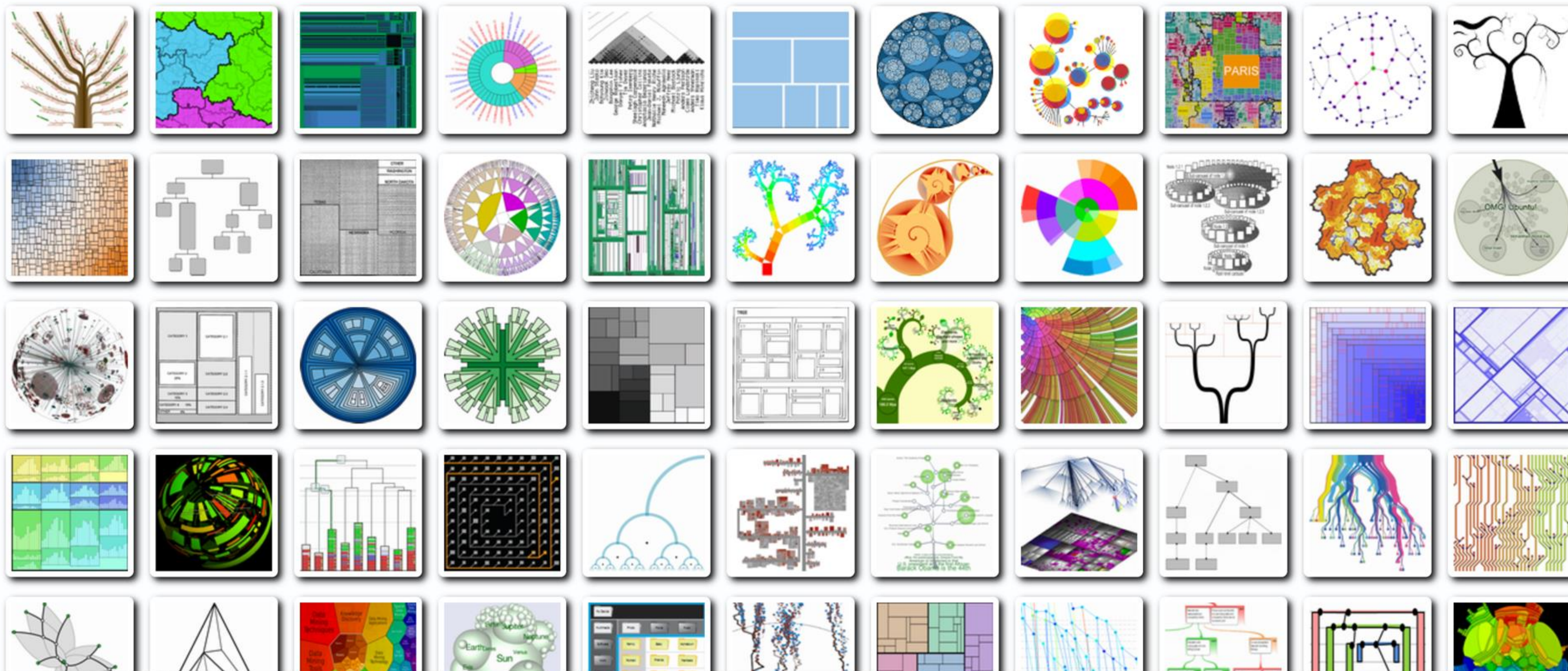
Alignment



Fulltext Search

Techniques Shown

292



# For Next Time

[neu-ds-4200-f23.github.io/schedule/](https://neu-ds-4200-f23.github.io/schedule/)

Look at the upcoming assignments and deadlines

- Textbook, Readings, & Reading Quizzes—Variable days
- In-Class Activities—If due, they are due 11:59pm the same day as class

Everyday Required Supplies:

- 5+ colors of pen or marker
- White paper
- Laptop and charger

Use Slack for general questions, email [codydunne-and-tas@ccs.neu.edu](mailto:codydunne-and-tas@ccs.neu.edu) for questions specific to you.

| Week 6: Networks and Trees; Spatial, 3D, and SciVis  |  |
|--|--|
| <b>Tue, Oct 10</b><br><i>Networks and Trees</i><br>Required Readings:<br><br>1 VAD Chapter 9—Arrange Networks and Trees  | <b>Fri, Oct 13</b><br><i>Spatial, 3D, and scientific visualization</i><br>Required Readings:<br><br>1 VAD Chapter 8—Arrange Spatial Data<br><br><b>A5—Altair interactive charts</b> due at 11:59pm |
| Week 7: Midterm  |  |
| <b>Tue, Oct 17</b><br><i>Midterm Q&amp;A and Study Session</i>   | <b>Fri, Oct 20</b><br><i>MIDTERM EXAM</i>  |
| Week 8: Storytelling, Validation   |  |
| <b>Tue, Oct 24</b><br><i>Storytelling, how to give a talk</i><br>Required Readings:<br><br>1 <b>Storytelling: The Next Step for Visualization by Robert Kosara and Jock Mackinlay (2013)</b> | <b>Fri, Oct 27</b><br><i>Validation and evaluation</i><br>Required Readings:<br><br>1 VAD Chapter 4—Analysis: Four Levels for Validation   |