



UNIVERSITÀ DEGLI STUDI DI PADOVA

Network Science

A.Y. 23/24

ICT for Internet & multimedia, Data science, Physics of data

Graph layout

Visualizing a graph



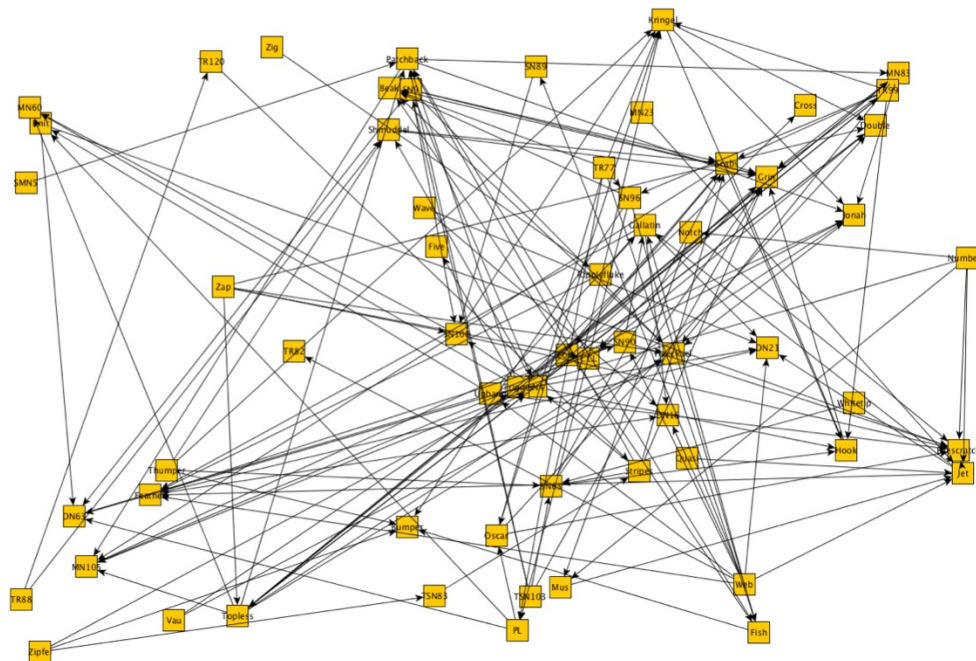
General layout problem

which aesthetic criteria would you optimize?

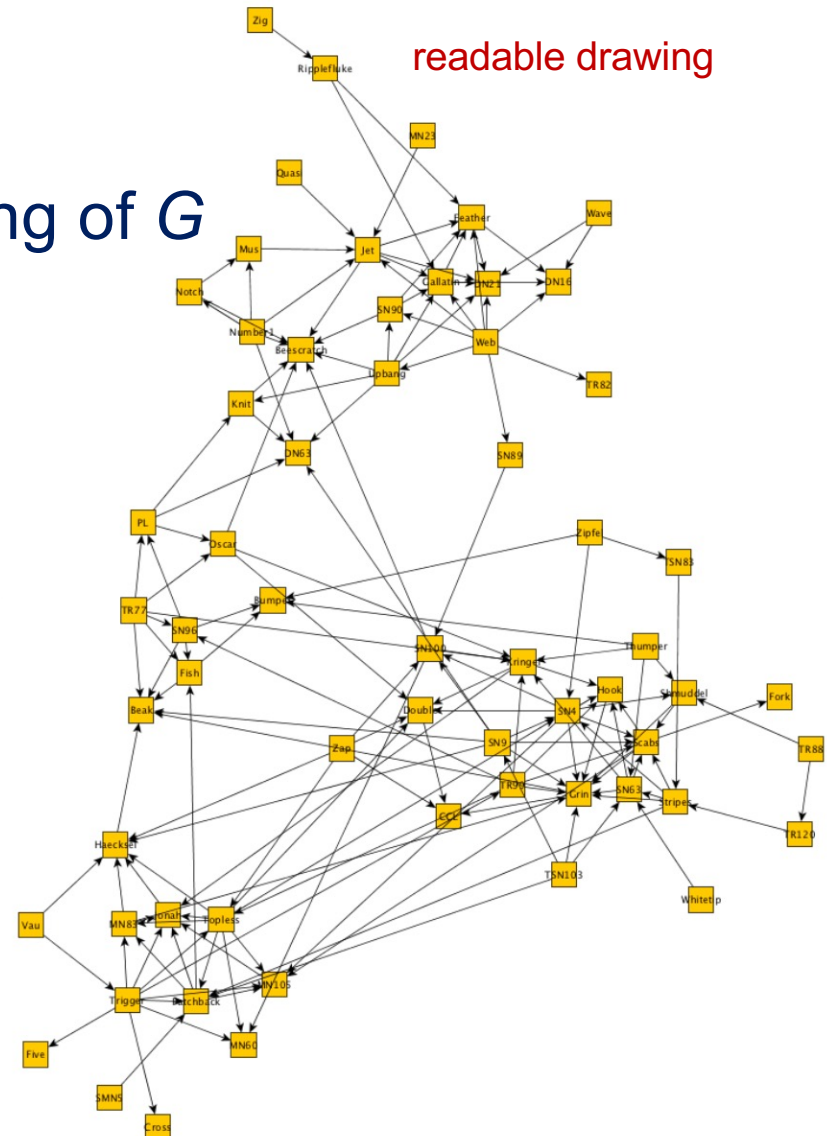
Problem:

Given a graph $G = (V, E)$

Find a clear and readable drawing of G



messy drawing



readable drawing



- ☐ adjacent nodes are close
- ☐ non-adjacent far apart
- ☐ edges length proportional to their weight
- ☐ densely connected parts (clusters) to form communities
- ☐ as few crossings as possible
- ☐ nodes distributed evenly



... but optimization criteria partially contradict each other



Before

- ❑ always based on some properties: tree, series-parallel graph, planar graph
- ❑ and on some additional information: ordering of the vertices, decompositions into SP-components
- ❑ **NP-hard** even in simple scenarios
 - edge lengths $\{1, 2\}$ (Saxe, '80)
 - planar drawing with unit edge lengths (Eades, Wormald, '90)

Today

- ❑ more direct and intuitive method based on physical analogies : **force directed algorithms**
- ❑ the methods are very popular: intuitiveness, easy to program, generality, fairly satisfactory results,...

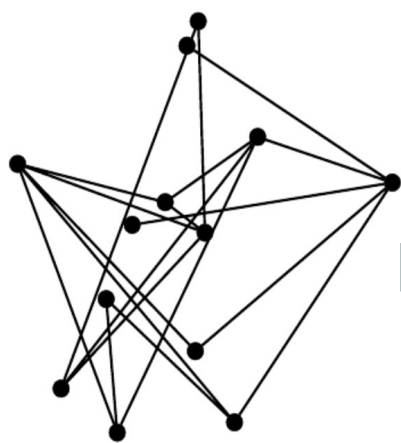
Force directed layout

a physical analogy for graphs

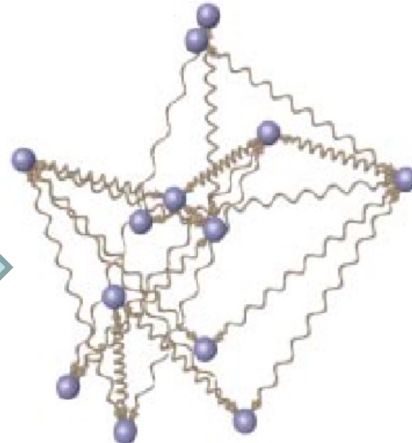
Springer embedder algorithm

Eades, "A heuristic for graph drawing" (1984)

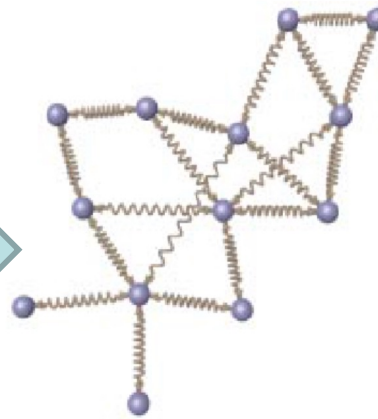
“To embed a graph we replace the vertices by steel rings and replace each edge with a spring to form a mechanical system . . . The vertices are placed in some initial layout and let go so that the spring forces on the rings move the system to a minimal energy state.”



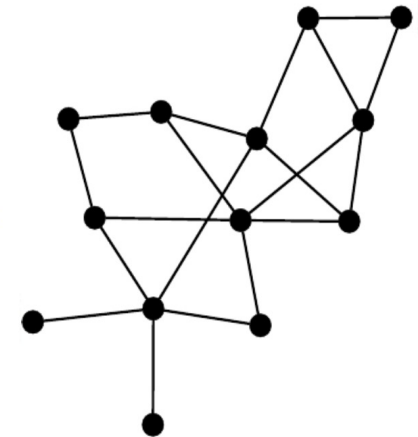
messy graph



convert it into a
mechanical system,
and let it run



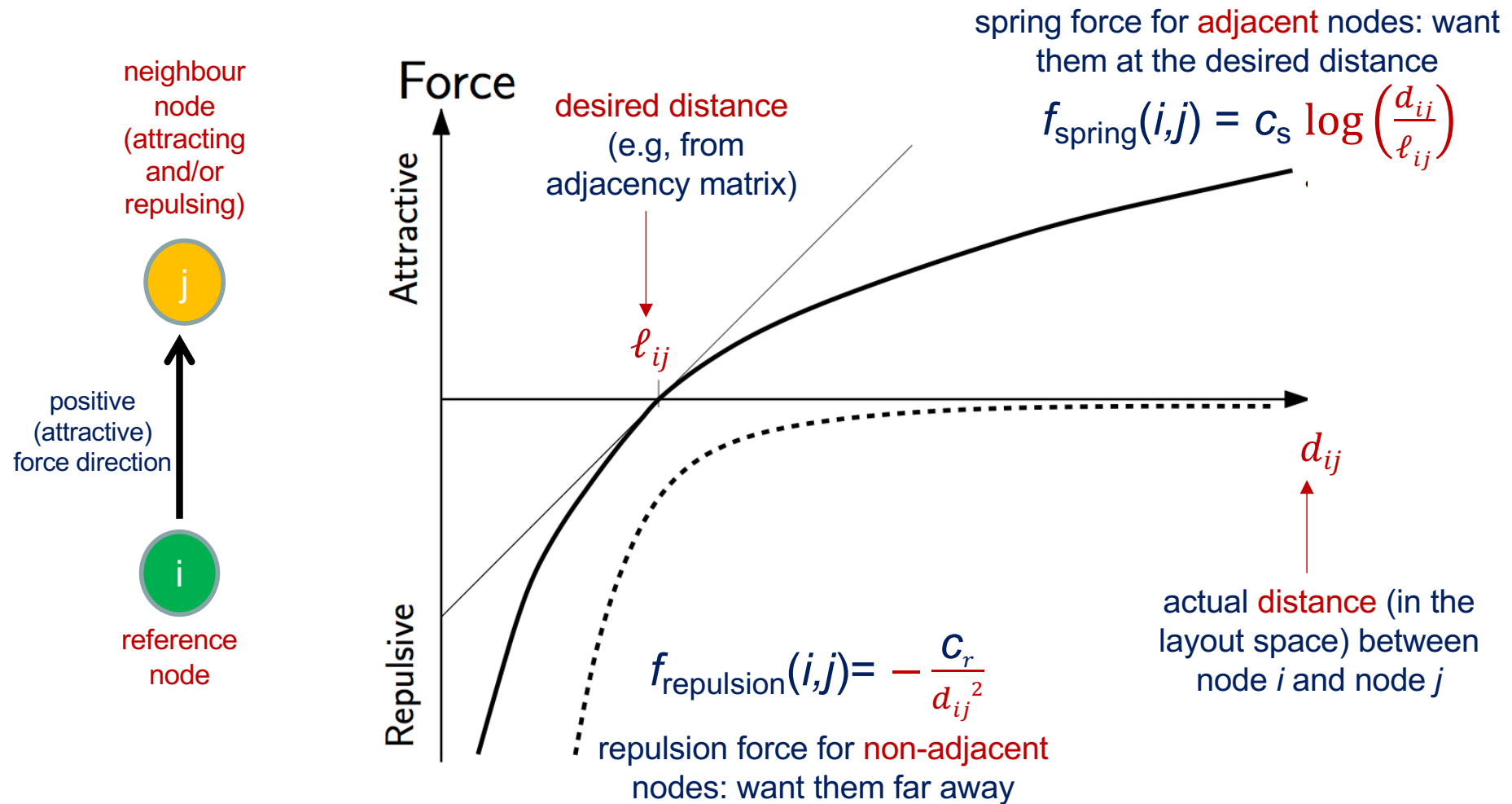
at steady state
forces are balanced
and nodes evenly
distributed



readable graph



Repulsive and attractive forces in a spring embedder





The algorithm

iteratively reaching the steady state

- Evaluate the **force** contribution on the i th node

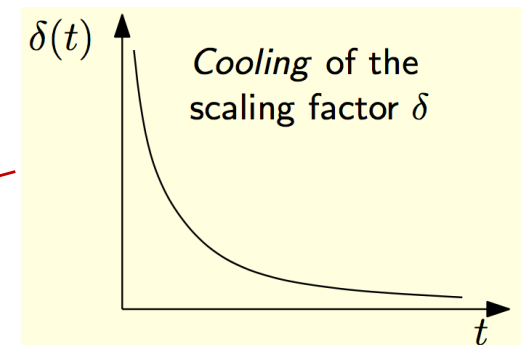
$$\begin{aligned} \mathbf{F}_i &= \sum_{j \in N_i} f_{\text{spring}}(i,j) (\mathbf{p}_j - \mathbf{p}_i) + \sum_{j \notin N_i} f_{\text{repulsion}}(i,j) \cdot (\mathbf{p}_j - \mathbf{p}_i) \\ &= c_s \sum_{j \in N_i} (\mathbf{p}_j - \mathbf{p}_i) \log \left(\frac{\|\mathbf{p}_j - \mathbf{p}_i\|}{\ell_{ij}} \right) - c_r \sum_{j \notin N_i} \frac{\mathbf{p}_j - \mathbf{p}_i}{\|\mathbf{p}_j - \mathbf{p}_i\|^2} \end{aligned}$$

position (in the layout space) of node i

direction from i to j

- **Update** the nodes position by applying forces

$$\mathbf{p}_i^+ = \mathbf{p}_i + \delta \mathbf{F}_i$$



- **Iterate** until the forces are small enough: $\max_i \|\mathbf{F}_i\| < \varepsilon$



Advantages

- ❑ very simple algorithm
- ❑ good results for small and medium-sized graphs
- ❑ good representation of symmetry/structure

Disadvantages

- ❑ system is **not stable** at the end
- ❑ converging to **local minima**
- ❑ **not scalable** - complexity is $O(N^2)$

Influence

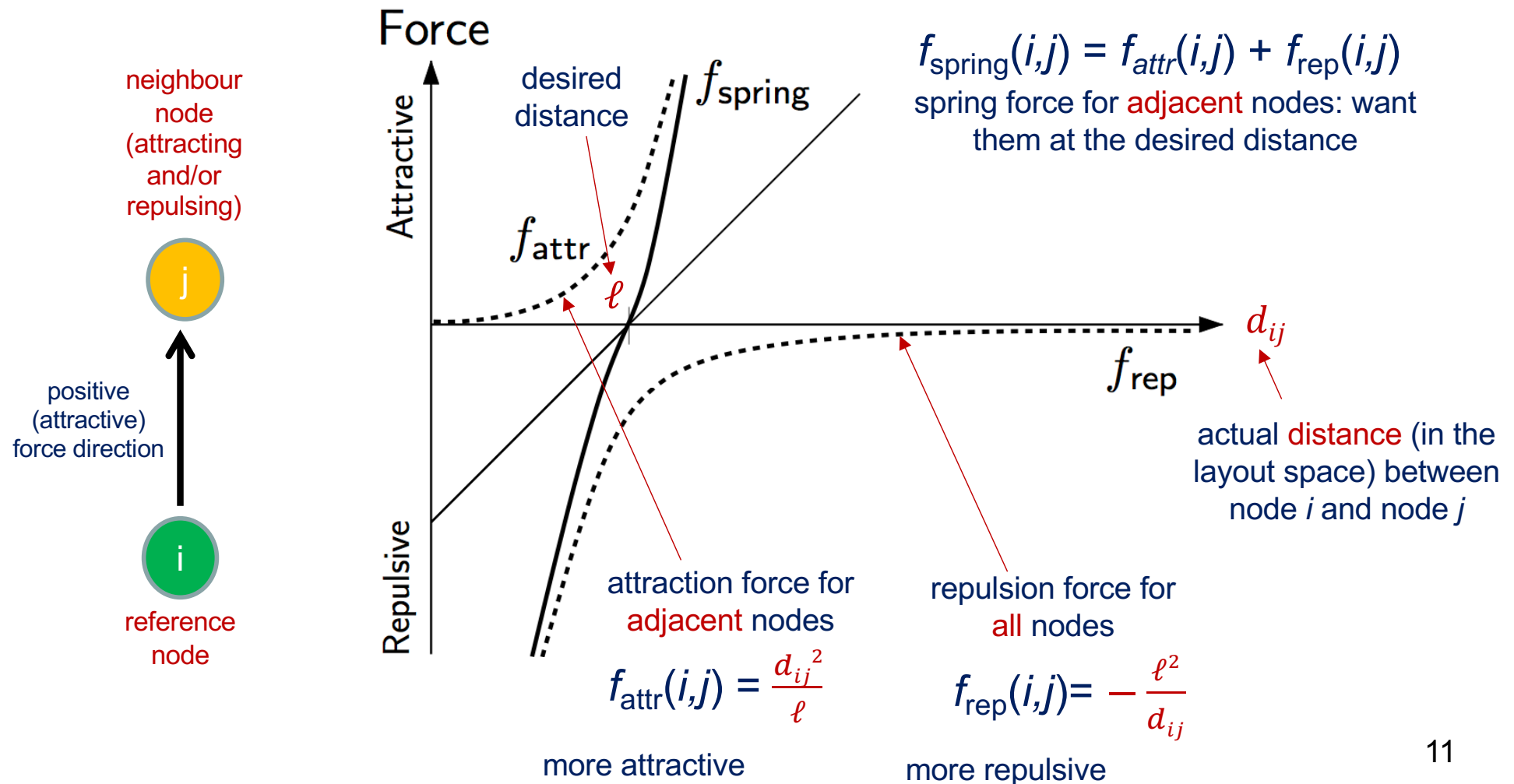
- ❑ basis for many further ideas



Fruchterman and Reingold

Fruchterman & Reingold, Graph drawing by
force-directed placement (1991)

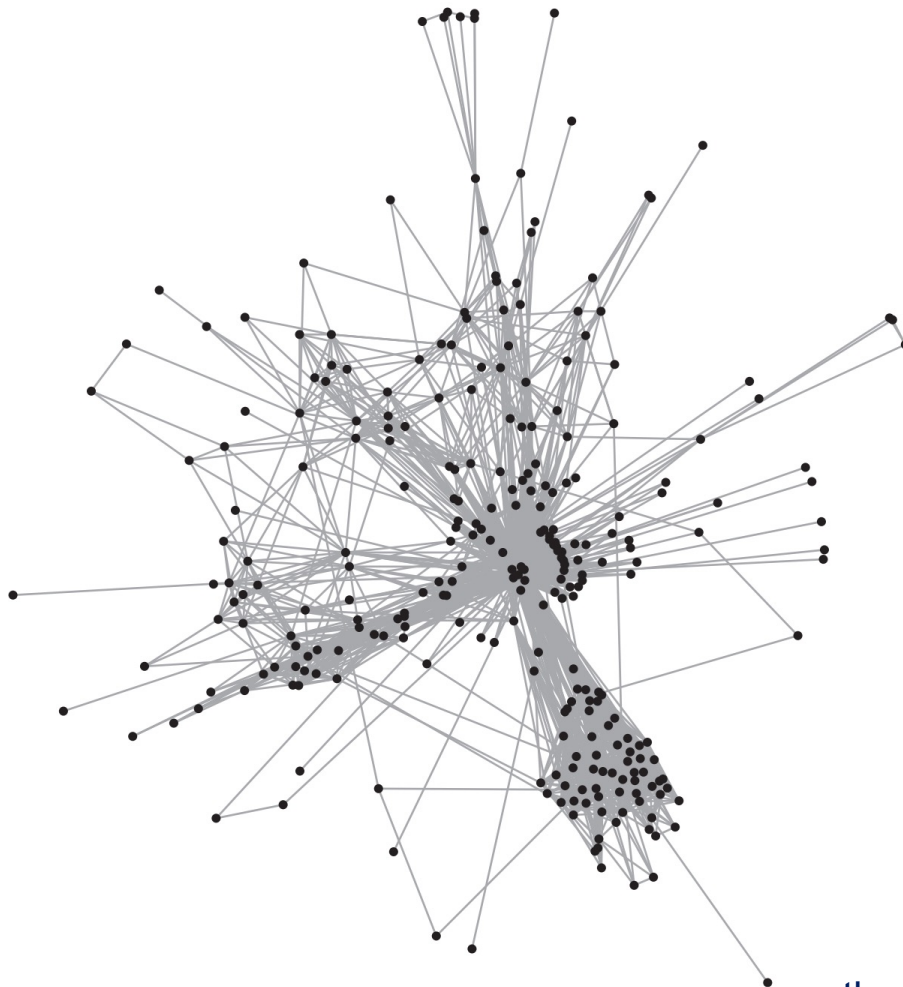
http://www.mathe2.uni-bayreuth.de/axel/papers/reingold:graph_drawing_by_force_directed_placement.pdf



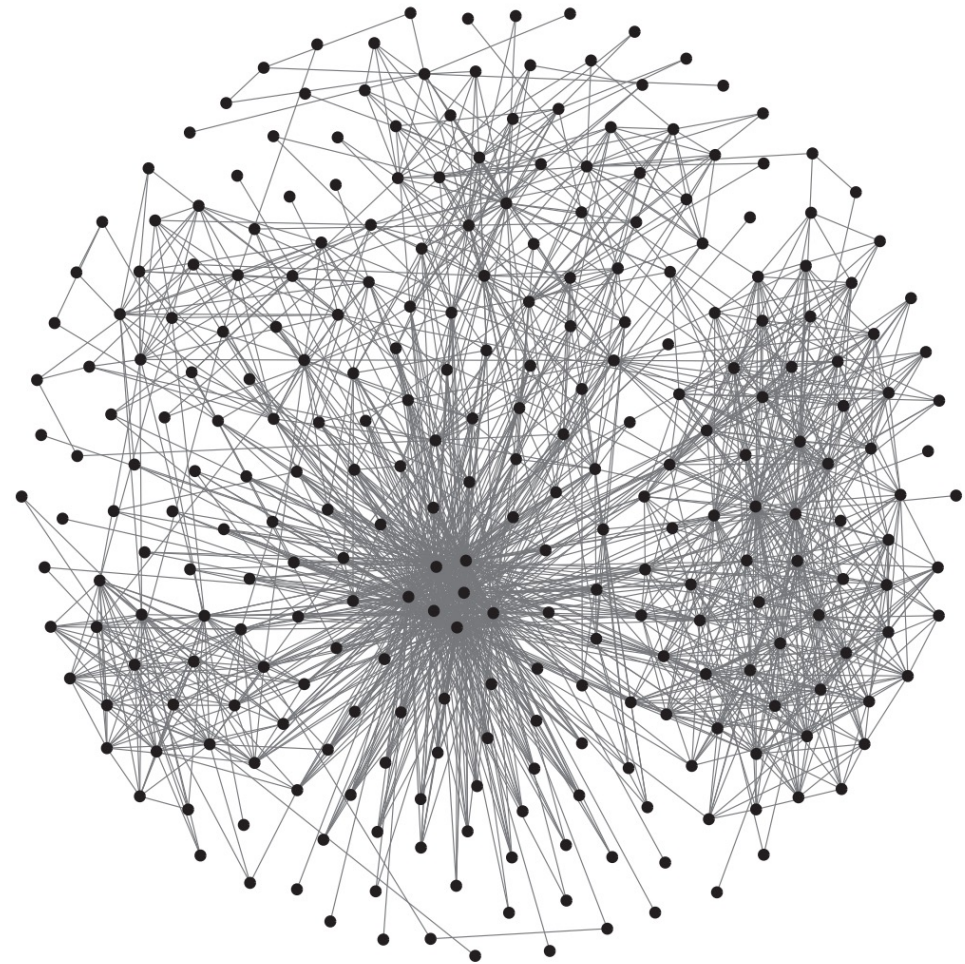


A visual example protein interaction network

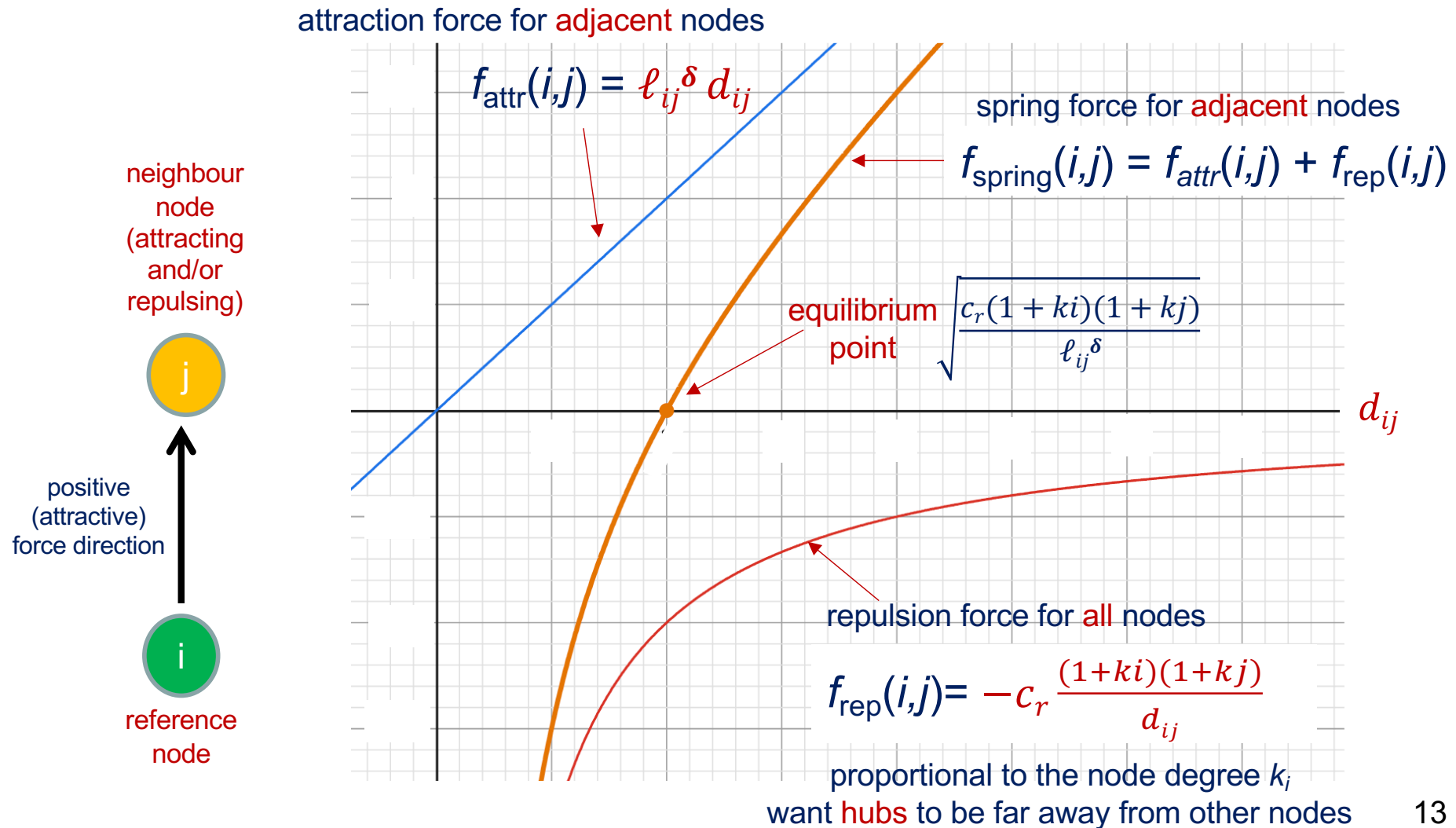
spring embedder



Fruchterman-Reingold



there exists a relation between Fruchterman-Reingold
layout and the communities found by modularity





Alternative modes

for attractive and repulsive forces

attraction force for
adjacent nodes

$$f_{\text{attr}}(i,j) = \begin{cases} \ell_{ij}^{\delta} d_{ij} \\ \frac{\ell_{ij}^{\delta} d_{ij}}{1+ki} \\ \ell_{ij}^{\delta} \log(1 + d_{ij}) \end{cases}$$

dissuade hubs

linlog mode

less attraction for
authorities (will have
more space)

weaker
dependence
on distance

repulsion force for
all nodes

$$f_{\text{rep}}(i,j) = -c_r \frac{(1+ki)(1+kj)}{d_{ij}}$$

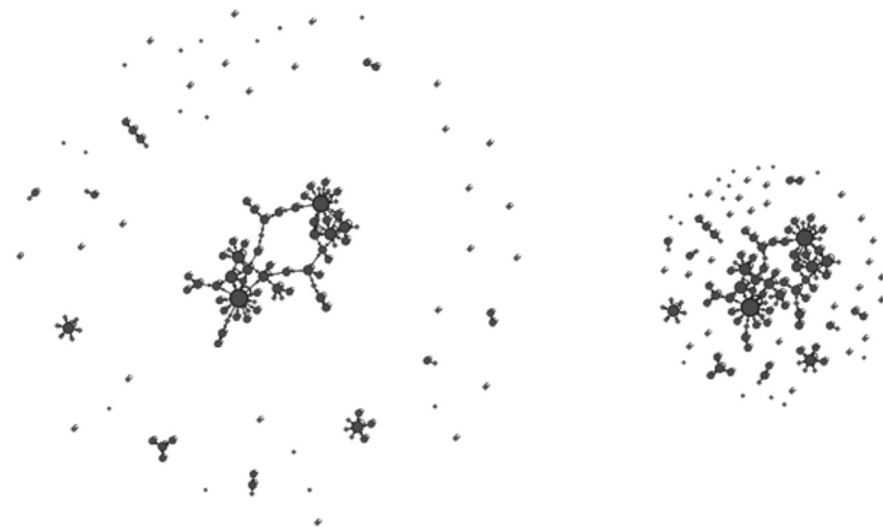
resized distance taking into
account node sizes

Prevent overlapping mode $d'_{ij} = [d_{ij} - \text{size}_i - \text{size}_j]^+$

$$d'_{ij} = 0 \quad \Rightarrow \quad \begin{aligned} f_{\text{attr}}(i,j) &= 0 \\ f_{\text{rep}}(i,j) &= -c'_r(1 + ki)(1 + kj) \end{aligned}$$



attracts nodes to the centre of the spatialisation. Its main purpose is to compensate repulsion for nodes that are far away from the centre



gravity force for
all nodes

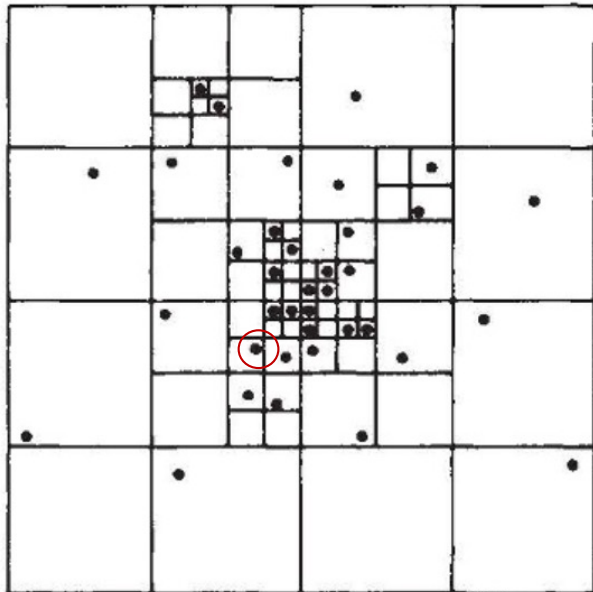
$$f_{\text{gravity}}(i) = \begin{cases} c_g (1 + ki) \\ c_g (1 + ki) k_i \end{cases} \quad \text{strong gravity}$$

applied towards the baricenter $\mathbf{p}_{\text{bary}} = \frac{1}{N} \sum_i \mathbf{p}_i$

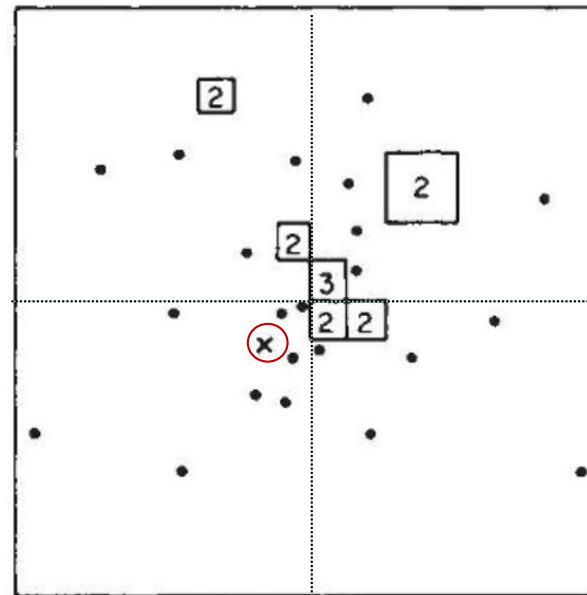


an heuristic to circumvent the $O(N^2)$ complexity of calculating
repulsion forces (2D example)

binary partition the space,
minimum partition to isolate nodes



nodes far away from x are
condensed in a single entity
located at the baricenter

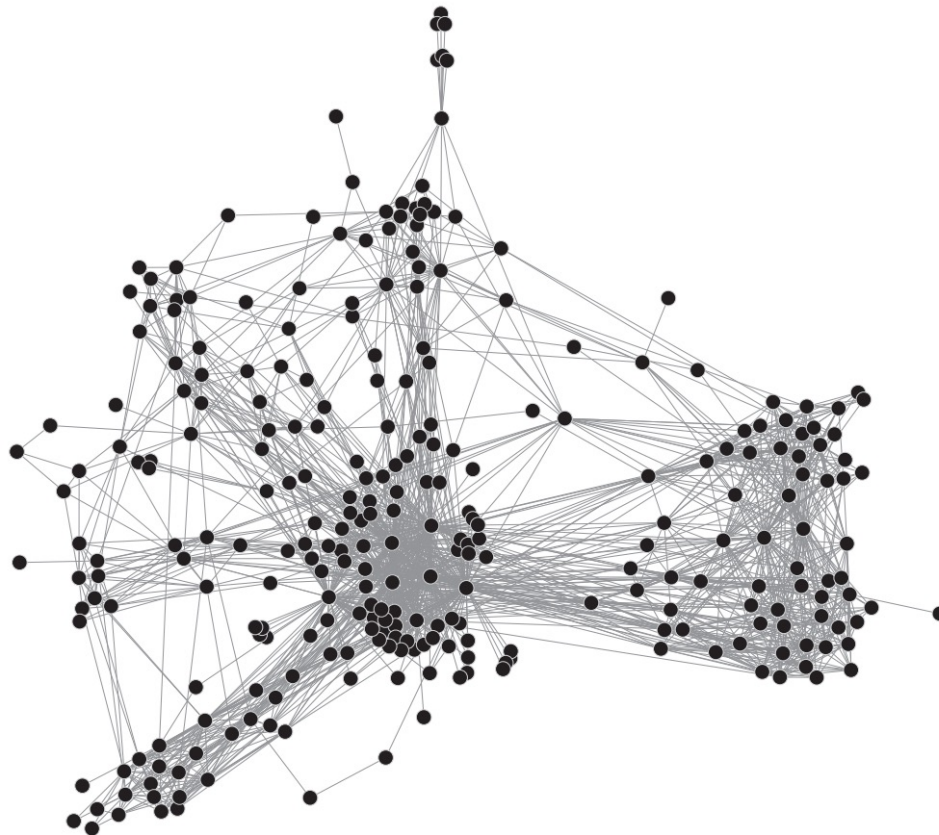


$$\begin{aligned}
 & \text{baricenter} \\
 p_b &= \frac{1}{C} \sum_{j \in C} p_j \\
 & \text{approximate force} \\
 f_{ic} &= \sum_{j \in C} f_{ij} \cdot (p_j - p_i) \\
 & \cong \sum_{j \in C} f_{ib} \cdot (p_j - p_i) \\
 & = f_{ib} C (p_b - p_i)
 \end{aligned}$$



A visual example protein interaction network

Force Atlas 2



more clearly separate communities,
compact layout

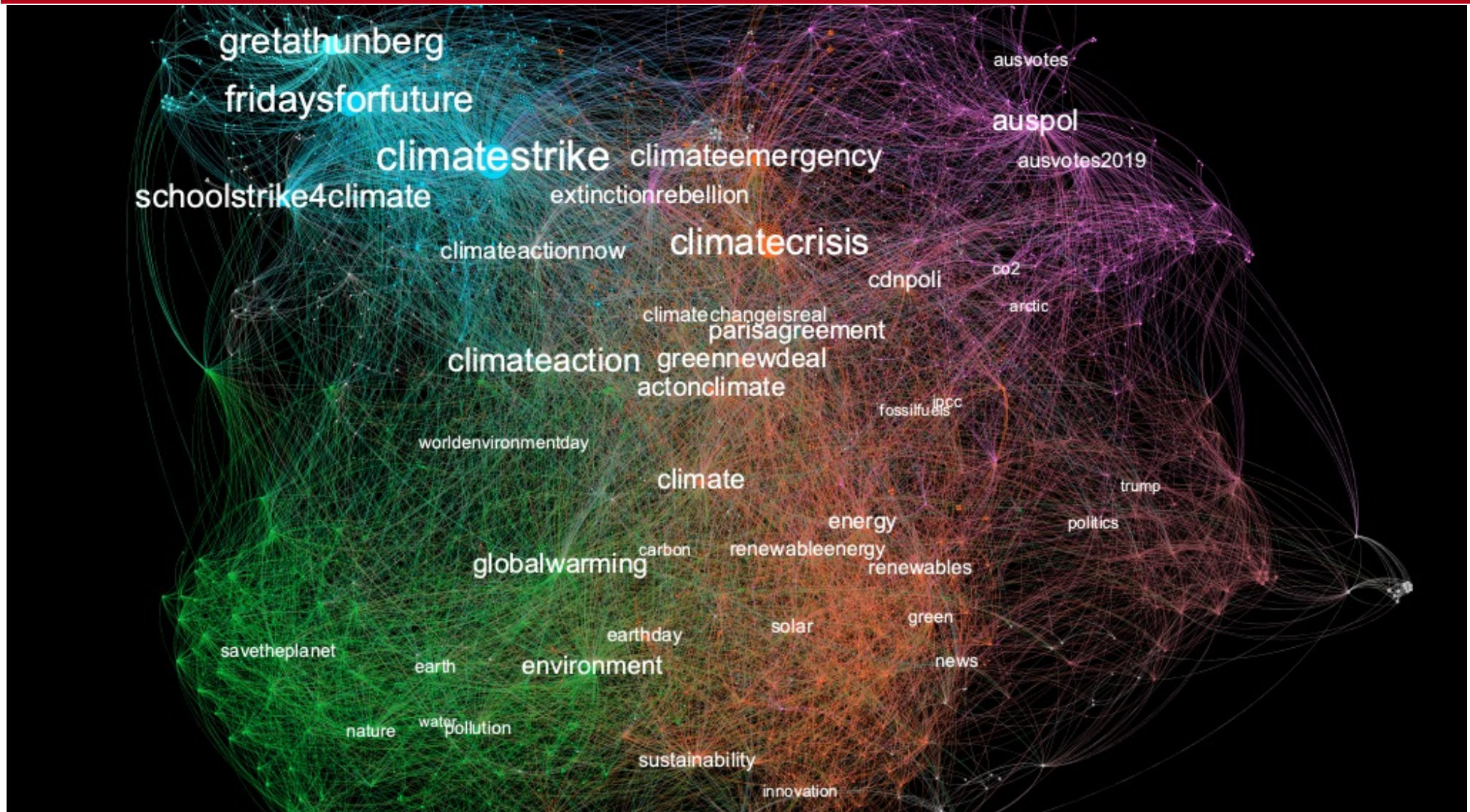
Force Atlas 2 – linlog mode



much greater spacing with linlog mode



A visual example semantic network on #climateaction



but to get such a nice output you also need manual intervention !!!



attraction force for **adjacent** nodes
(k nearest ones only)

$$f_{\text{attr}}(i,j) = \frac{a d_{ij}^{2(b-1)}}{1+d_{ij}^2} w_{ij}$$

active above a minimum distance

$$f_{\text{spring}}(i,j) = f_{\text{attr}}(i,j) + f_{\text{rep}}(i,j)$$

spring force for **adjacent** nodes
(k nearest ones only)

equilibrium point

d_{ij}

repulsion force for **sampled** nodes

this controls complexity

$$f_{\text{rep}}(i,j) = - \frac{(1-w_{ij})}{(\epsilon+d_{ij}^2)(1+a d_{ij}^{2b})}$$

zero above a minimum distance

neighbour node
(attracting and/or repulsing)



positive
(attractive)
force direction



reference node

the link weight w_{ij}
is derived heuristically from
desired distances ℓ_{ij}



assume a **desired link distance** ℓ_{ij} is available;
set $\ell_{ij} = \infty$ for nodes that are not connected

← can also be
extracted from
an adjacency
matrix

link weight model

$$w_{ij} = \exp\left(-\frac{\ell_{ij} - \ell_{i \min}}{\sigma_i}\right)$$

← $\ell_{i \min} = \min_j \ell_{ij}$

← identify σ_i by solving

$$\sum_{j \in K_i} \exp\left(-\frac{\ell_{ij} - \ell_{i \min}}{\sigma_i}\right) = \log_2(k)$$

k nearest neighbours to node i

hyperparameters model

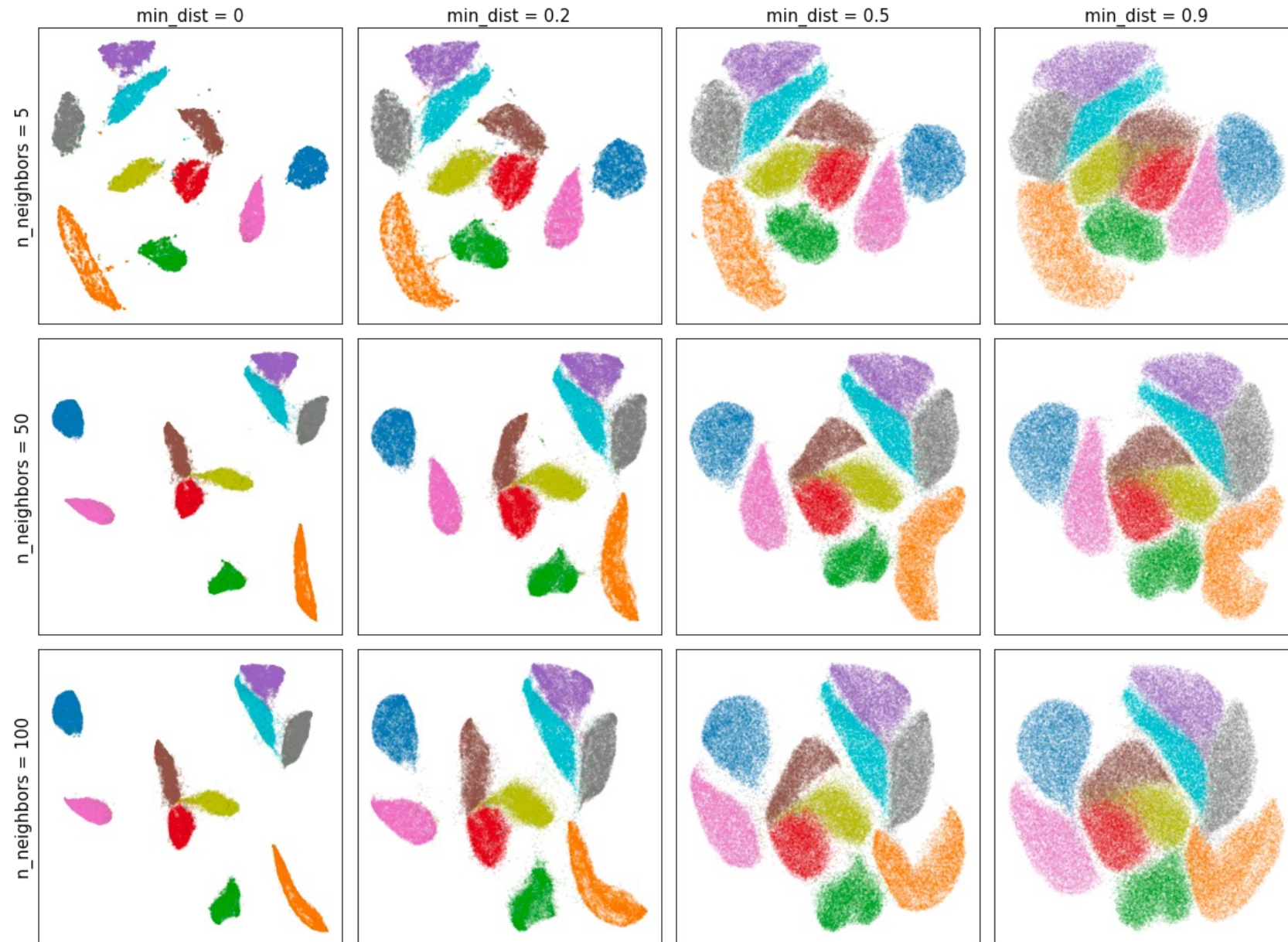
$$(a, b) = \operatorname{argmin}_{a, b} \sum_{i, j} \left(\frac{1}{1 + a \, dij^{2b}} - e^{-[d_{ij} - d_{\min}]^+} \right)^2$$

← non-linear
fitting to a
smooth
function

minimum distance
parameter



Parameters selection on the role of the minimum distance



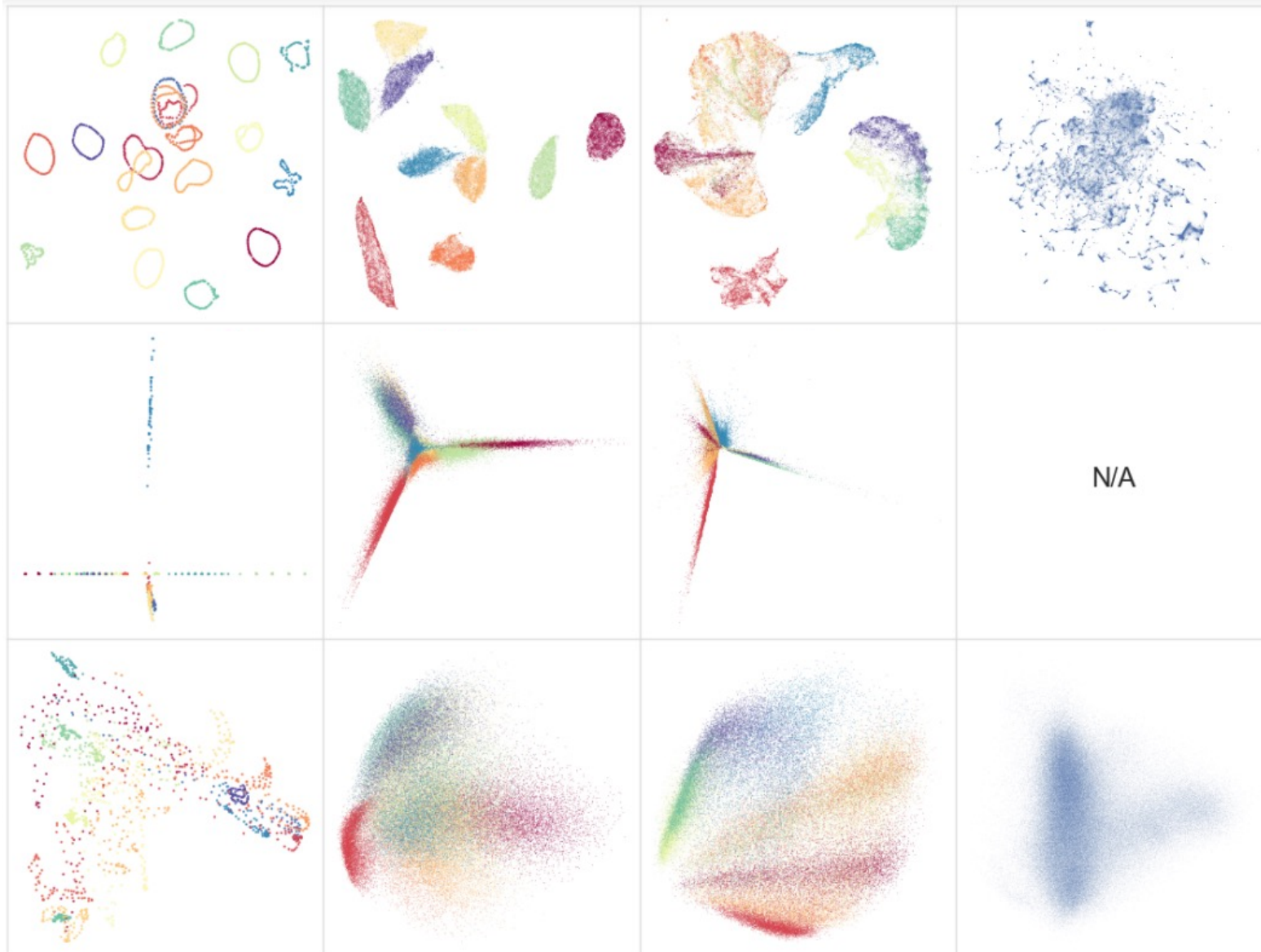


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Comparison with spectral approaches

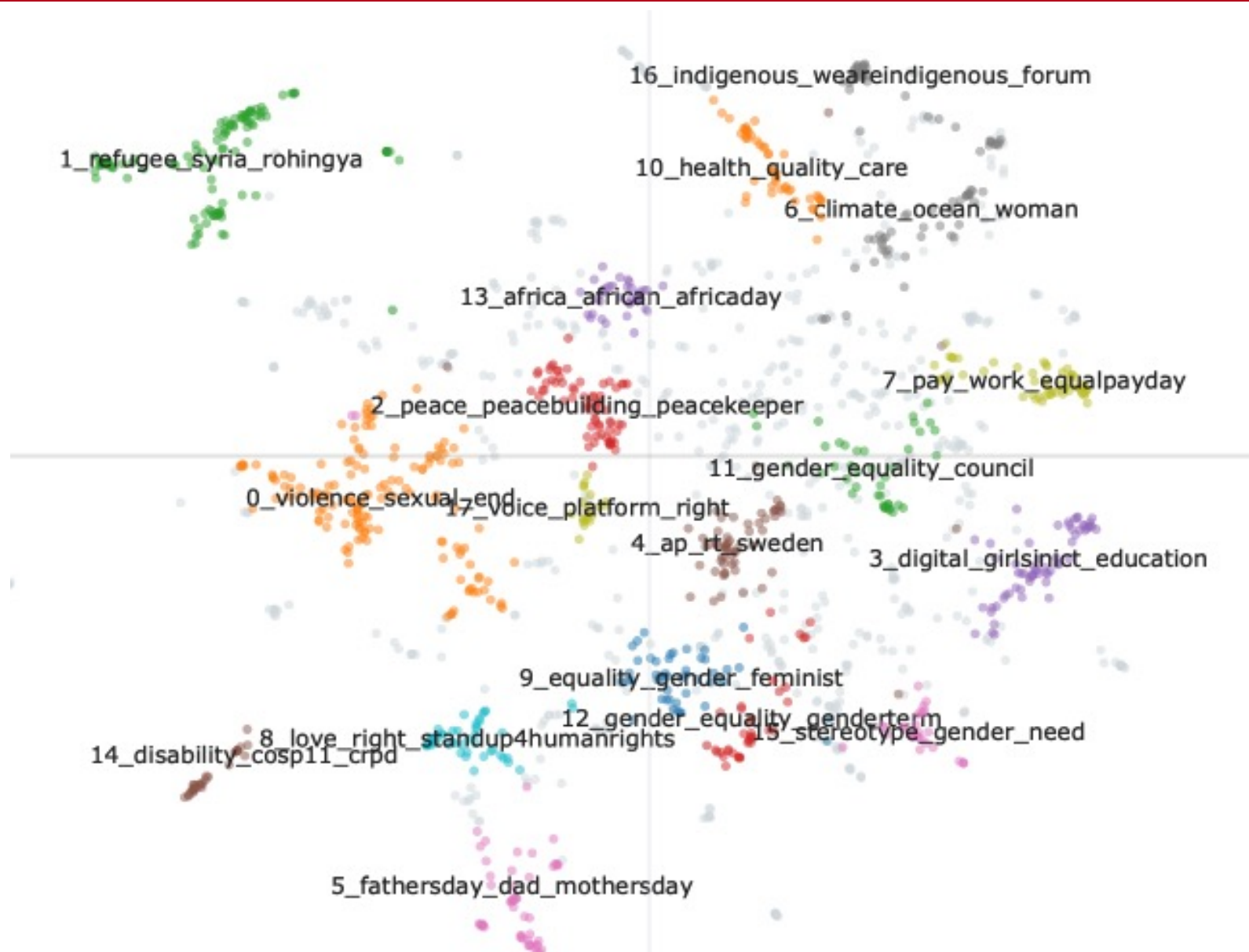
on the superior performance of force-directed algorithms

UMAP
Laplacian
eigenvector
PCA





A visual example document network using BERTopic





- ❑ easily understandable and implementable
- ❑ depending on the graphs (small and sparse)
- ❑ **amazingly good** layouts
- ❑ easily adaptable and configurable
- ❑ **robust**
- ❑ **scalable** (if wisely implemented)

But...

- ❑ quality mostly depends on the data (e.g., how to identify the **desired** link **distance** ℓ_{ij} ... might be challenging from an adjacency matrix)
- ❑ fine-tuning can be done by experts
- ❑ might need manual intervention

Software tools

for force-directed layouts

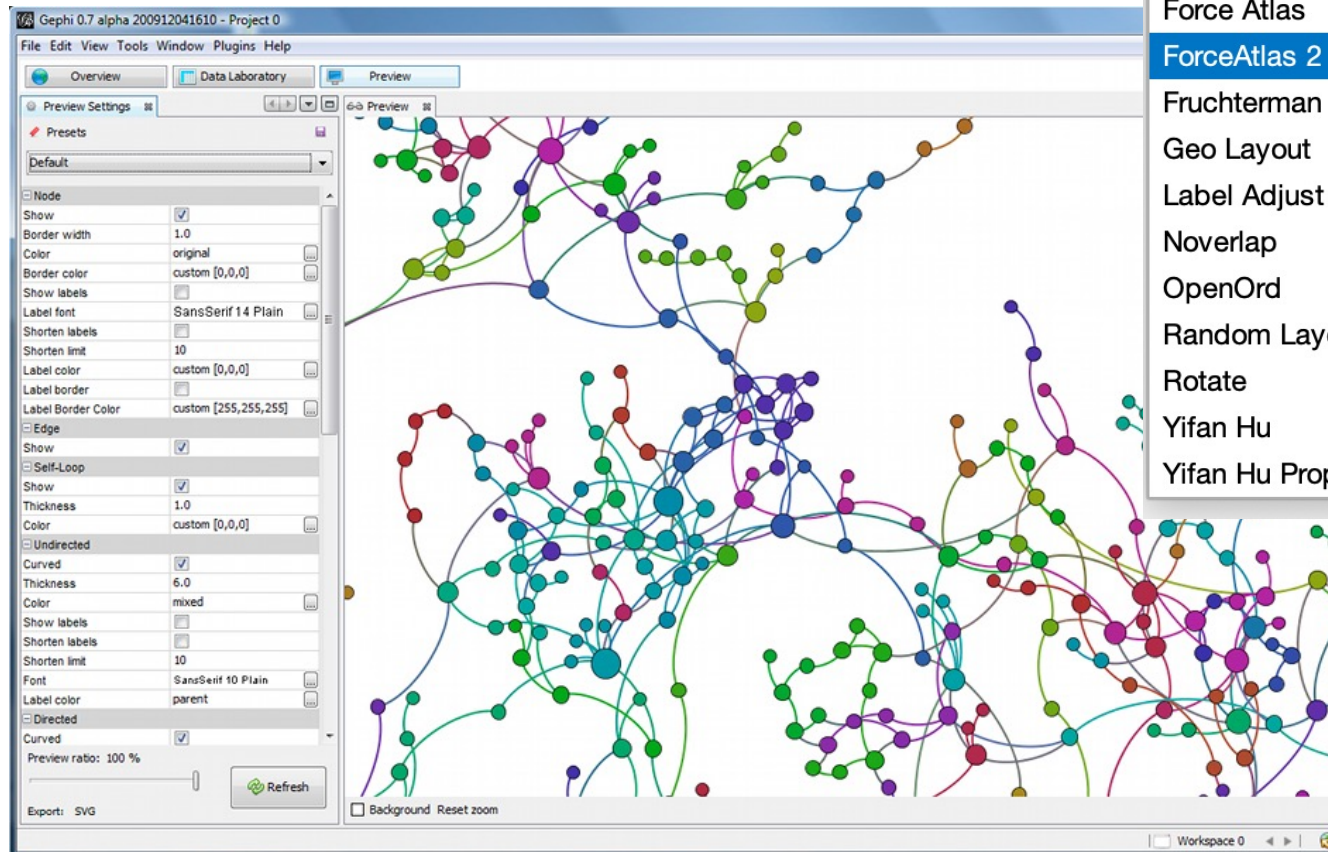


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Gephi

<https://gephi.org/>

Will be using it, **install** it
in your PCs **asap** !!!



Layout ×

---Choose a layout

- Contraction
- Expansion
- Force Atlas
- ForceAtlas 2**
- Fruchterman Reingold
- Geo Layout
- Label Adjust
- Noverlap
- OpenOrd
- Random Layout
- Rotate
- Yifan Hu
- Yifan Hu Proportional

Layout ×

ForceAtlas 2

Run

Threads

Threads number	3
----------------	---

Performance

Tolerance (speed)	1.0
Approximate Repuls	☒
Approximation	1.2

Tuning

Scaling	2.0
Stronger Gravity	☐
Gravity	1.0

Behavior Alternatives

Dissuade Hubs	☐
LinLog mode	☐
Prevent Overlap	☐
Edge Weight Influen	1.0
Normalize edge weig	☐
Inverted edge weigh	☐



❑ NetworkX

networkx.org/documentation/stable/index.html

kamada_kawai_layout

spring_layout → Fruchterman Reingold

spectral_layout

pydot_layout, graphviz_layout

❑ iGraph

python.igraph.org/en/stable/

layout_drl

layout_fruchterman_reingold

layout_graphopt

layout_kamada_kawai

layout_lgl, layout_mds

layout_umap → experimental 😊

❑ UMAP

<https://umap-learn.readthedocs.io/en/latest/>

UMAP



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UMAP example in R ☺ enhancing the role of communities

```
umap( $A \cdot (1 + \alpha C^T C)$ , n_neighbors=30, metric='cosine', min_dist=0.5)
```

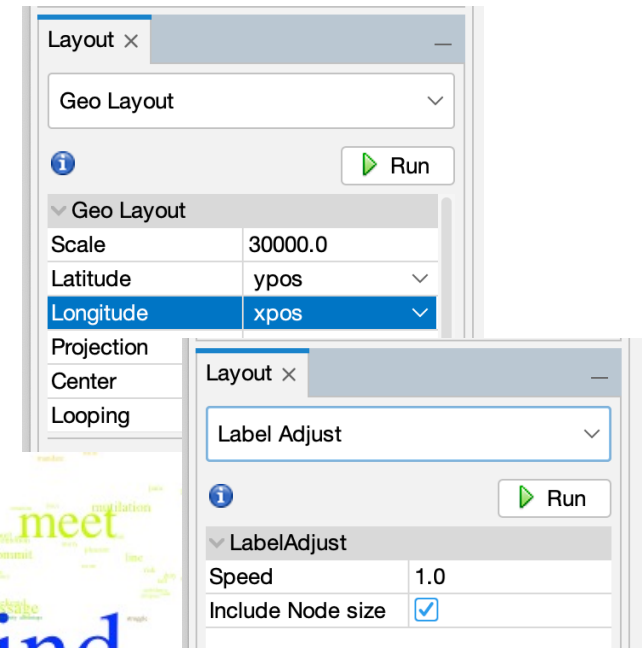
$\alpha = 9$

adjacency
matrix A
whose
elements
inside a
community
are multiplied
by $(1+\alpha)$;
communities
 C identified
by Louvain





Geo layout + Label adjust + Contraction/Expansion





- ❑ many layout algorithms are available in **Python**
- ❑ **UMAP** seems the best, but you never know
- ❑ **Gephi** is an alternative useful tool, but largely based on manual intervention
- ❑ use a combination of the two for best results
- ❑ a good project has a readable network (or wordcloud) clearly showing the role of communities