# Sparsity in Practice a bit of introspection





Felix Reidl Dagstuhl 2021

#### Part I

# The work so far



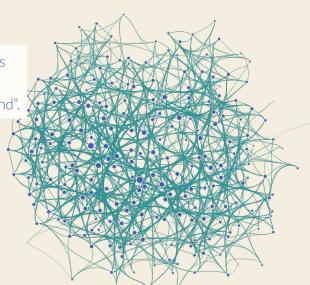


#### **Residence hall**

Student in ANU Hall

Friendship

Collected via interviews by Cynthia Webster, ranked as "best friend", "close friend", and "friend".

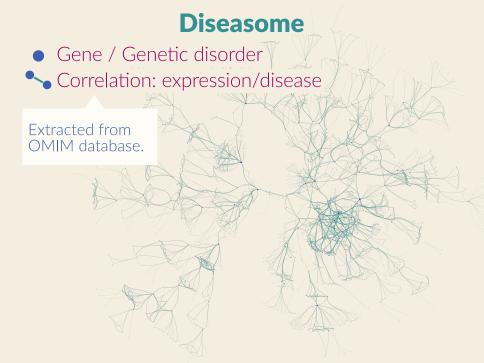


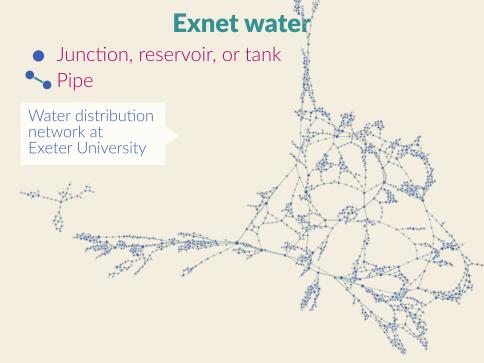


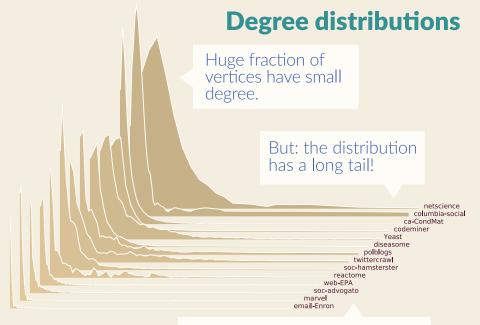
Proteins of Brewer's yeast

Interaction

Proteins interact *in vitro*, combination of several datasets.







More extreme in large networks!

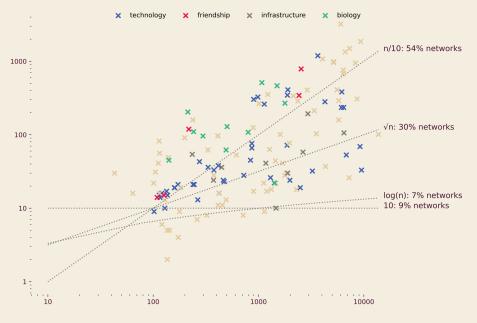
#### **Real vertex cover**



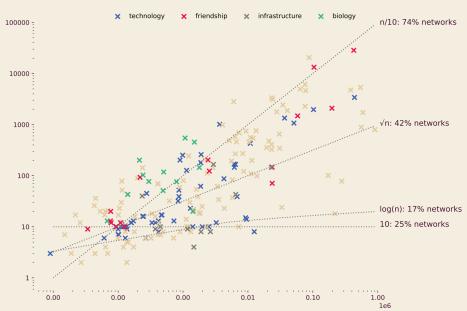
#### **Real treewidth**



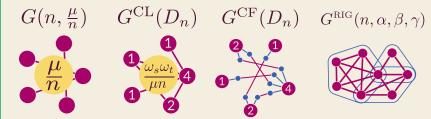
#### Real treedepth

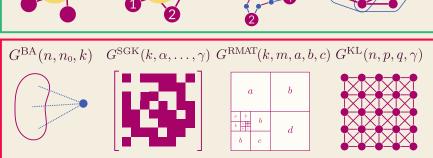


#### Real $wcol_3$

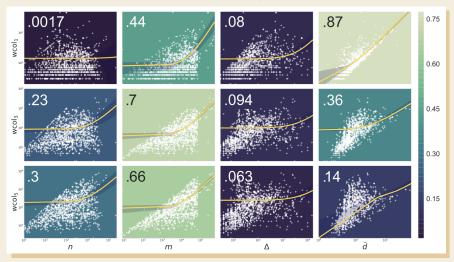


#### **Random model sparsity**





#### Real structural sparseness



W. Nadara, M. Pilipczuk, R. Rabinovich, FR, S. Siebertz: Empirical Evaluation of Approximation Algorithms for Generalized Graph Coloring and Uniform Quasi-Wideness. SEA 2018: 14:1-14:16

#### A hard-learnt lesson



Nešetřil J, de Mendez PO, Wood DR. Characterisations and examples of graph classes with bounded expansion. European Journal of Combinatorics. 2012 Apr 30:33(3):350-73.

Demaine ED, FR, Rossmanith P, Sánchez Villaamil F, Sikdar S, Sullivan BD. Structural sparsity of complex networks: Bounded expansion in random models and real-world graphs. Journal of Computer and System Sciences, 2019 May 24.

Farrell M, Goodrich TD, Lemons N, FR, Sánchez Villaamil F, Sullivan BD. Hyperbolicity, degeneracy, and expansion of random intersection graphs. Inhternational Workshop on Algorithms and Models for the Web-Graph 2015 Dec 10 (pp. 29-41). Springer, Cham.

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# No one cares.

\*not enough people

# **Part II**

# Reflection





# A false dichotomy









**NETWORK SCIENCE** 

Reality\*









\*Highly subjective opinion



#### Solvers vs Solutions





#### Solvers vs Solutions



Communities



#### Solvers vs Solutions



Diffuse problems

Competing views

Often sceptical of new approaches

Concrete problems

Single perspective

Appreciate new approaches





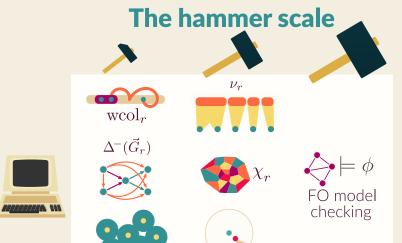
Must be very user-friendly

High maintenance

Usable for stakeholder

Maintenance by stakeholder









splitter games

r-Dominating

Set approx.



**Big hammers don't implement** 



#### Medium hammers don't scale

O'Brien MP, Sullivan BD.

Experimental evaluation of counting subgraph isomorphisms in classes of bounded expansion.

arXiv preprint arXiv:1712.06690. 2017 Dec 18.



#### Small hammers might just work!

Nadara W, Pilipczuk M, Rabinovich R, Reidl F, Siebertz S. Empirical evaluation of approximation algorithms for generalized graph coloring and uniform quasi-wideness. Journal of Experimental Algorithmics (JEA). 2019 Dec 10;24:1-34.

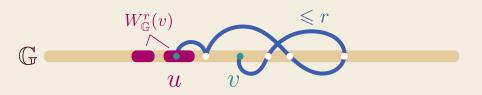
Brown CT, Moritz D, O'Brien MP, Reidl F, Reiter T, Sullivan BD. Exploring neighborhoods in large metagenome assembly graphs using spacegraphcats reveals hidden sequence diversity. Genome biology. 2020 Dec;21(1):1-6.

Reidl F, Sullivan BD. A color-avoiding approach to subgraph counting in bounded expansion classes. arXiv preprint arXiv:2001.05236. 2020 Jan 15.

github.com/ spacegraphcats/ spacegraphcats

github.com/ theoryinpractice/ mandoline

#### Weak colouring & bounded expansion

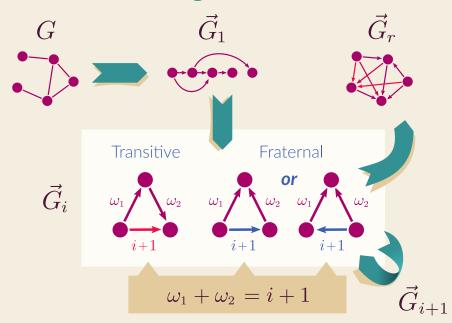


 $oldsymbol{u}$  is weakly r-reachable from v if there exists a path from v to  $oldsymbol{u}$  of length at most r such that  $oldsymbol{u}$  is the path's leftmost vertex.

$$\operatorname{wcol}_r(G) := \min_{\mathbb{G} \in \Pi(G)} \max_{v \in G} |W_{\mathbb{G}}^r(v)|$$

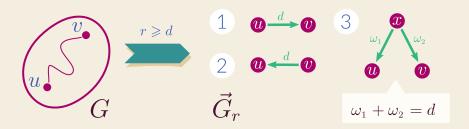


#### dtf-augmentations



#### **Distances under dtf-augmentations**

Let u and v be at distance d in G:



Pairs at distance at most r in the original graph have distance at most two in the r<sup>th</sup> augmentation.

#### **B.E.** & dtf-augmentations

There exist two (horrible) polynomials P and Q such that:

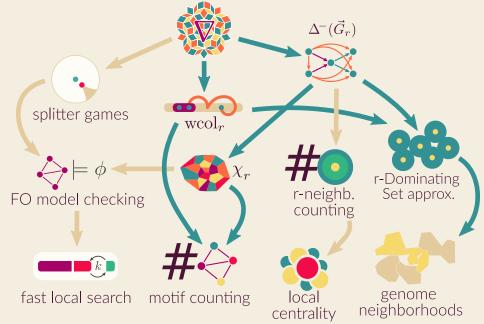
$$\chi_r(G) \leqslant P(\tilde{\nabla}_{(2\log r)^r}(G))$$
  
$$\Delta^-(\vec{G}_r) \leqslant Q(\tilde{\nabla}_r(G)\Delta^-(\vec{G}_1))$$



A graph class has bounded expansion iff it is  $\Delta^-(\vec{G}_r)$ -bounded.

We can compute dtf-augmenations in linear time (in bounded expansion classes)

### **Applications & Algorithms**

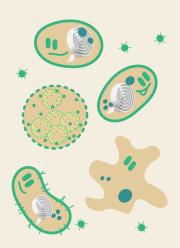


# Exhibit A CATLAS



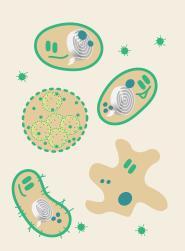
Metagenome exploration using hierarchical domination of de-Bruijn graphs

Joint work with C. Titus Brown, Dominik Moritz, Michael P. O'Brien, Taylor Reiter, Blair D. Sullivan





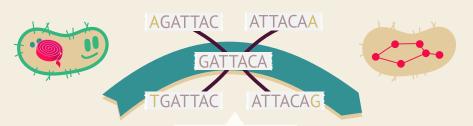






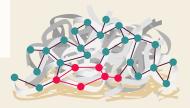


### De-Bruijn graphs

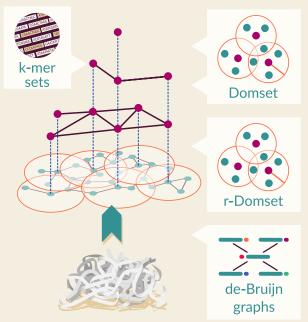




Bounded degree



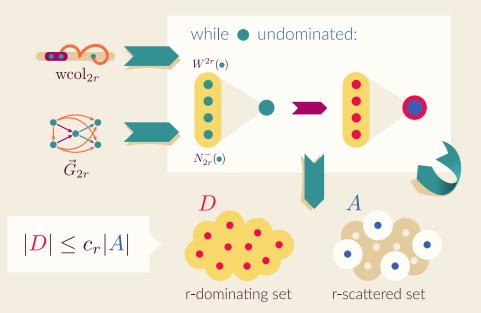
# **CATLAS Overview**

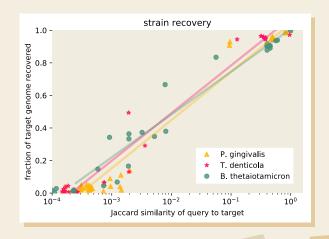


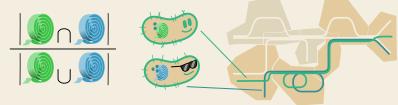




# Reminder: Dvořák's algorithm







# What really matters



The efficient computation of dominating sets will open up a whole new range of possibilities in bioinformatics.

C. Titus Brown, Associate Professor at UC Davis

spacegraphcats will transform the way biologists interact with genome assemblies.

It allows us to access previously discarded sequencing information thereby allowing more robust functional characterization.



Taylor Reiter, his much more eloquent post-doc

# **Engineering: efficiency**

**Lesson:** Stick to the 'sparsity methodology'.

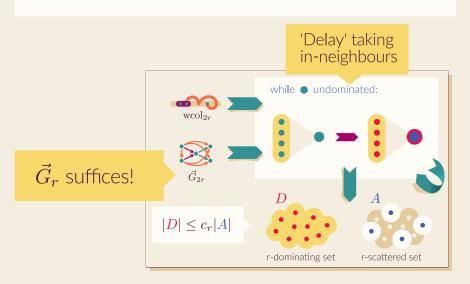
## Example: computing partition from r-domset





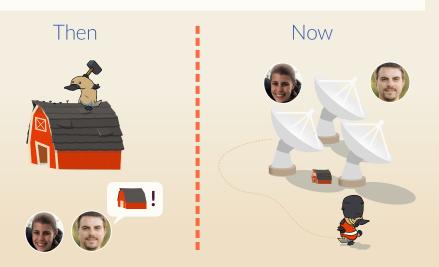
# **Engineering: efficiency**

**Lesson:** Practical issues inspire theoretical questions.



### **Collaboration**

**Lesson:** Expect roles to shift over time.



## **Next steps**

- 1 Faster language Believe it or not, so far this is all done in Python!
- 2 Improve network partition E.g. consider additional constraints
- 3 Whatever our collaborators need!



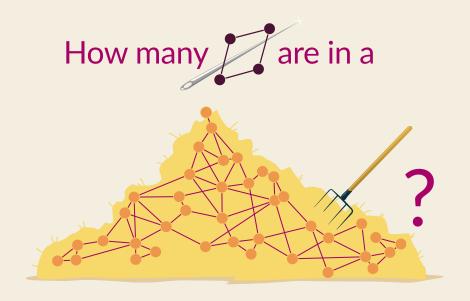
? Contracted de-Bruijn graphs These graphs have bounded degree, but probably much more structure that we could exploit!





Motif counting using generalized colourings

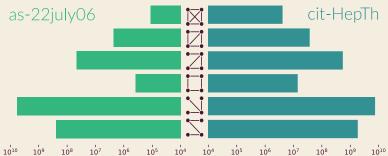
Joint work with Blair D. Sullivan





# **Graphlets**

We want to count all (connected) induced subgraphs up to a given size.



The graphlet degree distribution or the graphlet degree can be used to compare networks.

Pržulj N, Corneil DG, Jurisica I.

Modeling interactome: scale-free or geometric?. Bioinformatics. 2004 Jul 29:20(18):3508-15

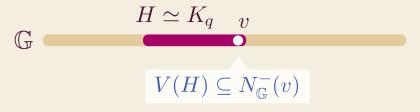
Pržulj N. Biological network comparison using graphlet degree distribution.

Bioinformatics, 2007 Jan 15:23(2):e177-83.

### Let's start with something easy!

We count cliques in a d-degenerate graph.

**Observation**: every clique is contained in the left-neighbourhood of its *last* vertex.



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We count cliques in a d-degenerate graph.

**Observation**: every clique is contained in the left-neighbourhood of its *last* vertex.

$$H \simeq K_q \quad v$$

$$V(H) \subseteq N_{\mathbb{G}}^-(v)$$

Therefore we can enumerate all cliques by enumerating all cliques in  $N^-(v)$  for all  $v \in G$ !

$$O(2^d n)$$
 time!

#### **Does it blend?**

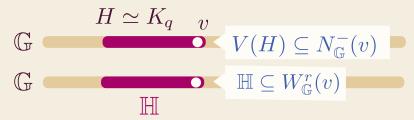
Can we 'lift' this algorithm to wcol?



1) What is the 'last' vertex of H? Enumerate all orderings  $\mathbb{H}$  of H.

#### **Does it blend?**

Can we 'lift' this algorithm to wcol?

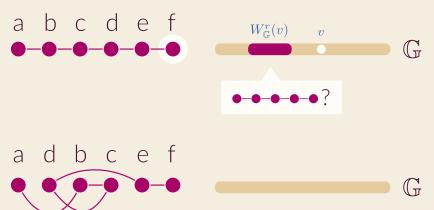


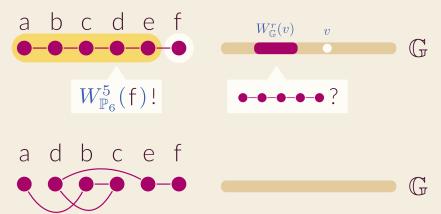
- 1) What is the 'last' vertex of H? Enumerate all orderings  $\mathbb{H}$  of H.
- 2 Does  $\mathbb{H} \subseteq W^r_{\mathbb{G}}(v)$  actually hold? Only sometimes!

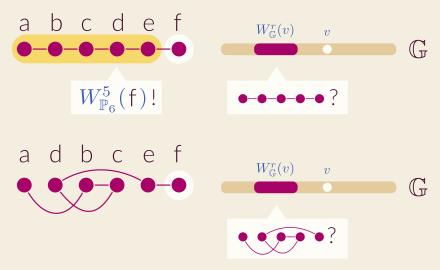


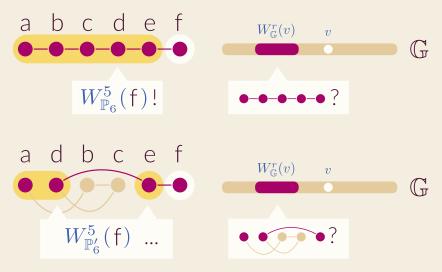


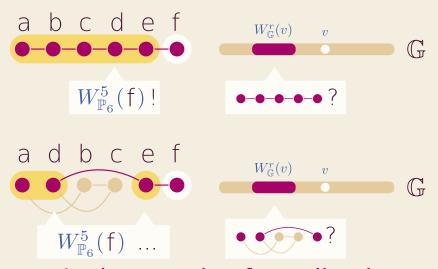






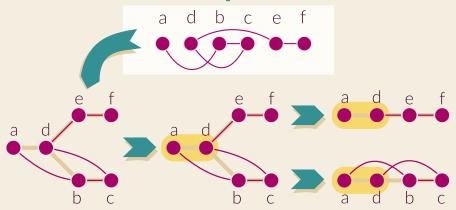






Is there a nice formalization of this property?

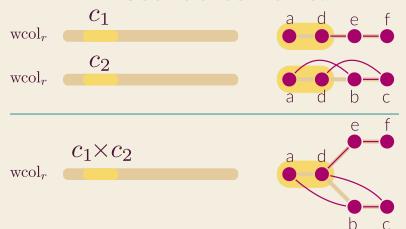
# **Decomposition!**



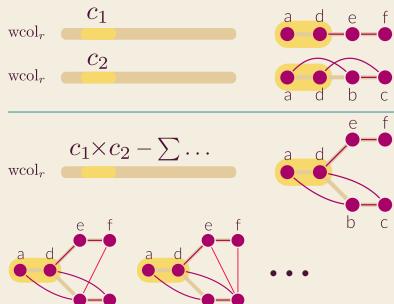
We can count linear pieces!

Progress! These pieces are linear!

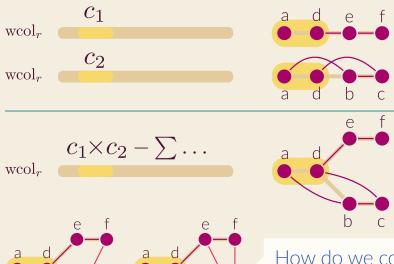
#### **Count & combine!**



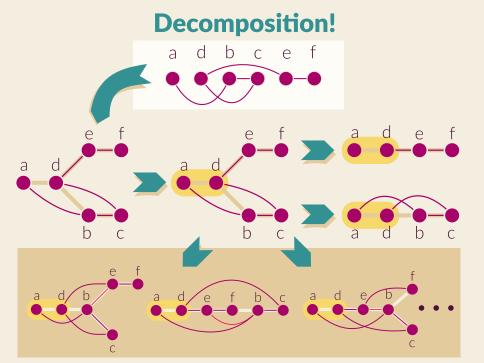
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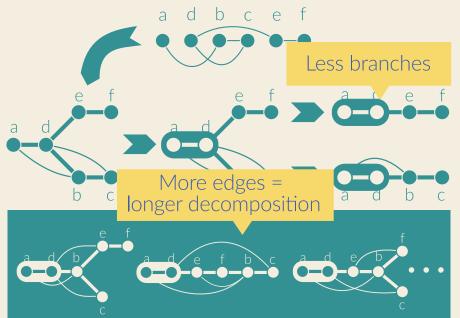
#### **Count & combine!**



How do we count these graphs?



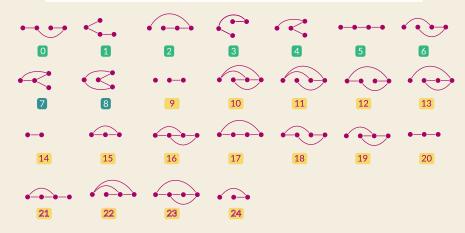
# **Decomposition!**



## Counting $P_4$ s using $wcol_3$

**Lemma 6.** Let  $\mathbf{H} \in \mathcal{H}$  be a (non-linear) pattern relaxation and let  $\mathbf{H}_1 \oplus_{\bar{x}} \mathbf{H}_2 = \mathbf{H}$ . Fix an ordered vertex set  $\bar{y} \in \mathbb{G}$  such that  $\mathbf{H}[\bar{x}] \simeq \mathbb{G}[\bar{y}]$ . Then

$$\underset{\bar{x} \mapsto \bar{y}}{\#}(\mathbf{H}, \mathbb{G}) = \underset{\bar{x} \mapsto \bar{y}}{\#}(\mathbf{H}_1, \mathbb{G}) \underset{\bar{x} \mapsto \bar{y}}{\#}(\mathbf{H}_2, \mathbb{G}) - \sum_{\mathbf{D} \in \mathcal{D}(\mathbf{H}_1, \mathbf{H}_2)} \frac{\underset{\bar{x} \mapsto \bar{x}}{\#}(\mathbf{H}, \mathbf{D} \mid \mathbf{H}_1, \mathbf{H}_2) \underset{\bar{x} \mapsto \bar{y}}{\#}(\mathbf{D}, \mathbf{D})}{\underset{\bar{x} \mapsto \bar{x}}{\#}(\mathbf{D}, \mathbf{D})}.$$



### Counting P<sub>4</sub>s using wcol<sub>3</sub>

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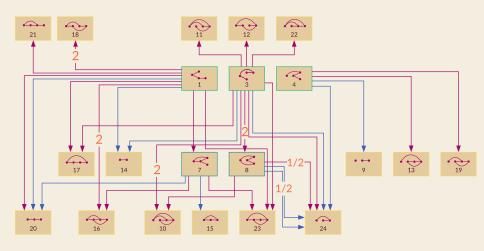
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$$4 \quad \bigcirc = \bigcirc \times \bigcirc \times \bigcirc - \bigcirc + \bigcirc \bigcirc$$

$$7 \quad \bullet \bullet = \bullet \quad \times \bullet \quad - \left( \bullet \bullet + \bullet \bullet \right)$$

$$= \frac{1}{2} \left( \underbrace{\phantom{0}}_{24} \times \underbrace{\phantom{0}}_{24} \right) - \left( \underbrace{\phantom{0}}_{10} + \underbrace{\phantom{0}}_{24} \right)$$

# Counting $P_4$ s using $wcol_3$



This 'counting-DAG' has bounded depth

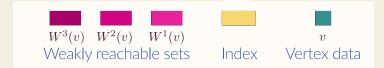
Subgraph counting using  $wcol_r$ Count linear patterns directly  $\frac{2\pi}{1000} = \frac{1}{2} (\bullet - \bullet \times \bullet - \bullet) - (\bullet - \bullet + 2 \bullet - \bullet)$ Compute composite Aggregate pattern counts

## **Engineering: memory**

**Lesson:** We cannot ignore memory locality. But this is really hard when working with graphs.

1 Flatten everything





# **Engineering: memory**

**Lesson:** We cannot ignore memory locality. But this is really hard when working with graphs.

2 Be as specific as possible

Vertex ids

(int16)  $\approx 60 \mathrm{K}$ 

int32  $\approx 4.3 \text{Mio}$ 

Node ids

3 bit

4 bit

k

n

Tough design choices

int64

All we could ever need

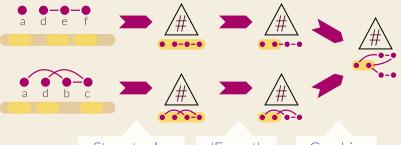
char

All we could ever need

## **Engineering: data structures**

**Lesson:** Common data structures not enough

3 Design special-purpose data structures



Store tuples

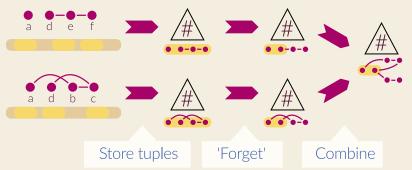
'Forget'

Combine

# **Engineering: data structures**

**Lesson:** Common data structures not enough

3 Design special-purpose data structures



Our solution: specialized trie. Better options?

#### **Next steps**

1 More features

Count coloured graphs, sum-of-weights, other types of embeddings (homomorphisms, non-induced, etc.)

2 Implement for  $col_r(\mathbb{G})$ 

We know how to do this in theory and  $\operatorname{col}_r(\mathbb{G})$  seems to be smaller in practice.

? Preprocessing!

Simple preprocessing rules (based e.g. min-degree of pattern) help a lot. Can we generate more elaborate, pattern-dependent rules?



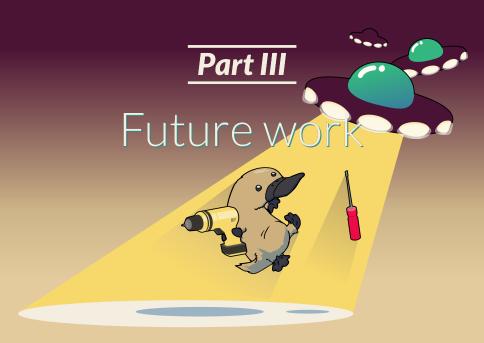
# Solvers vs Solutions











# The big open question

# **THANKS**





#### Exhibit X





Local centrality measures via neighbourhood surveys (Secret slides)

# **Close-to-Closeness Centralities**

C(v) r-Local version  $\left(\sum_{i=1}^{n} d_{i+1}(v,v)\right)^{-1}$ 

Closeness  $\left(\sum_{u \in G} \operatorname{dist}(u, v)\right)^{-1} \quad \left(\sum_{u \in N^r[v]} \operatorname{dist}(v, u)\right)^{-1}$ Harmonic  $\sum_{u \in G} \operatorname{dist}(u, v)^{-1} \quad \sum_{u \in N^r[v]} \operatorname{dist}(v, u)^{-1}$ 

Harmonic  $\sum_{u \in G} \operatorname{dist}(u, v)^{-1} \qquad \sum_{u \in N^r[v]} \operatorname{dist}(v, u)^{-1}$   $|\{u \mid \operatorname{dist}(u, v) < \infty\}|^2 \qquad |N^r[v]|^2$   $\sum_{u \in N^r[v]} \operatorname{dist}(v, u)$ 

 $u \in N^r[v]$ 

All three measures can be computed quickly if we know  $|N^d(v)|$  for  $1 \le d \le r$ .

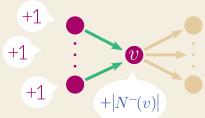
Can we compute this quickly in sparse graphs?

 $dist(u,v) < \infty$ 

# Warm-up: Counting with degeneracy

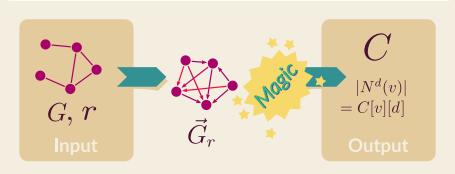
Let G be (d-1)-degenerate.

- 1 Compute orientation  $\vec{G}$  with  $\Delta^-(\vec{G}) \leqslant d$  in linear time.
- 2 Initialize counter C[v] = 0 for all  $v \in G$ .
- 3 For every  $v \in G$ , increment C[v] and C[u] for every in-neighbour  $u \in N^-(v)$ .



# **Degeneracy to dtf-augmentations**

**Thm.** Given a graph G and an integer r, we can compute the size of  $|N^d(v)|$  for all  $v \in G$  and  $1 \le d \le r$  in total time  $O(2^{\Delta^-(\vec{G}_r)}n)$ .



We compute the size of the r<sup>th</sup> nbhds:

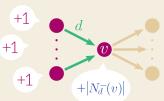
1 Compute dtf-augm.  $\vec{G}_r$  with small  $\Delta^-(\vec{G}_r)$  in linear time.

We compute the size of the r<sup>th</sup> nbhds:

- 1 Compute dtf-augm.  $\vec{G}_r$  with small  $\Delta^-(\vec{G}_r)$  in linear time.
- 2 Initialize counter C[v][d] = 0 for all  $v \in G$  and  $d \leq r$ .

We compute the size of the rth nbhds:

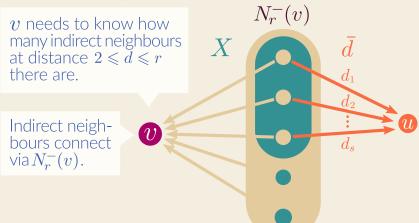
- 1 Compute dtf-augm.  $\vec{G}_r$  with small  $\Delta^-(\vec{G}_r)$  in linear time.
- 2 Initialize counter C[v][d] = 0 for all  $v \in G$  and  $d \leq r$ .
- 3 For every  $v \in G$ , increment C[v][d] and C[u][d] for every in-neighbour  $u \in N_d^-(v)$ .

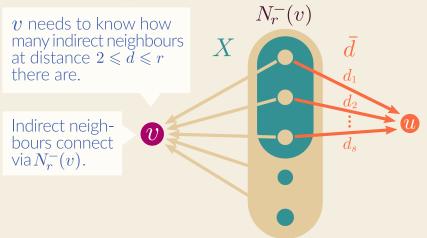




The counting so far takes care of the first two cases, but what about the *indirect* neighbours?

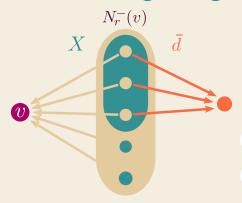
This is where the algorithm becomes **interesting**.





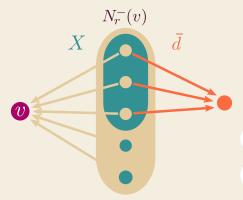
We compute the distance between v,u as follows:

$$dist(u, v) = \min(dist(v, X) + dist(u, X))$$



We need to compute for every set  $X\subseteq N_r^-(v)$  and every possible dist.-vector  $\bar{d}\in [r]^{|X|}$  the number of vertices u such that:

- 1  $N_r^-(u) \cap N_r^-(v) = X$
- 2 dist $(u, X) = \bar{d}$



We need to compute for every set  $X\subseteq N_r^-(v)$  and every possible dist.-vector  $\bar{d}\in [r]^{|X|}$  the number of vertices u such that:

- $1 \quad N_r^-(u) \cap N_r^-(v) = X$
- $2 \operatorname{dist}(u, X) = \bar{d}$

Let us call this number  $c(v, X, \bar{d})$ . Our first goal is to compute it for every vertex.

#### A data structure for $c(v, X, \overline{d})$

1 For every  $v \in \vec{G}_r, X \subseteq N_r^-(v)$  and  $\bar{d} \in [r]^{|X|}$ , initialize  $R[X][\bar{d}] = 0$ .

# A data structure for c(v, X, d)

by one.

- 1 For every  $v \in \vec{G}_r, X \subseteq N_r^-(v)$  and  $\bar{d} \in [r]^{|X|},$  initialize  $R[X][\bar{d}] = 0.$
- initialize R[X][d] = 0. 2 For every  $v \in \vec{G}_r, X \subseteq N_r^-(v)$ , increment  $R[X][\mathrm{dist}(v,X)]$

# A data structure for $c(v, X, \overline{d})$

- 1 For every  $v \in \vec{G}_r, X \subseteq N_r^-(v)$  and  $\bar{d} \in [r]^{|X|},$  initialize  $R[X][\bar{d}] = 0.$
- 2) For every  $v \in \vec{G}_r, X \subseteq N_r^-(v)$ , increment  $R[X][\operatorname{dist}(v,X)]$  by one.

#### Claim.

$$c(v, X, \bar{d}) = \sum_{X \subseteq Y \subseteq N_r^-(v)} (-1)^{|Y \setminus X|} \sum_{\bar{d}': \bar{d}'|_X = \bar{d}} R[Y][\bar{d}'].$$

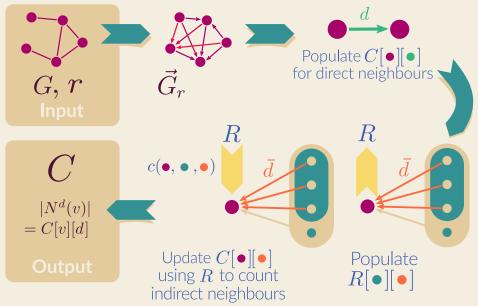
Given  $c(v, \bullet, \bullet)$  we can now count the number of indirect neighbours of v. For every subset  $X \subseteq N_r^-(v)$  and distance-vector  $\bar{d} \in [r]^{|X|}$ , apply the update:

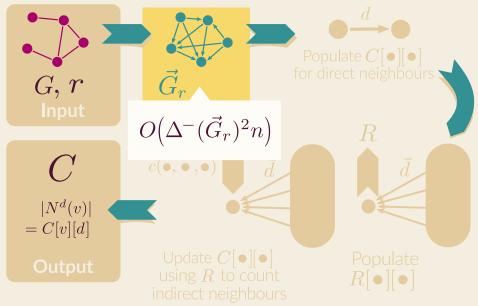
$$C[v][\min(\bar{d} + \operatorname{dist}(v, X))] \ += \ c\left(v, X, \bar{d}\right)$$

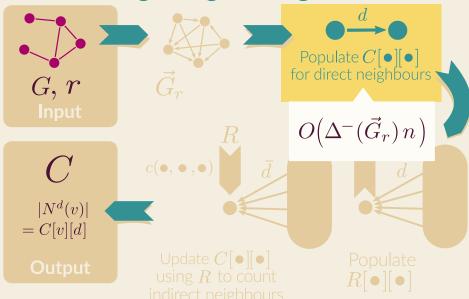
Since the above counts  $\,v\,$  as a neighbour of itself, we apply the following correction:

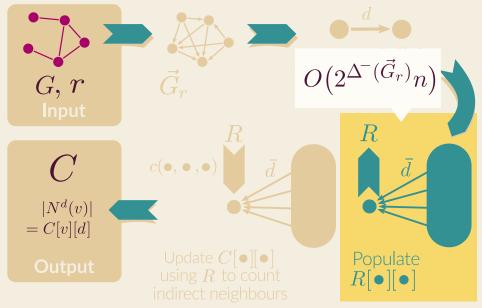
$$C[v][\min(\operatorname{dist}(v,X) + \operatorname{dist}(v,X))] = 1$$

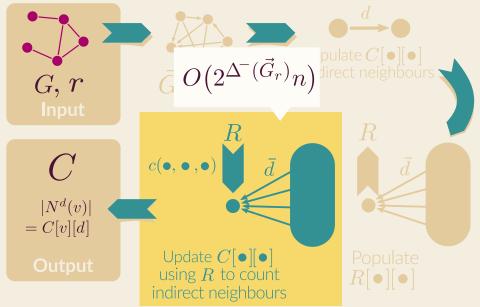
There are a few more corrections concerning direct neighbours, see paper.











**Thm.** Given a graph G and an integer r, we can compute the size of  $|N^d(v)|$  for all  $v \in G$  and  $1 \le d \le r$  in total time  $O(2^{\Delta^-(\vec{G}_r)}n)$ .

- Exponential vs quadratic?
- Does not scale to nowhere dense graphs!



Can we do **better**?

#### Can we do better?

#### CLOSED 2-NEIGHBOURHOOD SIZES

Input: A graph G.

Output:  $|N^2[v]|$  for every  $v \in G$ .

**Thm.** Unless SETH fails, 2-CNBS cannot be solved in time

- $\bullet O(|G|^{2-\varepsilon})$
- **2**  $O(2^{o(\Delta^{-}(\vec{G}_{2}))}n^{2-\varepsilon})$

# **Engineering: compromises**

**Lesson:** We cannot be *too* idealistic. Some vertices will have large in-neighbourhoods.

Mix algorithms

Conduct regular bfs for vertices with large inneighbourhood, use dtf-magic for everything else.

Luckily, these two approaches mix!

#### **Next steps**

1 More features

We can easily count the weights of neighbourhoods or neighbourhoods restricted to certain colours.

2 Optimize for small r

Most real-world cases will be for small distances. In particular for r=2 we can simplify the algorithm.

- ? Nowhere dense?
- Combine with neighbourhood complexity, handle large intersections differently.
- ? Approximate?

The Möbius inversion is the bottleneck. Sacrifice precision for speed?