
CSE 269 Approx. Algs

CSE 142 Machine Learning

CSE 185E Tech. Writing

ECE 175 Engg. CMT + Geo.



Vasjos

Approximation Algorithms

NP-hard algorithms, polynomial time, where not known

We do a presentation, read papers, and this is under ^{theoretical} CS

Approximating efficient values

class is reading papers, examining problems, presenting

Approximate solution for drive to SS.

Given $G(V, E, w)$
 $w > 0$

$A \xrightarrow{\text{stuff}} B$

BFS is a soln.

← what does this mean. w means weight

Dijkstra generalizes BFS $V = \# \text{ vertices}, M = \# \text{ edges}$

$V = \binom{n}{n}, M = \binom{E}{n}$

→ $O(m \log n)$

↳ 3 properties ↯ if $n \ll m$ then $\hookrightarrow \approx \tilde{O}(m)$

Polynomial Time, computes optimal solution, Always correct

↳ "efficient algorithm" $O(n), O(n^2), n \log n, n^3$ not 2^n or n^n

You can relax all 3 properties.

like 1 → slight exponential time: $1.3^n \ll 2^n$

For 2 → maybe approximate the optimal solution,

APX → 120, opt = 100

One method for the multiplicative factor appx : $\text{appx} \leq \kappa(\text{opt})$ $\text{appx/opt} = 1.2$
 \hookrightarrow common \hookrightarrow multiplicative factor

↳ additive factor appx : $\text{appx} \leq \text{opt} + B$ \hookrightarrow additive factor
 \hookrightarrow not scalar, not popular

③ Relaxes the probability of being correct. Related to analysis like worst-case ($O(n)$)
 \hookrightarrow Dijkstra $\forall G \leq O(m \log n)$ \downarrow any graph

But most places care for specific, not generic, instances.

Average case analysis, on random graphs. (heads or tails on each node/edge existing). $G(V, E, w)$. Cryptography looks at average case, & many algorithms care about this.

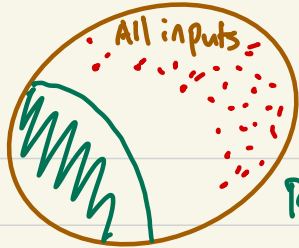
Some situations have high clustering coefficients $A \triangle B C$ (social network)
 while random graphs are $1/2$ (heads/tails exists)

How many friends in networks, for best algo, $E(\text{deg}) = p(n-1)$

This is power scaling, popular will have higher degree.

So for some products avg or worst case might not be best.

Beyond worst-case analysis (WCA) uses assumptions for inputs to creating algorithms.



Very bad cases (NP hard/complete)

Real world input space, typical inputs.

We're using this information for approx algorithms

These are called 'planted' solutions, like a password to brute force

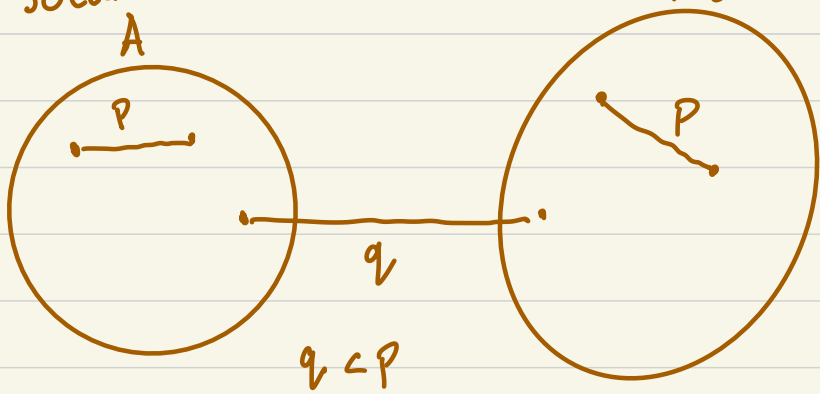
(example)

A popular model is called stochastic block model for community detection. Can you recover the 2 communities?

$P = \text{friends}$

soccer fans

Tennis fans



Can you recover the communities, and can you do it in ^(effici.) polynomial time?

$G(n, P)$ $G(n, p)$
 \hookrightarrow edge present w/ probability P

Recovery means finding all nodes inside A, and all nodes in B.

Exact recovery would be finding all the people in each community based on connections ($p=0.8, q=0.6$). for example. (for an outsider guessing person in A or B).

Approx recovery of communities would label a subset and find it. ^($x < .2$?)
algs exist such that if $p-q = x$, then can find solution
we want to recover w/ better probability than $n/2$ ($1/2$ guess).

What is satisfiability?

3 SAT $(x_1 \vee \bar{x}_2 \vee x_3) \wedge (x_7 \vee x_8 \vee \bar{x}_9) \wedge \dots$
 $\uparrow_{\text{or}} \quad \quad \quad \uparrow_{\text{and}}$

\hookrightarrow NP-complete, All NP-complete are a 3 SAT special case, vice-versa.

Read $1/2$ papers, write a report, do research.
and present to the class, and then a final exam.

Lecture 2, Oct 2nd

bet add. code?

Beyond WCA?

How does google search work?

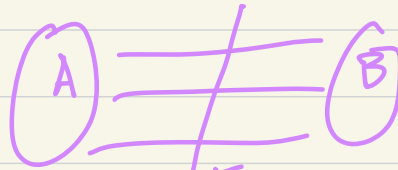
Hierarchical clustering.

Find partition of dataset into similar looking data points.

Clustering. Given points $S = \{x_1, x_2, \dots, x_n\}$
Find similar points.

Max-cut is a clustering problem. Points have a distance, separating into 2 groups is based on distance

$G(V, E, w)$



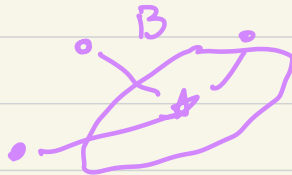
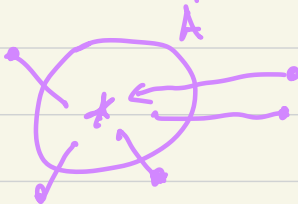
max $w(\text{cut})$

weight is maximizing the cut of connections

K-center, k median, k-means, k-shape optimize different objective functions for slightly different things.

means:

minimize all points to their center of group



K-median. puts it closer to most points

K-means comes about distance squared

K can be any number.

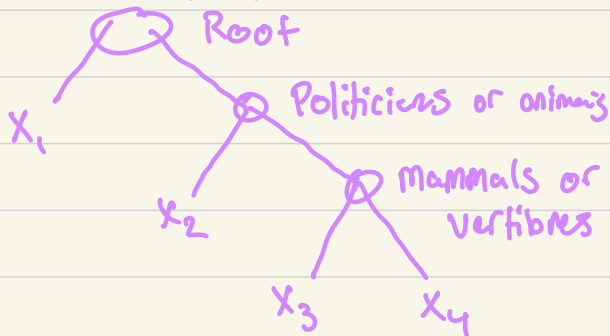
Hierarchical clustering goal is to get hierarchy of data points
How do you rank nice shoes? Instead of 4 clusters for sandals, running shoes, formal, kids, etc (like graph partitioning, or k-means) we use hierarchical clustering.



How do you find the # of clusters effectively \uparrow 3 vs 5 vs 100.

Hierarchical clusters

$n: x_1, x_2, x_3, \dots, x_n$



What do you do to split this effectively, what objective function (having similarity score).

Q (objective function):

$$\text{cost}(T) = \max_{x_i, x_j} \left(\frac{\text{dissimilarity}(x_i, x_j)}{\# \text{ hops}} \right)$$

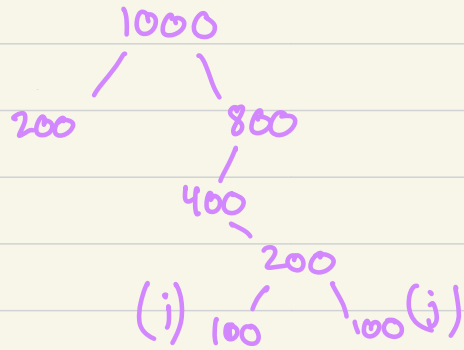
Everything is relative to rest of hierarchical, my idea was x_2 (adds all dissimilarity scores) compared to x_3 .

A good/popular cost function: → from STOC '16 journal

penalize if similar things (papers) were separated early on the tree versus later on the tree.

$$\text{cost}(T) = w_{ij} \left| \frac{\text{\# data points present when } i, j \text{ got split}}{n} \right|$$

similarity of pair $\cdot \left(\frac{200}{1000} \right)$
 split @ 200



$$\text{cost}(T) = \sum_{i,j \in E} w_{ij} \frac{|\text{\# present}|}{n}$$

$i, j \in E$ ↓
 similarity score (positive)

fraction of points present in split

Two types of clustering algorithms (popular) ↓

- linkage algorithms (single linkage, average linkage, complete linkage).

All points are separate, and merge documents that are highly similar. Bottom-up approach

- Divisive (top-down). Take all points together, and find the best 'cut' / 'split' to separate into clusters (balance-cut, sparsest-cut).

Represent \rightarrow as a convex optimization problem

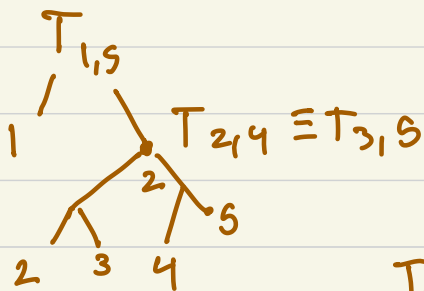
$$\text{cost}(T) = \sum_{i,j \in V} w_{i,j} \left| \frac{\text{\# points present during cut}}{n} \right|$$

Setting: $G(V, E, w \geq 0)$ $|V| = n$

vertices edges weights

Find (binary) tree minimize: $\text{cost}(T) = \sum_{i,j \in E} w_{i,j} \left| \frac{\text{\# Trees of } T_{i,j}}{T_{i,j}} \right|$

Penalty



$$\sum_{i,j \in E} w_{i,j} |T_{i,j}|$$

\hookrightarrow want the tree as low as possible when similarity is high (w).

$T_{i,j}$ is the subtree rooted at the LCA(i,j)

How do we write the objective function as vectors and use convex optimization relaxation.

(continuous optimization is easier than discrete optimization)

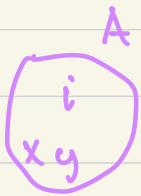
$$\sum_{i,j \in E} w_{ij} |T_{ij}|$$

Convex Relaxations

1) Assign a boolean variable for each pair of nodes
 $x_{ij} = \{0, 1\}$

2) (assign vectors)
All data points are now vectors w/ their features

Aside: easier example. Variable that says whether 2 points are separated or not.



$$x_{ij} = 1, \quad x_{x_y} = \emptyset$$

max-cut maximizes weight of edges cut.

↳ Output is partition to split in 2.

We're given $\{1$ if x_{ij} split from cut, \emptyset if not-cut $\}$

This obj function cuts it:

$$\max_{\text{assign} \rightarrow x} \sum_{i,j \in E} w_{ij} \cdot x_{ij}$$

x_{ij} or 1

$$x_{ij} = \begin{cases} 1 & \text{if split} \\ 0 & \text{if not split} \end{cases}$$

my idea to implement \uparrow for the main problem

T_{ij} = subtree minimize size

• similarity

$x_{ij} = \begin{cases} 1 & \text{split} \\ 0 & \text{nosplit} \end{cases}$

$$\text{minimize: } \left[\begin{array}{l} x_{ij} \cdot T_{ij} \cdot (1 - \text{similarity}) \\ \text{split} \end{array} \right]$$

Solution:



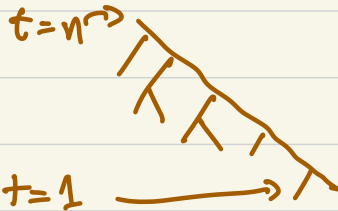
T_{ij} = Lowest common ancestor

• lets say here i & j split here

• we need 1 variable per level to describe this pair i, j

worst case of binary tree is $(n-1) \cdot \frac{1}{2} \cdot \frac{1}{4} \cdot \frac{1}{8} \dots \frac{1}{n}$, best is $\log(n)$?

we define 'level' $t \rightarrow$



x_{ij}^t is $\{0, 1\}$ if

i, j are together at level t .
(1 if split)

T_{ij} = lowest common ancestor

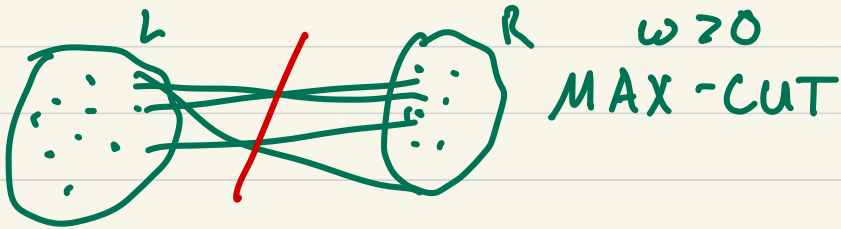
$t=n$	0	: \leftarrow once 1 always 1.
$t=n-1$	0	
$t=T_{ij}$	1	
$t=1$	1	

Lecture Oct 6 (late)

Talked about relaxation / NP-hard
Hierarchical clusters & semi-definite programming.

(2) graphs of vertices, edges, weights -

$G(V, E, w)$ given this split into L & R so the edges cut between both groups is maximized

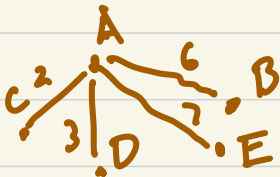


Given vertex u we know to keep on left or right side.

$$A: V \rightarrow \{0, 1\} \quad u \begin{cases} \rightarrow L \\ \rightarrow R \end{cases}$$

$$\max \sum_{u, v \in E} w_{uv} \mathbb{1}_{\{A(u) \neq A(v)\}}$$

\hookrightarrow what does this & u mean?



A goes to left, C & D go left, B on right $u \in E$

We used to use greedy

If all edges are cut, then bipartite then:
(all separated?)

max function (sum of all weights) = 2.

↳ back then greedy would take A & go from there
w/ opt $\geq 1/2$

Even worst case on any graph would mean
result in a 50% split (random split.)
Bipartite graph splits everything (100%)
50% of splitting or not cutting

Semi-definite programming was to improve the random split

One idea was to rewrite MAX-CUT:

For every vertex $v \in V$: a variable $x_v \in \{0, 1\}$ ↙ left or right
For every edge $e \in E$: $z_e \in \{0, 1\}$ ↗ cut edge y/n

How to rewrite using \uparrow weight \rightarrow cut y/n

$$\max \sum_{u, v \in E} w_{u, v} \cdot z_e$$

This would cut everything
so we need to have constraints so
is not = 1

Constraints for partitions:

1) $z_{u, v}$ is less than $x_u + x_v$, so if both $u \& v$

are on the same side then their connection is not cut:

$$Z_{uv} \leq x_u + x_v \quad \times 6 \text{ } \{0,1\}$$

$$Z_{uv} \leq 2 - x_u - x_v$$

People thought to solve over $[0,1]$ instead of $\{0,1\}$ ^{interval} ^{set.}
This is called the Linear Programs (LP)

↳ linear objectives & constraints for optimal solution

what is relaxation

But bc a cut is not 0 & 1 but can be 0.7 or 0.3 then the solution a weakness: (L/R)

You can set all variables to 0.5 & all z 's to 1 (cut)
which gives us the same response for all graphs.
LPs are not useful for MAX-CUT / this program.

We'll try another formulation to lead us to ^{Semi Definite} programming.

∀ vertex v encodes left or right.

$y_v \in \{-1, 1\}$ ← additional variable

$$\max \sum_{u,v \in E} w_{u,v} \cdot \left(\frac{1 - y_u \cdot y_v}{2} \right)$$

Discrete Problem

if same side cut $1 - (-1)(-1) = 0$ ✓ 0
if different sides then $1 - (1)(-1) = 2$
then divide by 2 = 1

For every vertex we add a vector of n dimensions

↳ we allow $y_v \in \mathbb{R}^2$, but must be unit vector

(l^2 norm = 1) with constraint:

What is an inner product

→ $\vec{y}_v \cdot \vec{y}_u = 1, \forall v \in V \quad \|\vec{y}_v\|_2 = 1$

max $\sum_{u,v \in E} w_{u,v} \cdot \left(\frac{1 - \vec{y}_u \cdot \vec{y}_v}{2} \right)$ w/ unit vector if same dir = 1, if diff = -1

These are SDPs (semi def prog).

matrix \times constraint

$A \vec{x} = \lambda \vec{x}$ Then λ is eigen value & \vec{x} is eigen vector

we can rewrite $\vec{y}_u \cdot \vec{y}_v$ to find eigen values easily & see if pos/neg.

Gaussian elimination runs in n^3 for n matrix to find eigen values quickly & see if all are positive.

inner product

$A_{uv} = y_u \cdot y_v$

$\begin{bmatrix} 1 & & \\ & \ddots & \\ & & 1 \end{bmatrix}$

pos only.

A_{uv} must have diagonals (1) & non-neg eigen values.
↳ something gives us this eff.

$A = \begin{bmatrix} 1 & & & & \\ & v_2 \cdot u_2 & & & \\ & & \ddots & & \\ & & & v_i \cdot u_i & \\ & & & & \ddots \end{bmatrix}$

$A = v_i \cdot u_i$

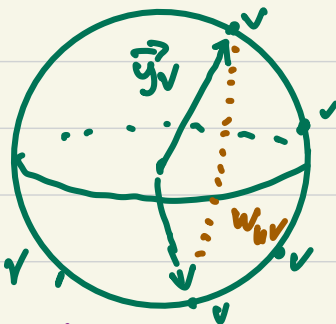
This is SDP and the best approx existing for max-cut.

anything more proves $P=NP$

here it is SDP:

"embed"

Put all vertices onto unit sphere of \mathbb{R}^n dimensions



if weights large then opposite sides of sphere. if opposite sides then split.

CVX solver in python

↳ or also: Random hyperplane rounding algorithm

Take a random hyperplane to cut the sphere to cut sphere. With pos or neg norm:

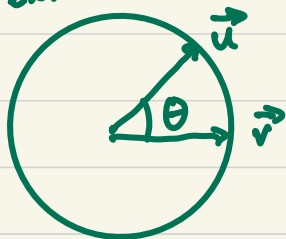
$$\{>0\}$$

$$\{<0\}$$



Algorithm $\geq \sim 0.878$ (optimal)

↳ This algorithm creates ~ 0.878 or better of optimal



Find $\Pr(u \text{ splits from } v)$
↳ $(2 \cdot \theta / 360)$ or $\frac{\theta}{\pi}$

$$w_{uv} \cdot \frac{\theta_{uv}}{\pi}$$

A_{uv} from SDP is $\cos(\theta_{uv})$ so $\frac{w_{uv} \cdot (1 - \cos \theta_{uv})}{2}$

$$W_{uv} \cdot \frac{\Theta_{uv}}{\pi} \geq 0.878 \cdot \left(W_{uv} \cdot \frac{(1 - \cos \Theta_{uv})}{2} \right)$$

$$Pr[i|j] = \frac{\Theta_{ij}}{\pi} \geq 0.878 \cdot \frac{1 - \cos \Theta}{2}$$

Missed loss class, now Oct 13th.

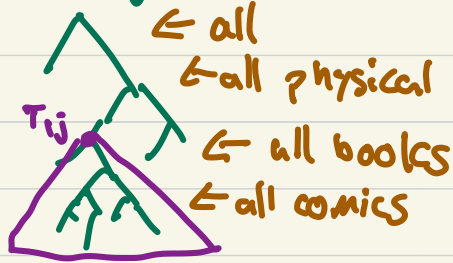
Hierarchical clustering

$$\max \sum_{ij \in E} w_{ij} (n - |T_{ij}|)$$

$n =$ everything (nodes)

outside

$T_{ij} =$ inside tree



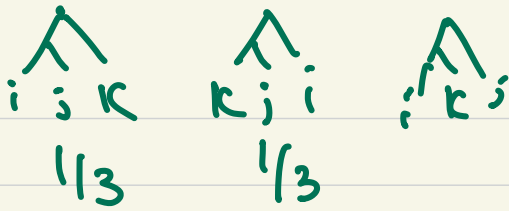
Compared to Max-cut, which is how 2 are separated, Here HC, everything is eventually split.

What is an indicator function? 11

When k is separated from i, j (dolphin, cat, dog)

$$(n - |T_{ij}|) = \sum_{k \neq i, j} \left(\mathbb{1} \left(k \text{ was first to separate from } i, j, k \right) \right)$$

$$\max \sum_{ij \in E} \sum_{k \neq i, j} \left(\mathbb{1} \left\{ \begin{array}{l} k \text{ was first to} \\ \text{be separated among} \\ i, j, k \end{array} \right\} \right)$$

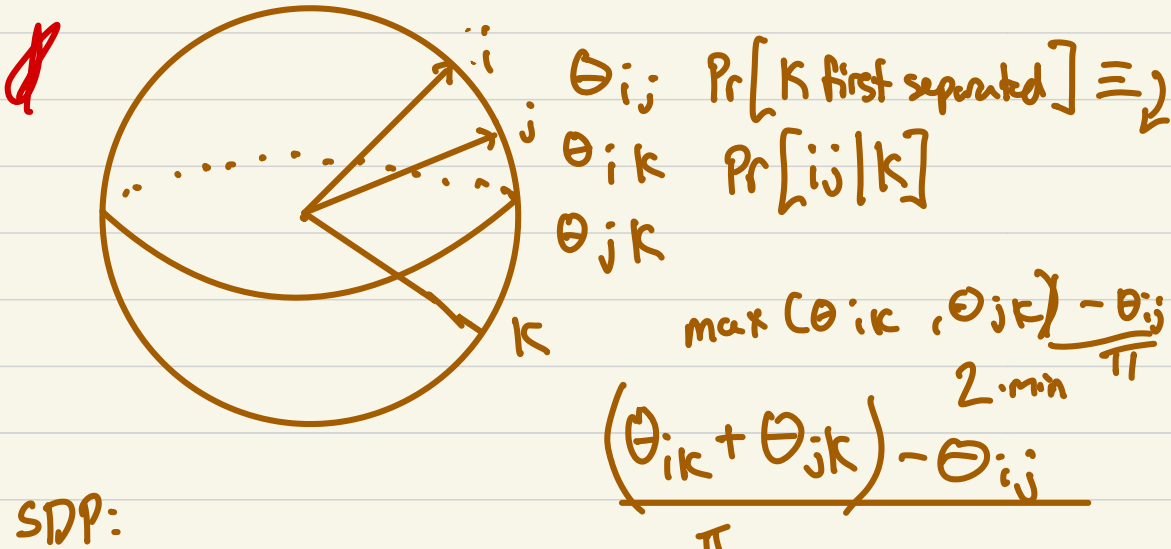


Summation over all weights

$$\text{random} = \frac{1}{3} \sum_{i,j \in E} (n-2)$$

Using similar method to max-cut SDP method

- 1) solve SDP for hierarchical clustering
- 2) Draw hyperplane through origin
- 3) This beats random



SDP:

$$\times \max(\theta_{ij}, \theta_{jk}) - \theta_{ij} / \pi$$

$$\frac{\Theta_{ik} + \Theta_{jk} - \Theta_{ij}}{2\pi} \quad \left[\begin{array}{l} \text{is prob of splitting} \\ k \text{ from } i \text{ \& } j \end{array} \right]$$

$$\left[\begin{array}{l} \Pr[ij|k] = x \\ \Pr[ik|i] = y \\ \Pr[jk|i] = z \end{array} \right. \quad \begin{array}{l} \nearrow ik \text{ got separated} \\ x + y = \frac{\Theta_{ik}}{\pi} \end{array}$$

3 by 3 linear system

$$y + z = \Theta_{ij} / \pi$$

$$x + y = \Theta_{jk} / \pi$$

$$x + z = \Theta_{ik} / \pi$$

Semi-definite Program:

$$\text{SDP : } \max \sum_{i,j \in E} \sum_{t=1}^n w_{ij} (1 - x_{ij}^t)$$

edges

levels

whether x_{ij} separated at level t .

$$x_{ij}^t = \frac{1}{2} \left\| v_i^t - v_j^t \right\|_2^2$$

↳ every vertex (i) has n vectors for each level.

After SDP we get n vectors for each vertex for

each size at most t

level t looks like:



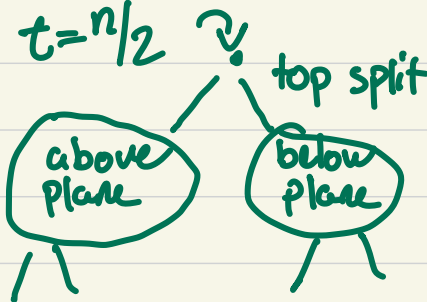
→ all clusters, we've cut

at n , we've cut a ton, and at 1 still a large amount of graph isn't split.

A good candidate is $n/2$ since we have already split a good amount w/o losing structure.

Then taking the $n/2$ ^{→ level} resulting vector from SDP, we run random hyperplane approx. algorithm.

so after [↑] we look at $t = n/2$ [?]



This results in an SDP graph.

cont. again →

This is like still better.

.3336 · Opt

max cut's SDP

Max bisection algorithm is an SDP that is $0.6 \cdot \text{opt}$. each level needs to have exactly half and half resulting output of graph.

1) run SDP 2) hyperplane rounding @ $t = n/2$

The opt soln is either:

$$1) \text{opt} < \underbrace{(1-\epsilon)}_{\substack{\text{some } 80\% \\ \text{or whatever}}} \underbrace{(n-2) \sum_{i,j \in E} w_{ij}}_{\substack{\text{cut cut more than} \\ \text{edges of graph}}}$$

↳

How can we beat $1/3$ approx?

$$\text{Random} = \frac{(n-2) \left(\sum w_{ij} \right) (1-\epsilon)}{3(1-\epsilon)}$$

↳ beats opt in cases where opt is not large
in these cases just choose $1/3$.

2) How about high opt solutions $\text{opt} \geq (1-\epsilon) \sum_{i,j \in E} w_{ij}$
Then we use what we made?

lect ure oct 15

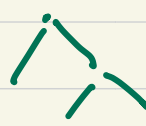
We use objective, SDP, Algos, Geometry, Analysis

Goal is to beat $1/3$ greedy baselines = $1/3$

This soln gets ≥ 0.336 optimal
which proves we can improve

$$G(V, E, w, z_0) \leq \sum_{ij \in E} w_{ij} (n - |T_{ij}|)$$

Equal to $\sum_{ij \in E}$

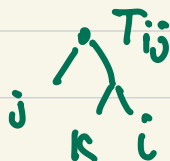


$$\sum_{ij \in E} w_{ij} \leq \sum_{k \neq i, j} \mathbb{1}_{\{k \text{ not a leaf of } T_{ij}\}}$$

only if k got separated first

$$y_{ijk} = \left[w_{ij} \mathbb{1}_{\{k \text{ not a leaf of } T_{ij}\}} \right] y_{ij} = \sum_{k \neq i, j} y_{ijk}$$

random variable



Alg, (k got separated first) $(i, j, k) = \Pr[ij|k] = 1/3$

Random partition recursive 

Alg 2 SDP first random next

- 1) solve SDP for hierarchical clustering
↳ gives vectors for all levels $\{x^t\}$
- 2) Look at vectors at $\lfloor n/2 \rfloor = t^* = 1$.
 $x_{ij}^* = x_{ij}^{t^*}$
- 3) Do the hyperplane rounding for partition (S, \bar{S})
- 4) Run random always (S)

so SDP/hyperplane first, then random always (Alg 1)
is a subroutine to run affr. $\text{Alg}_2 \geq 0.336 \text{ opt}$
 $\text{Alg}_1 \geq 0.393 \text{ opt}$

Max uncut bisection is a more optimal solution.

We have (spreading) and (monotonicity)

x_{ij}^t . if 1, then ij separated @ level t .
level t has at most clusters of size t .

$$\text{SDP obj: } \max \sum_{t=1}^{n-1} \sum_{ij \in E} w_{ij} (1 - x_{ij}^t)$$

spreading: $\sum_{j \neq i} x_{ij}^t \leq t$

$$\sum_{j \neq i} x_{ij}^t \geq n - t, \forall i, \forall t$$

monotonicity: $x_{ij}^{t+1} \leq x_{ij}^t \quad \forall ij \in E, \forall t, x_{ij}^t = 1$

$$\text{SDP obj: } \max \sum_{t=1}^n \sum_{ij \in E} w_{ij} (1 - x_{ij}^t)$$

$$\hookrightarrow x_{ij}^t = \frac{1}{2} \|v_i^t - v_j^t\|_2^2$$

$$\text{OPT} < (1 - \epsilon_1) \left[(n-2) \sum_{ij \in E} w_{ij} \right] \leftarrow w$$

max of all weights (valid upper bound).

$$\text{Random Always} = \frac{1}{3} (n-2) \sum_{ij \in E} w_{ij} = \frac{1}{3} \frac{(1 - \epsilon_1)}{(1 - \epsilon_2)} (w)$$

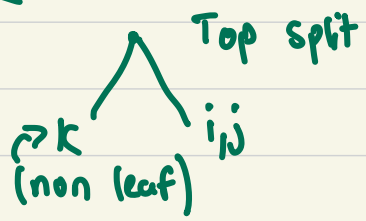
$$\text{so random always} > \frac{1}{3(1 - \epsilon_1)} \text{OPT}$$

now when using: $\text{SDP}_{\text{rel}} \geq \text{OPT} \geq (1 - \epsilon_1) w$

Analysis: Events to analyze: $E_{ij} = ij$ together aft. 1st cut.
 understand top cut? $\rightarrow E_{ijk} = ijk$ together aft. 1st cut
 $E_{ijk} = ij$ together aft 1st cut and k separated

$E[y_{ij|k}] = \frac{w_{ij} \cdot \Pr[\mathcal{E}_{ijk}]}{3} + w_{ij} \cdot \Pr[\mathcal{E}_{ij|k}]$

everyone remains to get w (top split didn't cut) if k is leaf or i or j split first then this is zero



$E[y_{ij|k}] = \frac{w_{ij}}{3} \Pr[\mathcal{E}_{ij}] + \frac{2w_{ij}}{3} \Pr[\mathcal{E}_{ij|k}]$

prob ij not split is $ijk + ij|k$

so:

$E[y_{ij}] = \sum_{k \neq ij} E(y_{ijk}) = \frac{w_{ij}}{3} [(n-2) \Pr[\mathcal{E}_{ij}] + 2 \sum_{k \neq ij} \Pr[\mathcal{E}_{ij|k}]]$

$\Pr[\mathcal{E}_{ij}] = 1 - \theta_{ij} / \pi$

hyperplane

$\alpha_{ij}^b = \cos \theta_{ij}$, the vertices together are

$\theta_{ij} \leq \bar{\theta}$ $\theta = \arccos(1 - \epsilon_2)$

$$\min: \sum_{k \neq j} P_r[\varepsilon_{ij}|k]$$

$$\min: \frac{1}{2\pi} \sum_{k \neq j} \theta_{ik} + \theta_{jk} - \bar{\theta} \geq (n-2) \left(\frac{1}{4} - \frac{\bar{\theta}}{2n} \right)$$

$$\text{Subj to: } \sum_{k \neq i} \cos \theta_{ik} \leq n/2 - 1$$

$$\sum_{k \neq j} \cos \theta_{jk} \leq n/2 - 1$$

20-25 min presentations \rightarrow week 8 or 9
10-15 mins technical, start w/ introduction to prob.
Don't jump directly into technical specifics.

slides for presentation. Present paper, part of proof (in a self contained way).

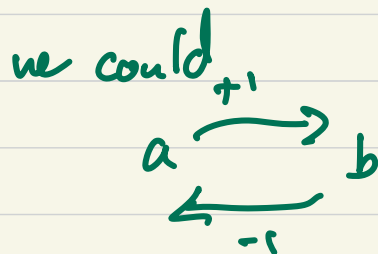
We have a stochastic model:

rankings, correlation clustering, hierarchical clusters

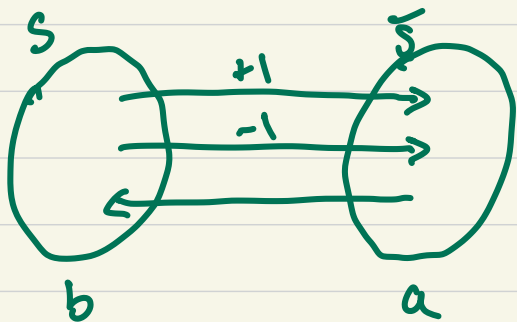
sample 2 items: $a ? b$: $\begin{cases} (1-\epsilon) \text{ correct} \\ \epsilon \text{ wrong} \end{cases}$
which is smaller

3 items: $(a, b, c) \rightarrow 1-\epsilon \text{ right}$
 $\epsilon \text{ (2 wrong answers)}$

For solving $a < b$,
set it up as

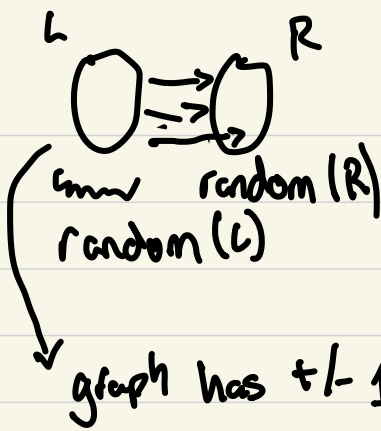


lets setup weighted directed max-cut:



we will only care about left to right edges, then include both weights

Alg: max cut first on s & \bar{s} , then random permutations of s & \bar{s} .



Maximize satisfied constraints is goal, how does $w(L,R)$ get to that?

\rightarrow max # SAT constraints.

we have a total of m constraints,

Beyond
worst
case

m_s is satisfied by first cut (L,R)

m_v violated by L,R

m_u unaffected

$\swarrow -1$ $\searrow +1$

weight of cut $w(L,R) = m_s - m_v$

$$Alg = m_s + \frac{1}{2} m_u$$

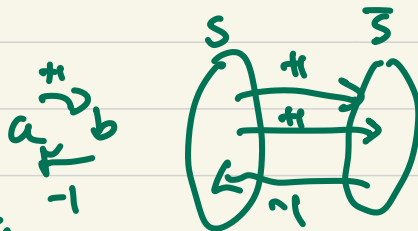
$$= m_s + \frac{1}{2} (m - m_s - m_v)$$

$$= \frac{1}{2} m + \frac{1}{2} m_s - \frac{1}{2} m_v = \frac{m}{2} + \frac{1}{2} w(L,R)$$

weight of any cut in graph
 \nwarrow
NP hard Problem

opt cut \geq median cut $\geq m/2$

How about approx alg?



The graph (total) has \emptyset wgt

We're doing Max-cut w/ Directed & negative weights.

$E(\text{SDP} + \text{hyperplane rounding}) \geq 0.878 \text{ opt}$ (typically)

• one solution, typically bad but good w/ we can add a coeff to make everything positive (no neg weights).

Only for regular max cut,
for directed + neg

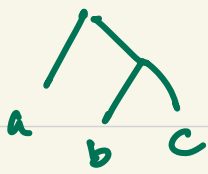
$\rightarrow E(\text{SDP hyper round}) \geq 0.857 \text{ opt} - 0.143 |w|$

if un-directed tm

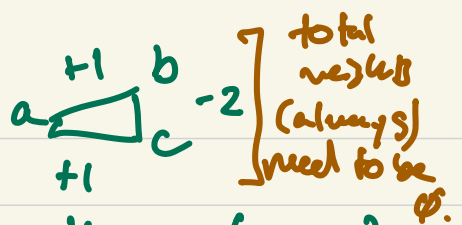
$E(\text{SDP hyp}) = 0.878 \text{ opt} - 0.122 |w|$

$\geq 0.857 \text{ opt} - 0.143 m \geq \frac{0.857 m}{2} - 0.143 m$

$= \frac{0.571 m}{2} = 0.286 m$



For these cases encode as:



Alg should be $\geq \frac{1}{3}m$ (random)

$$w(LR) = 2m_s - m_v$$

$$\text{Alg} = m_s + \frac{1}{3}(m - m_s - m_v)$$

$$= \frac{1}{3}m + \frac{1}{3}(2m_s - m_v)$$

$$= \frac{1}{3}m + \frac{1}{3}w(LR)$$

Nov 5th → wk 6

→ wk 7?

Nov 17th Monday (ian sokan, Adi)

~ 20 minutes with slide decks

Draw graphs too

Show actual self-contained proofs

How Hard is Inference for Structured Prediction

describe in our own words → Vaggos's advisor.

Embeddings, preserve comparison.

$$\delta(x, y) \leq \delta(w, z)$$

This is enough.

we can embed points in \mathbb{R}^d where $d \leq n$
 $\& n/2 \leq d \leq n$

Euclidean norm assumed

$$\| \dots \|_2 = \sqrt{x_1^2 + \dots + x_d^2}$$

$$\|x\|_1 = |x_1| + |x_2| + \dots + |x_d|$$

\hookrightarrow l_1 norm

conv. way to write δ \uparrow $\text{sign}(\vec{x}) \cdot \vec{x}$

$$\| \psi(x) - \psi(y) \|_2 \leq \delta$$

$$\| \psi(w) - \psi(z) \|_2 \leq \delta$$

$$\|x\|_\infty = \max_{i \in [d]} |x_i|$$

L_∞ norm

$$(1, -1) \cdot (x_1, x_2)$$

$\hookrightarrow x_1 - x_2$

\hookrightarrow So to preserve data we need minimum $n/2$ dimensions $\&$ n max dim.

The # ordinal embeddings is less than potential things trying to capture $\&$ ord less than $n/2$

Terminal embeddings
 (ord. embed.) (VIP)

How do we keep only the important embed. / nodes.

$T = \{t_1, t_2, \dots, t_k\}$ Terminal nodes
↳ k terminals

$VIT = \{v_1, v_2, \dots, v_{n-k}\}$
all other nodes

We want to keep T and preserve those our VIT .

↳ $(t, ?) \triangleleft (t', ?)$ We want to compare distances
of this form.
↓ my node
 $t, t' \in T$

Upper bound is k dimensions, these are our VIP nodes (lets say k is small)

if anyone is VIP , you need to preserve every comparison.

How do we preserve key embeddings? → terminal nodes
↳ by putting each one in its own dimension norm w/ L_2

Idea: Every vertex $v(\cdot, \cdot, \cdot, \dots)$ k coordinates
each coordinate is dedicated to a terminal node

terminal 1: $t_1(1, 0, 0 \dots 0)$
 $t_2(0, 1, 0 \dots 0)$

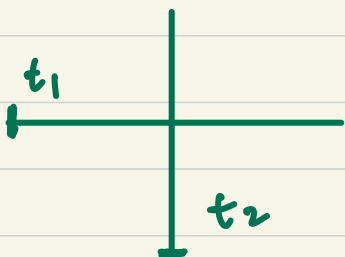
instead of doing 2 do $-M$ (large negative)

$$t_1 (-M, 0, 0, \dots, 0)$$

$$t_2 (0, -M, 0, 0, \dots, 0)$$

Pick $M = k^3 n^2$ (large number)

$t_i = -M \vec{e}_i$ lets say k is pretty small.



(creating embeddings based on 'Rank' of vertex to terminal (1 if closest, 2 if 2nd)

$$\text{rank } r \left\{ \begin{matrix} t_i \\ \downarrow \\ t_i, v \end{matrix} \right\} \in \{1, 2, \dots, k \cdot n\}$$

specifying order in which distance t_i, v appears w.r.t other vertices.

So what is the embedding for a vertex? (not terminal)

$$\varphi(v) = (r(t_1, v), r(t_2, v), \dots, r(t_k, v))$$

compare:

Pair (t, v) & (t', v')

$$\hookrightarrow \left\| \varphi(t) - \varphi(v) \right\|_2^2 = \sum_{i \neq t} r(i, v)^2 + (r(t, v) + M)^2$$

\downarrow $[0, 0, \dots, -M, 0, \dots]$ $i = t$

$$= \sum_i r(i, v)^2 + 2M \cdot r(i, v) + M^2$$

so here

$$\sum_i r(i, v) + 2M \cdot r(i, v) + M^2$$

l_1 = manhattan \rightarrow



l_2 norm = euclidean \rightarrow

Application: Finding l_1 diameter of a set P of points $\in \mathbb{R}^d$

$\hookrightarrow x_1, x_2, \dots, x_n \in \mathbb{R}^d$

Find pair $p, q \in P$ where $\|p - q\|_1 = \max_{p, q} \|p' - q'\|_1$

Naive is checking all diameters w/ distance calc:

(d is small, n is large) Naive: $O(n^2 \cdot d)$

Better: $O(n \cdot d \cdot 2^d)$

each $f_3(p) = \vec{s} \cdot \vec{p}$
 $f(p_1) = [\dots]$

$f(p_2) = [\dots]$

\vdots

$f(p_n) = [\dots]$

$\underbrace{\hspace{10em}}_{2^d \text{ coords}}$

? isometry?

$$\vec{s} = \{ -1, +1 \}^d$$

$$\|p - q\|_1 = \|f(p) - f(q)\|_\infty$$

$$f(p) = \vec{s} \cdot \vec{p}$$

$$L_1 \rightarrow L_\infty$$

isometry from $L_1^d \rightarrow L_\infty^d$

$$f(p_i) = [\dots]$$

Look at only 1 coordinate & do max minus min & compute for all.

$$f(p_n) = [\dots]$$

ordinal embedding:

Taking embeddings from 1 space to another while maintaining connections b/w nodes.

↳ matrix space (euclidean^{dist.} space or tree space)

let $X = ([n], \delta)$ be any matrix space. $\varphi = \text{phi}$

We say $\varphi: X \rightarrow \mathbb{R}^d$ is an ordinal embed. if for every $x, y, z \in X$ we have:

$$\delta(x, y) < \delta(z, w) \Leftrightarrow \|\varphi(x) - \varphi(y)\| < \|\varphi(w) - \varphi(z)\|$$

Distortion: if 1 dist becomes 17 dist, then 17 is distortion.

$\varphi(x) \xrightarrow{1} \varphi(y)$

$\varphi(z) \xrightarrow{17} \varphi(w)$

Def: Ordinal relaxation:

A relaxation of 10 means a dist mult. by 10 is still less than other distances, but keep & don't mess it up:

$$\alpha \downarrow \delta(i,j) < \delta(k,l) \Rightarrow \delta'(i,j) < \delta'(k,l)$$

'significantly different distances should be preserved'

Some constraints we $n > \text{dimensions} > n/2$ for $X = [n, d]$
? & # of ord embed of d vs # distinct metrics
(2)

if you go from higher dim to low, we want α to be larger to have less constraints

relaxation balances dimensions & points of comparison.

if $\alpha = 1$ then same, higher allows for some info lost.

Theorem tradeoffs a vs d

For every int d , every int n :

\exists Matrix space T on n points

'there is a matrix space'

such that the triplet relaxation

↳ special case (i, j) vs (i, k)

of any ordinal embedding of T into d dimension euclidean space.
matrix space

is atleast:

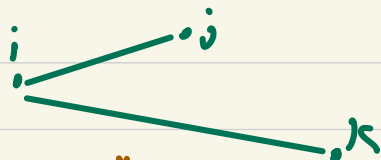
$$\frac{\log n}{\log d + \log(\log n) + \gamma} - 1$$

any embedding will have this loss

'max relaxation'. worst case bound.

↳ getting rid of this is still an open problem?

we want to preserve this relationship: in the new space



we will screw up some embeddings

How do we prove this?

1) we need a family of significantly different metrics spaces

we don't have constraints on close stuff

like w_i $\log(n)$ or something?

2) # ord embed of dim d

\ll # family of sig. diff metrics spaces

Girth G of a graph is shortest cycle



edges in shortest cycle, here is 4. A triangle is 3.

in a pentagon S



if it has a triangle in it

then 3:



it's important to construct high girth graphs to solve/make this.



Delete edges to make a big family of high girth graphs. Like before A to B

was 1, after edge gone, then $g-1$.

it doesn't have to be a, b it could be x, y .

we count probability of getting higher girth cycles after cut. we choose graph to be high girth.

vertex nodes, edges

1) Pick high girth graph $G(V, E)$ $|V|=n$

'how dense can you make graph while still being high girth.

$$\rightarrow m = |E| \geq \frac{1}{4} n^{1+1/g}$$

g : girth, E is edges
 $100 \geq \frac{1}{4} 20^{1+1/g}$? \rightarrow shortest cycle.

Pick g : $\frac{\log n}{\log d + \log \log n + 8}$ then $|E| > 16 \cdot n \cdot d \cdot \log n$

(at least $n d \log n$ edges & has high girth)

idk why \rightarrow

2) Subsample edges of G :

large # $N: [G_1, G_2, \dots, G_N]$
 ← this property we're making
 (*) $\forall G_i, G_j: \exists v \in V: E(v) \setminus E(v) \neq \emptyset$ and $E_{G_i}(v) \setminus E_{G_j}(v) \neq \emptyset$
 ↳ for any pair \leftarrow true exists

what this means is when is v unhappy if we go from G_i to G_j :



How many G do we choose? too high is hard for ↑
 to low doesn't guarantee:

odd embed of dim d \ll # family of sig. diff metrics spaces

so we sample $1/2$ of each edge $G_1: m$
 $G_2: m$

N pick $N = 2^{bm}$, for $b < 1/2 \log_2 4/3$
 G_N

Proof: simplifying assumptions:

- 1) K -regular graph G

G	G_1	G_2
$v \setminus 2$	$v \setminus 3$	$v \setminus 2$
$v \setminus 3$	v	v
$v \setminus 4$		
- 2) independence on v

being a witness for C_{ij} : C_{ij} being for any? α

Ordinal embeddings $\rightarrow n$ points

$$\text{distance}(a,b) < \text{distance}(a,c)$$

emb $F \in \mathbb{R}^d$

$$\|F(a) - F(b)\| < \|F(a) - F(c)\|$$

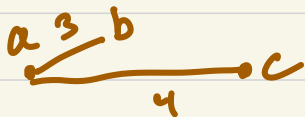
lets take care of contrastive triplets. m triplets

(anchor, pos, neg)

constraints: m triplets of this form

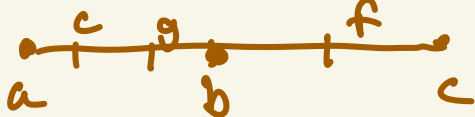
Realizable instances: no paradoxes like $a > b > c > a$
 \rightarrow Focus on this.

Goal: do this embeddings for all triplets w/ dim $O(\sqrt{m})$
we can always do w/ $O(m)$ or $O(n)$ so how?



1) Construct a graph: Tells how ^{points} triplets are related to each other. So if I have \dots then add a point, I need to ensure constraints are still related.

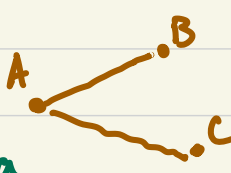
$(a, b, c) \rightarrow$ embed points
 (e, g, f)



B is fixed, since we need to ensure points are not violated (realizable instances).

$\| \cdot \|_2 \rightarrow$ euclidean.

1) (a, b, c) construct a

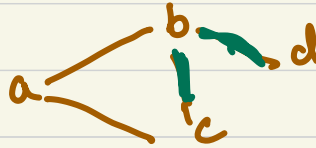


undirected & unweighted

dependency graph, set of n vertices.

Then place in (b, c, d)

bc, bd



Then place in (a, b, d)

ab, ad

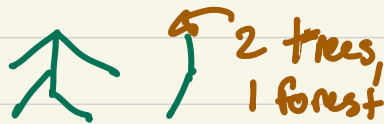


\rightarrow edges

$|E| = 2m$ edges. minimum

Arboricity of G : The \wedge # of forests in which edges can be partitioned (density).

Forest - disconnected tree.

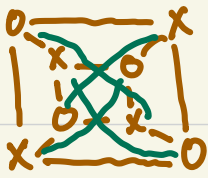


2 trees, 1 forest

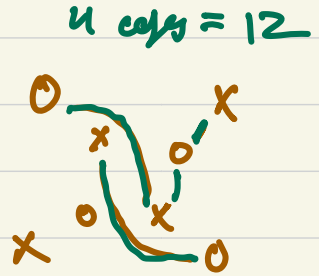
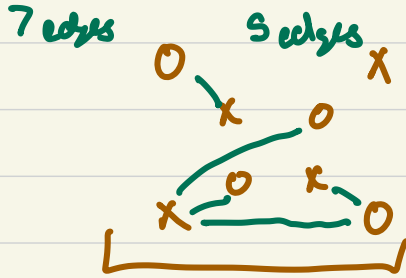
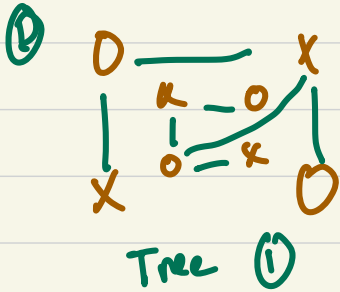
if we get arboricity of q , then $\text{emb } F \in \mathbb{R}^{d=4q}$

Example: Tree $r=1$ (how many trees to overlay to get).
 cycle $r=2$ \rightarrow $(+)$

The bipartite klick of 4,4 nodes: $K_{4,4}$
 klick on circles vs x's.



$K_{4,4}$
 \hookrightarrow at most 7 edges so
 $|Tree| \leq 7$



Does not have to be connected, can be 2 trees or 1 forest (acyclic graph)


Tree can have ≤ 7 edges or else it would be cyclic. We want a forest which is okay being disconnected, but not cyclic

if aboricity low, then tree like, high means likely cycles or dense.

Aboricity: maximize s $\left\{ \frac{|\text{Edges in subgraph } s|}{|\text{Vertices in subgraph } s| - 1} \right\}$

$s =$ subgraphs of G

$$\max_s \left\{ \frac{|E_s|}{|V_s| - 1} \right\}$$

Colorable:  \leftarrow 2 colorable (bc not cyclic).

Facts: 1) G of arb. $r \rightarrow$ is $2r$ colorable ↗

$$2) r \leq 2 \sqrt{|E_G|} \leq O(\sqrt{m})$$

3) \exists ordering $x_1, x_2, x_3, \dots, x_n$

There exists a ↙ such that every node has at most

$$|N^-(x_i)| \leq 2r-1, \forall i$$

↳ backwards order

where every previous node has at most $(2r-1)$ adjacency when looking back

$$\|F(a) - F(b)\|_2^2 < \|F(a) - F(c)\|_2^2$$

↓

$$\|F(a)\|_2^2 + \|F(b)\|_2^2 - 2F(a)F(b) < \|F(a)\|_2^2 + \|F(c)\|_2^2 - 2F(a)F(c)$$

1) Construct a graph ↗ 2) Linear algebra trick

By setting to unit circle then

$\|F(b)\|_2^2 \rightarrow$ goes to 1, same with $\|F(c)\|_2^2$

$$1 - 2F(a)F(b) < 1 - 2F(a)F(c)$$

$$(x_1, y_1) \cdot (x_2, y_2) = x_1 x_2 + y_1 y_2 \quad \curvearrowright$$

$$F(a) \cdot F(b) \leq \underbrace{F(a) \cdot F(c)}_{= \binom{n}{2}}$$

I got lost.

$$F: V \rightarrow \mathbb{R}^{4r} : F(x) = \begin{pmatrix} \overset{1}{x} & \overset{0}{x} \\ \underset{2r}{x} & \underset{2r}{x} \end{pmatrix}$$

if dependency graph:

$$\hat{x} \cdot \hat{y} \approx \text{rank of distance } d(x, y)$$

Order $\{x, y\}$ pairs in descending order, in terms of ^{distance}

if a linear system has no solution, it is inconsistent.
 Being linear system, it has a det = 0 and therefore
 adding slight noise to each of the variables (polynomials
 will make a ^(fiddle) solution since it will
 'escape the roots'.

$$x_1 \quad x_2 \quad x_3 \quad x_4 \quad x_5 - x_6$$

inner prod (like 3,4) is given by ranking.

