Approximate Gomory-Hu Tree in ~Max-Flow Time

Jason Li (CMU)

joint work with Debmalya Panigrahi (Duke)

STOC 2021

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This talk: approximations

- Approximate all-pairs mincut in O(n²) time [AKT FOCS'20]
- Nothing better known for GH tree
- This work: (1+E)-approximate GH tree and SSMC in exact max-flow time

Outline

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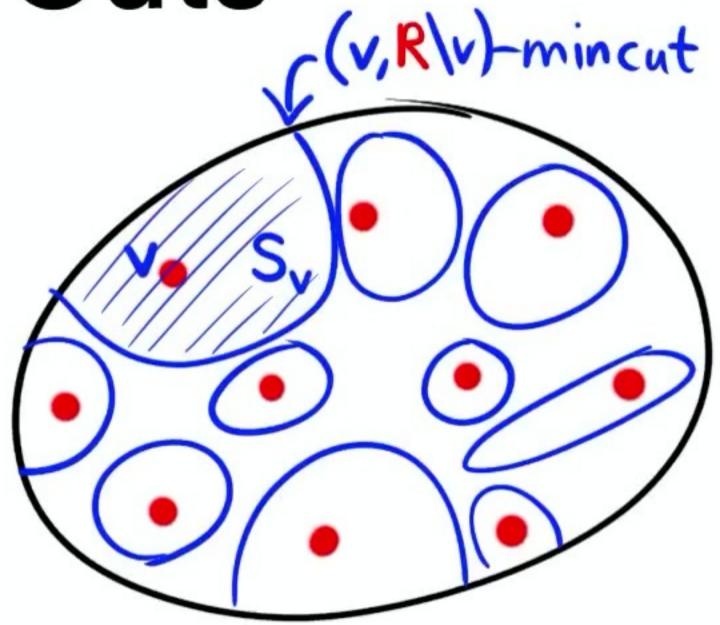
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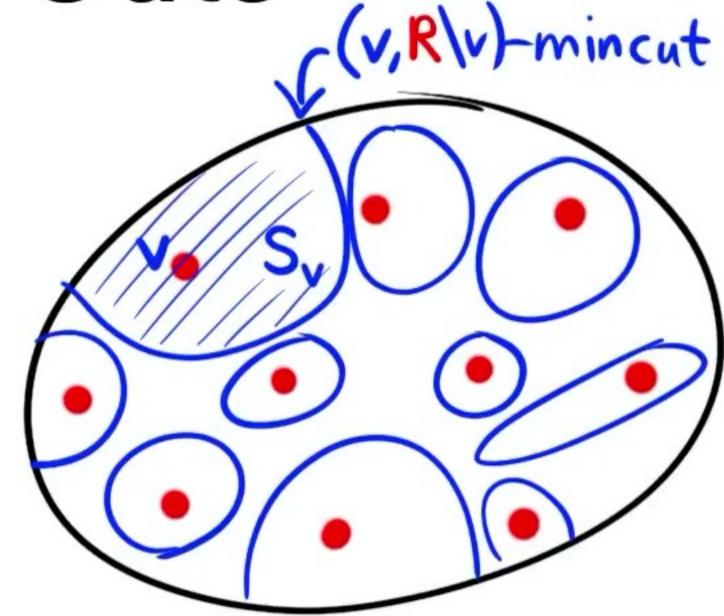
Cut Threshold => approximate single-source mincut (=> approximate all-pairs mincut [AKT20])

Cut Threshold => approximate Gomory-Hu tree

Given a graph and a set R of terminals, find, for each terminal v, the mincut S_v that isolates that terminal



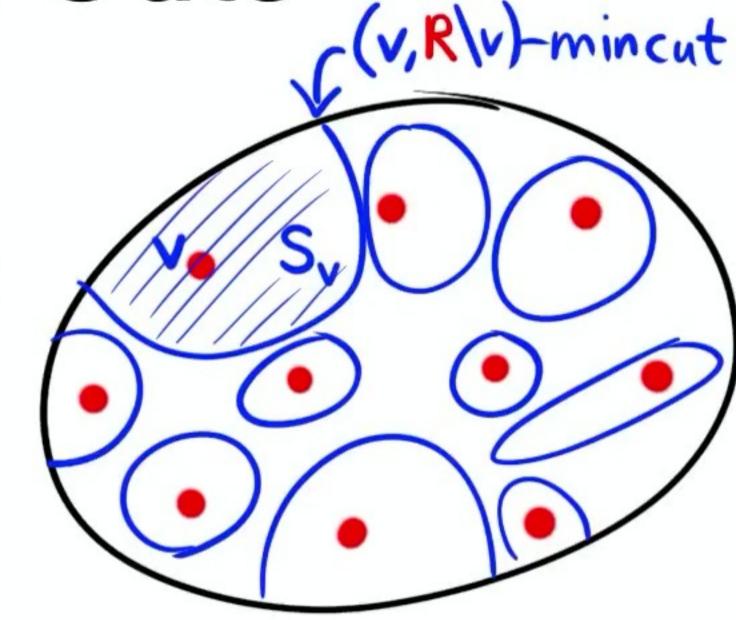
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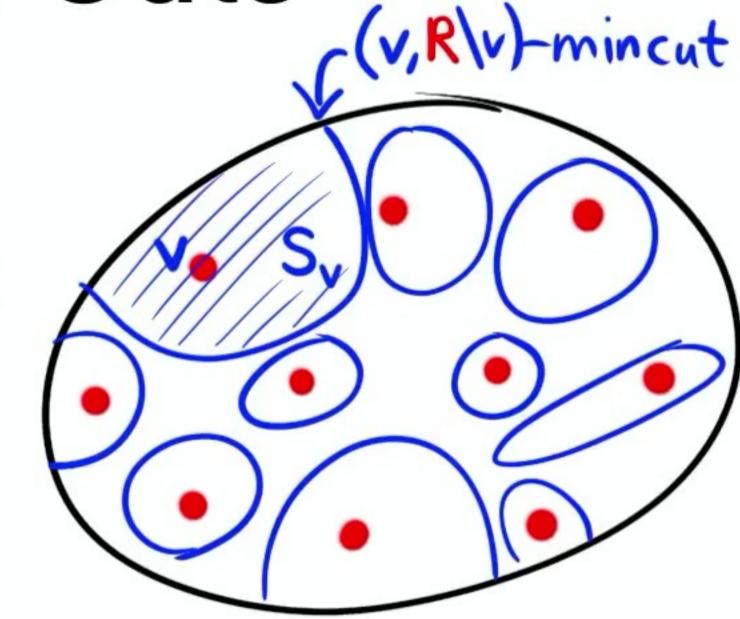


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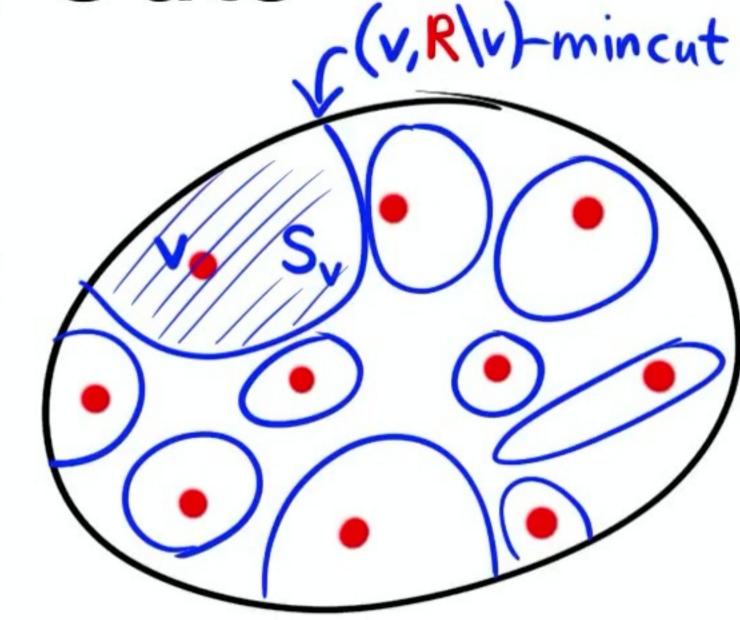


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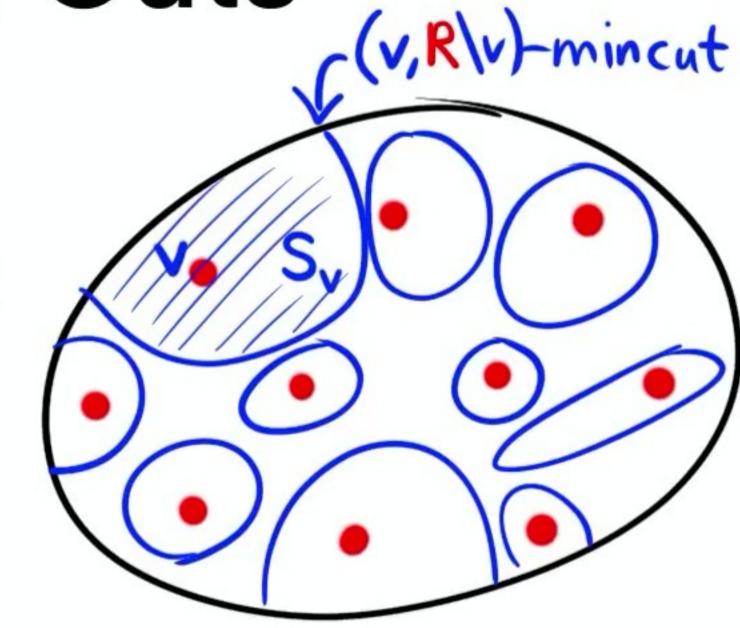


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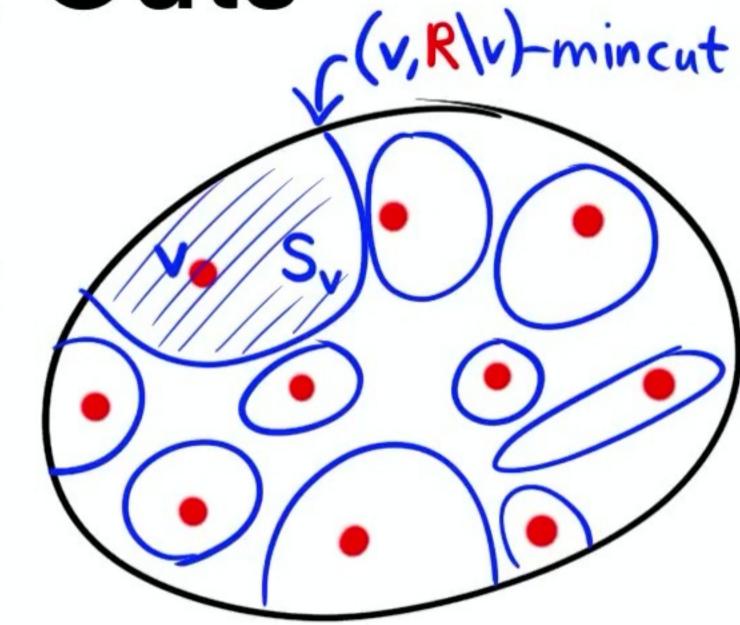
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Warm-up: global Steiner mincut (separate at least two terminals)

Sample at rate 1/2, 1/4, 1/8, ...

If sample at rate ~ \(\frac{1508}{1508}\), then constant prob. success

Given s and λ ("cut threshold"), output all v ∈ V with mincut(s,v) ≤ λ

Given s and λ ("cut threshold"), output all $v \in \mathbb{R}$ with mincut(s,v) $\leq \lambda$

Algorithm CutThresholdStep(G, s, R):

- For each i=1,2,...,logn
 - Sample in R with rate $p=1/2^{I}$ except s is always sampled
 - Compute Minimum Isolating Cuts with sampled set RI
 - For each set $S_v: v \in R^i \setminus s$, if $|\partial s_v| \le \lambda$ then certified mincut(s,u) $\le \lambda$ for all $u \in S_v$

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Algorithm CutThreshold: repeat O(log²n) times: call CutThresholdStep(G,s,R) where R = {still uncertified}

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with prob. ~1/IS∩RI, certify IS∩RI vertices => E[#certify]≈1
over all v certifiable, E[#certify]≈#certifiable

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Claim: Certify $\Omega(1/\log n)$ fraction in expectation

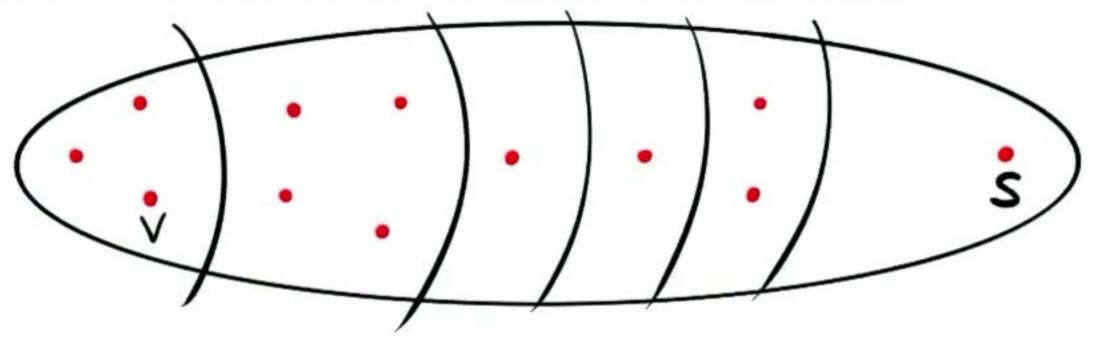
Intuition: fix $v \in R$ certifiable (mincut(s,v) $\leq \lambda$) overcounting? with prob. ~1/IS^RI, certify IS^RI vertices => E[#certify] ≈ 1 over all v certifiable, E[#certify] \approx #certifiable

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Claim: each vertex overcounted O(logn) times w.h.p.

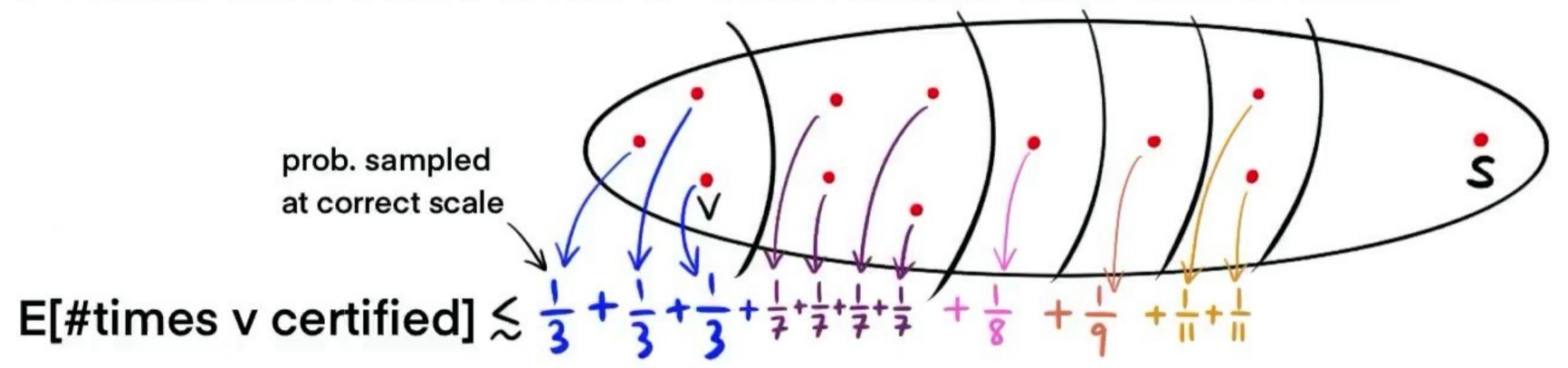
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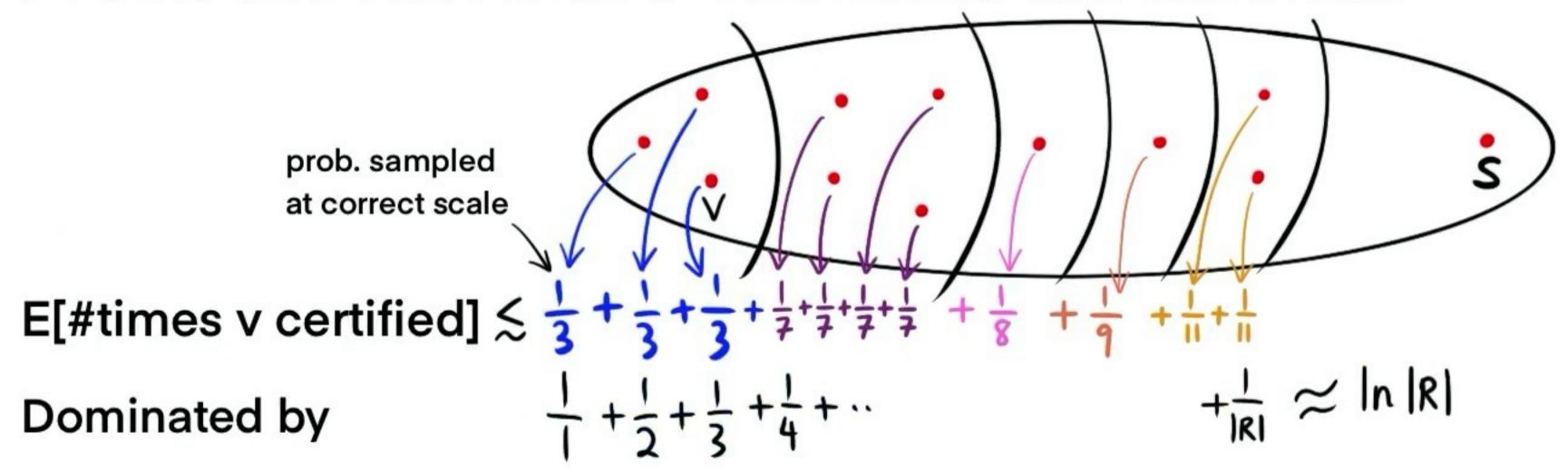
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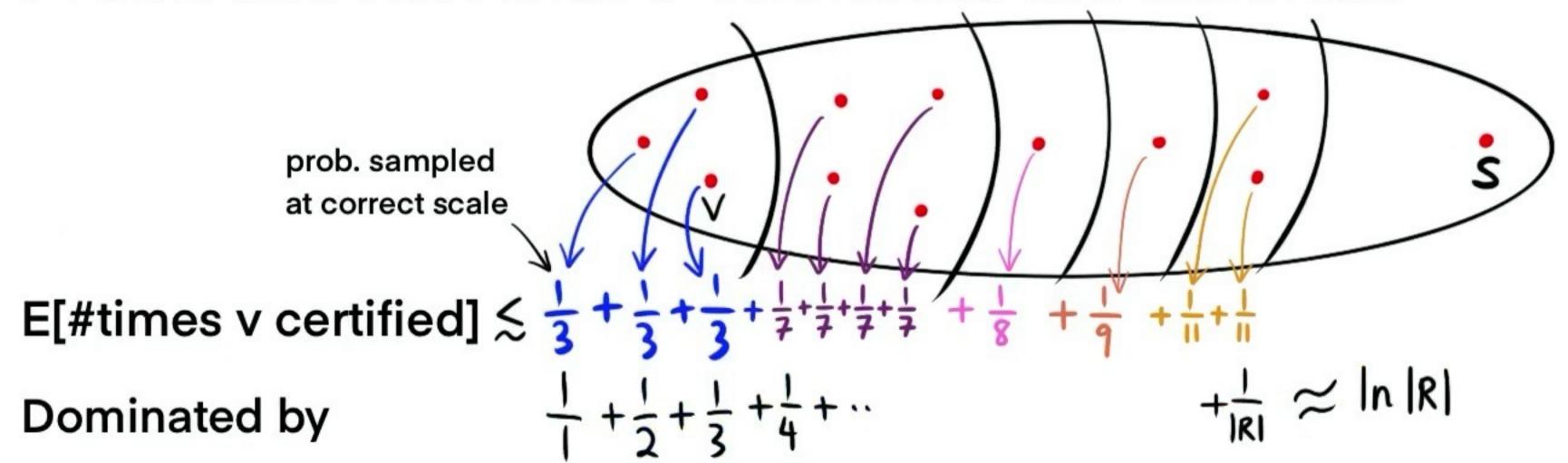
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E[#times v certified] \lesssim O(log IRI). Chernoff bound: O(log n) w.h.p.

Approximate SSMC and GH Tree

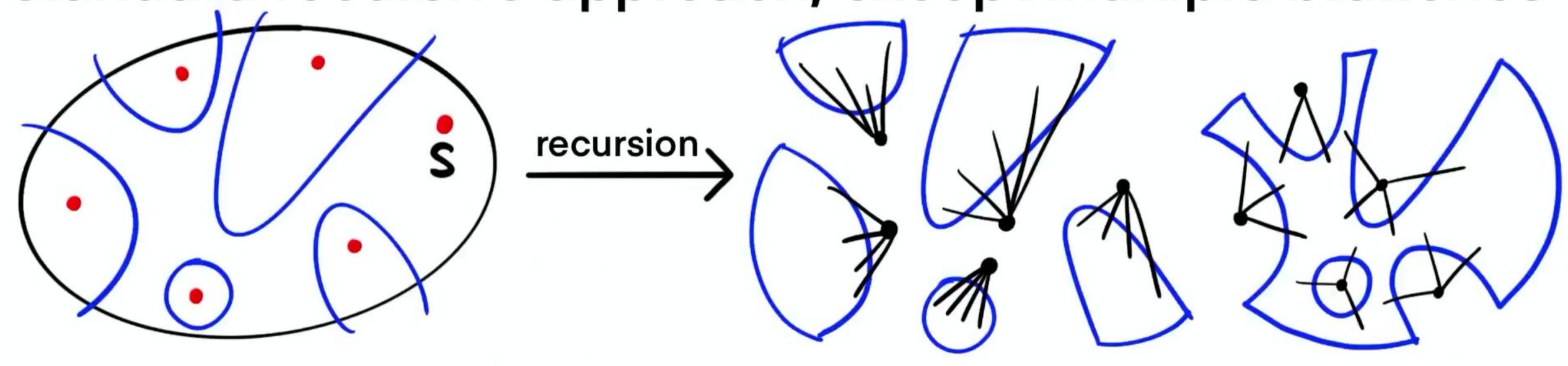
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Approximate SSMC and GH Tree

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standard recursive approach, except multiple branches

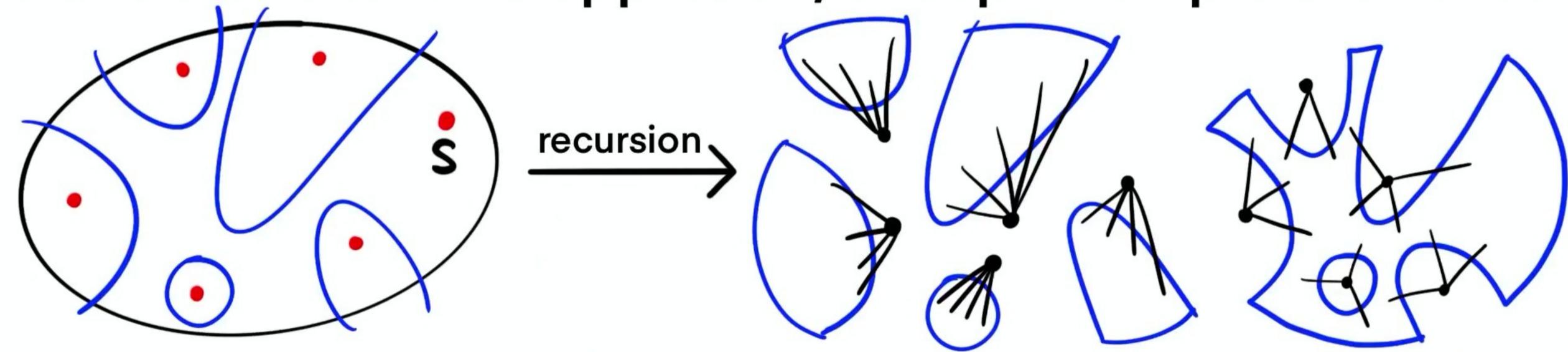


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select s uniformly at random [AKT'20]

Control the approximation factor (nontrivial)

Conclusion and Open Questions

Algorithms for Cut Threshold, (1+8)-SSMC, (1+8)-GHTree in ~max-flow time

Main tools: minimum isolating cuts + random sampling

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Open questions:

- Faster exact GH tree? Reduces to exact SSMC values
- SSMC values faster than n-1 max-flows?
- Approximate GH tree in near-linear time?
- More applications of Minimum Isolating Cuts?