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Minimum Spanning Tree

DPHPC

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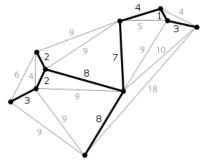
The MST Problem Use cases

Problem definition - reminder



The MST problem

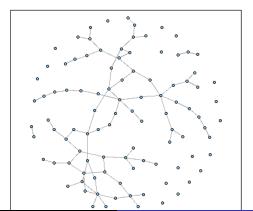
A minimum spanning tree (MST) or minimum weight spanning tree is a subset of the edges of a connected, edge-weighted (un)directed graph that connects all the vertices together, without any cycles and with the minimum possible total edge weight.





Input sets: G(n, p)

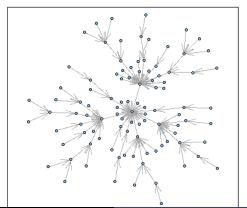
G(100, 0.02)





Input sets: PA(n, m)

PA(100, 1)





Input sets: 9th DIMACS challenge dataset

USA Roads

Name	Description	# nodes	# arcs
USA	Full USA	23,947	,347 58,333,344
CTR	Central USA	14,081	,816 34,292,496
w	Western USA	6,262	,104 15,248,146
E	Eastern USA	3,598	,623 8,778,114
LKS	Great Lakes	2,758	,119 6,885,658
CAL	California and Nevada	1,890	,815 4,657,742
NE	Northeast USA	1,524	,453 3,897,636
NW	Northwest USA	1,207	,945 2,840,208
FLA	Florida	1,070	,376 2,712,798
COL	Colorado	435	,666 1,057,066
BAY	San Francisco Bay Area	321	,270 800,172
NY	New York City	264	,346 733,846



Software Hardware

Setup



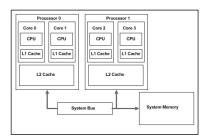
Software

- OMP
- Intel Threading Building Blocks (TBB)
- Intel Parallel Standard Template Library (PSTL)



EULER

- 1 node limitation (OMP)
- 2 sockets filled with 18 cores up to 3.7Ghz
- Inter-sockets bus speed: 10.4 GT/s





ase serial algorithms arallel improvements

Algorithms and parallel implementations



Baselines

- Boost Kruskal
- Boost Prim





Sollin

- For each connected component, find adjacent edge with minimum weight
- 2 Add edge to mst (each edge add at most twice)
- Merge connected components



Kruskal

```
1: A = \emptyset
 2: for all v \in G.V do
      MAKE-SET(v)
 4: end for
5: Sort (asc.) (weight(u, v))_{(u,v)\in G.E}
6: for all (u, v) in G.E ordered by weight do
      if FIND-SET(u) \neq FIND-SET(v) then
 7:
        A = A \cup (u, v)
8:
        UNION(u, v)
      end if
10:
11: end for
```

12: return A

Parallel sorting on Kruskal

m is the number of edges, p is the number of cores, T(m) is the runtime with m edges.

- Sequential: $O(m \ln(m) + m)$
- Parallel sort (Intel TBB): $O\left(\frac{m\ln(m)}{p} + m\right)$



- If $m \le threshold$ then solve with Kruskal
- Find pivot for edges (weight)
- **3** Partition E into $E_{<}, E_{>}$

- \bullet $A_{>} = filterKruskal(E_{>})$
- **?** Return $merge(A_{\leq}, A_{>})$



- **1** If $m \le threshold$ then solve with Kruskal
- ② Find pivot for edges (weight): O(1)
 - Concurrently sample 256 elements from the list (OpenMP)
 - Sort the list and return the median (TBB)
- **3** Partition E into $E_{\leq}, E_{>}$

- \bullet $A_{>} = filterKruskal(E_{>})$
- **?** Return $merge(A_{\leq}, A_{>})$



- **1** If $m \le threshold$ then solve with Kruskal
- 2 Find pivot for edges (weight): O(1)
- **3** Partition E into $E_{\leq}, E_{>}$: $O\left(\frac{m}{p}\right)$
 - Partition function (Intel Parallel STL)

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- **1** Return $merge(A_{\leq}, A_{>})$



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- **1** If $m \le threshold$ then solve with Kruskal
- 2 Find pivot for edges (weight): O(1)
- **3** Partition E into $E_{\leq}, E_{>}$: $O\left(\frac{m}{p}\right)$

- $\bullet A_{>} = filterKruskal(E_{>}): O(T(\frac{m}{2}))$
 - In practice way less than $\frac{m}{2}$
- **?** Return $merge(A_{\leq}, A_{>})$



- **1** If $m \le threshold$ then solve with Kruskal
- ② Find pivot for edges (weight): O(1)
- **3** Partition E into $E_{\leq}, E_{>}$: $O\left(\frac{m}{p}\right)$
- $A_{\leq} = filterKruskal(E_{\leq}): \ T\left(\frac{m}{2}\right)$

- **?** Return $merge(A_{\leq}, A_{>})$: O(1)



- Worst case: $T(m) \le O\left(\frac{m}{p}\right) + 2T\left(\frac{m}{2}\right)$
- Best case: $T(m) \le O\left(\frac{m}{p}\right) + T\left(\frac{m}{2}\right)$



- Worst case: $T(m) \le O\left(\frac{m}{p}\right) + 2T\left(\frac{m}{2}\right)$ $T(m) = O\left(\frac{m\ln(m)}{p} + m\right)$
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- Best case: $T(m) \le O\left(\frac{m}{p}\right) + T\left(\frac{m}{2}\right)$ $T(m) = O\left(\frac{m}{p} + m\right) = O(m)$
- $O(m) \leq T(m) \leq O\left(\frac{m\ln(m)}{p} + m\right)$



Kruskal vs Sollin

- Same complexity, but Sollin computing steps are more fine-grained
- Sollin is entirely paralelizable
- Filter Kruskal can be adapted to Filter Sollin

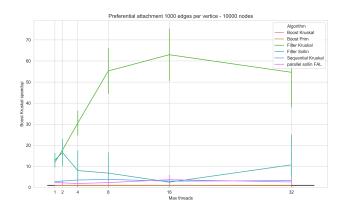


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Results overview

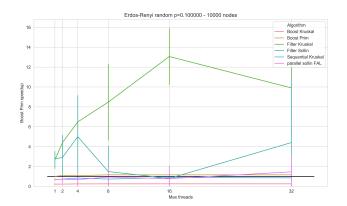


Speedup compared to Boost Kruskal

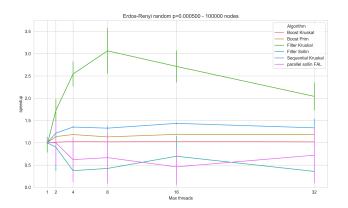




Speedup compared to Boost Prim



Speedups



Amdahl's law

 S_p is the speedup with p cores, f is the part of the program that is sequential.

$$S_p = \frac{1}{\frac{1-f}{p} + f}$$
$$f = \frac{\frac{p}{S_p} - 1}{p - 1}$$



Amdahl's law: Kruskal

Graph: Erdos-Renyi (100,000 nodes, p = 0.0005)

Cores	Median speed-up	Standard deviation	f
1	1	0.0129805395	-
2	1.1881513396	0.0400305172	0.6832872491
4	1.3340592415	0.0130798641	0.6661225318
8	1.3134515984	0.01048841	0.7272602975
16	1.4399618642	0.0088220116	0.6740936918
32	1.3877640643	0.0442081933	0.7115701488



Amdahl's law: Filter Kruskal

Graph: Preferential attachment (10,000 nodes, 1,000 edges per vertex)

Cores	Median speed-up	Standard deviation	f
1	1	0.174182691	-
2	1.6268639022	0.2472258016	0.2293591353
4	2.669168898	0.5213238241	0.1661979381
8	5.3584384444	0.9757941767	0.0704246098
16	5.7447285937	1.108109986	0.1190108026
32	5.6322979489	1.497976882	0.1510166972



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