Minimum Spawning Tree

DPHPC

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- Problem definition
 - Concepts
 - Use cases
- 2 Algorithms
 - Prim
 - Kruskal
 - Borůvka (Sollin)
 - Others
- 3 Environment
- 4 Benchmarking
 - Reference, baseline, tools



Problem definition



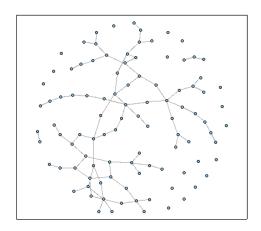
The MST problem



Concepts

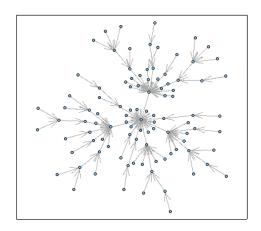


Input sets: G(100, 0.02)





Input sets: PA(100)





Problem definition Algorithms Environment Benchmarking

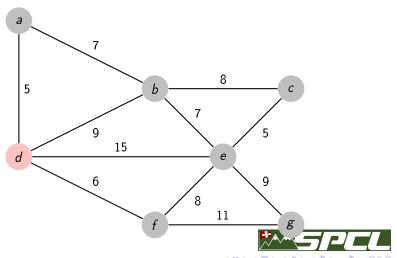
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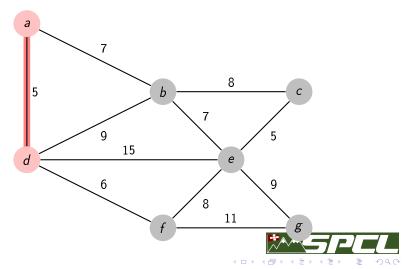
Algorithms

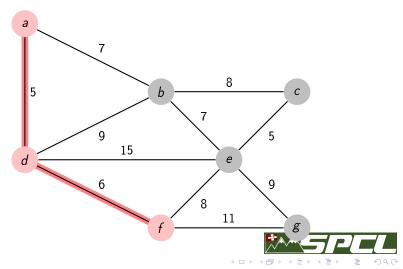


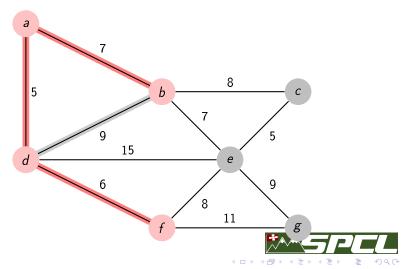
- Initialise a tree with a single random vertex.
- Among all the edges connecting the tree to another vertex, find the minimum-weighted one and transfer it to the tree.
- Repeat until all vertices are in the tree.

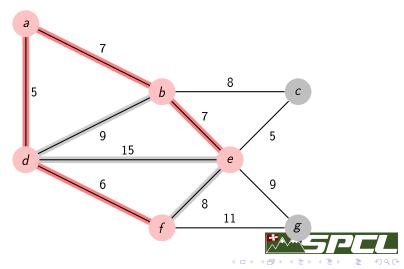


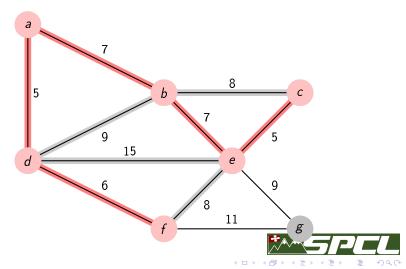


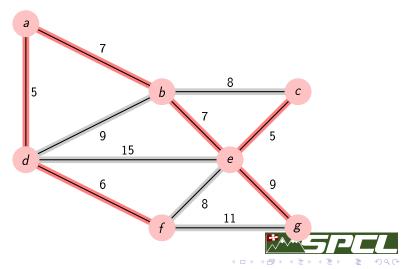












Sequential and parallel Prim

Sequential complexity:

- $O(V^2)$: Adjacency matrix representation
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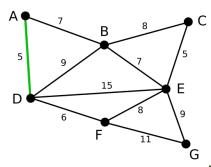
We can parallelize the search of the edge of minimum weight by dividing the vertices and edges between processors to compute local minima.



- Sort all edges by growing weight.
- For each edge: Add it to the MST if it doesn't create a cycle.

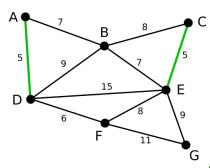


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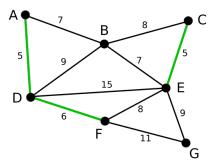


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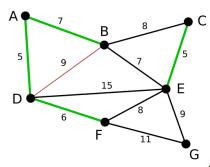


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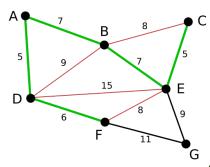


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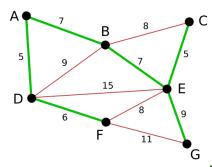


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- Create: Every vertex is in its own component
- Find(x): Find the component of this vertex.
 - If $parent[x] \neq x$, then parent[x] = find(parent[x])
 - ② Return parent[x].
- Union(x, y): Unite two components.
 - parent[find(x)] = find(y).



Sequential and parallel Kruskal

Sequential complexity:

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- O(E) (in practice): For each edge: Add it to the MST if it doesn't create a cycle.



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We can parallelize the sort on $O(\log E)$ processors: O(E) (in practice).



A better parallel approach: Filter-Kruskal

Similar to a quick sort¹:

- \bullet If E < threshold, solve using classical Kruskal
- Choose a pivot (edge)
- **3** Partition in two sets E_{\leq} , $E_{>}$ (weight)
- Recursive call to solve problem with E_{\leq}
- \odot Filter out the edges of $E_{>}$ that connect two vertices of the same component
- **1** Recursive call to solve problem with $E_{>}$



¹Osipov, Sanders, and Singler 2009.

A better parallel approach: Filter-Kruskal

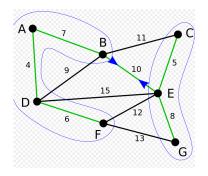
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Better for parallelization since we can distribute the edges for filtering and partitioning.

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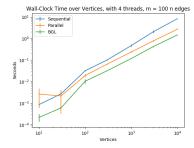
Borůvka (Sollin)



- 1) Init V as independent sets.
- 2) Initialize MST as empty.
- 3) While #sets > 1, do:
 - a) Find closest E from this set to another.
 - b) Add this E to MST if not already added.
- 4) Return MST.



- Prim (Parallel & Seq)
- Kruskal (Parallel & Seq), Kruskal filter
- Sollin (Parallel & Seq)
- Randomization
- Correctness





Problem definition Algorithms Environment Benchmarking

Environment



Architecture



EULER Cluster

Xeon
$$Ex, x \in \{3, 5, 7\}$$

x86_64 architecture

Source: https://scicomp.ethz.ch/wiki/Euler



Tools



- CMake v3.3+
- C++11 GCC v4.9.2+
- OpenMPI (shared memory) v1.6.5+



eference, baseline, tool:

Benchmarking



Tools

- Measures : LibSciBench library
- Interpretation :
 - LibSciBench's R scripts
 - (Custom python scripts)

Ref : https://spcl.inf.ethz.ch/Research/Performance/LibLSB/

Baseline



Borůvka's serial algorithm $O(E \cdot log(V))$

https://en.wikipedia.org/wiki/Otakar_Bor%C5%AFvka





Osipov, V., P. Sanders, and J. Singler (2009). "The Filter-Kruskal Minimum Spanning Tree Algorithm". In: *Proceedings of the Meeting on Algorithm Engineering & Expermiments*. New York, New York: Society for Industrial and Applied Mathematics, pp. 52–61.

