DP_Kruskal: a concurrent algorithm to maintain dynamically minimum spanning trees

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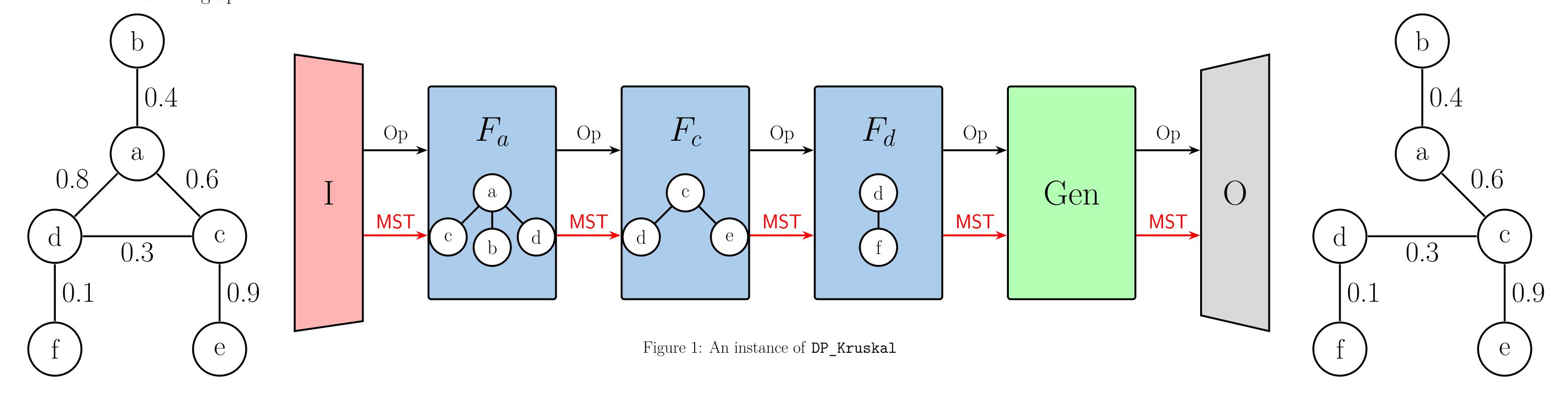


Introduction

We introduce DP_Kruskal: a parallel and concurrent Kruskal-based algorithm, defined on the Dynamic Pipeline Model (DPM) [5], to **solve** the fundamental Minimum Spanning Tree (MST) problem. DP_Kruskal outperforms other algorithms and is particularly effective for dense graphs.

DP_Kruskal

A dynamic pipeline (DP) is a sequence of **connected** (via communication channels) concurrent, stateful stages. DP_Kruskal distributes the input graph G = (V, E)along a DP with event and graph channels carrying events and MSTs, respectively.



Instance of DP_Kruskal

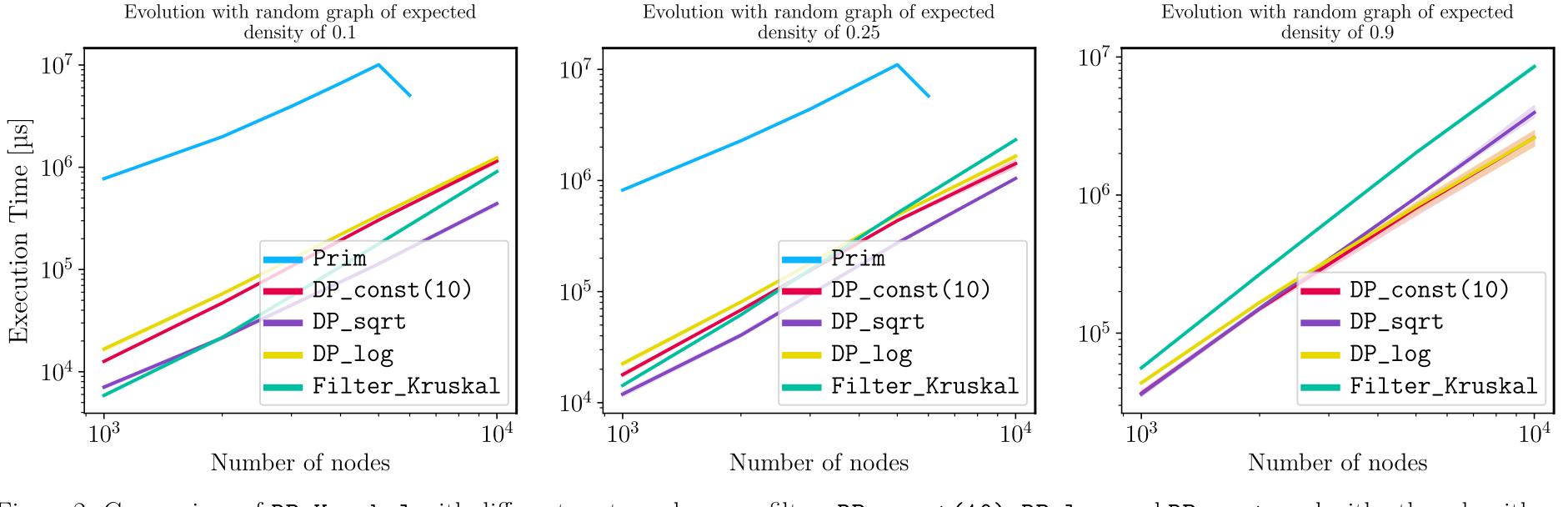
Figure 1 illustrates an instance of DP Kruskal. It consists of:

- Input (I): Receives events of the form (op, e) and passes them through the event channel; if op = mst, sends an empty set via the graph channel.
- Filter (F_v) : v is a vertex, called the root of the filter. When an event $(\mathsf{op}, \mathsf{e})$ arrives: (i) If $\mathsf{op} \in \{\mathsf{insert}, \mathsf{update}, \mathsf{delete}\}$ and e is incident to v, processes the event; otherwise, passes it to the next stage. (ii) If op = mst, computes a partial MST from the union of the arriving MST and the tree in F(v), then passes it to the next stage.
- Generator (Gen): If $op \in \{insert, update\}$ arrives, spawns a new Filter stage F(v), adding it to the pipeline between the last filter and Gen.

Source Code

• Output (O): If op = mst arrives, outputs the MST_G .

Single core and Static random graphs



We compared DP_Kruskal against (i) Filter_Kruskal [4], and (ii) a message-passing Prim [2] implementation on a single core and randomly generated graphs of n vertices studying the **effect** of the number of roots in each filter of DP_Kruskal three options were considered: (a) a constant number of roots DP_const, (b) $\log(n)$ roots DP_log, and (c) \sqrt{n} roots DP_sqrt.

Although for small graphs, Filter_Kruskal is faster; as graph size and density increase, DP_sqrt becomes the best option (see Figure 2).

Figure 2: Comparison of DP_Kruskal with different root numbers per filter: DP_const(10), DP_log, and DP_sqrt; and with other algorithms on a single core over random graphs of n vertices. For each size and probability 20 random graphs were generated

Multi core and Dynamic real graphs

We compare DP_Kruskal and Filter_Kruskal on realistic dynamic graphs obtained from DynGraphLab [1]. We observe that:

- DP_Kruskal is effective for maintaining the MST (See Table 1).
- DP_Kruskal has excellent performance in a parallel environment, showcasing **significant scalability** (See Figure 3).

Dataset	n	op.	Filter_Kruskal	DP_Kruskal
as-caida	31379	119468	1h 30min	1h 19min
movielens10m	49847	384585	1h 39min	1h 20min
simplewiki	100312	889016	17h 8min	11h 29min

Table 1: Time with 1 core and 10 roots per filter

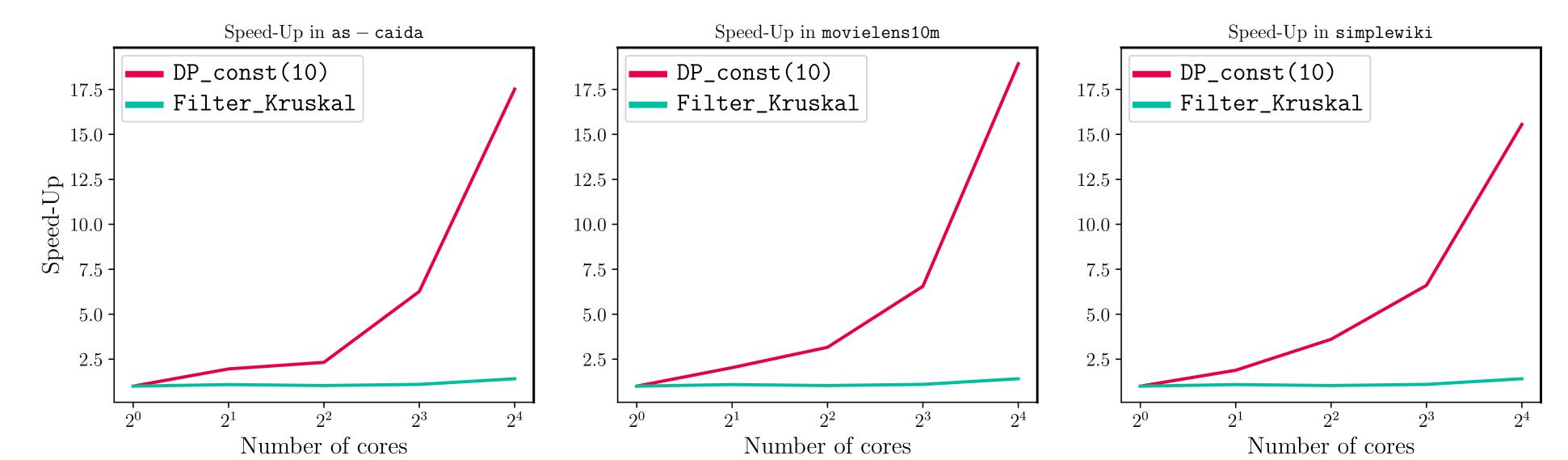


Figure 3: Multicore comparison of speed-ups

Final Remarks

Experiments on random and real graphs demonstrate that the DP_Kruskal algorithm is particularly competitive for dense graphs, showing improved performance over the Filter_Kruskal algorithm for graphs with over $2 \cdot 10^3$ vertices. It also **scales well**, enhancing efficiency and speed up to 16 cores. Future work includes for instance to compare DP_Kruskal against a MapReducebased competitor [3]; to evaluate other MST algorithms within the DP_Kruskal model; to assess different implementation languages; to experiment with larger datasets and other dynamic graph models.

References

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- [2] Lončar, V., Škrbić, S., Balaž, A.: Parallelization of minimum spanning tree algorithms using distributed memory architectures (2014)
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- [4] Osipov, V., Sanders, P., Singler, J.: The filter-kruskal minimum spanning tree algorithm (2009)
- [5] Pasarella, E., Vidal, M.E., Zoltan, C., Royo Sales, J.P.: A computational framework based on the dynamic pipeline approach (2024)