

# Assignment Project Exam Help

Distributed MST

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① Minimum spanning trees

② Prim MST

③ Kruskal MST

④ Borůvka MST

⑤ Discussions – Prim, Kruskal, Borůvka

⑥ Distributed MST (Sync)

⑦ Sync MST – Level 0 Components

⑧ Sync MST – Level 1 Components

⑨ Sync MST – Level 2 Components

⑩ Memento

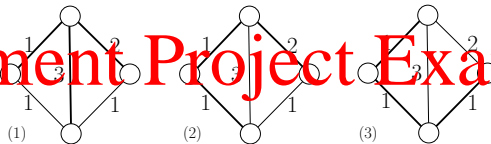
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# Spanning trees (ST)

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For (1) and (2), consider that the top node is the root

- (1) : min-height ST  
here also BFS ST (cf. [sync](#) Echo)
- (2) : shortest paths ST (cf. [sync](#)/[async](#) Bellman-Ford)
- (3) : minimum ST (cf. [sync](#)/[async](#) GHS)  
here also DFS ST (cf. [sync](#)/[async](#) Cidon)
- (1,2,3,...) : arbitrary ST (cf. [async](#) Echo)

# Minimum spanning tree (MST) algorithms

- If edges have different weights, then there is a **unique** MST

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- Prim (1957), Jarník (1930), Dijkstra (1959):  $O(M \log N)$  or  $O(M + N \log N)$
- Kruskal (1956):  $O(M \log N)$ ; Reverse-Kruskal:  $O(M \log N \dots)$
- Borůvka (1926); Choquet (1936); Fredrik, Łukasiewicz, Perkal, Steinhaus, and Zubrzycki (1951); Sollin (1965)<sup>1</sup>:  $O(M \log N)$
- faster algorithms – almost linear: Chazelle (2000); randomization: integer weights; ...
- Distributed MST (**sync, async**): GHS - Gallager, Humblet and Spira (1983):  $O(N \log N)$ ; improvements linear  $O(N)$ ; or even sub-linear

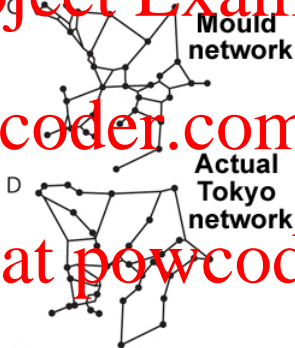
<sup>1</sup> “Because Sollin was the only computer scientist in this list living in an English speaking country, this algorithm is frequently called Sollin’s algorithm” (Wiki)

## Side-bar: Minimum spanning networks in biology

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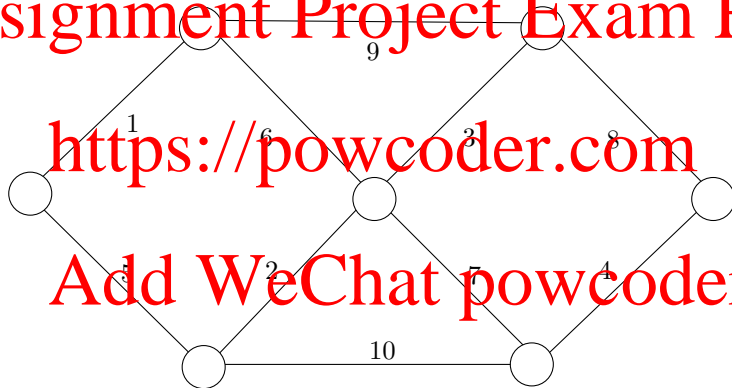


## Prim

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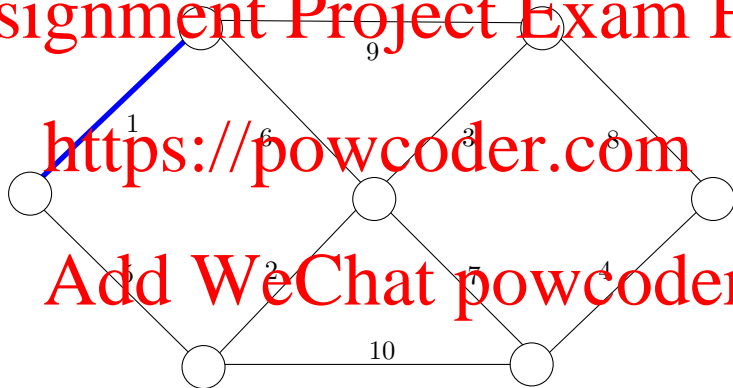


# Prim

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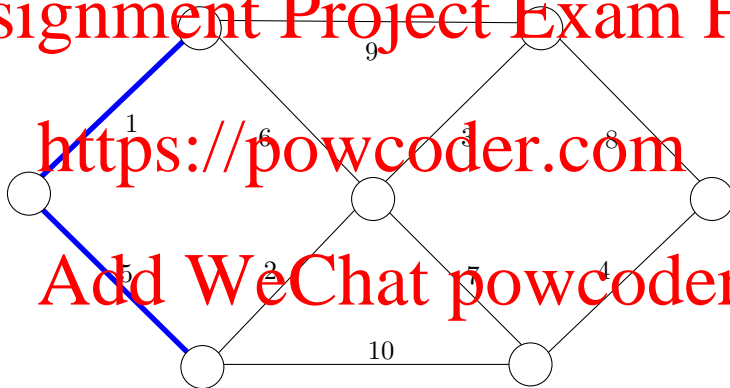


## Prim

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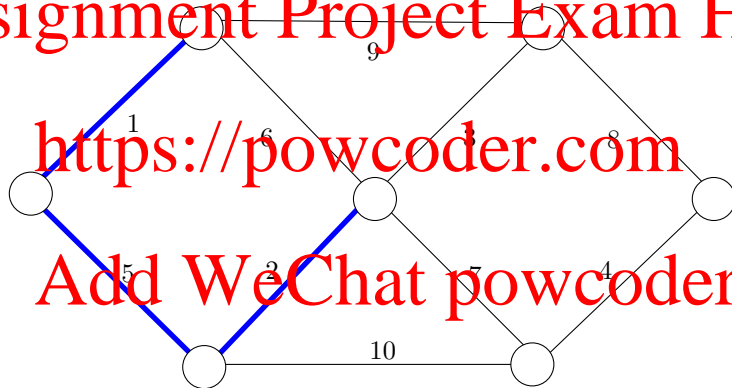


# Prim

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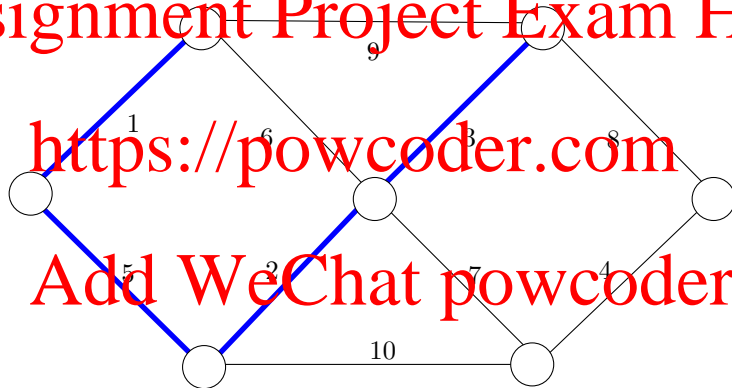


# Prim

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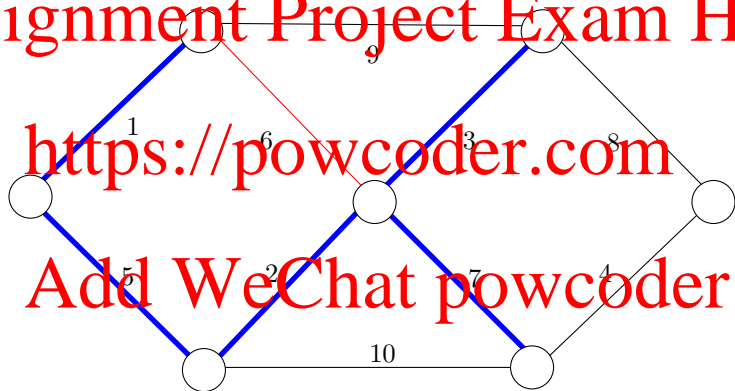


## Prim

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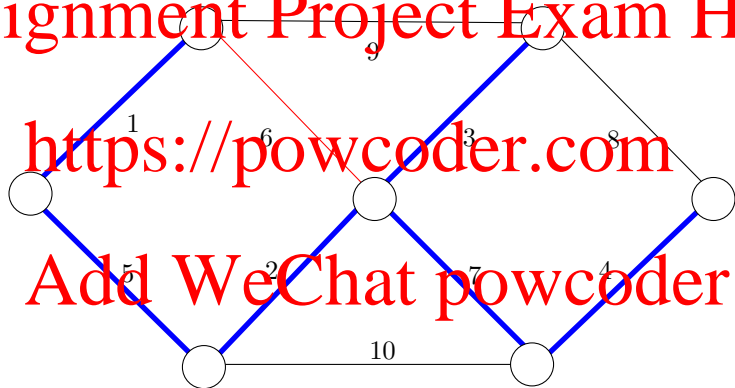


## Prim

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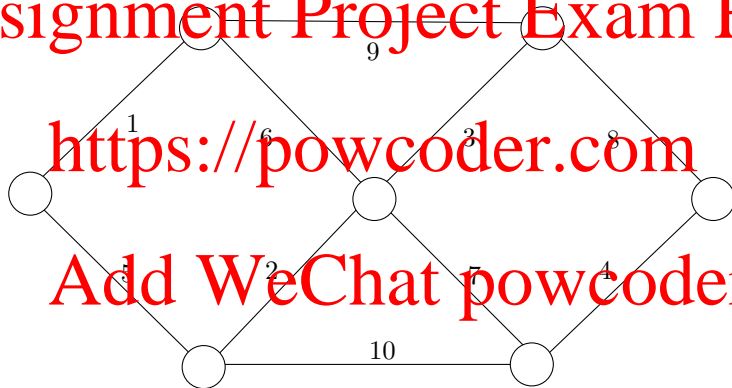


# Kruskal

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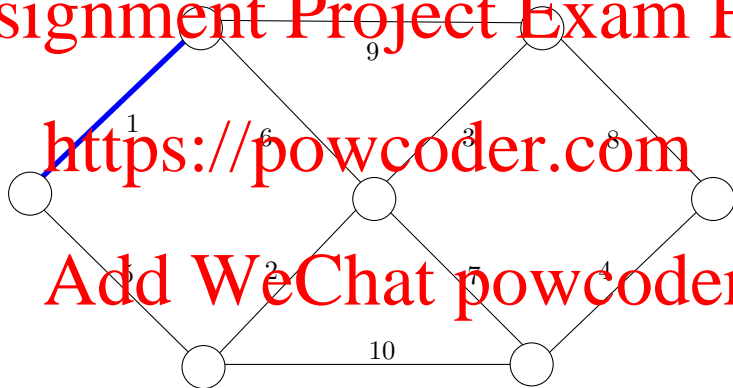


# Kruskal

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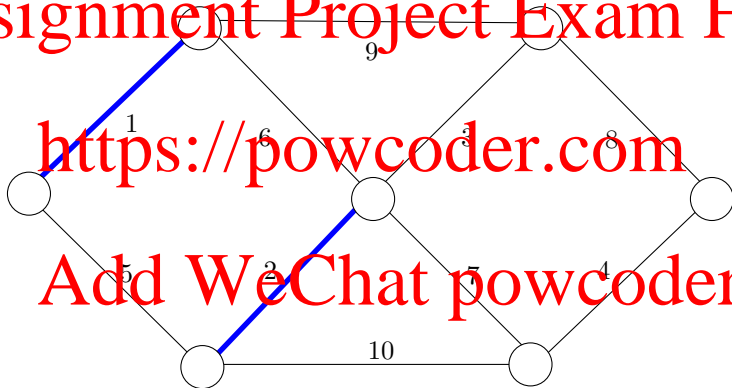


# Kruskal

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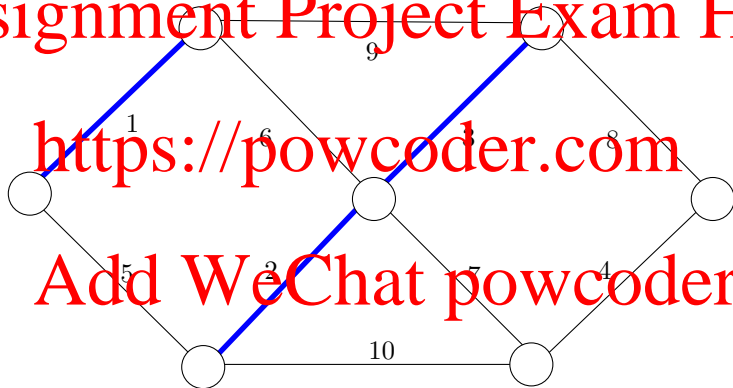


## Kruskal

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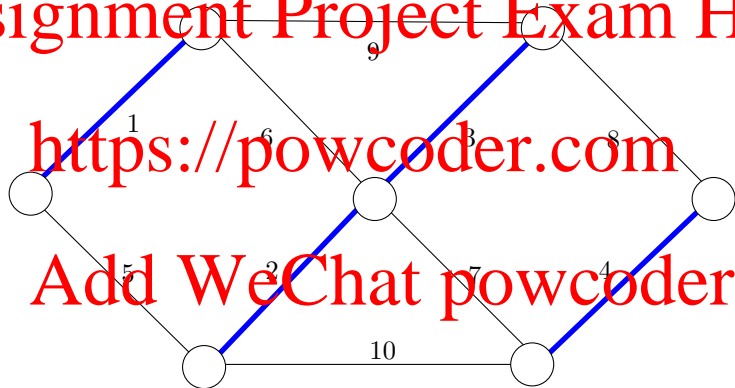


# Kruskal

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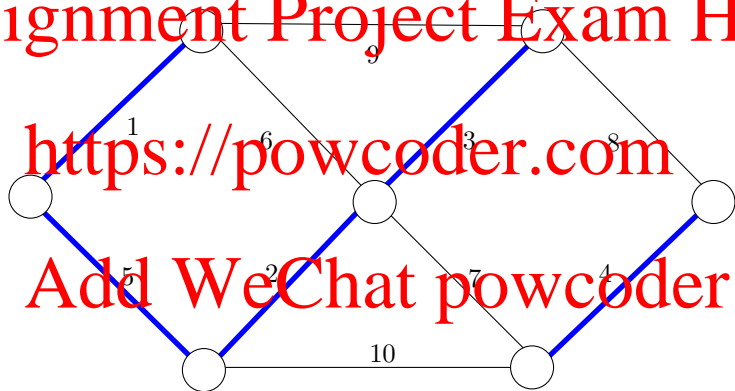


## Kruskal

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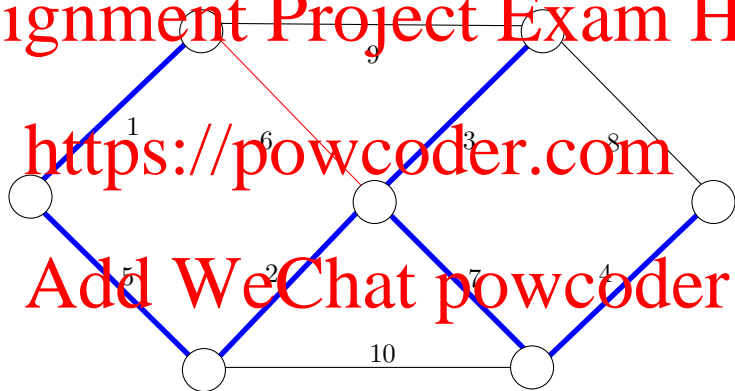


## Kruskal

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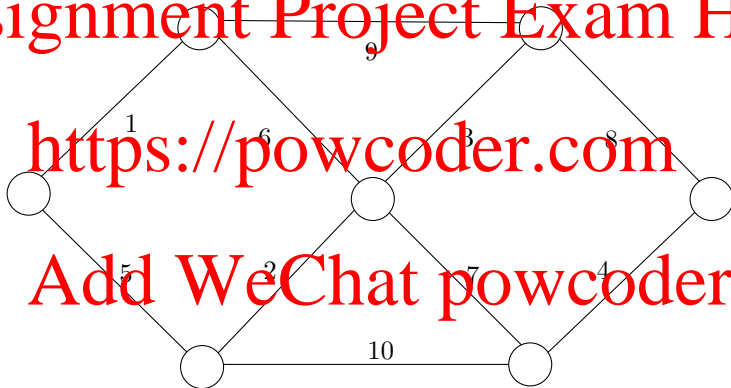


Borůvka – red: Common MWOE (min-weight out edge)

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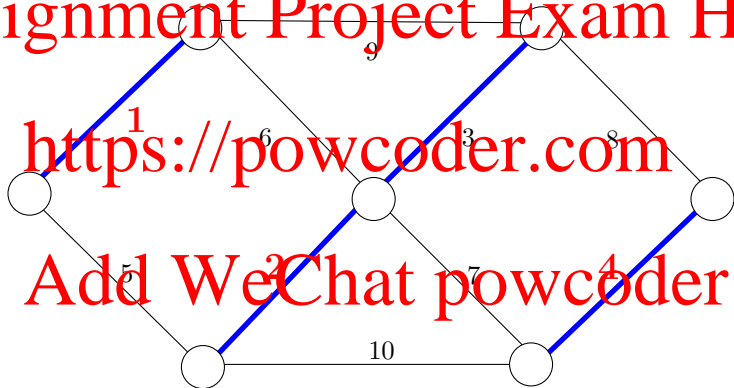


Borůvka – **red**: Common MWOE (min-weight out edge)

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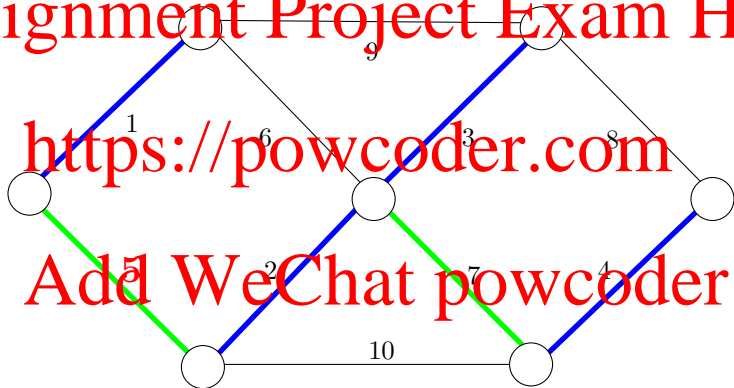


Borůvka – red: Common MWOE (min-weight out edge)

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# Prim and Kruskal as particular cases of Boruvka

- **Kruskal** is a restricted Boruvka, where we only use the **overall lowest-cost MWOE** (not necessarily common) - thus, at any given step, we only merge 2 trees (one single 2-way merge)
- **Prim** is a restricted Boruvka, where we only use the **MWOE of the chosen tree** - thus, at any given step, we merge the chosen tree with one node (a "level 0" tree)
- **Boruvka** deals with **all MWOEs** at the same "step" (called "level" in the distributed case) - thus, we can perform several multi-way merges "simultaneously" (at the same "step")
- Quotation marks indicate that many things may happen at the same "step" or "level" ...
- The distributed MST versions exploit this ability of Boruvka

## All unrooted trees (edge shapes) with 4 & 5 nodes

- Looking for MWEs (minimal working examples) where Kruskal is essentially different from Borůvka: 4 or 5 nodes?

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- Here the selected roots have minimum eccentricities
- WLG (without loss of generality), are these all that need consideration?



# Five nodes – Kruskal essentially different from Boruvka?

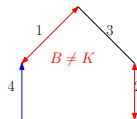
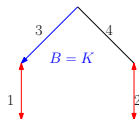
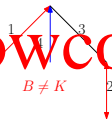
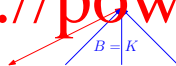
- B step 1: collects all red (Common MWOE) and blue (one way MWOE) edges

- K collects in order: red edges (costs 1, 2), then costs 3, 4

- Figure codes:  $B \neq K$  : essential differences,  $B = K$  : NO essential differences

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## Four nodes

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- On 4 nodes, Kruskal is NOT essentially different from Boruvka

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## Further discussions

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Prove, for Borůvka's algorithm:

- In any multi-way merge there is always one Common MWOE
- The Common MWOE is unique

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Prove that these hold for:

- Level 0 trees, i.e. initial nodes (size 1 trees)
- Level  $k$  trees, for any  $k \geq 0$

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≡!

# Distributed MST (Sync)

- Based on Borůvka; see Lynch §4.4

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- Nodes have **unique** IDs

- Nodes know their **adjacent neighbours**

- Nodes know the **graph size,  $N$**  (required for synchronisation)

- could be obtained by a preliminary phase, based on **Echo**

- Edges have **unique** weights or same weight edges can be ordered

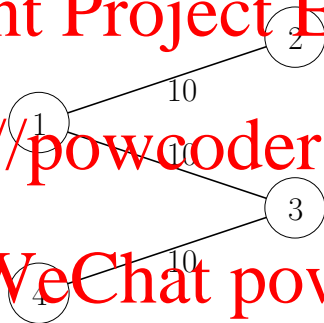
- e.g, by **lexicographical** comparisons, where each edge  $\{i, j\}$  is represented by an ordered triple  $(w, v, v')$ ,  $v < v'$ , where  $w$  = edge weight,  $v, v'$  = ID's of  $i, j$

# Sync MST – Comparing weights with equal weights

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$$\{1, 2\} = (10, 1, 2) < \{1, 3\} = (10, 1, 3) < \{4, 3\} = (10, 3, 4)$$

## Distributed MST (Sync)

- Time complexity:  $O(N \log N)$

Message complexity:  $O((N + M) \log N)$

- Levels:**  $O(\log N)$ ; each level defines a **spanning forest**
- Level 0 components are the individual nodes
- Level  $k+1$  builds larger components by merging 2 or more level  $k$  components into new components
- Thus, level  $k \geq 0$  has component subtrees of size  $2^k$  at least
- Each component has a distinguished **leader**; the leader ID identifies the component
- For connected graphs, the algorithm ends with a **MST** (unique, if edges have different weights)

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## Distributed MST (Sync)

- Details may vary, with slightly different performance...

The following diagrams use a set of suggestive, but not exactly necessary, messages

- *initiate*

- *test*

- *accept, reject*

- *report*

- *connect (implies a change-root), connect!*

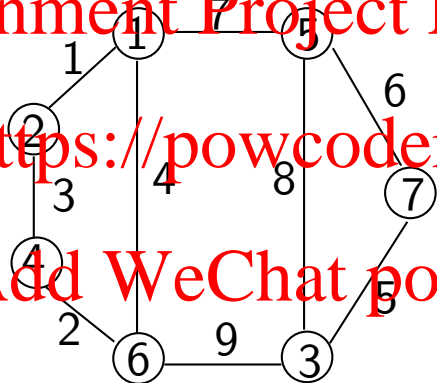
- To ensure correct component identification, each level  $k > 1$  takes a predefined number of steps,  $O(N)$  – nodes may need to stay idle until this count is completed
  - depending on the actual algorithm details, this may happen in different ways

## Sync MST – Level 0 Components

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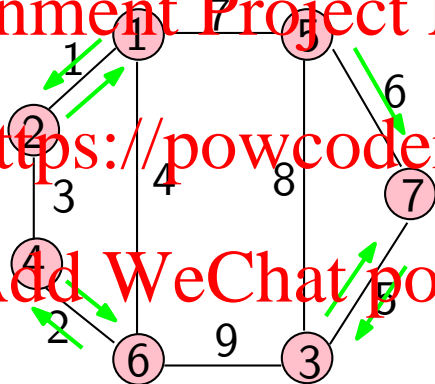


## Sync MST – Level 0 Components

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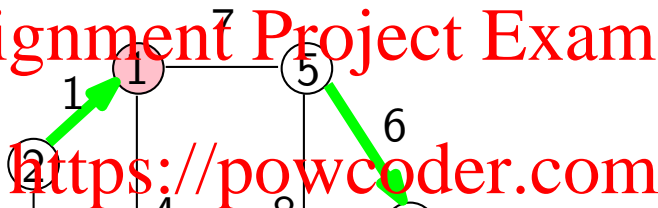
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→ connect!

## Sync MST – Level 1 Components

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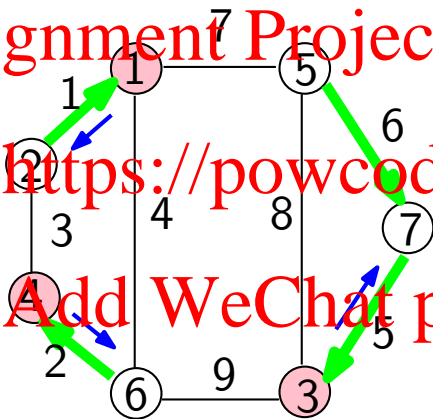
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## Sync MST – Level 1 Components

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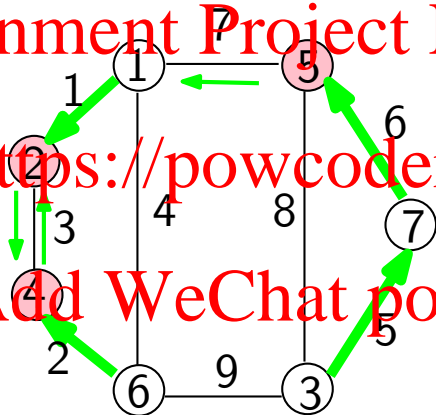
→ *initiate*

## Sync MST – Level 1 Components

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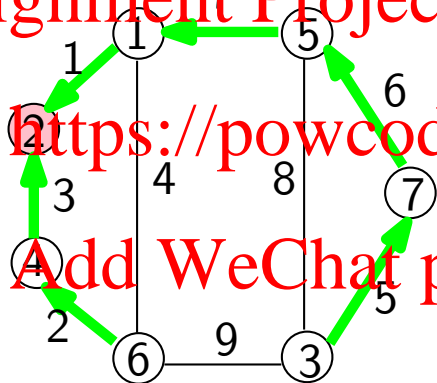
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→ *connect!*

## Sync MST – Level 2 Components

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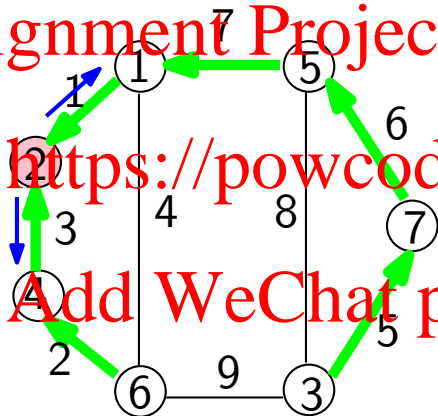
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## Sync MST – Level 2 Components

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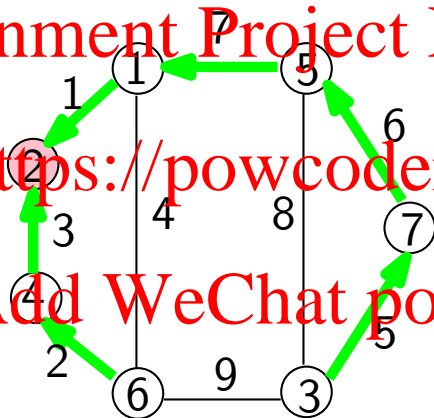
→ *initiate*

## Sync MST – Level 2 Components

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## Sync MST – Memento

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- Distributed Sync MST is based on Borůvka
- Borůvka can be viewed as **extension** of both Prim and Kruskal
- At each given **step**: Prim grows **one** single tree; Borůvka grows **all** trees simultaneously
- At each given **step**: Kruskal merges exactly **two** trees; Borůvka merges **all** trees simultaneously (2-way or n-way unions)
- The **steps** of sequential Borůvka correspond to more complex **levels** (phases) in distributed Sync MST



## Sync MST – Memento

- The MST is **unique**, if weights are pairwise distinct (but this can always be arranged, e.g. by lexicographic comparisons)

- There is one **single** common MWOE, aka the **core**, in all tree unions in the algorithm (obvious for 2-way unions, but needs proof for more)

- The leader of a union of trees is one of the two **endpoints** of the **core**, i.e. of the single common MWOE (this also ensures that the root stays reasonably close to the leaves)

- A level  $k$  union tree has size  $\geq 2^k$ , thus tree sizes grow **exponentially** and there are at most  $\log N$  phases

- To ensure required **synchronicity**, each level requires  $O(N)$  steps (this is ok, as there are only few levels)

## Sync MST – Memento

- The **accept** and **reject** messages are not really needed; i.e. the **test** messages may hold enough info for this decision

- The **report** messages are evaluated on-the fly, at each node, and only the **best** MWOE's details are forwarded up, towards the leader

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- The **report** messages leave behind a **route** to the node holding the best MWOE of that subtree

- The **connect** messages are **routed** on the tree path, which goes from the leader to the node holding the overall best MWOE

- The **connect** messages **reshape** the tree, by transporting the leadership, i.e. resetting parent and child pointers along their path

# Async MST – GHS and variants

Specific difficulties of the **async** version - not present in the **sync** version:

- Two neighbours may be part of the same component tree, but they have not learned this yet (logical error)
- Not all component trees may have a guaranteed size  $> 2^k$ ; some may grow much faster than others (complexity issue)
- More generally, component trees may be at different levels... (logical and complexity issues)

Read more in Lynch's textbook:

- §4.4 : **sync** GHS
- §15.5 : **async** GHS, plus summary revision of **sync** GHS
- §15.3 : **async** STtoLeader (on unrooted STs)