

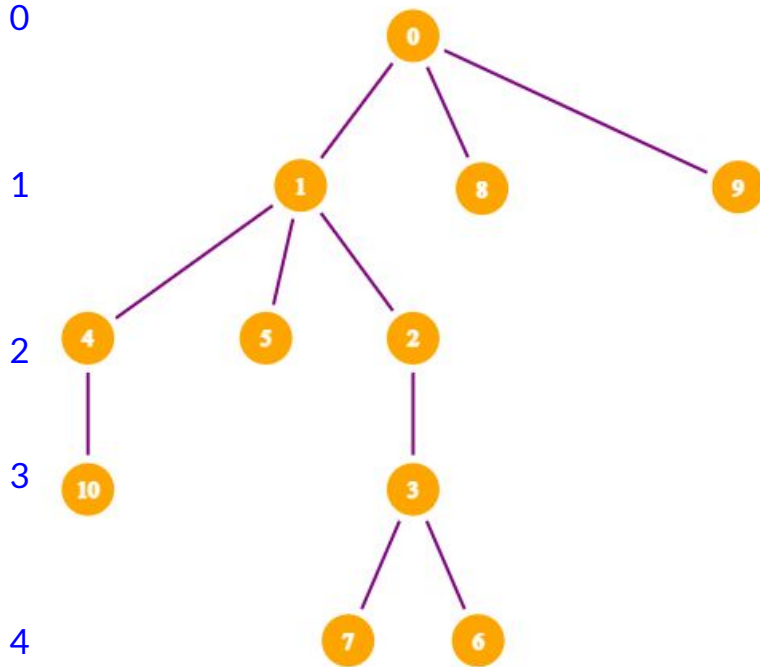


*Before we embark on a new adventure...*





What is the lowest common ancestor?



Given two nodes on a **TREE** (rooted at node 0), their LCA is the node that is a *parent* (ancestor) of each node AND has the maximal depth.

It is the 'best' meeting point between the two nodes, if you can only travel up the tree.

What is the LCA of node 4 and 6?

Solve queries: find LCA of nodes  $a$  &  $b$  quickly



Binary  
Jumping

Which algorithm to  
choose?

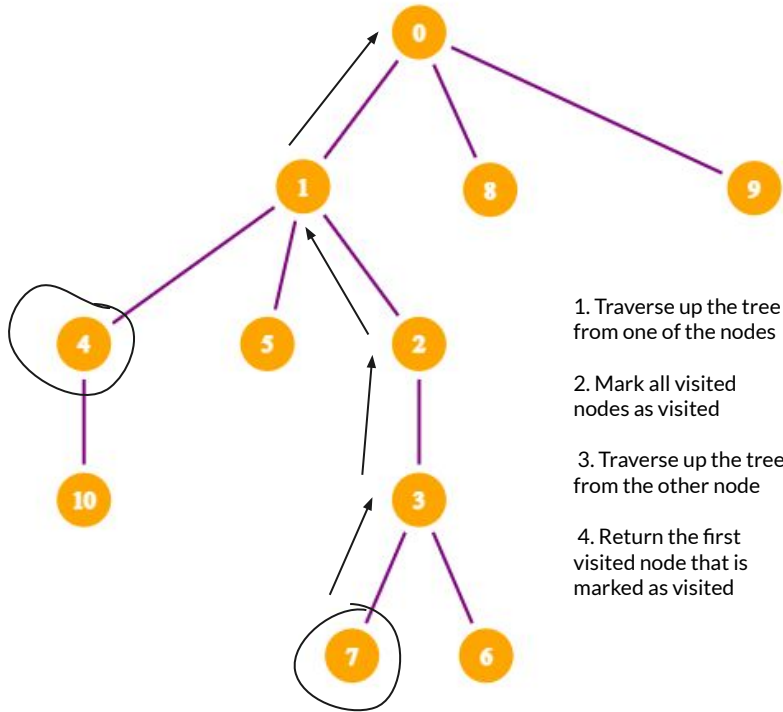
Euler  
Tour

~~Naive  
solution~~



## NAIVE SOLUTION

$O(N)$  time per query



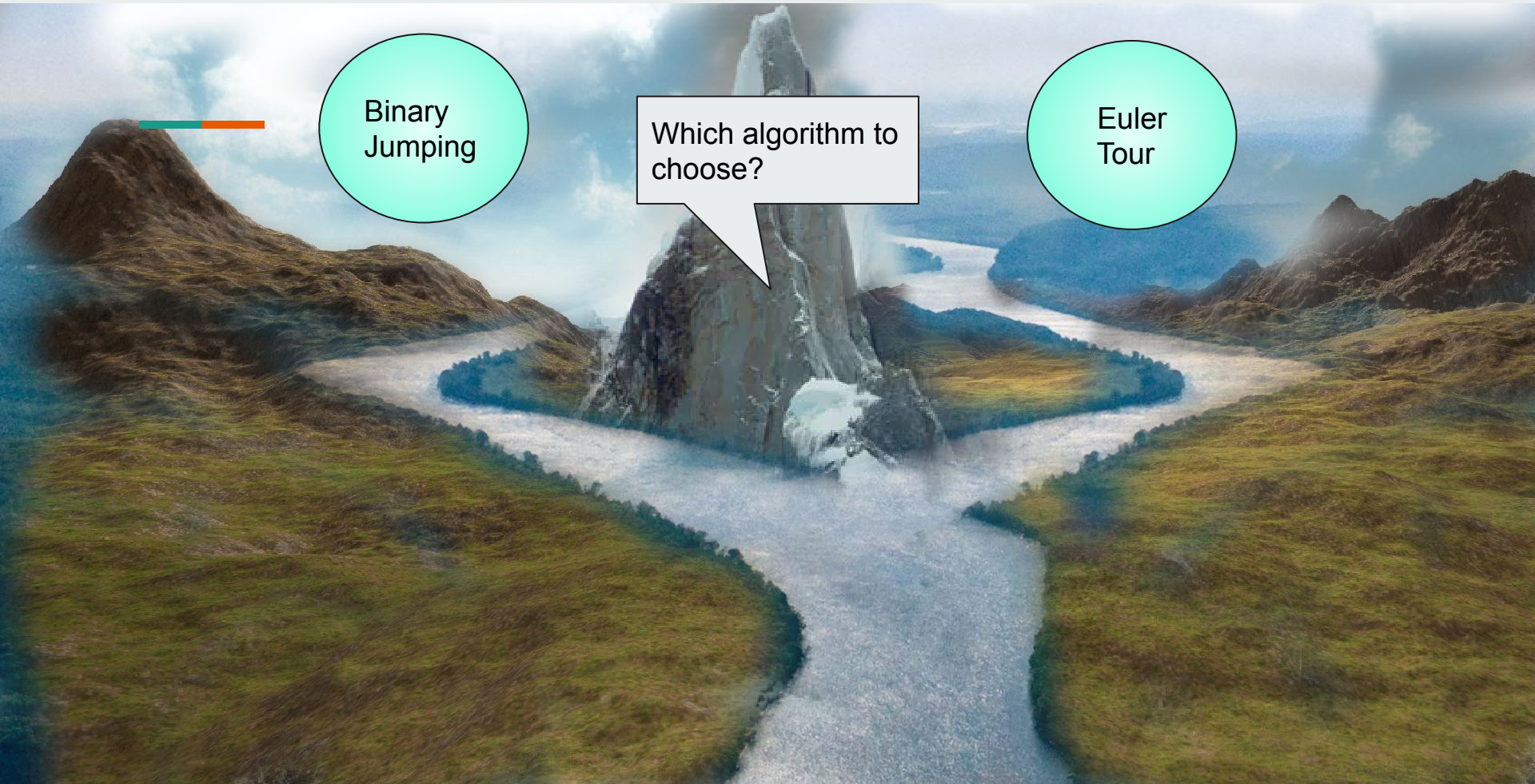
1. Traverse up the tree from one of the nodes

2. Mark all visited nodes as visited

3. Traverse up the tree from the other node

4. Return the first visited node that is marked as visited





Binary  
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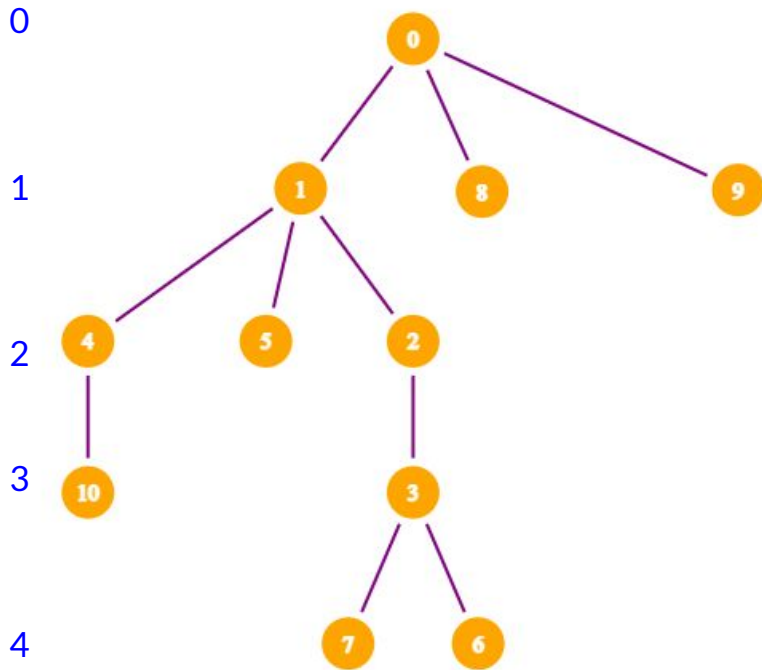
# Binary Jumping

Preprocess:  $O(N \log N)$

Query:  $O(\log N)$



## Algorithm using Sparse table and Binary jumping



- 1) Preprocessing using sparse table:  
Calculate for each node the  $2^k$  th parent

```

for (I = 1; I <= N; I++)
  for (J = 1; (1 << J) < N; J++)
    if (sparse_table[I][J-1] != -1)
      sparse_table[I][J] =
        sparse_table[sparse_table[I][J-1]][J-1];

```

NOTE\* Set  $\text{sparse\_table}[i][j] = -1$  for out of bounds jumps.

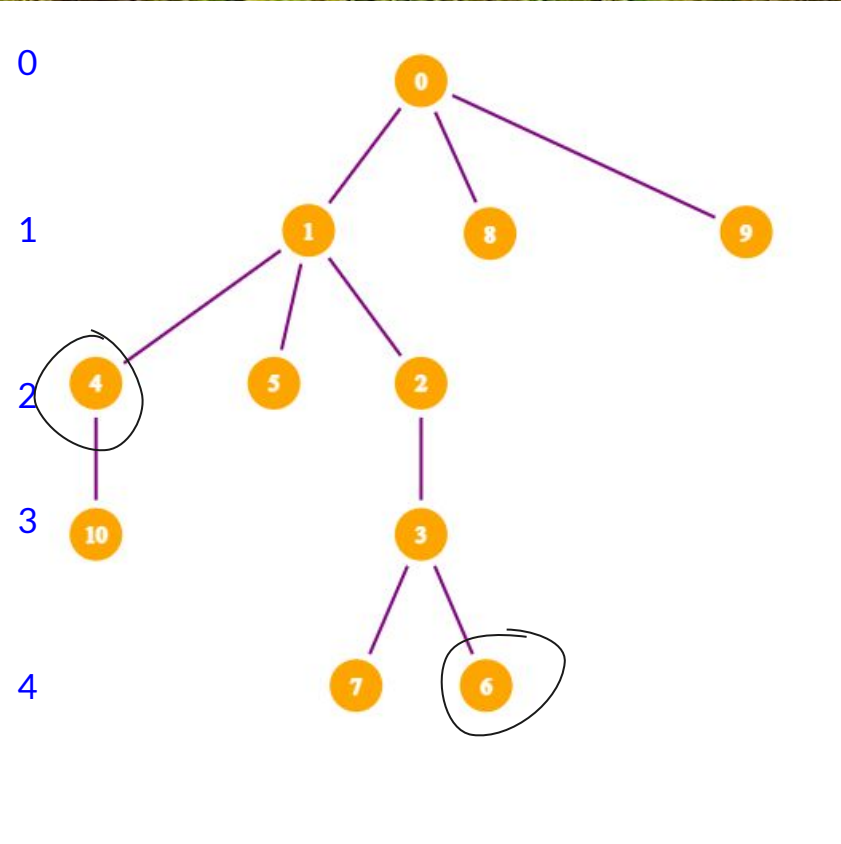
Example-if node  $u$  has to climb 37 edges to reach the LCA:  
 $37 = 32 + 4 + 1 = 100101_2$

Then we make a jump of 32. Set this to node  $u$   
 Make a jump of 4. Set this to node  $u$ .  
 Make a jump of 1. Set this to node  $u$ .

- 1\*) Run a simple DFS to get depth of each node



Algorithm using Sparse table and Binary jumping



2) if depth a  $\neq$  depth b:  
Move the deeper one up until depth a = depth b  
Use binary jump for  $\log N$  time

Move the deeper one up until depth  $a = \text{depth } b$   
Use binary jump for  $\log N$  time

3) Implementation detail:  
After 2) if  $a == b$  then we have found our LCA  
find a number **log** such that  $2^{(\text{log}+1)} > \text{de}$

After 2) if  $a == b$  then we have found our LCA

find a number **log** such that  $2^{(\log+1)} > \text{depth}[a]$

```
4) for(I = log; I>=0; I--)
    if(sparse_table[a][I] != -1 && sparse_table[a][I]
    != sparse_table[b][I])
        a = sparse_table[a][I];
        b = sparse_table[b][I];
```

```
if(sparse_table[a][I] != -1 && sparse_table[a][I]
```

```
!= sparse_table[b][I])
```

```
a = sparse_table[a][I];
```

```
b = sparse_table[b][I];
```

Intuitively:

- Start with the biggest jumps  $2^l$  and make the jumps smaller and smaller
- If we overshoot our LCA, just decrease our jump to  $2^{l-1}$
- Otherwise the  $2^l$  th ancestor of a and b are NOT equal  
So we set a and b to be their respective ancestors  
Also decrease our jump to size  $2^{l-1}$

Our jump will eventually reach size  $2^0 = 1$  in which case we find our LCA



Euler  
Tour

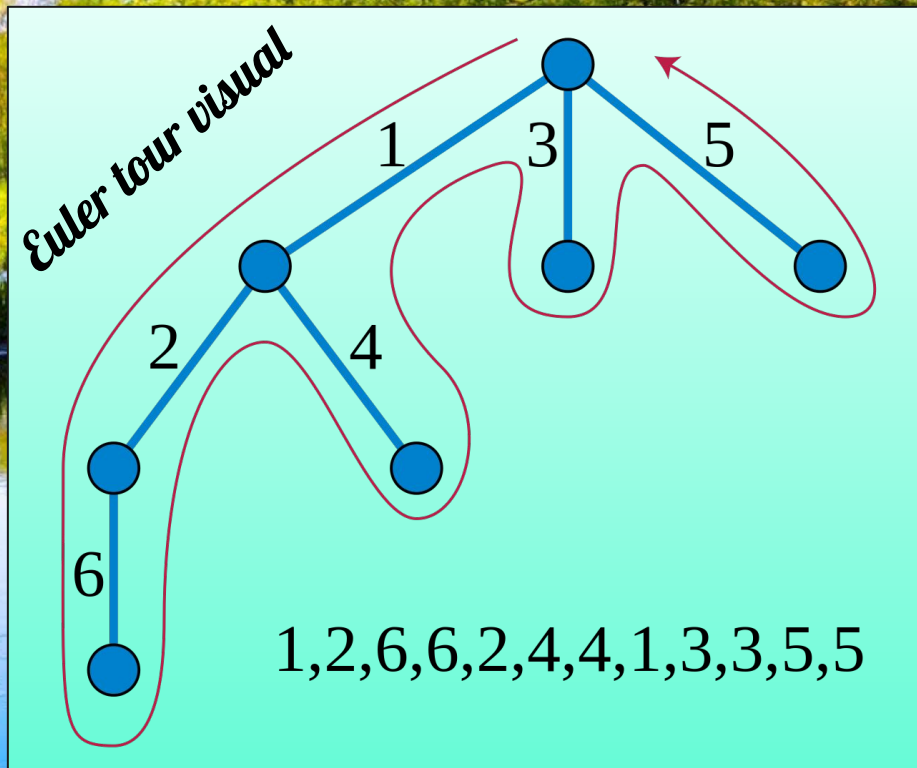


Finally ... We have  
found peace

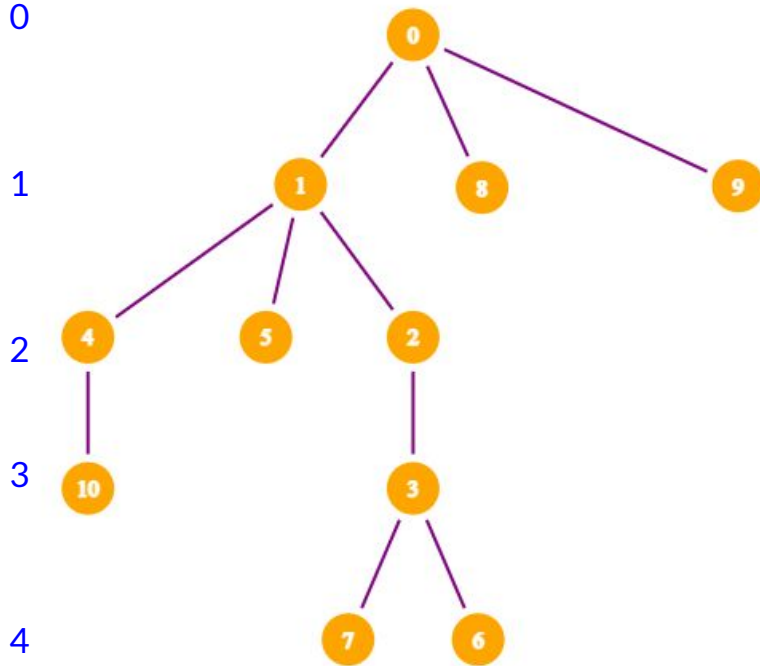
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**Euler tour  
technique**

Preprocess:  $O(N \log N)$

Query:  $O(\log N)$



## Algorithm using Euler tour array and RMQ



- 1) Run DFS on the tree :  
Add a node to the array *euler* whenever the node is called (time in) and whenever the dfs backtracks from one of its children.

*Pseudocode:*

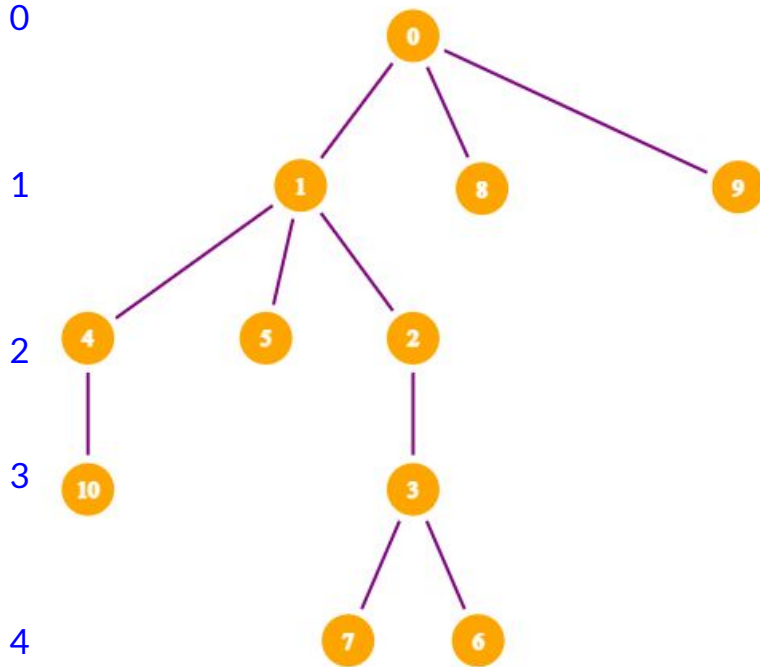
```

Dfs (current_node, parent_node):
    euler.push( current_node)
    For each child of current_node:
        Dfs(child , current_node)
    euler.push(current_node)
  
```

To note : each node is added to the euler  $c + 1$  times, where  $c$  is the number of children of the node. Therefore,  $\text{euler.size()} = 2n - 1$



## Algorithm using Euler tour array and RMQ



We also want to keep track of at least one occurrence for each node in the euler array. For simplicity, we record the first time for each node in the tin array

*Pseudocode:*

**timer = 0**

Dfs (current\_node, parent\_node):

euler.push( current\_node)

**tin[current\_node] = timer**

**timer ++**

For each child of current\_node:

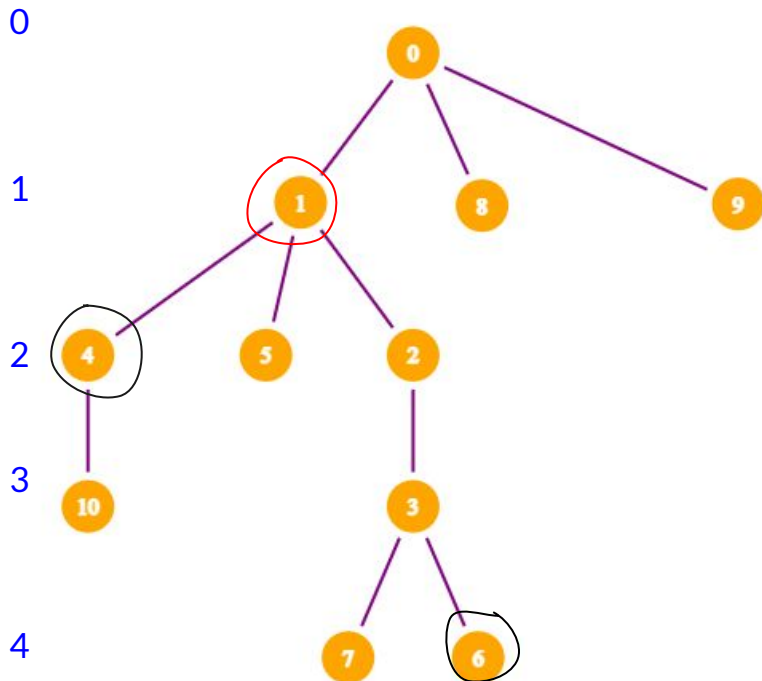
Dfs(child , current\_node)

euler.push(current\_node)

**timer ++**

0 1 4 10 4 1 5 1 2 3 7 3 6 3 2 1 0 8 0 9 0

## Algorithm using Euler tour array and RMQ



2) Find LCA of node a and b:

- ★ Pick any node a in the euler array (call its index  $i_a$ )
- ★ Pick any node b in the euler array (call its index  $i_b$ )
- ★ In the range  $[i_a, i_b]$  the LCA is the node with minimal depth (closest to the root)

Concrete example:

Find LCA node 4 and node 6

We can use our *tin* array:

$\text{tin}[4] = 2, \text{tin}[6] = 12$

From all the nodes between indices 2 and 12 the one closest to the root is node 1.

(Implementation detail : instead of looking at **depths** of nodes we can alternatively check **time in** of each node

**If  $\text{depth}[a] < \text{depth}[b]$  then  $\text{tin}[a] < \text{tin}[b]$**

So makes no difference! )

0	1	4	10	4	1	5	1	2	3	7	3	6	3	2	1	0	8	0	9	0
---	---	---	----	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---



But how can we find the minimal node in this range efficiently???

**RMQ = Range Minimum Query**



Here we go again



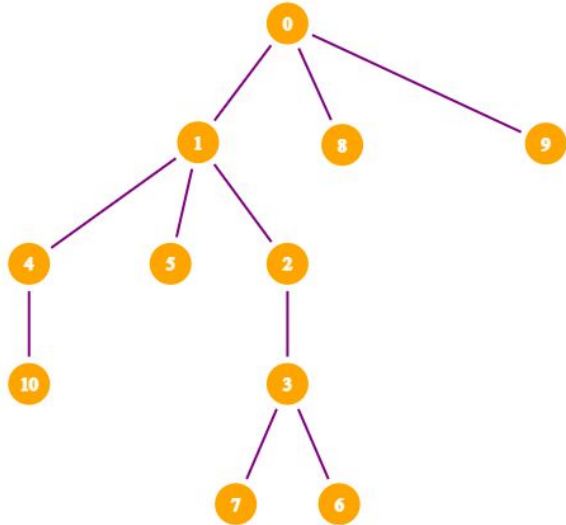


EULER

0	1	4	10	4	1	5	1	2	3	7	3	6	3	2	1	0	8	0	9	0
---	---	---	----	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---

Time in

0	1	2	3	2	1	6	1	8	9	10	9	12	9	8	1	0	17	0	19	0
---	---	---	---	---	---	---	---	---	---	----	---	----	---	---	---	---	----	---	----	---



3.) Build segment tree using array of *tin* values (or *depth* values)

Then call

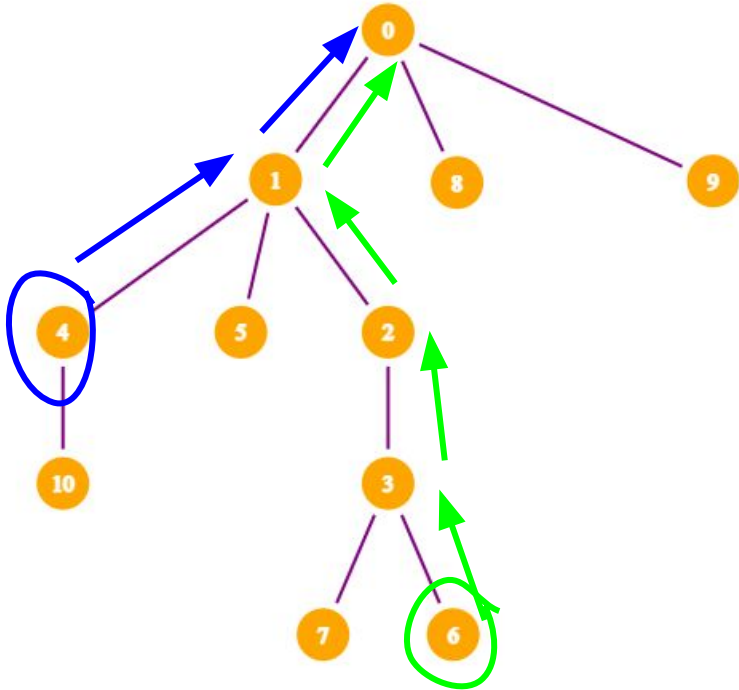
**query(tin[a], tin[b])**

Where query is segment tree query function.

It will return tin[LCA], which you can use to find LCA.

Each query takes  $O(\log N)$  time.

# One application of LCA



Distance between two nodes in a tree

If  $c = \text{LCA}(a, b)$

$ab = \text{dist}(a) + \text{dist}(b) - 2 * \text{dist}(c)$

Where  $\text{dist}(\text{node})$  is distance from node to root



# Thank you







# References & Resources

<https://codeforces.com/blog/entry/77451#:~:text=Definition%3A,edge%20or%20the%20sum%2C%20etc.> (Binary Jumping)

[https://saco-evaluator.org.za/presentations/2019%20Camp%202/Lowest%20Common%20Ancestor%20\(Andi%20Qu\).pdf](https://saco-evaluator.org.za/presentations/2019%20Camp%202/Lowest%20Common%20Ancestor%20(Andi%20Qu).pdf) (Andi's presentation)

[https://saco-evaluator.org.za/presentations/2018%20Camp%202/Range%20Queries%20and%20Fenwick%20Trees%20\(Yaseen%20Mowzer\).pdf](https://saco-evaluator.org.za/presentations/2018%20Camp%202/Range%20Queries%20and%20Fenwick%20Trees%20(Yaseen%20Mowzer).pdf) (Segment Trees)