

# Generating all distributions of objects to bins

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## Presented By

GROUP-3

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## Submitted To

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# Problem Statement

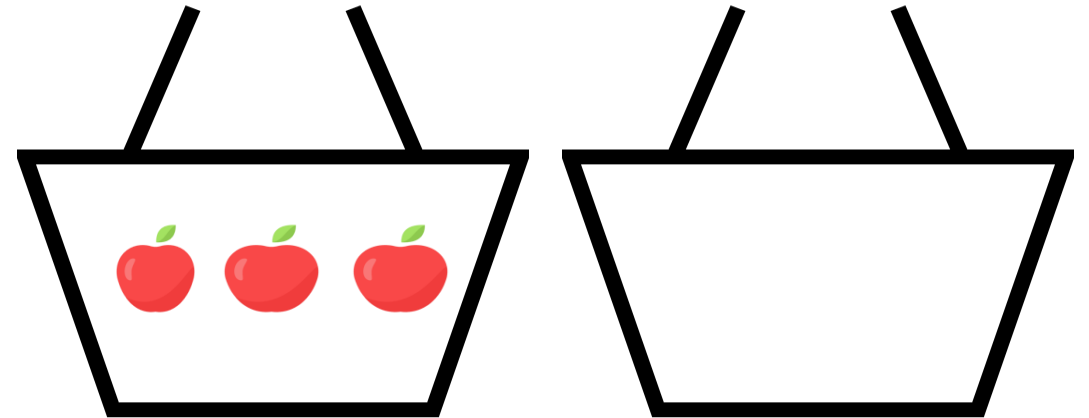
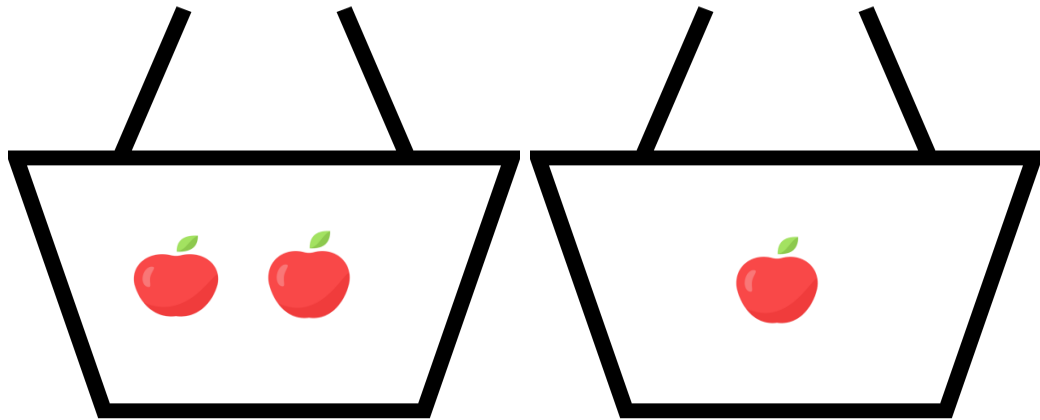
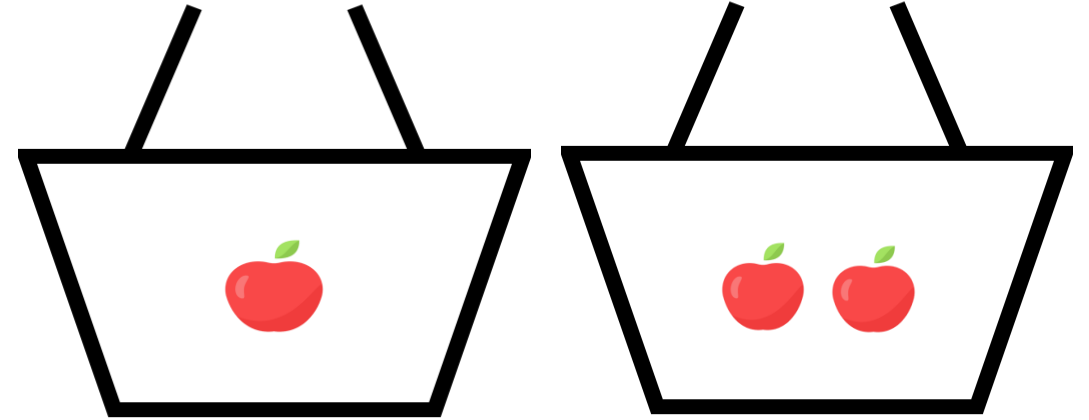
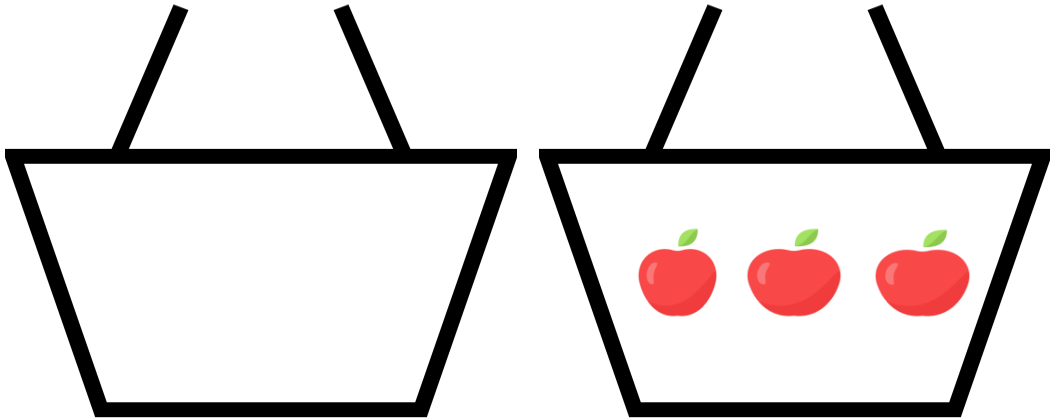
**Can we generate all distributions of identical objects to bins  
in  $O(1)$  ?**

# Real world example of Distributing Object into bins

Number of distributions =  $(n + m - 1)! / n!(m - 1)!$

m= Number of bins = 2

n= Number of objects = 3



# The problems with Klingsberg's algorithm

- Cannot generate each solution in  $O(1)$
- Generates solutions in constant time on average
- Klingsberg's method requires searching for the second non-zero element in the sequence for solutions that have a non-zero first elements
- Inefficient in the generation process

# The Efficient Algorithm

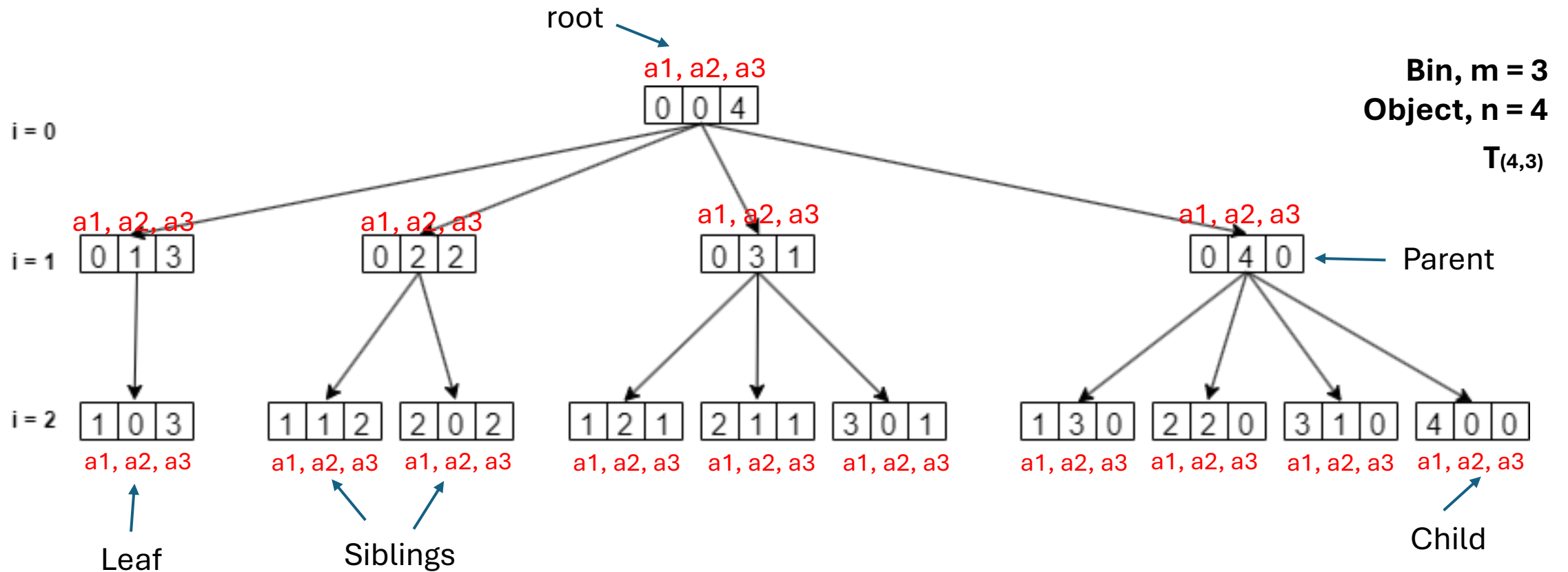
- **Constant Time Generation:** Generates each distribution in constant time (in ordinary sense)
- **Efficient Traversal:** Efficient tree traversal method to ensure each distribution is generated quickly.
- **Space Complexity:**  $O(m \lg n)$
- **Optimization:** Reduce non-generation steps
- **No repetition.**
- **Generates in specific order.**

# Application of the algorithm

- **Automation in Machines**
- **Computer Networks**
- **Client-Server Architecture**
- **Combinatorial Problems**

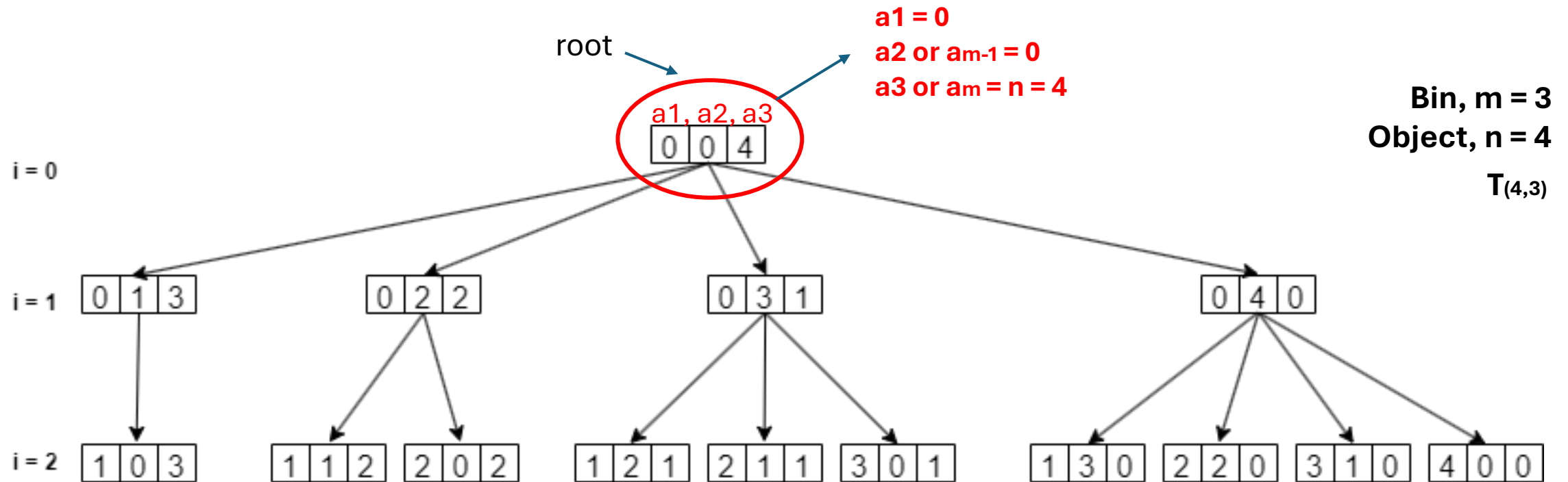
# The family tree of distributions

We define a tree structure  $T_{n,m}$  among the distributions in  $D(n, m)$ . Each node of  $T_{n,m}$  represents a distribution  $(a_1, a_2, \dots, a_m) \in D(n, m)$ . If there are  $m$  bins then there are  $m$  levels  $T_{n,m}$ .



# The family tree of distributions (cont.)

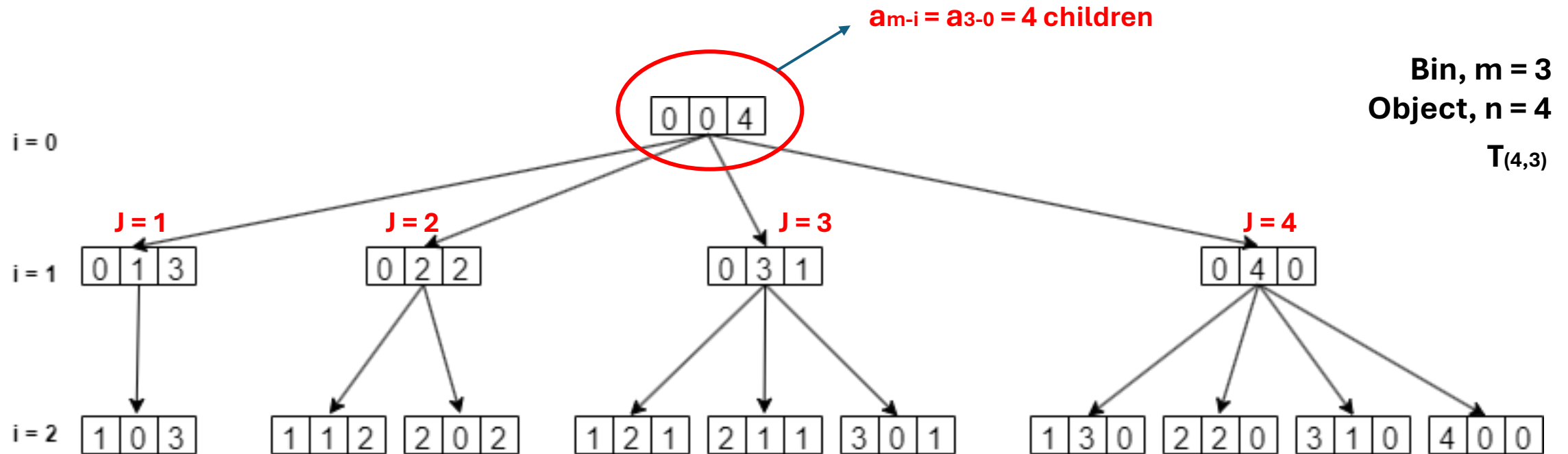
$T_{n,m}$  is a rooted tree we need a root, and the root is a node at level 0. we can observe that a node is at level 0 in  $T_{n,m}$  and  $a_1 = a_2 = \dots = a_{m-1} = 0$  and  $a_m = n$ .





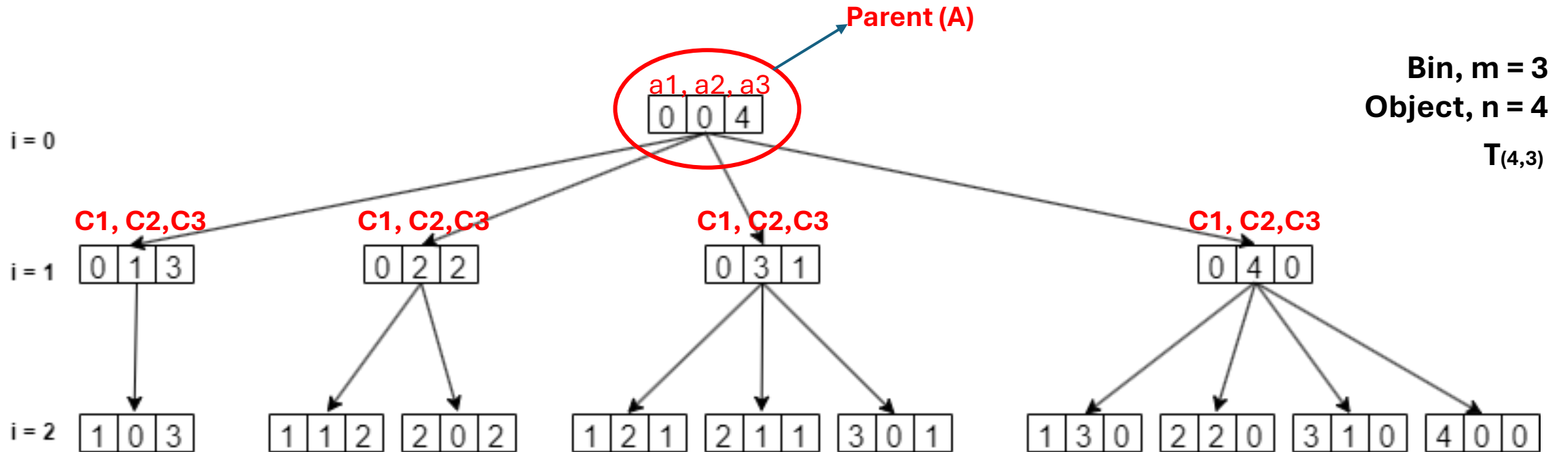
# Parent-child relationship

The number of children a parent has is equal to  $a_{m-i}$



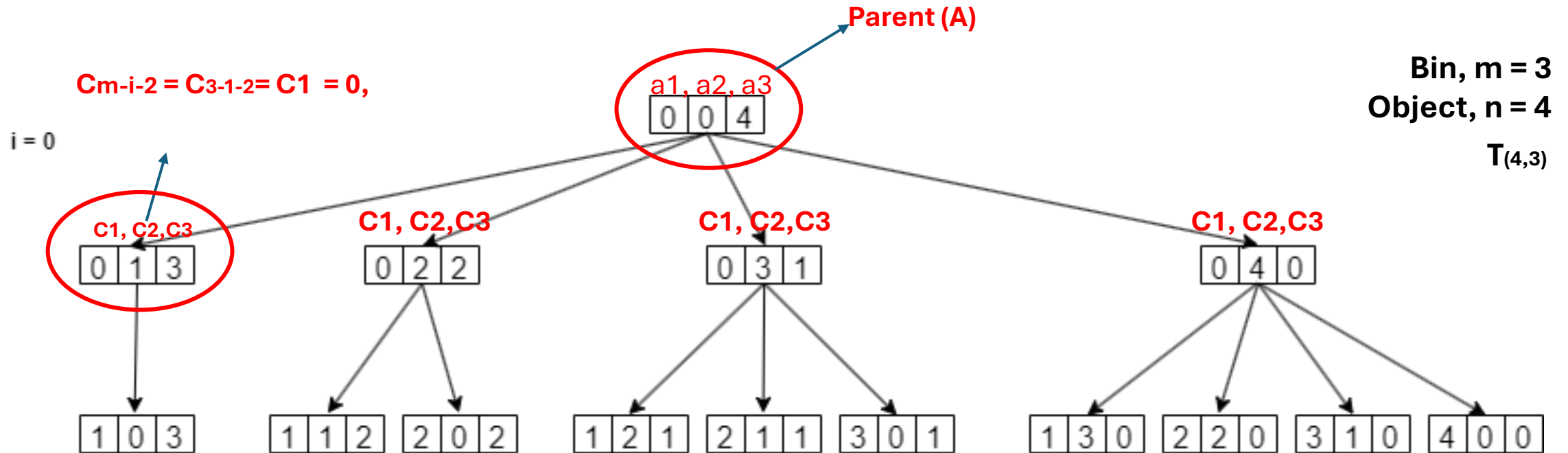
# Parent-child relationship (cont.)

Let  $C_j(A) \in D(n, m)$  be the sequence of  $j$ th child,  $1 \leq j \leq a_{m-i}$  of  $A$ . Note that  $A$  is in level  $i$  of  $T_{n,m}$  and  $C_j(A)$  will be in level  $i+1$  of  $T_{n,m}$ . We define the sequence for  $C_j(A)$  as  $(c_1, c_2, \dots, c_{m-i-1}, c_{m-i}, \dots, c_m)$ , where  $0 \leq i < m$ ,  **$c_1 = c_2 = \dots = c_{m-i-2} = 0$  and  $c_{m-i-1} = j$ ,  $c_{m-i} = a_{m-i} - j$  and  $c_k = a_k$  for  $m-i+1 \leq k \leq m$**



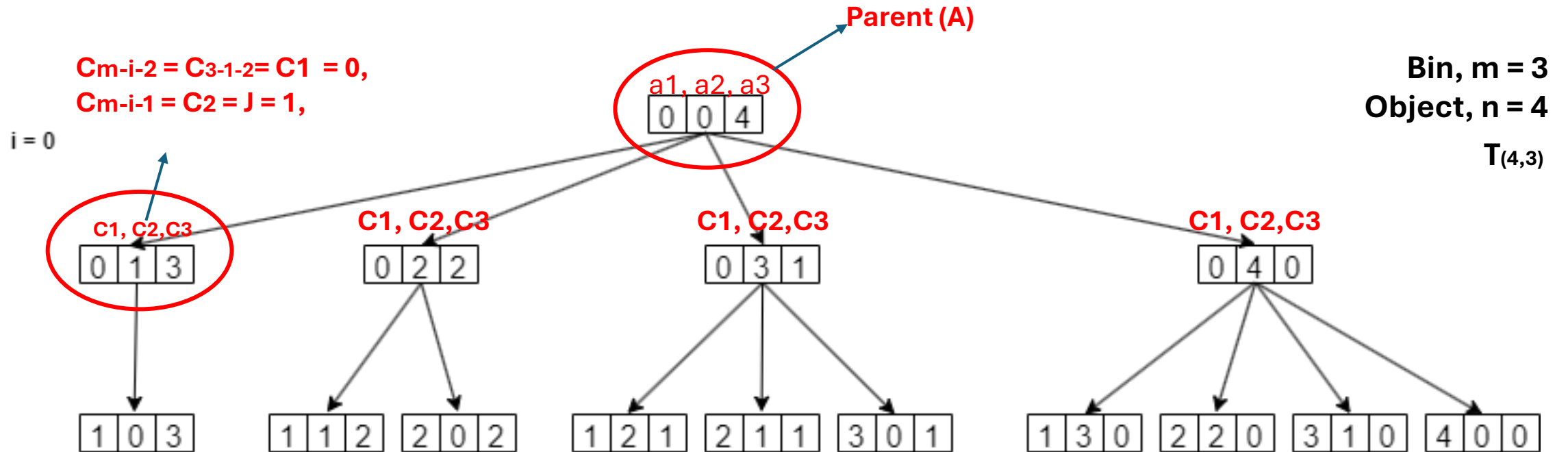
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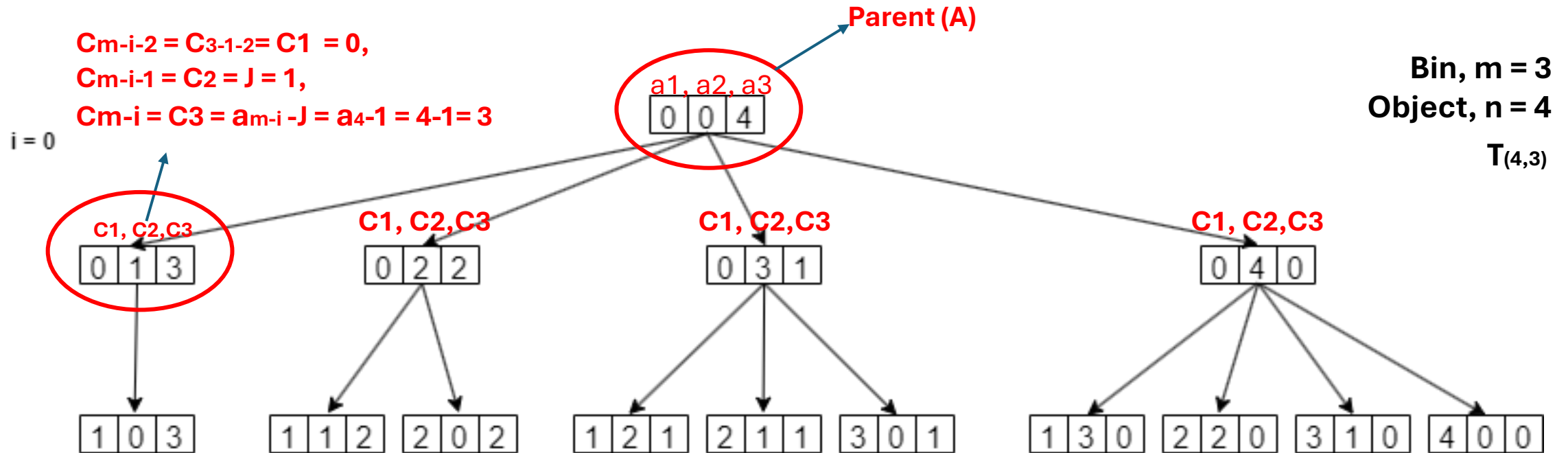
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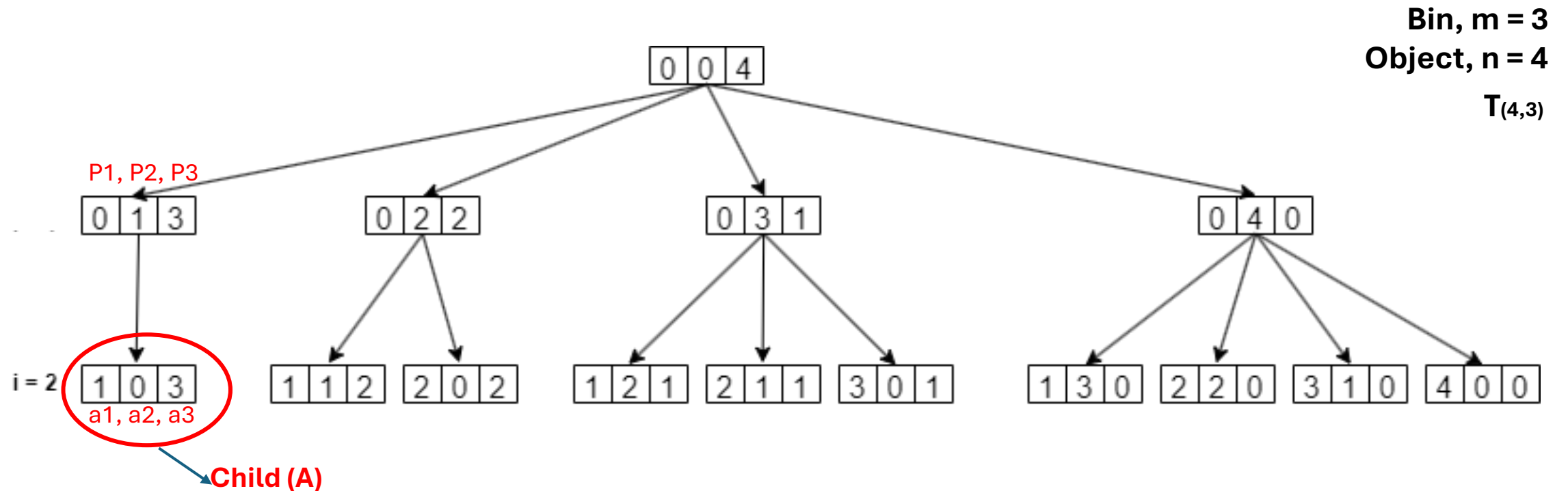
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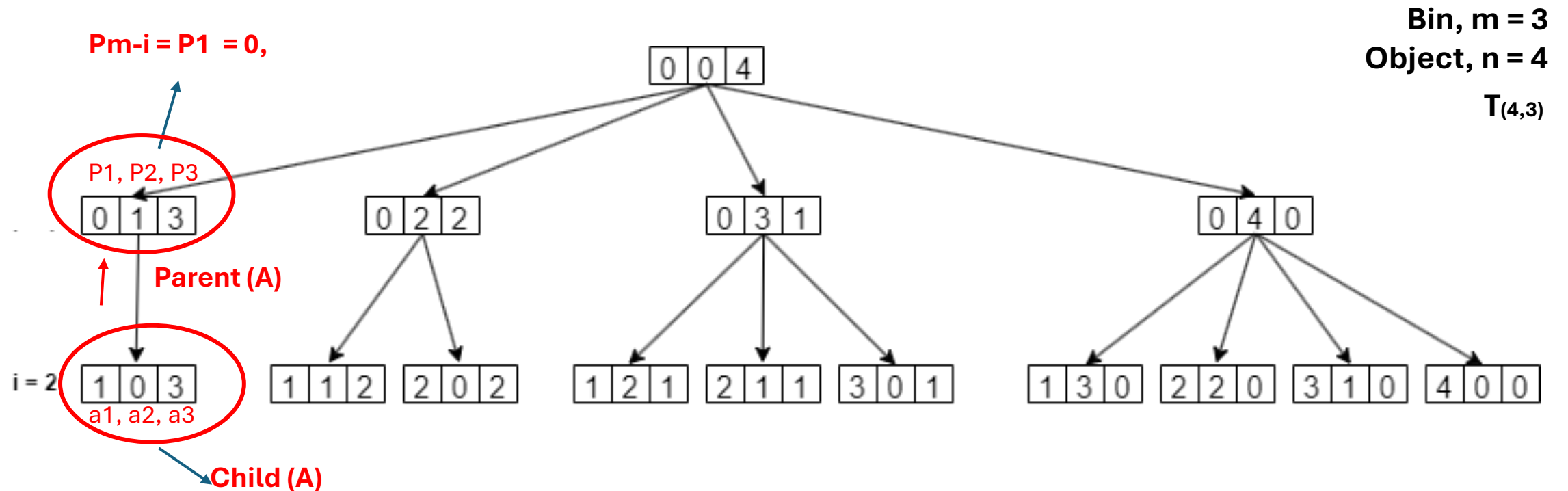
# Child–parent relationship

The child–parent relation is just the reverse of parent–child relation. Let  $P(A) \in D(n, m)$  be the parent sequence of  $A$ . We define the sequence for  $P(A)$  as  $(p_1, p_2, \dots, p_{m-i}, p_{m-i+1}, \dots, p_m)$  where  $1 \leq i < m$ ,  **$p_1 = p_2 = \dots = p_{m-i} = 0$** ,  **$p_{m-i+1} = a_{m-i} + a_{m-i+1}$**  and  **$p_j = a_j$  for  $m - i + 1 < j \leq m$** .



# Child-parent relationship(cont.)

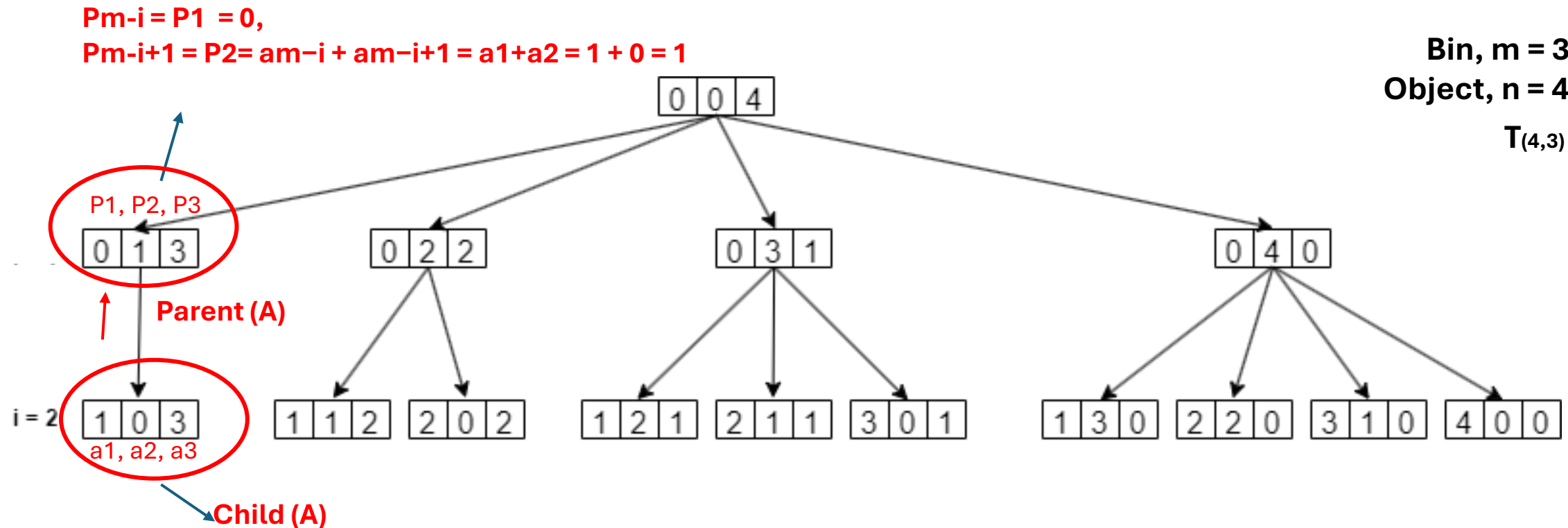
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# Child-parent relationship(cont.)

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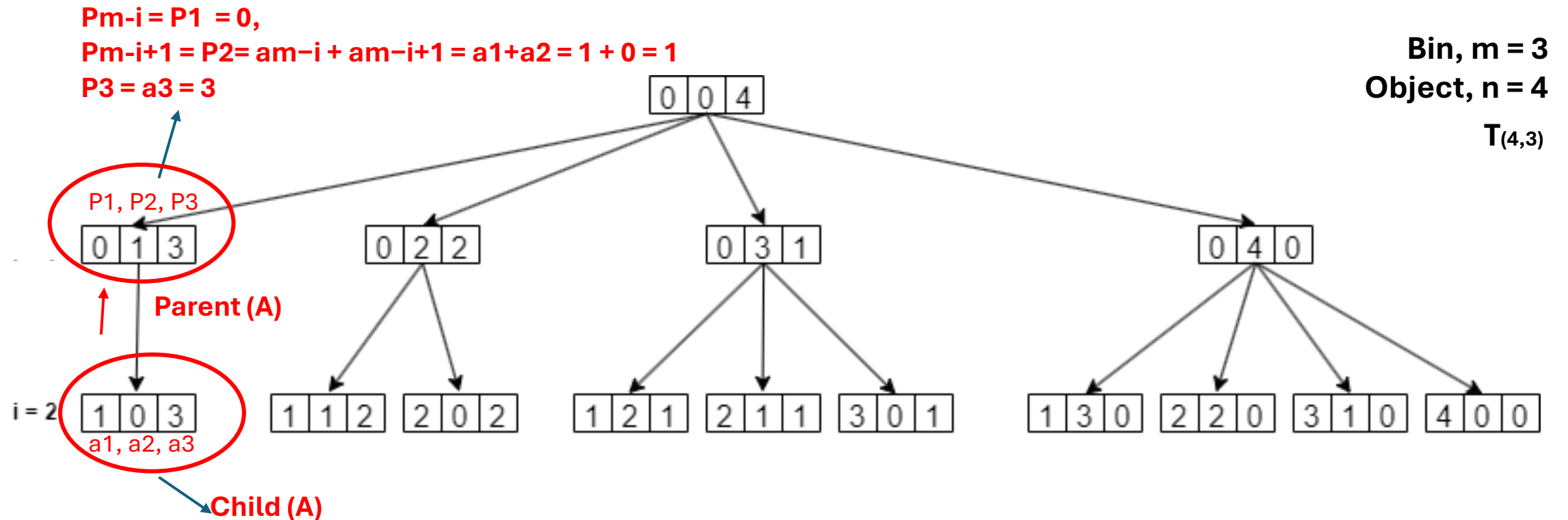
Bin,  $m = 3$   
Object,  $n = 4$   
 $T_{(4,3)}$





# Child-parent relationship(cont.)

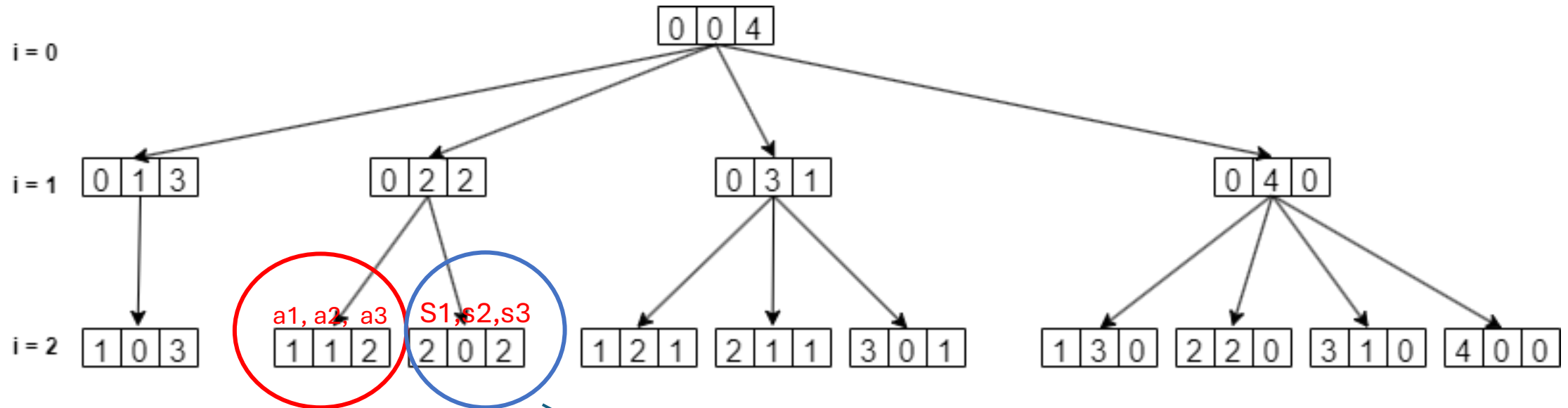
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# Efficient tree traversal $O(1)$ time

## Relationship between left sibling and right sibling

Right sibling  $A_s \in D(n,m)$  of node  $A$  exists if  $a_{m-i+1} \neq 0$  at level  $i$  of  $T_{n,m}$ . the sequence for  $A_s$  as  $(s_1, s_2, \dots, s_{m-i}, s_{m-i+1}, \dots, s_m)$ ,  $1 \leq i < m$  where  $s_1 = s_2 = \dots = s_{m-i-1} = 0$ ,  $s_{m-i} = a_{m-i+1}$ ,  $s_{m-i+1} = a_{m-i+1} - 1$  and  $s_j = a_j$  for  $m-i+2 \leq j \leq m$ .



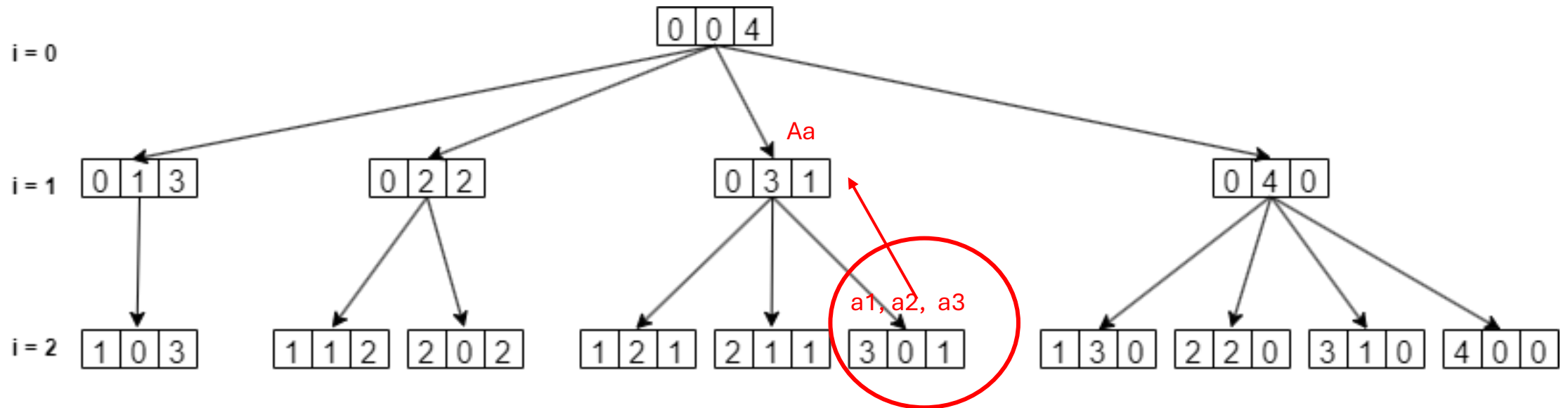
A

A<sub>s</sub>

- $m-i+2 \leq j \leq m \Rightarrow 3-2+2 \leq j \leq 3 \Rightarrow 3 \leq j \leq 3$   
 $\Rightarrow s_3 = a_3 \Rightarrow \mathbf{s_3 = 2}$
- $s_{m-i+1} = a_{m-i+1} - 1 \Rightarrow \mathbf{s_2 = a_2 - 1 = 1 - 1 = 0}$
- $s_{m-i} = a_{m-i+1} - 1 \Rightarrow \mathbf{s_1 = a_1 + 1 = 1 + 1 = 2}$

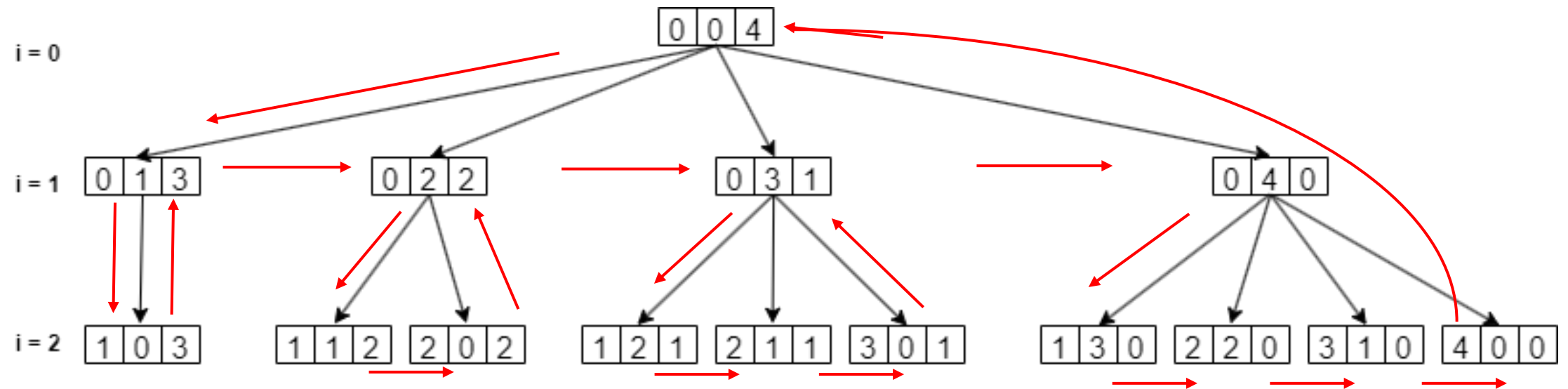
# Leaf-ancestor relationship

Al = rightmost leaf, Aa= Nearest ancestor which has right siblings, k = number of consecutive 0's, Aa ∈ D(n,m) of node Al exists if a2 = 0. Nearest ancestor is obtained by swapping a1 and ak+1 at level m-1-k



Al K=1,  
 $i = m-1-k = 3-1-1 = 1$   
 Aa = swapping a1 and ak+1 then Aa = 0,3,1

# Efficient tree traversal



# Pseudo Code

## Algorithm Find-All-Distributions( $n, m$ )

{  $A_r$  is the root sequence,  $S$  indicates the current stack }

**begin**

**Find-All-Child-Distributions**(  $A_r = (0, \dots, 0, n), 0, S$  );

**end.**



**Call the function by root**

## Procedure Find-All-Child-Distributions( $A = (a_1, a_2, \dots, a_m), i, S$ )

{  $A$  is the current sequence,  $i$  indicates the current level,  $A_c$  is the child sequence,  $A_s$  is the right sibling sequence,  $A_a$  is the ancestor sequence and  $S$  indicates the current stack }

**begin**

    Output  $A$  { Output the difference from the previous distribution }

**if**  $a_1 = 0$  **then**

**begin**

            {  $A$  has child }

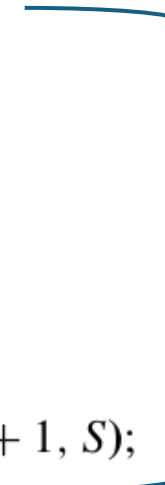
**if**  $a_{m-i} - 1 = 0$  **then**

**if**  $a_{m-i+1} \neq 0$  **then** **Push**(1,  $S$ );

**else** **Top**( $S$ )=**Top**( $S$ )+1;

**Find-All-Child-Distributions**(  $A_c = (a_1, a_2, \dots, a_{m-i-2}, 1, (a_{m-i} - 1), \dots, a_m), i + 1, S$ );

**end**



**Find Child**

```

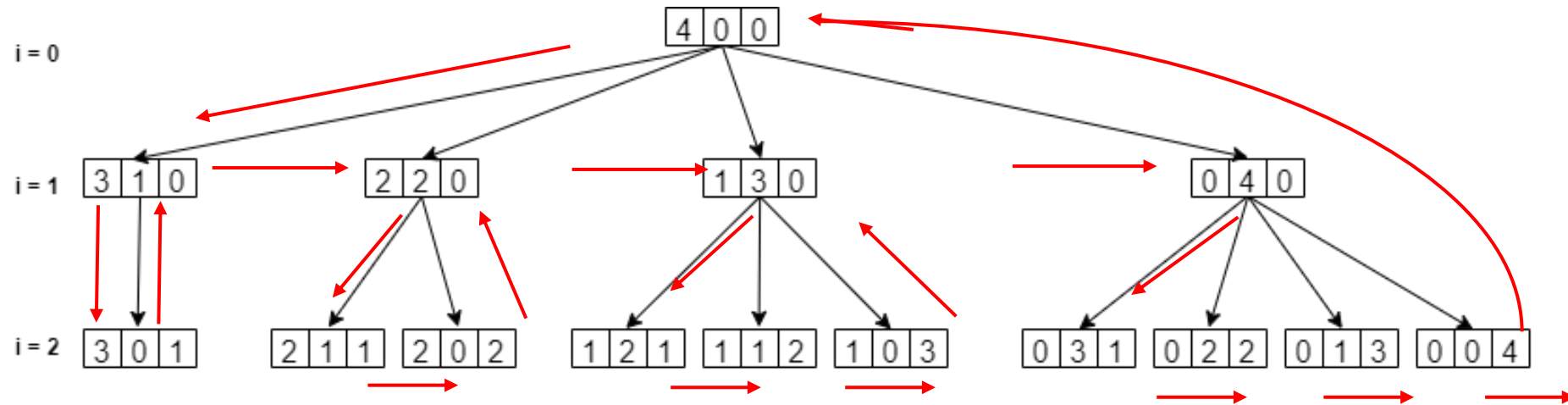
else if  $a_2 \neq 0$  then
  begin
    {  $A$  has right sibling }
    if  $a_2 - 1 = 0$  then
      if  $a_3 \neq 0$  then Push(1,  $S$ );
      else Top( $S$ )=Top( $S$ )+1;
    Find-All-Child-Distributions(  $A_s = ((a_1 + 1), (a_2 - 1), \dots, a_m), i, S$ )
  end
else
  begin
     $k$ =Pop( $S$ );
    Swap( $a_1, a_{k+1}$ ); {Generate the ancestor  $A_a$  of  $A$ }
    if  $k = m - 1$  then return ; {  $A_a$  is the root }
    else
      begin
        {  $A_a$  has right sibling }
        if  $a_{k+2} - 1 = 0$  then
          if  $a_{k+3} \neq 0$  or  $k + 2 = m$  then Push(1,  $S$ );
          else Top( $S$ )=Top( $S$ )+1;
        Find-All-Child-Distributions( $A_{as} = (a_1, a_2, \dots, (a_{k+1} + 1), (a_{k+2} - 1), \dots, a_m), m - 1 - k, S$ );
      end
    end
  end
end;

```

**Find Right Sibling**

**Find Ancestor's  
Right Sibling**

# Anti-lexicographical Order



# Conclusion

This paper presents a simple, efficient algorithm for generating all distributions in  $D(n, m)$  with specified order, including anti-lexicographic, operating in constant time.



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Thank You

