

# Data Structure

# Growth of Functions

input size : n

each basic operation takes constant time

+ - \* / % = [i] func(x) return val comparison

An example :

- How do we analyze an algorithm?
  - Count the number of basic operations

sum(A, n)

```
1 tempsum = 0
2 for i = 0 to n-1
3     tempsum += A[i]
4 return tempsum
```

Number of basic operations

1  
2<sup>n</sup> + 2  
2<sup>n</sup>  
1  
Total:  $4n + 4$

Why  $2^n + 2$ ? The actual effect of the loop is:  
for (i=0; i<n; i++)

We assign 0 to i once, increment i for n times, compare i < n 1 times

Why  $2^n$ ? Accessing A[i] n times, addition to tempsum n times

It is annoying to count the detailed number of basic operations, we will show how to simplify the counting process later (Page 27).

e.g. linear search

for i=0 to n-1

if array[i] == target:

return i

return -1

$2n + 1$

n

1

1

total =  $4n + 4$

e.g. binary search analysis

```
left = 0    right = n-1          2  
while left <= right:           log2n + 1  
    mid = (left + right) // 2   3 · log2n + 3  
    if target < array[mid]:    2 · log2n + 2  
        right = mid + 1       2 · log2n + 2  
    else if target > array[mid]: 2 · log2n + 2  
        left = mid + 1        2 · log2n + 2  
    else:                      1  
        return mid             1  
return -1
```

$$\text{total: } 12 \cdot (\log_2(n) + 1)$$

e.g. max sub array sum

```
maxSum = array[0]          2  
for i=0 to n-1              ≥ n+2  
    subSumFromI = 0           n  
    for j=i to n-1           (2n+2) + (2(n-1)+2) + ... + (2 · 1 + 2)  
        subSumFromI += array[j] 2(n + n-1 + n-2 + ... + 1)  
        if subSumFromI > maxSum  n + n-1 + n-2 + ... + 1  
            maxSum = subSumFromI  n + n-1 + n-2 + ... + 1  
    return maxSum             1
```

$$\text{total: } 3n^2 + 8n + 5$$

## Big-Oh Notation

def  $g(n) = O(f(n))$  iff exist a constant  $c$  and no s.t.  $g(n) \leq c \cdot f(n)$  for all  $n > n_0$ .

BP  $g(n) \leq f(n)$  为上界

describe running time of the algorithms using Big-Oh / Omega / Theta notations

linear search:  $g(n) = 4n + 4 = O(n)$  time complexity of the linear search algorithm is  $O(n)$   
any linear function:  $g(n) = a \cdot n + b = O(n)$

log functions with different base  $a > 1$  and  $b > 1$ :

$$g(n) = \log_a n = O(\log_b n) = O(\log n)$$

↑ can assume  $\log n = \log_a n$  or  $\log n = \log_b n$

Practice: Binary Search: prove that  $g(n) = 12 \log_2 n + 17 = O(\log n)$

can find  $c = 2^9$ ,  $n_0 = 2$  such that

$$g(n) = 12 \log_2 n + 17 \leq 12 \log_2 n + 17 \log_2 n = 29 \log_2 n \text{ for } n \geq 2$$

thus  $g(n) = O(\log_2 n) = O(\log n)$

$$g(n) = 1,000,000 = O(1)$$

can find  $c = 1,000,000$ ,  $n_0 = 1$  s.t.

$$g(n) = 1,000,000 \leq 1,000,000 \cdot 1 \text{ for } n \geq 1$$

thus  $g(n) = O(1)$

Disprove that  $g(n) = 2n^2 = O(n)$

find a constant  $c, n_0$ , s.t.

$$2n^2 \leq cn \text{ for all } n > n_0 \Leftrightarrow n \leq \frac{c}{2} \text{ for all } n > n_0$$

b.c.  $n$  can reach infinite, thus doesn't exist such constant  $c$

- **Transitivity:** Prove that if  $g(n) = O(f_1(n))$ ,  $f_1(n) = O(f_2(n))$ , then  $g(n) = O(f_2(n))$

**Proof:** According to the definition, we know that:

There exist positive constants  $c_1$  and  $n_1$  for  $g(n)$  such that

$$g(n) \leq c_1 \cdot f_1(n) \text{ for } n \geq n_1$$

and  $c_2, n_2$  for  $f_1(n)$  such that:

$$f_1(n) \leq c_2 \cdot f_2(n) \text{ for } n \geq n_2$$

Then, we know for  $n \geq \max(n_1, n_2)$ :

$$g(n) \leq c_1 f_1(n) \leq c_1 \cdot c_2 \cdot f_2(n).$$

Therefore  $g(n) = O(f_2(n))$  according to the definition, which finishes the proof.

**Big-Oh Rule** (i) Given  $a > b > 0$ ,  $n^a$  grows faster than  $n^b$

$$\lim_{n \rightarrow \infty} \frac{n^b}{n^a} = 0$$

(ii) Product property

$$g_1(n) = O(f_1(n)) \quad g_2(n) = O(f_2(n))$$

$$\text{Then, } g_1(n) \cdot g_2(n) = O(f_1(n) \cdot f_2(n))$$

(iii) Sum property only applies when the number of functions are constants

$$g_1(n) = O(f_1(n)), \quad g_2(n) = O(f_2(n))$$

$$\text{Then, } g_1(n) + g_2(n) = O(\max(f_1(n), f_2(n)))$$

(iv) Log functions grow slower than power functions

$$\lim_{n \rightarrow \infty} \frac{(\log n)^a}{n^b} \rightarrow 0 \quad \text{for } a, b > 0$$

(v) Exponential functions grow faster than power functions

$$\lim_{n \rightarrow \infty} \frac{n^k}{a^n} \rightarrow 0 \quad \text{if } a > 1 \text{ for any } k$$

Growth: power  $\rightarrow$  log  $\rightarrow$  constant

# Complexity Analysis Big-Omega and Big-Theta

Big-Omega def:  $g(n) = O(f(n))$  iff exist a constant  $c$  and no s.t.  $g(n) \geq c \cdot f(n)$  for all  $n > n_0$ .

即  $g(n) \leq k \cdot f(n)$  为下界

Big-Omega Rules: Big-Oh rules (i)-(v) all apply to Big-Omega

Big-Theta: Big-Oh & Big-Theta are not tight

Big-Theta def: If  $g(n) = O(f(n))$  and  $g(n) = \Omega(f(n))$   
 $g(n) = \Theta(f(n))$

# Linked List

Insertion

$\text{insert}(L, e)$ : add  $e$  at the end of the linked list

Step 1: identify the last node  $u_{\text{last}}$  stored in the linked link

Step 2: allocate a new node  $u_{\text{new}}$

Step 3:  $u_{\text{new}}.\text{element} = e$

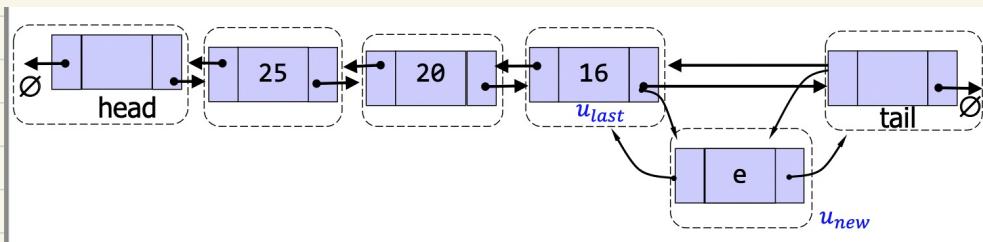
Step 4:  $u_{\text{new}}.\text{prev} = u_{\text{last}}$

Step 5:  $u_{\text{new}}.\text{next} = \text{tail}$

Step 6:  $u_{\text{last}}.\text{next} = u_{\text{new}}$

Step 7:  $\text{tail}.\text{prev} = u_{\text{new}}$

order can be changed



Insertion in Place:  $\text{insertInPlace}(L, p, e)$ : insert element  $e$  after node  $p$

Step 1: allocate a new node  $u_{\text{new}}$

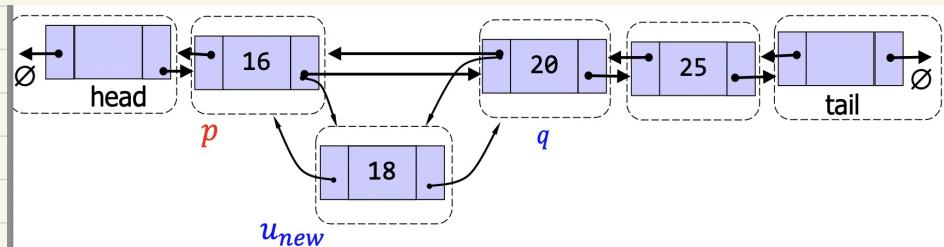
Step 2: set  $u_{\text{new}}.\text{element} = e$

Step 3: set  $u_{\text{new}}.\text{prev}$  to the address of  $p$

Step 4: set  $u_{\text{new}}.\text{next}$  to the address of  $q = p.\text{next}$

Step 5: set  $p.\text{next}$  to the address of  $u_{\text{new}}$

Step 6: set  $q.\text{prev}$  to the address of  $u_{\text{new}}$



Deletion : delete node p in the linked list.

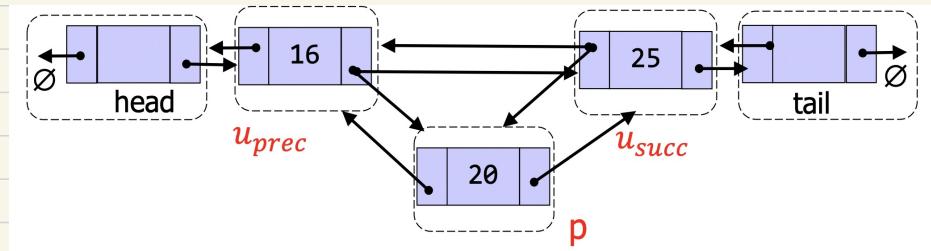
step 1 : identify the preceding node  $u_{prec}$  of node p

step 2 : identify the succeeding node  $u_{succ}$  of node p

step 3 : let  $u_{prec}.next$  points to  $u_{succ}$

Step 4 : let  $u_{succ}.prev$  points to  $u_{prec}$

Step 5 : free the memory of node p



# Stack

Last-In-First-Out

Balanced Symbol Checking

Evaluation of Expressions

Infix expression  $\Rightarrow$  Postfix Expressions



e.g. (1) infix:  $7 ( (2+3)*4 )$

postfix:  $7 2 3 + 4 *$



e.g. (2) infix:  $2 * ( 3 + 2 * 4 )$

postfix:  $2 3 2 + * 4 *$

# Queue First - In - First - Out

Implementation: Linked List

No capacity constraint

Array: circular structure

Better practical performance

Enqueue: change the rear position  
to the next position

Dequeue: change the front position  
to the next position

The next of capacity -1 is 0

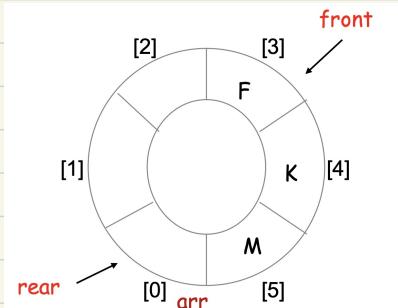
The next rear :

```
if rear != capacity - 1  
    rear = rear + 1  
else  
    rear = 0
```

Or:  $(\text{rear} + 1) \% \text{capacity}$

The next front.

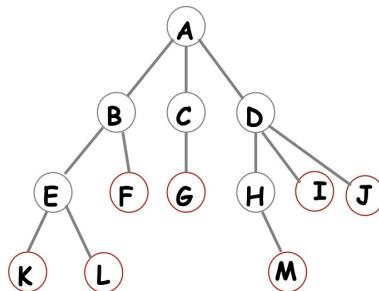
$(\text{front} + 1) \% \text{capacity}$



# Tree

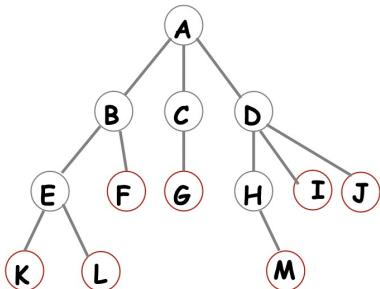
Terminology :

- **Node:** the elements plus the branches to each node.
- **Edge:** nodes are connected by **edges**.
- **Branch of a node:** the number of subtrees of a node
- **Leaf nodes (or external node):** nodes that have zero branch
- **Internal nodes:** nodes that don't belong to leaf nodes.



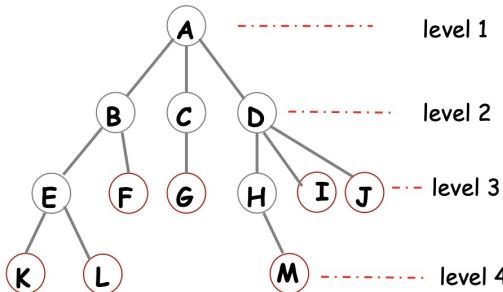
- **A** is the **root node**
- The **branch** of node **B** is 2
- **F, G, I, J, K, L, M** are **leaf nodes**
- **A, B, C, D, E, H** are **internal nodes**

- **Children:** the roots of the subtrees of a node X are the **children** of X
- **Parent:** X is the **parent** of its children
- **Siblings:** children of the same parent are said to be **siblings**.
- **Ancestors** of a node: parent, grand-parent, grand-grandparent,etc.
- **Descendants** of a node: child, grandchild, grand grand-child, etc.



- **E and F** are the **children** of **B**
- **B** is the **parent** of **E and F**
- **C** is the **sibling** of **B**
- **A, B, E** are the **ancestors** of **K**
- **H, I, J, M** are the **descendants** of node **D**

- The **level** of a node: defined by letting the root be at level one. If a node is at level  $l$ , then its children are at level  $l+1$ .
- Height (or depth)**: the maximum level of any node in the tree

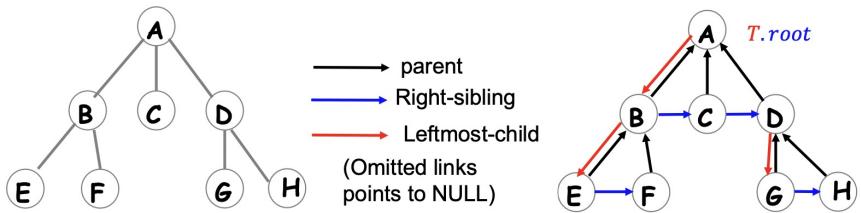


The height (depth) of the tree is 4

$$\# \text{ of nodes} = \# \text{ of edges} + 1$$

- Pointer representation of a tree  $T$

- Maintain  $\text{root}$  to store the address of the root node.
- For each node, it further maintains the address of its  $\text{parent}$ ,  $\text{leftmost-child}$ , and  $\text{right-sibling}$ .
  - For example, for node B
    - $B.\text{parent}$  stores the address of node A.
    - $B.\text{right-sibling}$  stores the address of node C.
    - $B.\text{leftmost-child}$  stores the address of node E.



# Binary Tree

maximum number of nodes on level  $i$  of a binary tree is  $2^{i-1}$  for  $i \geq 1$

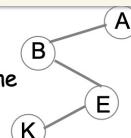
maximum number of nodes in a binary tree of height  $k$  is  $2^k - 1$  for  $k \geq 0$

height of a binary tree with  $n$  nodes falls in the range between  $\lceil \log_2(n+1) \rceil$  and  $n$

## Special Binary Tree

- **Skewed binary tree**

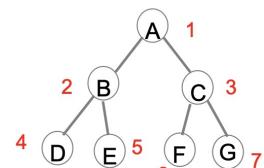
- All the nodes have only either one child or no child.



Skewed binary tree

- **Full binary tree**

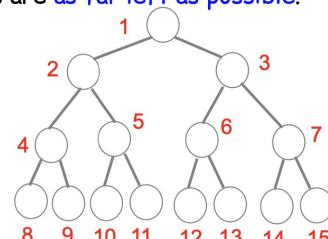
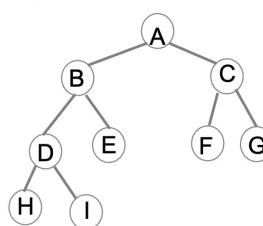
- every node other than the leaves has two children.
- If the height is  $k$ , the number of nodes is  $2^k - 1$  (i.e., the maximum we can have with height  $k$ )
- The nodes of a full binary tree can be numbered from 1 to  $2^k - 1$ .



Full binary tree

- **Complete binary tree**

- A binary tree in which every level, except possibly the last, is completely filled, and all nodes are as far left as possible.



- Its nodes correspond to the nodes numbered from 1 to  $n$  in the full binary tree of depth  $k$ .

- In this example, we have 9 nodes. The complete binary tree should correspond to the nodes numbered 1-9 in the full binary tree

# Binary Tree Implementation

Pointer :

node :

parent : ptr

leftchild : ptr

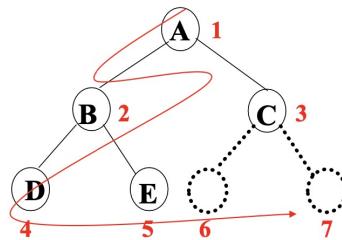
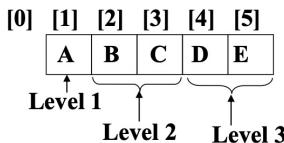
rightchild : ptr

element : value

Array : will be used for heap

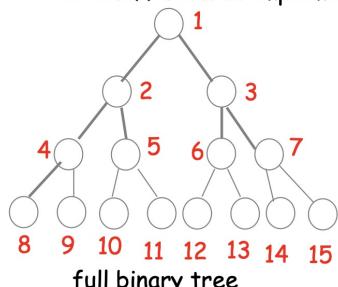
- An array representation

- Given a complete binary tree with  $n$  nodes. For any  $i$ -th node,  $1 \leq i \leq n$ ,
  - $\text{parent}(i)$  is  $\lfloor i/2 \rfloor$
  - $\text{leftChild}(i)$  is at  $2i$  if  $2i \leq n$ . Otherwise,  $i$  has no left child.
  - $\text{rightChild}(i)$  is at  $2i + 1$  if  $2i + 1 \leq n$ . Otherwise,  $i$  has no right child.



- An array representation

- Generalize to all binary trees
- Efficient for complete binary trees.
- But inefficient for skewed binary trees.
- Inefficient to implement the ADT



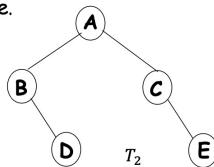
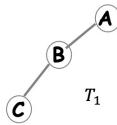
full binary tree

1	A	Level 1
2	B	Level 2
3	D	
4	E	
5	H	Level 3
6	I	
7	K	
8	L	
9		Level 4
10		
11		
12		
13		
14		
15		

e.g. Practice

- What are the array representation of the following binary trees?

- Show the content in the array.
- Hint: first obtain the ID for each node.



$T_1:$  [A | B | C | ]

$T_2:$  [A | B | C | D | E]

Inorder Traversal:

Inorder traverse its left-subtree

Then visit the root node

Inorder traverse its right-subtree

Postorder Traversal

Postorder traverse its left-subtree

Postorder traverse its right-subtree

Then visit the root node

Preorder Traversal

Visit the root node

Preorder traverse its left-subtree

Preorder traverse its right-subtree

- Calculate the height of a binary tree.

Algorithm: `height(btreetree)`

```
1 if isEmpty(btreetree)
2   return 0
3 lheight ← height(Lchild(btreetree))
4 rheight ← height(Rchild(btreetree))
5 return 1 + max(lheight, rheight)
```

1 ADT BinTree

```
2 BinTree create(bt);
3 Boolean isEmpty(bt)
4 BinTree MakeBT(bt1,
5 item, bt2);
6 BinTree Lchild(bt);
7 BinTree Rchild(bt);
8 element Data(bt);
```

- Sum over all the elements in the binary tree

Algorithm: `sum(btreetree)`

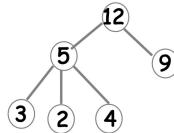
```
1 if isEmpty(btreetree)
2   return 0
3 lsum ← sum(Lchild(btreetree))
4 rsum ← sum(Rchild(btreetree))
5 return Data(btreetree) + lsum + rsum
```

# Heap

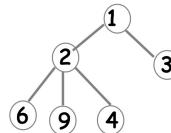
A special form of complete binary tree

e.g. max heap or min heap

- Max (resp. Min) heap is defined based on max (resp. min) tree and complete binary tree
- Max (resp. min) tree: A max (resp. min) tree is a tree in which the element, or call it the key value, in each node is no smaller (resp. no larger) than the key values in its children.



An example of a max tree

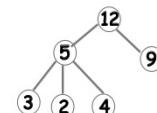


An example of a min tree

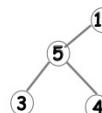
Max Heap

a complete binary tree that is also a max tree

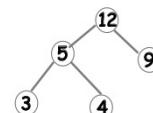
root contains the largest element  
easy to find the maximum



A max tree, but not a binary tree



A max tree, but not a complete binary tree



A max tree and also a complete binary tree

- A complete binary tree can be represented with arrays

- For any  $i$ -th node,  $1 \leq i \leq n$ ,
  - parent( $i$ ):  $\lfloor i/2 \rfloor$
  - leftChild( $i$ ):  $2i$  if  $2i \leq n$ . Otherwise, no left child.
  - rightChild( $i$ ):  $2i + 1$  if  $2i + 1 \leq n$ . Otherwise, no right child.

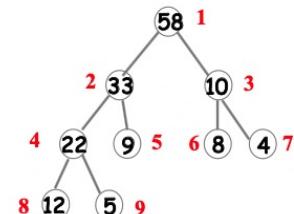
- A max heap can be represented as an array  $\text{arr}$  of elements such that

- $\text{arr}[i] \geq \text{arr}[2i]$  if  $2i \leq n$
- $\text{arr}[i] \geq \text{arr}[2i + 1]$  if  $2i + 1 \leq n$

[1] [2] [3] [4] [5] [6] [7] [8] [9]

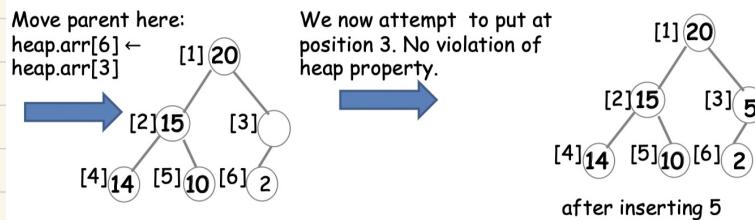
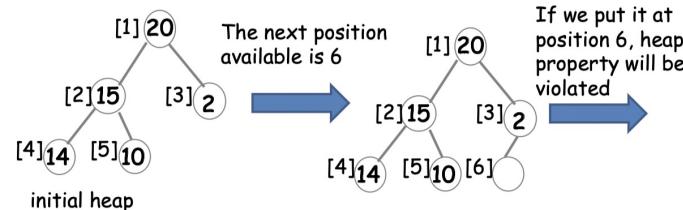
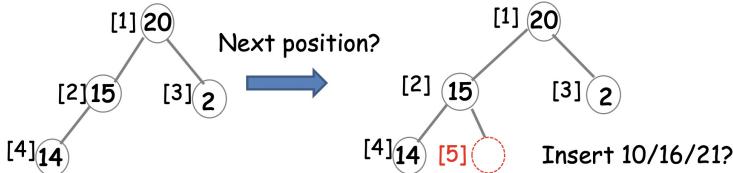
58 23 10 22 9 8 4 12 5

33

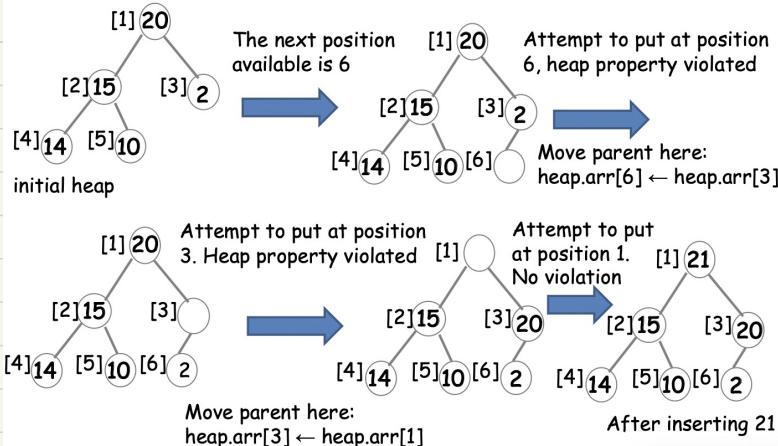


## • Questions:

- Where to insert?
  - After insertion, it must still be a complete binary tree.
  - If the current heap includes **x** nodes, then the new element should be inserted at the **(x+1)-th** position of the complete binary tree
- After insertion, does it violate the heap property?  
If so, can we resolve this?



Insert 21



## Heap Insertion Implementation

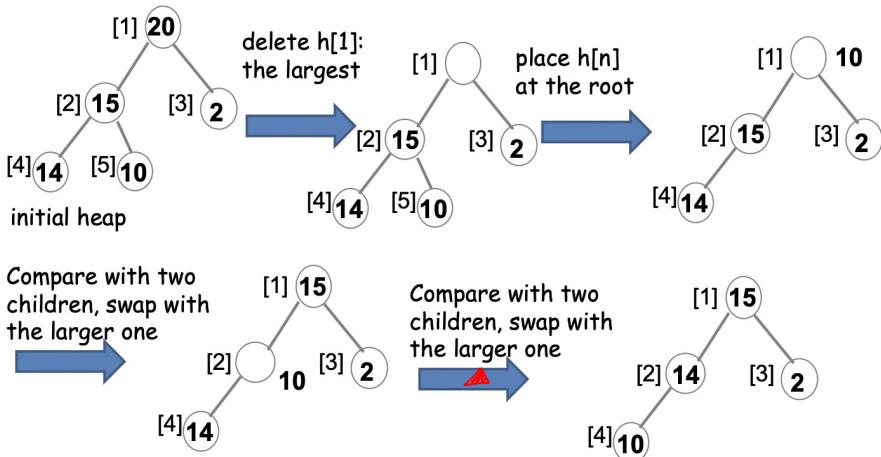
- Put it at the  $(\text{size}+1)$ -th position in the complete binary tree.
- But this may violate the heap property
  - Denote the current position as  $i$ . Its parent position is at  $i/2$
  - If the current position is at the root, just insert item to root
  - Otherwise, compare the parent and the item to be inserted.
    - If the parent has smaller key, move parent to current position, change the current position to  $i/2$ , and repeat above process.
    - Otherwise, insert the item to the current position

Algorithm: `InsertHeap(heap, item)`

```
1  if isFull(heap)
2      error "Heap full"
3  heap.size = heap.size+1
4  i = heap.size
5  While i!= 1 and item.key > heap.arr[i/2].key
6      heap.arr[i] = heap.arr[i/2]
7      i = i/2
8  heap.arr[i] = item
```

Heap Deletion  
 $O(\log n)$

- Return and delete the largest element,
  - Which one to delete?
    - According to the max-heap property, the largest element is at the root
  - Which one to fill at the root?
    - The tree must still be a complete binary tree: Use the last one to fill the root position
    - This obviously violates the heap property: how to resolve?



# Binary Search Tree

Heap, max tree, binary tree are inefficient for searching

In a sorted array:

Linear scan:  $O(n)$  cost

Binary search:  $O(\log n)$  cost

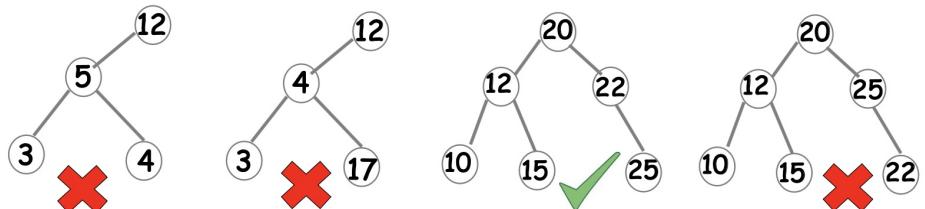
Main Idea: compare the searchnum with the middle element of the array, then we can reduce the search space by half

How to map to binary trees?

Binary Search Tree (BST) definition:

- A **binary search tree** is a binary tree that satisfies the following properties:
  - Every node has a **unique key**.
  - For each internal node  $x$ :
    - All nodes in a nonempty **left subtree** of  $x$  must have keys **smaller than the key of  $x$** .
    - All nodes in a nonempty **right subtree** of  $x$  must have keys **larger than the key of  $x$** .

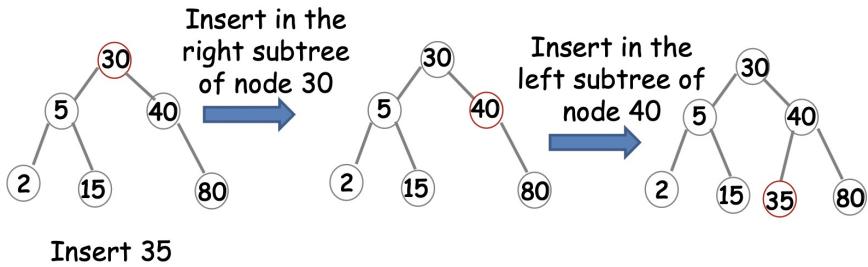
Are these trees binary search trees? Why?



Search cost: Denote  $h$  as the height of the BST, time complexity  $O(h)$

## Insertion of BST:

- No duplicate in the binary search tree
  - Ignore the insertion if it contains a key that exists in the binary search tree
- Where to insert?
  - After insertion, we should not violate the BST property

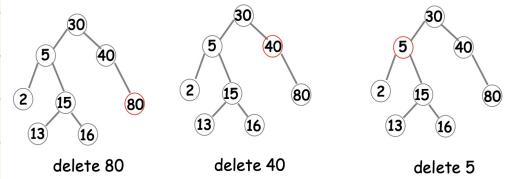


- Base case: the root is empty, insert the data at the root.
- Otherwise, follow the search path until we find a node  $x$  such that
  - (i)  $x.key = data.key$ , simply return since BST does not allow duplicate keys
  - (ii) node  $x$  has no right child and  $x.key < data.key$ 
    - Insert as the right child of  $x$
  - (iii) the node has no left child and  $x.key > data.key$ 
    - Insert as the left child of  $x$
- Pseudocode: left as self exercise
- Time complexity?
  - $O(h)$ , where  $h$  is the height of the BST

Deletion of BST : Case 1: delete a leaf node

Case 2: delete a node with only one child

Case 3: delete a node with two children

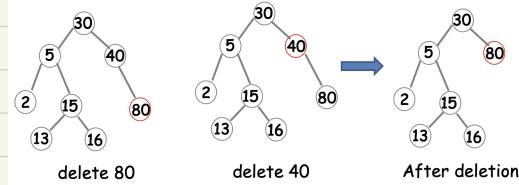


Case 1: delete leaf node

If the leaf node is a right (resp. left) child, just set the right (resp. left) child of its parent to NULL.

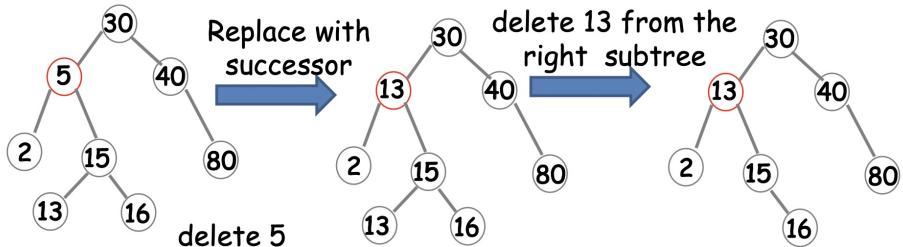
Case 2: delete a node with a single child

Put the single child at the place of the deleted.



Case 3: delete a node with two children

- We should replace a node here. But which node to put here?
- The **successor** of the node to be deleted
  - The successor of node 5 is node 13
- Replace the current node with its successor. Then delete the **successor** from the **right sub-tree**.
  - The deletion of successor falls into case 1 or case 2.
  - Think why?



time complexity: search, maximum / minimum/ successor / predecessor, insertion, deletion

$O(h)$ , where  $h$  is the height of the tree.

worst case =  $O(n)$ , where  $n$  is the # of nodes

# Balanced Binary Search Tree

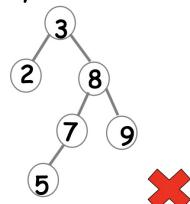
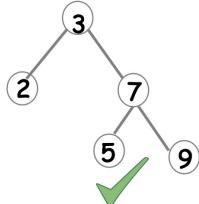
Goal: Keeping the height of the binary search tree low

Recap: A complete binary tree has height of  $O(\log n)$

Solution: relax the condition a little bit.

A binary search tree is balanced if:

- For every node in the tree, the height of the left subtree differs from the height of the right subtree by at most 1

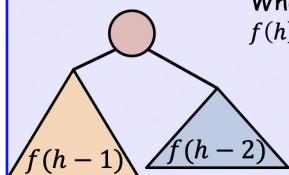


## Height of Balanced Binary Search Tree

Theorem 1: Given a balanced binary search tree  $T$  of  $n$  nodes, the height, or equivalently the depth, of  $T$  is  $O(\log n)$ .

Proof: Let  $f(h)$  be the minimum number of nodes of a balanced binary search tree of height  $h$ . Then, it is easy to verify that  $f(1) = 1, f(2) = 2$ .

For any  $h \geq 3$ , we have that  $f(h) = f(h-1) + f(h-2) + 1$



When  $h$  is even number:  
$$\begin{aligned}f(h) &> f(h-1) + f(h-2) \\&> 2f(h-2) \\&> 4f(h-4) \\&\dots \\&> 2^{\frac{h-1}{2}} \cdot f(2) = 2^{\frac{h}{2}}\end{aligned}$$

When  $h$  is odd number:  
$$\begin{aligned}f(h) &> f(h-1) \\&> 2^{\frac{h-1}{2}}$$

Therefore, given a balanced BST of  $n$  nodes of height  $h$ , we have:

$$n > 2^{\frac{h-1}{2}} \Rightarrow h < 2 \log_2 n + 1 \Rightarrow h = O(\log n)$$

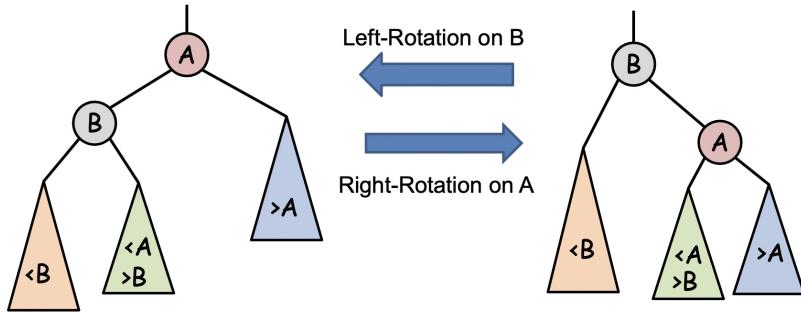
How to Keep a BST Balanced?

Dynamic Balancing: After updating the BST, we rebalance the BST using rotation

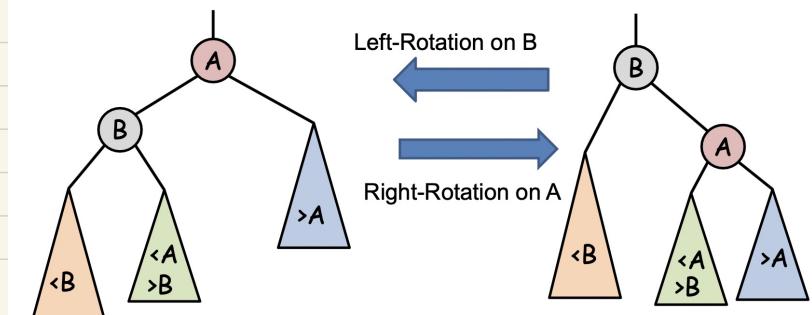
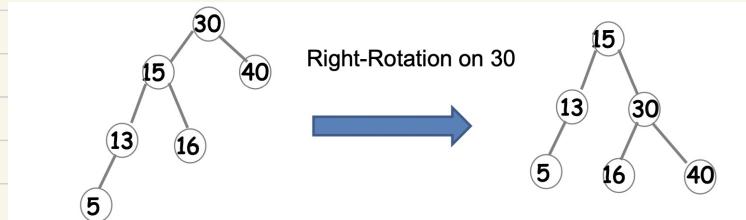


Rotation :

- Rotation allows us to change the structure without violating the binary search tree property



Rotation Example :



# Sorting Algothrim

## Selection sort

- Step 1: Scan all the  $n$  elements in the array to find the position  $i_{max}$  of the largest element  $maxnum$

$$i_{max} = 4, maxnum = 9$$

4	2	3	6	9	5
---	---	---	---	---	---

- Step 2: swap the position of the last one and  $maxnum$

4	2	3	6	5	9
---	---	---	---	---	---

- Step 3: We have a smaller problem: sorting the first  $n-1$  elements

4	2	3	6	5	sorted
---	---	---	---	---	--------

time complexity:

to find the maximum from  $n$  elements: a cost of  $n$

total cost for selection cost:

$$n + (n-1) + (n-2) + \dots + 1 = \frac{n(n+1)}{2}$$

time complexity:  $O(n^2)$

## Insertion sort:

- If we have a sorted array, and a new element comes, we **insert** the new element in the correct place

- If the first  $i$  elements in an array are sorted, we only need to insert the  $(i+1)$ -element to the right place to make the first  $(i+1)$  elements in the array are sorted

```
1 if arr.length <= 1 //Array is sorted if it has length <=1
2   return arr
3 for i from 1 to arr.length-1 //insert arr[i] to the right position
4   //move left if it is smaller than left element
5   for j from i downto 1
6     if arr[j] > arr[j-1]
7       break
8     temp ← arr[j]
9     arr[j] ← arr[j-1]
10    arr[j-1] ← temp
```

time complexity :  $O(n^2)$  (Worst case)

Let  $\text{arr}[i]$  to be left out of order (LOO) iff

$$\text{arr}[i] < \max\{\text{arr}[0], \text{arr}[1], \dots, \text{arr}[i-1]\}$$

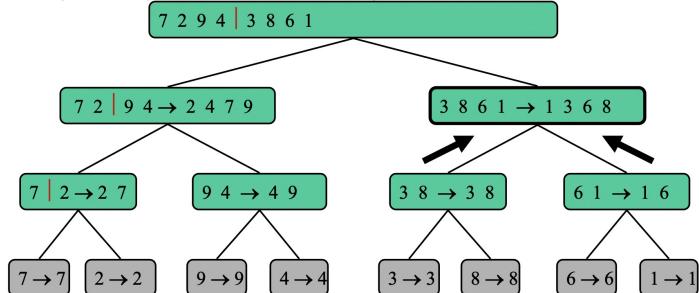
If we have  $K$  LOO array entries, the time complexity is  $O((k+1)n)$

Merge sort: High Level Idea

- Given  $n$  elements, we divide them into roughly two equal part
  - The first part:  $\lceil n/2 \rceil$  elements
  - The second part:  $\lfloor n/2 \rfloor$  elements
  - Sort the two subproblems recursively
    - Base case: array of size 1
- Then, combine the two sorted parts using an operation called merge.
  - After merge, the  $n$  elements are sorted

An example

- Step 1: Partition the initial array into two subarrays.
- Step 2: Sort each half
  - If the array only has one element, then it is sorted
  - Otherwise, return to Step 1 to sort it.
- Step 3: Merge the two sorted arrays



## Implementation:

- If the array only has one element, then it is sorted
- Otherwise, we partition them into two subarrays
  - Then using recursion for these two subarrays
  - Using the **left** and **right** variable to record the start and end positions of the subarray, respectively

Algorithm: `mergesort(arr, left, right)`

```
1 if left == right
2   return
3 middle ← (left+right)/2
4 mergesort(arr, left, middle)
5 mergesort(arr, middle+1, right)
6 merge(arr, left, middle, right)
```

## Merge Operation

- `merge(arr, 0, 3, 6)`
  - $\text{left}=0$ ,  $\text{middle}=3$ ,  $\text{right}=6$ , we use a **temparray** to store the merge result

$\text{left}=0 \quad \text{secondleft}=middle+1=4$

2	3	4	8	5	7	9
[0]	[1]	[2]	[3]	[4]	[5]	[6]

temparray

2						
[0]	[1]					

temparray

2	3					
[0]	[1]	[2]				

temparray

2	3	4				
[0]	[1]	[2]				

$\text{left}=1 \quad \text{secondleft}=4$

2	3	4	8	5	7	9
[0]	[1]	[2]	[3]	[4]	[5]	[6]

$\text{left}=2 \quad \text{secondleft}=4$

2	3	4	8	5	7	9
[0]	[1]	[2]	[3]	[4]	[5]	[6]

temparray

## Time Complexity:

- If we have  $n_1$  elements in the first subarray and  $n_2$  in the second subarray, what is the time complexity of merge?

$$O(n_1 + n_2)$$

$\text{left}=3 \quad \text{secondleft}=4$

2	3	4	8	5	7	9
[0]	[1]	[2]	[3]	[4]	[5]	[6]

temparray

2	3	4	5			
[0]	[1]	[2]	[3]			

...

temparray

2	3	4	5	7	8	9
[0]	[1]	[2]	[3]	[4]	[5]	[6]

## Merge Operation Implementation

Algorithm: `merge(arr, left, middle, right)`

```
1 n=right-left+1; start = left
2 temparray ← a new array of size n
3 secondleft = middle+1
4 filled = 0
5 while left <= middle and secondleft <=right //case 1: Neither of
6 the two subarray has reached the end
7     if arr[left] < arr[secondleft]
8         temparray[filled++] = arr[left++]
9     else
10        temparray[filled++] = arr[secondleft++]
11 while left <= middle //case 2: second subarray reached the end
12     temparray[filled++] = arr[left++]
13 while secondleft <= right //case 3: first subarray reached the end
14     temparray[filled++] = arr[secondleft++]
15 for i from 0 to n-1 //copy the sorted data back to arr
16     arr[start+i] = temparray[i]
17 free temparray // free the temparray
```

## Merge Operation Complexity Analysis

- Let  $f(n)$  be the number of basic operations executed by merge sort

Algorithm: `mergesort(arr, left, right)`

1	<code>if</code> left == right	$O(1)$
2	<code>return</code>	$O(1)$
3	<code>middle</code> ← (left+right)/2	$O(1)$
4	<code>mergesort(arr, left, middle)</code>	$f([n/2])$
5	<code>mergesort(arr, middle+1, right)</code>	$f([n/2])$
6	<code>merge(arr, left, middle, right)</code>	$O(n)$

What is the time complexity of merge sort? Which method shall we use to analyze the time complexity?

$$g(n) \leq a \cdot g\left(\left[\frac{n}{b}\right]\right) + O(n^\lambda)$$

If  $\log_b a < \lambda$ ,  $g(n) = O(n^\lambda)$

If  $\log_b a = \lambda$ ,  $g(n) = O(n^\lambda \cdot \log n)$

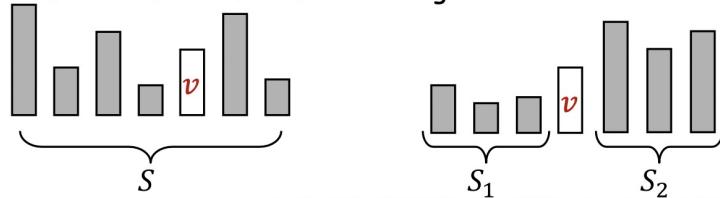
If  $\log_b a > \lambda$ ,  $g(n) = O(n^{\log_b a})$ .

$$f(n) \leq 2f\left(\left[\frac{n}{2}\right]\right) + c \cdot n$$

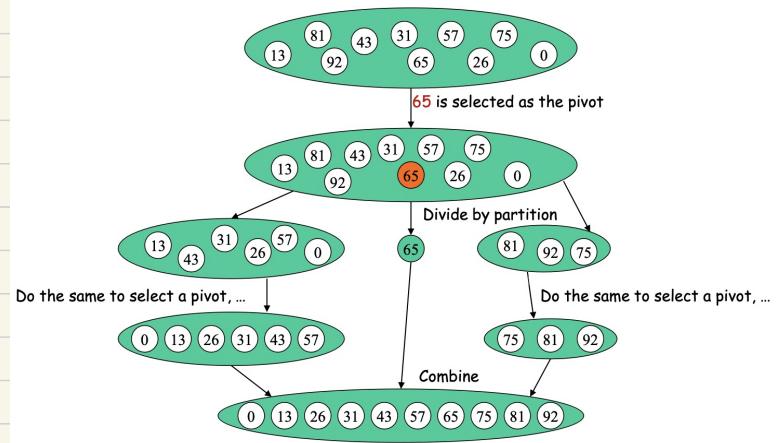
By master theorem:  $a = 2, b = 2, \lambda = 1$ .  $f(n) = O(n \cdot \log n)$

## Quick sort (divide and conquer)

- A randomized algorithm using divide-and-conquer
  - Time complexity:  $O(n \cdot \log n)$  expected running time
- High level idea: (we assume that elements are **distinct**)
  - Randomly pick an element, denoted as the **pivot**, and **partition** the remaining elements to three parts
    - The pivot
    - The elements in the left part: smaller than the pivot
    - The elements in the right part: larger than the pivot
    - For the left part and right part: repeat the above process if the number of elements is larger than 1



An example



Complexity Analysis :  $O(n \cdot \log n)$

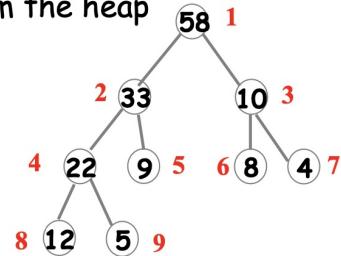
# Heap Sort & Sorting Lower Bound

Sorting with data structure

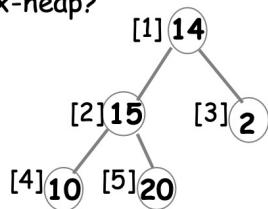
Max Heap : a max-tree + a complete binary tree

- To sort an array  $\text{arr}$ , we first create a heap  $H$  with a capacity of  $\text{arr.length}+1$
- Then, we insert  $\text{arr}[0]$  into  $H$ , insert  $\text{arr}[1]$  into  $H$ , ..., insert  $\text{arr}[\text{arr.length}-1]$  into  $H$ .
- Then, we repeatedly delete from the heap until the heap becomes empty

4	5	8	9	10	12	22	33	58
---	---	---	---	----	----	----	----	----



- In the previous solution, we need an additional array to create a max-heap, and then insert all numbers to this max-heap
  - Can we do faster to create a max heap?
  - Do we need to use two arrays?
- Reconsider an example of  $(14, 15, 2, 10, 20)$ . We can still view it as a complete binary tree.
  - Can we make a complete binary tree as a max-heap?

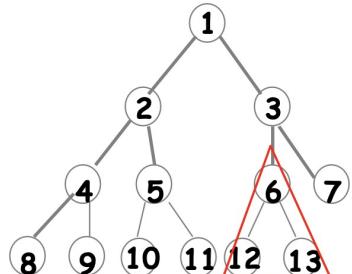


## ✓ 基于 Heap Adjust 实现 Heap sort

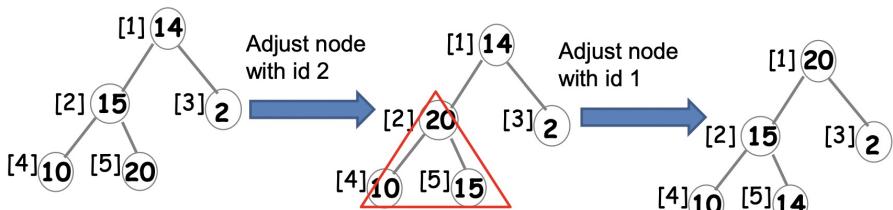
Heap Adjust: Given a node  $v$ , if its left-subtree and right-subtree are all max-tree, we can adjust the tree rooted at node  $v$  a max-tree

- Given node 6, both its left and right subtrees are max-trees (no violation). How to adjust to make the tree rooted as 6 a max-heap?
- Hint: How we do the heap adjustment after the deletion?

During the adjustment, swap the node to be adjusted with the larger child



- For a complete binary tree, we adjust from node with id  $\frac{n}{2}, \frac{n}{2} - 1, \frac{n}{2} - 2, \dots, 1$ .
- In the example of complete binary tree  $C_1$ , we only need to adjust from node with id 2 to id 1.



A complete binary tree  $C_1$

- The implementation is quite similar to heap deletion

## Heap Adjust Implementation

```

Algorithm 1: HeapAdjust(heap, nodeid, heapsize)
1. temp ← heap.arr[nodeid]
2. parent ← nodeid, child ← 2 * nodeid
3. While child <= heapsize
4.   if child < heapsize and heap.arr[child].key < heap.arr[child+1].key
5.     child ← child + 1 //choose the larger child
6.   if temp.key >= heap.arr[child].key
7.     break // no violation if we put at position 'parent'
8.   heap.arr[parent]←heap.arr[child]//swap larger child with parent
9.   parent ← child //attempt to insert at the child position
10.  child ← 2 * child
11.  heap[parent] ← temp
    
```

## Heap Sort Implementation

- First adjust the nodes from  $\frac{n}{2}, \frac{n}{2} - 1, \frac{n}{2} - 2, \dots, 1$  to make it a max-heap
  - Then, swap the largest number and the last element in the array, reduce the heap size by one and adjust the heap again.

Algorithm 1: *HeapSort(arr, n)*

```
1 //sort arr[1..n]
2 //adjust the complete binary tree so that it becomes a max-heap
3 for i ←  $\frac{n}{2}$  downto 1
4     adjust(arr, i, n)
5 //swap the root and last element, reduce the heap size, and adjust
6 for i ← n-1 downto 1 // i is the heap size
7     arr[i+1] ↔ arr[1] //swap the root and the last element
8     adjust(arr, 1, i) // adjust the heap of size i
```

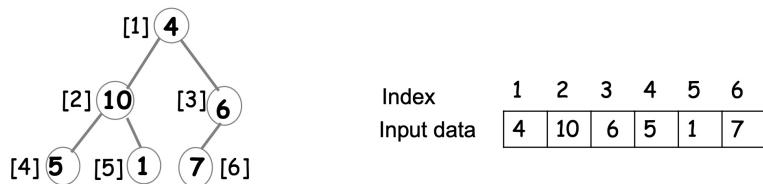
## Heap sort example

### Step 1

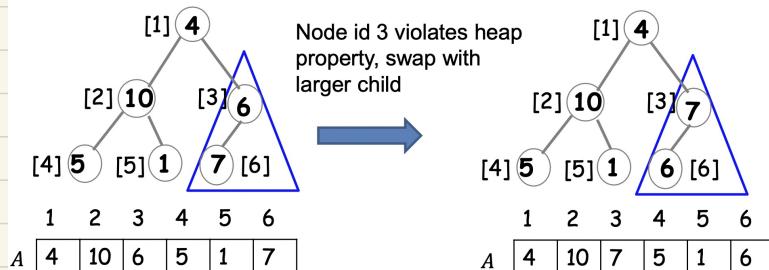
Sort the array  $A[1 \dots 6] = [4, 10, 6, 5, 1, 7]$  in ascending order by heap sort.

- Step 1: Adjust the heap to create a max heap. Given  $n$  nodes, we adjust from node  $n/2$ , then node  $n/2-1, \dots$ , and finally to node 1.

First, the complete binary tree corresponding to the array  $A[1 \dots 6] = [4, 10, 6, 5, 1, 7]$  is:

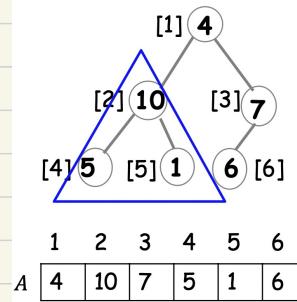


Adjust max heap from node id 3: Adjust the subtree rooted at node id 3

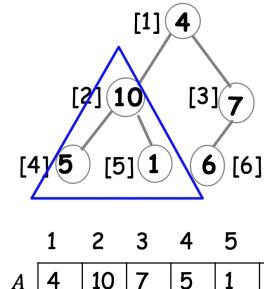


After adjust from node id 3: The subtree rooted at node 3 is a max heap, the array becomes:

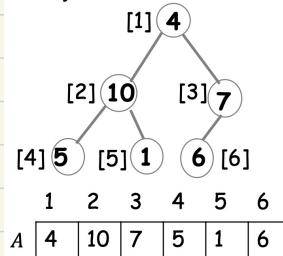
Adjust max heap from node id 2:  
Adjust the subtree rooted at node 2



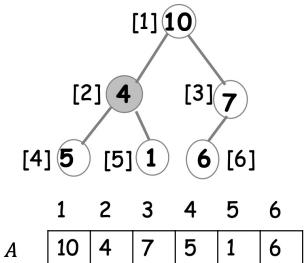
No violation



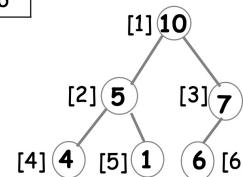
Adjust max heap from node id 1:  
Adjust the subtree rooted at node id 1



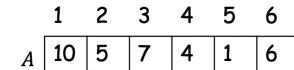
Node id 1 violates  
heap property, swap  
with larger child



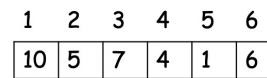
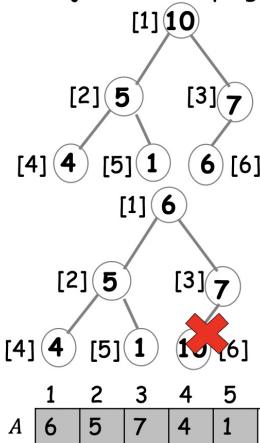
After swapping, node id 2  
violates the heap property,  
swap with larger child



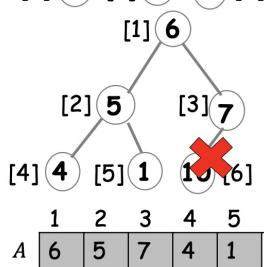
No violation. The  
array is now:



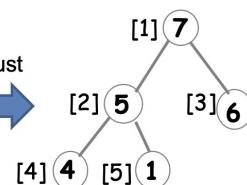
- Step 2: sorting. Repeatedly swap the root (the first element) and the last element in the array, reduce the heap size by one and adjust the heap again.



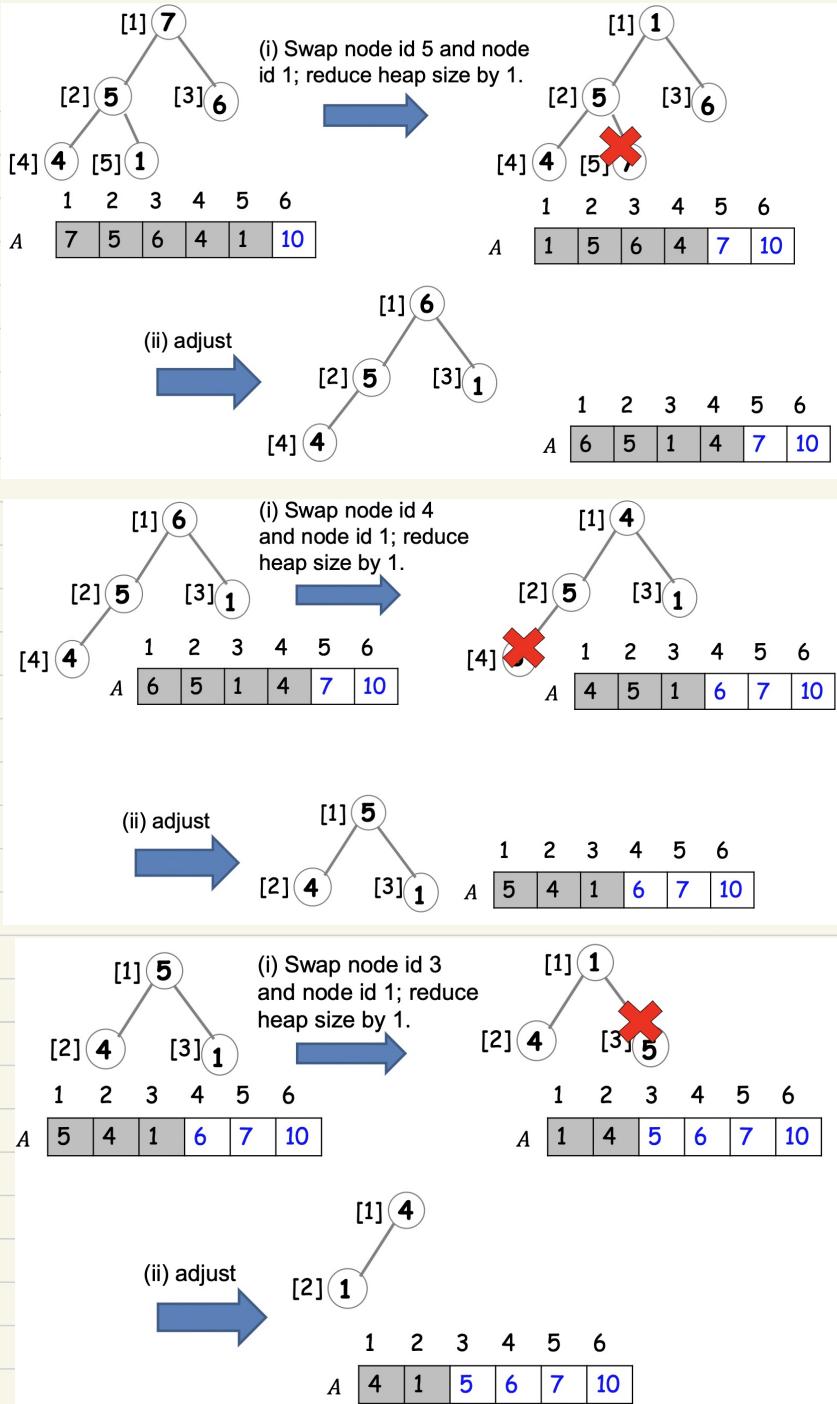
(i) Swap node id 6 and node id  
1; reduce heap size by 1.

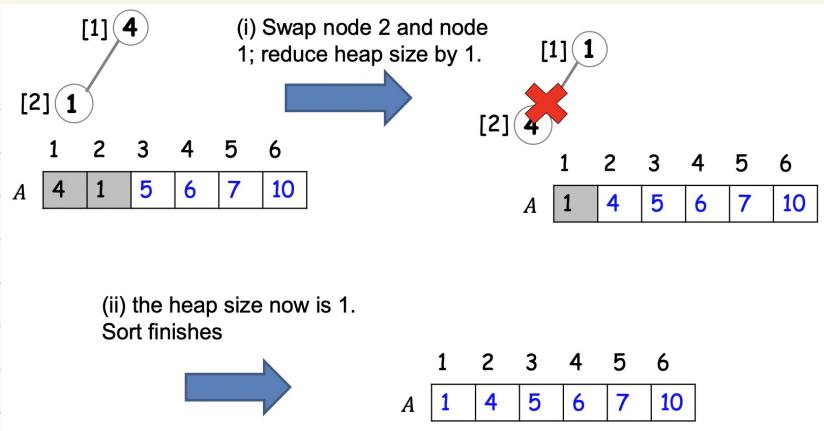


(ii) adjust



Step 2





### Heap Sort Complexity Analysis

adjust for  $\frac{n}{2}$  nodes to make the complete binary tree a max-heap:  
 $C_{\text{adjust}} = O(n \cdot \log n)$

then repeated swap the root and the last element , and do the adjustment:  
 $C_{\text{sort}} = O(n \cdot \log n)$

total cost:

$$C_{\text{adjust}} + C_{\text{sort}} = O(n \cdot \log n)$$

# Hashing

Searching cost analysis : current best time complexity :  $O(\log n)$

To preprocess the input (to sort / build the binary search tree)  
 $O(n \log n)$  if we have  $n$  records

To search the record  
 $O(\log n)$

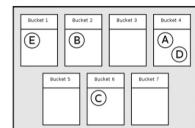
What hashing achieves ?

To preprocess the input  
 $O(n)$  if we have  $n$  records

To search the record  
 $O(1)$  in expectation

Hash Tables : The main idea

- In direct-addressing, the record with key  $k$  is stored in slot  $k$  of the array  $T$ , i.e.,  $T[k]$
- In hashing, the element is stored in slot  $h(k)$ , i.e.,  $T[h(k)]$ , where  $h$  is a hash function.
  - Assume keys are integers in the range of  $[0, U - 1]$
  - Denote by  $[x]$  the set of integers from  $0$  to  $x - 1$ .
  - A hash function  $h$  is a function from  $[U]$  to  $[m]$ 
    - For any integer  $k$ ,  $h(k)$  returns an integer in  $[m]$
    - The value  $h(k)$  is called the hash value of  $k$
    - $U > m$



Problem: hashing collision

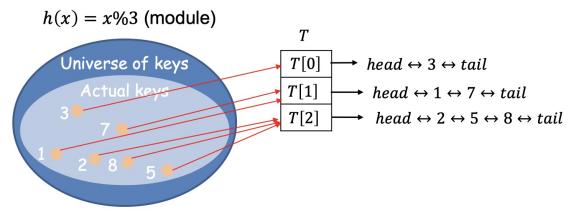
- If we have a set  $S$  of keys  $\{1, 2, 3, 5, 7, 8\}$ , and the hash function is  $h(x) = x \% 3$  (modulo)

- What are the hash values of the integers in  $S$ ?
  - $h(1) = 1 \% 3 = 1$
  - $h(2) = 2 \% 3 = 2$
  - $h(3) = 3 \% 3 = 0$
  - $h(5) = 5 \% 3 = 2$
  - $h(7) = 7 \% 3 = 1$
  - $h(8) = 8 \% 3 = 2$
- What are  $U$  and  $m$ ?

## Hashing Collision Resolution: Chaining & Open Addressing

### Resolution 1: Chaining

- Collision: two keys hash to the same slot
- Chaining: We place all elements that hash to the same slot into the same linked list



Hash Table Initialization w. Chaining  
 $\text{insert}(\text{ hashtable}, \text{key}, \text{record})$   
 $\text{search}(\text{ hashtable}, \text{key})$   
 $\text{delete}(\text{ hashtable}, \text{key})$

- Assume that the hash function  $h$  hashes keys to the range  $[0, m - 1]$
- We then create an array of size  $m$  where each entry stores the address of an empty linked list.

#### Algorithm: `createTable(sizem)`

```

1 hashtable ← allocate an array of size sizem
2 for i from 0 to sizem-1
3   hashtable[i] ← allocate an empty linkedlist
4 return hashtable

```

## Analysis of Hashing w. Chaining

- Load factor  $\alpha$ : the average number of elements stored in a chain
  - $\alpha = \frac{n}{m}$
- Assumption: Uniform hashing of  $h(k)$ 
  - Elements are equally likely to hash into any of the  $m$  slots.
  - $L_i$ : the linked list containing the elements hashes to  $i$ .

Theorem 1: In a hash table with collision resolved by chaining, the search/insertion/deletion takes  $O(1 + \alpha)$  time in expectation if  $h(k)$  is uniform hashing.

- By choosing  $m$  so that  $\frac{n}{m} = \alpha = O(1)$ , the query time is  $O(1)$ .

## Resolution 2: Open Addressing

- All elements are stored in the hash table.
  - Unlike chaining, no elements are stored outside the table.
  - The hash table may "fill up" such no insertion can be made
    - Load factor  $\alpha$  is always smaller than 1.
- For insertion, we examine, or **probe**, the hash table until an empty slot is found to put the key and records.
  - How to probe the hash table?
    - Linear probing
    - Quadratic Probing (Not discussed in the lecture)
    - Double Hashing
- Deletion: not efficiently supported. Think why?
- ADT Implementation : refer to our tutorial

### (a) Linear Probing

- For **insertion**, we probe  $h(k), h(k) + 1, h(k) + 2, \dots, m, 1, 2, \dots, h(k) - 1$  one by one until we find an empty slot, and insert the record to this slot.
  - More formally we probe  $h(k, i) = (h(k) + i) \% m$  from  $i = 0$  to  $i = m - 1$  until we find an empty slot to insert.
- When **searching** for a record with a certain key,
  - Compute  $h(k)$
  - Examine the hash table buckets in order  $T[h(k, i)]$  for  $0 \leq i \leq m - 1$  until one of the following happens:
    - $T[h(k, i)]$  has the record with key equal to  $k$ .
    - $T[h(k, i)]$  is empty, then no record contains key  $k$  in the hash table

### (b) Double Hashing

- Deficiency of Linear Probing
  - Long sequence of occupied slots, which degrades the query efficiency
- Double hashing
  - We have an additional hash function  $h' > 0$ .
  - **Insertion:** we probe  $h(k, i) = (h(k) + i \cdot h'(k)) \% m$  one by one for  $i$  from 0 to  $m - 1$  until an empty slot is found.
  - **Search:** we search  $h(k, i)$  for  $i$  from 0 to  $m - 1$  until one of the following happens:
    - $T[h(k, i)]$  has the record with key equal to  $k$ .
    - $T[h(k, i)]$  is empty, then no record contains key  $k$  in the hash table

## Analysis of Open Addressing

- Load factor  $\alpha$ : If the hash table  $T$  has size  $m$  and we store  $n$  elements in the hash table

$$\alpha = \frac{n}{m}$$

- Assumption: Uniform hashing of  $h(k, i)$

- The probing sequence  $\{h(k, 0), h(k, 1), h(k, 2), \dots, h(k, m-1)\}$  used to insert or search for each key  $k$  is equally likely to be any permutation of  $\{0, 1, 2, \dots, m-1\}$ .

Theorem 2: In a hash table with collision resolved by open addressing, the search takes  $O(\frac{1}{1-\alpha})$  time in expectation if  $h(k, i)$  is uniform hashing.

- If  $\alpha = \Theta(1)$ , the query time is  $O(1)$

## Chaining vs. Open Addressing

- Pros of Chaining

- Less sensitive to hash functions and load factors ( $\alpha$  can be larger than 1), while open addressing requires to avoid long probes, and its load factor  $\alpha < 1$ .
- Support deletion, while open addressing is difficult to support deletion (Why).

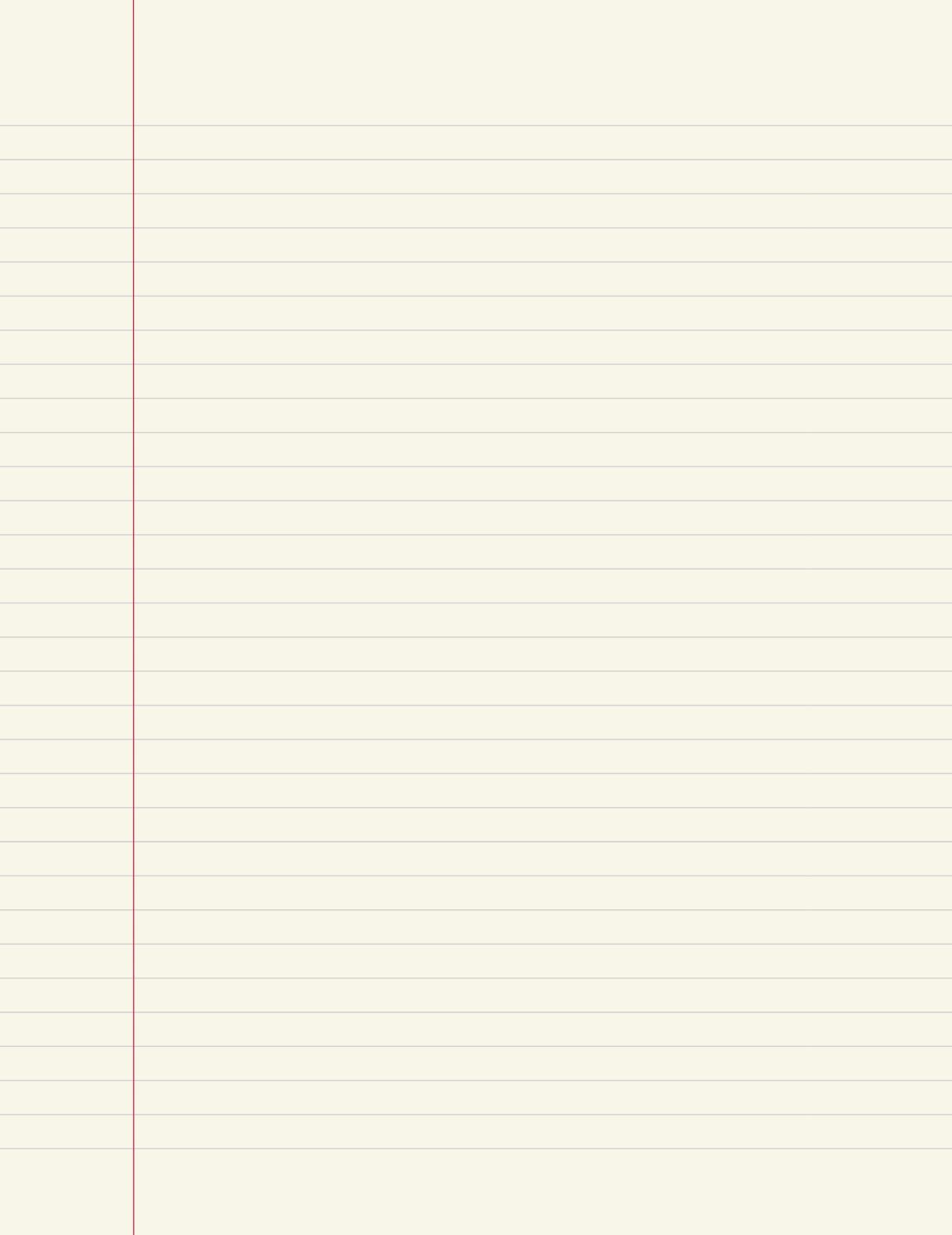
- Pros of Open addressing

- Usually much faster than chaining

## Designing Hash Functions

### Good Hash Function:

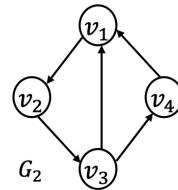
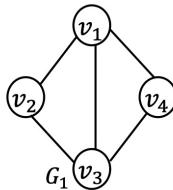
- A good hash function satisfies the uniform hashing property:
  - Each key is equally likely to hash to any of the  $m$  slots, independent of where other keys will hash to.
- We learn two hashing functions: Division and universal hashing
  - Division is effective in practice without any theoretical guarantee on  $O(1)$  query time.
  - Universal hashing provides theoretical guarantees on  $O(1)$  query time



# Graph Basics & BFS Traversal

Graph def:

- A graph  $G$  is defined as a pair of  $(V, E)$  where:
  - $V$  is the set of objects, each of which is called a **node** (or a **vertex**)
  - $E$  is the set of **edges**, where each edge is a pair of two node  $u$  and  $v$
- Notations
  - $n$ : the number of nodes, i.e.,  $|V|$
  - $m$ : the number of edges, i.e.,  $|E|$



undirected / directed

Terms : neighbor and degree

- Neighbor:  $v$  is called a **neighbor** of  $u$  if there is an edge between  $v$  and  $u$ .
  - In directed graphs,  $v$  is called the **out-neighbor** of  $u$  if there is an edge  $\langle u, v \rangle$ ;  $v$  is called the **in-neighbor** of  $u$  if there is an edge  $\langle v, u \rangle$ .
- **Degree**  $d(v)$  of a node  $v$ : the number of neighbors of this node  $v$ .
  - In directed graphs, the **out-degree**  $d_{out}(v)$  of a node  $v$  is the number of out-neighbors of this node; the **in-degree**  $d_{in}(v)$  of a node  $v$  is the number of in-neighbors of this node.

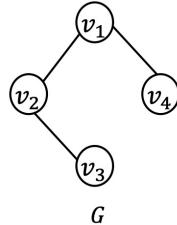
**Theorem 1:** In a undirected graph, the number  $m$  of edges is equal to  $\frac{1}{2} \sum_{v \in V} d(v)$ . In a directed graph,  $m = \sum_{v \in V} d_{out}(v) = \sum_{v \in V} d_{in}(v)$ .

path

- A **path** from node  $u$  to node  $v$  in a graph  $G$  is a sequence of nodes,  $(u, v_1, \dots, v_k, v)$  such that there exist a sequence of edges
  - $((u, v_1), (v_1, v_2), \dots, (v_k, v))$  if  $G$  is an undirected graph;
  - $((u, v_1), (v_1, v_2), \dots, (v_k, v))$  if  $G$  is a directed graph.
- A **simple path** is a path in which all nodes except the first and last are distinct.
- A **cycle** is a simple path in which the first and the last nodes are the same.

### Adjacency list for undirected graph

- Each node  $v \in V$  is associated with a linked list that stores all neighbors of  $v$ , we map an ID for each node.

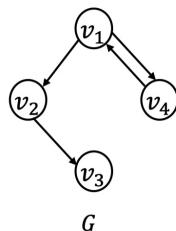


$v_1$	$\rightarrow$ head $\leftrightarrow v_2 \leftrightarrow v_4 \leftrightarrow tail$
$v_2$	$\rightarrow$ head $\leftrightarrow v_1 \leftrightarrow v_3 \leftrightarrow tail$
$v_3$	$\rightarrow$ head $\leftrightarrow v_2 \leftrightarrow tail$
$v_4$	$\rightarrow$ head $\leftrightarrow v_1 \leftrightarrow tail$

- Space:  $O(n + m)$ , where  $n$  is the number of nodes and  $m$  is the number of edges

### Adjacency list for directed graph

- Each node  $v \in V$  is associated with a linked list that stores all out-neighbors of  $v$



$v_1$	$\rightarrow$ head $\leftrightarrow v_4 \leftrightarrow v_2 \leftrightarrow tail$
$v_2$	$\rightarrow$ head $\leftrightarrow v_3 \leftrightarrow tail$
$v_3$	$\rightarrow$ head $\leftrightarrow tail$
$v_4$	$\rightarrow$ head $\leftrightarrow v_1 \leftrightarrow tail$

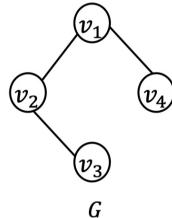
- Space:  $O(n + m)$ , where  $n$  is the number of nodes and  $m$  is the number of edges

Graph Representation  
Adjacency List

## Adjacency matrix for undirected graph

- A  $n \times n$  two dimensional matrix  $A$  where  $A[u][v] = 1$  if  $(u, v) \in E$ , or 0 otherwise.

Adjacency Matrix

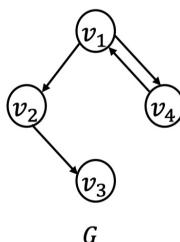


	$v_1$	$v_2$	$v_3$	$v_4$
$v_1$	0	1	0	1
$v_2$	1	0	1	0
$v_3$	0	1	0	0
$v_4$	1	0	0	0

- $A$  is symmetric
- Space:  $O(n^2)$  where  $n$  is the number of nodes.

## Adjacency matrix for directed graph

- A  $n \times n$  two dimensional matrix  $A$  where  $A[u][v] = 1$  if  $\langle u, v \rangle \in E$ , or 0 otherwise.



	$v_1$	$v_2$	$v_3$	$v_4$
$v_1$	0	1	0	1
$v_2$	0	0	1	0
$v_3$	0	0	0	0
$v_4$	1	0	0	0

- $A$  may not be symmetric
- Space:  $O(n^2)$  where  $n$  is the number of nodes

Comparison :

### Adjacency list:

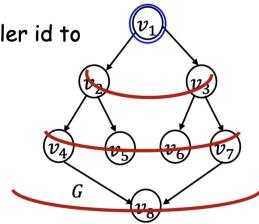
- Space:  $O(n + m)$ , save space if the graph is sparse, i.e.,  $m \ll n^2$
- Check the existence of an edge  $(u, v)$ :  $O(k)$  time where  $k$  is the number of neighbors of  $v$
- Retrieve the neighbors of a node:  $O(k)$  time.
- Add/delete a node:  $O(n)$
- Add/delete an edge:  $O(1)$

### Adjacency matrix:

- Space consumption:  $O(n^2)$
- Check the existence of an edge  $(u, v)$ :  $O(1)$  time.
- Retrieve the neighbors of a node:  $O(n)$  time
- Add/delete a node:  $O(n^2)$ , (create a new matrix)
- Add/delete an edge:  $O(1)$

## Graph Traversal : Breadth-First Search (BFS)

- Intuition of BFS
  - Given a source node  $s$ , always visit nodes that are closer to the source  $s$  first before visiting the others.
- The result is not unique, if we do not define an order among out-going edges from a node.
  - Possible results
    - $v_1, v_2, v_3, v_4, v_5, v_6, v_7, v_8$
    - $v_1, v_3, v_2, v_7, v_6, v_5, v_4, v_8$
  - We impose an order by going from smaller id to larger id. The result then will be unique.

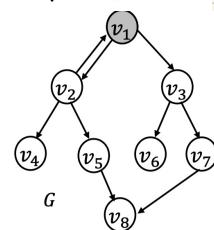


- At the beginning, color all nodes to be white
- Create a queue  $Q$ , enqueue the source  $s$  to  $Q$ , and color the source to be gray (meaning  $s$  is in the queue)

Assume the source is  $v_1$ .

Example :

$Q: (v_1)$



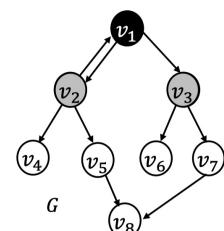
Repeat the following until queue  $Q$  is empty

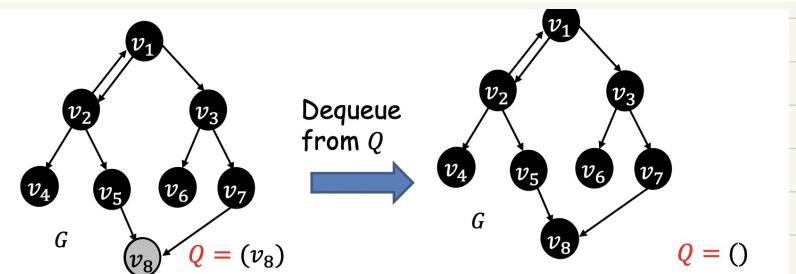
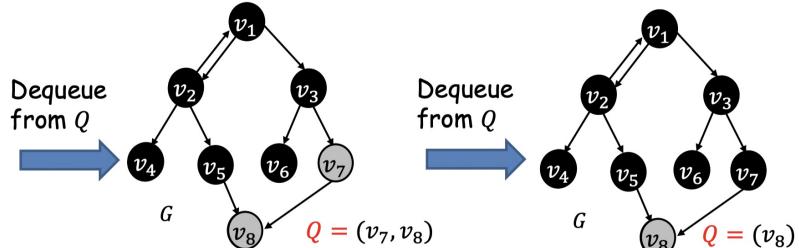
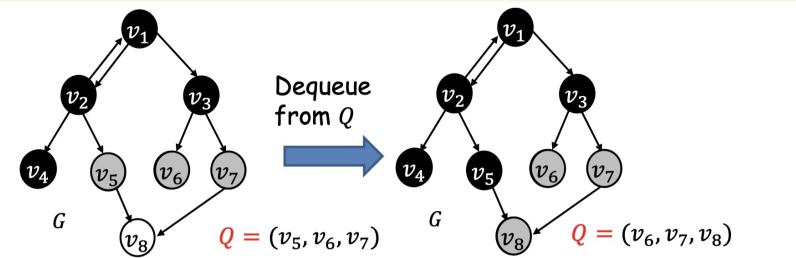
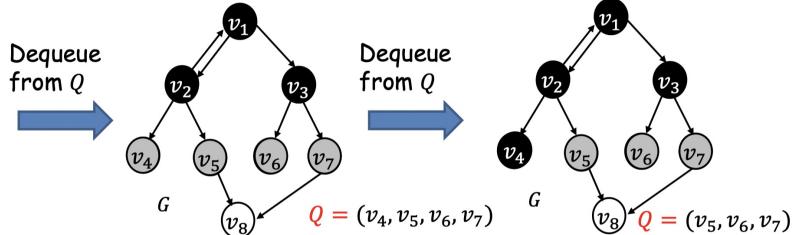
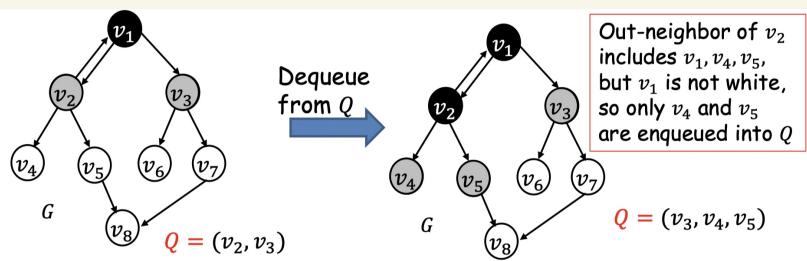
- Dequeue from  $Q$ , let the node be  $v$ .
- For every out-neighbor  $u$  of  $v$  that is still white
  - Enqueue  $u$  into  $Q$ , and color  $u$  to gray (to indicate  $u$  is in queue)
- Color  $v$  to be black (meaning  $v$  has finished).

$Q = (v_1)$

After dequeuing  $v_1$

$Q = (v_2, v_3)$





Now  $Q$  is empty  
BFS finishes

## Time complexity

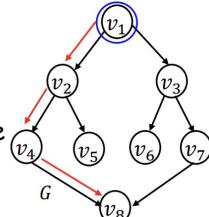
- When a node  $u$  is dequeued
  - We examine all of its neighbors (check their color), enqueue them and color them to gray if they are white.
  - After that, we color  $u$  as black
  - This incurs  $c(1 + d_{out}(u))$  costs for node  $u$ .
- Each node is dequeued at most once.
  - Why?
  - We enqueue a node at most once. If it is in the queue, its color is gray and we will not further enqueue it.
- Therefore, the total running time is
  - $\sum_{u \in V} c(1 + d_{out}(u)) = c(n + m) = O(n + m)$

# Depth-First Search

DFS:

- Going along one path until we cannot go further

- Imposing an order to make the traversal unique: from smaller id to larger id
  - Visiting order:  $v_1, v_2, v_4, v_8, v_5, v_3, v_6, v_7$



- We still focus on directed graph
  - Extension to undirected graph will be straightforward



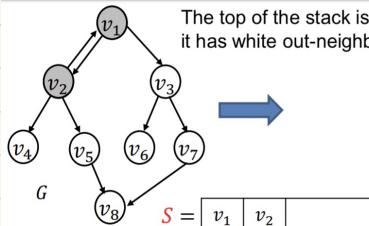
- Initialization:

- At the beginning, color all nodes to be white
- Create a stack  $S$ , push the source  $s$  to  $S$ , and color the source to be gray (meaning  $s$  is in the stack)

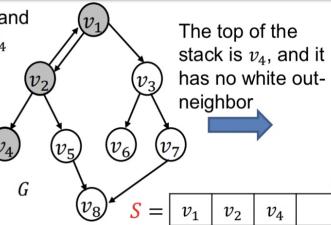
- Example:

- Assume that  $v_1$  is the source

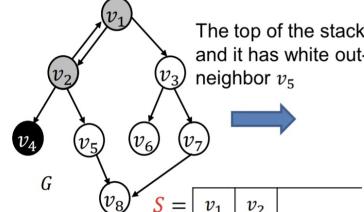
$$S = \boxed{v_1} \quad \boxed{\phantom{v_1}}$$



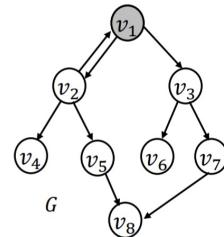
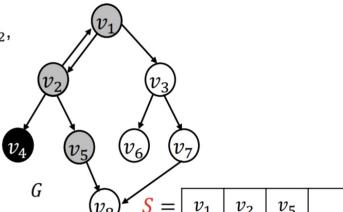
The top of the stack is  $v_2$ , and it has white out-neighbor  $v_4$

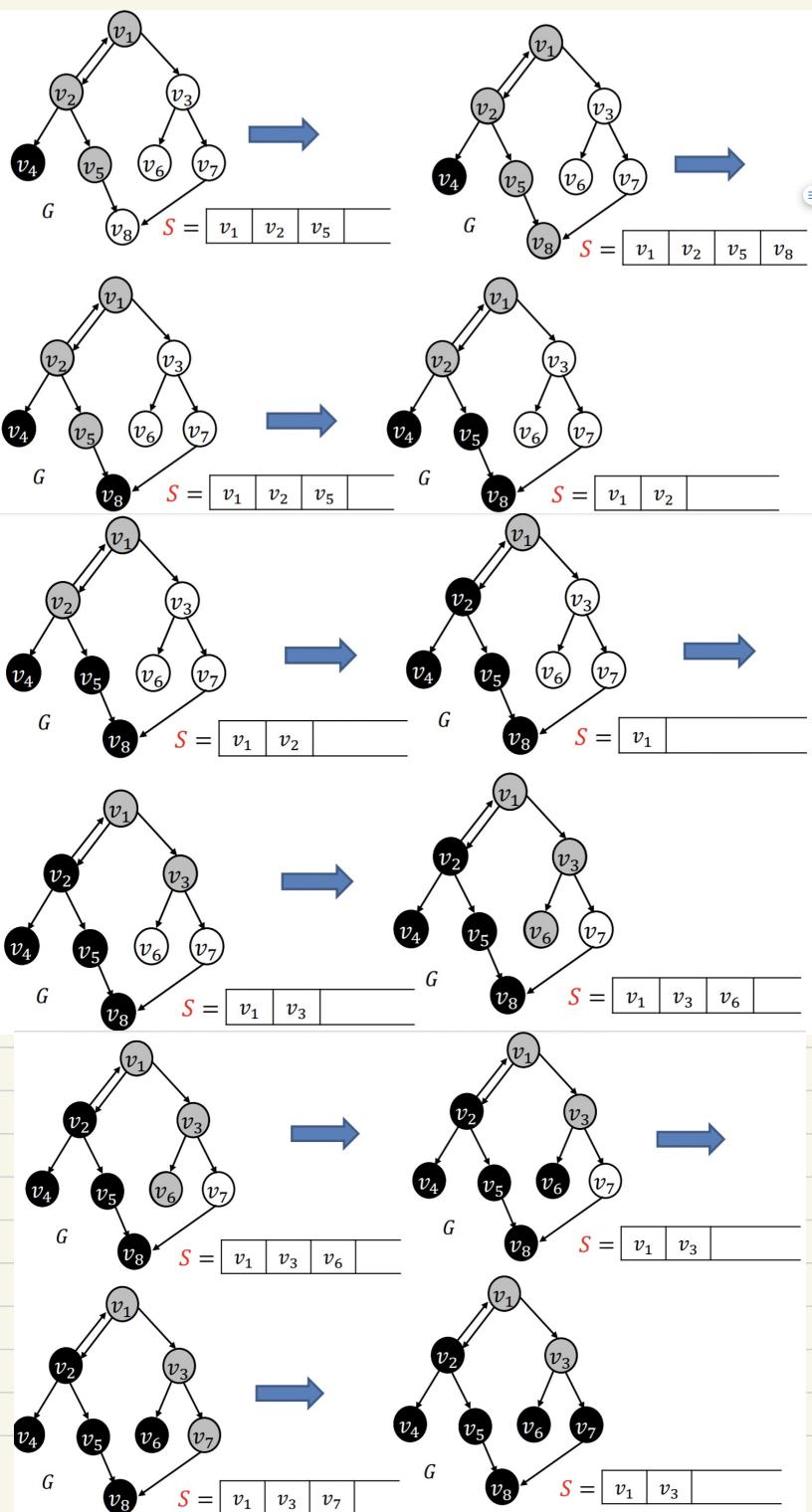


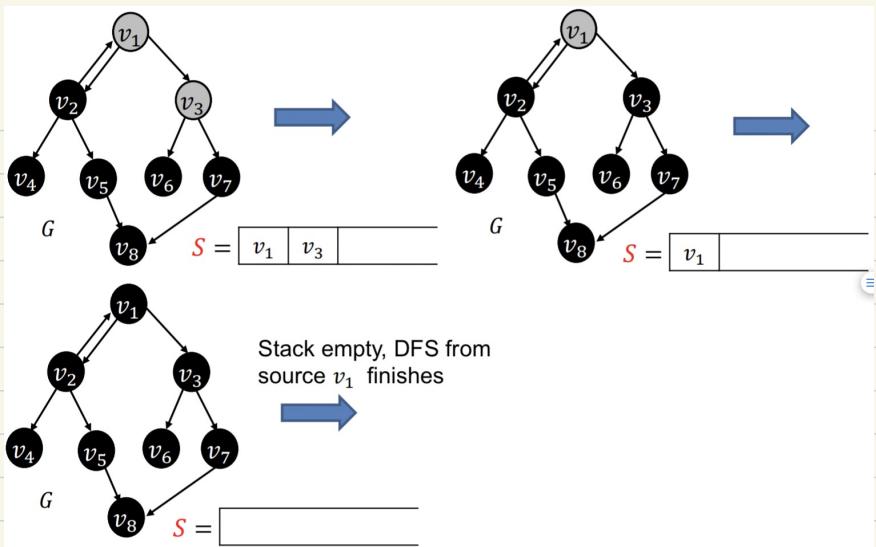
The top of the stack is  $v_4$ , and it has no white out-neighbor



The top of the stack is  $v_2$ , and it has white out-neighbor  $v_5$







### Complexity Analysis:

- When a node  $v$  get popped from the stack?
  - None of its out-neighbors is a white node
  - We may need to repeated check if node  $v$  has white out-neighbor. If  $v$  has  $d_{out}(v)$  out-neighbors, we need to check  $d_{out}(v)$  times in the worst case.
    - Cost:  $d_{out}(v) \cdot d_{out}(v)$ ?
    - Can we do better?
      - We record the position checked last time. All nodes in previous positions will not be white.
      - Cost:  $O(d_{out}(v))$
- Each node is popped at most once.
- Therefore, the total running time is
  - $\sum_{u \in V} c(1 + d_{out}(u)) = c(n + m) = O(n + m)$