PRECAIT FAST TRACK BATCH OM22

Data Structure SunBeam Institute of Information & Technology, Hinjwadi, Pune & Karad. SACHIN PAWAR



PreCAT Scope: Data Structure

- In Section B, 7 Questions are reserved for this subject and mostly all questions are pseudocode oriented (i.e. theory based/concepts).
- In this course the main focus is on **basic data structure's algorithms** and **introduction to an advanced data structures** to build a base which is required to learn and implement advanced data structures and algorithms in CDAC courses.



+ Introduction

- Data structure
- Algorithm and analysis of an algorithm

+ Array

- Concept & definition
- Searching Algorithms:
 - 1. Linear Search
 - 2. Binary Search

- Sorting Algorithms:

- 1. Selection Sort
- 2. Bubble Sort
- 3. Insertion Sort
- 4. Quick Sort
- 5. Merge sort



+ Linked List

- Concept & definition
- Types of Linked List
- Operations on Linked List
- Differnce between an array and linked list

+ Stack

- Concept & definition
- Implementation of stack data structure
- Stack applications algorithms:
- 1. Conversion of infix expression into its equivalent prefix
- 2. Conversion of infix expression into its equivalent postfix
- 3. Conversion of prefix expression into its equivalent postfix
- 4. Postfix expression evalution



+ Queue

- Concept & definition
- Types of queue
- Implementation of queue data structure
- Applications of queue

+ Introduction to an advanced data structure

- Tree
- Graph



Q. What is Data Structure?

Data Structure is a way to store data elements into the memory (i.e. into the main memory) in an organized manner so that operations like addition, deletion, traversal, searching, sorting etc... can be performed on it efficiently.



Two types of **Data Structures** are there:

- 1. Linear/Basic Data Structures: data elements gets stored into the memory in a linear manner (e.g. sequentially) and hence can be accessed linearly/sequentially.
 - Array
 - Structure & Union
 - Linked List
 - Stack
 - Queue
- 2. Non-linear/Advanced Data Structures: data elements gets stored into the memory in a non-linear manner (e.g. hierarchical) and hence can be accessed non-linearly.
 - Tree (Hierarchical)
 - Graph
 - Hash Table
 - Binary Heap



Array: It is a basic/linear data structure which is a collection/list of logically related similar type of elements in which data elements gets stored into the memory at contiguos locations.

Structure: It is a basic/linear data structure which is a collection/list of logically related similar and disimmilar type of elements gets stored into the memory collectively (as a single entity/record).

Size of of the structure = sum of size of all its members.

Union: Union is same like structure, except, memory allocation i.e. size of union is the size of max size member defined in it and that memory gets shared among all its members for effective memory utilization (can be used in a special case only).



Q. What is a Program?

- A program is a finite set of instructions written in any programming language (like C, C++, Java, Python, Assembly etc...) given to the machine to do specific task.

Q. What is an Algorithm?

- An algorithm is a finite set of instructions written in human understandable language (like english), if followed, acomplishesh a given task.
- An algorithm is a finite set of instructions written in human understandable language (like english) with some programming constraints, if followed, acomplishesh a given task, such an algorithm also called as pseudocode.
- An algorithm is a template whereas a program is an implementation of an algorithm.



Example: An algorithm to do sum of all array elements

```
Algorithm ArraySum(A, n)//whereas A is an array of size n

{
    sum=0;//initially sum is 0
    for( index = 1; index <= size; index++) {
        sum += A[ index ];//add each array element into the sum
    }
    return sum;
}
```

- In this algorithm, **traversal/scanning** operation is applied on an array. Initially sum is 0, each array element gets added into to the sum by traversing array sequentially from the first element till last element and final result is returned as an output.



- **Analysis of an algorithm** is a work of determining how much **time** i.e. computer time and **space** i.e. computer memory it needs to run to completion.
- There are two measures of an analysis of an algorithms:
- **1. Time Complexity** of an algorithm is the amount of time i.e. computer time required for it to run to completion.
- **2. Space Complexity** of an algorithm is the amount of space i.e. computer memory required for an algorithm to run to completion.
- Asymptotic Analysis: It is a mathematical way to calculate time complexity and space complexity of an algorithm without implementing it in any programming language.
- In this type of analysis, analysis can be done on the basis of **basic operation** in that algorithm.
- e.g. in searching & sorting algorithms comparison is the basic operation and hence analysis gets done on the basis of no. of comparisons, in addition of matrices algorithms addition is the basic operation and hence on the basis of addition operation.



- "Best case time complexity": if an algo takes min amount of time to run to completion then it is referred as best case time complexity.
- "Worst case time complexity": if an algo takes max amount of time to run to completion then it is referred as worst case time complexity.
- "Average case time complexity": if an algo takes neither min nor max amount of time to to run to completion then it is referred as an average case time complexity.

"Asympotic Notations":

- 1. Big Omega (Ω): this notation is used to denote best case time complexity also called as asymptotic lower bound
- 2. Big Oh (O): this notation is used to denote worst case time complexity also called as asymptotic upper bound
- 3. Big Theta (θ) : this notation is used to denote an average case time complexity also called as **asymptotic tight bound**



1. Linear Search/Sequential Search:

```
Step-1: accept key from the user
```

Step-2: compare the value of key with each array element sequentially by traversing it from the first element till either key is found or maximum till the last element. If key is found then return true otherwise return false.

```
Algorithm LinearSearch( A, size, key){
  for( index = 1 ; index <= size ; index++ ){
   if( key == A[ index ] )
    return true;
  }
  return false;
}</pre>
```



Best Case: If key is found at very first position in only 1 no. of comparison then it is considered as a best case and running time of an algorithm in this case is O(1) => and hence time complexity = $\Omega(1)$

Worst Case: If either key is found at last position or key does not exists, maximum \mathbf{n} no. of comparisons takes place, it is considered as a worst case and running time of an algorithm in this case is O(n) => and hence time complexity = O(n)

Average Case: If key is found at any in between position it is considered as an average case and running time of an algorithm in this case is O(n/2) => and hence time complexity $= \Theta(n)$



2. Binary Search/Logarithmic Search:

- This algorithm follows divide-and-conquer approach.
- To apply binary search on an array prerequisite is that array elements must be in a sorted manner.

Step-1: accept key from the user

Step-2: calculate mid position of an array by the formula, mid = (left+right)/2 (by means of calculating mid position big size array has been divided logically into two subarrays, from left to mid-1 = left subarray & mid+1 to right = right subarray)

Step-3: compare the value of key with element which is at mid position. if key matches with element at mid position means key is found and return true.

Step-4: if key do not matches then check, is the value of key is less than element which is at mid position, if yes then goto search key only into the left subarray by skipping whole right subarray otherwise (means of the value of key is greater than element which is at mid position) goto search key only into the right subarray by skipping whole left subarray.

Step-5: repeat Step-2, Step-3 & Step-4 till either key is found or max till the subarray is valid, if subarray becomes invalid means key is not found and hence return false in this case.



```
Algorithm BinarySearch(A, n, key) //A is an array of size "n", and key to be search
  left = 1;
  right = n;
  while( left <= right )</pre>
    //calculate mid position
    mid = (left+right)/2;
    //compare key with an ele which is at mid position
    if( key == A[ mid ] )//if found return true
      return true;
    //if key is less than mid position element
    if( key < A[ mid ] )</pre>
      right = mid-1; //search key only in a left subarray
    else//if key is greater than mid position element
      left = mid+1;//search key only in a right subarray
  }//repeat the above steps either key is not found or max any subarray is valid
  return false;
```



- as in each iteration one comparison takes place and search space is getting reduced by half.

```
n => n/2 => n/4 => n/8 \dots
after iteration-1 => n/2 + 1 => T(n) = (n/2^1) + 1
after iteration-2 => n/4 + 2 => T(n) = (n/2^2) + 2
after iteration-3 => n/8 + 3 => T(n) = (n/2^3) + 3
after k iterations => T(n) = (n/2k) + k \dots (equation-1)
let us assume,
=> n = 2k
=> \log n = \log 2^k (by taking log on both sides)
=> \log n = k \log 2
=> \log n = k
=> k = log n
By substituting value of n & k in equation-I, we get
=> T(n) = (n / 2^k) + k
=> T(n) = (2k / 2k) + \log n
=> T(n) = 1 + \log n => T(n) = \log n => T(n) = O(\log n).
```



Best Case: if the key is found in very first iteration at mid position in only 1 no. of comparison it is considered as a best case and running time of an algorithm in this case is $O(1) = \Omega(1)$.

Worst Case: if either key is not found or key is found at leaf position it is considered as a worst case and running time of an algorithm in this case is $O(\log n) = O(\log n)$.

Average Case: if key is found at non-leaf position it is considered as an average case and running time of an algorithm in this case is $O(\log n) = \Theta(\log n)$.



1. Selection Sort:

- In this algorithm, in first iteration, first position gets selected and element which is at selected position gets compared with all its next position elements, if selected position element found greater than any other position element then swapping takes place and in first iteration smallest element gets setteled at first position.
- In the second iteration, second position gets selected and element which is at selected position gets compared with all its next position elements, if selected position element found greater than any other position element then swapping takes place and in second iteration second smallest element gets setteled at second position, and so on in maximum (n-1) no. of iterations all array elements gets arranged in a sorted manner.



| iteration-1 | iteration-2 | iteration-3 | iteration-4 | iteration-5 |
|---|---|---|---|---|
| 30 20 60 50 10 40 0 1 2 3 4 5 sel_pos pos | 0 1 2 3 4 5 sel_pos pos | 0 1 2 3 4 5 sel_pos pos | 0 1 2 3 4 5 sel_pos pos | 0 1 2 3 4 5 sel_pos pos |
| 0 1 2 3 4 5 sel_pos pos | 0 1 2 3 4 5 sel_pos pos | 0 1 2 3 4 5 sel_pos pos | 0 1 2 3 4 5 sel_pos pos | 10 20 30 40 50 60 0 1 2 3 4 5 |
| 0 1 2 3 4 5 sel_pos pos | 10 30 60 50 20 40 0 1 2 3 4 5 sel_pos pos | 0 1 2 3 4 5 sel_pos pos | 10 20 30 40 60 50 0 1 2 3 4 5 | |
| 0 1 2 3 4 5 sel_pos pos | 0 1 2 3 4 5 sel_pos pos | 10 20 30 60 50 40 0 1 2 3 4 5 | | |
| 0 1 2 3 4 5 sel_pos pos | 10 20 60 50 30 40 0 1 2 3 4 5 | | | |
| 10 30 60 50 20 40 0 1 2 3 4 5 | | | | |



Best Case : $\Omega(n^2)$

Worst Case : O(n²)

Average Case : $\theta(n^2)$

2. Bubble Sort:

- In this algorithm, in every iteration elements which are at two consecutive positions gets compared, if they are already in order then no need of swapping between them, but if they are not in order i.e. if prev position element is greater than its next position element then swapping takes place, and by this logic in first iteration largest element gets setteled at last position, in second iteration second largest element gets setteled at second last position and so on, in max (n-1) no. of iterations all elements gets arranged in a sorted manner.



| iteration-1 | iteration-2 | iteration-3 | iteration-4 | iteration-5 |
|---|---|---|---|----------------------------------|
| 30 20 60 50 10 40 | 20 30 50 10 40 60 | 20 30 10 40 50 60 | 20 10 30 40 50 60 | 10 20 30 40 50 60 |
| 0 1 2 3 4 5 pos pos+1 | 0 1 2 3 4 5 pos pos+1 | 0 1 2 3 4 5 pos pos+1 | 0 1 2 3 4 5 pos pos+1 | 0 1 2 3 4 5 pos pos+1 |
| 20 30 60 50 10 40 0 1 2 3 4 5 | 20 30 50 10 40 60 0 1 2 3 4 5 | 20 30 10 40 50 60 0 1 2 3 4 5 | 0 1 2 3 4 5 | 10 20 30 40 50 60 0 1 2 3 4 5 |
| 20 30 60 50 10 40 0 1 2 3 4 5 pos pos+1 | 20 30 50 10 40 60 0 1 2 3 4 5 | 20 10 30 40 50 60 0 1 2 3 4 5 | 10 20 <u>30 40 50 60</u> 0 1 2 3 4 5 | |
| 20 30 50 60 10 40 0 1 2 3 4 5 | 20 30 10 50 40 60 0 1 2 3 4 5 pos pos+1 | 20 10 30 40 50 60 0 1 2 3 4 5 | | |
| 20 30 50 10 60 40 0 1 2 3 4 5 pos pos+1 | 20 30 10 40 50 60 0 1 2 3 4 5 | | | |
| 20 30 50 10 40 60 0 1 2 3 4 5 | | | | |



Best Case : $\Omega(n)$ - if array elements are already arranged in a sorted manner.

Worst Case : O(n²)

Average Case : $\theta(n^2)$

3. Insertion Sort:

- In this algorithm, in every iteration one element gets selected as a **key element** and key element gets inserted into an array at its appropriate position towards its left hand side elements in a such a way that elements which are at left side are arranged in a sorted manner, and so on, in max **(n-1)** no. of iterations all array elements gets arranged in a sorted manner.
- This algorithm works efficiently for already sorted input sequence by design and hence running time of an algorithm is O(n) and it is considered as a best case.



Best Case : $\Omega(n)$ - if array elements are already arranged in a sorted manner.

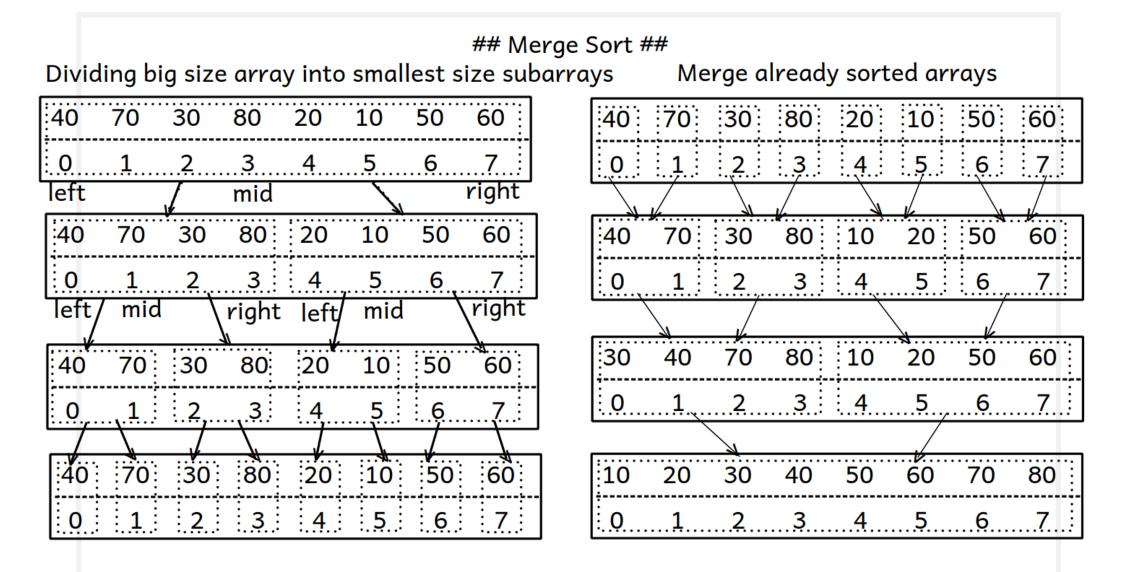
Worst Case : O(n²) Average Case: Θ(n²)

- Insertion sort algortihm is an efficient algorithm for smaller input size array.

4. Merge Sort:

- This algorithm follows divide-and-conquer approach.
- In this algorithm, big size array is divided logically into smallest size (i.e. having size 1) subarrays, as if size of subarray is 1 it is sorted, after dividing array into sorted smallest size subarray's, subarrays gets merged into one array step by step in a sorted manner and finally all array elements gets arranged in a sorted manner.
- This algorithm works fine for **even** as well **odd** input size array.
- This algorithm takes extra space to sort array elements, and hence its space complexity is more.







Best Case : $\Omega(n \log n)$

Worst Case : O(n log n)

Average Case : $\theta(n \log n)$

5. Quick Sort:

- This algorithm follows **divide-and-conquer** approach.
- In this algorithm the basic logic is a partitioning.
- **Partitioning:** in parititioning, pivot element gets selected first (it may be either leftmost or rightmost or middle most element in an array), after selection of pivot element all the elements which are smaller than pivot gets arranged towards as its left as possible and elements which are greater than pivot gets arranged as its right as possible, and big size array is divided into two subarray's, so after first pass pivot element gets settled at its appropriate position, elements which are at left of pivot is referred as **left partition** and elements which are at its right referred as a **right partition**.



Best Case : $\Omega(n \log n)$

Worst Case: O(n²) - worst case rarely occures

Average Case : $\theta(n \log n)$

- Quick sort algortihm is an efficient sorting algorithm for larger input size array.



- Limitations of an array data structure:
- **1. Array is static**, i.e. size of an array is fixed, its size cannot be either grow or shrink during runtime.
- 2. Addition and deletion operations on an array are not efficient as it takes O(n) time, and hence to overcome these two limitations of an Array Linked List data structure has been designed.

Linked List: It is a collection/list of logically related similar type of elements in which,

- an address of first element in a collection/list is stored into a pointer variable referred as a head pointer.
- each element contains data and an address of its next (as well as its previous element).
- An element in a Linked List is also called as a **Node.**
- Four types of linked lists are there: Singly Linear Linked List, Singly Circular Linked List, Doubly Linear Linked List and Doubly Circular Linked List.



- Basically we can perform **addition**, **deletion**, **traversal** etc... operations on linked list data structure.
- We can add and delete node by three ways: we can add node into the linked list at last position, at first position and at any specific position, simillarly we can delete node from linked list which is at first position, last position and at any specific position.
- 1. Singly Linear Linked List: It is a linked list in which
- head always contains an address of first element, if list is not empty.
- each node has two parts:
- i. data part: contains data of any primitive/non-primitive type.
- ii. pointer part(next): contains an address of its next element/node.
- last node points to NULL, i.e. next part of last node contains NULL.

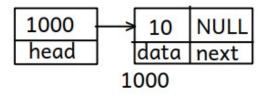


SINGLY LINEAR LINKED LIST

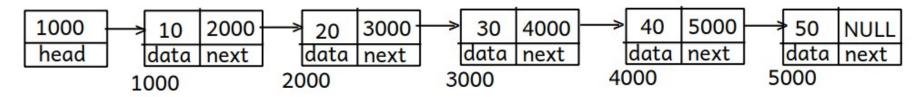
1) singly linear linked list --> list is empty



2) singly linear linked list --> list contains only one node



3) singly linear linked list --> list contains more than one nodes





Limitations of Singly Linear Linked List:

- Add node at last position & delete node at last position operations are not efficient as it takes O(n) time.
- We can starts traversal only from first node and can traverse the SLLL only in a forward direction.
- Previous node of any node cannot be accessed from it.
- Any node cannot be revisited to overcome this limitation Singly Circular Linked List has been designed.

2. Singly Circular Linked List: It is a linked list in which

- head always contains an address of first element, if list is not empty.
- each node has two parts:
- i. data part: contains data of any primitive/non-primitive type.
- ii. pointer part(next): contains an address of its next element/node.
- last node point to first node, i.e. next part of last node contains an address of first node.

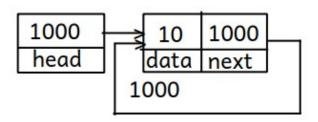


SINGLY CIRCULAR LINKED LIST

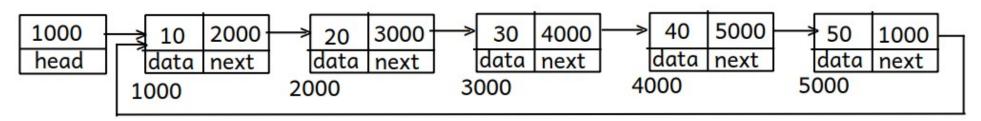
1) singly circular linked list --> list is empty



2) singly circular linked list --> list contains only one node



3) singly circular linked list --> list contains more than one nodes





Limitations of Singly Circular Linked List:

- Add last, delete last & add first, delete first operations are not efficient as it takes O(n) time.
- We can starts traversal only from first node and can traverse the SCLL only in a forward direction.
- Previous node of any node cannot be accessed from it to overcome this limitation Doubly Linear Linked List has been designed.
- 3. Doubly Linear Linked List: It is a linked list in which
- head always contains an address of first element, if list is not empty.
- each node has three parts:
- i. data part: contains data of any primitive/non-primitive type.
- ii. pointer part(next): contains an address of its next element/node.
- iii .pointer part(prev): contains an address of its previous element/node.
- next part of last node & prev part of first node point to NULL.

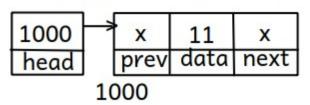


DOUBLY LINEAR LINKED LIST

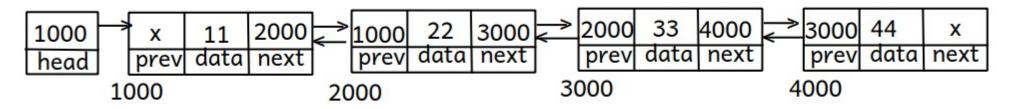
1. doubly linear linked list --> list is empty



2. doubly linear linked list --> list is contains only one node



3. doubly linear linked list --> list is contains more than one nodes



Limitations of Doubly Linear Linked List:

- Add last and delete last operations are not efficient as it takes O(n) time.
- We can starts traversal only from first node, and hence to overcome these limitations Doubly Circular Linked List has been designed.

4. Doubly Circular Linked List: It is a linked list in which

- head always contains an address of first node, if list is not empty.
- each node has three parts:
- i. data part: contains data of any primitive/non-primitive type.
- ii. pointer part(next): contains an address of its next element/node.
- iii .pointer part(prev): contains an address of its previous element/node.
- next part of last node contains an address of first node & prev part of first node contains an address of last node.

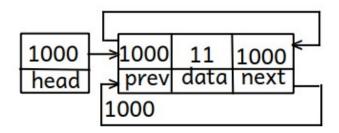


DOUBLY CIRCULAR LINKED LIST

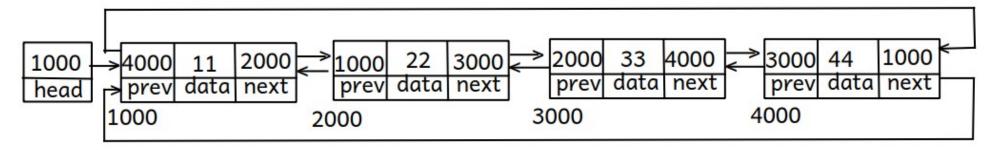
1. doubly circular linked list --> list is empty



2. doubly circular linked list -> list is contains only one node



3. doubly circular linked list --> list is contains more than one nodes





Advantages of Doubly Circular Linked List:

- DCLL can be traverse in forward as well as in a backward direction.
- Add last, add first, delete last & delete first operations are efficient as it takes O(1) time and are convenient as well.
- Traversal can be start either from first node or from last node.
- Any node can be revisited.
- Previous node of any node can be accessed from it

Array v/s Linked List:

- Array is **static** data structure whereas linked list is dynamic data structure.
- Array elements can be accessed by using **random access** method which is efficient than linked list elements which can be accessed by **sequential access** method.
- Addition & Deletion operations are efficient on linked list than on an array.
- Array elements gets stored into the **stack section**, whereas linked list elements gets stored into **heap section**.
- In a linked list extra space is required to maintain link between elements, whereas in an array to maintain link between elements is the job of **compiler**.

