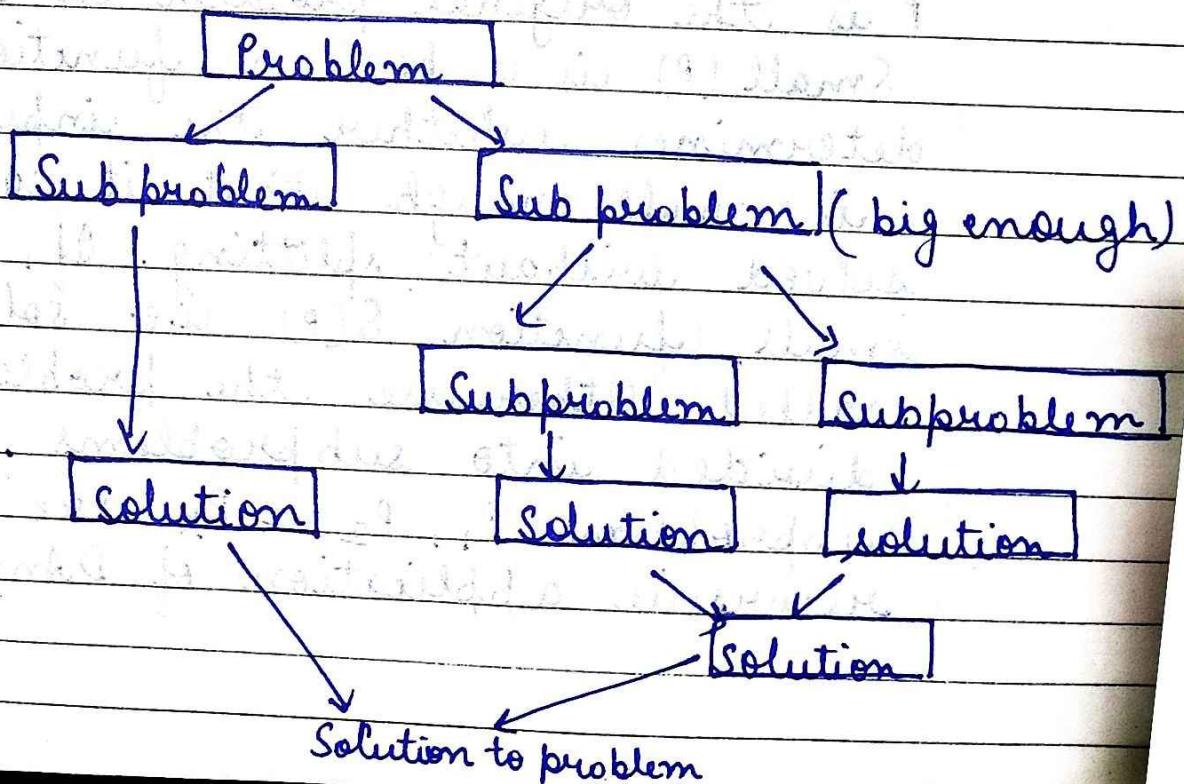


Divide & conquer:

General method:

Given a function to compute n -inputs the divide & conquer strategy says splitting the inputs into K distinct subsets, $1 \leq K \leq n$, results into K sub-problems. Now, these subproblems are solved separately & then a method must be found to combine subproblems into a solution. If the subproblems are large, then the divide and conquer can be reapplied. The subproblems resulted from big problems are of same type as the original problem. The small & small subproblems are generated until the small subproblems are small enough to be solved without splitted are produced.



Divide & conquer principle is generally implemented using recursive algorithm.

Algorithm DAndC(P)

```
if small( $P$ )
    return  $S(P)$ 
```

```
else
    {
        divide  $P$  into subproblems  $P_1, P_2, \dots, P_k$ ;
        Apply DAndC to each subproblem;
        return (combine (DAndC( $P_1$ ), DAndC( $P_2$ ),
                          ... DAndC( $P_k$ )));
    }
```

Time complexity for DAndC algorithm :-
 subproblem is solved separately & if needed they are again divided into more subproblems. After that, combine solution to P using the solutions to k sub problems. If the size of problem P is n , size of each sub problem is $n_1, n_2, n_3 \dots n_k$. DAndC is called for each sub problem & then solution is found & then combined into one solution.

$$T(n) = \begin{cases} g(n) \\ T(n_1) + T(n_2) + T(n_3) \dots T(n_k) + f(n) \end{cases}$$

where $T(n) \rightarrow$ Time for DAndC algorithm for input size ' n '.

$g(n) \rightarrow$ Time to compute answer directly from subproblems

$T(n_1), T(n_2) \dots T(n_k) \rightarrow$ Time for sub problems.

$f(n) \rightarrow$ It is the time for dividing P & combining the solutions to subproblems.

→ If the size of two subproblems are same, then time complexity,

divide P into subproblems P_1, P_2, \dots, P_k are solved by recursive application of DAndC. Each

$$T(n) = \begin{cases} g(n) \\ 2T(n/2) + f(n) \end{cases}$$

Now, if problem is divided into 'a' subproblems & size of each subproblem after dividing 'n' is n/b

$$T(n) = \begin{cases} T(1) & \text{if } n=1 \\ aT(n/b) + f(n) & n>1. \end{cases}$$

where a & b are constants

$$E.g. \quad a=2, b=2, f(n)=n, T(1)=2$$

where $n \rightarrow$ number of elements
 $x \rightarrow$ element to be searched.

$$\begin{aligned} T(n) &= 2(T(n/2)) + n \\ &= 2(2T(n/4) + n/2) + n \\ &= 2^2(2T(n/8) + n/4) + 2n \\ &= 2^3T(n/16) + 2n/2 \\ &= 2^3T(n/16) + 8n \end{aligned}$$

$$T(n) = 2^i T(n/2^i) + 3n.$$

So, $T(n) = 2^i T(n/2^i) + in$.
 where $\log_2 n \geq i \geq 1$.

$$T(n) = 2^{\log_2 n} T(n/2^{\log_2 n}) + n \log_2 n.$$

If $i = \log_2 n$

$$T(n) = n \cdot T(1) + n \log_2 n$$

$$= n \cdot 2 + n \log_2 n$$

$$= 2n + \log_2 n \cdot n$$

Binary search :

Divide & conquer can be used to solve this problem. Let $\text{Small}(P)$ be true if $n=1$. $S(P)$ will take value x if $x=a_i$, otherwise it will take value 0.

Recursive binary search :

```
Algorithm BoxSearch (a, u, l, x)
{
```

```
    if (u==l) then {
        if (a[u]==x)
            return u;
    }
```

```
    else
```

```
        return 0;
```

```
}
```

```
else {
```

```
    mid = [u+l]/2;
```

```
    if (x == a[mid])
```

```
        return i;
```

else if ($a[\text{mid}] > x$)
return $\text{BinSearch}(a, i, \text{mid}-1, x)$;

else
return $\text{BinSearch}(a, \text{mid}+1, l, x)$;

}

}

Iterative Binary Search :

Algorithm BinSearch :

low = 0;

high = $n-1$;

do {

mid = $(\text{low} + \text{high})/2$;

if ($x < a[\text{mid}]$) then

end = mid - 1;

else if ($x > a[\text{mid}]$) then

start = mid + 1;

else

return mid;

} while ($\text{low} \leq \text{high}$)

return 0; // if element is not found,

}

it will return 0.

$l \cdot g \div$

0 1 2 3 4 5 6 7 8

-15 -6 0 7 9 10 12 16 18

let $x = 12$

low = 0, high = 8
mid = $0 + 8 = 8 = 4$

2 8

Now, $x > a[\text{mid}]$
 $12 > a[4]$

$12 > 9$

then, high = high
start = mid + 1 = $4 + 1 = 5$

5 6 7 8
10 12 16 18

high > low, 8 > 5

start = 5, end ≠ high = 8.

$$\text{mid} = \frac{5+8}{2} = \frac{13}{2} = 6.5 \Leftarrow = 7$$

Now, $x < a[\text{mid}]$
 $12 < 16$

high = mid - 1 = $7 - 1 = 6$.

6 > 5

$$\text{mid} = \frac{6+5}{2} = \frac{11}{2} = 5.5 = 6.$$

Now, $x == a[\text{mid}]$,

$12 == a[6]$,

$12 == 12$.

return mid.

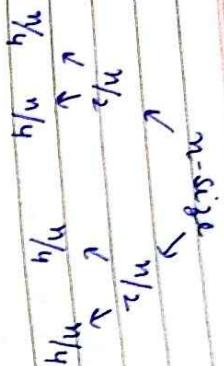
so, element found at index 6.

Time complexity :

Best-case : $\Omega(1)$

Worst-case : $O(\log n)$.

if ($a[i] < \min$) then
 $\min = a[i]$;



So, every time array is divided into $\frac{n}{2^k}$

$$\frac{n}{2^k} = n$$

$$k \log_2 = \log_2 n$$

$$K = \log_2 n$$

Average-case: $\Theta(\log n)$

Find the maximum & minimum :

The problem is to find the minimum & maximum items in a set of n elements.

StraightMaxMin requires $\Theta(n-1)$ elements comparisons in the best, average & worst case.

Algorithm StraightMaxMin(a, n, max, min)

{

 max = min = a[1];

 for i = 2 to n do

 if (a[i] > max) then

 max = a[i];

But, there is a problem, $a[i] < \min$ is only required when $a[i] > \max$ is false. So, we can replace if with elseif.

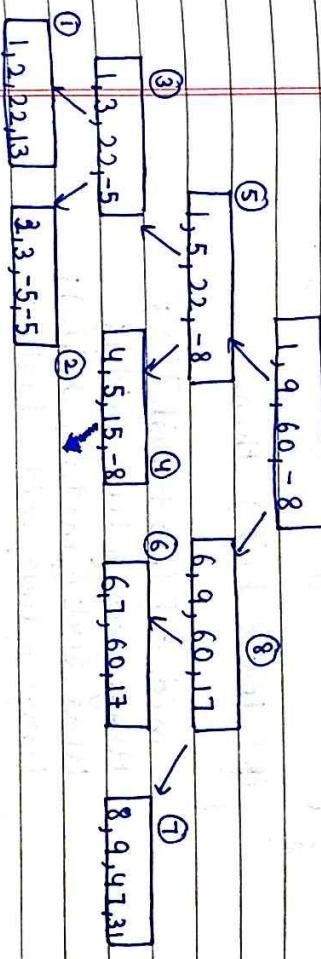
```
if ( a[i] > max) then
    max = a[i]
else if ( a[i] < min) then
    min = a[i].
```

This problem can be implemented using recursion i.e. divide & conquer rule. Let $P = (n, a[i] \dots a[j])$ where n is number of elements in list $a[i] \dots a[j]$. Let Small(P) be true when $n \leq 2$. If $n=1$, $a[i]$ is the maximum & minimum. If $n=2$, it means, problem can be solved by making one comparison.

If list has more than 2 elements, P has to be divided into smaller instances. For example, when P is divided into two parts like $P_1 = (n/2, a[1] \dots a[n/2])$ &

$$P_2 = (n - n/2), a[n/2 + 1], \dots, a[n]).$$

We calculate $\text{MAX}(P_1) \& \text{MAX}(P_2)$ & similarly $\text{MIN}(P_1), \text{MIN}(P_2)$. Now maximum of P_1 is compared with maximum of P_2 & the largest element is largest element from P .



Algorithm: $\text{MAXMIN}(i, j, \text{max}, \text{min})$

```

{
    if (i == j) then
        max = min = a[i];
    else if (i = j - 1) then
        {
            if a[i] < a[j] then
                max = a[i];
            else
                min = a[i];
        }
    else
        {
            max = a[i];
            min = a[i];
            for (j = i + 1; j <= j - 1; j++)
                if a[i] < a[j] then
                    max = a[i];
                else
                    min = a[i];
            }
}

```

Merge sort :

$$T(n) = \begin{cases} T(n/2) + T(n/2) + 2n & n > 2 \\ 0 & n = 1 \end{cases}$$

The Merge Sort algorithm is a sorting algorithm that is based on divide & conquer paradigm. In this algorithm, array is divided into two halves & they are combined into sorted manner.

This algorithm continuously splits the array in half until it can't be further divided. This means that if the array is empty or has one element only left, the dividing will stop. If the elements are divided more than one, recursively merge sort is applied on both of halves. Finally, when arrays are sorted, the merge operation is applied. Merge operation is the process of taking two smaller sorted arrays & combine them to large one.

Algorithm MergeSort (a, low, high)

```
{
    if (low < high) then
    {
        mid = [low + high]/2;
        MergeSort (a, low, mid);
        MergeSort (a, mid+1, high);
        Merge (a, low, mid, high);
    }
}
```

```

y
{
    Merge (a, low, mid, high);
    MergeSort (a, mid+1, high);
    Merge (a, low, mid, high);
}

```

```

n1 = mid - low + 1;
n2 = high - mid;

```

11. Declare two arrays arr1 & arr2

of size n1 and n2.

```
for i=0 to n1 do
    arr1[i] = a[low+i];
```

```
for j=0 to n2 do
    arr2[j] = a[1+mid+j];
```

```
i=j=0;
while i < n1 and j < n2 do
```

```
    if arr1[i] < arr2[j] then
        a[k] = arr1[i], i++;

```

```
    else
        a[k] = arr2[j], j++;
```

```
j++;
k++;

```

```
while i < n1 do
```

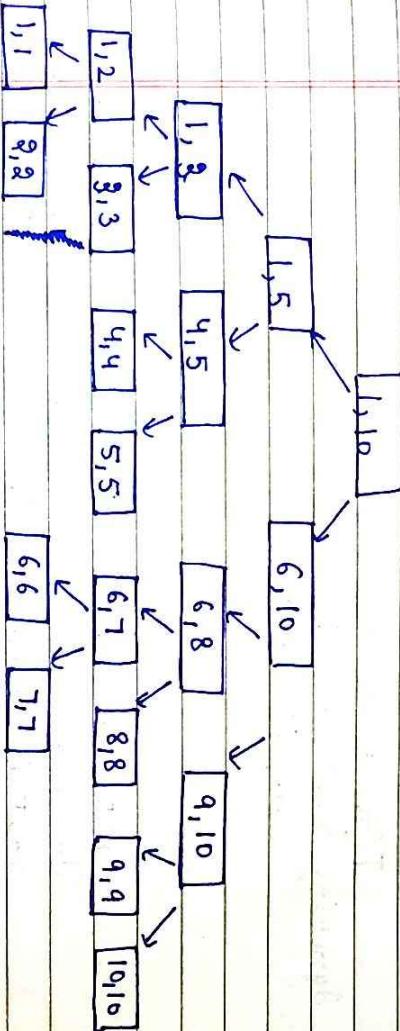
```
    a[k] = arr1[i];
    i++;

```

```
j++;
```

while j < n2 do

a[k] = arr2[j];
 k++;



Time complexity $\frac{n}{2}$

$T(n) = \begin{cases} a & n=1, a \text{ is constant} \\ 2T(n/2) + cn & n>1, c \text{ is constant} \end{cases}$

When n is power of 2, $n=2^k$.

$$T(n) = 2T(n/2) + cn$$

$$= 2[2T(n/4) + cn/2] + cn$$

$$= 4T(n/4) + 2cn$$

$$= 4[4T(n/8) + cn/4] + 2cn$$

$$= 8T(n/8) + 4cn + 2cn$$

$$\begin{aligned}
 & \rightarrow 2^K T(n/n) + Kcn \\
 & \Rightarrow 2^K T(1) + Kcn. \\
 & [2^K = n, T(1) = a] \\
 & = an + cn \log n
 \end{aligned}$$

!

$$\begin{aligned}
 T(n) &= an + cn \log n. \\
 \text{Ignoring constants, } T(n) &= n + n \log n. \\
 \text{So, } T(n) &= O(n \log n)
 \end{aligned}$$

Strassen's matrix multiplication

Let A and B be true $n \times n$ matrices.

The product matrix $C = AB$, is also $n \times n$ matrix whose i, j th element is formed by taking elements in the i th row & the j th column of B & multiplying them to get,

$$C(i,j) = \sum_{k=1}^{k=n} A(i,k) \cdot B(k,j)$$

$$\begin{bmatrix} A \\ I \\ \vdots \\ I \\ B \end{bmatrix} \times \begin{bmatrix} B \\ I \\ \vdots \\ I \\ C \end{bmatrix} = \begin{bmatrix} C \\ I \\ \vdots \\ I \\ I \end{bmatrix}$$

No. of rows of B = No. of columns of A.

Algorithm for multiplication:

for $i = 0$ to n do

 for $j = 0$ to n do

$c[i,j] = 0;$

```

for k = 0 to n do
    c[i,j] += A[i,k] * B[k,j];
end of for loop.
end of for loop.
end for loop.

```

To compute C_{ij} we need ' n ' multiplications. As the matrix C has n^2 elements, the time complexity for resulting algorithm is $O(n^3)$.

The divide & conquer strategy suggests another way to compute the product of true $n \times n$ matrices. We assume that n is a power of 2, i.e. $n = 2^k$ where k is non-negative integer. In case, if n is not a power of 2, then enough rows & columns of zeros can be added to both A & B so that resulting dimensions are of power of two.

Suppose $n = 4$, then A and B are each partitioned into four square submatrices, each submatrix having dimensions $n/2 \times n/2$. AB can be computed by using formula $\sum A(i,k) B(k,j)$ for product of 2×2 matrices.

$$\begin{bmatrix} A_{11} & A_{12} \\ A_{21} & A_{22} \end{bmatrix} \begin{bmatrix} B_{11} & B_{12} \\ B_{21} & B_{22} \end{bmatrix} = \begin{bmatrix} C_{11} & C_{12} \\ C_{21} & C_{22} \end{bmatrix}$$

$$\begin{aligned}
 C_{11} &= A_{11}B_{11} + A_{12}B_{21} \\
 C_{12} &= A_{11}B_{12} + A_{12}B_{22} \\
 C_{21} &= A_{21}B_{11} + A_{22}B_{21}
 \end{aligned}$$

$$C_{22} = A_{21}B_{12} + A_{22}B_{22}$$

If $n=2$, the above 4 formulas are computed using multiplication operation for elements of A and B . If $n > 2$, elements of C can be calculated using matrix multiplication & addition operations applied to matrix size of $n/2 \times n/2$.

Since n is power of 2, the matrix products can be successively computed by the same algorithm. This algo. will continuously applying to itself until n becomes relatively small.

$$A = \begin{bmatrix} a_{11} & a_{12} & a_{13} & a_{14} \\ a_{21} & a_{22} & a_{23} & a_{24} \\ a_{31} & a_{32} & a_{33} & a_{34} \\ a_{41} & a_{42} & a_{43} & a_{44} \end{bmatrix} \quad X = \begin{bmatrix} b_{11} & b_{12} & b_{13} & b_{14} \\ b_{21} & b_{22} & b_{23} & b_{24} \\ b_{31} & b_{32} & b_{33} & b_{34} \\ b_{41} & b_{42} & b_{43} & b_{44} \end{bmatrix}$$

$$T(n) = \begin{cases} b & n \leq 2 \\ 8 T(n/2) + cn^2 & n > 2 \end{cases}$$

$$8 \text{ Matrix multiplication call for } n/2$$

Addition of $n=2$, requires 4 additions - one i.e. n^2 .

Algorithm SMM (A, B, n)

```

if (n ≤ 2)
    C11 = A11 B11 + A12 B21
    C12 = A11 B12 + A12 B22
    C21 = A21 B11 + A22 B21
    C22 = A21 B12 + A22 B22
}
else
    mid = n/2.
    SMM(A11, B11, n/2) + SMM(A12, B21, n/2)
    SMM(A11, B12, n/2) + SMM(A12, B22, n/2)
    SMM(A21, B11, n/2) + SMM(A22, B21, n/2)
    SMM(A21, B12, n/2) + SMM(A22, B22, n/2)
}
}
```

Recurrence relation for above algorithm,

$$A_{11} = \begin{bmatrix} a_{11} & a_{12} \\ a_{21} & a_{22} \end{bmatrix} \quad A_{12} = \begin{bmatrix} a_{13} & a_{14} \\ a_{23} & a_{24} \end{bmatrix}$$

$$A_{21} = \begin{bmatrix} a_{31} & a_{32} \\ a_{41} & a_{42} \end{bmatrix} \quad A_{22} = \begin{bmatrix} a_{33} & a_{34} \\ a_{43} & a_{44} \end{bmatrix}$$

Strassen's has discovered a way to compute $C(i,j)$ using 7 multiplications & 18 additions or subtractions.

- 1.) Firstly, compute the seven $n/2 \times n/2$ matrices, P, Q, R, S, T, U and V .
- 2.) Then compute $C(i,j)$.

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P, Q, R, S, T, U and V is calculated using 7 matrix multiplications and 10 matrix additions or subtractions.

Then, C_{ij} requires additional 8 additions or subtractions.

$$P = (A_{11} + A_{22})(B_{11} + B_{22})$$

$$Q = (A_{21} + A_{22})B_{11}$$

$$R = A_{11}(B_{12} - B_{22})$$

$$S = A_{22}(B_{21} - B_{11})$$

$$T = \cancel{B_{11}}(A_{11} + A_{12})B_{22}$$

$$U = (A_{21} - A_{11})(B_{11} + B_{21})$$

$$V = (A_{12} - A_{22})(B_{12} + B_{22})$$

$$C_{11} = P + S - T + V$$

$$C_{12} = R + T$$

$$C_{21} = Q + S$$

$$C_{22} = P + R - Q + U$$

So, resulting recurrence relation,

$$T(n) = \begin{cases} b & n \leq 2 \\ 7T(n/2) + an^2 & n > 2 \end{cases}$$

where a & b are constants.

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$$\text{So, } O(n^{\log_2 7}) = O(n^{2.81}) \approx O(n^2).$$

Quicksort :

Quicksort is a sorting algorithm which is based on divide & conquer paradigm. In merge sort $a[1:n]$ was divided at its mid point into subarrays which were independently sorted & later merged. In quicksort, the division into two subarrays is made so that the sorted subarrays need not be merged.

The elements in $a[1:n]$ is rearranged such that $a[i] \leq a[j]$ for all i between l & m and for all j between $m+1$ & n .

So $a[l:m]$ & $a[m+1:n]$ is sorted independently, thus need not be merged.

The rearrangement is done in such a manner that, we ~~had~~ picked one element of $a[]$, say pivot, $\text{pivot} = a[s]$, and then reordering the other elements so that all elements before pivot in $a[1:n]$ are less than or equal to pivot & all elements appearing after pivot are greater than equal to pivot. This rearranging is referred to as partitioning.

Algorithm QuickSort (a, p, q)

{

if ($p < q$) then

loc = partition (a, p, q);

quicksort ($a, p, loc-1$);

quicksort ($a, loc+1, q$);

// There is no need to combine solution

}

Algorithm partition (a, p, q)

```

pivot = a[p];
low = p;
high = q;
repeat
{
    repeat
    {
        low++;
    } until (a[low] < a[pivot])
}

```

repeat
high--;

```

until (a[high] > a[pivot])
if (low < high)
    interchange (a, low, high);
} until (low <= high);

```

```

Interchange (a, high, pivot);
return high;
}
```

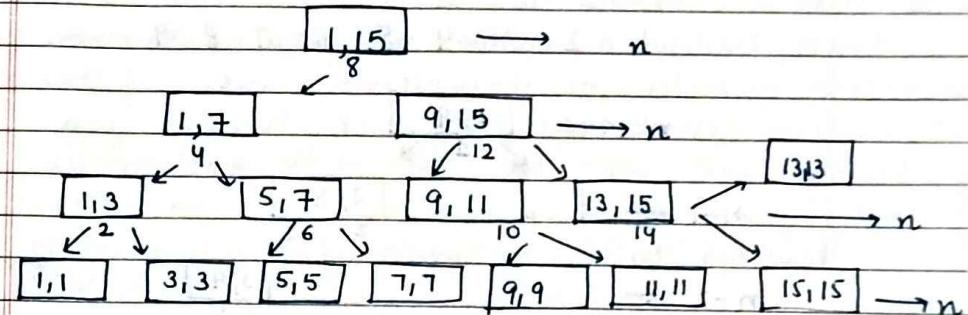
Algorithm interchange (a, i, j) {

```

p = a[i];
a[i] = a[j];
a[j] = p;
}
```

Time complexity :-

1. Best case :- The partition is always done at the middle.



Algorithm quicksort (a, p, q)

```

if (p < q) then
    n ← loc = partition (a, p, q);
    quicksort (a, p, loc-1);
    quicksort (a, loc+1, q);
}

```

The partition function does 'n' comparison to divide the array.

The height of tree = $\frac{n}{2}$ [The 'n' is splitted to half everytime]

$$\frac{n}{2^k}$$

$$n = 2^k$$

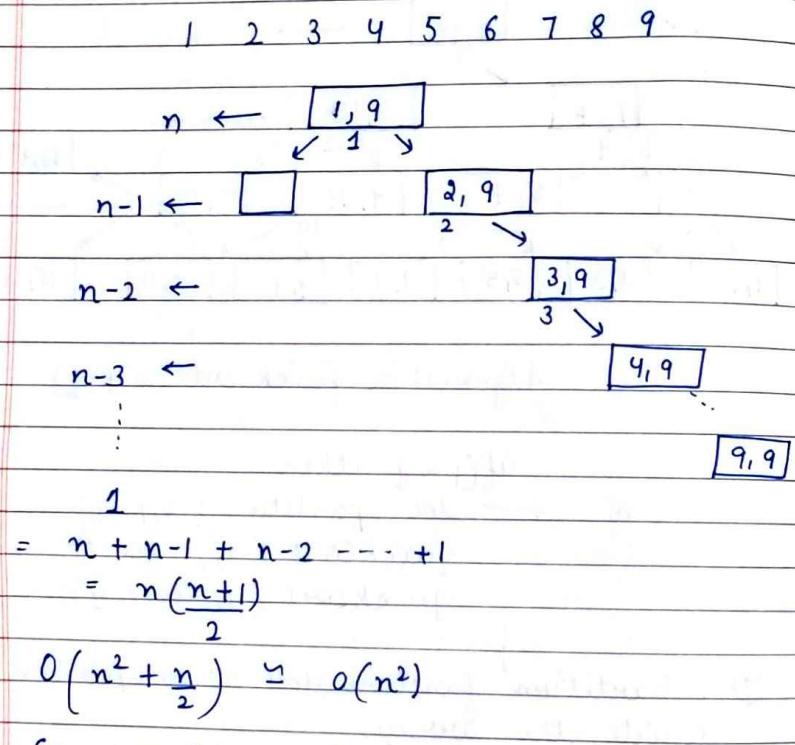
$$\log_2 n = k$$

until 1 element left

So, the height of tree = $\log n$
Time complexity = $n \log n$

So, best-case complexity is $\Theta(n \log n)$

2. Worst-Case: It happens when array is already sorted.



So, worst-case time complexity is $O(n^2)$.

- (i) If the pivot element is the middle element of array, the worst-case can be converted into best-case.
- (ii) Select ~~middle~~ random element as pivot.

Space complexity = $\log n$ to n .

greedy method:

The general method:

In greedy method, a problem has 'n' inputs & problem is stored in such a manner that we obtain a subset that satisfies some constraints or selection criteria. Any subset that satisfies these constraints is called a feasible solution. The feasible solution that maximizes or minimizes the given objective function is called optimal solution.

The greedy method suggests that, algorithm can divide the algorithm into stages, considering one input at a time. At each stage, a decision is made that a particular input will give a optimal solution or not. This is done by considering the inputs in the order determined by selection procedure. If the inclusion of next input into partially constructed optimal solution will result in an infeasible solution, then the input is not added to partial solution. Otherwise, it is added. The selection procedure is also based on some optimization measure. Different optimization measures are possible for a given solution/problem.

Greedy algorithm satisfies two paradigm i.e. subset & ordering paradigm.

Subset paradigm satisfies that only those inputs should be included in the

set that can satisfy the objective function or lead to feasible solution.

Ordering paradigm is used to select the inputs in the set in such a order that can meet the objective function. Each decision is made using optimisation criteria that can be computed using decision already made.

Algorithm Greedy (a, n)

// a[0:n] contain n inputs

{

solution = \emptyset ; // Initialize solution
for i = 0 to n do

{

x = select(a);

if feasible (solution, x) then

solution = union (solution, x);

y

return solution;

}

The function Select() selects an input from a[] & removes it. The selected input value is assigned to x. Feasible is a Boolean-valued ~~to~~ function that determines whether x can be included into the solution vector. If it returns True, then the function Union combines x with the solution & update the solution vector. The function Greedy

describes the essential way that a greedy algorithm will look.

Knapsack Problem:

In Knapsack problem, we are given 'n' objects & a knapsack or a bag having capacity 'm'. Object 'i' has a weight w_i & profit p_i . The objective is to obtain a filling the knapsack that maximizes the total profit earned. Since, the knapsack capacity is m, we require that total weight of all chosen objects to be at most m.

If weight of all given items is less than or equal to m, all are chosen. If the total weight exceeds the capacity, we can't pack them all. In that case, we choose subset of items/object that can fit in knapsack with maximum profit.

So, Knapsack problem can be stated as,

$$\text{maximize } \sum_{1 \leq i \leq n} p_i$$

$$\text{subject to } \sum_{1 \leq i \leq n} w_i \leq m.$$

The profits & the weights are positive numbers.

There are three greedy strategies to obtain feasible solutions,

- (i) First, trying filling the knapsack by including object with maximum profit.

ii) Secondly, considering the objects in order of non-decreasing weights w_i .

iii) last, is to achieve a balance between the rate at which profit increases & the rate at which capacity is used.

So, we include object which has maximum profit per unit of capacity used.

So, objects are considered in order of p_i/w_i .

Each of above strategy requires $O(n)$ time.

e.g. $i = 7$

$$\{p_1, \dots, p_7\} = \{10, 5, 15, 7, 6, 18, 3\}$$

$$\{w_1, \dots, w_7\} = \{2, 3, 5, 7, 1, 4, 1\}$$

Knapsack capacity (m) = 15.

Object	1	2	3	4	5	6	7
P_i	10	5	15	7	6	18	3
w_i	2	3	5	7	1	4	1
P_i/w_i	5	1.67	3	1	6	4.5	3

1. Selecting objects on basis of maximum profit :

Obj	P_i	w_i	Remaining Weight
6	18	4	$15-4=11$

3	15	5	$11-5=6$
1	10	2	$6-2=4$
4	4	$4 \times 1=4$	$4-4=0$

$$\Sigma = 47$$

2. Selecting object on basis of increasing weight or with minimum weight :

Obj	P_i	w_i	Remaining weight
5	6	1	$15-1=14$
7	3	1	$14-1=13$
1	10	2	$13-2=11$
2	5	3	$11-3=8$
6	18	4	$8-4=4$
3	12	4	$4-4=0$

$$\Sigma = 54$$

3. Selecting object on basis of maximum P_i/w_i :

Obj	P_i	w_i	Remaining Wt.
5	6	1	$15-1=14$
1	10	2	$14-2=12$
6	18	4	$12-4=8$
3	15	5	$8-5=3$
7	3	1	$3-1=2$
2	2×1.67	2	$2-2=0$

$$\Sigma = 55.34$$

So, the maximum profit is taken when objects are included on basis of maximum P_i/w_i .

Algorithm MinWeight (wt, pf, n)

```

{
    // W is capacity of knapsack
    // wt array of decreasing order.
    // pf corresponding profits
    // n number of values in
    // array.

    // n is initialized to size of array.

    if (n == 0 || W == 0) then
        return 0;
    else
        return max ((pf[n-1] + knapsack(W-wt[n-1],
                                         wt, pf, n-1)), knapsack(W,
                                         wt, pf, n-1))
}

```

Algorithm MaxRatio (W, wt, pf, n)

```

{
    for i=1 to n do
        x[i] = 0
    weight = 0; profit = 0;
    for i=n to 1 do {
        if weight + w[i] ≤ W then
            x[i] = pf[i] / w[i]
            weight = weight + w[i]
        else
            profit = profit + pf[i]
        x[i] = pf[i] / w[i]
        profit = profit + (W-weight)*x[i]
        break
    }
    return profit;
}

```

In above algorithm, arrays are sorted in increasing order of P_i/W_i ratio.

Algorithm MaxProfit (W, wt, pf, n)

```

{
    weight = 0
    profit = 0.
    for i=n to 1 do
        if weight + w[i] ≤ W then
            weight = weight + w[i]
            profit = profit + pf[i]
        else
            x = Pf[i] / w[i]
            profit = profit + (W-weight)*x.
            break.
    return profit
}

```

Job sequencing with deadlines:

In this problem, there are set of n jobs. Associated with job ' i ' is an integer deadline $d_i \geq 0$ and a profit $P_i > 0$. For any job ' i ', the profit P_i is earned if the job is executed by its deadline. To complete a job, one has to process the job on a machine for one unit of time. Only one machine is available for processing jobs. A subset J of jobs is a feasible solution such that each job is completed by its deadline. The value of a feasible solution

J is sum of profits of jobs in T . An optimal solution is a feasible solution with maximum value.

Since, the problem involves the identification of a subset, it fits the subset paradigm.

Algorithm JS (d, p, n)

{
 $d_{\max} = \max(d)$;

 for $i=1$ to n do

 set $K = \min(d_{\max}, \text{deadline}(i))$

 while $K \geq 1$ do

 if $\text{timeslot}[K] == \text{empty}$ then

$\text{timeslot}[K] = \text{job}[i]$

 break

 end if

 Set $K = K - 1$

 end while.

 for $i=1$ to d_{\max} then

$\text{profit} = \text{profit} + p[i]$.

 return timeslot

}

E.g. →

Job	Deadline	Profit
A ₁	2	60
A ₂	1	100
A ₃	3	20
A ₄	2	40
A ₅	1	20

Sorting the array in decreasing order of max. profit,

Job	A ₂	A ₁	A ₄	A ₅	A ₃
Deadline	1	2	2	1	3
Profit	100	60	40	20	20

(i)

$d_{\max} = 3$.

$K = \min(3, 1)$

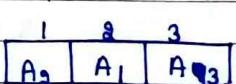
$K = 1$

while $1 \geq 1$ do.

$\text{timeslot}[1] = \text{empty, true}$

$\text{timeslot}[1] = \text{job}[1]$

$K = 0$



(ii)

$K = \min(2, 3)$

$K = 2$

while $2 \geq 1$ do

$\text{timeslot}[2] = \text{empty, true}$

$\text{timeslot}[2] = \text{job}[2]$

$K = 1$

while $1 \geq 1$ do

$\text{timeslot}[1] = \text{empty, false}$

$K = 0$.

(iii)

$K = \min(3, 3)$

$K = 3$

while $3 \geq 1$ do

$\text{timeslot}[3] = \text{empty, then}$

$\text{timeslot}[3] = \text{job}[5]$

$K = 2$, $\text{timeslot}[2] = \text{empty, false}$

$K = 1$, $\text{timeslot}[1] = \text{empty, false}$.

Q.

Job	Deadline	Profit
A ₁	2	100
A ₄	1	87
A ₃	2	15
A ₂	1	10

(i)

$$i=1, d_{\max} = 2$$

1	2
A ₄	A ₁

$$K = (2, 2)$$

$$K = 2$$

while $2 \geq 1$ then

if timeslot[2] = empty then

timeslot[2] = job[1]

~~K=1~~ break.

(ii)

$$i=2$$

$$K = \min(1, 2)$$

$$K = 1$$

while $1 \geq 1$ do

if timeslot[1] = empty, true

timeslot[1] = job[2]

break.

$$K = 0$$

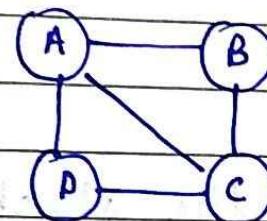
Spanning Tree:

A spanning tree of a graph is just a subgraph that contains all the vertices & is tree. That is, a spanning tree of a connected graph or contains all the vertices & has the edges which connect all the vertices.

So, the number of edges is 1 less than the number of vertices.



$G(V, E)$

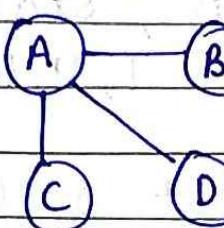


$G'(V', E')$

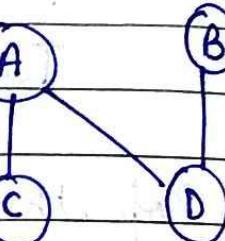
$V' = V$

$E' \subset E$

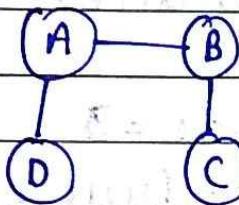
$E' = |V| - 1$



a.)



b.)



c.)

All the trees have
edges 1 less than
vertices.

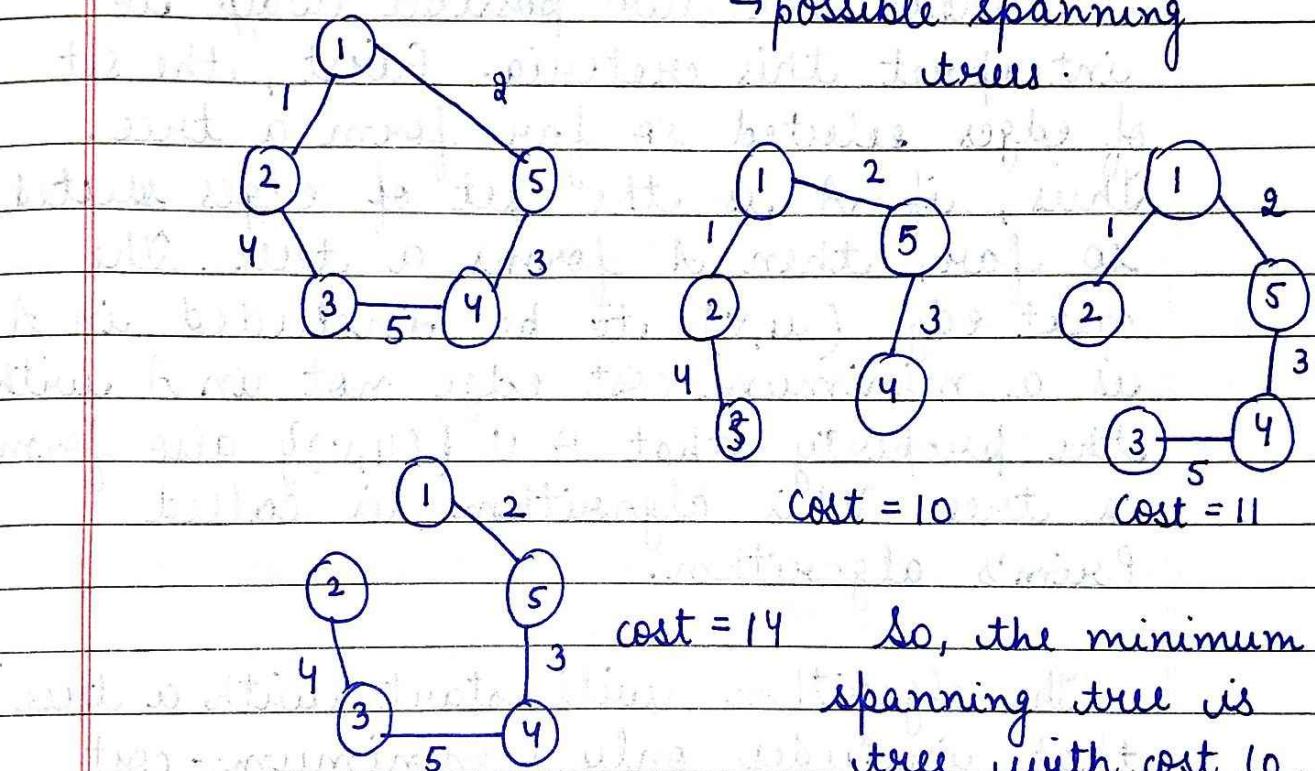
If the graph is not connected, i.e. graph with n -vertices has $n-1$ edges, spanning tree is not possible.

Minimum spanning tree:

When, a connected weighted graph, then to create a spanning tree T for G , such that the weight of edges in T is as small as possible. Such a tree is called minimum spanning tree.

E.g.

→ possible spanning trees



The minimum spanning tree should follow below rules:-

1. If G is a graph with vertices V and E . Then, minimum spanning tree G' should be, $G'(V', E')$ such that,
- $$V' = V$$
- $$E' \subseteq E \text{ or } |E'| = |V| - 1$$
2. The minimum spanning tree shouldn't create a cycle.
 3. The MST should be connected.

edge by edge. The next edge to include chosen according to some optimization criteria. The simplest criteria is to choose an edge that results in minimum increase in the sum of the costs of edges so far included.

There are two possible ways to interpret this criterion. First, the set of edges selected so far form a tree. Thus, if A is the set of edges selected so far, then A forms a tree. The next edge (u, v) to be included in A is a minimum-cost edge not in A with the property that $A \cup \{(u, v)\}$ also form a tree. This algorithm is called Prims algorithm.

The algorithm will start with a tree that includes only a minimum-cost edge of G . Then, edges are added to this tree one by one. The next edge (u, v) to be added is such that ' v ' is a vertex already included in tree & v is not included yet (v is adjacent to u) and the cost of $\text{cost}[u, v]$ is minimum among all edges (K, L) such that K is in tree & vertex L is not in tree.

To determine this edge (u, v) efficiently,

Algorithm Prims(E, cost, n, t)

```
1.  $E \rightarrow \text{Set of edges}$ 
2.  $\text{cost}[1:n, 1:n] \rightarrow \text{cost adjacency matrix}$ 
   of an  $n$  vertex graph such that  $\text{cost}[i, j]$ 
   is either positive number or  $\infty$  if no
```

edge exists.

3. The minimum spanning tree is stored as set of edges in the array $t[1:n-1, 1:n]$ such that $(t[i, 1], t[i, 2])$ is an edge in minimum spanning tree.

The final cost 'mincost' is returned.

Let (K, L) be an edge of minimum cost in E ,

$$\text{mincost} = \text{cost}[K, L];$$

1. $t[1, 1] = K; t[1, 2] = L;$
2. for $i = 1$ to n do $\|$ initialize near

if $\text{cost}[u, L] < \text{cost}[u, K]$

$\text{near}[L] = u;$

else

 we associate ' u ' with each vertex v not yet included in tree with value $\text{near}[v]$. The value $\text{near}[v]$ is a vertex in a tree such that $\text{cost}[v, \text{near}[v]]$ is minimum among all choices for $\text{near}[v]$.

The $\text{near}[v]$ is set to 0 for all vertices v that are included in tree. The next edge to be included is defined by i such that $\text{near}[i] \neq 0 \& \text{cost}[i, \text{near}[i]]$ is minimum.

3. Set $\text{near}[K] = \text{near}[L] = 0$,

4. for $i=2$ to $n-1$ do

{

 1) Find $n-i$ edges for tree.

 2) If j be index where $\text{near}[j] \neq 0$ &

$\text{cost}[i,j, \text{near}[j]]$ is minimum:

$t[i,j] = j$; $\text{near}[i,j] = \text{near}[j]$;

$\text{near}[j] = 0$;

$\text{near}[i] = \text{near}[i] + 1$;

 for $K=1$ to n do

 if ($\text{near}[K] \neq 0$ and $\text{cost}(K, \text{near}[K]) <$

$\text{cost}(K, j)$)

 then $\text{near}[K] = j$;

5. return near ;

}

The time complexity of Prim's algorithm is

$O(n^2)$. [as, the step 4 has complexity

$O(n)$ and nested loop has $O(n)$].

So, total complexity is $O(n^2)$.

Kruskal's algorithm:

In Kruskal's algorithm, edges are considered in non-decreasing order i.e.

increasing order of cost. In this algo, the set of edges selected at the end will be

result into a tree i.e. it is possible to complete 't' into a tree. Thus it may not be a tree at all stages of algorithm.

It will generally only be a forest & not of edges t can be converted into a tree if there are no cycles in t. This method is known as Kruskal algorithm.

1. $t = \emptyset$;

while ((t has less than $n-1$ edges) & $E \neq \emptyset$) do

{

 1). choose an edge (v_i, v_j) from E of

 lowest cost;

 2. Delete (v_i, v_j) from E;

 3. if (v_i, v_j) doesn't form a cycle in t

 then add (v_i, v_j) unto t;

 else

 discard (v_i, v_j) ;

}

The time complexity of above method

is $O(|V||E|)$.

As $|V| = n$, $|E| = n-1$

So, $O(n(n-1)) = O(n^2) + O(n)$

$= O(n^2)$

If the edges are maintained as a minheap, then the next edge to consider can be obtained in $O(\log |E|)$ time. The construction of heap itself takes $O(|E|)$ time.

To determine whether an edge inclusion form a cycle, the vertices in G should be grouped together in such a way that one can easily determine whether the vertices v and w are already connected by earlier selection edge. If they are, then the edge (v, w) be discarded. If they aren't, the edge

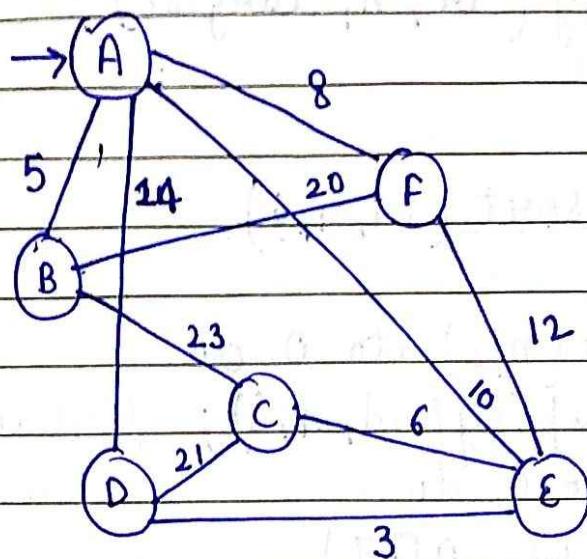
(v, w) is added in t . One possible way of grouping is to place all the vertices in same ^{connected} component of it into a set. So, two vertices v & w , if connected are in same set. E.g. \rightarrow The edge $(2, 6)$ is to be considered, the sets are $\{1, 2\}$, $\{3, 4, 6\}$ & $\{5\}$. Vertices 2 & 6 are not in same set, so these sets are combined $\rightarrow \{1, 2, 3, 4, 6\}$ & $\{5\}$. The next edge to be considered is $(1, 4)$. Since 1 & 4 are in same set, the edge is rejected. The next edge $(3, 5)$ connects vertices in two sets into single set & result in formation of minimum spanning tree.

```

# Algorithm kruskal( $\epsilon$ , cost, n, t) {
    Construct a heap out of edge costs using
    heapify.
    for  $i=1$  to  $n$  do ; parent[i] = -1;
        // Each vertex is in different set.
         $i=0$ ; mincost = 0.0;
        while ( $i < n-1$  & (heap not empty)) do
            { Delete a minimum cost edge  $(u, v)$  from
            heap & reheapify using Adjust;
             $j = \text{Find}(u)$ ;  $k = \text{Find}(v)$ ;
            if ( $j \neq k$ ) {
                 $i = i+1$ ;
                 $t[i, 1] = u$ ;  $t[i, 2] = v$ ;
                mincost = mincost + cost[u, v];
                union(j, k);
            }
            if ( $i \neq n-1$ ) then write "No MST";
            else { return mincost; } }
```

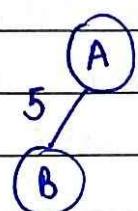
11 Examples of Prim's Algo & Kruskal's algo;

1.



Prim's algo:

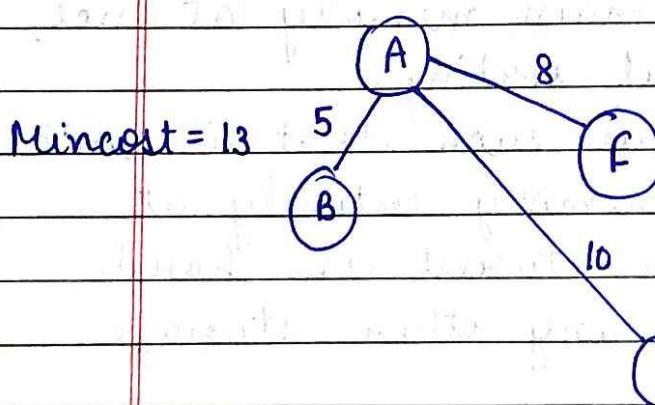
a) Choose mincost edge.



Neighbors of A $\rightarrow B, F, E, D$
 B $\rightarrow A, F, C$.

Mincost = 5

b) A $\rightarrow F$ cost is minimum among all the choices present.



Mincost = 13

Neighbors of F $\rightarrow E, B$
 (12), (20)

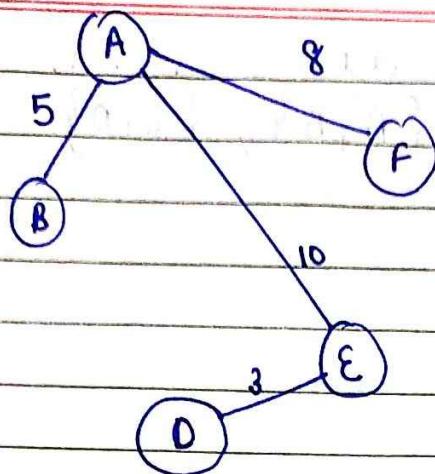
c) A $\rightarrow E, D$
 (10), (20)

Minimum cost is

Mincost = 23 AE, so it is selected.

d) $E \rightarrow C = 6, E \rightarrow D = 3$

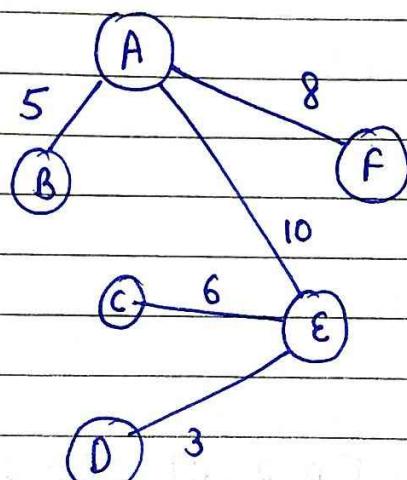
So, ED edge is selected.



$\text{Mincost} = 26$

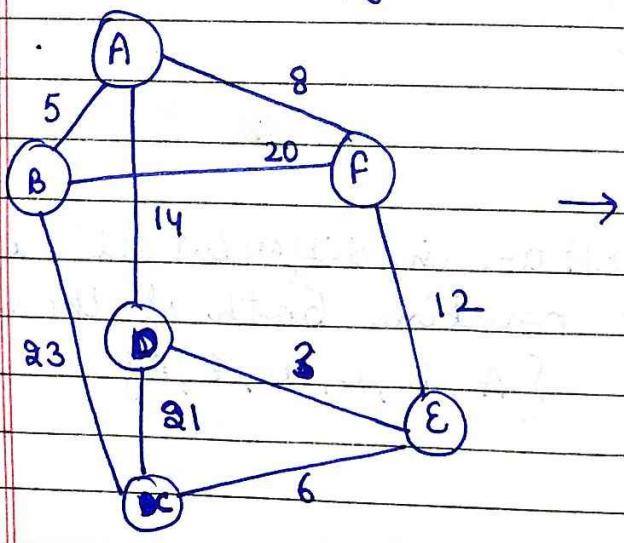
e.) $D \rightarrow C = 21$, $E \rightarrow C = 6$, $B \rightarrow C = 23$, $A \rightarrow C = \infty$,
 $F \rightarrow C = \infty$.

So, edge EC is selected.

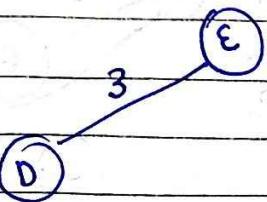


$\text{Mincost} = 32$

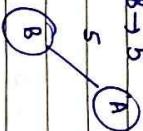
Kruskal's algorithm:



(ij) $DE \rightarrow 3$



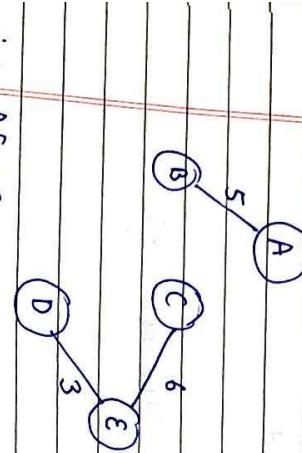
(ii) $AB \rightarrow 5$ $\{A, B\}, \{E, D\}$



$\{A, B\}, \{C, E, D\}$

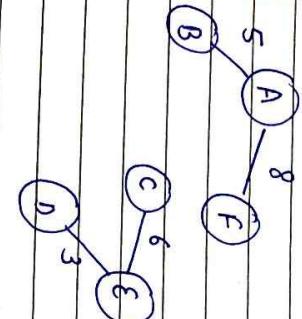
(iii) $EC \rightarrow 6$

$\{A, B\}, \{C, E, D\}$



(iv) $AF \rightarrow 8$

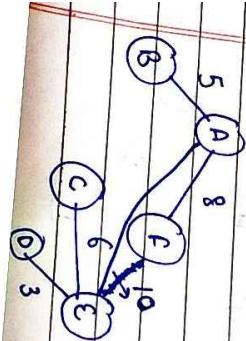
$\{A, B, F\}, \{C, E, D\}$



v) $EA \rightarrow 10$

[EA are in different set, so we combine both of them]

$\{A, B, C, D, E, F\}$



Single source shortest path:

In directed graph, $G = (V, E)$, a weighting function $w(e)$ for edges of G , vertex v_0 referred as source vertex. The problem is to determine the shortest path from v_0 to all remaining vertices of G called destination.

The greedy way to generate shortest path from v_0 to remaining vertices in its generate these paths in increasing order. First, a shortest path of nearest vertex is generated. Then, a shortest path to next nearest vertex is generated & so on.

So, we determine

(i) The next vertex to which a shortest path is generated.

(ii) A shortest path to that vertex.
Let S be set of vertices to which shortest path is already determined. For vertex w not in S , let $dist[w]$ be length of shortest path starting from w , going through only those vertices that are in S & ending at w .

Algorithm ShortestPath(v , $cost$, $dist$, n)

1. $dist[j], 1 \leq j \leq n$, is set to length to path

for vertex v i.e. source to j in graph G with n vertices. $dist[v]$ is set to zero. G is represented by

cost adjacency matrix cost [1:n, 1:n]

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```
for u = 1 to n do {
    set S[i] = false ;
    dist[i] = cost[v, i] ;
```

```
S[v] = true ; dist[v] = 0.0 ;
// As source vertex is included in
set S.
```

```
for num 2 to n-1 do
```

Choose u from among those

vertices not in S & dist[u] is

minimum.

```
S[u] = true; //Put u in S.
for each w adjacent to u, S[w] = false
do
```

```
if (dist[w] > (dist[u] + cost[u,w])) then
    dist[w] = dist[u] + cost[u,w];
```

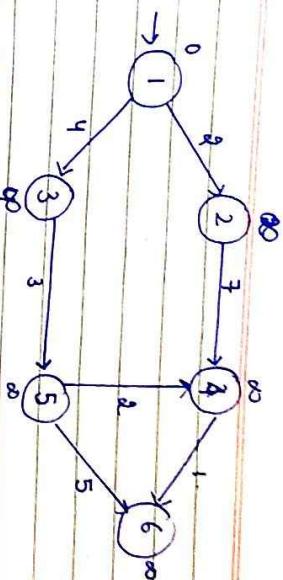
```
// Update distances.
```

```
}
```

} Time complexity for single ~~sinked~~ source
shortest path algorithm is $O(n^2)$

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L9:



$$S = \{1, 3\}$$

$$\text{dist}[1] = 0;$$

Vertex 1 is the source vertex.
So, $\text{dist}[2] = 2$

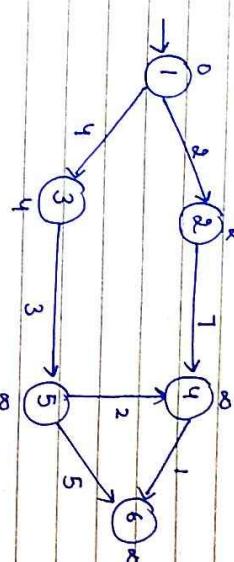
$$\text{dist}[3] = 4.$$

$$S = \{1, 2, 3\}$$

$$\text{dist}[2] > \text{dist}[1] + \text{cost}[1, 2]$$

$$\infty > 0 + 2; \infty > 2. \text{ Yes,}$$

$$\text{dist}[2] = 2;$$



$$\text{if } (\text{dist}[w] > (\text{dist}[u] + \text{cost}[u,w])) \text{ then}$$

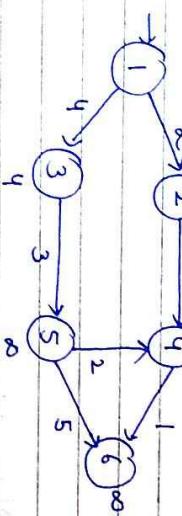
(iii)

$$S = \{1, 2, 3\}, \text{ let } w=4, u=2$$

$$\text{if } (\text{dist}[w] > \text{dist}[u] + \text{cost}[u,w])$$

$$\infty > 2 + 7; \infty > 9. \text{ Yes,}$$

$$\text{dist}[w] = 9;$$

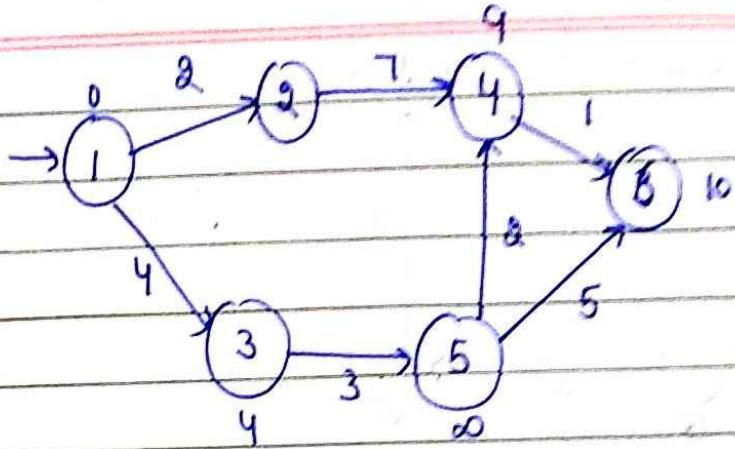


$$(iii) S = \{1, 2, 3, 4\}$$

$$u = 4, N = 6$$

$$\text{dist}[6] > 9+1; \infty > 10.$$

$$\text{So, } \text{dist}[6] = 10.$$



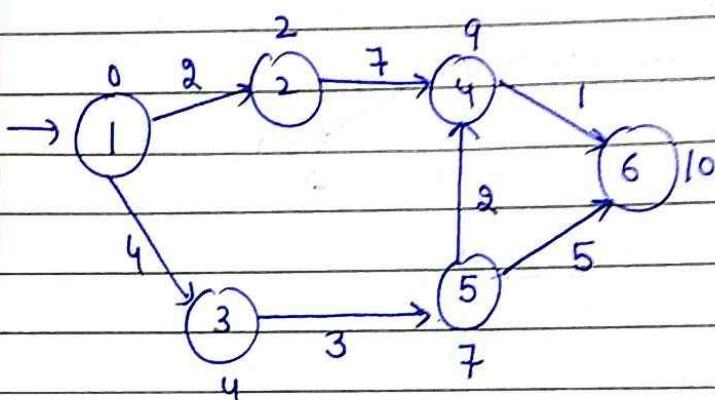
$$S = \{1, 2, 3, 4, 6\}$$

let $u = 3, w = 5$

$\text{dist}[5] > \text{dist}[3] + \text{cost}[3, 5]$

$\infty > 4 + 3; \infty > 7$

So, $\text{dist}[5] = 7$.



let $u = 5, w = 4$

if $\text{dist}[4] > \text{dist}[5] + \text{cost}[5, 4]$

$9 > 7 + 2$, No.

let $u = 5, w = 6$

if $\text{dist}[6] > \text{dist}[5] + \text{cost}[5, 6]$

$10 > 7 + 5$

$10 > 12$, No.

So, final shortest path.

