

LECTURENOTES

DESIGN & ANALYSIS OF ALGORITHM

B.Tech,4THSemester,CSE

Preparedby:

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COURSECONTENT

DESIGN & ANALYSIS OF ALGORITHM

B.Tech,4THSemester,CSE

➤ Notation of algorithm

{PageNo. 1}

Growth of functions, Recurrences: The Master method, The Substitution method, The Iteration method, Asymptotic Notations and Basic Efficiency Classes (Use of Big O, θ , etc.) in analysis of algorithms, Mathematical Analysis of few Non-Recursive and Recursive Algorithms.

➤ Sorting and searching techniques

{Page No. 42}

Selection Sort, Bubble Sort, Insertion Sort, Sequential Search, Binary Search, Depth First Search and Breadth First Search, Balanced Search Trees, AVL Trees, Red-Black Trees, Heaps and Heap Sort, Disjoint Set and their Implementation, Divide and Conquer Paradigm of problem solving, Complexity analysis and understanding of Merge Sort, Quick Sort, Binary Search Trees..

➤ Greedy techniques

{PageNo. 68}

Prim's Algorithm, Kruskal's Algorithm, Dijkstra's and Bellman Ford Algorithm, Huffman Trees, Knapsack problem.

Dynamic Programming Paradigm: Floyd-Warshall Algorithm, Optimal Binary Search trees, Matrix Chain Multiplication Problem, Longest Common Subsequence Problem, 0/1 Knapsack Problem, Maximum Network Flow Problem.

➤ String matching Algorithms

{Page No. 96}

Naive string-matching algorithm, The Rabin-Karp Algorithm, string matching with Finite Automata, Knuth Morris Pratt string matching algorithm.

Backtracking: n-Queen's problem, Hamiltonian Circuit problem, Subset-Sum problem, State Space Search Tree for these problems

➤ Branch and Bound

{Page No. 112}

Travelling Salesman Problem and its State Space Search Tree.

Introduction to Computability: Polynomial-time verification, NP-Completeness and Reducibility, NP- Complete problems. Approximation Algorithms: Vertex Cover Problem.

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B.Tech,4THSemester,CSE

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- [1] Computer Algorithms/C++, Ellis Horowitz, SatrajSahni and Rajasekaran, 2nd Edition, 2014, Universities Press..
- [2] Introduction to the Design and Analysis of Algorithms, Anany Levitin: 2nd Edition, 2009. Pearson.
- [3] R. S. Salaria, Khanna, “Data Structure & Algorithms”, Khanna Book Publishing Co. (P) Ltd.T.H. Cormen, C.E. Leiserson, R.L. Rivest and C. Stein, “Introduction to Algorithms”, PHI Publication. 2007.
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UNITI:

Introduction: Algorithm, Psuedo code for expressing algorithms, Performance Analysis- Space complexity, Time complexity, Asymptotic Notation- Big Oh notation, Omega notation, Theta notation and Little oh notation, Probabilistic analysis, Amortized analysis.

Divide and conquer: General method, applications- Binary search, Quicksort, Mergesort, Strassen's matrix multiplication.

INTRODUCTION TO ALGORITHM

History of Algorithm

- The word algorithm comes from the name of a Persian author, Abu Ja'far Mohammed ibn Musa al Khwarizmi (c. 825 A.D.), who wrote a textbook on mathematics.
- He is credited with providing the step-by-step rules for adding, subtracting, multiplying, and dividing ordinary decimal numbers.
- When written in Latin, the name became Algorismus, from which algorithm is but a small step.
- This word has taken on a special significance in computer science, where "algorithm" has come to refer to a method that can be used by a computer for the solution of a problem.
- Between 400 and 300 B.C., the great Greek mathematician Euclid invented an algorithm for finding the greatest common divisor (gcd) of two positive integers.
- The gcd of X and Y is the largest integer that exactly divides both X and Y.
- Eg., the gcd of 80 and 32 is 16.
- The Euclidian algorithm, as it is called, is considered to be the first non-trivial algorithm ever devised.

What is an Algorithm?

Algorithm is a set of steps to complete a task.

For example,

Task: to make a cup of tea.

Algorithm:

- add water and milk to the kettle,
- boil it, add tea leaves,
- Add sugar, and then serve it in a cup.

“a set of steps to accomplish or complete a task that is described precisely enough that a computer can run it”.

Described precisely: very difficult for a machine to know how much water, milk to be added etc. in the above tea making algorithm.

These algorithms run on computers or computational devices..For example, GPS in our smartphones, Google hangouts.

GPS uses shortest path algorithm. **Online shopping** uses cryptography which uses RSA algorithm.

- Algorithm Definition 1:

- An algorithm is a finite set of instructions that, if followed, accomplishes a particular task. In addition, all algorithms must satisfy the following criteria:

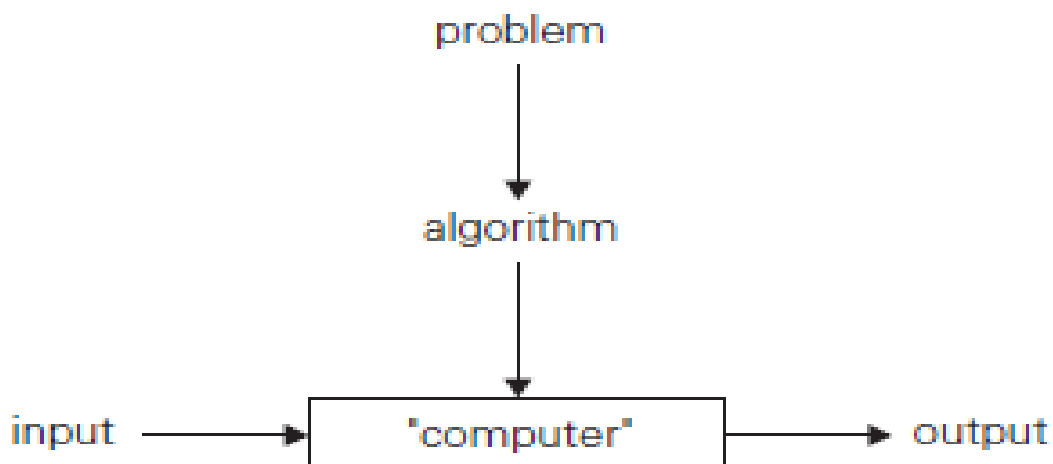
- Input. Zero or more quantities are externally supplied.
- Output. At least one quantity is produced.
- Definiteness. Each instruction is clear and unambiguous.
- Finiteness. The algorithm terminates after a finite number of steps.
- Effectiveness. Every instruction must be very basic enough and must be feasible.

- Algorithm Definition 2:

- An algorithm is a sequence of unambiguous instructions for solving a problem, i.e., for obtaining a required output for any legitimate input in a finite amount of time.

- Algorithms that are definite and effective are also called computational procedures.

- A program is the expression of an algorithm in a programming language



- Algorithms for Problem Solving

The main steps for Problem Solving are:

1. Problem definition
2. Algorithm design/Algorithm specification
3. Algorithm analysis
4. Implementation
5. Testing
6. [Maintenance]

- Step1.ProblemDefinition

What is the task to be accomplished?

Ex: Calculate the average of the grades for a given student

- Step2.AlgorithmDesign / Specifications:

Describe: in natural language/pseudo-code/diagrams/ etc

- Step3.Algorithm analysis

Space complexity-How much space is required

Time complexity-How much time does it take to run the algorithm

Computer Algorithm

An algorithm is a procedure (a finite set of well-defined instructions) for accomplishing some tasks which, given an initial state, terminate in a defined end-state

The computational complexity and efficient implementation of the algorithm are important in computing, and this depends on suitable data structures.

- Steps 4,5,6: Implementation, Testing, Maintenance

- Implementation:

Decide on the programming language to use C, C++, Lisp, Java, Perl, Prolog, assembly, etc., etc.

Write clean, well documented code

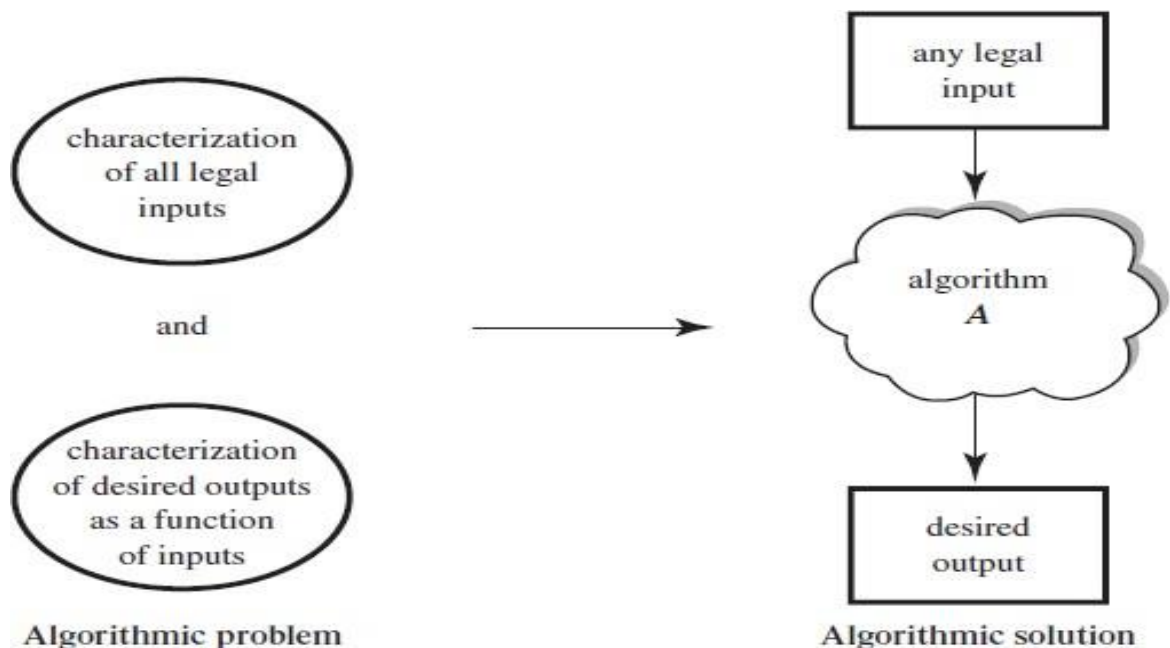
- Test, test, test

Integrate feedback from users, fix bugs, ensure compatibility across different versions

- Maintenance.

Release Updates, fix bugs

Keeping illegal inputs separate is the responsibility of the algorithmic problem, while treating special classes of fun, unusual or undesirable inputs is the responsibility of the algorithm itself.



- **4 Distinct areas of study of algorithms:**

- How to devise algorithms. → Techniques – Divide & Conquer, Branch and Bound, Dynamic Programming
- How to validate algorithms.
- Check for Algorithm that it computes the correct answer for all possible legal inputs. → algorithm validation. → First Phase
- Second phase → Algorithm to Program → Program Proving or Program Verification → Solution bestated into two forms:
- First Form: Program which is annotated by a set of assertions about the input and output variables of the program → predicate calculus
- Second form: is called as specification
- 4 Distinct areas of study of algorithms (..Contd)
- How to analyze algorithms.
- Analysis of Algorithms or performance analysis refers to the task of determining how much computing time & storage an algorithm requires
- How to test a program → 2 phases →
- Debugging - Debugging is the process of executing programs on sampled datasets to determine whether faulty results occur and, if so, to correct them.
- Profiling or performance measurement is the process of executing a correct program on data sets and measuring the time and space it takes to compute the results

PSEUDOCODE:

- Algorithm can be represented in Text mode and Graphic mode
- Graphical representation is called Flowchart
- Text mode most often represented in close to any High level languages such as C, Pascal → Pseudocode
- **Pseudocode: High-level description of an algorithm.**
- → More structured than plain English.
- → Less detailed than a program.
- → Preferred notation for describing algorithms.
- → Hides program design issues.
- **Example of Pseudocode:**
- To find the max element of an array

```

Algorithm arrayMax(A, n)
Input array A of n integers
Output maximum element of A
currentMax ← A[0]
for i ← 1 to n - 1 do
    if A[i] > currentMax then
        currentMax ← A[i]
    
```


return <i>currentMax</i>

- Control flow
- if ... then... [else ...]
- while ... do ...
- repeat...until...
- for... do ...
- Indentation replaces braces
- Method declaration
- Algorithm *method(arg[,arg...])*
- Input...
- Output ...
- Method call
- *var.method(arg[,arg...])*
- Return value
- *return expression*
- Expressions
- Assignment (equivalent to =)
- Equality testing (equivalent to ==)
- n^2 Superscripts and other mathematical formatting allowed

PERFORMANCE ANALYSIS:

- What are the Criteria for judging algorithms that have a more direct relationship to performance?
- computing time and storage requirements.
- **Performance evaluation** can be loosely divided into two major phases:
- a priori estimates and
- a posteriori testing.
- ➔ refers as performance analysis and performance measurement respectively
- The space complexity of an algorithm is the amount of memory it needs to run to completion.
- The time complexity of an algorithm is the amount of computer time it needs to run to completion.

Space Complexity:

- Space Complexity Example:
- Algorithm *abc(a,b,c)*
{
 return $a+b++*c+(a+b-c)/(a+b) + 4.0$;
}

➔ The Space needed by each of these algorithms is seen to be the sum of the following component.

1. A fixed part that is independent of the characteristics (eg: number, size) of the inputs and outputs.

The part typically includes the instruction space (ie. Space for the code), space for simple variable and fixed-size component variables (also called aggregate) space for constants, and so on.

2. A variable part that consists of the space needed by component variables whose size is dependent on the particular problem instance being solved, the space needed by referenced variables (to the extent that it depends on instance characteristics), and the recursion stack space.

The space requirements $S(p)$ of any algorithm p may therefore be written as, $S(P) =$

$c + Sp(\text{Instance characteristics})$

Where 'c' is a constant.

Example 2:

Algorithm sum(a, n)

```
{  
s=0.0;  
for I=1 to n do s=  
s+a[I]; return  
s;  
}
```

- The problem instances for this algorithm are characterized by n, the number of elements to be summed. The space needed by 'n' is one word, since it is of type integer.
- The space needed by 'a' is the space needed by variables of type array of floating point numbers.
- This is at least 'n' words, since 'a' must be large enough to hold the 'n' elements to be summed.
- So, we obtain $S_{\text{sum}}(n) \geq (n + s)$
- [n for a[], one each for n, I, a & s]

Time Complexity:

- The time $T(p)$ taken by a program P is the sum of the compile time and the run time (execution time)
- The compile time does not depend on the instance characteristics. Also we may assume that a compiled program will be run several times without recompilation. This run time is denoted by $tp(\text{instance characteristics})$.
- The number of steps any problem statement is assigned depends on the kind of statement.
- For example, comments are 0 steps.
Assignment statements are 1 step.

[Which does not involve any calls to other algorithms]

Interactive statements such as for, while & repeat-until are control parts of the statement.

We introduce a variable, count into the program statement to increment count with initial value 0. Statement to increment count by the appropriate amount are introduced into the program.

This is done so that each time a statement in the original program is executed, count is incremented by the step count of that statement.

Algorithm:

AlgorithmSum(a,n)

```
{
s=0.0;
count=count+1; for
I=1 to n do
{
count=count+1;
s=s+a[I];
count=count+1;
}
count=count+1;
count=count+1;
return s;
}
```

→ If the count is zero to start with, then it will be $2n+3$ on termination. So each invocation of sum executes a total of $2n+3$ steps.

2. The second method to determine the step count of an algorithm is to build a table in which we list the total number of steps contributed by each statement.

→ First determine the number of steps per execution (s/e) of the statement and the total number of times (ie., frequency) each statement is executed.

→ By combining these two quantities, the total contribution of all statements, the step count for the entire algorithm is obtained.

Statement	Steps per execution	Frequency	Total
1. AlgorithmSum(a,n)	0	-	0
2. {	0	-	0
3. S=0.0;	1	1	1
4. for I=1 to n do	1	$n+1$	$n+1$
5. s=s+a[I];	1	$n+1$	$n+1$
6. return s;	1	-	0
7. }	0		
Total			$2n+3$

How to analyse an Algorithm?

Let us form an algorithm for Insertion sort (which sorts a sequence of numbers). The pseudo code for the algorithm is given below.

Pseudo code for Insertion Algorithm:

Identify each line of the pseudocode with symbols such as C1, C2 ..

Pseudo code for Insertion Algorithm	Line Identification
for j=2 to A length	C1
key=A[j]	C2
//Insert A[j] into sorted Array A[1. j-1]	C3
i=j-1	C4
while i>0 & A[i]>key	C5
A[i+1]=A[i]	C6
i=i-1	C7
A[i+1]=key	C8

Let C₃ be the cost of this line. Since comment lines will not incur any cost C₃=0.

Cost	No. Of times Executed
C1	N
C2	n-1
C3=0	n-1
C4	n-1
C5	$\sum_{j=2}^{n-1} t_j$
C6	$\sum_{j=2}^n t_j - 1$
C7	$\sum_{j=2}^n t_j - 1$
C8	n-1

Running time of the algorithm is:

$$T(n) = C_1 n + C_2 (n-1) + 0(n-1) + C_4 (n-1) + C_5 \left(\sum_{j=2}^{n-1} t_j \right) + C_6 \left(\sum_{j=2}^n t_j - 1 \right) + C_7 \left(\sum_{j=2}^n t_j - 1 \right) + C_8 (n-1)$$

Best case:

It occurs when Array is sorted. All t_j

values are 1.

$$T(n) = C_1n + C_2(n-1) + 0(n-1) + C_4(n-1) + C_5\left(\sum_{j=2}^{n-1} 1\right) + C_6\left(\sum_{j=2}^n 1-1\right) + C_7\left(\sum_{j=2}^n 1-1\right) + C_8(n-1)$$

$$= C_1n + C_2(n-1) + 0(n-1) + C_4(n-1) + C_5 + C_8(n-1)$$

$$= (C_1 + C_2 + C_4 + C_5 + C_8)n - (C_2 + C_4 + C_5 + C_8)$$

· Which is of the form $an + b$.

→ Linear function of n .

→ So, linear growth.

Worstcase:

It occurs when Array is reverse sorted, and $tj = j$

$$T(n) = C_1n + C_2(n-1) + 0(n-1) + C_4(n-1) + C_5\left(\sum_{j=2}^{n-1} j\right) + C_6\left(\sum_{j=2}^n j-1\right) + C_7\left(\sum_{j=2}^n j-1\right) + C_8(n-1)$$

$$= C_1n + C_2(n-1) + C_4(n-1) + C_5\left(\frac{n(n-1)}{2} - 1\right) + C_6\left(\sum_{j=2}^n \frac{n(n-1)}{2}\right) + C_7\left(\sum_{j=2}^n \frac{n(n-1)}{2}\right) + C_8(n-1)$$

which is of the form $an^2 + bn + c$

Quadratic function. So in worst case insertion sort grows in n^2 . Why we

concentrate on worst-case running time?

· The worst-case running time gives a guaranteed upper bound on the running time for any input.

· For some algorithms, the worst case occurs often. For example, when searching, the worst case often occurs when the item being searched for is not present, and searches for absent items may be frequent.

· Why not analyze the average case? Because it's often about as bad as the worst case.

Order of growth:

It is described by the highest degree term of the formula for running time. (Drop lower-order terms. Ignore the constant coefficient in the leading term.)

Example: We found out that for insertion sort the worst-case running time is of the form $an^2 + bn + c$.

Drop lower-order terms. What remains is an^2 . Ignore constant coefficient. It results in n^2 . But we cannot say that the worst-case running time $T(n)$ equals n^2 . Rather it grows like n^2 . But it doesn't equal n^2 . We say that the running time is $\Theta(n^2)$ to capture the notion that the order of growth is n^2 .

We usually consider one algorithm to be more efficient than another if its worst-case running time has a smaller order of growth.

Complexity of Algorithms

The complexity of an algorithm M is the function $f(n)$ which gives the running time and/or storage space requirement of the algorithm in terms of the size ' n ' of the input data. Mostly, the storage space required by an algorithm is simply a multiple of the data size ' n '.

Complexity shall refer to the running time of the algorithm.

The function $f(n)$, gives the running time of an algorithm, depends not only on the size ' n ' of the input data but also on the particular data. The complexity function $f(n)$ for certain cases are:

1. **Best Case:** The minimum possible value of $f(n)$ is called the best case.
2. **Average Case:** The expected value of $f(n)$.
3. **Worst Case:** The maximum value of $f(n)$ for any key possible input.

ASYMPTOTIC NOTATION

→ Formal way notation to speak about functions and classify them

The following notations are commonly used notations in performance analysis and used to characterize the complexity of an algorithm:

1. Big-O (O),
2. Big-OMEGA (Ω),
3. Big-THETA (Θ) and
4. Little-O (o)

Asymptotic Analysis of Algorithms:

Our approach is based on the *asymptotic complexity* measure. This means that we don't try to count the exact number of steps of a program, but how that number grows with the size of the input to the program. That gives us a measure that will work for different operating systems, compilers and CPUs. The asymptotic complexity is written using big-O notation.

- It is a way to describe the characteristics of a function in the limit.
- It describes the rate of growth of functions.
- Focus on what's important by abstracting away low-order terms and constant factors.
- It is a way to compare "sizes" of functions: $O \approx$

$\leq$

$\Omega \approx \geq$
 $\Theta \approx =$
 $o \approx <$
 $\omega \approx >$

Time complexity	Name	Example
$O(1)$	Constant	Adding an element to the front of a linked list
$O(\log n)$	Logarithmic	Finding an element in a sorted array
$O(n)$	Linear	Finding an element in an unsorted array
$O(n \log n)$	Linear	Logarithmic Sorting items by 'divide-and-conquer' - Mergesort
$O(n^2)$	Quadratic	Shortest path between two nodes in a graph
$O(n^3)$	Cubic	Matrix Multiplication
$O(2^n)$	Exponential	The Towers of Hanoi problem

Big 'oh': the function $f(n) = O(g(n))$ iff there exist positive constants c and n_0 such that $f(n) \leq c \cdot g(n)$ for all $n, n \geq n_0$.

Omega: the function $f(n) = \Omega(g(n))$ iff there exist positive constants c and n_0 such that $f(n) \geq c \cdot g(n)$ for all $n, n \geq n_0$.

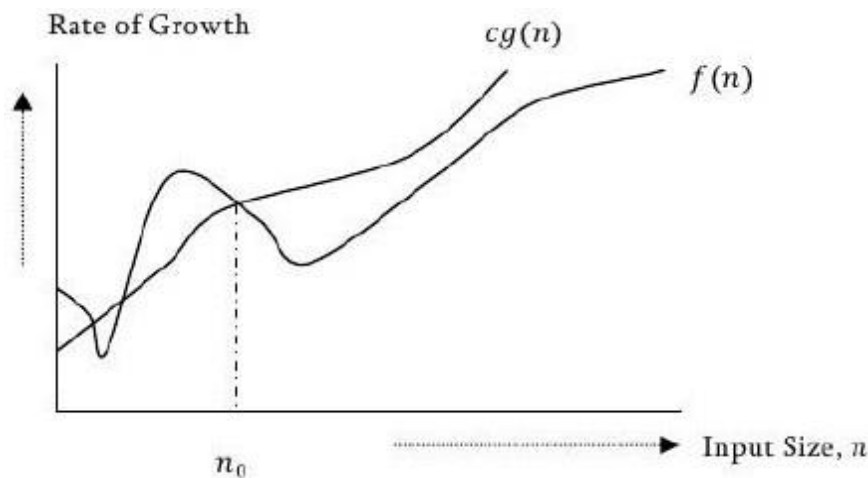
Theta: the function $f(n) = \Theta(g(n))$ iff there exist positive constants c_1, c_2 and n_0 such that $c_1 \cdot g(n) \leq f(n) \leq c_2 \cdot g(n)$ for all $n, n \geq n_0$.

Big-O Notation

This notation gives the tight upper bound of the given function. Generally we represent it as $f(n) = O(g(n))$. That means, at larger values of n , the upper bound of $f(n)$ is $g(n)$. For example, if $f(n) = n^4 + 100n^2 + 10n + 50$ is the given algorithm, then n^4 is $g(n)$. That means $g(n)$ gives the maximum rate of growth for $f(n)$ at larger values of n .

O—notation defined as $O(g(n)) = \{f(n) : \text{there exist positive constants } c \text{ and } n_0 \text{ such that } 0 \leq f(n) \leq c \cdot g(n) \text{ for all } n \geq n_0\}$. $g(n)$ is an asymptotic tight upper bound for $f(n)$. Our objective is to give some rate of growth $g(n)$ which is greater than given algorithm's rate of growth $f(n)$.

In general, we do not consider lower values of n . That means the rate of growth at lower values of n is not important. In the below figure, n_0 is the point from which we consider the rate of growth for a given algorithm. Below n_0 the rate of growths may be different.



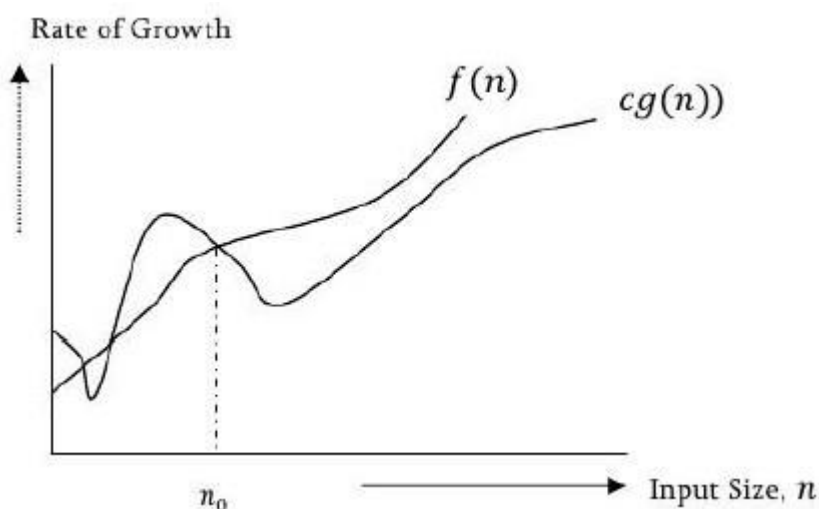
Note Analyze the algorithms at large values of n only. What this means is, below now we do not care for rates of growth.

Omega— Ω notation

Similar to above discussion, this notation gives the tighter lower bound of the given algorithm and we represent it as $f(n) = \Omega(g(n))$. That means, at large values of n , the tighter lower bound of $f(n)$ is g .

For example, if $f(n) = 100n^2 + 10n + 50$, $g(n)$ is $\Omega(n^2)$.

The Ω notation as be defined as $\Omega(g(n)) = \{f(n) : \text{there exist positive constants } c \text{ and } n_0 \text{ such that } 0 \leq cg(n) \leq f(n) \text{ for all } n \geq n_0\}$. $g(n)$ is an asymptotic lower bound for $f(n)$. $\Omega(g(n))$ is the set of functions with smaller or same order of growth as $f(n)$.

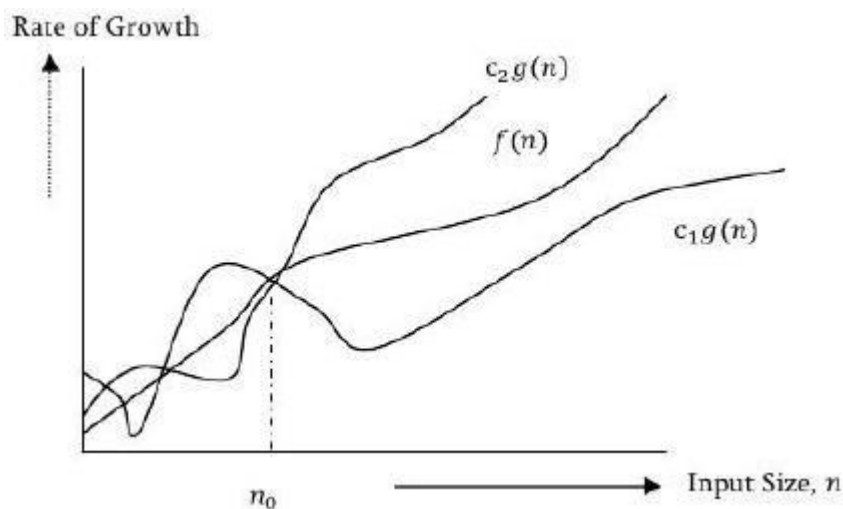


Theta- Θ notation

This notation decides whether the upper and lower bounds of a given function are same or not. The average running time of an algorithm is always between lower bound and upper bound.

If the upper bound (O) and lower bound (Ω) gives the same result then Θ notation will also have the same rate of growth. As an example, let us assume that $f(n) = 10n + n$ is the expression. Then, its tight upper bound $g(n)$ is $O(n)$. The rate of growth in best case is $g(n) = O(n)$. In this case, rate of growths in best case and worst are same. As a result, the average case will also be same.

None: For a given function (algorithm), if the rate of growths (bounds) for O and Ω are not same then the rate of growth Θ case may not be same.



Now consider the definition of Θ notation. It is defined as $\Theta(g(n)) = \{f(n) : \text{there exist positive constants } C_1, C_2 \text{ and } n_0 \text{ such that } C_1 g(n) \leq f(n) \leq C_2 g(n) \text{ for all } n \geq n_0\}$. $g(n)$ is an asymptotic tight bound for $f(n)$. $\Theta(g(n))$ is the set of functions with the same order of growth as $g(n)$.

Important Notes

For analysis (best case, worst case and average) we try to give upper bound (O) and lower bound (Ω) and average running time (Θ). From the above examples, it should also be clear that, for a given function (algorithm) getting upper bound (O) and lower bound (Ω) and average running time (Θ) may not be possible always.

For example, if we are discussing the best case of an algorithm, then we try to give upper bound (O) and lower bound (Ω) and average running time (Θ).

In the remaining chapters we generally concentrate on upper bound (O) because knowing lower bound (Ω) of an algorithm is of no practical importance and we use Θ notation if upper bound (O) and lower bound (Ω) are same.

Little Oh Notation

The little Oh is denoted as o . It is defined as: Let, $f(n)$ and $g(n)$ be the nonnegative functions then

$$\lim_{n \rightarrow \infty} \frac{f(n)}{g(n)} = 0$$

such that $f(n) = o(g(n))$ i.e. $f(n)$ is little O of $g(n)$.

$f(n) = o(g(n))$ if and only if $f(n) = O(g(n))$ and $f(n) \neq \Theta(g(n))$

PROBABILISTIC ANALYSIS

Probabilistic analysis is the use of probability in the analysis of problems.

In order to perform a probabilistic analysis, we must use knowledge of, or make assumptions about, the distribution of the inputs. Then we analyze our algorithm, computing an average-case running time, where we take the average over the distribution of the possible inputs.

Basics of Probability Theory

Probability theory has the goal of characterizing the outcomes of natural or conceptual “experiments.” Examples of such experiments include tossing a coin ten times, rolling a die three times, playing a lottery, gambling, picking a ball from an urn containing white and red balls, and so on

Each possible outcome of an experiment is called a sample point and the set of all possible outcomes is known as the sample space S . In this text we assume that S is finite (such a sample space is called a discrete sample space). An event E is a subset of the sample space S . If the sample space consists of n sample points, then there are 2^n possible events.

Definition—Probability: The probability of an event E is defined to be $\frac{|E|}{|S|}$ where S is the sample space.

Then the **indicator random variable** $I\{A\}$ associated with the event A is defined as

$I\{A\} = 1$ if A occurs ;
0 if A does not occur

The probability of event E is denoted as $\text{Prob.}[E]$. The complement of E , denoted E^c , is defined to be $S - E$. If E_1 and E_2 are two events, the probability of E_1 or E_2 or both happening is denoted as $\text{Prob.}[E_1 \cup E_2]$. The probability of both E_1 and E_2 occurring at the same time is denoted as $\text{Prob.}[E_1 \cap E_2]$. The corresponding event is $E_1 \cap E_2$.

Theorem 1.5

1. $\text{Prob.}[E^c] = 1 - \text{Prob.}[E]$.
2. $\text{Prob.}[E_1 \cup E_2] = \text{Prob.}[E_1] + \text{Prob.}[E_2] - \text{Prob.}[E_1 \cap E_2]$
 $\leq \text{Prob.}[E_1] + \text{Prob.}[E_2]$

Expected value of a random variable

The simplest and most useful summary of the distribution of a random variable is the average of the values it takes on. The **expected value** (or, synonymously, **expectation** or **mean**) of a discrete random variable X is

$$E[X] = \sum x P\{X=x\}$$

which is well defined if the sum is finite or converges absolutely.

Consider a game in which you flip two fair coins. You earn \$3 for each head but lose \$2 for each tail. The expected value of the random variable X representing

your earnings is

$$\begin{aligned} E[X] &= 6 \cdot \Pr\{2H's\} + 1 \cdot \Pr\{1H, 1T\} - 4 \cdot \Pr\{2T's\} \\ &= 6(1/4) + 1(1/2) - 4(1/4) \\ &= 1 \end{aligned}$$

Any one of these first i candidates is equally likely to be the best-qualified so far. Candidate i has a probability of $1/i$ of being better qualified than candidates 1 through $i-1$ and thus a probability of $1/i$ of being hired.

$$E[X_i] = 1/i$$

So,

$$E[X] = E\left[\sum_{i=1}^n X_i\right]$$

$$\begin{aligned} &= \sum_{i=1}^n E[X_i] \\ &= \sum_{i=1}^n 1/i \end{aligned}$$

AMORTIZED ANALYSIS

In an **amortized analysis**, we average the time required to perform a sequence of data structure operations over all the operations performed. With amortized analysis, we can show that the average cost of an operation is small, if we average over a sequence of operations, even though a single operation within the sequence might be expensive. Amortized analysis differs from average-case analysis in that probability is not involved; an amortized analysis guarantees the *average performance of each operation in the worst case*.

Three most common techniques used in amortized analysis:

1. **Aggregate Analysis** - in which we determine an upper bound $T(n)$ on the total cost of a sequence of n operations. The average cost per operation is then $T(n)/n$. We take the average cost as the amortized cost of each operation
2. **Accounting method** - When there is more than one type of operation, each type of operation may have a different amortized cost. The accounting method overcharges some operations early in the sequence, storing the overcharge as "prepaid credit" on specific objects in the data structure. Later in the sequence, the credit pays for operations that are charged less than they actually cost.

3. **Potential method** - The potential method maintains the credit as the “potential energy” of the data structure as a whole instead of associating the credit with individual objects within the data structure. The potential method, which is like the accounting method in that we determine the amortized cost of each operation and may overcharge operations early on to compensate for undercharges later

DIVIDE AND CONQUER

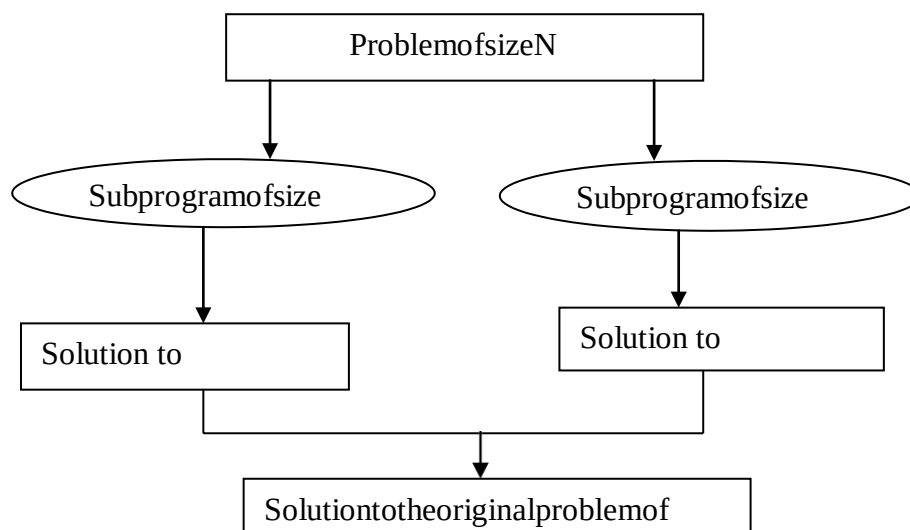
General Method

In divide and conquer method, a given problem is,

- i) Divided into smaller subproblems.
- ii) These subproblems are resolved independently.
- iii) Combining all the solutions of subproblems into a solution of the whole.

If the subproblems are large enough then divide and conquer is reapplied.
The generated subproblems are usually of some type as the original problem.

Hence recursive algorithms are used in divide and conquer strategy.



Pseudocode Representation of Divide and conquer rule for problem “P”

```

Algorithm DAndC(P)
{
  if small(P) then return S(P) else {
    divide P into smaller instances P1, P2, P3 ... Pk;
    apply DAndC to each of these subprograms; // means DAndC(P1), DAndC(P2) .....
    DAndC(Pk)
    return combine(DAndC(P1), DAndC(P2) ..... DAndC(Pk));
  }
}

// P → Problem
// Here small(P) → Boolean value function. If it is true, then the function S is
// invoked

```

Time Complexity of DAndC Algorithm:

$$T(n) = T(1) \text{ if } n=1$$

$$aT(n/b) + f(n) \text{ if } n > 1$$

$a, b \rightarrow$ constants.

This is called the **general divide and conquer recurrence**.

Example for GENERAL METHOD:

As an example, let us consider the problem of computing the sum of n numbers $a_0, \dots, a_{n-1}$.

If $n > 1$, we can divide the problem into two instances of the same problem. They are sum of the first $\lfloor n/2 \rfloor$ numbers

Compute the sum of the first $\lfloor n/2 \rfloor$ numbers, and then compute the sum of another $n/2$ numbers. Combine the answers of two $n/2$ numbers sum.

i.e.,

$$a_0 + \dots + a_{n-1} = (a_0 + \dots + a_{n/2}) + (a_{n/2} + \dots + a_{n-1})$$

Assuming that size n is a power of b , to simplify your analysis, we get the following recurrence for the running time $T(n)$.

$$T(n) = aT(n/b) + f(n)$$

This is called the **general divide and conquer recurrence**.

$f(n) \rightarrow$ is a function that accounts for the times spent on dividing the problem into smaller ones and on combining their solutions. (For the summation example, $a = b = 2$ and $f(n) = 1$).

Advantages of DAndC:

The times spent on executing the problem using DAndC is smaller than other method. This technique is ideally suited for parallel computation.

This approach provides an efficient algorithm in computer science.

Master Theorem for Divide and Conquer

In all efficient divide and conquer algorithms we will divide the problem into subproblems, each of which is some part of the original problem, and then perform some additional work to compute the final answer. As an example, if we consider merge sort [for details, refer Sorting chapter], it operates on two problems, each of which is half the size of the original, and then uses $O(n)$ additional work for merging. This gives the running time equation:

$$T(n) = 2T\left(\frac{n}{2}\right) + O(n)$$

The following theorem can be used to determine the running time of divide and conquer algorithms. For a given program or algorithm, first we try to find the recurrence relation for the problem. If the recurrence is of below form then we directly give the answer without fully solving it.

If the recurrence is of the form $T(n) = aT\left(\frac{n}{b}\right) + \Theta(n^k \log^p n)$, where $a \geq 1, b > 1, k \geq 0$ and p is a real number, then we can directly give the answer as:

- 1) If $a > b^k$, then $T(n) = \Theta(n^{\log_b a})$
- 2) If $a = b^k$
 - a. If $p > -1$, then $T(n) = \Theta(n^{\log_b a} \log^{p+1} n)$
 - b. If $p = -1$, then $T(n) = \Theta(n^{\log_b a} \log \log n)$
 - c. If $p < -1$, then $T(n) = \Theta(n^{\log_b a})$
- 3) If $a < b^k$
 - a. If $p \geq 0$, then $T(n) = \Theta(n^k \log^p n)$
 - b. If $p < 0$, then $T(n) = O(n^k)$

Applications of Divide and conquer rule or algorithm:

- Binary search,
- Quicksort,
- Mergesort,
- Strassen's matrix multiplication.

Binary search or Half-interval search algorithm:

1. This algorithm finds the position of a specified input value (the search "key") within an [array sorted by key value](#).
2. In each step, the algorithm compares the search key value with the key value of the middle element of the array.
3. If the keys match, then a matching element has been found and its index, or position, is returned.
4. Otherwise, if the search key is less than the middle element's key, then the algorithm repeats its action on the sub-array to the **left** of the middle element or, if the search key is greater, then the algorithm repeats on sub array to the **right** of the middle element.
5. If the search element is less than the minimum position element or greater than the maximum position element then this algorithm returns not found.

Binary search algorithm by using recursive methodology:

Program for binary search (recursive)	Algorithm for binary search (recursive)
int binary_search(int A[], int key, int min, int max)	Algorithm binary_search(A, key, imin, imax)

<pre> { if(imax<imin) returnarrayisempty; if(key<imin K>imax) returnelementnotinarraylistelse { intimid=(imin+imax)/2; if (A[imid] > key) returnbinary_search(A,key,imin,imid-1); else if (A[imid] < key) returnbinary_search(A,key,imid+1,imax); else returnimid; } } </pre>	<pre> { if (imax<imin) then return“arrayisempty”; if(key<imin K>imax) then return“elementnotinarraylist” else { imid=(imin+imax)/2; if(A[imid]>key)then returnbinary_search(A,key,imin,imid-1); else if (A[imid] < key) then returnbinary_search(A,key,imid+1,imax); else returnimid; } } </pre>
---	--

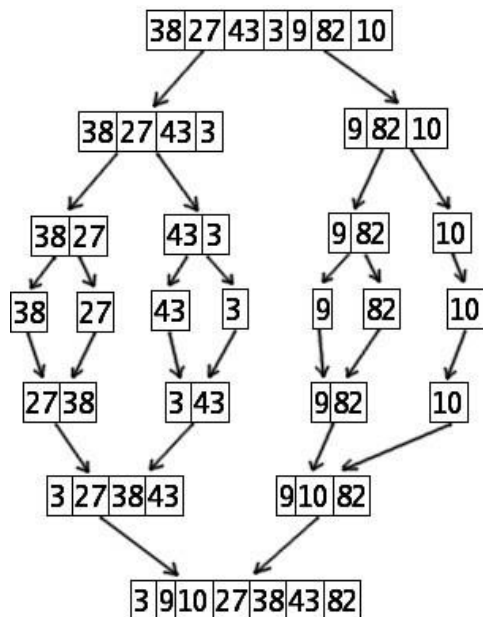
TimeComplexity:	
Datastructure:-Array	
For successful search	Unsuccessful search
Worst case → $O(\log n)$ or $\theta(\log n)$	$\theta(\log n)$:-for all cases.
Average case → $O(\log n)$ or $\theta(\log n)$	
Best case → $O(1)$ or $\theta(1)$	

Binary search algorithm by using iterative methodology:

Binary search program by using iterative methodology:	Binary search algorithm by using iterative methodology:
<pre> int binary_search(int A[], int key, int imin, int imax) { while(imax >= imin) { intimid = midpoint(imin, imax); if(A[imid] == key) returnimid; elseif(A[imid] < key) imin = imid + 1; else imax = imid - 1; } } </pre>	<pre> Algorithm binary_search(A, key, imin, imax) { While<(imax >= imin)>do { intimid = midpoint(imin, imax); if(A[imid] == key) returnimid; elseif(A[imid] < key) imin = imid + 1; else imax = imid - 1; } } </pre>

Merge Sort:

The merge sort splits the list to be sorted into two equal halves, and places them in separate arrays. This sorting method is an example of the DIVIDE-AND-CONQUER paradigm i.e. it breaks the data into two halves and then sorts the two half data sets recursively, and finally merges them to obtain the complete sorted list. The merge sort is a comparison sort and has an algorithmic complexity of $O(n \log n)$. Elementary implementations of the merge sort make use of two arrays - one for each half of the dataset. The following image depicts the complete procedure of merge sort.



Advantages of Merge Sort:

1. Marginally faster than the heap sort for larger sets
2. Merge Sort always does less number of comparisons than Quick Sort. Worst case for merge sort does about 39% less comparisons against quick sort's average case.
3. Merge sort is often the best choice for sorting a linked list because the slow random-access performance of a linked list makes some of the other algorithms (such as quick sort) perform poorly, and others (such as heap sort) completely impossible.

Program for Merge sort:

```
#include<stdio.h>
#include<conio.h>
int n;
void main(){
int i,low,high,z,y;
int a[10];
void mergesort(int a[10],int low,int high);
void display(int a[10]);
clrscr();
printf("\n\t\t\tmergesort\n");
printf("\n\t\t\tenter the length of the list:");
scanf("%d",&n);
printf("\n\t\t\tenter the list elements");
for(i=0;i<n;i++)
scanf("%d",&a[i]);
low=0;
high=n-1;
mergesort(a,low,high);
display(a);
getch();
}
void mergesort(int a[10],int low, int high)
```



```

{
int mid;
voidcombine(inta[10],intlow,intmid,inthigh);
if(low<high)
{
mid=(low+high)/2;
mergesort(a,low,mid);
mergesort(a,mid+1,high);
combine(a,low,mid,high);
}
}
voidcombine(inta[10],intlow,intmid,inthigh){ int
i,j,k;
inttemp[10];
k=low;
i=low;
j=mid+1;
while(i<=mid&& j<=high){
if(a[i]<=a[j])
{
temp[k]=a[i];
i++;
k++;
}
else
{
temp[k]=a[j];
j++;
k++;
}
}
while(i<=mid){
temp[k]=a[i];
i++;
k++;
}

while(j<=high){
temp[k]=a[j];
j++;
k++;
}
for(k=low;k<=high;k++)
a[k]=temp[k];
}
voiddisplay(inta[10]){ int
i;
printf("\n\nthesortedarrayis\n"); for(i=0;i<n;i++)
printf("%d\t",a[i]);}

```

Algorithm for Mergesort:

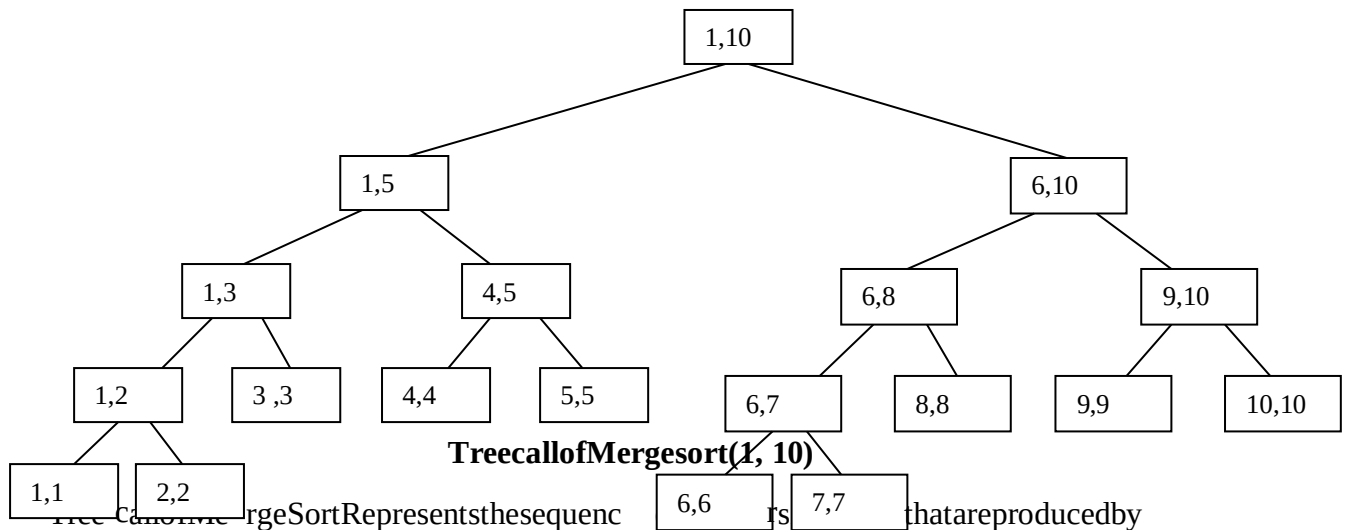
```
Algorithm mergesort(low, high)
{
  if (low < high) then
  {
    mid = (low + high) / 2;
    mergesort(low, mid);
    mergesort(mid + 1, high); // Solve the sub-problems
    Merge(low, mid, high); // Combine the solution
  }
}

void Merge(low, mid, high) {
  k = low;
  i = low;
  j = mid + 1;
  while (i <= mid && j <= high) do {
    if (a[i] <= a[j]) then
    {
      temp[k] = a[i];
      i++;
      k++;
    }
    else
    {
      temp[k] = a[j];
      j++;
      k++;
    }
  }
  while (i <= mid) do {
    temp[k] = a[i];
    i++;
    k++;
  }
  while (j <= high) do {
    temp[k] = a[j];
    j++;
    k++;
  }
  for (k = low to high) do
    a[k] = temp[k];
}
```

Tree of Merge Sort

Consider an example: (From text book)

$A[1:10] = \{310, 285, 179, 652, 351, 423, 861, 254, 450, 520\}$



Tree of Merge Sort represents the sequence of merges that are produced by mergesort.

“Once observe the explained notes in class room”

Computing Time for Merge Sort:

The time for the merging operation is proportional to n , then the computing time for merge sort is described by using recurrence relation.

$$T(n) = \begin{cases} a & \text{if } n=1; \\ 2T(n/2) + cn & \text{if } n>1 \end{cases}$$

Here $c, a \rightarrow$ Constants.

If n is power of 2, $n=2^k$

Form recurrence relation

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

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

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

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

$$2^3T(n/8) + 3cn$$

$$2^4T(n/16) + 4cn$$

$$2^k$$

$$T(1) + kcn + c$$

$n(\log n)$

By representing it by in the form of Asymptotic notation O is

$$T(n) = O(n \log n)$$

QuickSort

Quick Sort is an algorithm based on the DIVIDE-AND-CONQUER paradigm that selects a pivot element and reorders the given list in such a way that all elements smaller to it are on one side and those bigger than it are on the other. Then the sublists are recursively sorted until the list gets completely sorted. The time complexity of this algorithm is $O(n \log n)$.

- Auxiliary space used in the average case for implementing recursive function calls is $O(\log n)$ and hence proved to be a bit space costly, especially when it comes to large data sets.
- Its worst case has a time complexity of $O(n^2)$ which can prove very fatal for large datasets. Competitive sorting algorithms

Quicksort program

```
#include<stdio.h>
#include<conio.h>
int n,j,i;
void main(){
    int i,low,high,z,y;
    int a[10],kk;
    void quick(int a[10],int low,int high);
    int n;
    clrscr();
    printf("\n\t\tmergesort\n");
    printf("\n enter the length of the list:");
    scanf("%d",&n);
    printf("\n enter the list elements");
    for(i=0;i<n;i++){
        scanf("%d",&a[i]);
        low=0;
        high=n-1;
        quick(a,low,high);
        printf("\n sorted array is:");
        for(i=0;i<n;i++){
            printf("%d",a[i]);
            getch();
        }

        int partition(int a[10],int low,int high){
            int i=low,j=high;
            int temp;
            int mid=(low+high)/2;
            int pivot=a[mid];
            while(i<=j)
            {
                while(a[i]<=pivot)
                    i++;
```

```

while(a[j]>pivot)
j--;
if(i<=j){
    temp=a[i];
    a[i]=a[j];
    a[j]=temp;
    i++;
    j--;
}
return j;
}
void quick(int a[10], int low, int high)
{
    int m=partition(a, low, high);
    if(low<m)
        quick(a, low, m);
    if(m+1<high)
        quick(a, m+1, high);
}

```

Algorithm for Quicksort

```

Algorithm quickSort(a, low, high){
    If(high>low) then{
        m=partition(a, low, high);
        if(low<m) then quick(a, low, m);
        if(m+1<high) then quick(a, m+1, high);
    }
}

Algorithm partition(a, low, high){
    i=low, j=high;
    mid=(low+high)/2;
    pivot=a[mid];
    while(i<=j) do{
        while(a[i]<=pivot)
            i++;
        while(a[j]>pivot)
            j--;
        if(i<=j){
            temp=a[i];
            a[i]=a[j];
            a[j]=temp;
            i++;
            j--;
        }
    }
    return j;
}

```

Name	Time Complexity			Space Complexity
	Bestcase	Average Case	Worst Case	
Bubble	$O(n)$	-	$O(n^2)$	$O(n)$
Insertion	$O(n)$	$O(n^2)$	$O(n^2)$	$O(n)$
Selection	$O(n^2)$	$O(n^2)$	$O(n^2)$	$O(n)$

Quick	$O(\log n)$	$O(n \log n)$	$O(n^2)$	$O(n + \log n)$
Merge	$O(n \log n)$	$O(n \log n)$	$O(n \log n)$	$O(2n)$
Heap	$O(n \log n)$	$O(n \log n)$	$O(n \log n)$	$O(n)$

Comparison between Merge and Quick Sort:

- Both follow Divide and Conquer rule.
- Statistically both merge sort and quick sort have the same average case time i.e., $O(n \log n)$.
- Merge Sort Requires additional memory. The pros of merge sort are: it is a stable sort, and there is no worst case (means average case and worst case time complexity is same).
- Quick sort is often implemented in place thus saving the performance and memory by not creating extra storage space.
- But in Quick sort, the performance falls on already sorted / almost sorted list if the pivot is not randomized. Thus why the worst case time is $O(n^2)$.

Randomized Sorting Algorithm: (Random quicksort)

- While sorting the array $a[p:q]$ instead of picking $a[m]$, pick a random element (from among $a[p]$, $a[p+1]$, $a[p+2]$ --- $a[q]$) as the partition elements.
- The resultant randomized algorithm works on any input and runs in an expected $O(n \log n)$ times.

Algorithm for Random Quicksort
<pre> Algorithm RquickSort(a, p, q) { If (high > low) then { If ((q - p) > 5) then Interchange(a, Random() mod (q - p + 1) + p, p); m = partition(a, p, q + 1); quick(a, p, m - 1); quick(a, m + 1, q); } } </pre>

Strassen's Matrix Multiplication:

Let A and B be two $n \times n$ Matrices. The product matrix $C = AB$ is also an $n \times n$ matrix whose i, j^{th} element is formed by taking elements in the i^{th} row of A and j^{th} column of B and multiplying them to get

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

Here $1 \leq i \leq n$ means i and j are in between 1 and n .

To compute $C(i, j)$ using this formula, we need n multiplications.

The divide and conquer strategy suggests another way to compute the product of two $n \times n$ matrices. For simplicity assume n is a power of 2 that is $n = 2^k$ Here $k \rightarrow$ any nonnegative integer.

If n is not a power of two, then enough rows and columns of zeros can be added to both A and B , so that resulting dimensions are a power of two.

Let A and B be $2n \times n$ Matrices. Imagine that A & B are each partitioned into four square sub matrices. Each sub matrix having dimensions $n/2 \times n/2$.

The product of AB can be computed by using previous formula. If AB

is product of 2×2 matrices then

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

$$C_{11} = A_{11}B_{11} + A_{12}B_{21}$$

$$C_{12} = A_{11}B_{12} + A_{12}B_{22}$$

$$C_{21} = A_{21}B_{11} + A_{22}B_{21}$$

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

Here 8 multiplications and 4 additions are performed.

Note that Matrix Multiplication are more Expensive than matrix addition and subtraction.

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

Volkerstrassen has discovered a way to compute the $C_{i,j}$ of above using 7 multiplications and 18 additions or subtractions.

For this first compute 7 $n/2 \times n/2$ matrices P, Q, R, S, T, U & V

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

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

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

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

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

$$U = (A_{21} - A_{11})(B_{11} + B_{12})$$

$$V = (A_{12} - A_{22})(B_{21} + B_{22})$$

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

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

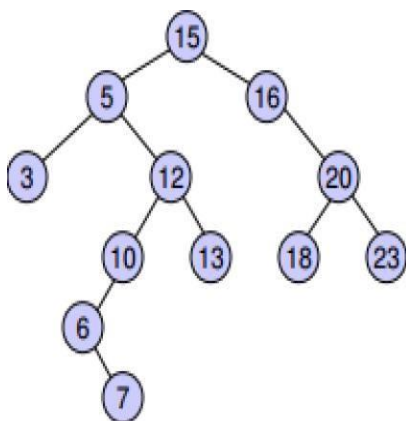
$$C_{21} = Q + S$$

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

$T(n) = \begin{cases} b & \text{if } n \leq 2; \\ 7T(n/2) + cn^2 & \text{if } n > 2 \end{cases}$
--

UNITIII:

Searching and Traversal Techniques: Efficient non - recursive binary tree traversal algorithm, Disjoint set operations, union and find algorithms, Spanning trees, Graph traversals - Breadth first search and Depth first search, AND / OR graphs, game trees, ConnectedComponents,Bi-connectedcomponents.DisjointSets-disjointsetoperations, unionandfindalgorithms,spanningtrees,connected components and biconnected components.



Efficientnonrecursivetreetraversalalgorithms

in-order:(left,root,right)

3,5,6,7,10,12,13

15,16,18,20,23

pre-order:(root,left,right) 15,

5, 3, 12, 10, 6, 7,

13,16,20,18,23

post-order:(left,right,root)

3, 7, 6, 10, 13, 12, 5,

18,23,20,16,15

NonrecursiveInordertraversal algorithm

1. Startfromtheroot.let'sitiscurrent.
2. IfcurrentisnotNULL. pushthenodeontostack.
3. Movetoleftchildofcurrentandgotostep2.
4. IfcurrentisNULL,andstackisnotempty,popnodefromthetack.
5. Printthenodevalueandchangecurrenttorightchildofcurrent.
6. Goto step 2.

So we go on traversing all left node. as we visit the node. we will put that node into stack.rememberneedto visit parent afterthechildandasWewillencounterparentfirstwhen start from root.it'scasefor LIFO:)and hencethe stack). Oncewereach NULLnode. wewill take the node at the top of the stack. last node which we visited. Print it.

Check if there is right child to that node. If yes. move right child to stack and again start traversing left child node and put them on to stack. Once we have traversed all node. ourstack will be empty.

Nonrecursivepostordertraversalalgorithm

Leftnode.rightnode and lastparentnode.

Createanemptystack

DoFollowingwhilerootis not NULL

- a) Pushroot'srightchildandthenroottostack.

b) Set root as root's left child.

Pop an item from stack and set it as root.

a) If the popped item has a right child and the right child is at top of stack, then remove the right child from stack, push the root back and set root as root's right child.

1a) Else print root's data and set root as NULL.

Repeat steps 2.1 and 2.2 while stack is not empty.

Disjoint Sets: If S_i and $S_j, i \neq j$ are two sets, then there is no element that is in both S_i and S_j .

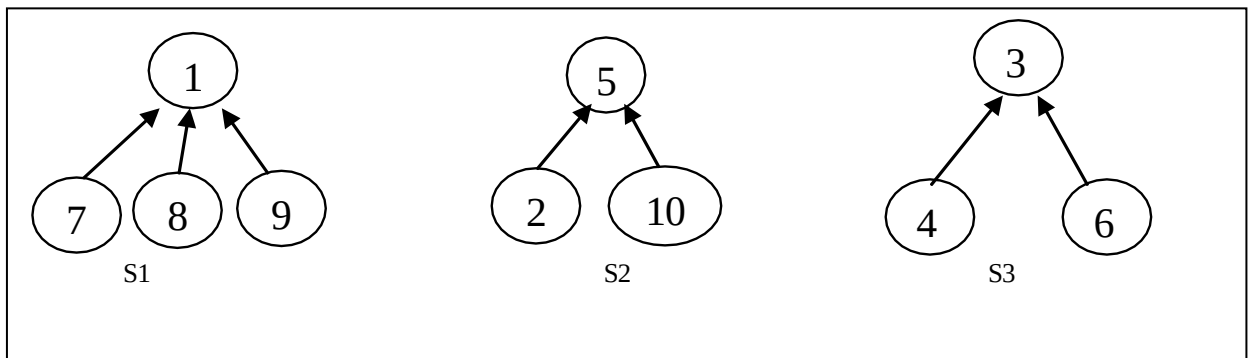
For example: $n=10$ elements can be partitioned into three disjoint sets,

$S_1 = \{1, 7, 8, 9\}$

$S_2 = \{2, 5, 10\}$

$S_3 = \{3, 4, 6\}$

Treerepresentationofsets:

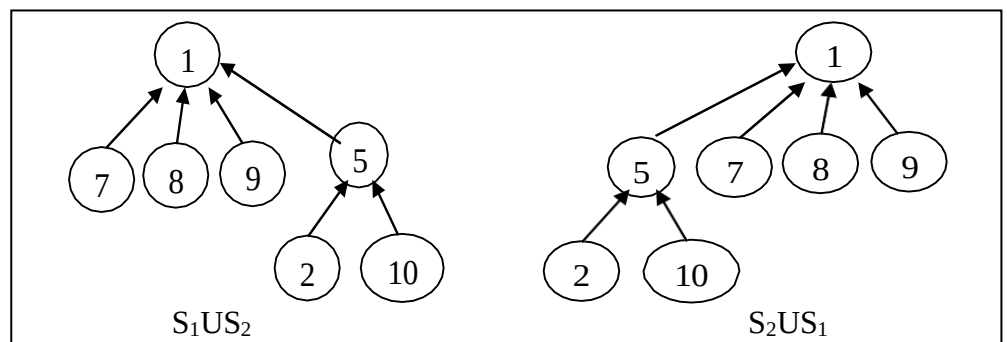


Disjoint set Operations:

- Disjoint set Union
- Find(i)

Disjoint set Union: Means Combination of two disjoint set elements. From above example $S_1 \cup S_2 = \{1, 7, 8, 9, 5, 2, 10\}$

For $S_1 \cup S_2$ treerepresentation, simply make one of the tree is a subtree of the other.



Find: Given element i , find the set containing i .

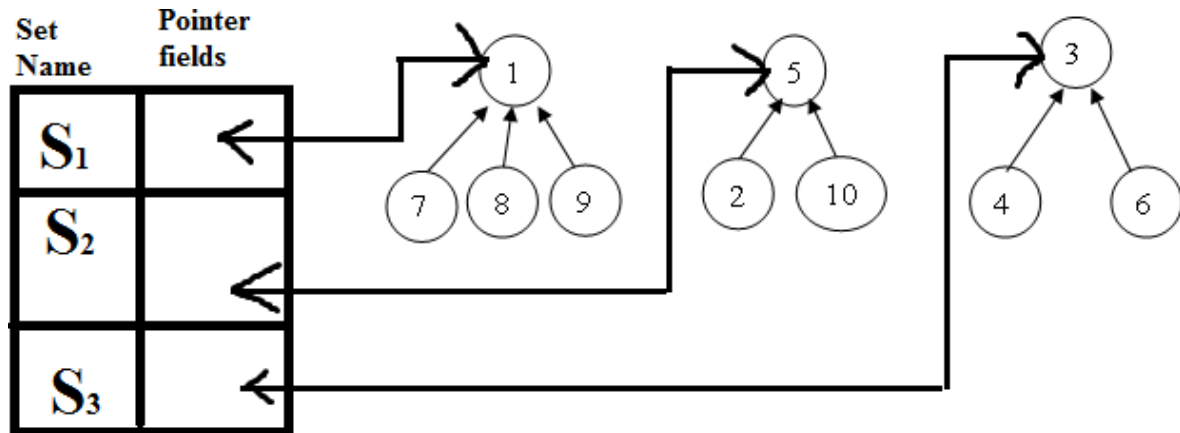
From above example:

$\text{Find}(4) \rightarrow S_3$

Find(1) → S₁
Find(10) → S₂

Data representation of sets:

Tress can be accomplished easily if, with each set name, we keep a pointer to the root of the tree representing that set.



For presenting the union and find algorithms, we ignore the set names and identify sets just by the roots of the trees representing them.

For example: if we determine that element 'i' is in a tree with root 'j' has a pointer to entry 'k' in the set name table, then the set name is just **name[k]**

For unite(adding or combine) to a particular set we use FindPointer function.

Example: If you wish to unite S_i and S_j then we wish to unite the tree with roots FindPointer(S_i) and FindPointer(S_j)

FindPointer → is a function that takes a set name and determines the root of the tree that represents it. For determining operations:

Find(i) → 1st determine the root of the tree and find its pointer to entry in set name table. Union(i, j) → Means union of two trees whose roots are i and j.

If set contains numbers 1 through n, we represent tree node

P[1:n].

n → Maximum number of elements. Each

node represent in array

i	1	2	3	4	5	6	7	8	9	10
P	-1	5	-1	3	-1	3	1	1	1	5

Find(1) by following the indices, starting at 1 and we reach a node with parent value -1.

Example: Find(6) start at 6 and then moves to 6's parent. Since P[3] is negative, we reached the root.

Algorithm for finding Union(i, j):	Algorithm for find(i)
<pre> Algorithm SimpleUnion(i, j) { P[i] := j; // Accomplish the union } </pre>	<pre> Algorithm SimpleFind(i) { While (P[i] ≥ 0) do i := P[i]; return i; } </pre>

If n numbers of roots are there then the above algorithms are not useful for union and find. For union of n trees $\rightarrow$ Union(1,2), Union(2,3), Union(3,4), ..., Union($n-1$, n).
 For Find in n trees $\rightarrow$ Find(1), Find(2), ..., Find(n).

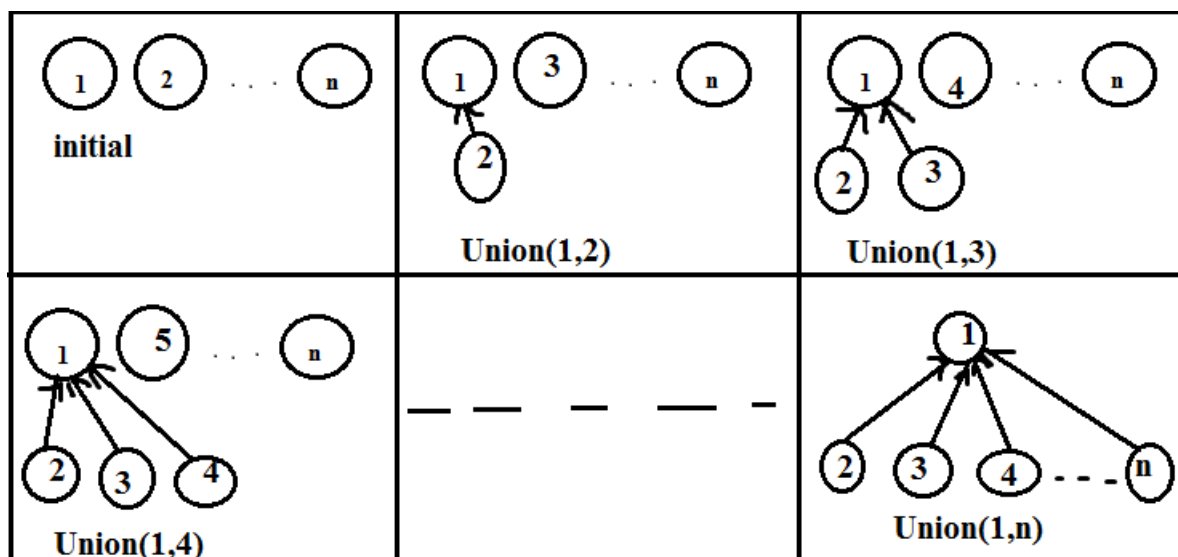
Time taken for the union (simple union) is $\rightarrow O(1)$ (constant).
 For then-1 unions $\rightarrow O(n)$.

Time taken for the find for an element at level i of a tree is $\rightarrow O(i)$.
 For n finds $\rightarrow O(n^2)$.

To improve the performance of our union and find algorithms by avoiding the creation of degenerate trees. For this we use a weighting rule for union(i , j)

Weighting rule for Union(i , j):

If the number of nodes in the tree with root ' i ' is less than the tree with root ' j ', then make ' j ' the parent of ' i '; otherwise make ' i ' the parent of ' j '.



Tree obtained using the weighting rule

AlgorithmforweightedUnion(i, j)

```
AlgorithmWeightedUnion(i,j)
//Unionsetswithrootsiandj, i≠j
//Theweightingrule, p[i]= -count[i] andp[j]= -count[j].
{
temp := p[i]+p[j];
if(p[i]>p[j]) then
{//ihasfewernodes. P[i]:=j;
P[j]:=temp;
}
else
{//jhasfewerorequalnodes. P[j] :=
i;
P[i] := temp;
}
}
```

For implementing the weighting rule, we need to know how many nodes there are in every tree.

For this we maintain a count field in the root of every tree. $i \rightarrow$ root node

$\text{count}[i] \rightarrow$ number of nodes in the tree.

Time required for this above algorithm is $O(1)$ + time for remaining unchanged is determined by using **Lemma**.

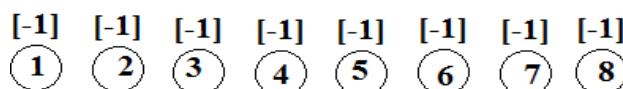
Lemma:- Let T be a tree with **m** nodes created as a result of a sequence of unions each performed using WeightedUnion. The height of T is no greater than $\lceil \log_2 m \rceil + 1$.

Collapsing rule: If 'j' is a node on the path from 'i' to its root and $p[i] \neq \text{root}[i]$, then set $p[j]$ to $\text{root}[i]$.

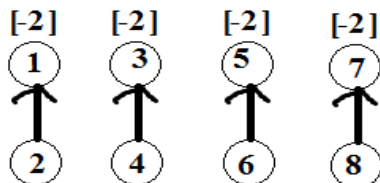
AlgorithmforCollapsingfind.
<pre> AlgorithmCollapsingFind(i) //Findtherootofthetreecontainingelementi. //collapsingruletocollapseallnodesfromitorotheroot. { r:=i; while(p[r]>0) do r := p[r]; //Find the root. While(i≠r)do//Collapse nodes from it to root r. { s:=p[i]; p[i]:=r; i:=s; } returnr; } </pre>

Collapsingfindalgorithmisusedtoperformfindoperationonthetreecreatedby WeightedUnion.

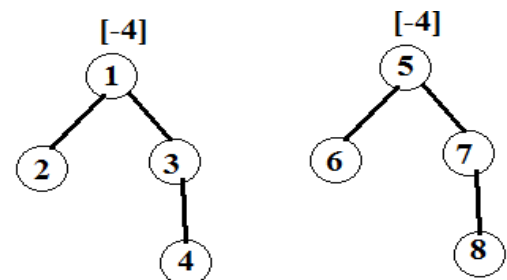
Forexample:TreecreatedbyusingWeightedUnion



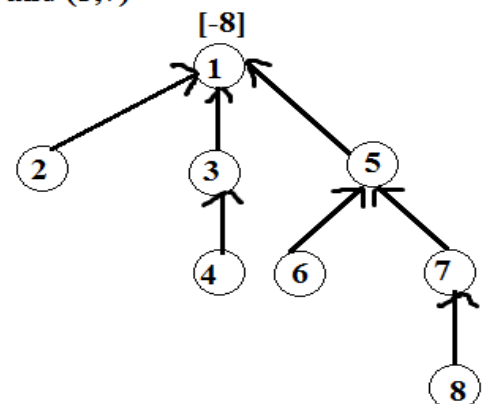
(a) initial height -1 tree



(b) Height -2 trees following Union(1,2),(3,4),(5,6),(7,8)



(c) Height -3 trees following Union (1,3) and (5,7)



(d) Height -4 tree Following Union(1,5)

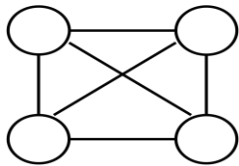
Nowprocessthefollowingeightfinds:Find(8), Find(8),..... Find(8)

If SimpleFind is used, each Find(8) requires going up three parent link fields for a total of 24 moves to process all eight finds.

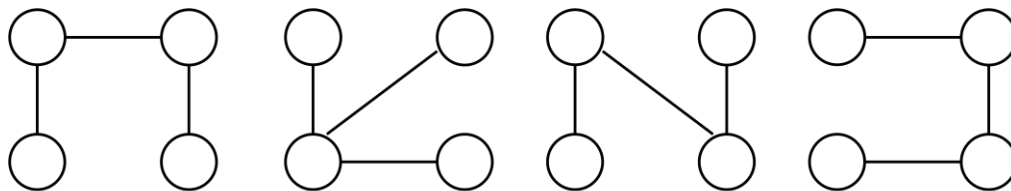
When Collapsing Find is used the first Find(8) requires going up three links and then resetting two links. Total 13 moves required for process all eight finds.

Spanning Tree:-

Let $G=(V,E)$ be an undirected connected graph. A sub graph $t=(V,E^1)$ of G is a spanning tree of G iff t is a tree.



A connected,
undirected graph



Four of the spanning trees of the graph

Spanning Trees have many applications.

Example:-

It can be used to obtain an independent set of circuit equations for an electric network.

Any connected graph with n vertices must have at least $n-1$ edges and all connected graphs with $n-1$ edges are trees. If nodes of G represent cities and the edges represent possible communication links connecting two cities, then the minimum number of links needed to connect the n cities is $n-1$.

There are two basic algorithms for finding minimum-cost spanning trees, and both are greedy algorithms

- Prim's Algorithm
- Kruskal's Algorithm

Prim's Algorithm: Start with any one node in the spanning tree, and repeatedly add the cheapest edge, and the node it leads to, for which the node is not already in the spanning tree.

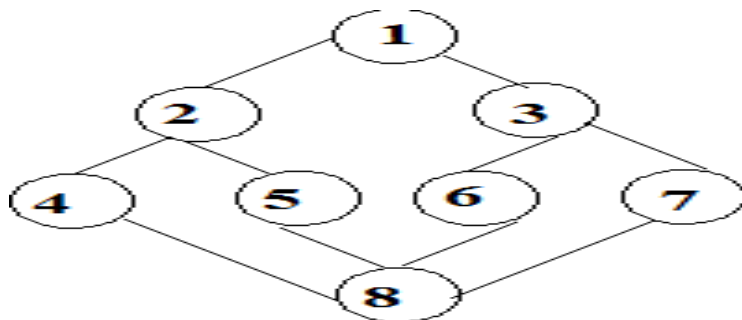
Kruskal's Algorithm: Start with *no* nodes or edges in the spanning tree, and repeatedly add the cheapest edge that does not create a cycle.

Connected Component:

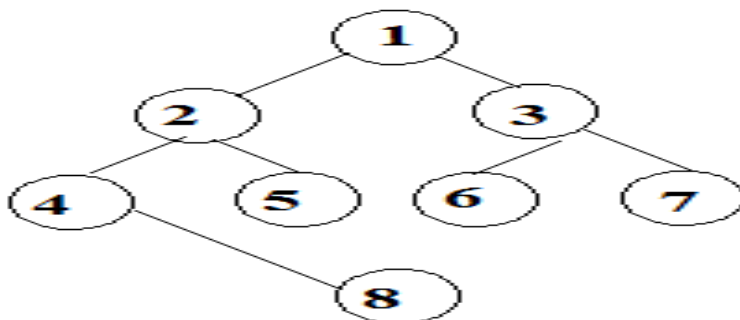
Connected component of a graph can be obtained by using BFST (Breadth first search and traversal) and DFST (Depth first search and traversal). It is also called the spanning tree.

BFST (Breadth first search and traversal):

- In BFS we start at a vertex V mark it as reached (visited).
- The vertex V is at this time said to be unexplored (not yet discovered).
- A vertex is said to be explored (discovered) by visiting all vertices adjacent from it.
- All unvisited vertices adjacent from V are visited next.
- The first vertex on this list is the next to be explored.
- Exploration continues until no unexplored vertex is left.
- These operations can be performed by using Queue.



Undirected Graph G



BFS Spanning tree

This is also called connected graph or spanning tree.

Spanning trees obtained using BFS then it called breadth first spanning trees.

Algorithm for BFS to convert undirected graph G to Connected component or spanning tree.

```

Algorithm BFS(v)
//abfsof G is begin at vertex v
//for any node I, visited[i]=1 if I has already been visited.
//the graph G, and array visited[] are global
{

```



```

U:=v;//qisaqueueofunexploredvertices.
Visited[v]:=1;
Repeat{
ForallverticeswadjacentfromUdo If
(visited[w]=0) then
{
Addwtoq;//wisunexploredVisited[w]:=1;
}
Ifqisemptythenreturn;//Nounexploredvertex. Delete U
from q; //Get 1st unexplored vertex.
} Until(false)
}

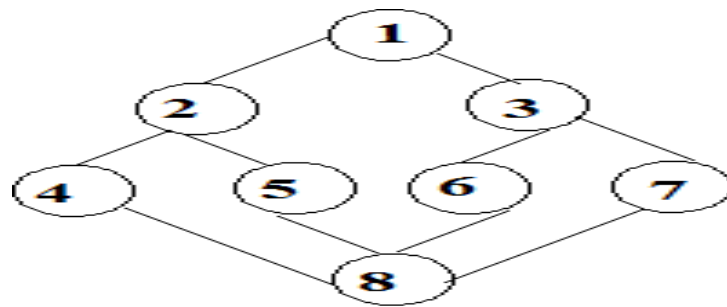
```

MaximumTimecomplexityandspacecomplexityofG(n,e),nodesareinadjacencylist. $T(n, e)=\theta(n+e)$
 $S(n, e)=\theta(n)$

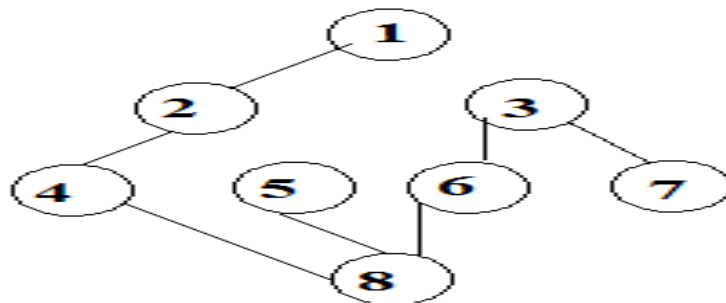
Ifnodesareinadjacencymatrixthen
 $T(n, e)=\theta(n^2)$
 $S(n, e)=\theta(n)$

DFST(Deptfirstsearchandtraversal).:

- Dfsdifferentfrombfs
- Theexplorationofavertexvissuspended(stopped)asoonas anewvertexisreached.
- In this the exploration of the new vertex (example v) begins; this new vertex has been explored, the exploration of v continues.
- Note: exploration start at the new vertex which is not visited in other vertex exploring and choose nearest path for exploring next or adjacent vertex.



Undirected Graph G



DFS(1) Spanning tree

Algorithm for DFS to convert undirected graph G to Connected component spanning tree.

```

Algorithm DFS(v)
//a DFS of G is begun at vertex v
//initially an array visited[] is set to zero.
//this algorithm visits all vertices reachable from v.
//the graph G, and array visited[] are global
{
  Visited[v] := 1;
  For each vertex w adjacent from v do
  {
    If (visited[w] = 0) then DFS(w);
  }
  Add w to Q; // w is unexplored
  Visited[w] := 1;
}

```

Maximum Time complexity and space complexity of $G(n, e)$, nodes are in adjacency list. $T(n, e) = \theta(n + e)$

$S(n, e) = \theta(n)$

If nodes are in adjacency matrix then

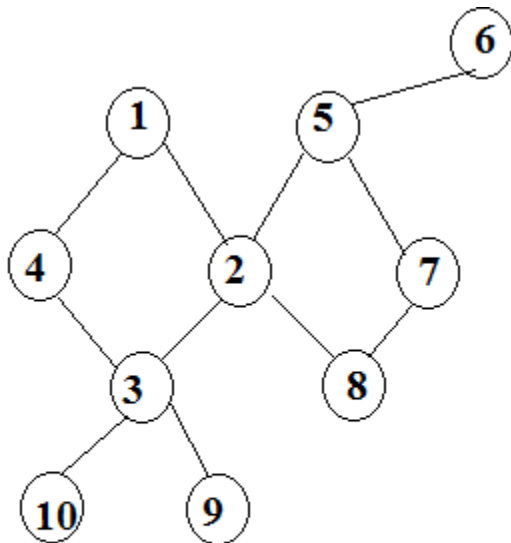
$T(n, e) = \theta(n^2)$

$S(n, e) = \theta(n)$

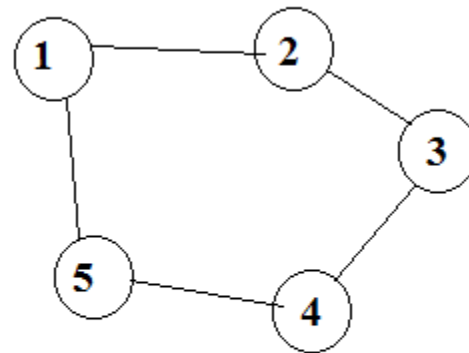
Bi-connected Components:

A graph G is biconnected, iff (if and only if) it contains no articulation point (joint or junction).

A vertex v in a connected graph G is an articulation point, if and only if (iff) the deletion of vertex v together with all edges incident to v disconnects the graph into two or more non empty components.



Graph G1



**Graph Gb
Biconnected Graph**

Not a biconnected graph

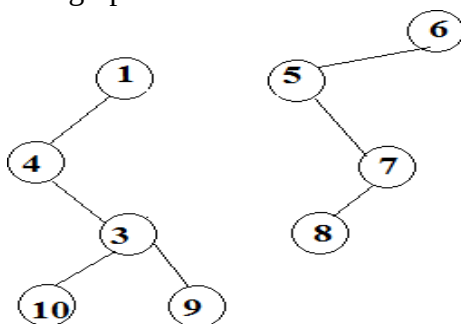
The presence of articulation points in a connected graph can be an undesirable (unwanted) feature in many cases.

For example

if $G1 \rightarrow$ Communication network with
 Vertex $\rightarrow$ communication stations.
 Edges $\rightarrow$ Communication lines.

Then the failure of a communication station that is an articulation point, then we lose the communication in between other stations. F

Form graph $G1$

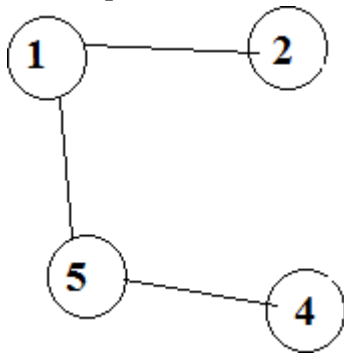


After deleting vertex (2)

(Here 2 is articulation point)

If the graph is bi-connected graph (means no articulation point) then if any station i fails, we can still communicate between every two stations not including station i .

From Graph Gb



There is an efficient algorithm to test whether a connected graph is biconnected. In the case of graphs that are not biconnected, this algorithm will identify all the articulation points. Once it has been determined that a connected graph G is not biconnected, it may be desirable (suitable) to determine a set of edges whose inclusion makes the graph biconnected.

UNIT III:

Greedy method: General method, applications - Job sequencing with deadlines, 0/1 knapsack problem, Minimum cost spanning trees, Single source shortest path problem.

Dynamic Programming: General method, applications - Matrix chain multiplication, Optimal binary search trees, 0/1 knapsack problem, All pairs shortest path problem, Travelling sales person problem, Reliability design.

Greedy Method:

The greedy method is perhaps (maybe or possible) the most straightforward design technique, used to determine a feasible solution that may or may not be optimal.

Feasible solution: - Most problems have inputs and its solution contains a subset of inputs that satisfies a given constraint (condition). Any subset that satisfies the constraint is called feasible solution.

Optimal solution: To find a feasible solution that either maximizes or minimizes a given objective function. A feasible solution that does this is called optimal solution.

The greedy method suggests that an algorithm works in stages, considering one input at a time. At each stage, a decision is made regarding whether a particular input is in an optimal solution.

Greedy algorithms neither postpone nor revise the decisions (ie., no back tracking).

Example: Kruskal's minimal spanning tree. Select an edge from a sorted list, check, decide, and never visit it again.

Application of Greedy Method:

- Job sequencing with deadline
- 0/1 knapsack problem
- Minimum cost spanning trees
- Single source shortest path problem.

Algorithm for Greedy method

```
Algorithm Greedy(a,n)
//a[1:n] contains the inputs.
{
  Solution := 0;
  For i = 1 to n do
  {
    X := select(a);
    If Feasible(solution, x) then
      Solution := Union(solution, x);
  }
  Return solution;
}
```

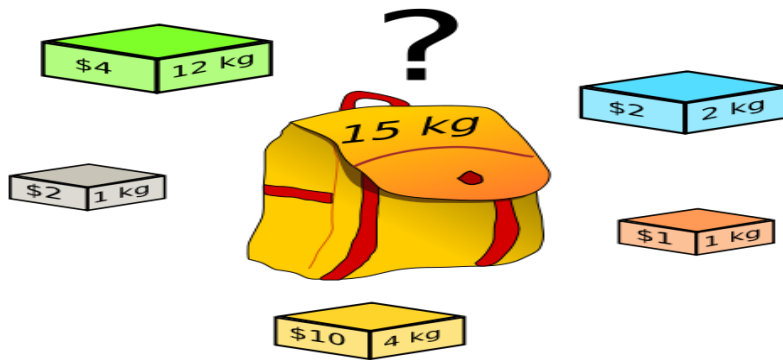
Selection → Function, that selects an input from a[] and removes it. The selected input's value is assigned to x.

Feasible → Boolean-valued function that determines whether x can be included into the solution vector.

Union → function that combines x with solution and updates the objective function.

Knapsack problem

The **knapsack problem** or **rucksack (bag) problem** is a problem in combinatorial optimization: Given a set of items, each with a mass and a value, determine the number of each item to include in a collection so that the total weight is less than or equal to a given limit and the total value is as large as possible



There are two versions of the problems

1. 0/1 knapsack problem
2. Fractional Knapsack problem
 - a. Bounded Knapsack problem.
 - b. Unbounded Knapsack problem.

Solutions to knapsack problems

- **Brute-force approach**: - Solve the problem with a straightforward algorithm
- **Greedy Algorithm**: - Keep taking most valuable items until maximum weight is reached or taking the largest value of each item by calculating $v_i = \text{value}_i / \text{Size}_i$
- **Dynamic Programming**: - Solve each sub problem once and store their solutions in an array.

0/1 knapsack problem:

Let there be n items, z_1 to z_n where z_i has a value v_i and weight w_i . The maximum weight that we can carry in the bag is W . It is common to assume that all values and weights are nonnegative. To simplify the representation, we also assume that the items are listed in increasing order of weight.

$$\text{Maximize } \sum_{i=1}^n v_i x_i \quad \text{subject to } \sum_{i=1}^n w_i x_i \leq W, \quad x_i \in \{0, 1\}$$

Maximize the sum of the values of the items in the knapsack so that the sum of the weights must be less than the knapsack's capacity.

Greedy algorithm for knapsack

Algorithm GreedyKnapsack(m, n)

// $p[1:n]$ and $w[1:n]$ contain the profits and weights respectively

// if the n objects are ordered such that $p[i]/w[i] \geq p[i+1]/w[i+1]$, $m \rightarrow$ size of knapsack and $x[1:n] \rightarrow$ the solution vector

```
{
  For  $i := 1$  to  $n$  do  $x[i] := 0.0$   $U := m$ ;
  For  $i := 1$  to  $n$  do
  {
    if ( $w[i] > U$ ) then break;
     $x[i] := 1.0$ ;
     $U := U - w[i]$ ;
  }
  If ( $i \leq n$ ) then  $x[i] := U/w[i]$ ;
}
```

Ex: - Consider 3 objects whose profits and weights are defined

as $(P_1, P_2, P_3) = (25, 24, 15)$

$(W_1, W_2, W_3) = (18, 15, 10)$

$n = 3 \rightarrow$ number of objects

$m = 20 \rightarrow$ Bag capacity

Consider a knapsack of capacity 20. Determine the optimum strategy for placing the objects into the knapsack. The problem can be solved by the greedy approach where the inputs are arranged according to the selection process (greedy strategy) and solve the problem in stages. The various greedy strategies for the problem could be as follows.

(x_1, x_2, x_3)	$\sum x_i w_i$	$\sum x_i p_i$
(1, 2/15, 0)	$18 \times 1 + \frac{2}{15} \times 15 = 20$	$25 \times 1 + \frac{2}{15} \times 24 = 28.2$
(0, 2/3, 1)	$\frac{2}{3} \times 15 + 10 \times 1 = 20$	$\frac{2}{3} \times 24 + 15 \times 1 = 31$

$(0, 1, \frac{1}{2})$	$1 \times 15 + \frac{1}{2} \times 10 = 20$	$1 \times 24 + \frac{1}{2} \times 15 = 31.5$
$(\frac{1}{2}, \frac{1}{3}, \frac{1}{4})$	$\frac{1}{2} \times 18 + \frac{1}{3} \times 15 + \frac{1}{4} \times 10 = 16.5$	$\frac{1}{2} \times 25 + \frac{1}{3} \times 24 + \frac{1}{4} \times 15 = 12.5 + 8 + 3.75 = 24.25$

Analysis: - If we do not consider the time considered for sorting the input then all of the three greedy strategies complexity will be $O(n)$.

Job Sequence with Deadline:

There is set of n -jobs. For any job i , is a integer deadling $d_i \geq 0$ and profit $P_i > 0$, the profit P_i is earned iff the job completed by its deadline.

To complete a job one had to process the job on a machine for one unit of time. Only one machine is available for processing jobs.

A feasible solution for this problem is a subset J of jobs such that each job in this subset can be completed by its deadline.

The value of a feasible solution J is the sum of the profit of the jobs in J , i.e., $\sum_{i \in J} P_i$

An optimal solution is a feasible solution with maximum value.

The problem involves identification of a subset of jobs which can be completed by its deadline. Therefore the problem suites the subset methodology and can be solved by the greedy method.

Ex:-Obtain the optimal sequence for the following jobs.

$$(P_1, P_2, P_3, P_4) = \begin{matrix} j_1 j_2 & j_3 j_4 \\ (100, 10, 15, 27) \end{matrix}$$

$$\begin{matrix} (d_1, d_2, d_3, d_4) \\ n=4 \end{matrix} = (2, 1, 2, 1)$$

Feasible solution	Processing sequence	Value
$j_1 j_2$ (1,2)	(2,1)	$100+10=110$
(1,3)	(1,3) or (3,1)	$100+15=115$
(1,4)	(4,1)	$100+27=127$
(2,3)	(2,3)	$10+15=25$
(3,4)	(4,3)	$15+27=42$
(1)	(1)	100
(2)	(2)	10
(3)	(3)	15
(4)	(4)	27

In the example solution '3' is the optimal. In this solution only jobs 1&4 are processed and the value is 127. These jobs must be processed in the order j_4 followed by j_1 . the process of job 4 begins at time 0 and ends at time 1. And the processing of job 1 begins at time 1 and ends at time 2. Therefore both the jobs are completed within their deadlines. The optimization measure for determining the next job to be selected in to the solution is according to the profit. The next job to include is that which increases $\sum p_i$ the most, subject to the constraint that the resulting "j" is the feasible solution. Therefore the greedy strategy is to consider the jobs in decreasing order of profits.

The greedy algorithm is used to obtain an optimal solution.

We must formulate an optimization measure to determine how the next job is chosen.

```

algorithm js(d, j, n)
//d → deadline, j → subset of jobs, n → total number of jobs
//d[i] ≥ 1 1 ≤ i ≤ n are the deadlines,
//the jobs are ordered such that p[1] ≥ p[2] ≥ ... ≥ p[n]
//j[i] is the ith job in the optimal solution 1 ≤ i ≤ k, k → subset range
{
d[0] = j[0] = 0;
j[1] = 1;
k = 1;
for i = 2 to n do {
r = k;
while ((d[j[r]] > d[i]) and (d[j[r]] ≠ r)) do
r = r - 1;
if ((d[j[r]] ≤ d[i]) and (d[i] > r)) then
{
for q = k to (r + 1) set p[q - 1] = j[q];
j[r + 1] = i;
k = k + 1;
}
}
return k;
}

```

Note: The size of subset j must be less than or equal to maximum deadline in given list.

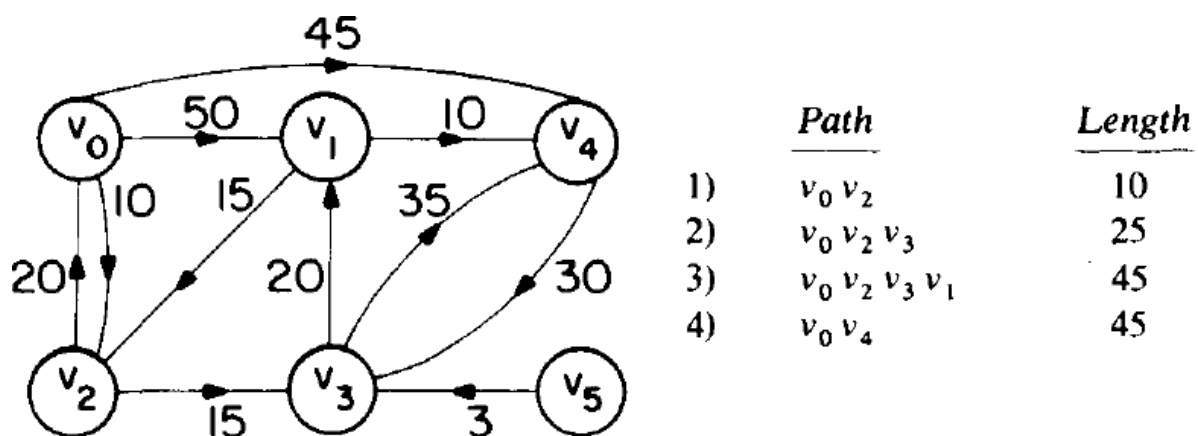
Single Source Shortest Paths:

- Graphs can be used to represent the highway structure of a state or country with vertices representing cities and edges representing sections of highway.
- The edges have assigned weights which may be either the distance between the 2 cities connected by the edge or the average time to drive along that section of highway.
- For example if A motorist wishing to drive from city A to B then we must answer the following questions
 - Is there a path from A to B
 - If there is more than one path from A to B which is the shortest path
- The length of a path is defined to be the sum of the weights of the edges on that path.

Given a directed graph $G(V, E)$ with weight edge $w(u, v)$. we have to find a shortest path

from source vertex $S \in V$ to every other vertex $v \in V - S$.

- To find SSSP for directed graphs $G(V, E)$ there are two different algorithms.
 - Bellman-Ford Algorithm
 - Dijkstra's algorithm
- Bellman-Ford Algorithm: -allow -ve weighted edges in input graph. This algorithm either finds a shortest path from source vertex $S \in V$ to other vertex $v \in V$ or detects a -ve weight cycle in G , hence no solution. If there is no negative weight cycles are reachable from source vertex $S \in V$ to every other vertex $v \in V$
- Dijkstra's algorithm: -allow only +ve weighted edges in the input graph and finds a shortest path from source vertex $S \in V$ to every other vertex $v \in V$.



Graph and shortest paths from v_0 to all destinations

- Consider the above directed graph, if node 1 is the source vertex, then shortest path from 1 to 2 is 1,4,5,2. The length is $10+15+20=45$.
- To formulate a greedy based algorithm to generate the shortest paths, we must conceive of a multistage solution to the problem and also of an optimization measure.
- This is possible by building the shortest paths one by one.
- As an optimization measure we can use the sum of the lengths of all paths so far generated.
- If we have already constructed 'i' shortest paths, then using this optimization measure, the next path to be constructed should be the next shortest minimum length path.
- The greedy way to generate the shortest paths from V_0 to the remaining vertices is to generate these paths in non-decreasing order of path length.
- For this 1st, a shortest path of the nearest vertex is generated. Then a shortest path to the 2nd nearest vertex is generated and so on.

Algorithm for finding Shortest Path

```
AlgorithmShortestPath(v, cost, dist, n)
//dist[j],  $1 \leq j \leq n$ , is set to the length of the shortest path from vertex v to vertex j in graph G with n-vertices.
//dist[v] is zero
{
  for i=1 to n do{
    s[i]=false;
    dist[i]=cost[v,i];
  }
  s[v]=true;
  dist[v]=0.0; // put v in s
  num=2 to n do{
    //determine n-1 paths from v
    choose u from among those vertices not in s such that dist[u] is minimum. s[u]=true; // put u in s
    for (each w adjacent to u with s[w]=false) do if (dist[w] > (dist[u] + cost[u, w])) then
      dist[w] = dist[u] + cost[u, w];
  }
}
```

Minimum Cost Spanning Tree:

SPANNING TREE: -A Subgraph 'n' of graph 'G' is called as a spanning tree if

- (i) It includes all the vertices of 'G'
- (ii) It is a tree

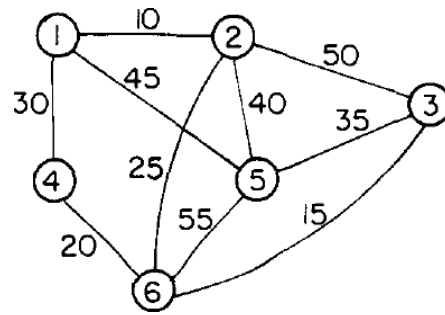
Minimum cost spanning tree: For a given graph 'G' there can be more than one spanning tree. If weights are assigned to the edges of 'G' then the spanning tree which has the minimum cost of edges is called as minimal spanning tree.

The greedy method suggests that a minimum cost spanning tree can be obtained by contacting the tree edge by edge. The next edge to be included in the tree is the edge that results in a minimum increase in the sum of the costs of the edges included so far.

There are two basic algorithms for finding minimum-cost spanning trees, and both are greedy algorithms

- Prim's Algorithm
- Kruskal's Algorithm

Prim's Algorithm: Start with any one node in the spanning tree, and repeatedly add the cheapest edge, and the node it leads to, for which the node is not already in the spanning tree.



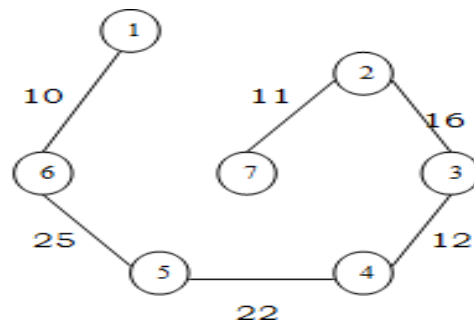
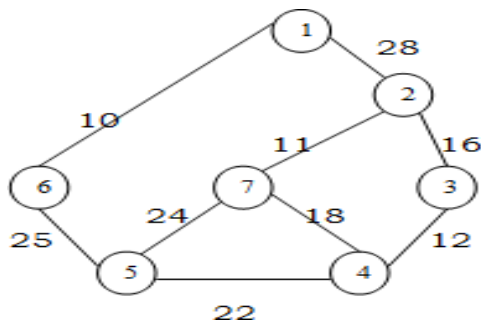
Edge	Cost	Spanning tree
(1,2)	10	
(2,6)	25	
(3,6)	15	
(6,4)	20	
(1,4)	reject	
(3,5)	35	

Stages in Prim's Algorithm

PRIM'S ALGORITHM:-

- Select an edge with minimum cost and include it into the spanning tree.
- Among all the edges which are adjacent with the selected edge, select the one with minimum cost.
- Repeat step 2 until n vertices and $(n-1)$ edges are included. And the sub-graph obtained does not contain any cycles.

Notes: - At every state a decision is made about an edge of minimum cost to be included into the spanning tree. From the edges which are adjacent to the last edge included in the spanning tree i.e. at every stage the sub-graph obtained is a tree.



Prim's minimum spanning tree algorithm

```

Algorithm Prim(E, cost, n, t)
// E is the set of edges in G. Cost(1:n, 1:n) is the
// Cost adjacency matrix of a n vertex graph such that
// Cost(i, j) is either a positive real no. or  $\infty$  if no edge (i, j) exists.
// A minimum spanning tree is computed and
// Stored in the array T(1:n-1, 2).
// (t(i, 1), t(i, 2)) is an edge in the minimum cost spanning tree. The final cost is returned
{
    Let (k, l) be an edge with min cost in E
    Min cost := Cost (x, l);
    T(1, 1) := k; t(1, 2) := l;
for i := 1 to n do // initialize near
    if (cost (i, l) < cost (i, k)) then near (i) := l; else
    near (i) := k;
    near (k) := near (l) := 0;
    for i := 2 to n-1 do
    { // find n-2 additional edges for t
    let j be an index such that near (i)  $\neq$  0 & cost (j, near (i)) is minimum; t (i, 1) := j +
    (i, 2) := near (j);
    min cost := Min cost + cost (j, near (j));
    near (j) := 0;
    for k := 1 to n do // update near[]
    if ((near (k)  $\neq$  0) and (cost {k, near (k)} > cost {k, j}))
    then near Z(k) := j
    }
return min cost;
}

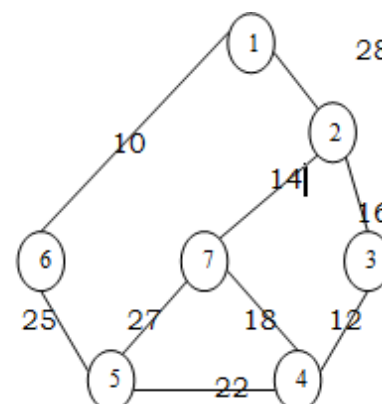
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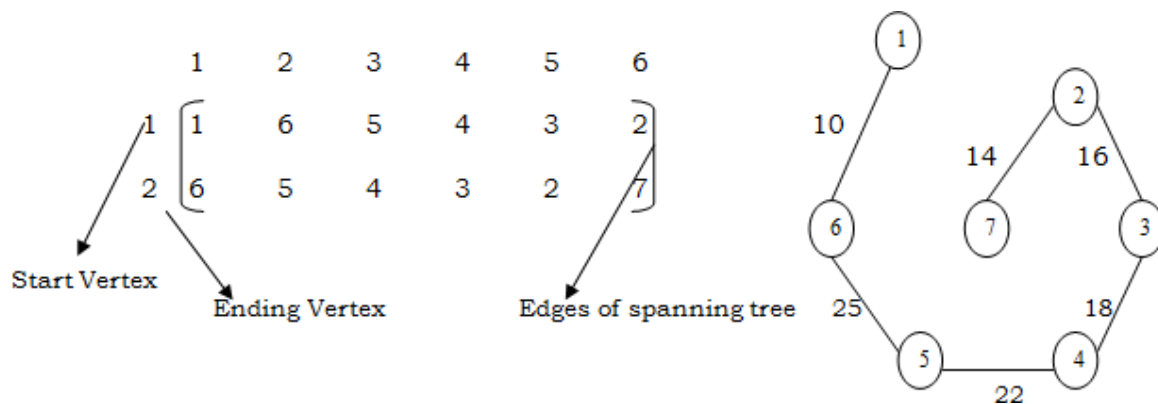
The algorithm takes four arguments E: set of edges, cost is $n \times n$ adjacency matrix cost of (i, j) = +ve integer, if an edge exists between i & j otherwise infinity. 'n' is no. of vertices. 't' is a $(n-1) \times 2$ matrix which consists of the edges of spanning tree.

$E = \{(1, 2), (1, 6), (2, 3), (3, 4), (4, 5), (4, 7), (5, 6), (5, 7), (2, 7)\}$

$n = \{1, 2, 3, 4, 5, 6, 7\}$

Cost	1	2	3	4	5	6	7
1	α	28	α	α	α	10	α
2	28	α	16	α	α	α	14
3	α	10	α	12	α	α	α
4	α	α	12	α	22	α	18
5	α	α	α	22	α	25	24
6	10	α	α	α	25	α	α
7	α	14	α	18	24	α	α





- The algorithm will start with a tree that includes only minimum cost edge of G . The edges are added to this tree one by one.
- The next edge (i, j) to be added is such that i is a vertex which is already included in the tree and j is a vertex not yet included in the tree and cost of i, j is minimum among all edges adjacent to ' i '.
- With each vertex ' j ' next yet included in the tree, we assign a value $\text{near}'j$ '. The value $\text{near}'j$ represents a vertex in the tree such that $\text{cost}(j, \text{near}(j))$ is minimum among all choices for $\text{near}(j)$.
- We define $\text{near}(j) := 0$ for all the vertices ' j ' that are already in the tree.
- The next edge to include is defined by the vertex ' j ' such that $\text{near}(j) \neq 0$ and cost of $(j, \text{near}(j))$ is minimum.

Analysis:-

The time required by the Prim's algorithm is directly proportional to the number of vertices. If a graph ' G ' has ' n ' vertices then the time required by Prim's algorithm is $O(n^2)$.

Kruskal's Algorithm: Start with *no* nodes or edges in the spanning tree, and repeatedly add the cheapest edge that does not create a cycle.

In Kruskal's algorithm for determining the spanning tree we arrange the edges in the increasing order of cost.

- i) All the edges are considered one by one in that order and deleted from the graph and are included into the spanning tree.
- ii) At every stage an edge is included; the sub-graph at a stage need not be a tree. Infact it is a forest.
- iii) At the end if we include 'n' vertices and n-1 edges without forming cycles then we get a single connected component without any cycles i.e. a tree with minimum cost.

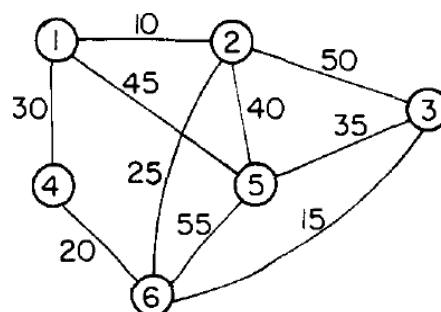
At every stage, as we include an edge in to the spanning tree, we get disconnected trees represented by various sets. While including an edge into the spanning tree we need to check it does not form cycle. Inclusion of an edge (i,j) will form a cycle if i,j both are in same set. Otherwise the edge can be included into the spanning tree.

Kruskal minimum spanning tree algorithm

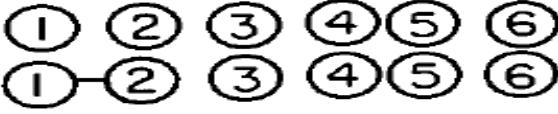
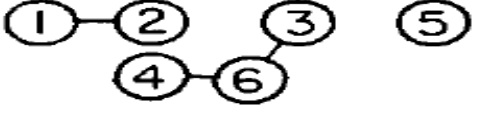
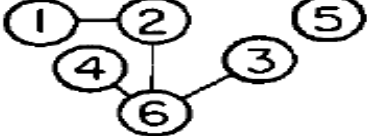
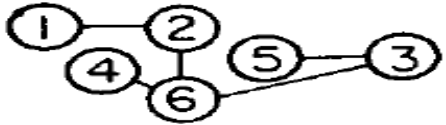
```

Algorithm Kruskal(E, cost, n, t)
// E is the set of edges in G. 'G' has 'n' vertices
// Cost{u,v} is the cost of edge (u,v) t is the set
// of edges in the minimum cost spanning tree
// The final cost is returned
{construct a heap out of the edge costs using heapify;
  for i:=1 to n do parent(i):=-1 // place in different sets
// each vertex is in different set      {1}{1}{3}
  i:=0; min cost:=0.0;
  While(i<n-1) and (heap not empty) do
  {
  Delete a minimum cost edge (u,v) from the heaps; and reheapify using adjust; j:=find
  (u); k:=find (v);
  if (j≠k) then
  {i:=i+1;
    +(i,1)=u; +(i,2)=v;
    min cost:=min cost+cost(u,v);
    Union (j,k);
  }
  }
  if (i≠n-1) then write("No spanning tree"); else
    return min cost;
  }

```



Consider the above graph of, Using Kruskal's method the edges of this graph are considered for inclusion in the minimum cost spanning tree in the order (1,2), (3,6), (4,6), (2,6), (1,4), (3,5), (2,5), (1,5), (2,3), and (5,6). This corresponds to the cost sequence 10, 15, 20, 25, 30, 35, 40, 45, 50, 55. The first four edges are included in T. The next edge to be considered is (1,4). This edge connects two vertices already connected in T and so it is rejected. Next, the edge (3,5) is selected and that completes the spanning tree.

<u>Edge</u>	<u>Cost</u>	<u>Spanning Forest</u>
(1,2)	10	
(3,6)	15	
(4,6)	20	
(2,6)	25	
(1,4)	30	(reject)
(3,5)	35	

Stages in Kruskal's algorithm

Analysis:- If the number of edges in the graph is given by E , then the time for Kruskal's algorithm is given by $O(|E| \log |E|)$.

Dynamic Programming

Dynamic programming is a name, coined by Richard Bellman in 1955. Dynamic programming, as greedy method, is a powerful algorithm design technique that can be used when the solution to the problem may be viewed as the result of a sequence of decisions. In the greedy method we make irrevocable decisions one at a time, using a greedy criterion. However, in dynamic programming we examine the decision sequence to see whether an optimal decision sequence contains optimal decision subsequence.

When optimal decision sequences contain optimal decision subsequences, we can establish recurrence equations, called *dynamic-programming recurrence equations*, that enable us to solve the problem in an efficient way.

Dynamic programming is based on the principle of optimality (also coined by Bellman). The principle of optimality states that no matter whatever the initial state and initial decision are, the remaining decision sequence must constitute an optimal decision sequence with regard to the state resulting from the first decision. The principle implies that an optimal decision sequence is comprised of optimal decision subsequences. Since the principle of optimality may not hold for some formulations of some problems, it is necessary to verify that it does hold for the problem being solved. Dynamic programming cannot be applied when this principle does not hold.

The steps in a dynamic programming solution are:

- Verify that the principle of optimality holds
- Set up the dynamic-programming recurrence equations
- Solve the dynamic-programming recurrence equations for the value of the optimal solution.
- Perform a traceback step in which the solution itself is constructed.

5.1 MULTISTAGE GRAPHS

A multistage graph $G = (V, E)$ is a directed graph in which the vertices are partitioned into $k \geq 2$ disjoint sets $V_i, 1 \leq i \leq k$. In addition, if $\langle u, v \rangle$ is an edge in E , then $u \in V_i$ and $v \in V_{i+1}$ for some $i, 1 \leq i < k$.

Let the vertex 's' be the source, and 't' the sink. Let $c(i, j)$ be the cost of edge $\langle i, j \rangle$. The cost of a path from 's' to 't' is the sum of the costs of the edges on the path. The multistage graph problem is to find a minimum cost path from 's' to 't'. Each set V_i defines a stage in the graph. Because of the constraint on E , every path from 's' to 't' starts in stage 1, goes to stage 2, then to stage 3, then to stage 4, and so on, and eventually terminates in stage k .

A dynamic programming formulation for a k -stage graph problem is obtained by first noticing that every s to t path is the result of a sequence of $k-2$ decisions. The i th

decision involves determining which vertex in $v_{i+1}, 1 \leq i \leq k-2$, is to be on the path. Let $c(i, j)$ be the cost of the path from source to destination. Then using the forward approach, we obtain:

$$\text{cost}(i, j) = \min_{\substack{l \in V_{i+1} \\ \langle j, l \rangle \in E}} \{c(j, l) + \text{cost}(i + 1, l)\}$$

ALGORITHM:

Algorithm Fgraph(G, k, n, p)

// The input is a k-stage graph $G = (V, E)$ with n vertices // indexed in order of stages. E is a set of edges and $c[i, j]$ is the cost of (i, j). p[1 : k] is a minimum cost path.

```
{
    cost[n] := 0.0;
    for j := n - 1 to 1 step - 1 do
    {
        // compute cost[j]
        let r be a vertex such that (j, r) is an edge of G
        and c[j, r] + cost[r] is minimum; cost[j] := c
        [j, r] + cost[r];
        d[j] := r;
    }
    p[1] := 1; p[k] := n;
    for j := 2 to k - 1 do p[j] := d[p[j - 1]];
}
```

The multistage graph problem can also be solved using the backward approach. Let $bp(i, j)$ be a minimum cost path from vertex s to vertex j in V_i . Let $Bcost(i, j)$ be the cost of $bp(i, j)$. From the backward approach we obtain:

$$Bcost(i, j) = \min_{\substack{l \in V_i \\ \langle l, j \rangle \in E}} \{Bcost(i-1, l) + c(l, j)\}$$

Algorithm Bgraph(G, k, n, p)

```
// Same function as Fgraph {
    Bcost[1] := 0.0; for j := 2 to n do { // Compute Bcost
        [j].
        Let r be such that (r, j) is an edge of
        G and Bcost[r] + c[r, j] is minimum;
        Bcost[j] := Bcost[r] + c[r, j];
        D[j] := r;
    } // find a minimum cost path
    p[1] := 1; p[k] := n;
    for j := k - 1 to 2 do p[j] := d[p[j + 1]];
}
```

Complexity Analysis:

The complexity analysis of the algorithm is fairly straightforward. Here, if G has $\sim E$ edges, then the time for the first for loop is $CJ(V \sim \sim E)$.

EXAMPLE 1:

Find the minimum cost path from s to t in the multistage graph of five stages shown below. Do this first using forward approach and then using backward approach.

FORWARD APPROACH:

We use the following equation to find the minimum cost path from s to t : $\text{cost}(i,$

$$j) = \min \{ c(j, l) + \text{cost}(i+1, l) \}$$

$$l \in V_{i+1}$$

$$1 \leq j, l \leq cE$$

$$\begin{aligned} \text{cost}(1, 1) &= \min \{ c(1, 2) + \text{cost}(2, 2), c(1, 3) + \text{cost}(2, 3), c(1, 4) + \text{cost}(2, 4), c(1, 5) + \\ &\quad \text{cost}(2, 5) \} \\ &= \min \{ 9 + \text{cost}(2, 2), 7 + \text{cost}(2, 3), 3 + \text{cost}(2, 4), 2 + \text{cost}(2, 5) \} \end{aligned}$$

Now first starting with,

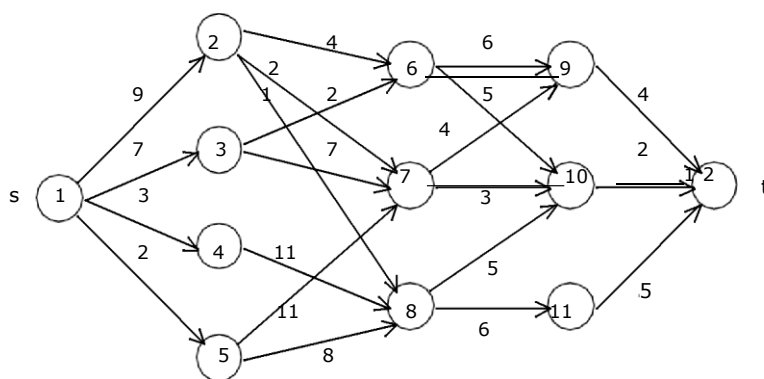
$$\text{cost}(2, 2) = \min \{ c(2, 6) + \text{cost}(3, 6), c(2, 7) + \text{cost}(3, 7), c(2, 8) + \text{cost}(3, 8) \} = \min \{ 4 + \text{cost}(3, 6), 2 + \text{cost}(3, 7), 1 + \text{cost}(3, 8) \}$$

$$\begin{aligned} \text{cost}(3, 6) &= \min \{ c(6, 9) + \text{cost}(4, 9), c(6, 10) + \text{cost}(4, 10) \} \\ &= \min \{ 6 + \text{cost}(4, 9), 5 + \text{cost}(4, 10) \} \end{aligned}$$

$$\text{cost}(4, 9) = \min \{ c(9, 12) + \text{cost}(5, 12) \} = \min \{ 4 + 0 \} = 4 \text{ cost}(4,$$

$$10) = \min \{ c(10, 12) + \text{cost}(5, 12) \} = 2$$

$$\text{Therefore, cost}(3, 6) = \min \{ 6 + 4, 5 + 2 \} = 7$$

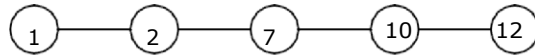


$$\begin{aligned} \text{cost}(3, 7) &= \min \{ c(7, 9) + \text{cost}(4, 9), c(7, 10) + \text{cost}(4, 10) \} \\ &= \min \{ 4 + \text{cost}(4, 9), 3 + \text{cost}(4, 10) \} \end{aligned}$$

$$\text{cost}(4,9)=\min\{c(9,12)+\text{cost}(5,12)\}=\min\{4+0\}=4$$

10)=

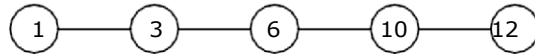
The path is



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or

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(10,

$$2)+\text{cost}(5,12)\}=\min\{2+0\}=2 \text{ Therefore, } \text{cost}(3,7)=\min\{4+4,3$$

$$+2\}=\min\{8,5\}=5$$

$$\begin{aligned} \text{cost}(3,8) &= \min\{c(8,10)+\text{cost}(4,10), c(8,11)+\text{cost}(4,11)\} \\ &= \min\{5+\text{cost}(4,10), 6+\text{cost}(4,11)\} \end{aligned}$$

$$\text{cost}(4,11)=\min\{c(11,12)+\text{cost}(5,12)\}=5$$

$$\text{Therefore, } \text{cost}(3,8)=\min\{5+2, 6+5\}=\min\{7, 11\}=7$$

$$\text{Therefore, } \text{cost}(2,2)=\min\{4+7, 2+5, 1+7\}=\min\{11, 7, 8\}=7$$

$$\begin{aligned} \text{Therefore, } \text{cost}(2,3) &= \min\{c(3,6)+\text{cost}(3,6), c(3,7)+\text{cost}(3,7)\} \\ &= \min\{2+\text{cost}(3,6), 7+\text{cost}(3,7)\} \\ &= \min\{2+7, 7+5\}=\min\{9, 12\}=9 \end{aligned}$$

$$\begin{aligned} \text{cost}(2,4) &= \min\{c(4,8)+\text{cost}(3,8)\}=\min\{11+7\}=18 \\ \text{cost}(2,5) &= \min\{c(5,7)+\text{cost}(3,7), c(5,8)+\text{cost}(3,8)\}=\min\{11+5, 8+7\} \\ &= \min\{16, 15\}=15 \end{aligned}$$

$$\text{Therefore, } \text{cost}(1,1)=\min\{9+7, 7+9, 3+18, 2+15\}=\min\{16, 16, 21, 17\}=16$$

The minimum cost path is 16.

BACKWARD APPROACH:

We use the following equation to find the minimum cost path from i to j : $B_{\text{cost}}(i, j) = \min$

$$\{B_{\text{cost}}(i-1, l) + c(l, j)\}$$

$$\begin{aligned} & l \in V-1 \\ & \langle l, j \rangle \in E \end{aligned}$$

$$\begin{aligned} B_{\text{cost}}(5, 12) &= \min\{B_{\text{cost}}(4, 9) + c(9, 12), B_{\text{cost}}(4, 10) + c(10, 12), \\ & B_{\text{cost}}(4, 11) + c(11, 12)\} \\ &= \min\{B_{\text{cost}}(4, 9) + 4, B_{\text{cost}}(4, 10) + 2, B_{\text{cost}}(4, 11) + 5\} \end{aligned}$$

$$\begin{aligned} B_{\text{cost}}(4, 9) &= \min\{B_{\text{cost}}(3, 6) + c(6, 9), B_{\text{cost}}(3, 7) + c(7, 9)\} \\ &= \min\{B_{\text{cost}}(3, 6) + 6, B_{\text{cost}}(3, 7) + 4\} \end{aligned}$$

$$\begin{aligned} B_{\text{cost}}(3, 6) &= \min\{B_{\text{cost}}(2, 2) + c(2, 6), B_{\text{cost}}(2, 3) + c(3, 6)\} \\ &= \min\{B_{\text{cost}}(2, 2) + 4, B_{\text{cost}}(2, 3) + 2\} \end{aligned}$$

$$\begin{aligned} \text{Bcost}(2,2) &= \min\{\text{Bcost}(1,1)+c(1,2)\} = \min\{0+9\} = 9 \\ \text{Bcost}(2,3) &= \min\{\text{Bcost}(1,1)+c(1,3)\} = \min\{0+7\} = 7 \\ \text{Bcost}(3,6) &= \min\{9+4, 7+2\} = \min\{13,9\} = 9 \end{aligned}$$

$$\text{Bcost}(3,7) = \min\{\text{Bcost}(2,2)+c(2,7), \text{Bcost}(2,3)+c(3,7), \text{Bcost}(2,5)+c(5,7)\}$$

$$\text{Bcost}(2,5) = \min\{\text{Bcost}(1,1)+c(1,5)\} = 2$$

$$\begin{aligned} \text{Bcost}(3,7) &= \min\{9+2, 7+7, 2+11\} = \min\{11,14,13\} = 11 \\ \text{Bcost}(4,9) &= \min\{9+6, 11+4\} = \min\{15,15\} = 15 \end{aligned}$$

$$\text{Bcost}(4,10) = \min\{\text{Bcost}(3,6)+c(6,10), \text{Bcost}(3,7)+c(7,10), \text{Bcost}(3,8)+c(8,10)\}$$

$$\text{Bcost}(3,8) = \min\{\text{Bcost}(2,2)+c(2,8), \text{Bcost}(2,4)+c(4,8), \text{Bcost}(2,5)+c(5,8)\}$$

$$\text{Bcost}(2,4) = \min\{\text{Bcost}(1,1)+c(1,4)\} = 3$$

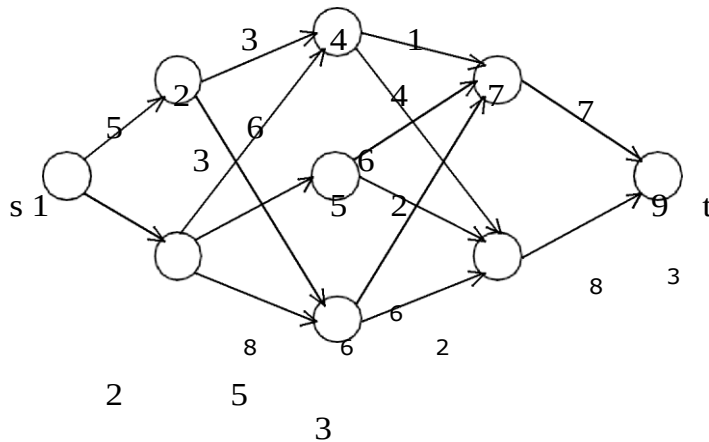
$$\text{Bcost}(3,8) = \min\{9+1, 3+11, 2+8\} = \min\{10,14,10\} = 10$$

$$\begin{aligned} \text{Bcost}(4,10) &= \min\{9+5, 11+3, 10+5\} = \min\{14,14,15\} = 14 \\ \text{Bcost}(4,11) &= \min\{\text{Bcost}(3,8)+c(8,11)\} = \min\{\text{Bcost}(3,8)+6\} = \min\{10+6\} = 16 \end{aligned}$$

$Bcost(5,12)=\min\{15+4,14+2,16+5\}=\min\{19,16,21\}=16$. **EXAMPLE**

2:

Find the minimum cost path from s to t in the multistage graph of five stages shown below. Do this first using forward approach and then using backward approach.



SOLUTION: FORWARD

APPROACH:

$$cost(i,j)=\min\{c(j,l)+cost(i+1,l)\}$$

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$$<J, l>EE$$

$$cost(1,1)=\min\{c(1,2)+cost(2,2),c(1,3)+cost(2,3)\}$$

$$=\min\{5+cost(2,2),2+cost(2,3)\}$$

$$cost(2,2)=\min\{c(2,4)+cost(3,4),c(2,6)+cost(3,6)\}$$

$$=\min\{3+cost(3,4),3+cost(3,6)\}$$

$$cost(3,4)=\min\{c(4,7)+cost(4,7),c(4,8)+cost(4,8)\}$$

$$=\min\{(1+cost(4,7),4+cost(4,8))\}$$

$$cost(4,7)=\min\{c(7,9)+cost(5,9)\}=\min\{7+0\}=7$$

$$cost(4,8)=\min\{c(8,9)+cost(5,9)\}=3$$

$$\text{Therefore, } cost(3,4)=\min\{8,7\}=7$$

$$cost(3,6)=\min\{c(6,7)+cost(4,7),c(6,8)+cost(4,8)\}$$

$$=\min\{6+cost(4,7),2+cost(4,8)\}=\min\{6+7,2+3\}=5$$

$$\text{Therefore, } cost(2,2)=\min\{10,8\}=8$$

$$cost(2,3)=\min\{c(3,4)+cost(3,4),c(3,5)+cost(3,5),c(3,6)+cost(3,6)\}$$

$$cost(3,5)=\min\{c(5,7)+cost(4,7),c(5,8)+cost(4,8)\}=\min\{6+7,2+3\}=5$$

Therefore, $\text{cost}(2,3)=\min\{13,10,13\}=10$

$\text{cost}(1,1)=\min\{5+8,2+10\}=\min\{13,12\}=12$

BACKWARD APPROACH:

$$\text{Bcost}(i,j)=\min_{\substack{1 \leq l \leq j-1 \\ \langle l,j \rangle \in E}} \{ \text{Bcost}(i,l) + c(l,j) \}$$

$$\begin{aligned} \text{Bcost}(5,9) &= \min\{ \text{Bcost}(4,7) + c(7,9), \text{Bcost}(4,8) + c(8,9) \} \\ &= \min\{ \text{Bcost}(4,7) + 7, \text{Bcost}(4,8) + 3 \} \end{aligned}$$

$$\begin{aligned} \text{Bcost}(4,7) &= \min\{ \text{Bcost}(3,4) + c(4,7), \text{Bcost}(3,5) + c(5,7), \\ &\quad \text{Bcost}(3,6) + c(6,7) \} \\ &= \min\{ \text{Bcost}(3,4) + 1, \text{Bcost}(3,5) + 6, \text{Bcost}(3,6) + 6 \} \end{aligned}$$

$$\begin{aligned} \text{Bcost}(3,4) &= \min\{ \text{Bcost}(2,2) + c(2,4), \text{Bcost}(2,3) + c(3,4) \} \\ &= \min\{ \text{Bcost}(2,2) + 3, \text{Bcost}(2,3) + 6 \} \end{aligned}$$

$$\text{Bcost}(2,2) = \min\{ \text{Bcost}(1,1) + c(1,2) \} = \min\{ 0 + 5 \} = 5$$

$$\text{Bcost}(2,3) = \min\{ \text{Bcost}(1,1) + c(1,3) \} = \min\{ 0 + 2 \} = 2$$

$$\text{Therefore, Bcost}(3,4) = \min\{ 5 + 3, 2 + 6 \} = \min\{ 8, 8 \} = 8$$

$$\text{Bcost}(3,5) = \min\{ \text{Bcost}(2,3) + c(3,5) \} = \min\{ 2 + 5 \} = 7$$

$$\text{Bcost}(3,6) = \min\{ \text{Bcost}(2,2) + c(2,6), \text{Bcost}(2,3) + c(3,6) \} = \min\{ 5 + 5, 2 + 8 \} = 10$$

$$\text{Therefore, Bcost}(4,7) = \min\{ 8 + 1, 7 + 6, 10 + 6 \} = 9$$

$$\begin{aligned} \text{Bcost}(4,8) &= \min\{ \text{Bcost}(3,4) + c(4,8), \text{Bcost}(3,5) + c(5,8), \text{Bcost}(3,6) + c(6,8) \} \\ &= \min\{ 8 + 4, 7 + 2, 10 + 2 \} = 9 \end{aligned}$$

$$\text{Therefore, Bcost}(5,9) = \min\{ 9 + 7, 9 + 3 \} = 12$$

pair shortest paths

In the all-pair shortest path problem, we are to find a shortest path between every pair of vertices in a directed graph G . That is, for every pair of vertices (i, j) , we are to find a shortest path from i to j as well as one from j to i . These two paths are the same when G is undirected.

When no edge has a negative length, the all-pairs shortest path problem may be solved by using Dijkstra's greedy single source algorithm n times, once with each of the n vertices as the source vertex.

The all-pair shortest path problem is to determine a matrix A such that $A(i, j)$ is the length of a shortest path from i to j . The matrix A can be obtained by solving n single-source

problems using the algorithm `shortestPaths`. Since each application of this procedure requires $O(n^2)$ time, the matrix A can be obtained in $O(n^3)$ time.

The dynamic programming solution, called Floyd's algorithm, runs in $O(n^3)$ time. Floyd's algorithm works even when the graph has negative length edges (provided there are no negative length cycles).

The shortest i to j path in G , $i \neq j$ originates at vertex i and goes through some intermediate vertices (possibly none) and terminates at vertex j . If k is an intermediate vertex on this shortest path, then the subpaths from i to k and from k to j must be shortest paths from i to k and k to j , respectively. Otherwise, the i to j path is not of minimum length. So, the principle of optimality holds. Let $A^k(i, j)$ represent the length of a shortest path from i to j going through no vertex of index greater than k , we obtain:

$$A^k(i, j) = \min \{ \min_{1 \leq k < n} \{ A^{k-1}(i, k) + A^{k-1}(k, j) \}, c(i, j) \}$$

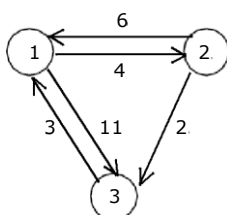
Algorithm AllPaths(Cost, A, n)

```
//cost[1:n, 1:n] is the cost adjacency matrix of a graph which
//n vertices; A[i, j] is the cost of a shortest path from vertex
//i to vertex j. cost[i, i] = 0.0, for 1 ≤ i ≤ n.
{
    for i := 1 to n do
        for j := 1 to n do
            A[i, j] := cost[i, j]; //copy cost into A.
        for k := 1 to n do
            for i := 1 to n do
                for j := 1 to n do
                    A[i, j] := min(A[i, j], A[i, k] + A[k, j]);
}
```

Complexity Analysis: A Dynamic programming algorithm based on this recurrence involves in calculating $n+1$ matrices, each of size $n \times n$. Therefore, the algorithm has a complexity of $O(n^3)$.

Example 1:

Given a weighted digraph $G=(V, E)$ with weight. Determine the length of the shortest path between all pairs of vertices in G . Here we assume that there are no cycles with zero or negative cost.



$$\text{Cost adjacency matrix}(A^0) = \begin{bmatrix} 0 & 6 & 4 \\ 3 & 0 & 2 \\ 11 & 0 & 0 \end{bmatrix}$$

General formula: $\min\{A^{k-1}(i,k) + A^{k-1}(k,j), c(i,j)\}$
 $1 < k < n$

Solve the problem for different values of $k=1, 2$

and **Step 1**: Solving the equation for, $k=1$;

$$\begin{aligned} A_1(1,1) &= \min\{(A^0(1,1) + A^0(1,1)), c(1,1)\} = \min\{0+0, 0\} = 0 \\ A_1(1,2) &= \min\{(A^0(1,1) + A^0(1,2)), c(1,2)\} = \min\{(0+4), 4\} = 4 \\ A_1(1,3) &= \min\{(A^0(1,1) + A^0(1,3)), c(1,3)\} = \min\{(0+11), 11\} = 11 \\ A_1(2,1) &= \min\{(A^0(2,1) + A^0(1,1)), c(2,1)\} = \min\{(6+0), 6\} = 6 \\ A_1(2,2) &= \min\{(A^0(2,1) + A^0(1,2)), c(2,2)\} = \min\{(6+4), 0\} = 0 \\ A_1(2,3) &= \min\{(A^0(2,1) + A^0(1,3)), c(2,3)\} = \min\{(6+11), 2\} = 2 \\ A_1(3,1) &= \min\{(A^0(3,1) + A^0(1,1)), c(3,1)\} = \min\{(3+0), 3\} = 3 \\ A_1(3,2) &= \min\{(A^0(3,1) + A^0(1,2)), c(3,2)\} = \min\{(3+4), 0\} = 0 \\ A_1(3,3) &= \min\{(A^0(3,1) + A^0(1,3)), c(3,3)\} = \min\{(3+11), 0\} = 0 \end{aligned}$$

$$A_{(1)} = \begin{matrix} \sim 0 & 4 & 11 \\ \sim & & \sim \\ \sim 6 & 0 & 2 \\ \sim L3 & 7 & 0 \sim U \end{matrix}$$

Step 2: Solving the equation for, $K=2$;

$$\begin{aligned} A_2(1,1) &= \min\{(A^1(1,2) + A^1(2,1)), c(1,1)\} = \min\{(4+6), 0\} = 0 \\ A_2(1,2) &= \min\{(A^1(1,2) + A^1(2,2)), c(1,2)\} = \min\{(4+0), 4\} = 4 \\ A_2(1,3) &= \min\{(A^1(1,2) + A^1(2,3)), c(1,3)\} = \min\{(4+2), 11\} = 6 \\ A_2(2,1) &= \min\{(A^1(2,2) + A^1(2,1)), c(2,1)\} = \min\{(0+6), 6\} = 6 \\ A_2(2,2) &= \min\{(A^1(2,2) + A^1(2,2)), c(2,2)\} = \min\{(0+0), 0\} = 0 \\ A_2(2,3) &= \min\{(A^1(2,2) + A^1(2,3)), c(2,3)\} = \min\{(0+2), 2\} = 2 \\ A_2(3,1) &= \min\{(A^1(3,2) + A^1(2,1)), c(3,1)\} = \min\{(7+6), 3\} = 3 \\ A_2(3,2) &= \min\{(A^1(3,2) + A^1(2,2)), c(3,2)\} = \min\{(7+0), 7\} = 7 \\ A_2(3,3) &= \min\{(A^1(3,2) + A^1(2,3)), c(3,3)\} = \min\{(7+2), 0\} = 0 \end{aligned}$$

$$A_{(2)} = \begin{matrix} \sim 0 & 4 & 6 \\ \sim & & \sim \\ \sim 6 & 0 & 2 \\ \sim L3 & 7 & 0 \sim \sim \end{matrix}$$

Step3: Solving the equation for, $k=3$;

$$\begin{aligned}
 A_3(1, 1) &= \min\{A^2(1,3) + A^2(3, 1), c(1, 1)\} = \min\{(6+3), 0\} = 0 \\
 A_3(1, 2) &= \min\{A^2(1,3) + A^2(3, 2), c(1, 2)\} = \min\{(6+7), 4\} = 4 \\
 A_3(1, 3) &= \min\{A^2(1,3) + A^2(3, 3), c(1, 3)\} = \min\{(6+0), 6\} = 6 \\
 A_3(2, 1) &= \min\{A^2(2,3) + A^2(3, 1), c(2, 1)\} = \min\{(2+3), 6\} = 5 \\
 A_3(2, 2) &= \min\{A^2(2,3) + A^2(3, 2), c(2, 2)\} = \min\{(2+7), 0\} = 0 \\
 A_3(2, 3) &= \min\{A^2(2,3) + A^2(3, 3), c(2, 3)\} = \min\{(2+0), 2\} = 2 \\
 A_3(3, 1) &= \min\{A^2(3,3) + A^2(3, 1), c(3, 1)\} = \min\{(0+3), 3\} = 3 \\
 A_3(3, 2) &= \min\{A^2(3,3) + A^2(3, 2), c(3, 2)\} = \min\{(0+7), 7\} = 7
 \end{aligned}$$

$$A_3(3,3) = \min\{A^2(3,3) + A^2(3,3), c(3,3)\} = \min\{(0+0), 0\} = 0$$

$$A_{(3)} = \begin{matrix} & \sim 0 & 4 & 6 \sim \\ & & 0 & \sim \\ & & & 2 \sim \\ \sim 5 \sim 3 & & 7 & 0 \sim \end{matrix}$$

TRAVELLINGSALESPERSON PROBLEM

Let $G=(V,E)$ be a directed graph with edge costs C_{ij} . The variable c_{ij} is defined such that $c_{ij} > 0$ for all i and j and $c_{ij} = a$ if $\langle i, j \rangle \in E$. Let $|V| = n$ and assume $n > 1$. A tour of G is a directed simple cycle that includes every vertex in V . The cost of a tour is the sum of the cost of the edges on the tour. The traveling sales person problem is to find a tour of minimum cost. The tour is to be a simple path that starts and ends at vertex 1.

Let $g(i, S)$ be the length of shortest path starting at vertex i , going through all vertices in S , and terminating at vertex 1. The function $g(1, V - \{1\})$ is the length of an optimal salesperson tour. From the principle of optimality it follows that:

$$g(1, V - \{1\}) = \min_{k \in V - \{1\}} \{c_{1k} + g(k, V - \{1, k\})\} \quad (1)$$

min

Generalizing equation 1, we obtain (for $i \in S$) (2)

$$g(i, S) = \min_{j \in S} \{c_{ij} + g(j, S - \{i, j\})\}$$

The Equation can be solved for $g(1, V - \{1\})$ if we know $g(k, V - \{1, k\})$ for all choices of k .

Complexity Analysis:

For each value of $|S|$ there are $n-1$ choices for i . The number of distinct sets S of

size k not including 1 and is $\binom{n-1}{k}$.

Hence, the total number of $g(i, S)$'s to be computed before computing $g(1, V - \{1\})$ is:

$$\sum_{k=0}^{n-2} \binom{n-1}{k} \approx \sum_{k=0}^{n-1} \binom{n-1}{k} = 2^{n-1}$$

To calculate this sum, we use the binomial theorem:

$$\sum_{i=0}^{n-1} \binom{n-1}{i} = \sum_{i=0}^{n-1} \frac{(n-1)!}{i!(n-1-i)!} = \frac{(n-1)!}{(n-1)!} \sum_{i=0}^{n-1} \frac{(n-1)!}{i!(n-1-i)!} = \sum_{i=0}^{n-1} \frac{(n-1)!}{i!(n-1-i)!} \approx 2^{n-1}$$

According to the binomial theorem:

$$\sum_{i=0}^{n-1} \binom{n-1}{i} = \sum_{i=0}^{n-1} \frac{(n-1)!}{i!(n-1-i)!} = \frac{(n-1)!}{(n-1)!} \sum_{i=0}^{n-1} \frac{(n-1)!}{i!(n-1-i)!} = \sum_{i=0}^{n-1} \frac{(n-1)!}{i!(n-1-i)!} \approx 2^{n-1}$$

Therefore,

$$\sum_{k=0}^{n-2} \binom{n-1}{k} \approx 2^{n-1}$$

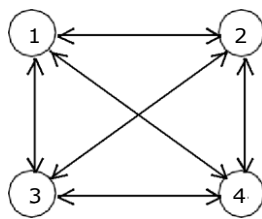
This is $\Phi(n^{2^{n-2}})$, so there are exponential number of calculate. Calculating one $g(i, S)$ require finding the minimum of at most n quantities. Therefore, the entire algorithm is $\Phi(n^{2^{n-2}})$. This is better than enumerating all $n!$ different tours to find the best one.

So, we have traded on exponential growth for a much smaller exponential growth.

The most serious drawback of this dynamic programming solution is the space needed, which is $O(n^2)$. This is too large even for modest values of n .

Example 1:

For the following graph find minimum cost tour for the traveling salesperson problem:



The cost adjacency matrix =

1	2	3	4
10	15	20	5
0	9	12	8
13	0	0	1
8	9	0	1

Let us start the tour from vertex 1:

$$g(1, V - \{1\}) = \min_{2 \leq k \leq n} \{c_{1k} + g(k, V - \{1, k\})\} \quad - \quad (1)$$

More generally writing:

$$g(i, s) = \min \{c_{ij} + g(j, s - \{j\})\} \quad - \quad (2)$$

Clearly, $g(i, T) = c_{i1}$, $1 \leq i \leq n$. So,

$$g(2, T) = c_{21} = 5$$

$$g(3, T) = c_{31} = 6$$

$$g(4, T) = c_{41} = 8$$

Using equation (2) we obtain:

$$g(1, \{2, 3, 4\}) = \min \{c_{12} + g(2, \{3, 4\}), c_{13} + g(3, \{2, 4\}), c_{14} + g(4, \{2, 3\})\}$$

$$g(2, \{3, 4\}) = \min \{c_{23} + g(3, \{4\}), c_{24} + g(4, \{3\})\} \\ = \min \{9 + g(3, \{4\}), 10 + g(4, \{3\})\}$$

$$g(3, \{4\}) = \min \{c_{34} + g(4, T)\} = 12 + 8 = 20$$

$$g(4, \{3\}) = \min \{c_{43} + g(3, T)\} = 9 + 6 = 15$$

Therefore, $g(2, \{3, 4\}) = \min\{9+20, 10+15\} = \min\{29, 25\} = 25$

$g(3, \{2, 4\}) = \min\{(c_{32} + g(2, \{4\})), (c_{34} + g(4, \{2\}))\}$

$g(2, \{4\}) = \min\{c_{24} + g(4, T)\} = 10 + 8 = 18$

$g(4, \{2\}) = \min\{c_{42} + g(2, \sim)\} = 8 + 5 = 13$

Therefore, $g(3, \{2, 4\}) = \min\{13+18, 12+13\} = \min\{41, 25\} = 25$

$g(4, \{2, 3\}) = \min\{c_{42} + g(2, \{3\}), c_{43} + g(3, \{2\})\}$

$g(2, \{3\}) = \min\{c_{23} + g(3, \sim)\} = 9 + 6 = 15$

$g(3, \{2\}) = \min\{c_{32} + g(2, T)\} = 13 + 5 = 18$

Therefore, $g(4, \{2, 3\}) = \min\{8+15, 9+18\} = \min\{23, 27\} = 23$

$g(1, \{2, 3, 4\}) = \min\{c_{12} + g(2, \{3, 4\}), c_{13} + g(3, \{2, 4\}), c_{14} + g(4, \{2, 3\})\} = \min\{10+25, 15+25, 20+23\} = \min\{35, 40, 43\} = 35$

The optimal tour for the graph has length = 35

The optimal tour is: 1, 2, 4, 3, 1.

OPTIMAL BINARY SEARCH TREE

Let us assume that the given set of identifiers is $\{a_1, \dots, a_n\}$ with $a_1 < a_2 < \dots < a_n$.

Let $p(i)$ be the probability with which we search for a_i . Let $q(i)$ be the probability that the identifier x being searched for is such that $a_i < x < a_{i+1}$, $0 \leq i \leq n$ (assume $a_0 = -\infty$ and $a_{n+1} = +\infty$).

We have to arrange the identifiers in a binary search tree in a way that minimizes the expected total access time.

In a binary search tree, the number of comparisons needed to access an element at depth d_i is $d_i + 1$, so if a_i is placed at depth d_i , then we want to minimize:

$$\sum_{i=1}^n p_i(1 + d_i).$$

Let $P(i)$ be the probability with which we shall be searching for a_i . Let $Q(i)$ be the probability of an unsuccessful search. Every internal node represents a point where a successful search may terminate. Every external node represents a point where an unsuccessful search may terminate.

The expected cost contribution for the internal node for a_i is:

$$P(i) * \text{level}(a_i).$$

Unsuccessful search terminates with

$= 0$ (if it is an external node). Hence the cost contribution for this node is:

$$Q(i) * \text{level}((E_i) - 1)$$

The expected cost of binary search tree is:

$$\sim \sum_{i=1}^n P(i) \cdot \text{level}(a_i) + \sum_{i=1}^n Q(i) \cdot \text{level}((E_i)-1)$$

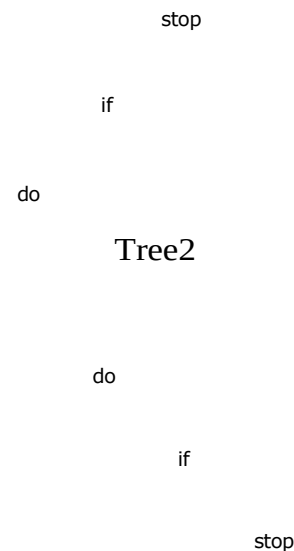
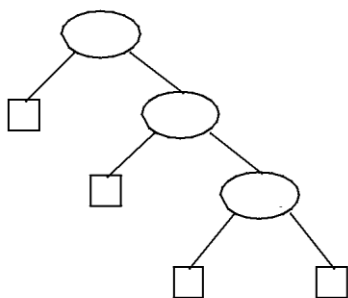
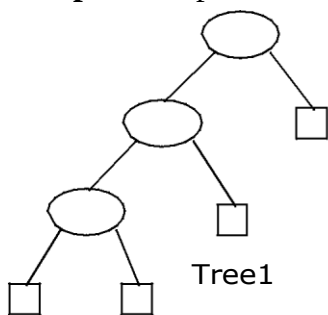
Given a fixed set of identifiers, we wish to create a binary search tree organization. We may expect different binary search trees for the same identifier set to have different performance characteristics.

The computation of each of these $c(i, j)$'s requires us to find the minimum of m quantities. Hence, each such $c(i, j)$ can be computed in time $O(m)$. The total time for all $c(i, j)$'s with $j-i=m$ is therefore $O(nm-m^2)$.

The total time to evaluate all the $c(i, j)$'s and $r(i, j)$'s is therefore:

$$\sim (nm - m^2) = O(n^3) \\ 1 < m < n$$

Example 1: The possible binary search trees for the identifier set $(a_1, a_2, a_3) = (\text{do}, \text{if}, \text{stop})$ are as follows. Given the equal probabilities $p(i) = Q(i) = 1/7$ for all i , we have:



$$\text{Cost}(\text{tree\#1}) = \left(\frac{1 \times 1}{7} + \frac{1 \times 2}{7} + \frac{1 \times 3}{7} \right) + \left(\frac{1 \times 1}{7} \right)$$

$$1 + 2 + 3 + 1 + 2 + 3 + 6 + 9 = 15$$

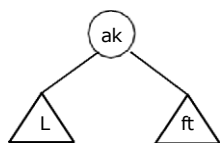
$$\begin{aligned} \text{Cost}(\text{tree\#3}) &= \frac{1}{7}x_1 + \frac{1}{7}x_2 + \frac{1}{7}x_3 + \frac{1}{7}x_4 + \frac{1}{7}x_5 + \frac{1}{7}x_6 + \frac{1}{7}x_7 + \frac{1}{7}x_8 + \frac{1}{7}x_9 + \frac{1}{7}x_{10} + \frac{1}{7}x_{11} + \frac{1}{7}x_{12} + \frac{1}{7}x_{13} + \frac{1}{7}x_{14} + \frac{1}{7}x_{15} \\ &= \frac{1+2+3}{7} + \frac{1+2+3+3}{7} + \frac{6+9}{7} + \frac{15}{7} \end{aligned}$$

$$\begin{aligned} \text{Cost}(\text{tree\#4}) &= \frac{1}{7}x_1 + \frac{1}{7}x_2 + \frac{1}{7}x_3 + \frac{1}{7}x_4 + \frac{1}{7}x_5 + \frac{1}{7}x_6 + \frac{1}{7}x_7 + \frac{1}{7}x_8 + \frac{1}{7}x_9 + \frac{1}{7}x_{10} + \frac{1}{7}x_{11} + \frac{1}{7}x_{12} + \frac{1}{7}x_{13} + \frac{1}{7}x_{14} + \frac{1}{7}x_{15} \\ &= \frac{1+2+3}{7} + \frac{1+2+3+3}{7} + \frac{6+9}{7} + \frac{15}{7} \end{aligned}$$

$$\begin{aligned} \text{Cost}(\text{tree\#2}) &= \frac{1}{7}x_1 + \frac{1}{7}x_2 + \frac{1}{7}x_3 + \frac{1}{7}x_4 + \frac{1}{7}x_5 + \frac{1}{7}x_6 + \frac{1}{7}x_7 + \frac{1}{7}x_8 + \frac{1}{7}x_9 + \frac{1}{7}x_{10} + \frac{1}{7}x_{11} + \frac{1}{7}x_{12} + \frac{1}{7}x_{13} + \frac{1}{7}x_{14} + \frac{1}{7}x_{15} \\ &= \frac{1+2+2}{7} + \frac{2+2+2+2}{7} + \frac{5+8}{7} + \frac{13}{7} \end{aligned}$$

Huffman coding tree solved by a greedy algorithm has a limitation of having the data only at the leaves and it must not preserve the property that all nodes to the left of the root have keys, which are less etc. Construction of an optimal binary search tree is harder, because the data is not constrained to appear only at the leaves, and also because the tree must satisfy the binary search tree property and it must preserve the property that all nodes to the left of the root have keys, which are less.

A dynamic programming solution to the problem of obtaining an optimal binary search tree can be viewed by constructing a tree as a result of sequence of decisions by holding the principle of optimality. A possible approach to this is to make a decision as to which of the a_i 's be arranged to the root node at 'T'. If we choose ' a_k ' then it is clear that the internal nodes for $a_1, a_2, \dots, a_{k-1}$ as well as the external nodes for the classes $E_0, E_1, \dots, E_{k-1}$ will lie in the left subtree, L, of the root. The remaining nodes will be in the right subtree, ft. The structure of an optimal binary search tree is:



$$\text{Cost}(L) = \sum_{i=1}^K P(i) * \text{level}(a_i) + \sum_{i=0}^K Q(i) * \text{level}(E_i) - 1$$

$$\text{Cost}(ft) = \sum_{i=K}^n P(i) * \text{level}(a_i) + \sum_{i=K}^n Q(i) * \text{level}(E_i) - 1$$

The $C(i, J)$ can be computed as:

$$C(i, J) = \min_{i < k < J} \{C(i, k-1) + C(k, J) + P(K) + w(i, K-1) + w(K, J)\}$$

$$= \min_{i < k < J} \{C(i, K-1) + C(K, J)\} + w(i, J) \quad \text{--} \quad (1)$$

$$\text{Where } W(i, J) = P(J) + Q(J) + w(i, J-1) \quad \text{--} \quad (2)$$

Initially $C(i, i) = 0$ and $w(i, i) = Q(i)$ for $0 \leq i \leq n$.

Equation (1) maybe solved for $C(0, n)$ by first computing all $C(i, J)$ such that $J - i = 1$. Next, we can compute all $C(i, J)$ such that $J - i = 2$, Then all $C(i, J)$ with $J - i = 3$ and soon.

$C(i, J)$ is the cost of the optimal binary search tree 'Tij' during computation we record the root $R(i, J)$ of each tree 'Tij'. Then an optimal binary search tree may be constructed from these $R(i, J)$. $R(i, J)$ is the value of 'K' that minimizes equation (1).

We solve the problem by knowing $W(i, i+1), C(i, i+1)$ and $R(i, i+1), 0 \leq i < 4$;

Knowing $W(i, i+2), C(i, i+2)$ and $R(i, i+2), 0 \leq i < 3$ and repeating until $W(0, n), C(0, n)$ and $R(0, n)$ are obtained.

The results are tabulated to recover the actual tree.

Example 1:

Let $n=4$, and $(a_1, a_2, a_3, a_4) = (\text{do, if, need, while})$. Let $P(1:4) = (3, 3, 1, 1)$ and $Q(0:4) = (2, 3, 1, 1, 1)$

Solution:

Table for recording $W(i, j), C(i, j)$ and $R(i, j)$:

Column Row	0	1	2	3	4
0	2, 0, 0	3, 0, 0	1, 0, 0	1, 0, 0,	1, 0, 0
1	8, 8, 1	7, 7, 2	3, 3, 3	3, 3, 4	
2	12, 19, 1	9, 12, 2	5, 8, 3		
3	14, 25, 2	11, 19, 2			
4	16, 32, 2				

This computation is carried out row-wise from row 0 to row 4. Initially, $W(i, i) = Q(i)$ and $C(i, i) = 0$ and $R(i, i) = 0, 0 \leq i < 4$. Solving for

$C(0, n)$:

First, computing all $C(i, j)$ such that $j-i=1$; $j=i+1$ and as $0 \leq i < 4$; $i=0, 1, 2$ and 3 ; $i < k \leq J$. Start with $i = 0$; so $j = 1$; as $i < k \leq j$, so the possible value for $k = 1$

$$W(0, 1) = P(1) + Q(1) + W(0, 0) = 3 + 3 + 2 = 8$$

$$C(0, 1) = W(0, 1) + \min\{C(0, 0) + C(1, 1)\} = 8$$

$$R(0, 1) = 1 \text{ (value of 'K' that is minimum in the above equation).}$$

Next with $i=1$; so $j=2$; as $i < k \leq j$, so the possible value for $k=2$

$$W(1, 2) = P(2) + Q(2) + W(1, 1) = 3 + 1 + 3 = 7$$

$$C(1, 2) = W(1, 2) + \min\{C(1, 1) + C(2, 2)\} = 7$$

$$R(1, 2) = 2$$

Next with $i=2$; so $j=3$; as $i < k \leq j$, so the possible value for $k=3$

$$W(2, 3) = P(3) + Q(3) + W(2, 2) = 1 + 1 + 1 = 3$$

$$C(2, 3) = W(2, 3) + \min\{C(2, 2) + C(3, 3)\} = 3 + [(0 + 0)] = 3$$

$$ft(2, 3) = 3$$

Next with $i=3$; so $j=4$; as $i < k \leq j$, so the possible value for $k=4$

$$W(3, 4) = P(4) + Q(4) + W(3, 3) = 1 + 1 + 1 = 3$$

$$C(3, 4) = W(3, 4) + \min\{[C(3, 3) + C(4, 4)]\} = 3 + [(0 + 0)] = 3$$

$$ft(3, 4) = 4$$

Second, Computing all $C(i, j)$ such that $j-i=2$; $j=i+2$ and as $0 \leq i < 3$; $i=0, 1, 2$; $i < k \leq J$. Start with $i=0$; so $j=2$; as $i < k \leq J$, so the possible values for $k=1$ and 2 .

$$W(0, 2) = P(2) + Q(2) + W(0, 1) = 3 + 1 + 8 = 12$$

$$C(0, 2) = W(0, 2) + \min\{(C(0, 0) + C(1, 2)), (C(0, 1) + C(2, 2))\} = 12$$

$$+ \min\{(0 + 7, 8 + 0)\} = 19$$

$$ft(0, 2) = 1$$

Next, with $i=1$; so $j=3$; as $i < k \leq j$, so the possible value for $k=2$ and 3 .

$$W(1, 3) = P(3) + Q(3) + W(1, 2) = 1 + 1 + 7 = 9$$

$$C(1, 3) = W(1, 3) + \min\{[C(1, 1) + C(2, 3)], [C(1, 3) + C(2, 2)]\}$$

$$= W(1, 3) + \min\{(0 + 3), (7 + 0)\} = 9 + 3 = 12$$

$$ft(1, 3) = 2$$

Next, with $i=2$; so $j=4$; as $i < k \leq j$, so the possible value for $k=3$ and 4 .

$$W(2, 4) = P(4) + Q(4) + W(2, 3) = 1 + 1 + 3 = 5$$

$$C(2, 4) = W(2, 4) + \min\{[C(2, 2) + C(3, 4)], [C(2, 3) + C(4, 4)]\}$$

$$= 5 + \min\{(0 + 3), (3 + 0)\} = 5 + 3 = 8$$

$$ft(2, 4) = 3$$

Third, Computing all $C(i, j)$ such that $J-i=3$; $j=i+3$ and as $0 \leq i < 2$; $i=0, 1$; $i < k \leq J$. Start with $i=0$; so $j=3$; as $i < k \leq j$, so the possible values for $k=1, 2$ and 3 .

$$W(0, 3) = P(3) + Q(3) + W(0, 2) = 1 + 1 + 12 = 14$$

$$C(0, 3) = W(0, 3) + \min\{[C(0, 0) + C(1, 3)], [C(0, 1) + C(2, 3)], [C(0, 2) + C(3, 3)]\}$$

$$+ 11 = 25$$

$$ft(0, 3) = 14 + \min\{(0 + 12), (8 + 3), (19 + 0)\} = 14$$

Start with $i=1$; $soj=4$; as $i < k \leq j$, so the possible values for $k=2, 3$ and 4 .

$$\begin{aligned}
 W(1,4) &= P(4) + Q(4) + W(1,3) = 1 + 1 + 9 = 11 = W \\
 C(1,4) &= (1,4) + \min\{[C(1,1) + C(2,4)], [C(1,3) + C(4,4)]\} \\
 &= 11 + \min\{(0+8), (7+3), (12+0)\} = 11 + 2 = 13 \\
 ft(1,4) &= 11 + 2 = 13
 \end{aligned}$$

Fourth, Computing all $C(i,j)$ such that $j-i=4$; $j=i+4$ and as $0 \leq i < 1$; $i=0$; $i < k \leq J$.

Start with $i=0$; so $j=4$; as $i < k \leq j$, so the possible values for $k=1, 2, 3$ and 4 .

$$\begin{aligned}
 W(0, 4) &= P(4) + Q(4) + W(0, 3) = 1 + 1 + 14 = 16 \\
 C(0, 4) &= W(0, 4) + \min\{[C(0, 0) + C(1, 4)], [C(0, 1) + C(2, 4)], \\
 &\quad [C(0, 2) + C(3, 4)], [C(0, 3) + C(4, 4)]\} \\
 &= 16 + \min[0+19, 8+8, 19+3, 25+0] = 16 + 16 = 32 \\
 ft(0, 4) &= 2
 \end{aligned}$$

From the table we see that $C(0,4)=32$ is the minimum cost of a binary search tree for (a_1, a_2, a_3, a_4) . The root of the tree 'T04' is 'a2'.

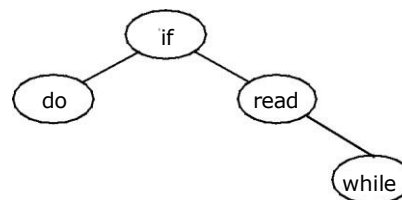
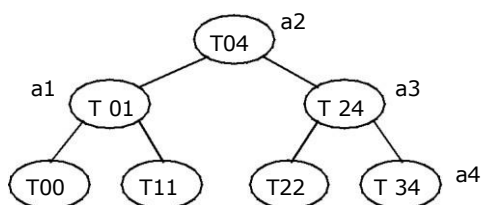
Hence the left subtree is 'T01' and right subtree is 'T24'. The root of 'T01' is 'a1' and the root of 'T24' is 'a3'.

The left and right subtrees for 'T01' are 'T00' and 'T11' respectively. The root of 'T01' is 'a1'.

The left and right subtrees for 'T24' are 'T22' and 'T34' respectively. The root of 'T24' is 'a3'.

The root of 'T22' is null.

The root of 'T34' is 'a4'.



Example2:

Consider four elements a_1, a_2, a_3 and a_4 with $Q_0 = 1/8, Q_1 = 3/16, Q_2 = Q_3 = Q_4 = 1/16$ and $p_1 = 1/4, p_2 = 1/8, p_3 = p_4 = 1/16$. Construct an optimal binary search tree. Solving for $C(0, n)$:

First, computing all $C(i,j)$ such that $j-i=1$; $j=i+1$ and as $0 \leq i < 4$; $i=0, 1, 2$ and 3 ; $i < k \leq J$. Start with $i = 0$; so $j = 1$; as $i < k \leq j$, so the possible value for $k = 1$

$$W(0,1) = P(1) + Q(1) + W(0,0) = 4 + 3 + 2 = 9$$

$$C(0,1)=W(0,1)+\min\{C(0,0)+C(1,1)\}=9+[(0+0)]=9$$

ft(0,1)=1 (value of 'K' that is minimum in the above equation).

Next with $i=1$; $soj=2$; $as i < k \leq j$, so the possible value for $k=2$

$$W(1,2)=P(2)+Q(2)+W(1,1)=2+1+3=6$$

$$C(1,2)=W(1,2)+\min\{C(1,1)+C(2,2)\}=6+[(0+0)]=6$$

ft(1,2)=2

Next with $i=2$; $soj=3$; $as i < k \leq j$, so the possible value for $k=3$

$$W(2,3) = P(3)+Q(3)+W(2,2) = 1+1+3+1=3$$

$$C(2,3) = W(2,3)+\min\{C(2,2)+C(3,3)\}=3+[(0+0)]=3$$

$$ft(2,3)=3$$

Next with $i=3$; $soj=4$; $asi < k \leq j$, so the possible value for $k=4$

$$W(3,4) = P(4) + Q(4) + W(3,3) = 1 + 1 + 1 = 3$$

$$C(3,4) = W(3,4) + \min\{[C(3,3) + C(4,4)]\} = 3 + [(0 + 0)] = 3$$

$$ft(3,4) = 4$$

Second, Computing all $C(i,j)$ such that $j-i=2$; $j=i+2$ and $0 \leq i < 3$; $i=0,1,2$; $i < k \leq J$

Start with $i=0$; $soj=2$; $asi < k \leq j$, so the possible values for $k=1$ and 2 .

$$W(0,2) = P(2) + Q(2) + W(0,1) = 2 + 1 + 9 = 12$$

$$C(0,2) = W(0,2) + \min\{(C(0,0) + C(1,2)), (C(0,1) + C(2,2))\} = 12 + \min\{(0+6, 9+0)\} = 12+6 = 18$$

$$ft(0,2)=1$$

Next, with $i=1$; $soj=3$; $asi < k \leq j$, so the possible value for $k=2$ and 3 .

$$W(1,3) = P(3) + Q(3) + W(1,2) = 1 + 1 + 6 = 8$$

$$C(1,3) = W(1,3) + \min\{[C(1,1) + C(2,3)], [C(1,2) + C(3,3)]\} \\ = W(1,3) + \min\{(0+3), (6+0)\} = 8+3 = 11$$

$$ft(1,3)=2$$

Next, with $i=2$; $soj=4$; $asi < k \leq j$, so the possible value for $k=3$ and 4 .

$$W(2,4) = P(4) + Q(4) + W(2,3) = 1 + 1 + 3 = 5$$

$$C(2,4) = W(2,4) + \min\{[C(2,2) + C(3,4)], [C(2,3) + C(4,4)]\} \\ = 5 + \min\{(0+3), (3+0)\} = 5+3 = 8$$

$$ft(2,4)=3$$

Third, Computing all $C(i,j)$ such that $J-i=3$; $j=i+3$ and $0 \leq i < 2$; $i=0,1$; $i < k \leq J$.

Start with $i=0$; $soj=3$; $asi < k \leq j$, so the possible values for $k=1,2$ and 3 .

$$W(0,3) = P(3) + Q(3) + W(0,2) = 1 + 1 + 12 = 14$$

$$C(0,3) = W(0,3) + \min\{[C(0,0) + C(1,3)], [C(0,1) + C(2,3)], [C(0,2) + C(3,3)]\}$$

$$= 14 + \min\{(0+11), (9+3), (18+0)\} = 14+11 = 25$$

$$ft(0,3)=1$$

Start with $i=1$; $soj=4$; $asi < k \leq j$, so the possible values for $k=2,3$ and 4 .

$$W(1,4) = P(4) + Q(4) + W(1,3) = 1 + 1 + 8 = 10$$

$$C(1,4) = W(1,4) + \min\{[C(1,1) + C(2,4)], [C(1,2) + C(3,4)], [C(1,3) + C(4,4)]\} \\ = 10 + \min\{(0+8), (6+3), (11+0)\} = 10+8 = 18$$

$$ft(1,4) = 10$$

Fourth, Computing all $C(i,j)$ such that $J-i=4$; $j=i+4$ and $0 \leq i < 1$; $i=0$;

$i < k \leq J$. Start with $i=0$; $soj=4$; $asi < k \leq j$, so the possible values for $k=1,2,3$ and 4 .

$$W(0,4) = P(4) + Q(4) + W(0,3) = 1 + 1 + 14 = 16$$

$$C(0,4) = W(0,4) + \min\{[C(0,0) + C(1,4)], [C(0,1) + C(2,4)], [C(0,2) + C(3,4)], [C(0,3) + C(4,4)]\}$$

$$=16+\min[0+18,9+8,18+3,25+0]=16+17=33$$

$$R(0,4)$$

Table for recording $W(i,j)$, $C(i,j)$ and $R(i,j)$

Column Row	0	1	2	3	4
0	2,0,0	1,0,0	1, 0, 0	1, 0, 0,	1,0,0
1	9,9,1	6,6,2	3, 3, 3	3, 3, 4	
2	12,18, 1	8,11,2	5, 8, 3		
3	14,25, 2	11,18, 2			
4	16,33, 2				

From the table we see that $C(0,4)=33$ is the minimum cost of a binary search tree for (a_1, a_2, a_3, a_4)

The root of the tree 'T04' is 'a2'.

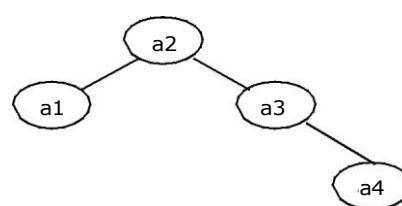
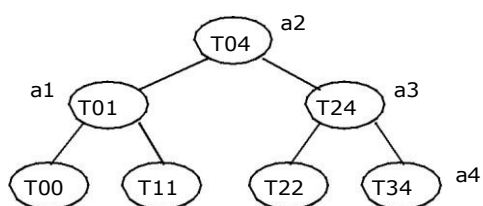
Hence the left subtree is 'T01' and right subtree is T24. The root of 'T01' is 'a1' and the root of 'T24' is 'a3'.

The left and right subtrees for 'T01' are 'T00' and 'T11' respectively. The root of T01 is 'a1'.

The left and right subtrees for T24 are T22 and T34 respectively. The root of T24 is 'a3'.

The root of T22 is null.

The root of T34 is a4.



0/1-KNAPSACK

We are given n objects and a knapsack. Each object i has a positive weight w_i and a positive value V_i . The knapsack can carry a weight not exceeding W . Fill the knapsack so that the value of objects in the knapsack is optimized.

A solution to the knapsack problem can be obtained by making a sequence of decisions on the variables $x_1, x_2, \dots, x_n$. A decision on variable x_i involves determining which of the values 0 or 1 is to be assigned to it. Let us assume that

decisions on the x_i are made in the order $x_n, x_{n-1}, \dots, x_1$. Following a decision on x_n , we may be in one of two possible states: the capacity remaining in $m - w_n$ and a profit of p_n has accrued. It is clear that the remaining decisions $x_{n-1}, \dots, x_1$ must be optimal with respect to the problem state resulting from the decision on x_n . Otherwise, $x_n, \dots, x_1$ will not be optimal. Hence, the principle of optimality holds.

$$F_n(m) = \max\{f_{n-1}(m), f_{n-1}(m - w_n) + p_n\} \quad \text{--} \quad 1$$

For arbitrary $f_i(y)$, $i > 0$, this equation generalizes to:

$$F_i(y) = \max\{f_{i-1}(y), f_{i-1}(y - w_i) + p_i\} \quad \text{--} \quad 2$$

Equation-2 can be solved for $f_n(m)$ by beginning with the knowledge $f_0(y) = 0$ for all y and $f_i(y) = -\infty$, $y < 0$. Then $f_1, f_2, \dots, f_n$ can be successively computed using equation-2.

When the w_i 's are integers, we need to compute $f_i(y)$ for integers y , $0 \leq y \leq m$. Since $f_i(y) = -\infty$ for $y < 0$, these function values need not be computed explicitly. Since each f_i can be computed from f_{i-1} in $\Theta(m)$ time, it takes $\Theta(mn)$ time to compute f_n . When the w_i 's are real numbers, $f_i(y)$ is needed for real numbers y such that $0 < y \leq m$. So, f_i cannot be explicitly computed for all y in this range. Even when the w_i 's are integers, the explicit $\Theta(mn)$ computation of f_n may not be the most efficient computation. So, we explore **an alternative method for both cases**.

The $f_i(y)$ is an ascending step function; i.e., there are a finite number of y 's, $0 = y_1 < y_2 < \dots < y_k$, such that $f_i(y_1) < f_i(y_2) < \dots < f_i(y_k)$; $f_i(y) = -\infty$, $y < y_1$; $f_i(y) = f_i(y_k)$, $y \geq y_k$; and $f_i(y) = f_i(y_j)$, $y_j \leq y < y_{j+1}$. So, we need to compute only $f_i(y_j)$, $1 \leq j \leq k$. We use the ordered set $S^i = \{(f(y_j), y_j) \mid 1 \leq j \leq k\}$ to represent $f_i(y)$. Each number of S^i is a pair (P, W) , where $P = f_i(y_j)$ and $W = y_j$. Notice that $S^0 = \{(0, 0)\}$. We can compute S^{i+1} from S^i by first computing:

$$S^{i+1} = \{(P, W) \mid (P - p_i, W - w_i) \in S^i\}$$

Now, S^{i+1} can be computed by merging the pairs in S^i and S^{i+1} together. Note that if S^{i+1} contains two pairs (P_j, W_j) and (P_k, W_k) with the property that $P_j \leq P_k$ and $W_j > W_k$, then the pair (P_j, W_j) can be discarded because of equation-2. Discarding or purging rules such as this one are also known as dominance rules. Dominated tuples get purged. In the above, (P_k, W_k) dominates (P_j, W_j) .

Reliability Design

The problem is to design a system that is composed of several devices connected in series. Let r_i be the reliability of device D_i (that is, r_i is the probability that device i will function properly) then the reliability of the entire system is $\prod_{i=1}^n r_i$. Even if the individual devices are very reliable (the r_i 's are very close to one), the reliability of the system may not be very good. For example, if $n = 10$ and $r_i = 0.99$, $1 \leq i \leq 10$, then $\prod_{i=1}^{10} r_i = .904$. Hence, it is desirable to duplicate devices. Multiply copies of the same device type are connected in parallel.

If stage i contains m_i copies of device D_i . Then the probability that all m_i have a malfunction is $(1 - r_i)^{m_i}$. Hence the reliability of stage i becomes $1 - (1 - r_i)^{m_i}$.

i

The reliability of stage ' i ' is given by a function $f_i(m_i)$.

Our problem is to use device duplication. This maximization is to be carried out under a cost constraint. Let c_i be the cost of each unit of device i and let C be the maximum allowable cost of the system being designed.

We wish to solve:

$$\begin{aligned} & \text{Maximize } \prod_{i=1}^n f_i(m_i) \\ & \text{Subject to } \sum_{i=1}^n c_i m_i \leq C \\ & m_i \geq 1 \text{ and integer, } 1 \leq i \leq n \end{aligned}$$

Assume each $c_i > 0$, each m_i must be in the range $1 \leq m_i \leq u_i$, where

$$u_i = \left\lfloor \frac{C}{c_i} \right\rfloor$$

The upper bound u_i follows from the observation that $m_j \geq 1$. An

optimal solution $m_1, m_2, \dots, m_n$ is the result of a sequence of decisions, one decision for each m_i .

Let $f_i(x)$ represent the maximum value of $\sum_{j=1}^n q_j m_j$

subject to the constraints:

$$\sum_{j=1}^i c_j m_j \leq x \quad \text{and } 1 \leq m_j \leq u_j, 1 \leq j \leq i$$

The last decision made requires one to choose m_n from $\{1, 2, 3, \dots, u_n\}$

Once a value of m_n has been chosen, the remaining decisions must be such as to use the remaining funds $C - c_n m_n$ in an optimal way.

The principle of optimality holds on

$$f_n(x) = \max_{1 \leq m_n \leq u_n} \{ c_n(m_n) f_{n-1}(x - c_n(m_n)) \}$$

for any $f_i(x)$, $i \geq 1$, this equation generalizes to

$$f_n(x) = \max_{1 \leq m_i \leq u_i} \{ c_i(m_i) f_{i-1}(x - c_i(m_i)) \}$$

clearly, $f_0(x) = 1$ for all x , $0 \leq x \leq C$ and $f(x) = -\infty$ for all $x < 0$. Let

S^i consist of tuples of the form (f, x) , where $f = f_i(x)$.

There is at most one tuple for each different 'x', that result from a sequence of decisions on $m_1, m_2, \dots, m_n$. The dominance rule (f_1, x_1) dominates (f_2, x_2) iff $f_1 \geq f_2$ and $x_1 \leq x_2$. Hence, dominated tuples can be discarded from S^i .

Example 1:

Design a three-stage system with device types D1, D2 and D3. The costs are \$30, \$15 and \$20 respectively. The cost of the system is to be no more than \$105. The reliability of each device is 0.9, 0.8 and 0.5 respectively.

Solution:

We assume that if stage i has m_i devices of type i in parallel, then $O_i(m_i) = 1 - (1 - r_i)^{m_i}$

Since, we can assume each $c_i > 0$, each m_i must be in the range $1 \leq m_i \leq u_i$. Where:

$$u_i = \left\lfloor \frac{C - \sum_{j=1}^{i-1} c_j}{c_i} \right\rfloor$$

Using the above equation compute u_1, u_2 and u_3 .

$$u_1 = \frac{105 + 30 - (30 + 15 + 20)}{30} = \frac{70}{30} = 2$$

$$u_2 = \frac{105 + 15 - (30 + 15 + 20)}{15} = \frac{55}{15} = 3$$

$$u_3 = \frac{105 + 20 - (30 + 15 + 20)}{20} = \frac{60}{20} = 3$$

We use S^i : stage number and J : no. of devices in stage $i = m_i S^o$

$= \{f_o(x), x\}$ initially $f_o(x) = 1$ and $x = 0$, so, $S^o = \{1, 0\}$

Compute S^1, S^2 and S^3 as follows:

S_1 depends on u_1 value, as $u_1 = 2$, so

$$S_1 = \{S_1^1, S_1^2\}$$

S_2 depends on u_2 value, as $u_2 = 3$, so

$$s_2 = \{S_2^1, S_2^2, S_2^3\}$$

S_3 depends on u_3 value, as $u_3 = 3$, so

$$S_3 = \{S_3^1, S_3^2, S_3^3\}$$

Now find $S^1(x)$, $S^2(x)$, $S^3(x)$

$f_1(x) = \{01(1)f_o, 01(2)f_o\}$ With devices $m_1 = 1$ and $m_2 = 2$ Compute $\emptyset_1(1)$

and $\emptyset_1(2)$ using the formula: $\emptyset_i(m_i) = 1 - (1 - r_i)^{m_i}$

$$\emptyset_1(1) = 1 - (1 - 0.9)^1 = 0.9$$

$$\emptyset_1(2) = 1 - (1 - 0.9)^2 = 0.99$$

$$S_1^1 = \{f_1, x\} \sim (0.9, 30)$$

S_2

$$S_2^1 = \{10.99, 30 + 30\} = (0.99, 60)$$

Therefore, $S^1 = \{(0.9, 30), (0.99, 60)\}$

Next find $S_1^2(x)$, $S_2^2(x)$, $S_3^2(x)$

$f_2(x) = \{02(1)*f_1, 02(2)*f_1, 02(3)*f_1\}$

$$\sim 2 \sim 1 \sim \sim 1 \sim \sim 1 \sim r I_{\sim \sim \sim} \cdot 1 - (1 - 0.8) = 1 \cdot 0.2 = 0.8$$

$$2^{2^{2^{1^{1^{0.8^2}}}}} = 0.96$$

$$O_2(3)=1-(1-0.8)^3=0.992$$

$$\begin{aligned}
 &= \{(0.8(0.9), 30+15), (0.8(0.99), 60+15)\} = \{(0.72, 45), (0.792, 75)\} = \\
 &\quad \{(0.96(0.9), 30+15+15), (0.96(0.99), 60+15+15)\} \\
 &= \{(0.864, 60), (0.9504, 90)\}
 \end{aligned}$$

$$\begin{aligned} &= \{(0.992(0.9), 30+15+15+15), (0.992(0.99), 60+15+15+15)\} \\ &= \{(0.8928, 75), (0.98208, 105)\} \end{aligned}$$

$$S_2 = \{S_1^2, S_2^2, S_3^2\}$$

By applying Dominance rule to S^2 :

Therefore, $S_2 = \{(0.72, 45), (0.864, 60), (0.8928, 75)\}$ DominanceRule:

If S^i contains two pairs (f_1, x_1) and (f_2, x_2) with the property that $f_1 \geq f_2$ and $x_1 \leq x_2$, then (f_1, x_1) dominates (f_2, x_2) , hence by dominance rule (f_2, x_2) can be discarded. Discarding or pruning rules such as the one above is known as dominance rule. Dominating tuples will be present in S^i and Dominated tuples has to be discarded from S^i .

Case1:iff1≤f2andx1>x2thendiscard(f1,x1)

Case2:iff1≥f2andx1<x2thediscard(f2,x2)

Case3:otherwise simply write(f1,x1)

$$S2=\{(0.72,45),(0.864,60),(0.8928,75)\}$$

$$\emptyset 3(1)=1 \sim 1_{rI \sim mi}=1-(1-0.5)^1=1-0.5=0.5$$

$$\frac{\theta_2 \sim 2 \sim 1 \sim 1}{\sim S} = 0.75$$

$$S13 = \{(0.5(0.72), 45+20), (0.5(0.864), 60+20), (0.5(0.8928), 75+20)\}$$

$$S_{13} = \{(0.36, 65), (0.437, 80), (0.4464, 95)\}$$

$$S_2^2 = \{(0.75 (0.72), 45 + 20 + 20), (0.75 (0.864), 60 + 20 + 20), (0.75(0.8928), 75+20+20)\}$$

$$=\{(0.54,85),(0.648,100),(0.6696,115)\}$$

$$S_3 = \{ \square 0.875(0.72), 45+20+20+20, \quad \square 0.875(0.864), 60+20+20+20, \\ \square 0.875(0.8928), 75+20+20+20 \quad \square \}$$

S_3

$3 = \{(0.63, 105), (1.756, 120), (0.7812, 135)\}$
If cost exceeds 105, remove that tuples

$$S_3 = \{(0.36, 65), (0.437, 80), (0.54, 85), (0.648, 100)\}$$

The best design has a reliability of 0.648 and a cost of 100. Tracing back for the solution through S^i , we can determine that $m_3=2, m_2=2$ and $m_1=1$.

Other Solution:

According to the principle of optimality:

$$f_n(C) = \max_{mn < un} \{ \sim n(mn). f_{n-1}(C - C_{nmn}) \text{ with } f_0(x) = 1 \text{ and } 0 \leq x \leq C; 1 \sim$$

Since, we can assume each $c_i > 0$, each m_i must be in the range $1 \leq m_i \leq u_i$. Where:

$$S_2 = \{ (0.75(0.72), 45 + 20 + 20), (0.75(0.864), 60 + \sim u_{20+20}), i \\ = \sim i C + C_i \sim C_j r / C_i I \sim$$

Using the above equation compute u_1, u_2 and u_3 .

$$u_1 = \frac{105 \square \square 30 \square +}{30} \sim \frac{70}{30} = 2 \\ u_2 = \frac{105 \square 15 \square 30 \square +}{15} \sim \frac{55}{15} = 3 \\ u_3 = \frac{105 \square \square 30 \square + 20 \square}{20} = \frac{60}{20}$$

$$f_3(105) = \max \{ \sim 3(m_3). f_2(105 - 20m_3) \} \quad 1 < m_3 \leq u_3$$

$$= \max \{ 3(1) f_2(105 - 20), \underline{63(2) f_2(105 - 20 \times 2)}, \sim 3(3) f_2(105 - 20 \times 3) \} = \max \{ 0.5 \\ f_2(85), 0.75 f_2(65), 0.875 f_2(45) \}$$

$$= \max \{ 0.5 \times 0.8928, 0.75 \times 0.864, 0.875 \times 0.72 \} = 0.648.$$

$$= \max \{ 2(m_2). f_1(85 - 15m_2) \} \\ 1! m_2! u_2$$

$$f_2(85) = \max \{ 2(1). f_1(85 - 15), \sim 2(2). f_1(85 - 15 \times 2), \sim 2(3). f_1(85 - 15 \times 3) \} = \\ \max \{ 0.8 f_1(70), 0.96 f_1(55), 0.992 f_1(40) \} \\ = \max \{ 0.8 \times 0.99, 0.96 \times 0.9, 0.99 \times 0.9 \} = 0.8928$$

$$f_1(70) = \max \{ \sim 1(m_1). f_0(70 - 30m_1) \} \\ 1! m_1! u_1 \\ = \max \{ \sim 1(1) f_0(70 - 30), t_1(2) f_0(70 - 30 \times 2) \}$$

$$=\max\{\sim 1(1)x_1, t_1(2)x_1\}=\max\{0.9, 0.99\}=0.99$$

$$f_1(55) = \max\{t_1(m_1).f_0(55-30m_1)\} \\ 1!m_1!u_1$$

$$=\max\{\sim 1(1)f_0(50-30), t_1(2)f_0(50-30x_2)\}$$

$$=\max\{\sim 1(1)x_1, t_1(2)x_{-oo}\}=\max\{0.9, -oo\}=0.9$$

$$f_1(40) = \max\{\sim 1(m_1).f_0(40-30m_1)\} \\ 1!m_1!u_1$$

$$=\max\{\sim 1(1)f_0(40-30), t_1(2)f_0(40-30x_2)\}$$

$$=\max\{\sim 1(1)x_1, t_1(2)x_{-oo}\}=\max\{0.9, -oo\}=0.9$$

$$f_2(65)=\max\{2(m_2).f_1(65-15m_2)\} \quad 1! \\ m_2!u_2$$

$$=\max\{2(1)f_1(65-15), \underline{62(2)f_1(65-15x_2)}, \sim 2(3)f_1(65-15x_3)\}=\max\{0.8f_1(50), \\ 0.96f_1(35), 0.992f_1(20)\}$$

$$=\max\{0.8x0.9, 0.96x0.9, -oo\}=0.864$$

$$f_1(50) = \max\{\sim 1(m_1).f_0(50 - 30m_1)\} \quad 1! \\ !m_1!u_1$$

$$=\max\{\sim 1(1)f_0(50-30), t_1(2)f_0(50-30x_2)\}$$

$$=\max\{\sim 1(1)x_1, t_1(2)x_{-oo}\}=\max\{0.9, -oo\}=0.9f_1(35)=\max$$

$$\sim 1(m_1).f_0(35-30m_1)\}$$

$$1!m_1!u_1$$

$$=\max\{\sim 1(1).f_0(35-30), \sim 1(2).f_0(35-30x_2)\}$$

$$=\max\{\sim 1(1)x_1, t_1(2)x_{-oo}\}=\max\{0.9, -oo\}=0.9$$

$$f_1(20)=\max\{\sim 1(m_1).f_0(20-30m_1)\} \quad 1! \\ m_1!u_1$$

$$=\max\{\sim 1(1)f_0(20-30), t_1(2)f_0(20-30x_2)\}$$

$$=\max\{\sim 1(1)x_-, \sim 1(2)x_{-oo}\}=\max\{-oo, -oo\}=-oo f_2$$

$$(45) = \max\{2(m_2).f_1(45-15m_2)\} \\ 1!m_2!u_2$$

$$=\max\{2(1)f_1(45-15), \sim 2(2)f_1(45-15x_2), \sim 2(3)f_1(45-15x_3)\}=\max\{0.8f_1(30),$$

$$0.96f_1(15), 0.992f_1(0)\}$$

$$=\max\{0.8x0.9, 0.96x-, 0.99x_{-oo}\}=0.72$$

$$\begin{aligned}
 f_1(30) &= \max\{f_1(m_1), f_0(30-30m_1)\} \quad 1 \leq m_1 \leq u_1 \\
 &= \max\{f_1(1)f_0(30-30), f_1(2)f_0(30-30 \times 2)\} \\
 &= \max\{f_1(1) \times 1, f_1(2) \times -\infty\} = \max\{0.9, -\infty\} = 0.9
 \end{aligned}$$

Similarly, $f_1(15) = -$, $f_1(0) = -$.

-.

The best design has a reliability = 0.648 and

$$\text{Cost} = 30x_1 + 15x_2 + 20x_3 = 100.$$

Tracing back for the solution through S^i , we can determine that: $m_3 = 2, m_2 = 2$ and $m_1 =$

1.

- Explicit constraints.
- Implicit constraints.

Explicit constraints: Explicit constraints are rules that restrict each x_i to take on values only from a given set.

Example: $x_i \geq 0$ or $s_i = \{\text{all nonnegative real numbers}\}$

$x_i = 0$ or 1 or $s_i = \{0, 1\}$

$l_i \leq x_i \leq u_i$ or $s_i = \{a: l_i \leq a \leq u_i\}$

The explicit constraint depends on the particular instance of the problem being solved. All tuples that satisfy the explicit constraints define a possible solution space for I .

Implicit Constraints:

The implicit constraints are rules that determine which of the tuples in the solution space of I satisfy the criterion function. Thus implicit constraints describe the way in which the x_i must relate to each other.

Applications of Backtracking:

- NQueens Problem
- Sum of subsets problem
- Graph coloring
- Hamiltonian cycles.

N-Queens Problem:

It is a classic combinatorial problem. The eight queen's puzzle is the problem of placing eight queens puzzle is the problem of placing eight queens on an 8×8 chessboard so that no two queens attack each other. That is so that no two of them are on the same row, column, or diagonal. The 8-queens puzzle is an example of the more general n-queens problem of placing n queens on an $n \times n$ chessboard.

	1	2	3	4	5	6	7	8
1				Q				
2						Q		
3								Q
4		Q						
5							Q	
6	Q							
7			Q					
8					Q			

One solution to the 8-queens problem

Here queens can also be numbered 1 through 8

Each queen must be on a different row

Assume queen 'i' is to be placed on row 'i'

All solutions to the 8-queens problem can therefore be represented as s -tuples $(x_1, x_2, x_3, \dots, x_8)$ where $x_i \rightarrow$ the column on which queen 'i' is placed

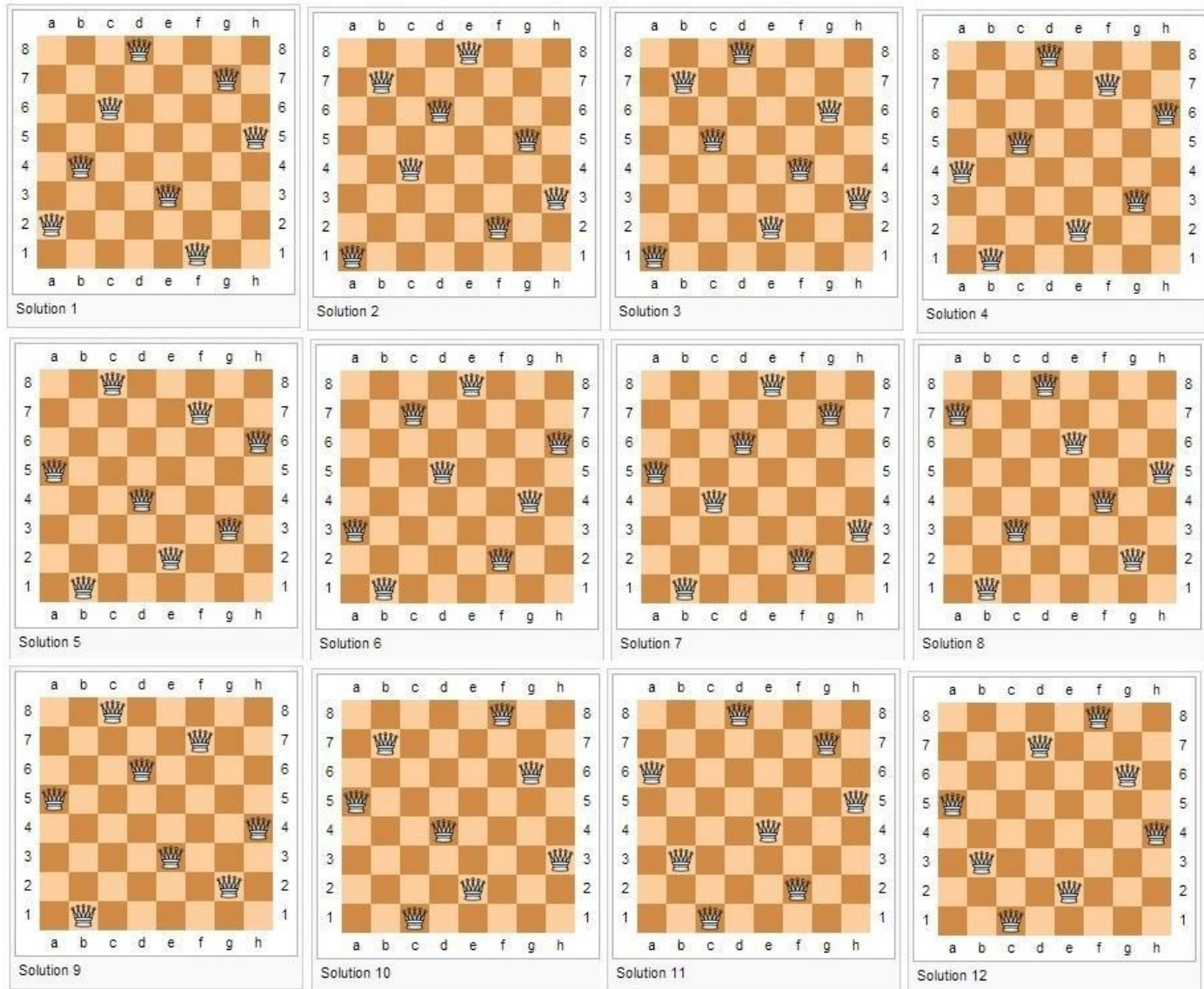
$s_i \rightarrow \{1, 2, 3, 4, 5, 6, 7, 8\}, 1 \leq i \leq 8$

Therefore the solution space consists of 8^8 s -tuples.

The implicit constraints for this problem are that not two x_i 's can be the same column and not two queens can be on the same diagonal.

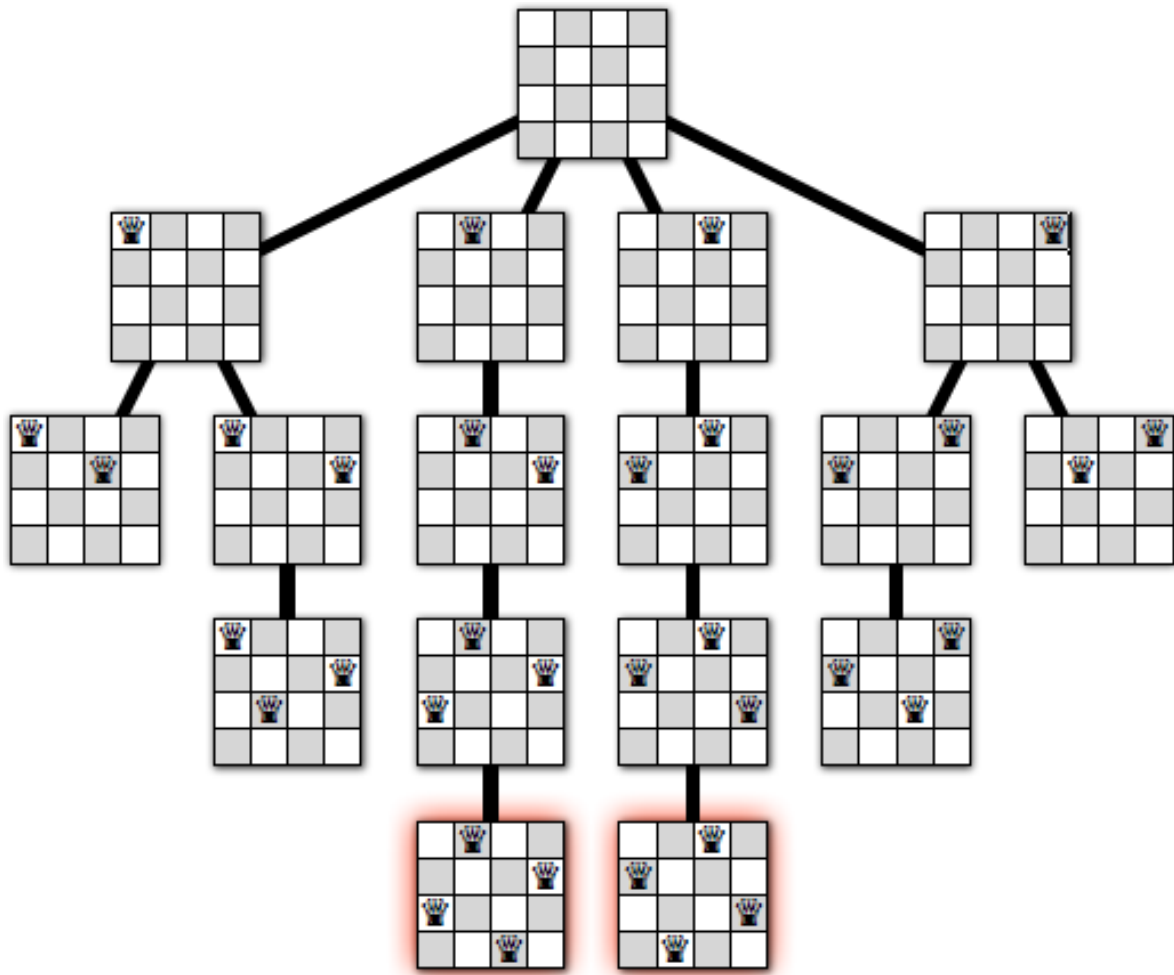
By these two constraints the size of solution space reduces from 8^8 tuples to $8!$ Tuples. For example $s_i(4, 6, 8, 2, 7, 1, 3, 5)$

In the same way for n-queens are to be placed on an $n \times n$ chessboard, the solution space consists of all $n!$ Permutations of n-tuples (1,2, -----n).



Some solution to the 8-Queens problem

Algorithm for new queen be placed	All solution to the n-queens problem
<pre> AlgorithmPlace(k,i) //Return true if a queen can be placed in kth row & ith column //Otherwise return false { forj:=1 tok-1 do if(x[j]=iorAbs(x[j]-i)=Abs(j-k))) then return false return true } </pre>	<pre> AlgorithmNQueens(k,n) //its prints all possible placementsof n- queens on an n×n chessboard. { fori:=1 to n do{ ifPlace(k,i)then { X[k]:=i; if(k==n)thenwrite(x[1:n]); else NQueens(k+1, n); } }} </pre>



The complete recursion tree for our algorithm for the 4 queens problem.

Sum of Subsets Problem:

Given positive numbers $w_1, w_2, \dots, w_n$, and m , here sum of subsets problem is finding all subset of w_i whose sums are m .

Definition: Given n distinct +ve numbers (usually called weights), desire (want) to find all combinations of these numbers whose sums are m . this is called sum of subsets problem. To formulate this problem by using either fixed sized tuples or variable sized tuples. Backtracking solution uses the fixed size tuple strategy.

For example:

If $n=4$ (w_1, w_2, w_3, w_4) = (11, 13, 24, 7) and $m=31$.

Then desired subsets are (11, 13, 7) & (24, 7).

These two solutions are described by the vectors (1, 2, 4) and (3, 4).

In general all solutions are k -tuples $(x_1, x_2, x_3, \dots, x_k)$ $1 \leq k \leq n$, different solutions may have different sized tuples.

- Explicit constraints requires $x_i \in \{j/j \text{ is an integer } 1 \leq j \leq n\}$
- Implicit constraints requires:
No two be the same & that the sum of the corresponding w_i 's be m
i.e., (1, 2, 4) & (1, 4, 2) represents the same. Another constraint is $x_i < x_{i+1}$ $1 \leq i \leq k$

$w_i \rightarrow$ weight of item i

$M \rightarrow$ Capacity of bag (subset)

$X_i \rightarrow$ the element of the solution vector is either one or zero.

X_i value depending on whether the weight w_i is included or not. If $X_i = 1$ then w_i is chosen.

If $X_i = 0$ then w_i is not chosen.

$$\underbrace{\sum_{i=1}^k W(i)X(i)}_{\text{Total sum till now}} + \underbrace{\sum_{i=k+1}^n W(i)}_{\text{Still there}} \geq M$$

The above equations specify that $x_1, x_2, x_3, \dots, x_k$ cannot lead to an answer if this condition is not satisfied.

$$\sum_{i=1}^k W(i)X(i) + W(k+1) > M$$

The equation cannot lead to solution.

$$B_k(X(1), \dots, X(k)) = \text{true iff} \left(\sum_{i=1}^k W(i)X(i) + \sum_{i=k+1}^n W(i) \geq M \text{ and } \sum_{i=1}^k W(i)X(i) + W(k+1) \leq M \right)$$

$$s = \sum_{j=1}^{k-1} W(j)X(j). \quad \text{and} \quad r = \sum_{j=k}^n W(j)$$

Recursive backtracking algorithm for sum of subsets problem

Algorithm SumOfSub(s, k, r)

{

$$//s = \sum_{j=1}^{k-1} W(j)X(j). \quad \text{and} \quad r = \sum_{j=k}^n W(j)$$

$X[k] = 1$

If $(S + w[k] = M)$ then write $(x[1:])$; // subset found.

Else if $(S + w[k] + w[k+1] \leq M)$

Then SumOfSub($S + w[k], k+1, r - w[k]$);

if $((S + r - w[k] \geq M) \text{ and } (S + w[k+1] \leq M))$ then

{

$X[k] = 0$;

SumOfSub($S, k+1, r - w[k]$);

}

}

Graph Coloring:

Let G be a undirected graph and 'm' be a given +ve integer. The graph coloring problem is assigning color to the vertices of an undirected graph with the restriction that not two adjacent vertices are assigned the same color yet only 'm' colors are used.

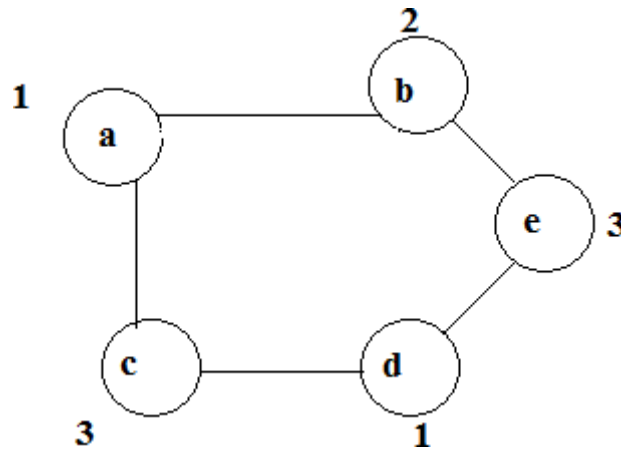
The optimization version calls for coloring a graph using the minimum number of coloring.

The decision version, known as K-coloring asks whether a graph is colourable using at most k- colors.

Note that, if 'd' is the degree of the given graph then it can be colored with 'd+1' colors.

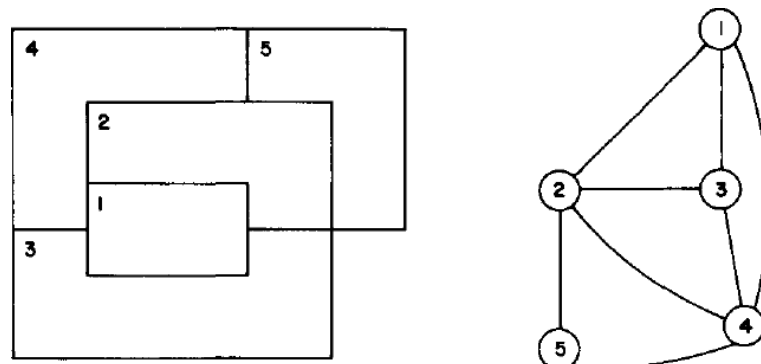
The m-colorability optimization problem asks for the smallest integer 'm' for which the graph G can be colored. This integer is referred as "**Chromatic number**" of the graph.

Example



- Above graph can be colored with 3 colors 1, 2, & 3.
- The color of each node is indicated next to it.
- 3-colors are needed to color this graph and hence this graph's Chromatic Number is 3.
- A graph is said to be planar if it can be drawn in a plane (flat) in such a way that not two edges cross each other.
- **M-Colorability decision problem** is the 4-color problem for planar graphs.
- Given any map, can the regions be colored in such a way that not two adjacent regions have the same color yet only 4-colors are needed?
- To solve this problem, graphs are very useful, because a map can easily be transformed into a graph.
- Each region of the map becomes a node, and if two regions are adjacent, then the corresponding nodes are joined by an edge.

○ Example:



○ **A map and its planar graph representation**

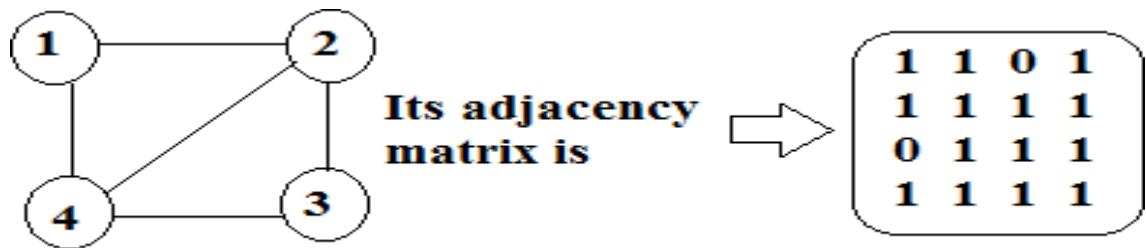
The above map requires 4 colors.

- Many years, it was known that 5-colors were required to color this map.

- After several hundred years, this problem was solved by a group of mathematicians with the help of a computer. They show that 4-colors are sufficient.

Suppose we represent a graph by its adjacency matrix $G[1:n, 1:n]$

Ex:



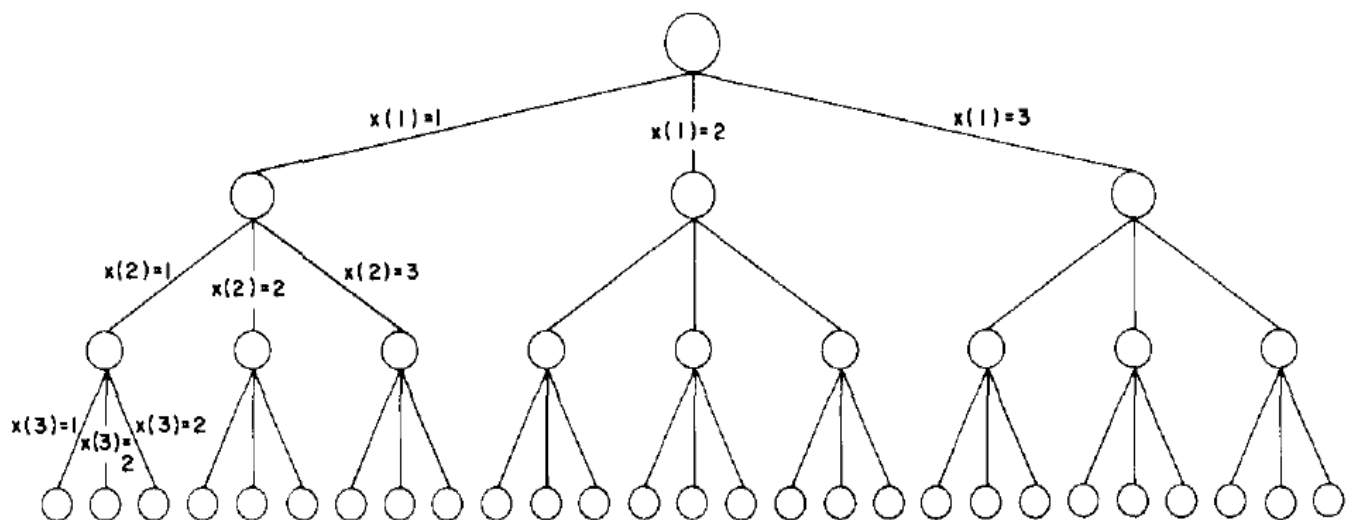
Here $G[i,j]=1$ if (i,j) is an edge of G , and $G[i,j]=0$ otherwise.

Colors are represented by the integers 1, 2, ..., m and the solutions are given by the n -tuple $(x_1, x_2, \dots, x_n)$ $x_i \rightarrow$ Color of node i .

State Space Tree for

$n=3 \rightarrow$ nodes

$m=3 \rightarrow$ colors



State space tree for M Coloring when $n = 3$ and $m = 3$

1st node coloured in 3-ways

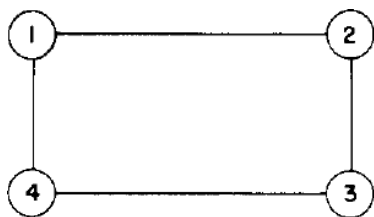
2nd node coloured in 3-ways

3rd node coloured in 3-ways

So we can colour in the graph in 27 possibilities of colouring.

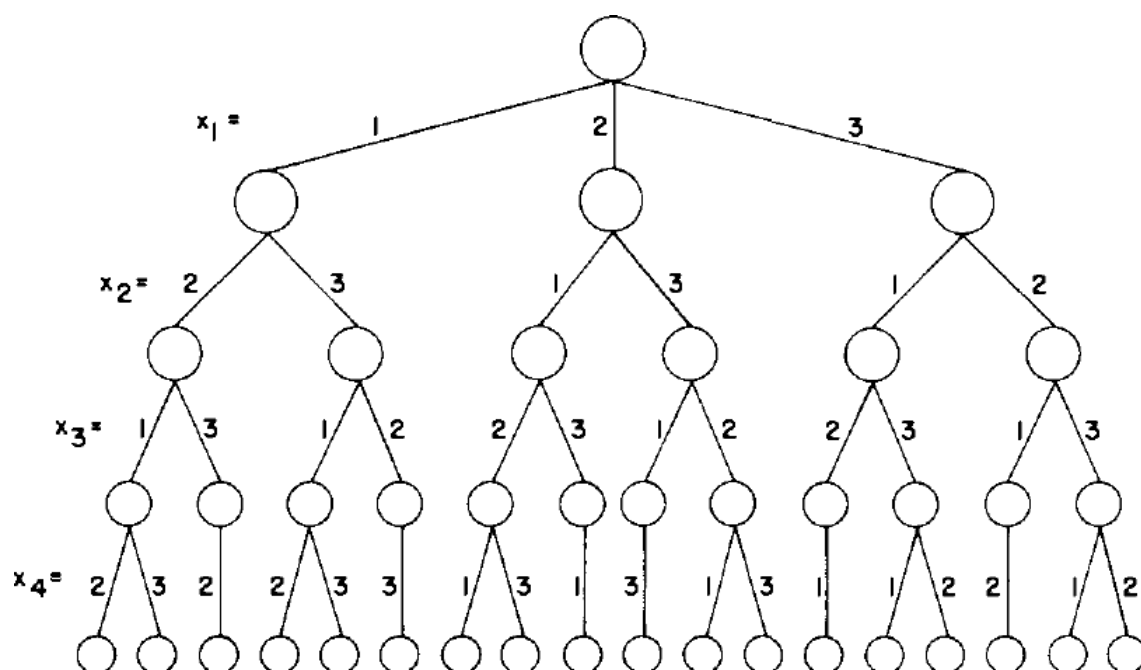
Finding all m-colorings of a graph	Getting next color
<pre> Algorithm mColoring(k){ //g(1:n,1:n) → boolean adjacency matrix. //k → index(node) of the next vertex to color. repeat{ nextvalue(k); // assign to x[k] a legal color. if(x[k]=0) then return; // no new color possible if(k=n) then write(x[1:n]); else mcoloring(k+1); } until(false) } </pre>	<pre> Algorithm NextValue(k){ //x[1],x[2],...x[k-1] have been assigned integer values in the range [1, m] repeat{ x[k]=(x[k]+1) mod (m+1); // next highest color if(x[k]=0) then return; // all colors have been used. for j=1 to n do { if((g[k,j]≠0) and (x[k]=x[j])) then break; } if(j=n+1) then return; // new color found } until(false) } </pre>

Previous paper example:



Adjacency matrix is

$$\begin{pmatrix} 1 & 1 & 0 & 1 \\ 1 & 1 & 1 & 0 \\ 0 & 1 & 1 & 1 \\ 1 & 0 & 1 & 1 \end{pmatrix}$$

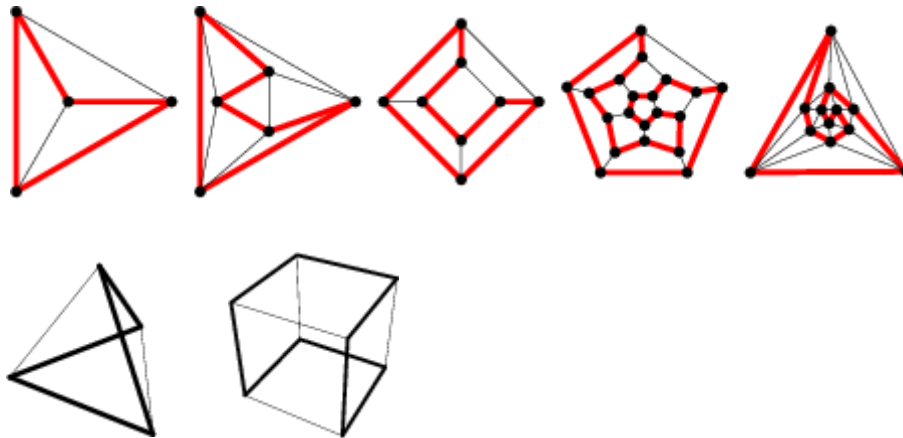


A 4 node graph and all possible 3 colorings

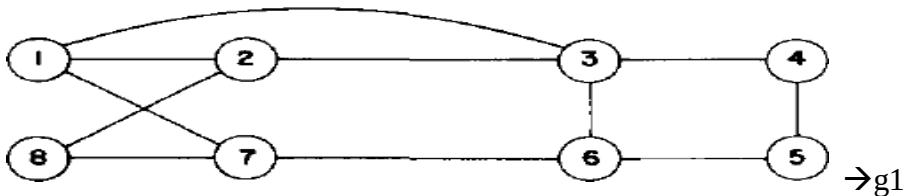
Hamiltonian Cycles:

- **Def:** Let $G=(V, E)$ be a connected graph with n vertices. A Hamiltonian cycle is a round trip path along n -edges of G that visits every vertex once & returns to its starting position.
- It is also called the Hamiltonian circuit.
- Hamiltonian circuit is a graph cycle (i.e., closed loop) through a graph that visits each node exactly once.
- A graph possessing a Hamiltonian cycle is said to be a Hamiltonian graph.

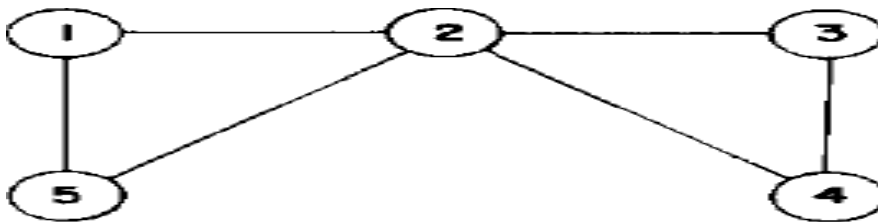
Example:



- In graph G , Hamiltonian cycle begins at some vertex $v_1 \in G$ and the vertices of G are visited in the order $v_1, v_2, \dots, v_{n+1}$, then the edges (v_i, v_{i+1}) are in E , $1 \leq i \leq n$.



The above graph contains Hamiltonian cycle: 1,2,8,7,6,5,4,3,1



The above graph contains no Hamiltonian cycles.

- There is no known easy way to determine whether a given graph contains a Hamiltonian cycle.
- By using backtracking method, it can be possible
 - Backtracking algorithm, that finds all the Hamiltonian cycles in a graph.
 - The graph may be directed or undirected. Only distinct cycles are output.
 - From graph1 backtracking solution vector = {1,2,8,7,6,5,4,3,1}
 - The backtracking solution vector $(x_1, x_2, \dots, x_n)$
 $x_i \rightarrow i^{\text{th}}$ visited vertex of proposed cycle.

- By using backtracking we need to determine how to compute the set of possible vertices for x_k if $x_1, x_2, x_3, \dots, x_{k-1}$ have already been chosen.

If $k=1$ then x_1 can be any of the n -vertices.

By using “Next Value” algorithm the recursive backtracking scheme to find all Hamiltonian cycles.

This algorithm is started by 1st initializing the adjacency matrix $G[1:n, 1:n]$ then setting $x[2:n]$ to zero & $x[1]$ to 1, and then executing Hamiltonian (2)

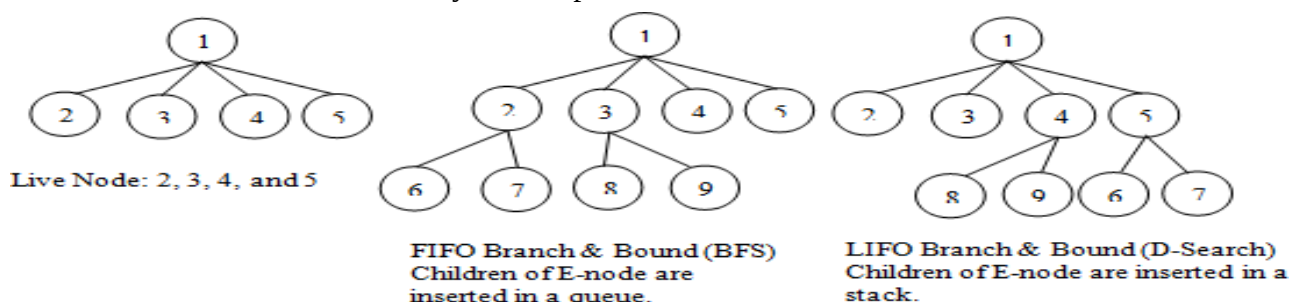
Generating Next Vertex	Finding all Hamiltonian Cycles
<pre> Algorithm NextValue(k) { //x[1:k-1] → is path of k-1 distinct vertices. //if x[k]=0, then no vertex has yet been assigned to x[k] Repeat{ X[k]=(x[k]+1)mod(n+1); //Next vertex If(x[k]=0) then return; If(G[x[k-1], x[k]] ≠ 0) then { For j:=1 to k-1 do if(x[j]=x[k]) then break; //Check for distinctness If(j=k) then //if true, then vertex is distinct If((k < n) or (k = n) and G[x[n], x[1]] ≠ 0)) Then return ; } } Until (false); }</pre>	<pre> Algorithm Hamiltonian(k) { Repeat{ NextValue(k); //assign a legal next value to x[k] If(x[k]=0) then return; If(k=n) then write(x[1:n]); Else Hamiltonian(k+1); } until (false) }</pre>

Branch & Bound

Branch & Bound (B & B) is a general algorithm (or Systematic method) for finding optimal solution of various optimization problems, especially in discrete and combinatorial optimization.

- The B&B strategy is very similar to backtracking in that a state space tree is used to solve a problem.
- The differences are that the B&B method
 - ✓ Does not limit us to any particular way of traversing the tree.
 - ✓ It is used only for optimization problems.
 - ✓ It is applicable to a wide variety of discrete combinatorial problems.
- B&B is a rather general optimization technique that applies where the greedy method & dynamic programming fail.
- It is much slower, indeed (truly), it often (rapidly) leads to exponential time complexities in the worst case.
- The term B&B refers to all state space search methods in which all children of the “E-node” are generated before any other “live node” can become the “E-node”
 - ✓ **Live node** → is a node that has been generated but whose children have not yet been generated.
 - ✓ **E-node** → is a live node whose children are currently being explored.

✓ **Dead node** → is a generated node that is not to be expanded or explored any further. All children of a dead node have already been expanded.



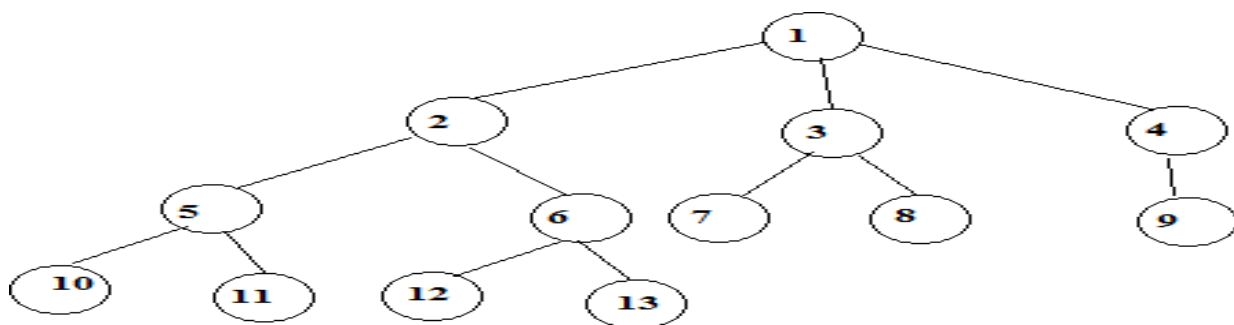
- Two graph search strategies, BFS & D-search (DFS) in which the exploration of a new node cannot begin until the node currently being explored is fully explored.
- Both BFS & D-search (DFS) generalized to B&B strategies.
- ✓ **BFS** → like state space search will be called FIFO (First In First Out) search as the list of live nodes is “First-in-first-out” list (or queue).
- ✓ **D-search (DFS)** → Like state space search will be called LIFO (Last In First Out) search as the list of live nodes is a “last-in-first-out” list (or stack).
- In backtracking, bounding function are used to help avoid the generation of sub-trees that do not contain an answer node.
- We will use 3 types of search strategies in branch and bound
 - 1) FIFO (First In First Out) search
 - 2) LIFO (Last In First Out) search
 - 3) LC (Least Count) search

FIFO B&B:

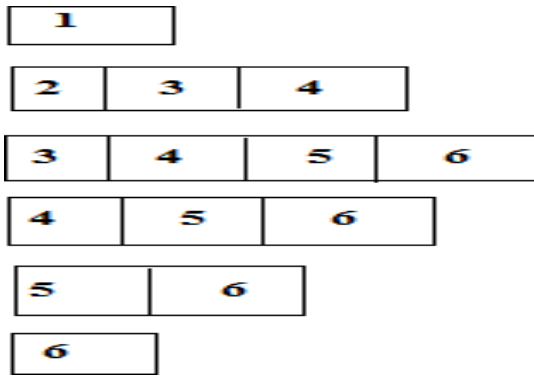
FIFO Branch & Bound is a BFS.

In this, children of E-Node (or Live nodes) are inserted in a queue. Implementation of list of live nodes as a queue

- ✓ **Least()** → Removes the head of the Queue
- ✓ **Add()** → Adds the node to the end of the Queue



Assume that node '12' is an answer node in FIFO search, 1st we take E-node has '1'



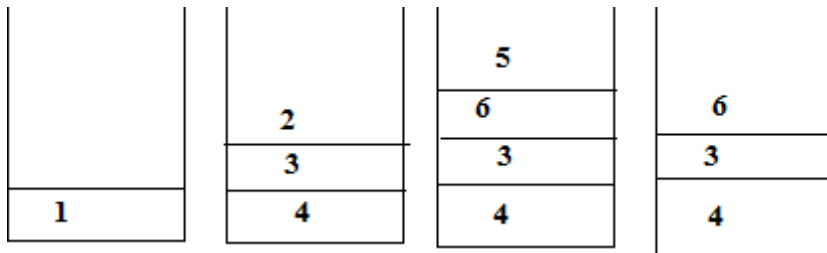
LIFOB&B:

LIFO Branch & Bound is a D-search (or DFS).

In this, children of E-node (live nodes) are inserted in a stack.

Implementation of List of live nodes as a stack

- ✓ Least() → Removes the top of the stack
- ✓ ADD() → Adds the node to the top of the stack.



Least Cost (LC) Search:

The selection rule for the next E-node in FIFO or LIFO branch and bound is sometimes “blind”. i.e., the selection rule does not give any preference to a node that has a very good chance of getting the search to an answer node quickly.

The search for an answer node can often be speeded by using an “intelligent” ranking function. It is also called an approximate cost function “ $\hat{C}$ ”.

Expanded node (E-node) is the live node with the best $\hat{C}$ value.

Branching: A set of solutions, which is represented by a node, can be partitioned into mutually (jointly or commonly) exclusive (special) sets. Each subset in the partition is represented by a child of the original node.

Lower bounding: An algorithm is available for calculating a lower bound on the cost of any solution in a given subset.

Each node X in the search tree is associated with a cost: $\hat{C}(X)$

C = cost of reaching the current node, X (E-node) from the root + The cost of reaching an answer node from X .

$$\hat{C} = g(X) + h(X).$$

Example:

8-puzzle

Cost function: $\hat{C} = g(x) + h(x)$

where $h(x)$ = the number of misplaced tiles

and $g(x)$ = the number of moves so far

Assumption: move one tile in any direction cost 1.

Travelling Salesman Problem:

Def:- Find a tour of minimum cost starting from a node S going through other nodes only once and returning to the starting point S .

Time Complexity of TSP for Dynamic Programming algorithm is $O(n^2 2^n)$

B&B algorithms for this problem, the worst case complexity will not be any better than $O(n^2 2^n)$ but good bounding functions will enable these B&B algorithms to solve some problem instances in much less time than required by the dynamic programming algorithm.

Let $G=(V,E)$ be a directed graph defining an instance of TSP. Let

$C_{ij} \rightarrow$ cost of edge $\langle i, j \rangle$

$C_{ij} = \infty$ if $\langle i, j \rangle \notin E$

$|V|=n \rightarrow$ total number of vertices.

Assume that every tour starts & ends at vertex 1.

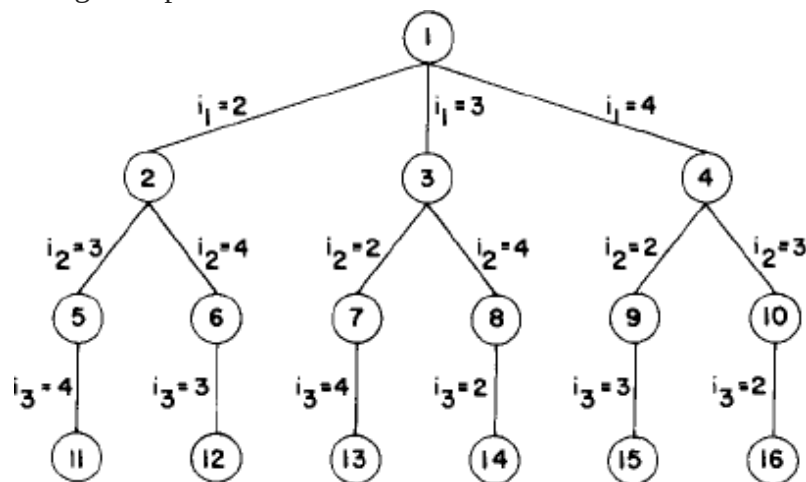
Solution Space $S = \{1, \Pi, 1/\Pi \mid \Pi \text{ is a permutation of } (2, 3, 4, \dots, n)\}$ then $|S| = (n-1)!$

The size of S reduced by restricting S

So that $(1, i_1, i_2, \dots, i_{n-1}, 1) \in S$ iff $\langle i_j, i_{j+1} \rangle \in E, 0 \leq j \leq n-1, i_0 = i_n = 1$

Can be organized into "Statespace tree".

Consider the following Example



Statespace tree for the travelling salesperson problem with $n=4$ and $i_0=i_4=1$. The

above diagram shows tree organization of a complete graph with $|V|=4$.

Each leaf node 'L' is a solution node and represents the tour defined by the path from the root to L.

Node 12 represents the tour.

$i_0=1, i_1=2, i_2=4, i_3=3, i_4=1$

Node 14 represents the tour.

$i_0=1, i_1=3, i_2=4, i_3=2, i_4=1$.

TSP solved by using LC Branch & Bound:

To use LCBB to search the travelling salesperson "Statespace tree" first define a cost function $C(.)$ and other 2 functions $\hat{C}(.)$ & $u(.)$

Such that $\hat{C}(r) \leq C(r) \leq u(r) \rightarrow$ for all nodes r .

Cost $C(.) \rightarrow$ is the solution node with least $C(.)$ corresponds to a shortest tour in G .

$C(A) = \{\text{Length of tour defined by the path from root to } A \text{ if } A \text{ is leaf}$

$\text{Cost of a minimum-cost leaf in the sub-tree } A, \text{ if } A \text{ is not leaf}\}$

From $\hat{C}_1(r) \leq C(r)$ then $\hat{C}(r) \rightarrow$ is the length of the path defined at node A .

From previous example the path defined at node 6 is $i_0, i_1, i_2 = 1, 2, 4$ & it consists of

$\langle 1, 2 \rangle$ & $\langle 2, 4 \rangle$

A better $\hat{C}(r)$ can be obtained by using the reduced cost matrix corresponding to G .

➤ A row (column) is said to be reduced iff it contains at least one zero & remaining entries are non negative.

➤ A matrix is reduced iff every row & column is reduced.

$$\begin{bmatrix} \infty & 20 & 30 & 10 & 11 \\ 15 & \infty & 16 & 4 & 2 \\ 3 & 5 & \infty & 2 & 4 \\ 19 & 6 & 18 & \infty & 3 \\ 16 & 4 & 7 & 16 & \infty \end{bmatrix}$$

(a) Cost Matrix

$$\begin{bmatrix} \infty & 10 & 17 & 0 & 1 \\ 12 & \infty & 11 & 2 & 0 \\ 0 & 3 & \infty & 0 & 2 \\ 15 & 3 & 12 & \infty & 0 \\ 11 & 0 & 0 & 12 & \infty \end{bmatrix}$$

(b) Reduced Cost Matrix

$$L = 25$$

Given the following cost matrix:

$$\begin{bmatrix} \text{inf} & 20 & 30 & 10 & 11 \\ 15 & \text{inf} & 16 & 4 & 2 \\ 3 & 5 & \text{inf} & 2 & 4 \\ 19 & 6 & 18 & \text{inf} & 3 \\ 16 & 4 & 7 & 16 & \text{inf} \end{bmatrix}$$

➤ The TSP starts from node 1: **Node 1**

➤ Reduced Matrix: To get the lower bound of the path starting at node 1

Row # 1: reduce by 10	Row #2: reduce by 2	Row #3: reduce by 2
$\begin{bmatrix} \text{inf} & 10 & 20 \\ 15 & \text{inf} & 16 \\ 3 & 5 & \text{inf} \\ 19 & 6 & 18 \\ 16 & 4 & 7 \end{bmatrix}$	$\begin{bmatrix} \text{inf} & 10 & 20 \\ 13 & \text{inf} & 14 \\ 3 & 5 & \text{inf} \\ 19 & 6 & 18 \\ 16 & 4 & 7 \end{bmatrix}$	$\begin{bmatrix} \text{inf} & 10 & 20 \\ 13 & \text{inf} & 14 \\ 1 & 3 & \text{inf} \\ 19 & 6 & 18 \\ 16 & 4 & 7 \end{bmatrix}$
Row # 4: Reduce by 3:	Row #5: Reduce by 4	Column 1: Reduce by 1

$\begin{bmatrix} \text{inf} & 10 & 20 \\ 13 & \text{inf} & 14 \\ 1 & 3 & \text{inf} & 0 \\ 16 & 3 & 15 & \text{inf} \\ 16 & 4 & 7 & 16 \end{bmatrix}$	$\begin{bmatrix} \text{inf} & 10 & 20 \\ 13 & \text{inf} & 14 \\ 1 & 3 & \text{inf} & 0 \\ 16 & 3 & 15 & \text{inf} \\ 12 & 0 & 3 & 12 \end{bmatrix}$	$\begin{bmatrix} \text{inf} & 10 & 20 \\ 12 & \text{inf} & 14 \\ 0 & 3 & \text{inf} & 0 \\ 15 & 3 & 15 & \text{inf} \\ 11 & 0 & 3 & 12 \end{bmatrix}$
Column2:Itisreduced.	Column 3: Reduceby3 $\begin{bmatrix} \text{inf} & 10 & 17 & 0 & 1 \\ 12 & \text{inf} & 11 & 2 & 0 \\ 0 & 3 & \text{inf} & 0 & 2 \\ 15 & 3 & 12 & \text{inf} & 0 \\ 11 & 0 & 0 & 12 & \text{inf} \end{bmatrix}$	Column4:Itisreduced. Column5:Itisreduced.

The reduced cost is: RCL=25
 So the cost of node 1 is: Cost(1)=25 The reduced matrix is:

$$\text{Cost (1) = 25}$$

$$\begin{bmatrix} \text{inf} & 10 & 17 & 0 & 1 \\ 12 & \text{inf} & 11 & 2 & 0 \\ 0 & 3 & \text{inf} & 0 & 2 \\ 15 & 3 & 12 & \text{inf} & 0 \\ 11 & 0 & 0 & 12 & \text{inf} \end{bmatrix}$$

➤ **Choosetogotovertex 2:Node2**

- Cost of edge <1,2> is: A(1,2) = 10
- Set row #1 = inf since we are choosing edge <1,2>
- Set column # 2 = inf since we are choosing edge <1,2>
- Set A(2,1) = inf
- The resulting cost matrix is:

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & 11 & 2 & 0 \\ 0 & \text{inf} & \text{inf} & 0 & 2 \\ 15 & \text{inf} & 12 & \text{inf} & 0 \\ 11 & \text{inf} & 0 & 12 & \text{inf} \end{bmatrix}$$

- The matrix is reduced:
- RCL=0
- The cost of node 2 (Considering vertex 2 from vertex 1) is:
Cost(2) = cost(1) + A(1,2) = 25 + 10 = 35

➤ Choosetogotovertex 3:Node3

- Cost of edge <1,3> is: $A(1,3) = 17$ (In the reduced matrix)
- Set row #1 = inf since we are starting from node 1
- Set column # 3 = inf since we are choosing edge <1,3>
- Set $A(3,1) = \text{inf}$
- The resulting cost matrix is:

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 12 & \text{inf} & \text{inf} & 2 & 0 \\ \text{inf} & 3 & \text{inf} & 0 & 2 \\ 15 & 3 & \text{inf} & \text{inf} & 0 \\ 11 & 0 & \text{inf} & 12 & \text{inf} \end{bmatrix}$$

Reduce the matrix: Rows are reduced

The columns are reduced except for column #1:

Reduce column 1 by 11:

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 1 & \text{inf} & \text{inf} & 2 & 0 \\ \text{inf} & 3 & \text{inf} & 0 & 2 \\ 4 & 3 & \text{inf} & \text{inf} & 0 \\ 0 & 0 & \text{inf} & 12 & \text{inf} \end{bmatrix}$$

The lower bound is: $RCL = 11$

The cost of going through node 3 is:

$$\text{cost}(3) = \text{cost}(1) + RCL + A(1,3) = 25 + 11 + 17 = 53$$

➤ Choosetogo to vertex 4: Node 4

Remember that the cost matrix is the one that was reduced at the starting vertex 1. Cost of

edge <1,4> is: $A(1,4) = 0$

Set row #1 = inf since we are starting from node 1

Set column #4 = inf since we are choosing edge <1,4> Set

$A(4,1) = \text{inf}$

The resulting cost matrix is:

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 12 & \text{inf} & 11 & \text{inf} & 0 \\ 0 & 3 & \text{inf} & \text{inf} & 2 \\ \text{inf} & 3 & 12 & \text{inf} & 0 \\ 11 & 0 & 0 & \text{inf} & \text{inf} \end{bmatrix}$$

Reduce the matrix: Rows are reduced

Columns are reduced The
lower bound is: RCL = 0
The cost of going through node 4 is:
 $\text{cost}(4) = \text{cost}(1) + \text{RCL} + A(1,4) = 25 + 0 + 0 = 25$

➤ **Choose to go to vertex 5: Node 5**

- Remember that the cost matrix is the one that was reduced at starting vertex 1
- Cost of edge <1,5> is: $A(1,5) = 1$
- Set row #1 = inf since we are starting from node 1
- Set column # 5 = inf since we are choosing edge <1,5>
- Set $A(5,1) = \text{inf}$
- The resulting cost matrix is:

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 12 & \text{inf} & 11 & 2 & \text{inf} \\ 0 & 3 & \text{inf} & 0 & \text{inf} \\ 15 & 3 & 12 & \text{inf} & \text{inf} \\ \text{inf} & 0 & 0 & 12 & \text{inf} \end{bmatrix}$$

Reduce the matrix:

Reduce rows:

Reduce row #2: Reduce by 2

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 10 & \text{inf} & 9 & 0 & \text{inf} \\ 0 & 3 & \text{inf} & 0 & \text{inf} \\ 15 & 3 & 12 & \text{inf} & \text{inf} \\ \text{inf} & 0 & 0 & 12 & \text{inf} \end{bmatrix}$$

Reduce row #4: Reduce by 3

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 10 & \text{inf} & 9 & 0 & \text{inf} \\ 0 & 3 & \text{inf} & 0 & \text{inf} \\ 12 & 0 & 9 & \text{inf} & \text{inf} \\ \text{inf} & 0 & 0 & 12 & \text{inf} \end{bmatrix}$$

Columns are reduced

The lower bound is: $\text{RCL} = 2 + 3 = 5$ The cost
of going through node 5 is:

$$\text{cost}(5) = \text{cost}(1) + \text{RCL} + A(1,5) = 25 + 5 + 1 = 31$$

In summary:

So the live nodes we have so far are:

- ✓ 2: cost(2)=35, path: 1->2
- ✓ 3: cost(3)=53, path: 1->3
- ✓ 4: cost(4)=25, path: 1->4
- ✓ 5: cost(5)=31, path: 1->5

Explore the node with the lowest cost: Node 4 has a cost of 25. Vertices to be explored from node 4: 2, 3, and 5

Now we are starting from the cost matrix at node 4 is:

Cost (4) = 25					
$\left[\begin{array}{ccccc} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 12 & \text{inf} & 11 & \text{inf} & 0 \\ 0 & 3 & \text{inf} & \text{inf} & 2 \\ \text{inf} & 3 & 12 & \text{inf} & 0 \\ 11 & 0 & 0 & \text{inf} & \text{inf} \end{array} \right]$					

➤ **Choose to go to vertex 2: Node 6 (path is 1->4->2)**

Cost of edge <4,2> is: $A(4,2) = 3$

Set row #4 = inf since we are reconsidering edge <4,2>

Set column #2 = inf since we are reconsidering edge <4,2> Set

$A(2,1) = \text{inf}$

The resulting cost matrix is:

$\left[\begin{array}{ccccc} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & 11 & \text{inf} & 0 \\ 0 & \text{inf} & \text{inf} & \text{inf} & 2 \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 11 & \text{inf} & 0 & \text{inf} & \text{inf} \end{array} \right]$
--

Reduce the matrix: Rows are reduced

Columns are reduced The

lower bound is: $RCL = 0$

The cost of going through node 2 is:

$\text{cost}(6) = \text{cost}(4) + RCL + A(4,2) = 25 + 0 + 3 = 28$

➤ **Chooseto go to vertex 3: Node7 (pathis 1->4->3)**

Costof edge<4,3>is: $A(4,3) = 12$

Setrow#4 =inf since weareconsideringedge<4,3>

Setcolumn#3=infsinceweareconsideringedge<4,3> Set

$A(3,1) = \text{inf}$

The resulting cost matrix is:

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 12 & \text{inf} & \text{inf} & \text{inf} & 0 \\ \text{inf} & 3 & \text{inf} & \text{inf} & 2 \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 11 & 0 & \text{inf} & \text{inf} & \text{inf} \end{bmatrix}$$

Reduce the matrix:

Reduce row #3: by 2:

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 12 & \text{inf} & \text{inf} & \text{inf} & 0 \\ \text{inf} & 1 & \text{inf} & \text{inf} & 0 \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 11 & 0 & \text{inf} & \text{inf} & \text{inf} \end{bmatrix}$$

Reduce column #1: by 11

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 1 & \text{inf} & \text{inf} & \text{inf} & 0 \\ \text{inf} & 1 & \text{inf} & \text{inf} & 0 \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 0 & 0 & \text{inf} & \text{inf} & \text{inf} \end{bmatrix}$$

The lower bound is: $RCL = 13$

So the RCL of node 7 (Considering vertex 3 from vertex 4) is: $\text{Cost}(7) = \text{cost}(4) + RCL + A(4,3) = 25 + 13 + 12 = 50$

➤ Choosetogoto vertex 5:Node 8(pathis 1->4->5)

Costofedge<4,5>is: $A(4,5) = 0$

Setrow#4 =inf since weareconsideringedge<4,5>

Setcolumn#5=infsinceweareconsideringedge<4,5> Set

$A(5,1) = \text{inf}$

The resulting cost matrix is:

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 12 & \text{inf} & 11 & \text{inf} & \text{inf} \\ 0 & 3 & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & 0 & 0 & \text{inf} & \text{inf} \end{bmatrix}$$

Reduce the matrix:

Reduced row 2 by 11

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 1 & \text{inf} & 0 & \text{inf} & \text{inf} \\ 0 & 3 & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & 0 & 0 & \text{inf} & \text{inf} \end{bmatrix}$$

Columns are reduced

The lower bound is: $RCL = 11$

So the cost of node 8 (Considering vertex 5 from vertex 4) is:

$\text{Cost}(8) = \text{cost}(4) + RCL + A(4,5) = 25 + 11 + 0 = 36$

In summary: So the live nodes we have so far are:

- ✓ 2: $\text{cost}(2) = 35$, path: 1->2
- ✓ 3: $\text{cost}(3) = 53$, path: 1->3
- ✓ 5: $\text{cost}(5) = 31$, path: 1->5
- ✓ 6: $\text{cost}(6) = 28$, path: 1->4->2
- ✓ 7: $\text{cost}(7) = 50$, path: 1->4->3
- ✓ 8: $\text{cost}(8) = 36$, path: 1->4->5
- Explore the node with the lowest cost: Node 6 has a cost of 28
- Vertices to be explored from node 6: 3 and 5
- Now we are starting from the cost matrix at node 6 is:

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & 11 & \text{inf} & 0 \\ 0 & \text{inf} & \text{inf} & \text{inf} & 2 \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 11 & \text{inf} & 0 & \text{inf} & \text{inf} \end{bmatrix}$$

➤ **Choosetogoto vertex3: Node9(pathis 1->4->2->3)**

Costof edge<2,3>is: $A(2,3) = 11$

Setrow#2 =inf since weareconsideringedge<2,3>

Setcolumn#3=infsinceweareconsideringedge<2,3> Set

$A(3,1) = \text{inf}$

The resulting cost matrix is:

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & 2 \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 11 & \text{inf} & \text{inf} & \text{inf} & \text{inf} \end{bmatrix}$$

Reducethematrix: Reducerow #3: by2

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & 0 \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 11 & \text{inf} & \text{inf} & \text{inf} & \text{inf} \end{bmatrix}$$

Reducecolumn #1: by11

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & 0 \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 0 & \text{inf} & \text{inf} & \text{inf} & \text{inf} \end{bmatrix}$$

The lower bound is: $\text{RCL} = 2 + 11 = 13$

So the cost of node9 (Considering vertex3 from vertex2) is: $\text{Cost}(9) = \text{cost}(6) + \text{RCL} + A(2,3) = 28 + 13 + 11 = 52$

➤ **Choosetogotovertex 5:Node10 (path is1->4->2->5)**

Costof edge<2,5>is: $A(2,5) = 0$

Setrow#2 =inf since weareconsideringedge<2,3>

Setcolumn#3=infsinceweareconsideringedge<2,3> Set

$A(5,1) = \text{inf}$

The resulting cost matrix is:

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 0 & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & 0 & \text{inf} & \text{inf} \end{bmatrix}$$

Reducethematrix: Rows reduced
Columnsreduced

The lower bound is: $\text{RCL} = 0$

Sothecostofnode10(Consideringvertex5fromvertex2)is:

$\text{Cost}(10) = \text{cost}(6) + \text{RCL} + A(2,3) = 28 + 0 + 0 = 28$

Insummary: **Sothelivenodeswehavesofarare:**

- ✓ 2: $\text{cost}(2)=35$, path:1->2
- ✓ 3: $\text{cost}(3)=53$, path:1->3
- ✓ 5: $\text{cost}(5)=31$, path:1->5
- ✓ 7: $\text{cost}(7)=50$, path:1->4->3
- ✓ 8: $\text{cost}(8)=36$, path:1->4->5
- ✓ 9: $\text{cost}(9)=52$, path: 1->4->2->3
- ✓ 10: $\text{cost}(2)=28$, path: 1->4->2->5
- Explorenodewiththelowest cost: Node10 has a cost of 28
- Verticestobe explored fromnode10: 3
- Nowwearestartingfromthecostmatrixat node 10is:

Cost (10)=28

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ 0 & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & 0 & \text{inf} & \text{inf} \end{bmatrix}$$

➤ **Choosetogo tovertex3: Node11(pathis1->4->2->5->3)**

Costof edge<5,3>is: $A(5,3) = 0$

Setrow#5 =inf since weareconsideringedge<5,3>

Setcolumn#3=infsinceweareconsideringedge<5,3> Set

$A(3,1) = \text{inf}$

Theresultingcostmatrixis:

$$\begin{bmatrix} \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \\ \text{inf} & \text{inf} & \text{inf} & \text{inf} & \text{inf} \end{bmatrix}$$

Reducethe matrix: Rows reduced

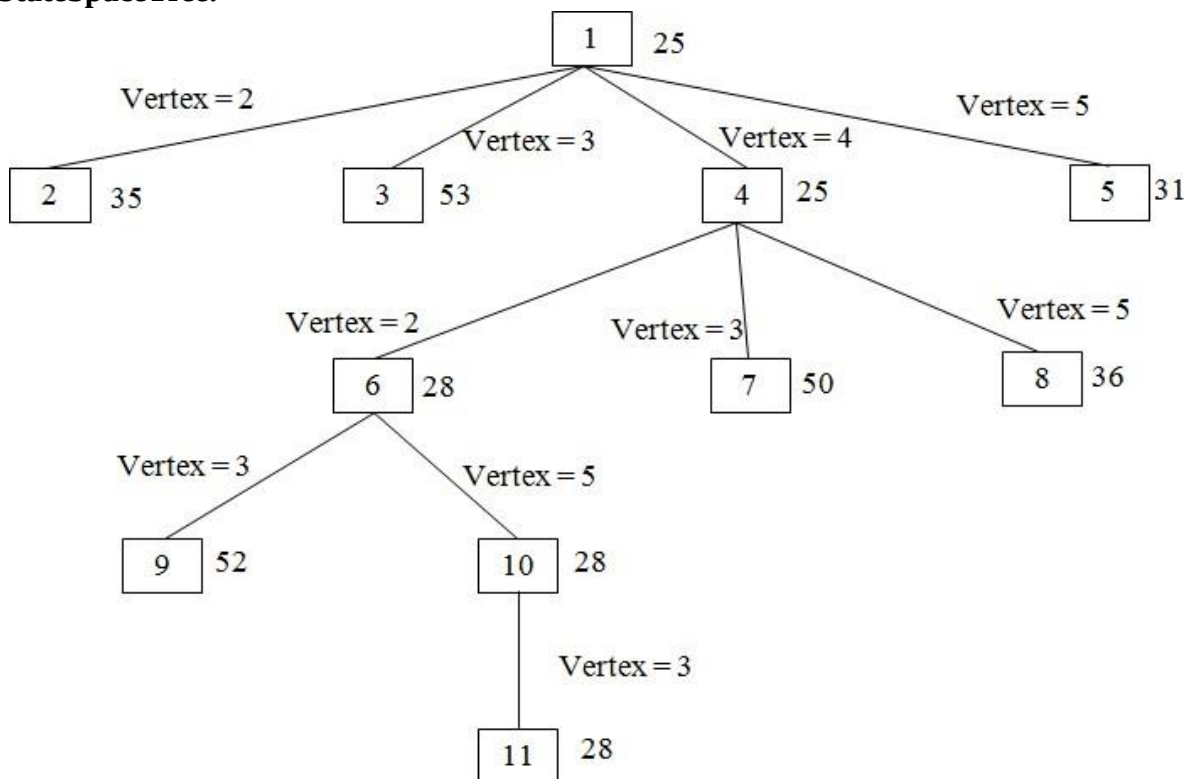
Columnsreduced

The lower bound is: $\text{RCL} = 0$

Sothecostofnode11(Consideringvertex5fromvertex3)is:

$\text{Cost}(11) = \text{cost}(10) + \text{RCL} + A(5,3) = 28 + 0 + 0 = 28$

StateSpaceTree:



O/1KnapsackProblem

What is Knapsack Problem: Knapsack problem is a problem in combinatorial optimization, Given a set of items, each with a mass & a value, determine the number of each item to include in a collection so that the total weight is less than or equal to a given limit & the total value is as large as possible.

O-1Knapsack Problem can formulate as. Let there be n items, Z_1 to Z_n where Z_i has value P_i & weight w_i . The maximum weight that can carry in the bag is M . All values and weights are non negative.

Maximize the sum of the values of the items in the knapsack, so that the sum of the weights must be less than the knapsack's capacity m .

The formula can be stated as

$$\begin{aligned} &\text{maximize } \sum_{1 \leq i \leq n} p_i x_i \\ &\text{subject to } \sum_{1 \leq i \leq n} w_i x_i \leq M \end{aligned}$$

$$x_i = 0 \text{ or } 1 \quad 1 \leq i \leq n$$

To solve 0/1 knapsack problem using B&B:

➤ Knapsack is a maximization problem

▪ Replace the objective function $\sum p_i x_i$ by the function $-\sum p_i x_i$ to make it into a minimization problem

▪ The modified knapsack problem is stated as

$$\begin{aligned} &\text{Minimize } -\sum_{i=1}^n p_i x_i \\ &\text{subject to } \sum_{i=1}^n w_i x_i < m, \\ &x_i \in \{0, 1\}, 1 \leq i \leq n \end{aligned}$$

➤ Fixed tuple sizes solution space:

○ Every leaf node in state space tree represents an answer for which

$$\sum_{1 \leq i \leq n} w_i x_i \leq m$$

is an answer node; other leaf nodes are infeasible

○ For optimal solution, define

$$c(x) = -\sum_{1 \leq i \leq n} p_i x_i$$

for every answer node x

➤ For infeasible leaf nodes, $c(x) = \infty$

➤ For non-leaf nodes

$$c(x) = \min\{c(\text{lchild}(x)), c(\text{rchild}(x))\}$$

➤ Define two functions $\hat{c}(x)$ and $u(x)$ such that for every node x ,

$$\hat{c}(x) \leq c(x) \leq u(x)$$

➤ Computing $\hat{c}(\cdot)$ and $u(\cdot)$

Let x be a node at level j , $1 \leq j \leq n + 1$

Cost of assignment: $-\sum_{1 \leq i < j} p_i x_i$

$$c(x) \leq -\sum_{1 \leq i < j} p_i x_i$$

We can use $u(x) = -\sum_{1 \leq i < j} p_i x_i$

Using $q = -\sum_{1 \leq i < j} p_i x_i$, an improved upper bound function $u(x)$ is

$$u(x) = \text{ubound}(q, \sum_{1 \leq i < j} w_i x_i, j - 1, m)$$

```
Algorithm ubound(cp, cw, k, m)
{
//Input:   cp: Current profit total
//Input:   cw: Current weight total
//Input:   k:  Index of last removed item
//Input: m:   Knapsack capacity
b=cp; c=cw;
for i:=k+1 to n do{
    if(c+w[i]≤m) then{
        c:=c+w[i]; b:=b-p[i];
    }
}
return b;
}
```

UNIT V:

NP-Hard and NP-Complete problems: Basic concepts, nondeterministic algorithms, NP- Hard and NP Complete classes, Cook's theorem.

Basic concepts:

NP: Nondeterministic Polynomial time

The problems have best algorithms for their solutions have "Computing times", that cluster into two groups

Group1	Group2
> Problems with solution time bound by a polynomial of a small degree. > It is also called "Tractable Algorithms" > Most Searching & Sorting algorithms are polynomial time algorithms > Ex: Ordered Search ($O(\log n)$), Polynomial evaluation $O(n)$ Sorting $O(n \log n)$	> Problems with solution times not bound by polynomial (simply non polynomial) > These are hard or intractable problems > None of the problems in this group has been solved by any polynomial time algorithm > Ex: Traveling Sales Person $O(n^2 2^n)$ Knapsack $O(2^{n/2})$

No one has been able to develop a polynomial time algorithm for any problem in the 2nd group (i.e., group 2)

So, it is compulsory to find algorithms whose computing times are greater than polynomial very quickly because such a vast amount of time to execute that even moderate size problems cannot be solved.

Theory of NP-Completeness:

Show that many of the problems with no polynomial time algorithms are computational time algorithms are computationally related.

There are two classes of non-polynomial time problems

1. NP-Hard
2. NP-Complete

NP Complete Problem: A problem that is NP-Complete can be solved in polynomial time if and only if (iff) all other NP-Complete problems can also be solved in polynomial time.

NP-Hard: Problem can be solved in polynomial time then all NP-Complete problems can be solved in polynomial time.

All NP-Complete problems are NP-Hard but some NP-Hard problems are not known to be NP-Complete.

Nondeterministic Algorithms:

Algorithms with the property that the result of every operation is uniquely defined are termed as deterministic algorithms. Such algorithms agree with the way programs are executed on a computer.

Algorithms which contain operations whose outcomes are not uniquely defined but are limited to a specified set of possibilities. Such algorithms are called nondeterministic algorithms.

The machine executing such operations is allowed to choose any one of these outcomes subject to a termination condition to be defined later.

To specify nondeterministic algorithms, there are 3 new functions.

Choice(S), arbitrarily chooses one of the elements of sets S

Failure () Signals an Unsuccessful completion

Success(). Signals a successful completion.

Example for NonDeterministicalgorithms:

<pre> Algorithm Search(x){ //Problem is to search an element x //output J, such that A[J]=x; or J=0 if x is not in A J:=Choice(1,n); if(A[J]:=x)then{ Write(J); Success(); } else{ write(0); failure(); } } </pre>	Whenever there is a set of choices that lead to a successful completion then one such set of choices is always made and the algorithm terminates. A Nondeterministic algorithm terminates unsuccessfully if and only if (iff) there exists no set of choices leading to a successful signal.
---	--

Nondeterministic Knapsack algorithm	
Algorithm DKP (p,w,n,m,r,x){ W:=0; P:=0; for i:=1 to n do { x[i]:=choice(0,1); W:=W+x[i]*w[i]; P:=P+x[i]*p[i]; } if ((W>m) or (P<r)) then Failure(); else Success(); }	p: given Profits w: given Weights n: Number of elements (number of items) m: Weight of bag limit P: Final Profit W: Final weight

The Classes NP-Hard & NP-Complete:

For measuring the complexity of an algorithm, we use the input length as the parameter. For example, an algorithm A is of polynomial complexity $O(p(n))$ such that the computing time of A is $O(p(n))$ for every input of size n.

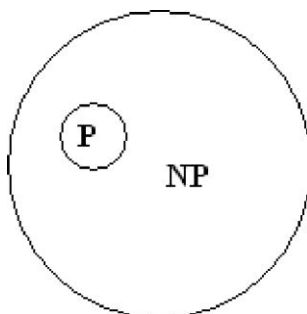
Decision problem/Decision algorithm: Any problem for which the answer is either zero or one is a decision problem. Any algorithm for a decision problem is termed a decision algorithm.

Optimization problem/Optimization algorithm: Any problem that involves the identification of an optimal (either minimum or maximum) value of a given cost function is known as an optimization problem. An optimization algorithm is used to solve an optimization problem.

P is the set of all decision problems solvable by deterministic algorithms in polynomial time.

NP is the set of all decision problems solvable by nondeterministic algorithms in polynomial time.

Since deterministic algorithms are just a special case of nondeterministic, by this we concluded that $P \subseteq NP$.



Commonly believed relationship between P & NP

The most famous unsolvable problem in Computer Science is Whether $P=NP$ or $P \neq NP$. In considering this problem, Cook formulated the following question.

If there is any single problem in NP, such that if we showed it to be in 'P' then that would imply that $P=NP$.

Cook answered this question with

Theorem: Satisfiability is in P if and only if (iff) $P=NP$

-) Notation of Reducibility

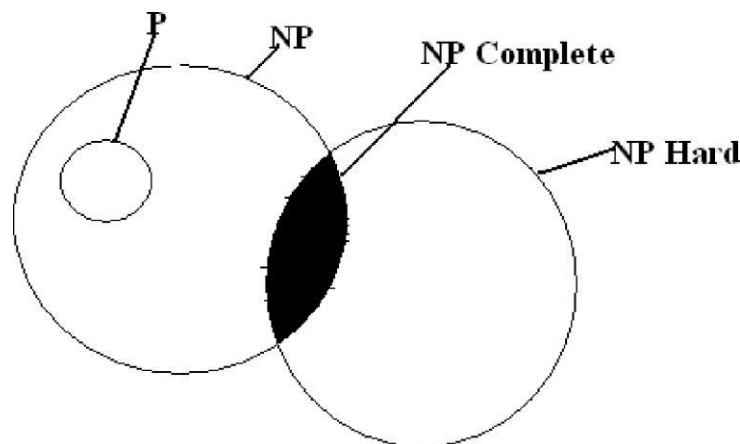
Let L_1 and L_2 be problems, Problem L_1 reduces to L_2 (written $L_1 \leq L_2$) iff there is a way to solve L_1 by a deterministic polynomial time algorithm using a deterministic algorithm that solves L_2 in polynomial time.

This implies that, if we have a polynomial time algorithm for L_2 , then we can solve L_1 in polynomial time.

Here $\leq$ is a transitive relation i.e., $L_1 \leq L_2$ and $L_2 \leq L_3$ then $L_1 \leq L_3$

A problem L is NP-Hard if and only if (iff) satisfiability reduces to L i.e., $\text{Satisfiability} \leq L$

A problem L is NP-Complete if and only if (iff) L is NP-Hard and $L \in NP$



Commonly believed relationship among P, NP, NP-Complete and NP-Hard

Most natural problems in NP are either in P or NP-complete.

Examples of NP-complete problems:

- > Packing problems: SET-PACKING, INDEPENDENT-SET.
- > Covering problems: SET-COVER, VERTEX-COVER.
- > Sequencing problems: HAMILTONIAN-CYCLE, TSP.
- > Partitioning problems: 3-COLOR, CLIQUE.
- > Constraint satisfaction problems: SAT, 3-SAT.
- > Numerical problems: SUBSET-SUM, PARTITION, KNAPSACK.

Cook's Theorem: States that satisfiability is in P if and only if $P=NP$. If $P=NP$ then satisfiability is in P.
If satisfiability is in P, then $P=NP$. To do this

> A-) Any polynomial time nondeterministic decision algorithm.

I-) Input of that algorithm

Then formula $Q(A, I)$, Such that Q is satisfiable iff A has a successful termination with Input I .

> If the length of I is ' n ' and the time complexity of A is $p(n)$ for some polynomial $p()$ then length of Q is $O(p^3(n) \log n) = O(p^4(n))$

The time needed to construct Q is also $O(p^3(n) \log n)$.

> A deterministic algorithm ' Z ' to determine the outcome of ' A ' on any input ' I '

Algorithm Z computes ' Q ' and then uses a deterministic algorithm for the satisfiability problem to determine whether ' Q ' is satisfiable.

> If $O(q(m))$ is the time needed to determine whether a formula of length ' m ' is satisfiable then the complexity of ' Z ' is $O(p^3(n) \log n + q(p^3(n) \log n))$.

> If satisfiability is ' p ', then ' $q(m)$ ' is a polynomial function of ' m ' and the complexity of ' Z ' becomes ' $O(r(n))$ ' for some polynomial ' $r()$ '.

> Hence, if satisfiability is in p , then for every nondeterministic algorithm A in NP , we can obtain a deterministic Z in p .

By this we show that satisfiability is in p then $P=NP$