mid term answers PRACTICE QUESTIONS FOR ETE

1. Analyse how different hashing methods can be applied to optimize search and sort operations.

Search Optimization with Hashing:

- **Hash Table:** Imagine a data structure like an indexed filing cabinet. Each item has a unique "key" (like an employee ID). A hash function takes that key and calculates an "address" (like a cabinet number) where the item is stored.
- **Fast Lookup:** By calculating the address for the item you're searching for, you can jump directly to that location in the hash table, bypassing the need to scan through the entire dataset. This is significantly faster than linear search, especially for large datasets.

Types of Hashing and Considerations:

- Collision: Multiple keys might map to the same address (collision). Techniques like separate chaining (storing multiple items at the same address) or open addressing (probing for the next available address) can handle collisions but add some overhead.
- **Hash Function Choice:** A good hash function should distribute keys uniformly across the available addresses to minimize collisions and optimize search time.
- 2. Analyse the performance and application scenarios of binomial and Fibonacci heaps.

Heap Type	Insertions/Merges	Other Operations	Applications		
Binomial	Faster than Binary Heaps	Slower than Fibonacci Heaps	Frequent merging/large insertions		
	Amortized constant time	-	High-speed insertions & merges (Dijkstra's algorithm)		
Investigate the disjoint set union data structure and its operations, evaluating its					

3. Investigate the disjoint set union data structure and its operations, evaluating its efficiency in different applications.

Disjoint Set Union (DSU) tracks separate groups (sets) of elements. It lets you:

- **Merge groups:** Combine two separate sets into one.
- Check group membership: See which group an element belongs to.

Super fast: DSU finds groups and merges them in near-constant time (average) on large datasets, making it very efficient.

Used for:

- Finding minimum spanning trees (electrical wires)
- Grouping friends in a social network
- Modeling connected areas (like fire spread)

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4. Compare and contrast Depth-First Search (DFS) and Breadth-First Search (BFS) in various contexts.

DFS vs. BFS

Feature	DFS	BFS
Traversal Strategy	Goes deep into one branch before backtracking	Expands outward level by level
Finding Specific	Slower for distant nodes, faster	Faster, especially for nodes
Node	for close ones	near the start
Connectivity Check	Efficient	Less efficient
Cycle Detection	More suitable due to revisiting nodes	Less suitable
Shortest Path (unweighted)	Not guaranteed	Guaranteed
Data Structure	Stack (LIFO)	Queue (FIFO)
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5. Evaluate shortest path algorithms, minimum spanning tree algorithms, and their applications in real-world problems.

Shortest path and minimum spanning tree (MST) algorithms are crucial in computer science with significant real-world applications.

Shortest Path Algorithms

- 1. **Dijkstra's Algorithm**: Finds shortest paths in weighted graphs with non-negative weights. Applications: GPS navigation, network routing, transportation planning.
- 2. **Bellman-Ford Algorithm**: Handles graphs with negative weights. Applications: Currency arbitrage, telecommunications routing.
- 3. **A* Algorithm**: Uses heuristics for efficient pathfinding. Applications: Video games, robotic motion planning.

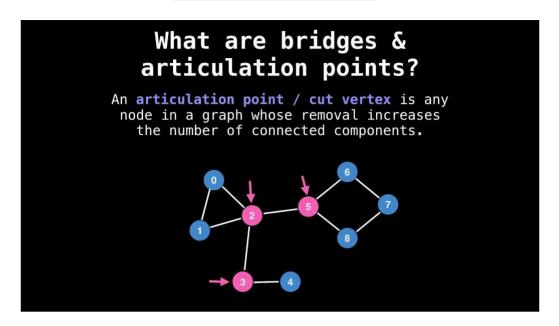
MST Algorithms

- 1. **Kruskal's Algorithm**: Builds MST by adding smallest edges without cycles. Applications: Network design, clustering.
- 2. **Prim's Algorithm**: Grows MST from an arbitrary vertex. Applications: Communication networks, road network optimization.

Applications

- 1. **Telecommunications**: Optimizing data routing and network costs.
- 2. **Transportation and Logistics**: Route optimization and infrastructure planning.
- 3. **Urban Planning**: Traffic flow optimization and utility planning.
- 4. **Computer Networks**: Efficient data transfer and network topology design.
- Investigate the significance of articulation points and bridges in network design and reliability.

mid term answers



Significance

1. Articulation Points:

- Critical nodes; their failure disrupts network connectivity.
- Enhances network robustness.

2. Bridges:

- o Critical connections; their failure disconnects parts of the network.
- Ensures path redundancy.

Applications

- 1. **Communication Networks**: Maintain connectivity and prevent partitioning.
- 2. **Transportation Networks**: Protect critical routes and ensure traffic flow.
- 3. **Utility Networks**: Safeguard service continuity.

7 Examine Strasson's matrix multiplication algorithm and compare it with conventional methods.

Strassen's Matrix Multiplication Algorithm

- Overview: Efficient algorithm that reduces matrix multiplication time complexity.
- **Key Idea**: Divides matrices into submatrices and performs only 7 multiplications instead of 8.
- Time Complexity: $O(nlog \frac{fo}{27}) \approx O(n2.81)O(n^{\{\log_2\{7\}\}}) \cdot O(n^{2.81})O(nlog 27) \approx O(n2.81)$
- **Best For**: Large matrices where computational efficiency is critical.
- **Drawbacks**: More complex and requires more memory.

Conventional Matrix Multiplication

- Overview: Standard method using three nested loops.
- **Time Complexity**: O(n3)O(n^3)O(n3)
- **Best For**: Small matrices due to simplicity and lower overhead.

mid term answers

• Advantages: Simpler implementation, less memory usage.

Comparison

- 1. Time Complexity:
 - o **Conventional**: O(n3)O(n^3)O(n3)
 - \circ Strassen's: $O(n2.81)O(n^{2.81})O(n2.81)$
- 2. Efficiency:
 - o Strassen's: Faster for large matrices.
 - Conventional: More practical for small matrices.
- 3. Memory Usage:
 - o Strassen's: Higher memory usage.
 - o Conventional: Lower memory usage.
- 4. Practical Use:
 - o **Strassen's**: Large-scale applications, theoretical computer science.
 - o **Conventional**: Simpler, straightforward for smaller tasks.
 - 1. Divide matrices A and B into submatrices A_{11} , A_{12} , A_{21} , A_{22} and B_{11} , B_{12} , B_{21} , B_{22} .
 - 2. Compute the 7 products:

$$egin{aligned} M_1 &= (A_{11} + A_{22})(B_{11} + B_{22}) \ M_2 &= (A_{21} + A_{22})B_{11} \ M_3 &= A_{11}(B_{12} - B_{22}) \ M_4 &= A_{22}(B_{21} - B_{11}) \ M_5 &= (A_{11} + A_{12})B_{22} \ M_6 &= (A_{21} - A_{11})(B_{11} + B_{12}) \ M_7 &= (A_{12} - A_{22})(B_{21} + B_{22}) \end{aligned}$$

3. Combine these products to get the resulting submatrices of the product matrix:

$$C_{11} = M_1 + M_4 - M_5 + M_7 \ C_{12} = M_3 + M_5 \ C_{21} = M_2 + M_4 \ C_{22} = M_1 - M_2 + M_3 + M_6$$

8 Analyse the time complexity and efficiency of algorithms based on divide and conquer, such as counting inversions and finding the closest pair of points.

Attribute	Counting Inversions	Closest Pair of Points
Algorithm	Counting Inversions	Closest Pair of Points
Problem	Counting inversions in an array	Finding the closest pair of points

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Key IdeaDivide the array, count inversions, Divide points, find closest pairs, merge results

Divide the array, count inversions, Divide points, find closest pairs,

Time $O(n\log f_0)n)O(n \log n)O(n\log n)$ $O(n\log f_0)n)O(n \log n)O(n\log n)$

Efficiency Efficient for large arrays Efficient for large point sets

Best Use Sorting-related tasks, data Geometric problems, clustering, analysis, algorithmic competitions computational geometry

9 Critically evaluate the effectiveness of universal hashing in various scenarios.

Universal hashing is a technique that improves the performance of hash functions by selecting one at random from a family of hash functions, minimizing the chance of collisions. Here's a summary of its effectiveness in various scenarios:

- 1. **General-purpose Hashing**: Highly effective for unknown input distributions, reducing worst-case collisions.
- 2. **Cryptographic Applications**: Not suitable due to lack of required security properties.
- 3. **Hash Tables and Data Structures**: Improves performance and reduces collisions, ensuring consistent O(1) operations.
- 4. **Adversarial Inputs**: Effective against collision attacks by making it hard to predict the hash function.
- 5. **Real-time Systems**: Improves performance predictability but may introduce slight delays due to random selection.
- 6. **Distributed Systems**: Enhances load balancing by ensuring even key distribution across nodes.
- 7. **Memory-limited Environments**: Additional memory and computational overhead may be a drawback.
- Judge the suitability of hashing methods for optimization problems and justify the chosen method.

• Simple Hashing

- **Suitability**: Small datasets, predictable distributions.
- **Pros**: Easy to implement, fast.
- Cons: Prone to collisions with larger or unpredictable datasets.

• Universal Hashing

• **Suitability**: Unknown or adversarial input distributions.

mid term answers

- **Pros**: Reduces collision probability, robust against worst-case scenarios.
- Cons: Slightly more complex due to random selection of hash functions.

• Perfect Hashing

- Suitability: Static datasets with known keys.
- **Pros**: O(1) lookup time without collisions.
- Cons: Only for static data; not suitable for dynamic datasets.

• Cuckoo Hashing

- Suitability: High-performance insertions and lookups, real-time systems.
- **Pros**: Guaranteed O(1) lookup time, handles high load factors.
- **Cons**: More complex to implement.

• Consistent Hashing

- **Suitability**: Distributed systems, load balancing.
- **Pros**: Minimizes remapping during resizing, balanced load distribution.
- **Cons**: Slightly more complex.

• Dynamic Perfect Hashing

- Suitability: Changing datasets over time.
- **Pros**: Efficient operations, adapts to dataset changes.
- **Cons**: Complex to implement.

Judge the effectiveness of shortest path algorithms and minimum spanning tree algorithms in solving real-world problems.

Shortest Path Algorithms

1. Dijkstra's Algorithm

- o **Pros**: Efficient for non-negative weighted graphs, used in network routing.
- o Cons: Ineffective with negative weights, slower for large graphs.

2. Bellman-Ford Algorithm

- o **Pros**: Handles negative weights, detects negative cycles, used in financial modeling.
- o Cons: Slower than Dijkstra's for non-negative weights.

3. A Algorithm*

- o **Pros**: Fast with good heuristics, used in AI and robotics for pathfinding.
- o **Cons**: Heuristic-dependent, unsuitable for negative weights.

4. Floyd-Warshall Algorithm

- **Pros**: Finds shortest paths for all pairs in dense graphs, simple implementation.
- Cons: High time complexity, impractical for large graphs.

Minimum Spanning Tree (MST) Algorithms

mid term answers

1. Kruskal's Algorithm

- o **Pros**: Simple, effective for sparse graphs, used in network design.
- Cons: Sorting edges can be time-consuming for large graphs.

2. Prim's Algorithm

- Pros: Efficient for dense graphs, used with priority queues in network construction.
- o **Cons**: Less intuitive in some applications.

3. Borůvka's Algorithm

- o **Pros**: Suitable for parallel processing, handles large graphs.
- o **Cons**: Complex implementation.

Real-World Applications

- **Shortest Path Algorithms**: Used in network routing, GPS navigation, and transportation logistics for optimal route planning.
- **Minimum Spanning Tree Algorithms**: Applied in network design, data clustering, and electrical grid infrastructure to minimize costs.

Justify the choice of graph algorithms based on problem constraints and requirements.

Shortest Path Algorithms

1. Dijkstra's Algorithm

- o **Use When**: You have a graph with non-negative weights.
- Why: It's efficient $(O(V^2))$ or $O(E + V \log V)$ with a priority queue) and straightforward for such graphs.
- **Example**: Finding the shortest route in a city map without negative distances.

2. Bellman-Ford Algorithm

- o Use When: You need to handle negative weights or detect negative cycles.
- Why: It works with negative weights and can identify negative cycles, though it's slower (O(VE)).
- o **Example**: Financial models where costs can decrease.

3. A Algorithm*

- o **Use When**: You need fast pathfinding with a known heuristic.
- Why: It uses heuristics to speed up the search, making it efficient in practice.
- o **Example**: GPS navigation systems considering real-time traffic.

4. Floyd-Warshall Algorithm

- Use When: You need shortest paths between all pairs of nodes in a dense graph.
- Why: It's simple and handles all pairs but has high time complexity $(O(V^3))$.
- Example: Analyzing all possible routes in a small, densely connected network.

Minimum Spanning Tree (MST) Algorithms

1. Kruskal's Algorithm

mid term answers

- o **Use When**: You have a sparse graph and can sort edges efficiently.
- Why: It's simple and works well for sparse graphs.
- o **Example**: Designing a cost-effective network with fewer connections.

2. Prim's Algorithm

- o **Use When**: You have a dense graph or prefer adjacency lists.
- Why: It's efficient for dense graphs and works well with priority queues.
- o **Example**: Building a network with many connections.

3. Borůvka's Algorithm

- Use When: You can use parallel processing and need to handle very large graphs.
- Why: It's suitable for parallel execution and large datasets.
- o **Example**: Large-scale network design requiring parallel computation.

Summary

- **Dijkstra** for non-negative weights.
- **Bellman-Ford** for negative weights.
- **A*** for fast pathfinding with heuristics.
- Floyd-Warshall for all-pairs shortest paths in dense graphs.
- Kruskal for sparse MSTs.
- **Prim** for dense MSTs.
- Borůvka for parallel processing and large graphs.
- 12 Create a system that dynamically selects the appropriate search method based on the dataset characteristics.

1. Input Analysis:

- o **Graph Type**: Determine if the graph is dense or sparse.
- o **Edge Weights**: Check if the graph contains negative weights.
- o **Path Requirements**: Identify if you need a single-source shortest path or all-pairs shortest paths.

2. Decision Criteria:

- Non-negative Weights:
 - Single Source: Use Dijkstra's algorithm.
 - All Pairs: Use Floyd-Warshall if the graph is dense.
- Negative Weights:
 - **Single Source**: Use Bellman-Ford.
- o **Heuristic-Based Search**: Use A* if a good heuristic is available.
- Minimum Spanning Tree (MST):
 - **Sparse Graph**: Use Kruskal's algorithm.
 - **Dense Graph**: Use Prim's algorithm.
 - **Parallel Processing**: Use Borůvka's algorithm.
- 3. **Dynamic Selection System** (Pseudocode):

```
python
Copy code
def select_algorithm(graph, heuristic=None):
    if is_mst_problem(graph):
        if is_dense(graph):
            return "Prim's Algorithm"
```

mid term answers

```
elif is parallel processing available():
            return "Borůvka's Algorithm"
           return "Kruskal's Algorithm"
    elif is shortest path problem(graph):
        if has negative weights (graph):
            return "Bellman-Ford Algorithm"
        elif heuristic:
            return "A* Algorithm"
        elif is all pairs required (graph):
           return "Floyd-Warshall Algorithm" if is dense(graph)
else "Johnson's Algorithm"
        else:
           return "Dijkstra's Algorithm"
        return "Algorithm not determined"
# Example Usage
selected algorithm = select algorithm(graph,
heuristic=some heuristic)
print(f"Selected algorithm: {selected algorithm}")
```

Design a robust hashing system that minimizes collisions and optimizes search and sort operations.

Designing a robust hashing system involves several key elements:

- 1. **Hash Function Selection**: Use a strong hash function like SHA-256 for good distribution of hash values.
- 2. **Collision Handling**: Implement techniques like chaining or open addressing to manage collisions efficiently.
- 3. **Load Balancing**: Use consistent hashing and dynamic resizing to balance the load on the hash table.
- 4. **Indexing and Sorting**: Employ hash indices or hash trees for fast search operations, and use sorting algorithms for sorting hash table entries.
- 5. **Parallelism and Concurrency**: Support concurrent access with locking mechanisms or lock-free data structures.
- 6. **Memory Management**: Optimize memory usage and consider caching techniques for performance improvement.
- 7. **Error Handling and Resilience**: Implement robust error handling and monitoring to ensure system integrity.
- 8. **Testing and Benchmarking**: Conduct thorough testing and benchmarking to evaluate performance metrics and identify areas for optimization.
- Design greedy algorithms tailored to solve specific optimization problems like activity selection and task scheduling.

Greedy algorithms provide efficient solutions to specific optimization problems such as activity selection and task scheduling.

• Activity Selection:

mid term answers

- o Greedily selecting activities based on their finish times maximizes the number of non-overlapping activities.
- The algorithm sorts activities by finish time and selects those that do not overlap with previously selected ones.

• Task Scheduling:

- Maximizing total profit by scheduling tasks within deadlines involves sorting tasks by profit and greedily selecting tasks for the latest available slots before their deadlines.
- The algorithm iterates through tasks, scheduling them in available slots if they do not conflict with previously scheduled tasks.

Greedy algorithms make locally optimal choices at each step, leading to efficient solutions. However, they may not always yield globally optimal results and require careful consideration of problem constraints to ensure desired optimization.

Apply the binary search technique to find the first occurrence of a number in a sorted array.

binary search technique to find the first occurrence of a number in a sorted array:

```
python
Copy code
def binary_search_first_occurrence(arr, target):
   low = 0
   high = len(arr) - 1
   result = -1 # Initialize result to -1 (not found)
   while low <= high:
        mid = low + (high - low) // 2
        if arr[mid] == target:
            result = mid # Update result to current index
            high = mid - 1  # Search for first occurrence on the
left side
        elif arr[mid] < target:</pre>
           low = mid + 1
        else:
           high = mid - 1
   return result
# Example usage
arr = [1, 2, 2, 2, 3, 4, 5]
target = 2
first occurrence = binary search first occurrence(arr, target)
print("First occurrence index:", first occurrence)
```

15 Apply the greedy technique to solve the activity selection problem.

mid term answers

```
def activity selection(start times, finish times):
  n = len(start_times)
  activities = [] # List to store selected activities
  activities.append(0) # Add the first activity (assuming sorted by finish times)
  prev finish = finish times[0]
  for i in range(1, n):
     if start times[i] >= prev finish:
       activities.append(i)
       prev finish = finish times[i]
  return activities
# Example usage
start\_times = [1, 3, 0, 5, 3, 5, 6, 8, 8, 2, 12]
finish_times = [4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14]
selected_activities = activity_selection(start_times, finish_times)
print("Selected activities:", selected_activities)
How would you apply stack operations to evaluate a postfix expression? Write
the function in C/C++/Java/Python.
def evaluate_postfix(expression):
  stack = []
  # Function to perform arithmetic operations
  def apply operator(op, operand1, operand2):
     if op == '+':
       return operand1 + operand2
     elif op == '-':
       return operand1 - operand2
     elif op == '*':
       return operand1 * operand2
     elif op == '/':
       return operand1 / operand2 # Assuming division is floating-point
  for token in expression.split():
     if token.isdigit():
       stack.append(int(token))
     else:
       operand2 = stack.pop()
       operand1 = stack.pop()
       result = apply_operator(token, operand1, operand2)
       stack.append(result)
  return stack.pop()
# Example usage
postfix expression = "3 4 + 2 *"
```

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mid term answers

result = evaluate_postfix(postfix_expression)
print("Result:", result)

Apply binary search tree operations to insert and find an element. Write the function in C/C++/Java/Python.

```
class Node:
   def __init__(self, key):
        \overline{\text{self.key}} = \text{key}
        self.left = None
        self.right = None
class BinarySearchTree:
    def init (self):
        \overline{\text{self.root}} = \text{None}
    def insert(self, key):
        self.root = self. insert recursive(self.root, key)
    def insert recursive (self, root, key):
        if root is None:
            return Node (key)
        if key < root.key:
            root.left = self. insert recursive(root.left, key)
        elif key > root.key:
             root.right = self. insert recursive(root.right, key)
        return root
    def find(self, key):
        return self. find recursive(self.root, key)
    def find recursive (self, root, key):
        if root is None or root.key == key:
            return root
        if key < root.key:</pre>
            return self. find recursive(root.left, key)
            return self. find recursive(root.right, key)
# Example usage
bst = BinarySearchTree()
bst.insert(50)
bst.insert(30)
bst.insert(20)
bst.insert(40)
bst.insert(70)
bst.insert(60)
bst.insert(80)
found node = bst.find(40)
if found node:
    print("Element found:", found node.key)
    print("Element not found")
```

Advanced Algorithmic Problem Solving mid term answers

Use BFS to implement a level order traversal of a binary tree. Write the function in C/C++/Java/Python.

```
class TreeNode:
  def __init__(self, val=0, left=None, right=None):
    self.val = val
    self.left = left
    self.right = right
def level_order_traversal(root):
  if root is None:
    return []
  result = []
  queue = [root]
  while queue:
    level = []
    level_size = len(queue)
    for _ in range(level_size):
       node = queue.pop(0)
       level.append(node.val)
       if node.left:
         queue.append(node.left)
       if node.right:
         queue.append(node.right)
    result.append(level)
  return result
# Example usage
root = TreeNode(1)
root.left = TreeNode(2)
root.right = TreeNode(3)
root.left.left = TreeNode(4)
root.left.right = TreeNode(5)
root.right.left = TreeNode(6)
root.right.right = TreeNode(7)
print("Level Order Traversal:", level_order_traversal(root))
```

Write a program that uses divide and conquer to find the closest pair of points in a 2D plane.

mid term answers

```
import math
class Point:
  def __init__(self, x, y):
     self.x = x
     self.y = y
def distance(point1, point2):
  return math.sqrt((point1.x - point2.x) ** 2 + (point1.y - point2.y) ** 2)
def brute force closest pair(points):
  min distance = float('inf')
  closest_pair = None
  for i in range(len(points)):
     for j in range(i + 1, len(points)):
       dist = distance(points[i], points[i])
       if dist < min_distance:
          min distance = dist
          closest_pair = (points[i], points[i])
  return min_distance, closest_pair
def closest pair divide conquer(points):
  sorted points = sorted(points, key=lambda point: point.x)
  return closest_pair_recursive(sorted_points)
def closest_pair_recursive(sorted_points):
  n = len(sorted\_points)
  if n \le 3:
     return brute_force_closest_pair(sorted_points)
  mid = n // 2
  left closest, left pair = closest pair recursive(sorted points[:mid])
  right closest, right pair = closest pair recursive(sorted points[mid:])
  min_dist = min(left_closest, right_closest)
  # Check for points crossing the midpoint
  strip = []
  for point in sorted_points:
     if abs(point.x - sorted_points[mid].x) < min_dist:
       strip.append(point)
  strip_closest, strip_pair = closest_pair_strip(strip, min_dist)
  if strip closest < min dist:
     return strip_closest, strip_pair
  elif left_closest < right_closest:</pre>
```

mid term answers

```
return left closest, left pair
     return right_closest, right_pair
def closest_pair_strip(strip, min_dist):
  min distance = min dist
  closest_pair = None
  strip.sort(key=lambda point: point.y)
  for i in range(len(strip)):
     for j in range(i + 1, len(strip)):
       if strip[j].y - strip[i].y < min_distance:
          dist = distance(strip[i], strip[j])
          if dist < min distance:
            min distance = dist
            closest_pair = (strip[i], strip[j])
       else:
          break
  return min_distance, closest_pair
# Example usage
points = [Point(2, 3), Point(12, 30), Point(40, 50), Point(5, 1), Point(12, 10),
Point(3, 4)
closest_distance, closest_pair = closest_pair_divide_conquer(points)
print("Closest Distance:", closest_distance)
print("Closest Pair:", closest_pair)
Write a program to implement dynamic programming to solve the 0/1 knapsack
problem and analyse the memory usage.
def knapsack 01(values, weights, capacity):
  n = len(values)
  dp = [[0] * (capacity + 1) for _ in range(n + 1)]
  for i in range(1, n + 1):
     for w in range(1, capacity + 1):
       if weights[i - 1] <= w:
          dp[i][w] = max(values[i-1] + dp[i-1][w - weights[i-1]], dp[i-1][w]
1][w]
          dp[i][w] = dp[i - 1][w]
  return dp[n][capacity]
# Example usage and memory analysis
values = [60, 100, 120]
weights = [10, 20, 30]
capacity = 50
```

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```
max_value = knapsack_01(values, weights, capacity)
print("Maximum value in knapsack:", max_value)

# Memory analysis
import sys
size_of_dp = sys.getsizeof(dp)
print("Size of DP table:", size_of_dp, "bytes")
```

Evaluate the efficiency of using a sliding window technique for a given dataset of temperature readings over brute force methods.

sliding window technique with brute force methods for analyzing temperature readings:

Aspect	Sliding Window Technique	Brute Force Methods
Time Complexity	$O(n)$ or $O(n \log n)$	O(n^2) or higher
Space Complexity	O(1) or O(n)	Higher than O(n)
Practical Efficiency	Efficient for large datasets and real-time processing	Less efficient for large datasets, more suitable for smaller datasets
Usage	Widely used in real-time data processing and window-based analytics	Used in scenarios with manageable problem sizes or smaller datasets

Given a number n, find sum of first *n* natural numbers. To calculate the sum, we will use a recursive function recur_sum().

```
def recur_sum(n):
    if n <= 0:
        return 0
    else:
        return n + recur_sum(n - 1)

# Example usage
n = 5
sum_of_natural_numbers = recur_sum(n)
print("Sum of first", n, "natural numbers:", sum_of_natural_numbers)</pre>
```

Implement a recursive algorithm to solve the Tower of Hanoi problem. Find its complexity also.

```
def tower_of_hanoi(n, source, target, auxiliary):
    if n == 1:
        print(f"Move disk 1 from {source} to {target}")
        return
    tower_of_hanoi(n-1, source, auxiliary, target)
        print(f"Move disk {n} from {source} to {target}")
        tower_of_hanoi(n-1, auxiliary, target, source)

# Example usage
    n = 3
    tower_of_hanoi(n, 'A', 'C', 'B')
```

he time complexity of the Tower of Hanoi problem using this recursive algorithm is $O(2^n)$

Given a Binary Search Tree and a node value X, find if the node with value X is present in the BST or not.

```
class TreeNode:
  def __init__(self, key):
     self.key = key
     self.left = None
     self.right = None
def search_bst(root, x):
  if root is None or root.key == x:
    return root is not None
  if x < root.key:
     return search_bst(root.left, x)
  else:
    return search_bst(root.right, x)
# Example usage
root = TreeNode(50)
root.left = TreeNode(30)
root.right = TreeNode(70)
root.left.left = TreeNode(20)
root.left.right = TreeNode(40)
root.right.left = TreeNode(60)
root.right.right = TreeNode(80)
x = 40
if search_bst(root, x):
  print(f"Node with value {x} is present in the BST.")
  print(f"Node with value \{x\} is not present in the BST.")
```

```
class TreeNode:
  def __init__(self, key):
     self.key = key
     self.left = None
     self.right = None
def in order traversal(root, result):
  if root is not None:
     in_order_traversal(root.left, result)
     result.append(root.key)
    in order traversal(root.right, result)
def find_common_nodes(root1, root2):
  common_nodes = []
  inorder_list1 = []
  inorder list2 = []
  in_order_traversal(root1, inorder_list1)
  in_order_traversal(root2, inorder_list2)
  i, j = 0, 0
  while i < len(inorder\_list1) and j < len(inorder\_list2):
     if inorder_list1[i] == inorder_list2[j]:
       common_nodes.append(inorder_list1[i])
       i += 1
       i += 1
     elif inorder_list1[i] < inorder_list2[j]:
       i += 1
     else:
       i += 1
  return common nodes
# Example usage
root1 = TreeNode(50)
root1.left = TreeNode(30)
root1.right = TreeNode(70)
root1.left.left = TreeNode(20)
root1.left.right = TreeNode(40)
root1.right.left = TreeNode(60)
root1.right.right = TreeNode(80)
root2 = TreeNode(50)
root2.left = TreeNode(30)
root2.right = TreeNode(70)
root2.left.left = TreeNode(20)
```

```
root2.left.right = TreeNode(40)
root2.right.left = TreeNode(60)
root2.right.right = TreeNode(90) # Different from root1
common nodes = find common nodes(root1, root2)
print("Common nodes in both BSTs:", common nodes)
Design and implement a version of quicksort that randomly chooses pivot
elements. Calculate the time and space complexity of algorithm.
import random
def randomized_quicksort(arr):
  if len(arr) \le 1:
    return arr
  pivot = random.choice(arr)
  left = [x for x in arr if x < pivot]
  equal = [x \text{ for } x \text{ in arr if } x == pivot]
  right = [x \text{ for } x \text{ in arr if } x > pivot]
  return randomized_quicksort(left) + equal + randomized_quicksort(right)
# Example usage
arr = [3, 6, 8, 10, 1, 2, 1]
sorted arr = randomized quicksort(arr)
print("Sorted array:", sorted_arr)
Design and implement an efficient algorithm to merge k sorted arrays.
import heapq
def merge_k_sorted_arrays(arrays):
  result = []
  heap = [(array[0], i, 0)] for i, array in enumerate(arrays) if array [(array[0], i, 0)] # (value,
array_index, element_index)
  heapq.heapify(heap)
  while heap:
     val, arr_idx, elem_idx = heapq.heappop(heap)
    result.append(val)
     if elem_idx + 1 < len(arrays[arr_idx]):
       next_val = arrays[arr_idx][elem_idx + 1]
       heapq.heappush(heap, (next val, arr idx, elem idx + 1))
  return result
# Example usage
arrays = [[1, 3, 5], [2, 4, 6], [0, 7, 8, 9]]
```

merged_array = merge_k_sorted_arrays(arrays)
print("Merged sorted array:", merged_array)

26

27

Given two strings str1 & str 2 of length n & m respectively, find the length of the longest subsequence present in both. A subsequence is a sequence that can be derived from the given string by deleting some or no elements without changing the order of the remaining elements. For example, "abe" is a subsequence of "abcde". Example :Input: n = 6, str1 = ABCDGH and m = 6, str2 = AEDFHR Output: 3 Explanation: LCS for input strings "ABCDGH" and "AEDFHR" is "ADH" of length 3.

```
def longest_common_subsequence(str1, str2):
    n = len(str1)
    m = len(str2)
    dp = [[0] * (m + 1) for _ in range(n + 1)]

for i in range(1, n + 1):
    for j in range(1, m + 1):
        if str1[i - 1] == str2[j - 1]:
            dp[i][j] = dp[i - 1][j - 1] + 1
        else:
            dp[i][j] = max(dp[i - 1][j], dp[i][j - 1])

return dp[n][m]

# Example usage
str1 = "ABCDGH"
str2 = "AEDFHR"
result = longest_common_subsequence(str1, str2)
print("Length of longest common subsequence:", result)
```

For the given example input, the output will be 3, which is the length of the longest common subsequence "ADH" between "ABCDGH" and "AEDFHR".

You are given an amount denoted by **value**. You are also given an array of coins. The **array** contains the **denominations** of the given coins. You need to find the **minimum number of coins** to make the change for **value** using the coins of given denominations. Also, keep in mind that you have **infinite supply** of the coins. Input:

```
value = 10
numberOfCoins = 4
coins[] = {2 5 3 6}
Output: 2

def min_coins(value, coins):
    n = len(coins)
    dp = [float('inf')] * (value + 1)
    dp[0] = 0

for i in range(1, value + 1):
    for j in range(n):
```

```
if coins[j] \le i:
                    dp[i] = min(dp[i], dp[i - coins[j]] + 1)
            return dp[value]
          # Example usage
          value = 10
          coins = [2, 5, 3, 6]
          result = min coins(value, coins)
          print("Minimum number of coins required:", result)
30
          Hashing is very useful to keep track of the frequency of the elements in a list.
          You are given an array of integers. You need to print the count of non-repeated
          elements in the array. Example: Input:1 1 2 2 3 3 4 5 6 7 Output:4
          def count_non_repeated_elements(arr):
            frequency = {}
            non\_repeated\_count = 0
            for num in arr:
               frequency[num] = frequency.get(num, 0) + 1
            for key, value in frequency.items():
               if value == 1:
                 non_repeated_count += 1
            return non_repeated_count
          # Example usage
          arr = [1, 1, 2, 2, 3, 3, 4, 5, 6, 7]
          result = count non repeated elements(arr)
          print("Count of non-repeated elements:", result)
31
          Given two arrays a[] and b[] of size n and m respectively. The task is to find the
          number of elements in the union between these two arrays. Union of the two
          arrays can be defined as the set containing distinct elements from both the arrays.
          If there are repetitions, then only one occurrence of element should be printed in
          the union. Input:1 2 3 4 5
                                     1 2 3
                                                Output: 5
          def count_union_elements(arr1, arr2):
            union_set = set(arr1) | set(arr2) # '|' operator computes the union of sets
            return len(union_set)
          # Example usage
          arr1 = [1, 2, 3, 4, 5]
          arr2 = [1, 2, 3]
          result = count union elements(arr1, arr2)
          print("Number of elements in the union:", result)
32
          Inorder traversal means traversing through the tree in a Left, Node, Right
```

manner. We first traverse left, then print the current node, and then traverse right. This is done recursively for each node. Given a BST, find its in-order traversal.

```
class TreeNode:
  def __init__(self, key):
     self.key = key
     self.left = None
     self.right = None
def inorder_traversal(root):
  result = []
  if root:
     result.extend(inorder_traversal(root.left))
     result.append(root.key)
     result.extend(inorder_traversal(root.right))
  return result
# Example usage
root = TreeNode(5)
root.left = TreeNode(3)
root.right = TreeNode(8)
root.left.left = TreeNode(2)
root.left.right = TreeNode(4)
root.right.left = TreeNode(6)
root.right.right = TreeNode(9)
inorder_result = inorder_traversal(root)
print("In-order traversal:", inorder_result)
```