Code:

1. Selection Sort

```
def selection_sort(arr):
    n = len(arr)
    for i in range(n):
        min_idx = i
        for j in range(i + 1, n):
        if arr[j] < arr[min_idx]:
            min_idx = j
        arr[i], arr[min_idx] = arr[min_idx], arr[i]

arr = [64, 25, 12, 22, 11]
selection_sort(arr)
print("Selection Sort Result:", arr)</pre>
```

Output:

```
Selection Sort Result: [11, 12, 22, 25, 64]
```

2. Prim's Algorithm

```
import heapq
def prim mst(graph):
  mst = []
  visited = set()
  start node = list(graph.keys())[0]
  visited.add(start node)
  edges = [(cost, start node, neighbor) for neighbor, cost in graph[start node]]
  heapq.heapify(edges)
  while edges:
     cost, u, v = heapq.heappop(edges)
     if v not in visited:
       visited.add(v)
       mst.append((u, v, cost))
       for neighbor, n_cost in graph[v]:
          if neighbor not in visited:
             heapq.heappush(edges, (n cost, v, neighbor))
  return mst
graph = {
  'A': [('B', 2), ('C', 3)],
  'B': [('A', 2), ('C', 4), ('D', 5)],
  'C': [('A', 3), ('B', 4), ('D', 1)],
  'D': [('B', 5), ('C', 1)],}
mst = prim_mst(graph)
print("Prim's Minimal Spanning Tree:", mst)
Output:
```

3. Kruskal's Algorithm:

```
def kruskal mst(graph):
  mst = []
  edges = []
  for node in graph:
     for neighbor, cost in graph[node]:
       edges.append((cost, node, neighbor))
  edges.sort()
  parent = {node: node for node in graph}
  def find(node):
     if parent[node] != node:
       parent[node] = find(parent[node])
     return parent[node]
  for cost, u, v in edges:
     if find(u) != find(v):
       mst.append((u, v, cost))
       parent[find(u)] = find(v)
  return mst
graph = {
  'A': [('B', 2), ('C', 3)],
  'B': [('A', 2), ('C', 4), ('D', 5)],
  'C': [('A', 3), ('B', 4), ('D', 1)],
  'D': [('B', 5), ('C', 1)],
}
mst = kruskal mst(graph)
print("Kruskal's Minimal Spanning Tree:", mst)
```

Output:

```
Kruskal's Minimal Spanning Tree: [('C', 'D', 1), ('A', 'B', 2), ('A', 'C', 3)]
```

4. <u>Dijkstra's Algorithm:</u>

import heapq

```
def dijkstra(graph, start):
    distances = {node: float('inf') for node in graph}
    distances[start] = 0
    queue = [(0, start)]

while queue:
    current_dist, node = heapq.heappop(queue)

if current_dist > distances[node]:
    continue
```

```
for neighbor, weight in graph[node]:
    distance = current_dist + weight
    if distance < distances[neighbor]:
        distances[neighbor] = distance
        heapq.heappush(queue, (distance, neighbor))

return distances
graph = {
    'A': [('B', 2), ('C', 5)],
    'B': [('A', 2), ('C', 1), ('D', 7)],
    'C': [('A', 5), ('B', 1), ('D', 3)],
    'D': [('B', 7), ('C', 3)],
}
start_node = 'A'
shortest_distances = dijkstra(graph, start_node)
print("Dijkstra's Shortest Distances from", start_node + ":", shortest_distances)
```

Output:

```
Dijkstra's Shortest Distances from A: {'A': 0, 'B': 2, 'C': 3, 'D': 6}
```

Time Complexity:

Selection Sort: $O(n^2)$, where n is the number of elements in the array. This is because, in the worst case, it needs to compare and swap elements for every possible pair.

Prim's Minimal Spanning Tree Algorithm: O(E log V), where E is the number of edges and V is the number of vertices. This complexity is for the binary heap-based implementation of Prim's algorithm.

Kruskal's Minimal Spanning Tree Algorithm: O(E log E), where E is the number of edges. Sorting the edges dominates the time complexity.

Dijkstra's Minimal Spanning Tree Algorithm (Single-Source Shortest Path): $O(V^2)$ for the naive implementation with an adjacency matrix or $O((V+E)\log V)$ using a binary heap-based priority queue.