Computational Linear Algebra: SPARSE MATRICES Homework

21 January 2025

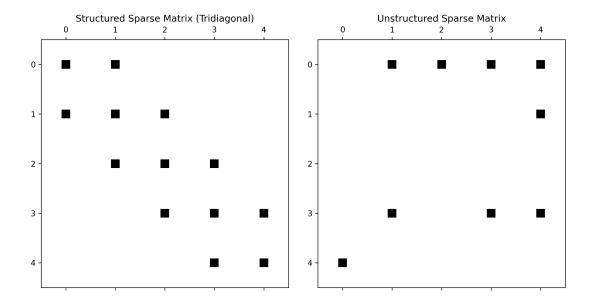
0. Importing Modules

```
[1]: import numpy as np
  import matplotlib.pyplot as plt
  import matplotlib.colors as mcolors
  from scipy.sparse import csr_matrix, csc_matrix, diags
  import networkx as nx
  from collections import deque
```

1. Introduction

Comparison of a structured and an unstructured sparse matrix

```
[2]: # Structured Sparse Matrix (Tridiagonal)
     diagonals = [np.ones(5), np.ones(4), np.ones(4)]
     A = diags(diagonals, [0, -1, 1], shape=(5, 5)).tocsr() # To Compressed Sparse Row
     # Unstructured Sparse Matrix
     B = csr_matrix(np.random.randint(0, 2, size=(5, 5)))
     plt.figure(figsize=(10, 5))
     plt.rcParams['axes.facecolor'] = 'white'
     # Subplot 1 (Structured Sparse Matrix (Tridiagonal))
     plt.subplot(1, 2, 1)
     plt.spy(A, marker='s', color='black')
     plt.title("Structured Sparse Matrix (Tridiagonal)")
     # Subplot 2 (Unstructured Sparse Matrix)
     plt.subplot(1, 2, 2)
     plt.spy(B, marker='s', color='black')
     plt.title("Unstructured Sparse Matrix")
     plt.tight_layout()
     plt.savefig("structuredVSunstructured.png", dpi=300)
     plt.show()
```



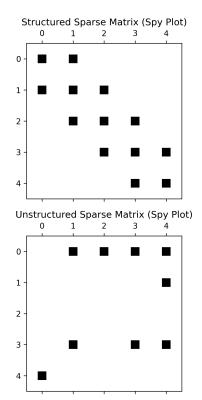
2. Graph Representations

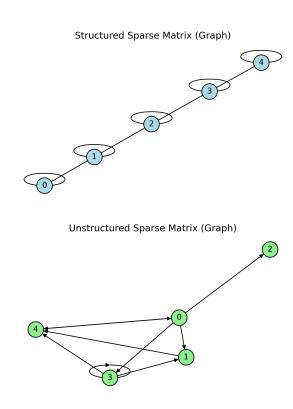
2.1 Graphs and Adjacency Graphs

Representation of structured and unstructured sparse matrices with respective spy plots and graphs

```
[3]: plt.figure(figsize=(10, 7))
    plt.rcParams['axes.facecolor'] = 'white'
     # Subplot 1 (Structured Matrix - Spy Plot)
    plt.subplot(2, 2, 1) # Use 2 rows, 2 columns
    plt.spy(A, marker='s', color='black')
    plt.title("Structured Sparse Matrix (Spy Plot)")
     # Subplot 2 (Structured Matrix - Graph)
    plt.subplot(2, 2, 2)
         # Create graph from unstructured matrix
    rows, cols = A.nonzero()
    edges = zip(rows.tolist(), cols.tolist())
    G_structured = nx.Graph()
    G_structured.add_nodes_from(range(A.shape[0])) # Add nodes explicitly
    G_structured.add_edges_from(edges)
         # Layout and visualization of the graph
    pos = nx.spring_layout(G_structured)
    nx.draw(G_structured, pos=pos, with_labels=True, node_size=400,__
     →node_color="lightblue", edgecolors="black", font_size=10)
    plt.title("Structured Sparse Matrix (Graph)")
```

```
# Subplot 3 (Unstructured Matrix - Spy Plot)
plt.subplot(2, 2, 3)
plt.spy(B, marker='s', color='black')
plt.title("Unstructured Sparse Matrix (Spy Plot)")
# Subplot 4 (Unstructured Matrix - Graph)
plt.subplot(2, 2, 4)
    # Create graph from unstructured matrix
rows, cols = B.nonzero()
edges = zip(rows.tolist(), cols.tolist())
G_unstructured = nx.DiGraph()
G_unstructured.add_nodes_from(range(B.shape[0])) # Add nodes explicitly
G_unstructured.add_edges_from(edges)
    # Layout and visualization of the graph
pos = nx.spring_layout(G_unstructured)
nx.draw(G_unstructured, pos=pos, with_labels=True, node_size=400,__
→node_color="lightgreen", edgecolors="black", font_size=10)
plt.title("Unstructured Sparse Matrix (Graph)")
plt.tight_layout()
plt.savefig("structured_unstructured_comparison.png", dpi=300)
plt.show()
```





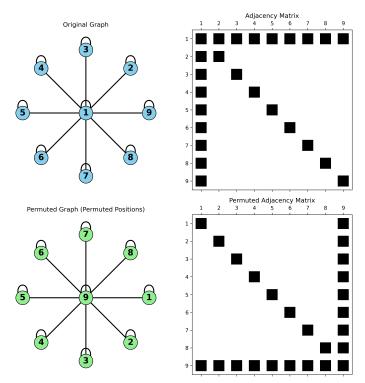
3. Permutations and Reorderings

3.1 Relations with the Adjacency Graph

```
[4]: # Create the ORIGINAL graph
     original_graph = nx.Graph() # Create an empty NetworkX graph object
     original_nodes = list(range(1, 10)) # Create a list of nodes from 1 to 9_{\square}
     \hookrightarrow (inclusive)
     original_graph.add_nodes_from(original_nodes) # Add the nodes to the graph
     for node in original_nodes: # Iterate through all nodes
         if node != 1: # If the node is not node 1
             original_graph.add_edge(1, node) # Add an edge between node 1 and the_
     \hookrightarrow current node
     for node in original_nodes:
         original_graph.add_edge(node, node) # Add a self-loop (edge to itself) for
      →each node
     # Calculate ORIGINAL positions
     original_positions = nx.spring_layout(original_graph) # Calculate node_
     ⇒positions using the spring layout algorithm
     original_positions[1] = (0.5, 0.5) # Manually set the position of node 1 to the_
     \rightarrow center of the plot (0.5, 0.5)
     for node in original_nodes:
         if node != 1:
             angle = (node - 1) * 2 * np.pi / 8 # Calculate the angle for the node
      →in a circular layout (8 nodes around the center)
             original_positions[node] = (0.5 + np.cos(angle) * 0.4, 0.5 + np.
      \rightarrowsin(angle) * 0.4) # Calculate x and y coordinates for circular placement
      \rightarrow around node 1
     # Create the PERMUTED graph
     permuted_nodes = list(range(9, 0, -1)) # Application of the permutation pi = {9, }
     \rightarrow 8, 7, 6, 5, 4, 3, 2, 1} (reverse order of nodes)
     node_mapping = dict(zip(original_nodes, permuted_nodes)) # Create a dictionary_
     → mapping original nodes to permuted nodes
     permuted_graph = nx.relabel_nodes(original_graph.copy(), node_mapping)
     # Create the NEW positions mapping
     permuted_positions = {}
     for original_node, permuted_node in node_mapping.items(): # Iterate through the_
     \rightarrow node mapping
         permuted_positions[permuted_node] = original_positions[original_node] #__
     →Assign the original position to the corresponding permuted node
     # Calculate ORIGINAL adjacency matrix
```

```
original_adjacency_matrix = nx.to_numpy_array(original_graph,_
→nodelist=original_nodes)
plt.figure(figsize=(10, 10))
# Subplot 1 - Plot the ORIGINAL graph
plt.subplot(2, 2, 1)
nx.draw_networkx_nodes(original_graph, original_positions, node_size=700,_
→node_color="skyblue", edgecolors="black") # Draw nodes in the original graph
→with skyblue color and black edges
nx.draw_networkx_edges(original_graph, original_positions, width=2) # Draw_1
\rightarrowedges in the original graph with a width of 2
nx.draw_networkx_labels(original_graph, original_positions, font_size=16,__
→font_weight="bold")
plt.title("Original Graph")
plt.gca().set_aspect('equal')
plt.axis('off')
# Subplot 2 - Plot the ORIGINAL adjacency matrix
plt.subplot(2, 2, 2)
inverted_adjacency_matrix = original_adjacency_matrix[::-1] # Invert the rows__
of the adjacency matrix to match the node order in the graph
plt.spy(inverted_adjacency_matrix, marker='s', markersize=20, color='black') #__
→Plot the adjacency matrix using square markers
plt.title("Adjacency Matrix") # Set the title for this subplot
plt.xticks(np.arange(len(permuted_nodes)), permuted_nodes[::-1]) # Set x-axis_u
→ labels (inverted order to match the matrix) with permuted node names
plt.yticks(np.arange(len(permuted_nodes)), permuted_nodes) # Set y-axis labels_
→with original node names
plt.gca().set_ylim([-0.5, len(permuted_nodes) - 0.5])
plt.gca().set_xlim([-0.5, len(permuted_nodes) - 0.5])
# Subplot 3 - Plot the PERMUTED graph
plt.subplot(2, 2, 3)
nx.draw_networkx_nodes(permuted_graph, permuted_positions, node_size=700,_
→node_color="lightgreen", edgecolors="black") # Draw nodes in the permuted_
→ graph with lightgreen color and black edges
nx.draw_networkx_edges(permuted_graph, permuted_positions, width=2) # Draw_
→edges in the permuted graph
nx.draw_networkx_labels(permuted_graph, permuted_positions, font_size=16,_
→font_weight="bold")
plt.title("Permuted Graph (Permuted Positions)")
plt.gca().set_aspect('equal')
plt.axis('off')
```

```
# Subplot 4 - Plot the PERMUTED adjacency matrix
plt.subplot(2, 2, 4)
permuted_adjacency_matrix = nx.to_numpy_array(permuted_graph,_
→nodelist=permuted_nodes) # Convert the permuted graph to a NumPy adjacency disconstant
→ matrix (based on permuted node order)
permuted_adjacency_matrix_transposed = permuted_adjacency_matrix.T # Transpose__
→ the adjacency matrix (for better visualization)
permuted_adjacency_matrix_inverted = permuted_adjacency_matrix_transposed[::-1, ::
\hookrightarrow:-1] # Invert both rows and columns of the transposed matrix
plt.spy(permuted_adjacency_matrix_inverted, marker='s', markersize=20,__
→color='black') # Plot the permuted adjacency matrix with square markers
plt.title("Permuted Adjacency Matrix")
# Configure Axes for the Adjacency Matrix Plot
plt.xticks(np.arange(len(permuted_nodes)), permuted_nodes[::-1])
plt.yticks(np.arange(len(permuted_nodes)), permuted_nodes[::-1])
plt.gca().set_ylim([len(permuted_nodes) - 0.5, -0.5])
plt.gca().set_xlim([-0.5, len(permuted_nodes) - 0.5])
# Finalize the plot
plt.tight_layout()
plt.savefig("graphs_and_matrices_2x2.png", dpi=300)
plt.show()
plt.close()
```



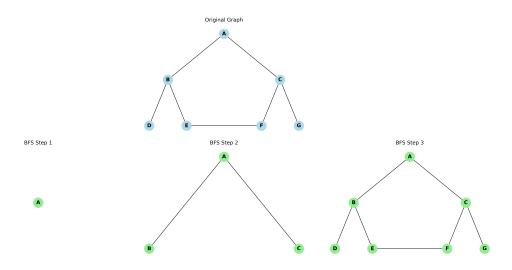
3.2 Common Reorderings

3.2.1 Level-set ordering

```
Breadth-First Search
[5]: def bfs(graph, start_node):
         if start_node not in graph: # Check if the starting node exists in the graph
             print(f"Error: Starting node '{start_node}' not found in the graph.") #_U
      →Print an error message if the start node is not found
             return None # Return None to indicate an error
         G_original = nx.from_dict_of_lists(graph) # Create a networkx graph from the
      \rightarrow graph dictionary
         # Fixed node positions for visualization
         pos = {
             'A': np.array([0.5, 0.8]),
             'B': np.array([0.2, 0.5]),
             'C': np.array([0.8, 0.5]),
             'D': np.array([0.1, 0.2]),
             'E': np.array([0.3, 0.2]),
             'F': np.array([0.7, 0.2]),
             'G': np.array([0.9, 0.2]),
         }
         s = deque([start_node]) # Initialize a deque for BFS, starting with the
      \rightarrowstart node
         seen = 1 # Initialize a counter to track the order in which nodes are
         pi = {seen: start_node} # Initialize a dictionary to map visit order to ⊔
      \rightarrownodes
         marked = {start_node} # Initialize a set to keep track of visited/marked_
      \rightarrownodes
         bfs_steps = [] # Initialize a list to store the graph at each BFS step
         while s: # While the queue is not empty
             s_new = deque() # Initialize a new deque for the next level of BFS
             current_step_graph = nx.Graph() # Create a new empty graph for the
      \rightarrow current BFS step
             current_step_graph.add_nodes_from(marked) # Add the already marked_
      →nodes to the current step graph
             for node in marked:
                 if node in graph:
                     for neighbor in graph[node]:
```

```
if neighbor in marked:
                       current_step_graph.add_edge(node, neighbor) # Add the_
→edge to the graph of the current step
       bfs_steps.append(current_step_graph) # Append the current state of the
\rightarrow graph to the list of steps
       for v in list(s): # Iterate through the nodes in the current queue
           if v in graph:
              for w in graph[v]: # Iterate through the neighbors of the
\hookrightarrow current node v
                   if w not in marked:
                       s_new.append(w) # Add the neighbor w to the new queue_
→ for the next level
                       marked.add(w) # Mark the neighbor w as visited
                       seen += 1 # Increment the visit order counter
                       pi[seen] = w # Map the visit order to the neighbor w in_
→ the pi dictionary
       s = s_new # Update the queue for the next level of BFS
   num_steps = len(bfs_steps) # Calculates the total number of BFS steps taken.
   fig, axes = plt.subplots(2, num_steps if num_steps > 0 else 1, figsize=(6 *
→(num_steps if num_steps > 0 else 1), 8))
   pos_dict = dict(pos) # Creates a dictionary pos_dict from the pos dictionary.
→ This is used to maintain node positions during plotting.
   # Plot the Original Graph (center of the first row)
   if num_steps > 0: # Check if there are any BFS steps to be plotted
       original_graph_ax = axes[0, num_steps // 2] # Selects the central_
→subplot in the first row to plot the original graph
       nx.draw(G_original, pos=pos_dict, with_labels=True,__
→ax=original_graph_ax) # Draws the original graph on the selected subplot with
→ specified visual properties
       original_graph_ax.set_title("Original Graph") # Sets the title for the
\rightarrow original graph subplot.
       # Hide the other axes in the first row
       for i in range(num_steps): # Iterates through all the subplots in the
\rightarrow first row
           if i != num_steps // 2: # Checks if the current subplot index is NOT_
\rightarrow the central one.
               axes[0, i].set_axis_off() # Turns off the axes (removes the_
→ spines, ticks, and labels) for the empty subplots in the first row, creating
\rightarrow whitespace
```

```
else: # This 'else' block handles the case where there are NO BFS steps,
 \rightarrow (only the original graph)
        original_graph_ax = axes[0] # Selects the first (and only) subplot in
\hookrightarrow the first row.
        nx.draw(G_original, pos=pos_dict, with_labels=True,__
→node_color='lightblue', node_size=500, font_size=12, font_weight='bold',
 →ax=original_graph_ax) # Draws the original graph on the subplot
        original_graph_ax.set_title("Original Graph") # Sets the title of the
\hookrightarrow subplot.
    # Plot BFS Steps (second row)
    for i, graph_step in enumerate(bfs_steps): # Iterates through the list of u
 →BFS steps, using enumerate to get both the index (i) and the graph for each
 \rightarrow step (graph_step)
        ax = axes[1, i] if num_steps > 0 else axes[1] # Select the current axis_
→to plot on. If there are multiple steps, selects the appropriate axis in the
 ⇒second row. If there is only one axis, selects that axis
        nx.draw(graph_step, pos=pos_dict, with_labels=True,_
→node_color='lightgreen', node_size=500, font_size=12, font_weight='bold',
 \rightarrowax=ax) # Draws the graph for the current BFS step on the corresponding subplot
→with specified visual properties
        ax.set_title(f"BFS Step {i+1}") # Sets the title for the current BFS_
\rightarrowstep subplot.
    plt.tight_layout() # Adjusts subplot parameters to provide padding and_
 →prevent overlapping of titles/labels.
    plt.savefig("BFS.png", dpi=300)
    plt.show() # Displays the plot in a new window.
    return pi, marked
# EXAMPLE
graph = {
    'A': ['B', 'C'],
    'B': ['D', 'E'],
    'C': ['F', 'G'],
    'D': [].
    'E': ['F'],
    'F': [],
    'G': ['C']
}
start_node = 'A'
pi, marked = bfs(graph, start_node)
```

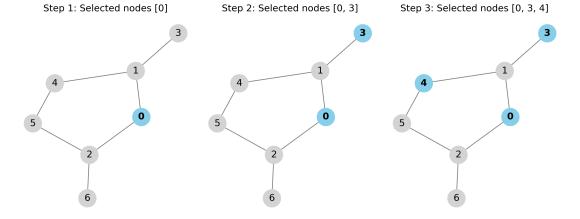


3.2.2 Greedy Algorithm for ISO

```
[6]: def adjacency_matrix_from_dict(graph):
         if not graph: # Check if the input graph is empty (None or {})
             return None, None # If the graph is empty, return None for both the
      →matrix and the node-to-index mapping
         nodes = sorted(list(graph.keys())) # Get a sorted list of the graph's nodes.
      \hookrightarrow Sorting ensures consistent matrix representation regardless of the \sqcup
      → dictionary's original order
         n = len(nodes) # Get the number of nodes in the graph
         node_to_index = {node: i for i, node in enumerate(nodes)} # Create a_
      →dictionary that maps each node to its corresponding index (0-based) in the
      \rightarrow matrix
         matrix = [[0] * n for _ in range(n)] # Initialize the adjacency matrix as a_{\sqcup}
      →list of lists, filled with zeros. This creates an n x n matrix
         for node, neighbors in graph.items():
             i = node_to_index[node]
             for neighbor in neighbors: # Iterate through each 'neighbor' of the
      →current 'node'
                 if neighbor in node_to_index:
                     j = node_to_index[neighbor] # Get the column index 'j' for the_
      → 'neighbor' from the node_to_index mapping
                     matrix[i][j] = 1  # Set the element at row 'i' and column 'j' of
      → the matrix to 1, indicating an edge between the 'node' and its 'neighbor'
         return matrix, node_to_index
```

```
[7]: def greedy_iso(adjacency_matrix, pos):
         n = len(adjacency_matrix) # Number of nodes in the graph
         S = [] # Initialize the independent set (empty at the beginning)
         marked = [False] * n # List to keep track of visited/marked nodes_
      \hookrightarrow (initially all False)
         original_graph = nx.from_numpy_array(np.array(adjacency_matrix)) # Create a_
      →networkx graph from the adjacency matrix
         subgraphs = [] # list of the subgraphs at each step
         for j in range(n):
             if not marked[j]: # If the current node j is not marked (not yet in the
      →independent set or a neighbor of one)
                 S.append(j) # Add node j to the independent set S
                 marked[j] = True # Mark node j as visited in the independent set
                 subgraph = original_graph.subgraph(S).copy() # Create a subgraph_
      \rightarrow with the nodes of S and copy it
                 subgraphs.append(subgraph)
                 for i in range(n): # Iterate through all other nodes to check for
      \rightarrowneighbors of j
                     if adjacency_matrix[j][i] == 1:
                         marked[i] = True # Mark node i as visited
         return S, original_graph, subgraphs
     # EXAMPLE
     graph = {
         'A': ['B', 'C'],
         'B': ['D', 'E'],
         'C': ['F', 'G'],
         'D': [],
         'E': ['F'],
         'F': [],
         'G': ['C']
     }
     matrix, node_to_index = adjacency_matrix_from_dict(graph)
     # Check if the adjacency matrix is not empty
     if matrix:
         adjacency_matrix = matrix # Assign the adjacency matrix for further_
      \hookrightarrowprocessing
         G = nx.from_numpy_array(np.array(adjacency_matrix)) # Create a NetworkX_
      → graph object from the adjacency matrix
```

```
pos = nx.spring_layout(G, seed=42)  # Generate node positions for_
 \hookrightarrow visualization using the spring layout algorithm
    independent_set, original_graph, subgraphs = greedy_iso(adjacency_matrix,_
    plt.figure(figsize=(10, 4 * (len(subgraphs) // 3 + 1)))
    for i, subgraph in enumerate(subgraphs):
        plt.subplot(len(subgraphs) // 3 + 1, 3, i + 1)
        plt.title(f"Step {i+1}: Selected nodes {list(subgraph.nodes)}")
        nx.draw(original_graph, pos, with_labels=True, node_color="lightgray", u
 →node_size=500, edge_color="gray")
        nx.draw(subgraph, pos, with_labels=True, node_color="skyblue",_
 →node_size=500, font_weight="bold")
    plt.tight_layout()
    plt.savefig("iso.png", dpi=300)
    plt.show()
# If the adjacency matrix was empty
else:
    print("Empty graph") # Print a message indicating an empty graph
```



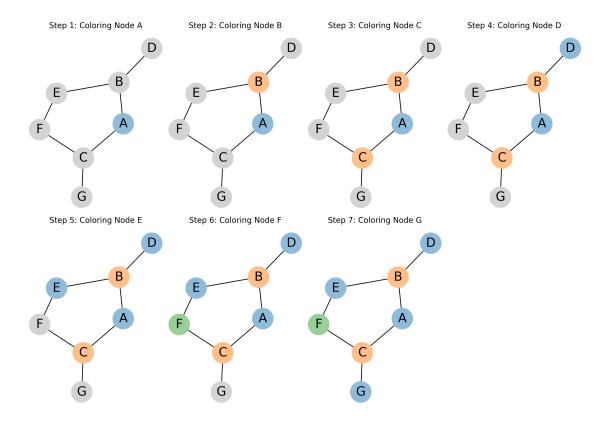
5 0

Step 4: Selected nodes [0, 3, 4, 6]

3.2.3 Greedy Multicoloring Algorithm

```
[8]: def greedy_multicoloring(graph, pos):
         if not graph:
             return None, None # Handle empty graph case
         G = nx.Graph(graph) # Convert adjacency list to NetworkX graph object for
      \rightarrow easier manipulation
         n = len(G.nodes) # Get the number of nodes in the graph
         color = {node: 0 for node in G.nodes} # Initialize a dictionary to store
      ⇒color assignments for each node
         node_colors = []
         light_colors = []
         for name, hex in mcolors.TABLEAU_COLORS.items():
             rgb = mcolors.hex2color(hex)
             light_rgb = [(c + 1) / 2 for c in rgb]
             light_colors.append(mcolors.rgb2hex(light_rgb))
         # Calculate the number of rows and columns needed for the plot (dynamically)
         rows = min(4, (n + 3) // 4)
         cols = min(4, n) if n < 4 else 4
         fig, axes = plt.subplots(rows, cols, figsize=(10, 7))
         axes = axes.flatten()
         for idx, i in enumerate(G.nodes):
             neighbor_colors = set() # Find the colors of neighboring nodes (already_
      \rightarrow assigned colors)
             for j in G.neighbors(i):
                 if color[j] != 0:
                     neighbor_colors.add(color[j])
             k = 1 # Find the next available color that doesn't conflict with
      \rightarrowneighbors
             while k in neighbor_colors:
                 k += 1
             color[i] = k
             node_colors.append(color.copy()) # Keep track of assigned colors for
      →visualization at each step
             ax = axes[idx] if idx < rows * cols else None # Get the corresponding_
      → subplot for the current node
```

```
if ax:
            node_color_map = []
            for node in G.nodes():
                # Assign color based on the color dictionary and a light color_{\sqcup}
\rightarrowscheme
                if node in color and color[node] != 0:
                    node_color_map.
 →append(light_colors[(color[node]-1)%len(light_colors)])
                else:
                    node_color_map.append("lightgrey") # Uncolored nodes
            nx.draw(G, pos=pos, with_labels=True, node_color=node_color_map,
                    node_size=700, font_size=16, ax=ax, font_color="black")
            ax.set_title(f"Step {idx+1}: Coloring Node {i}", fontsize=10)
            ax.set_axis_off()
        elif idx >= rows * cols:
            print(f"Number of nodes exceeds the plot grid ({rows}x{cols}), only_
 →the first {rows * cols} have been plotted")
            break
    # Hide unused subplots
    for i in range(idx + 1, rows * cols):
        if axes[i]:
            axes[i].axis('off')
    plt.tight_layout(pad=0.5)
    plt.savefig("color.png", dpi=300)
    plt.show()
# EXAMPLE
graph = {
    'A': ['B', 'C'],
    'B': ['D', 'E'],
    'C': ['F', 'G'],
    'D': [],
    'E': ['F'],
    'F': [],
    'G': []
}
G = nx.Graph(graph)
pos = nx.spring_layout(G, seed=42)
greedy_multicoloring(graph, pos)
```



4 Storage Schemes

4.1 Coordinate Format

```
[9]: def coordinate_format(matrix):
    # Find the positions of the non-zero elements
    non_zero_positions = matrix.nonzero()

# Extract the non-zero values from the matrix
AA = matrix[non_zero_positions]

# Get the row and column indices of the non-zero values
    JR = non_zero_positions[0]
    JC = non_zero_positions[1]

    return AA, JR, JC

def coo(matrix):
    coo = coo_matrix(matrix) # COO method
    AA = coo.data
    JR = coo.row
    JC = coo.col
```

```
return AA, JR, JC
     # EXAMPLE
     matrix = np.array([[1, 0, 0, 2, 0],
                        [3, 4, 0, 5, 0],
                        [6, 0, 7, 8, 9],
                        [0, 0, 10, 11, 0],
                        [0, 0, 0, 0, 12]])
     AA, JR, JC = coordinate_format(matrix)
     print(f"COO format representation of the matrix:\n{matrix}")
     print(f"Non-zero values (AA): {AA}")
     print(f"Row indices (JR): {JR}")
     print(f"Column indices (JC): {JC}")
     AA, JR, JC = coo(matrix)
     print(f"\nCOO format representation of the matrix:\n{matrix}")
     print(f"Non-zero values (AA): {AA}")
     print(f"Row indices (JR): {JR}")
     print(f"Column indices (JC): {JC}")
     COO format representation of the matrix:
     [[1 0 0 2 0]
      [3 4 0 5 0]
      [60789]
      [ 0 0 10 11 0]
      [000012]]
     Non-zero values (AA): [ 1 2 3 4 5 6 7 8 9 10 11 12]
     Row indices (JR): [0 0 1 1 1 2 2 2 2 3 3 4]
     Column indices (JC): [0 3 0 1 3 0 2 3 4 2 3 4]
     COO format representation of the matrix:
     [[1 0 0 2 0]
      [3 4 0 5 0]
      [6 0 7 8 9]
      [ 0 0 10 11 0]
      [000012]]
     Non-zero values (AA): [ 1 2 3 4 5 6 7 8 9 10 11 12]
     Row indices (JR): [0 0 1 1 1 2 2 2 2 3 3 4]
     Column indices (JC): [0 3 0 1 3 0 2 3 4 2 3 4]
     4.2 Compressed Sparse Row (CSR) Format
[10]: def csr_manual(matrix):
         rows, cols = matrix.shape # Get the number of rows and columns of the input
```

 \rightarrow matrix

```
nnz = np.count_nonzero(matrix) # Count the number of non-zero elements in_
\rightarrow the matrix
    AA = np.zeros(nnz) # Initialize the AA array (values) with zeros. Its size
\rightarrow is equal to the number of non-zero elements
    JA = np.zeros(nnz, dtype=int) # Initialize the JA array (column indices)
→with zeros
    IA = np.zeros(rows + 1, dtype=int) # Initialize the IA array (row pointers)
→with zeros
    index = 0 # Initialize an index to keep track of the current position in \Box
\hookrightarrow the AA and JA arrays.
    for i in range(rows):
        IA[i] = index # Set the i-th element of IA to the current index. This
→marks the beginning of the non-zero elements for row i in AA and JA.
        for j in range(cols):
            if matrix[i, j] != 0:
                 AA[index] = matrix[i, j] # If it's non-zero, store its value in_
\hookrightarrow AA at the current index
                 JA[index] = j # Store its column index in JA at the current
\hookrightarrow index
                 index += 1 # Increment the index to point to the next position⊔
\rightarrow in AA and JA
    IA[rows] = index # After processing all rows, set the last element of IA tou
→ the final index value. This marks the end of the non-zero elements in AA and
\hookrightarrow JA
   return AA, JA, IA
def csr(matrix):
    csr = csr_matrix(matrix) # CSR method
    AA = csr.data
    JA = csr.indices
    IA = csr.indptr
    return AA, JA, IA
# EXAMPLE
matrix = np.array([[1, 0, 0, 2, 0],
               [3, 4, 0, 5, 0],
               [6, 0, 7, 8, 9],
               [0, 0, 10, 11, 0],
               [0, 0, 0, 0, 12]])
AA, JA, IA = csr_manual(matrix)
```

```
print(f"CSR representation of the matrix:\n{matrix}")
      print(f"Non-zero values (AA): {AA}")
      print(f"Column indexes (JA): {JA}")
      print(f"Row indexes (IA): {IA}")
      AA, JA, IA = csr(matrix)
      print(f"\nCSR rrepresentation of the matrix:\n{matrix}")
      print(f"Non-zero values (AA): {AA}")
      print(f"Column indexes (JA): {JA}")
      print(f"Row indexes (IA): {IA}")
     CSR representation of the matrix:
     [[1 0 0 2 0]
      [3 4 0 5 0]
      [6 0 7 8 9]
      [ 0 0 10 11 0]
      [0 0 0 0 12]]
     Non-zero values (AA): [ 1. 2. 3. 4. 5. 6. 7. 8. 9. 10. 11. 12.]
     Column indexes (JA): [0 3 0 1 3 0 2 3 4 2 3 4]
     Row indexes (IA): [ 0 2 5 9 11 12]
     CSR rrepresentation of the matrix:
     [[1 0 0 2 0]
      [3 4 0 5 0]
      [60789]
      [ 0 0 10 11 0]
      [0 \ 0 \ 0 \ 0 \ 12]]
     Non-zero values (AA): [ 1 2 3 4 5 6 7 8 9 10 11 12]
     Column indexes (JA): [0 3 0 1 3 0 2 3 4 2 3 4]
     Row indexes (IA): [ 0 2 5 9 11 12]
     4.3 Compressed Sparse Column (CSC) Format
[11]: def csc_manually(matrix):
         rows, cols = matrix.shape # Get the number of rows and columns of the inputu
      \hookrightarrow matrix
         num_non_zero_elements = np.count_nonzero(matrix) # Count the number of_
      \rightarrownon-zero elements in the matrix
          # Allocate arrays for the CSC data
         data = np.zeros(num_non_zero_elements) # Initialize the 'data' array_
      → (values) with zeros
         row_indices = np.zeros(num_non_zero_elements, dtype=int) # Initialize the_
      → 'row_indices' array with zeros
         column_pointers = np.zeros(cols + 1, dtype=int) # Initialize the_
      → 'column_pointers' array with zeros
         current_index = 0
```

```
for col_index in range(cols):
        column_pointers[col_index] = current_index # Set the current element of:
 →'column_pointers' to the current index. This marks the beginning of the
 →non-zero elements for the current column
        for row_index in range(rows):
            if matrix[row_index, col_index] != 0:
                data[current_index] = matrix[row_index, col_index] # If it's_
→non-zero, store its value in 'data' at the current index
                row_indices[current_index] = row_index # Store its row index in_
 → 'row_indices' at the current index
                current_index += 1 # Increment the index to point to the next_
→position in 'data' and 'row_indices'
    column_pointers[cols] = current_index # After processing all columns, set_
→ the last element of 'column_pointers' to the final index value. This marks the
\rightarrow end of the non-zero elements
    return data, row_indices, column_pointers
def csc(matrix):
    # Convert to CSC format using SciPy
    csc = csc_matrix(matrix)
    # Extract data from the CSC matrix
    data = csc.data
    row_indices = csc.indices
    column_pointers = csc.indptr
   return data, row_indices, column_pointers
# EXAMPLE
matrix = np.array([[1, 0, 0, 2, 0],
              [3, 4, 0, 5, 0],
              [6, 0, 7, 8, 9],
              [0, 0, 10, 11, 0],
              [0, 0, 0, 0, 12]])
# Convert to CSC format manually
data, row_indices, column_pointers = csc_manually(matrix)
print(f"CSC representation of the matrix:\n{matrix}")
print(f"Non-zero values (data): {data}")
print(f"Row indices (row_indices): {row_indices}")
print(f"Column pointers (column_pointers): {column_pointers}")
# Convert to CSC format using SciPy
data, row_indices, column_pointers = csc(matrix)
print(f"\nCSC representation of the matrix:\n{matrix}")
print(f"Non-zero values (data): {data}")
```

```
print(f"Row indices (row_indices): {row_indices}")
print(f"Column pointers (column_pointers): {column_pointers}")
CSC representation of the matrix:
[[1 0 0 2 0]
[3 4 0 5 0]
[6 0 7 8 9]
[ 0 0 10 11 0]
[ 0 0 0 0 12]]
Non-zero values (data): [ 1. 3. 6. 4. 7. 10. 2. 5. 8. 11. 9. 12.]
Row indices (row_indices): [0 1 2 1 2 3 0 1 2 3 2 4]
Column pointers (column_pointers): [ 0 3 4 6 10 12]
CSC representation of the matrix:
[[1 0 0 2 0]
[3 4 0 5 0]
[6 0 7 8 9]
[ 0 0 10 11 0]
 [000012]]
Non-zero values (data): [ 1 3 6 4 7 10 2 5 8 11 9 12]
Row indices (row_indices): [0 1 2 1 2 3 0 1 2 3 2 4]
Column pointers (column_pointers): [ 0 3 4 6 10 12]
```