

Graphs and Graphing Algorithms

1. The Graph Abstract Data Type
2. Breadth-First Search
3. Depth-First Search
4. Shortest Path Problems
5. Minimum Spanning Tree

7.1 Introduction

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We will see in this chapter that once we have a good representation for a problem, we can use some standard graph algorithms to solve what otherwise might seem to be a very difficult problem!

A graph is just like a road map. If you have ever used one of the internet map sites, you know that a computer can find the shortest, quickest, or easiest path from one place to another.

7.2. Vocabulary and Definitions

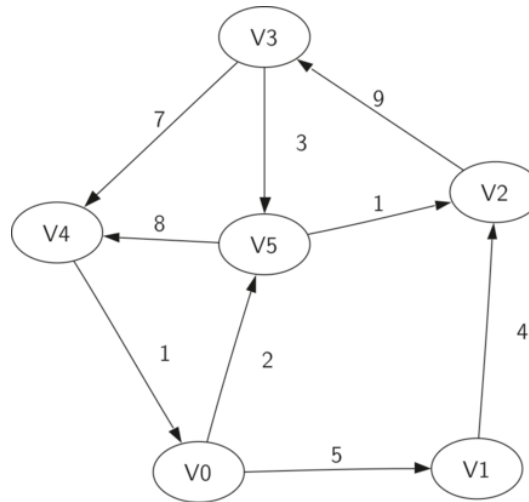
- Vertex: A vertex (also called a **node**) is a fundamental part of a graph. It can have a name, which we will call the **key**. A vertex may also have additional information. We will call this additional information the **value** or the payload.

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- Edge: An edge (also called an **arc**) is another fundamental part of a graph. An edge connects two vertices to show that there is a relationship between them. Edges may be one-way or two-way. If the edges in a graph are all one-way, we say that the graph is a directed graph, or a digraph.

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- Weight: Edges may be weighted to show that there is a cost to go from one vertex to another, which we call **edge cost**. For example, in a graph of roads that connect one city to another, the weight on the edge might represent the distance between the two cities.

A graph can be represented by $G = (V, E)$. For the graph G , V is a set of vertices and E is a set of edges. Each edge is a tuple where $e = (u, v)$. We can add a third component to the edge tuple to represent a weight. A subgraph is a set of edges and vertices such that $G' = (V', E')$ and $V' \subseteq V$ and $E' \subseteq E$.

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The above is a simple weighted digraph. Formally we can represent this graph as the set of six vertices:

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and the set of nine edges:

The example graph helps illustrate two other key graph terms:

- Path: A path in a graph is a sequence of distinct vertices such that for all i , i and $i+1$ are adjacent. The **unweighted path length** is the number of edges, $|E|$; the **weighted path length** is the sum of the weights of all edges in the path. For example, the path from s to t is $s \rightarrow a \rightarrow b \rightarrow t$. (A more general sequence that allows repeated vertices is called a **walk**.)

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- Path: A path in a graph is a sequence of distinct vertices such that for all i , v_i and v_{i+1} are adjacent. The **unweighted path length** is the number of edges, $|E|$; the **weighted path length** is the sum of the weights of all edges in the path. For example, the path from v_1 to v_4 is v_1, v_2, v_3, v_4 . (A more general sequence that allows repeated vertices is called a **walk**.)
- Cycle: A cycle is a closed sequence $v_1, v_2, \dots, v_k, v_1$ where $v_1, \dots, v_k$ are distinct vertices and consecutive vertices are connected by an edge. For example, v_1, v_2, v_3, v_1 is a cycle. A graph with no cycles is called an **acyclic graph**, and a directed graph with no cycles is called a **directed acyclic graph (DAG)**. We will see that many important problems can be solved efficiently if they can be modeled as a DAG!

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- Cycle: A cycle is a closed sequence where $v_1, v_2, \dots, v_n$ are distinct vertices and consecutive vertices are connected by an edge. For example, v_1, v_2, v_3, v_1 is a cycle. A graph with no cycles is called an **acyclic graph**, and a directed graph with no cycles is called a **directed acyclic graph (DAG)**. We will see that many important problems can be solved efficiently if they can be modeled as a DAG!

A tree is a connected, acyclic, undirected graph. In other words, there is a path between every pair of vertices, and the graph contains no cycles. We will explore different types of trees in detail in the next chapter!

7.3. The Graph Abstract Data Type

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- `add_edge(from_vert, to_vert)` : adds a new directed edge to the graph that connects two vertices.
- `add_edge(from_vert, to_vert, weight)` : adds a new weighted directed edge to the graph that connects two vertices.

- `get_vertex(vert_key)` : finds the vertex in the graph named `vert_key`.
- `get_vertices()` : returns the list of all vertices in the graph.
- `in` : returns `True` for a statement `vertex in graph` if the given vertex is in the graph, `False` otherwise.

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Now that we have looked at the definition for the graph ADT, there are several ways we can implement it. There are two well-known implementations of a graph, the adjacency matrix and the adjacency list.

7.4. An Adjacency Matrix

One of the easiest ways to implement a graph is to use a two-dimensional matrix. Each of the rows and columns represents a vertex in the graph. The value that is stored in the cell at the intersection of row and column indicates if there is an edge.

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	V0	V1	V2	V3	V4	V5
V0		5				2
V1			4			
V2				9		
V3					7	3
V4	1					
V5			1		8	

When two vertices are connected by an edge, we say that they are adjacent and the value in each cell represents the weight. The advantage of the adjacency matrix is that it is simple, and for small graphs it is easy to see which nodes are connected to other nodes. However, we can say that this matrix is **sparse**. A matrix is not a very efficient way to store sparse data.

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However, there are few real problems that approach this sort of connectivity. The problems we will look at in this chapter all involve graphs that are sparsely connected!

7.5. An Adjacency List

A more space-efficient way to implement a sparsely connected graph is to use an adjacency list. In this implementation, we keep a master list of all the vertices in the `Graph` object, and each vertex object in the graph maintains a list of the other vertices that it is connected to.

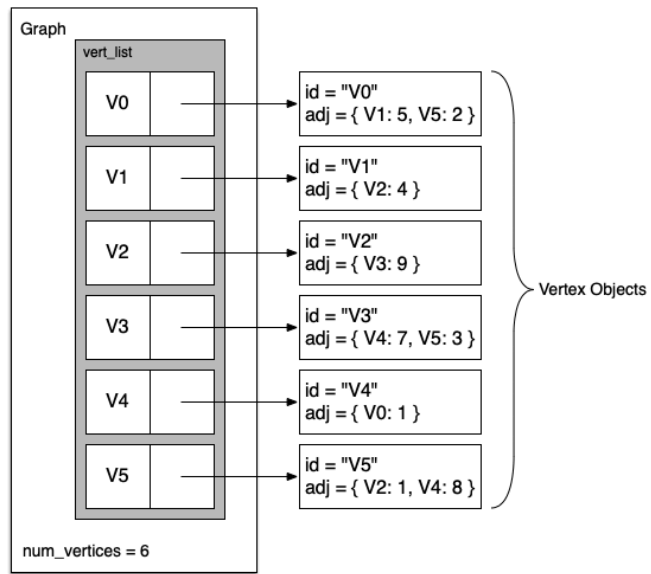
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The advantage of the adjacency list implementation is that it allows us to compactly represent a sparse graph. The adjacency list also allows us to easily find all the links that are directly connected to a particular vertex!



```
In [4]: display_quiz(quiz_path+"graph.json", max_width=800)
```

Regarding graph implementation methods, which of the following statements is true according to the materials?

- ☐ In an adjacency matrix, if two vertices are not connected, no memory is allocated for that pair.
- ☐ Finding all neighbors of a vertex is faster in an adjacency matrix than in an adjacency list.
- ☐ An adjacency matrix is the most space-efficient way to represent a sparse graph.
- ☐ An adjacency list is more space-efficient for graphs that are sparsely connected.

7.6. Implementation

In [5]:

```
class Vertex:
    def __init__(self, key):
        self.key = key
        self.neighbors = {}

    def get_neighbor(self, other):
        return self.neighbors.get(other, None)
    def set_neighbor(self, other, weight=0):
        self.neighbors[other] = weight

    def get_neighbors(self):
        return self.neighbors.keys()
    def get_key(self):
        return self.key

    def __repr__(self):
        return f"Vertex({self.key})"
    def __str__(self):
        return (str(self.key) + " connected to: " + str([x.key for x in
                                                         self.neighbors]))
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                                                            self.neighbors]))
```

Each `Vertex` uses a dictionary to keep track of the vertices to which it is connected and the weight of each edge. Note that the constructor simply initializes the key, which will typically be a string and the `get_neighbor()` method returns the weight of the edge from this vertex to the vertex passed as a parameter.

```
In [6]: class Graph:
    def __init__(self):
        self.vertices = {}

    def set_vertex(self, key):
        self.vertices[key] = Vertex(key)
    def add_edge(self, from_vert, to_vert, weight=0):
        if from_vert not in self.vertices:
            self.set_vertex(from_vert)
        if to_vert not in self.vertices:
            self.set_vertex(to_vert)
        self.vertices[from_vert].set_neighbor(self.vertices[to_vert], weight)

    def get_vertex(self, key):
        return self.vertices.get(key, None)
    def get_vertices(self):
        return self.vertices.keys()

    def __contains__(self, key):
        return key in self.vertices
    def __iter__(self):
        return iter(self.vertices.values())
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    def get_vertex(self, key):
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    def __contains__(self, key):
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    def __iter__(self):
        return iter(self.vertices.values())
```

The `Graph` class, also contains a dictionary that **maps vertex names to vertex objects**. Note that we have implemented the `__iter__` method to make it easy to iterate over all the vertex objects in a particular graph.

Now let us define the graph that we have seen earlier. First we create six vertices numbered 0 through 5. Then we display the vertex dictionary. Notice that for each key 0 through 5 we have created an instance of a `Vertex`:

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```
In [7]: g = Graph()
        for i in range(6):
            g.set_vertex(i)

        g.vertices
```

```
Out[7]: {0: Vertex(0),
         1: Vertex(1),
         2: Vertex(2),
         3: Vertex(3),
         4: Vertex(4),
         5: Vertex(5)}
```

Next, we add the edges that connect the vertices together. Finally, a nested loop verifies that each edge in the graph is properly stored:

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```
In [8]: g.add_edge(0, 1, 5)
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g.add_edge(1, 2, 4)
g.add_edge(2, 3, 9)
g.add_edge(3, 4, 7)
g.add_edge(3, 5, 3)
g.add_edge(4, 0, 1)
g.add_edge(5, 4, 8)
g.add_edge(5, 2, 1)

for v in g:
    for w in v.get_neighbors():
        print(f"({v.get_key()}, {w.get_key()}, {v.get_neighbor(w)})")
```

```
(0, 1, 5)
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(1, 2, 4)
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(4, 0, 1)
(5, 4, 8)
(5, 2, 1)
```

7.7. The Word Ladder Problem

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FOOL
POOL
POLL
POLE
PALE
SALE
SAGE

There are many variations of the word ladder puzzle. For example you might be given a particular number of steps in which to accomplish the transformation, or you might need to use a particular word. In this section, we are interested in figuring out the smallest number of transformations needed to turn the starting word into the ending word.

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Here is an outline of where we are going:

- Represent the relationships between the words as a graph.

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Here is an outline of where we are going:

- Represent the relationships between the words as a graph.
- Use the graph algorithm known as breadth-first search to find an efficient path from the starting word to the ending word.

7.8. Building the Word Ladder Graph

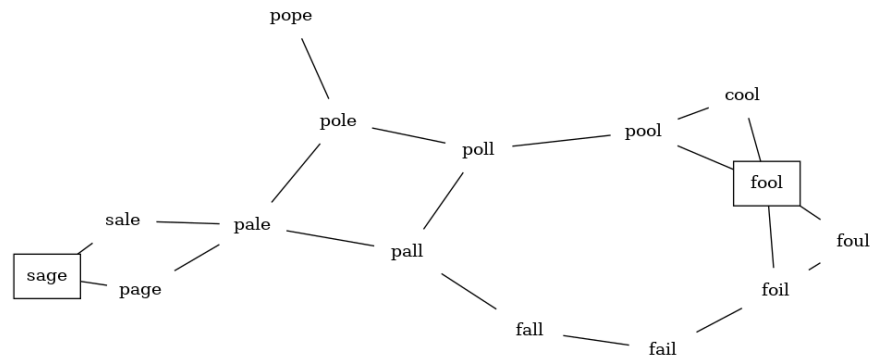
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For a small set of words that approach would work fine. However, let's suppose we have a list of **5,110 words**. Roughly speaking, comparing one word to every other word on the list is an algorithm. For 5,110 words, is more than 26 million comparisons!

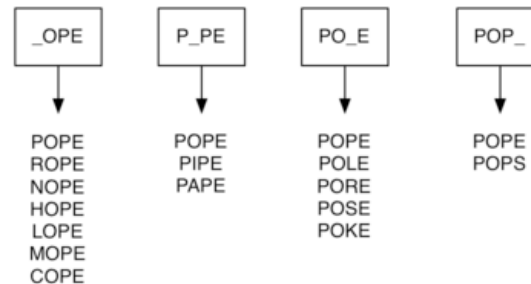
We can do much better by assuming that we have a number of buckets, each labeled with a four-letter word, except that one of the letters on the label has been replaced by an underscore. As we process a list of words, we compare each word with each bucket using the underscore (`_`) as a wildcard.

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Every time we find a matching bucket we put the word in that bucket, so that both `POPE` and `POPS` would both go into the `POP_` bucket. Once we have all the words in the appropriate buckets, we know that all the words in each bucket must be connected!

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```
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from python3.graphs import Graph
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```
In [10]: def build_graph(words):
        buckets = {}
        the_graph = Graph()
        all_words = words
        # create buckets of words that differ by 1 letter
        for line in all_words:
            word = line.strip()
            for i, _ in enumerate(word):
                bucket = f"{word[:i]}_{word[i + 1 :]}"
                buckets.setdefault(bucket, set()).add(word)

        # add edges between different words in the same bucket
        for similar_words in buckets.values():
            for word1 in similar_words:
                for word2 in similar_words - {word1}:
                    the_graph.add_edge(word1, word2)
        return the_graph
```

We can implement the scheme we have just described by using a dictionary. The labels on the buckets we have just described are the keys in our dictionary. The value stored for each key is a list of words.

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Since this is our first real-world graph problem, you might be wondering how sparse the graph is. The list of four-letter words we have for this problem is 5,110 words long. If we were to use an adjacency matrix, the matrix would have cells. The graph constructed by the `build_graph()` function has exactly 53,286 edges, so the matrix would have only 0.20% of the cells filled! That is a very sparse matrix indeed.

7.9. Implementing Breadth-First Search

With the graph constructed we can now turn our attention to the algorithm we will use to find the shortest solution to the word ladder problem. The graph algorithm we are going to use is called the breadth-first search (BFS), and it is one of the easiest algorithms for searching a graph!

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Given a starting vertex of a graph, a breadth first search proceeds by exploring edges in the graph to find all the vertices in for which there is a path from .

The remarkable thing about a breadth-first search is that it finds all the vertices that are a distance from before it finds any vertices that are a distance .

To keep track of its progress, BFS colors each of the vertices white, gray, or black. All the vertices are initialized to white when they are constructed. A white vertex is an undiscovered vertex. When a vertex is initially discovered it is colored gray, and when BFS has completely explored a vertex it is colored black.

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This means that once a vertex is colored black, it has no white vertices adjacent to it. A gray node, on the other hand, may have some white vertices adjacent to it, indicating that there are still additional vertices to explore.

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The `bfs()` shown below uses the adjacency list graph representation we developed earlier. In addition it uses a `Queue`, a crucial point as we will see, to decide which vertex to explore next.


```
In [11]: from pythonds3.basic import Queue
         from pythonds3.graphs import Graph # This is the extended version

def bfs(start):
    start.distance = 0
    start.previous = None
    vert_queue = Queue()
    vert_queue.enqueue(start)
    while vert_queue.size() > 0:
        current = vert_queue.dequeue()
        for neighbor in current.get_neighbors():
            if neighbor.color == "white":
                neighbor.color = "gray"
                neighbor.distance = current.distance + 1
                neighbor.previous = current
                vert_queue.enqueue(neighbor)
        current.color = "black"
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                 current.color = "black"
```

The BFS algorithm uses an extended version of the `Vertex` class that adds three new instance variables: `distance`, `previous`, and `color`. Each of these instance variables also has the appropriate getter and setter methods.

BFS begins at the starting vertex `start` and paints it gray. Two other values, the `distance` and the `previous`, are initialized to 0 and `None` respectively. Finally, `start` is placed on a `Queue`. The next step is to begin to systematically explore vertices at the front of the queue. We explore each new node at the front of the queue by iterating over its adjacency list.

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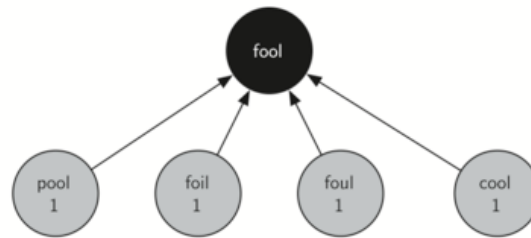
As each node on the adjacency list is examined, its color is checked. If it is white four things happen:

1. The new unexplored vertex `neighbor` is colored gray.
2. The predecessor of `neighbor` is set to the current node `current`.

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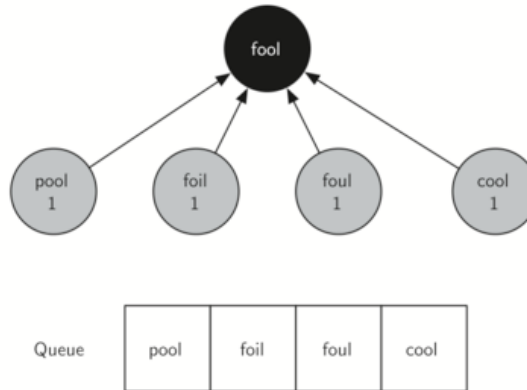
As each node on the adjacency list is examined, its color is checked. If it is white four things happen:

1. The new unexplored vertex `neighbor` is colored gray.
2. The predecessor of `neighbor` is set to the current node `current`.
3. The distance to `neighbor` is set to the distance to `current + 1`.
4. `neighbor` is added to the end of a queue. This effectively schedules this node for further exploration, but not until all the other vertices on the adjacency list of `current` have been explored!

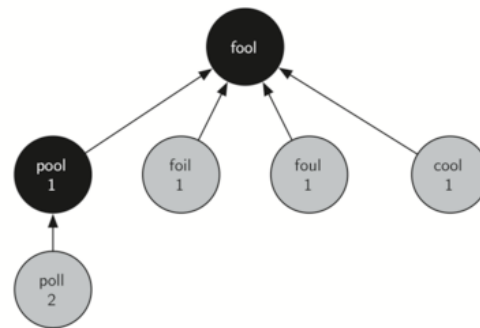


Queue

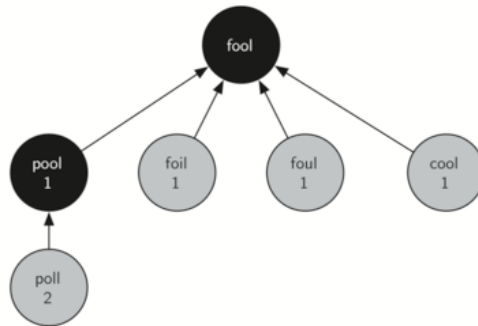
pool	foil	foul	cool
------	------	------	------



Starting from **F00L** we take all nodes that are adjacent to **F00L** and add them to the queue. The adjacent nodes include **P00L**, **FOIL**, **FOUL**, and **COOL**. Each of these nodes are new nodes to expand.



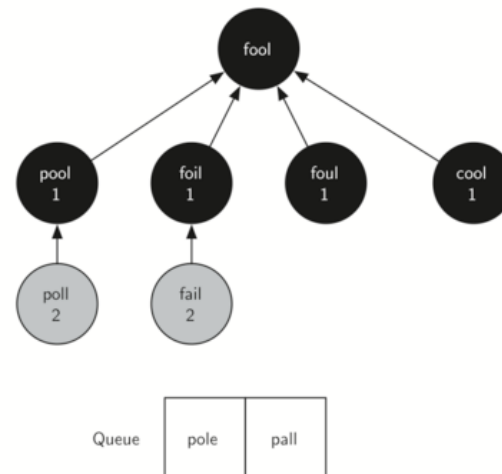
Queue				
	foil	foul	cool	poll

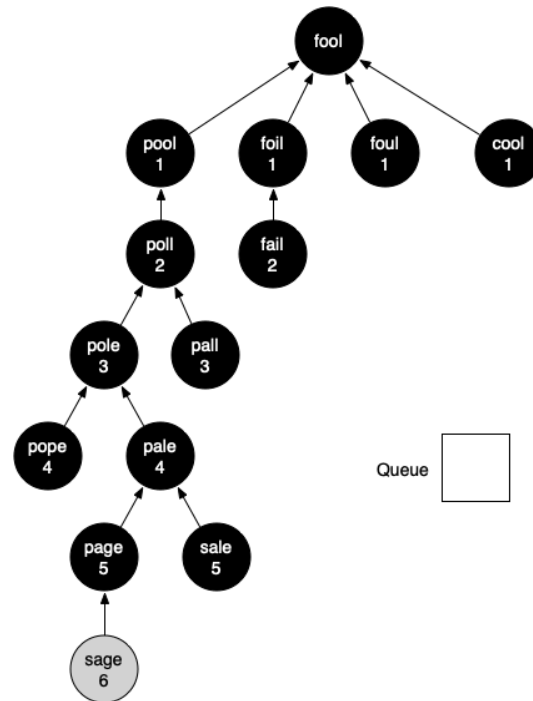


In the next step bfs removes the next node (**POOL**) from the front of the queue and repeats the process for all of its adjacent nodes.

The next vertex on the queue is **FOIL**. The only new node that **FOIL** can add to the tree is **FAIL**. As bfs continues to process the queue, neither of the next two nodes adds anything new to the queue or the tree. Below shows the tree and the queue after expanding all the vertices on the second level of the tree:

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We can start at any vertex in the breadth-first search tree and follow the predecessor arrows back to the root to find the shortest word ladder from any word back to `FOOL`.

```
In [12]: def traverse(starting_vertex):
          current = starting_vertex
          while current:
              print(current.key, end="")
              if current.previous:
                  print("->", end="")
                  current = current.previous
```

```
In [13]: g = build_graph(["fool", "cool", "pool", "poll", "pole", "pall", "fall",  
                          "fail", "foil", "foul", "pope", "pale", "sale", "sage", "page"])  
bfs(g.get_vertex("fool"))  
traverse(g.get_vertex("sage"))
```

sage->sale->pale->pall->poll->pool->fool


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In [13]: g = build_graph(["fool", "cool", "pool", "poll", "pole", "pall", "fall",
                        "fail", "foil", "foul", "pope", "pale", "sale", "sage", "page"])
bfs(g.get_vertex("fool"))
traverse(g.get_vertex("sage"))
```

sage->sale->pale->pall->poll->pool->fool

```
In [14]: header = f"{'Key':^8}|{'Color':^8}|{'Distance':^8}|{'Discover':^4}|{'Closing':^4}|{'Closing':^4}"
print(header)
for key in g.get_vertices():
    vertex = g.get_vertex(key)
    print(vertex)
```

Key	Color	Distance	Discover	Closing	Previous
fool	black	0	0	0	None
cool	black	1	0	0	fool
pool	black	1	0	0	fool
foil	black	1	0	0	fool
foul	black	1	0	0	fool
poll	black	2	0	0	pool
pall	black	3	0	0	poll
pole	black	3	0	0	poll
pale	black	4	0	0	pall
pope	black	4	0	0	pole
fall	black	3	0	0	fail
fail	black	2	0	0	foil
sale	black	5	0	0	pale
page	black	5	0	0	pale
sage	black	6	0	0	sale

7.10. Breadth-First Search Analysis

Let's analyze the run time performance of the breadth-first search algorithm. The first thing to observe is that the `while` loop is executed, at most, one time for each vertex in the graph (up to iterations). You can see that this is true because a vertex must be white before it can be examined and added to the queue.

Let's analyze the run time performance of the breadth-first search algorithm. The first thing to observe is that the `while` loop is executed, at most, one time for each vertex in the graph (up to V iterations). You can see that this is true because a vertex must be white before it can be examined and added to the queue.

This gives us $O(V)$ for the while loop. The `for` loop, which is nested inside the `while`, is executed at most once for each edge in the graph (up to E iterations). The reason is that every vertex is dequeued at most once and we examine an edge from node u to node v only when node u is dequeued.

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This gives us $O(E)$ for the `for` loop. Combining the two loops gives us $O(V + E)$.

Of course doing the breadth-first search is only part of the task. Following the links from the starting node to the goal node is the other part of the task.

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The worst case for this would be if the graph was a single long chain. In this case traversing through all of the vertices would be $O(V)$. The normal case is going to be some fraction of $O(V)$.

Finally, at least for this problem, there is the time required to build the initial graph.

7.11. The Knight's Tour Problem

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The object of the puzzle is to find a sequence of moves that allow the knight to visit every square on the board exactly once. One such sequence is called a **tour**.

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The object of the puzzle is to find a sequence of moves that allow the knight to visit every square on the board exactly once. One such sequence is called a **tour**.

The upper bound on the number of possible legal tours for an chessboard is known to be ; however, there are even more possible dead ends.

Although researchers have studied many different algorithms to solve the knight's tour problem, a graph search is one of the easiest to understand and program. Once again we will solve the problem using two main steps:

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- Represent the legal moves of a knight on a chessboard as a graph.

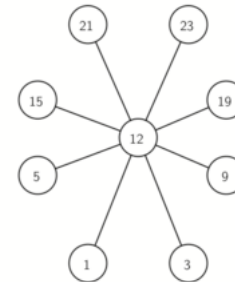
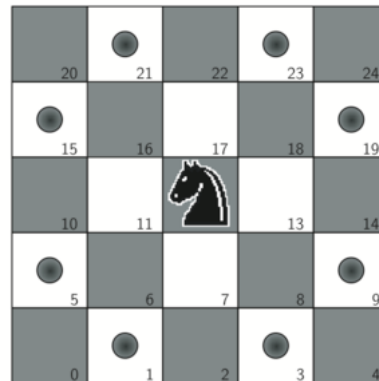
Although researchers have studied many different algorithms to solve the knight's tour problem, a graph search is one of the easiest to understand and program. Once again we will solve the problem using two main steps:

- Represent the legal moves of a knight on a chessboard as a graph.
- Use a graph algorithm to find a path of length n where every vertex on the graph is visited exactly once!

7.12. Building the Knight's Tour Graph

To represent the knight's tour problem as a graph we will use the following two ideas: each square on the chessboard can be represented as a node in the graph and each legal move by the knight can be represented as an edge in the graph.

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```
In [16]: from pythonds3.graphs import Graph

def knight_graph(board_size):
    kt_graph = Graph()
    for row in range(board_size):
        for col in range(board_size):
            node_id = row * board_size + col
            new_positions = gen_legal_moves(row, col, board_size)
            for row2, col2 in new_positions:
                other_node_id = row2 * board_size + col2
                kt_graph.add_edge(node_id, other_node_id)
    return kt_graph
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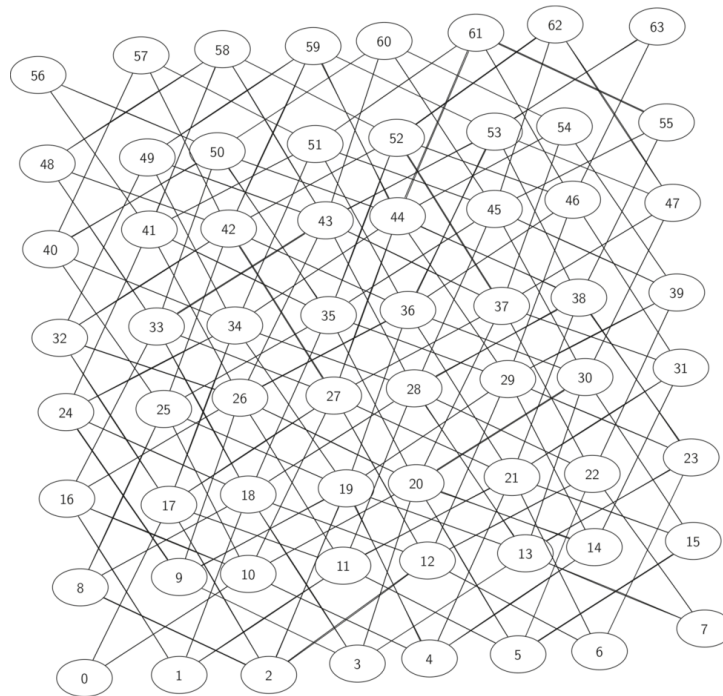
The `knight_graph()` function makes one pass over the entire board. At each square on the board the function calls a helper, `gen_legal_moves()`, to create a list of legal moves for that position on the board.

All legal moves are then converted into edges in the graph. Each location on the board is converted into a linear vertex number.

The `gen_legal_moves()` takes the position of the knight on the board and generates each of the eight possible moves, making sure those moves are still within the board:

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```
In [17]: def gen_legal_moves(row, col, board_size):
    new_moves = []
    move_offsets = [
        (-1, -2), # left-down-down
        (-1, 2),  # left-up-up
        (-2, -1), # left-left-down
        (-2, 1),  # left-left-up
        (1, -2),  # right-down-down
        (1, 2),   # right-up-up
        (2, -1),  # right-right-down
        (2, 1),   # right-right-up
    ]
    for r_off, c_off in move_offsets:
        if 0 <= row + r_off < board_size and 0 <= col + c_off < board_size:
            new_moves.append((row + r_off, col + c_off))
    return new_moves
```



The figure shows the complete graph of possible moves on an board. There are exactly 336 edges in the graph. Notice that the vertices corresponding to the edges of the board have fewer connections (legal moves) than the vertices in the middle of the board.

The figure shows the complete graph of possible moves on an 8x8 board. There are exactly 336 edges in the graph. Notice that the vertices corresponding to the edges of the board have fewer connections (legal moves) than the vertices in the middle of the board.

Once again we can see how sparse the graph is. If the graph was fully connected there would be 4,096 edges. Since there are only 336 edges, the adjacency matrix would be only 8.2 percent full!

7.13. Implementing Knight's Tour

The search algorithm we will use to solve the knight's tour problem is called depth-first search (DFS). Whereas the breadth-first search algorithm builds a **search tree** one level at a time, a depth-first search creates a search tree by exploring one branch of the tree as deeply as possible.

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We will look at two algorithms that implement DFS. The first algorithm we will look at specifically solves the knight's tour problem by explicitly **forbidding a node to be visited more than once**.

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We will look at two algorithms that implement DFS. The first algorithm we will look at specifically solves the knight's tour problem by explicitly **forbidding a node to be visited more than once**.

The second implementation is more general, but allows nodes to be visited more than once as the tree is constructed. The second version is used in subsequent sections to develop additional graph algorithms.

The depth-first exploration of the graph is exactly what we need in order to find a path through 64 vertices (one for each square on the chessboard) and 63 edges.

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The `knight_tour()` below takes four parameters: `n`, the current depth in the search tree; `path`, a `list` of vertices visited up to this point; `u`, the vertex in the graph we wish to explore; and `limit`, the number of nodes in the path.

```
In [18]: def knight_tour(n, path, u, limit):
    u.color = "gray"
    path.append(u)
    if n < limit:
        neighbors = sorted(list(u.get_neighbors()))
        i = 0
        done = False
        while i < len(neighbors) and not done:
            if neighbors[i].color == "white":
                done = knight_tour(n + 1, path, neighbors[i], limit)
                i = i + 1
            if not done: # prepare to backtrack
                path.pop()
                u.color = "white" # Not added to path yet return to unvisited
        else:
            done = True
    return done
```

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    return done
```

The function is recursive. When the function is called, it first checks the base case condition. If we have a path that contains 64 vertices, we return from `knight_tour()` with a status of `True`, indicating that we have found a successful tour. If the path is not long enough, we continue to explore one level deeper by choosing a new vertex to explore and calling it recursively for that vertex.

DFS also uses colors to keep track of which vertices in the graph have been visited. Unvisited vertices are colored white, and visited vertices are colored gray. If all neighbors of a particular vertex have been explored and we have not yet reached our goal length of 64 vertices, we have reached a dead end and must backtrack.

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Backtracking happens when we return from `knight_tour()` with a status of `False`. In the breadth-first search we used a queue to keep track of which vertex to visit next. Since depth-first search is recursive, we are implicitly using a **stack** to help us with our backtracking.

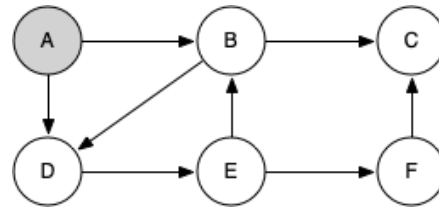
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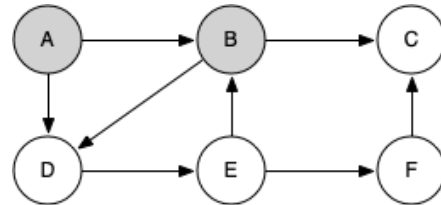
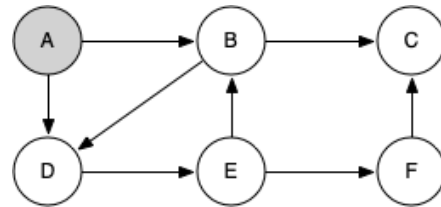
When we return from a call with a status of `False`, we remain inside the `while` loop and look at the next vertex in neighbors.

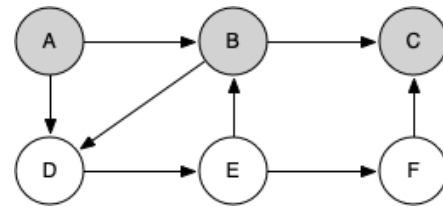
Let's look at a simple example of `knight_tour()` in action. You can refer to the figures below to follow the steps of the search. For this example, we will assume that the call to the `get_neighbors()` method on line 5 orders the nodes in alphabetical order. We begin by calling `knight_tour(1, path, A, 6)`.

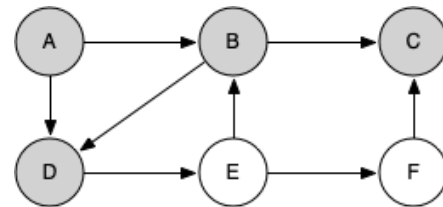
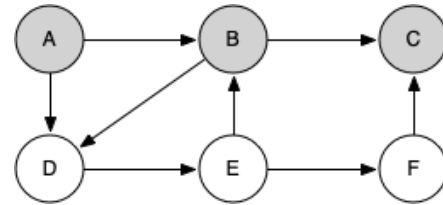
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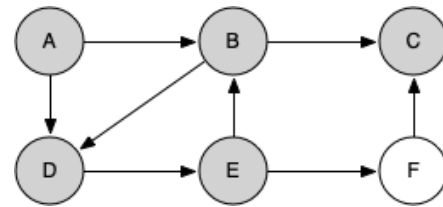


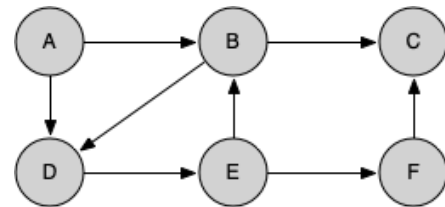
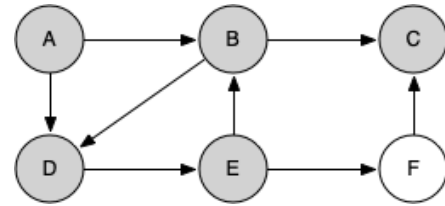
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```
In [19]: board_size = 8
kt_graph = knight_graph(board_size)
path = []
start = kt_graph.get_vertex(0)
finished = knight_tour(1, path, start, board_size * board_size)
if finished:
    solution = [(v.get_key() // board_size, v.get_key() % board_size)
                for v in path]
    print("One possible path:")
    print(solution)
else:
    print("No path found.")
```

One possible path:

```
[(0, 0), (1, 2), (0, 4), (1, 6), (2, 4), (0, 3), (1, 1), (2, 3), (0,
2), (1, 0), (2, 2), (0, 1), (1, 3), (0, 5), (1, 7), (2, 5), (0, 6),
(1, 4), (2, 6), (0, 7), (1, 5), (2, 7), (3, 5), (4, 3), (3, 1), (5,
0), (4, 2), (2, 1), (3, 3), (4, 1), (2, 0), (3, 2), (4, 0), (6, 1),
(5, 3), (3, 4), (4, 6), (6, 7), (7, 5), (5, 4), (7, 3), (6, 5), (7,
7), (5, 6), (3, 7), (4, 5), (5, 7), (3, 6), (4, 4), (6, 3), (7, 1),
(5, 2), (6, 0), (7, 2), (6, 4), (7, 6), (5, 5), (4, 7), (6, 6), (7,
4), (6, 2), (7, 0), (5, 1), (3, 0)]
```

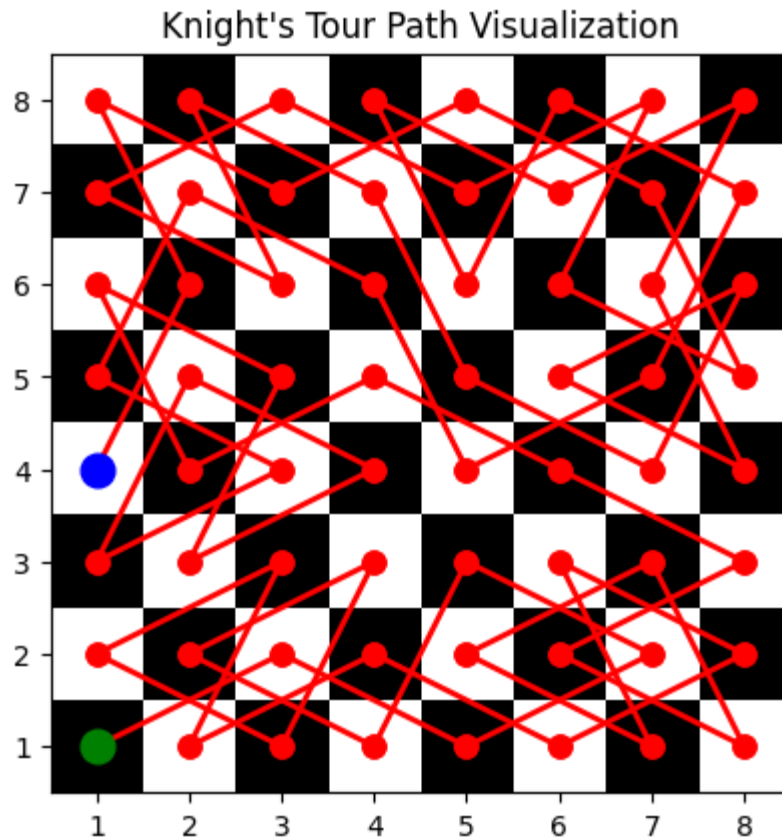
```

In [20]: import matplotlib.pyplot as plt
def visualize_knight_path(board_size, path):
    # Create a board with alternating colors
    board = [[(i+j) % 2 for j in range(board_size)] for i in range(board_size)]
    fig, ax = plt.subplots()
    ax.imshow(board, cmap='gray', interpolation='nearest')
    # Draw the path
    for i in range(len(path) - 1):
        start = path[i]
        end = path[i + 1]
        ax.plot([start[1], end[1]], [start[0], end[0]], 'ro-', linewidth=2, ma
    # Mark the start and end points
    start = path[0]
    end = path[-1]
    ax.plot(start[1], start[0], 'go', markersize=12) # start point in green
    ax.plot(end[1], end[0], 'bo', markersize=12) # end point in blue
    # Set up the plot grid, labels, and title
    ax.set_xticks(range(board_size))
    ax.set_yticks(range(board_size))
    ax.set_xticklabels(range(1, board_size + 1))
    ax.set_yticklabels(range(1, board_size + 1))
    ax.set_title("Knight's Tour Path Visualization")
    # Show the plot
    plt.gca().invert_yaxis() # Invert y-axis to match the chessboard represen
    plt.show()

```



```
In [21]: visualize_knight_path(board_size, solution) #NP-hard
```



```
In [22]: header = f"{'Key':^8}|{'Color':^8}|{'Distance':^8}|{'Discover':^4}|{'Closing':  
print(header)  
for key in kt_graph.get_vertices():  
    vertex = kt_graph.get_vertex(key)  
    print(vertex)
```

Key	Color	Distance	Discover	Closing	Previous
0	gray	inf	0	0	None
10	gray	inf	0	0	None
17	gray	inf	0	0	None
1	gray	inf	0	0	None
11	gray	inf	0	0	None
16	gray	inf	0	0	None
18	gray	inf	0	0	None
2	gray	inf	0	0	None
8	gray	inf	0	0	None
12	gray	inf	0	0	None
19	gray	inf	0	0	None
3	gray	inf	0	0	None
9	gray	inf	0	0	None
13	gray	inf	0	0	None
20	gray	inf	0	0	None
4	gray	inf	0	0	None
14	gray	inf	0	0	None
21	gray	inf	0	0	None
5	gray	inf	0	0	None
15	gray	inf	0	0	None
22	gray	inf	0	0	None
6	gray	inf	0	0	None
23	gray	inf	0	0	None
7	gray	inf	0	0	None
25	gray	inf	0	0	None
24	gray	inf	0	0	None
26	gray	inf	0	0	None
27	gray	inf	0	0	None
28	gray	inf	0	0	None
29	gray	inf	0	0	None

30	gray	inf	0	0	None
31	gray	inf	0	0	None
33	gray	inf	0	0	None
32	gray	inf	0	0	None
34	gray	inf	0	0	None
35	gray	inf	0	0	None
36	gray	inf	0	0	None
37	gray	inf	0	0	None
38	gray	inf	0	0	None
39	gray	inf	0	0	None
41	gray	inf	0	0	None
40	gray	inf	0	0	None
42	gray	inf	0	0	None
43	gray	inf	0	0	None
44	gray	inf	0	0	None
45	gray	inf	0	0	None
46	gray	inf	0	0	None
47	gray	inf	0	0	None
49	gray	inf	0	0	None
48	gray	inf	0	0	None
50	gray	inf	0	0	None
51	gray	inf	0	0	None
52	gray	inf	0	0	None
53	gray	inf	0	0	None
54	gray	inf	0	0	None
55	gray	inf	0	0	None
57	gray	inf	0	0	None
56	gray	inf	0	0	None
58	gray	inf	0	0	None
59	gray	inf	0	0	None
60	gray	inf	0	0	None

61		gray		inf		0		0		None
62		gray		inf		0		0		None
63		gray		inf		0		0		None

7.14. Knight's Tour Analysis

`knight_tour()` is very sensitive to the method you use to select the next vertex to visit. For example, on a board you can produce a path in about 1.5 seconds on a reasonably fast computer. But what happens if you try an board? In this case, depending on the speed of your computer, you may have to wait up to several minutes to get the results!

`knight_tour()` is very sensitive to the method you use to select the next vertex to visit. For example, on a 8x8 board you can produce a path in about 1.5 seconds on a reasonably fast computer. But what happens if you try an 10x10 board? In this case, depending on the speed of your computer, you may have to wait up to several minutes to get the results!

The reason for this is that the knight's tour problem as we have implemented it so far is an exponential algorithm of size $O(2^n)$, where n is the number of squares on the chess board, and c is a small constant.

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The reason for this is that the knight's tour problem as we have implemented it so far is an exponential algorithm of size $O(2^N)$, where N is the number of squares on the chess board, and c is a small constant.

We can use a tree to help us understand this point. The root of the tree represents the starting point of the search. From there the algorithm generates and checks each of the possible moves the knight can make.

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The reason for this is that the knight's tour problem as we have implemented it so far is an exponential algorithm of size $O(2^N)$, where N is the number of squares on the chess board, and 2 is a small constant.

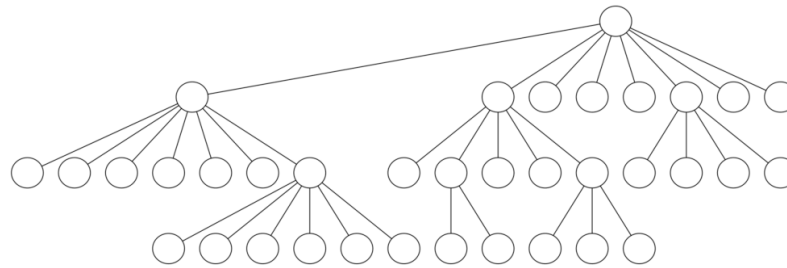
We can use a tree to help us understand this point. The root of the tree represents the starting point of the search. From there the algorithm generates and checks each of the possible moves the knight can make.

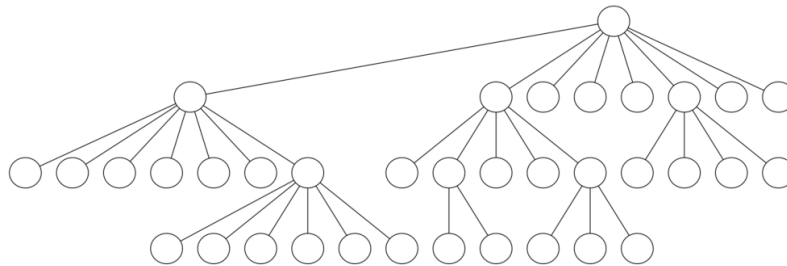
As we have noted before, the number of moves possible depends on the position of the knight on the board. In the corners there are only two legal moves, on the squares adjacent to the corners there are three, and in the middle of the board there are eight.

2	3	4	4	4	4	3	2
3	4	6	6	6	6	4	3
4	6	8	8	8	8	6	4
4	6	8	8	8	8	6	4
4	6	8	8	8	8	6	4
4	6	8	8	8	8	6	4
3	4	6	6	6	6	4	3
2	3	4	4	4	4	3	2

2	3	4	4	4	4	3	2
3	4	6	6	6	6	4	3
4	6	8	8	8	8	6	4
4	6	8	8	8	8	6	4
4	6	8	8	8	8	6	4
4	6	8	8	8	8	6	4
3	4	6	6	6	6	4	3
2	3	4	4	4	4	3	2

At the next level of the tree there are once again between two and eight possible next moves from the position we are currently exploring. The number of possible positions to examine corresponds to the number of nodes in the search tree.





For a tree with nodes that may have up to eight children, the number of nodes is large. Because the branching factor of each node is variable, we could estimate the number of nodes using an **average branching factor**. The important thing to note is that this algorithm is exponential: $O(b^d)$, where b is the average branching factor for the board.

Let's look at how rapidly this grows! For a board that is the tree will be 25 levels deep. The average branching factor is so the number of nodes in the search tree is or .

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For a board, , there are nodes, and for a regular chess board, , there are !

Of course, since there are multiple solutions to the problem we won't have to explore every single node, but the fractional part of the nodes we do have to explore is just a constant multiplier which does not change the exponential nature of the problem.

Luckily there is a way to speed up the case so that it runs in under one second. The `order_by_avail()` will be used in place of the call to `u.get_neighbors()`. The critical line in the function is line 5. This line ensures that we select the vertex that has the fewest available moves to go next.

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```
In [23]: def order_by_avail(n):  
         res_list = []  
         for v in n.get_neighbors():  
             if v.color == "white":  
                 c = 0  
                 for w in v.get_neighbors():  
                     if w.color == "white":  
                         c = c + 1  
                 res_list.append((c, v))  
         res_list.sort(key=lambda x: x[0])  
         return [y[1] for y in res_list]
```

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         res_list.sort(key=lambda x: x[0])
         return [y[1] for y in res_list]
```

You might think this is really counterproductive; why not select the node that has the most available moves?

The problem with using the vertex with the most available moves as your next vertex on the path is that it tends to have the knight visit the middle squares early on in the tour!

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When this happens it is easy for the knight to get stranded on one side of the board where it cannot reach unvisited squares on the other side of the board. On the other hand, visiting the squares with the fewest available moves first pushes the knight to visit the squares around the edges of the board first.

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When this happens it is easy for the knight to get stranded on one side of the board where it cannot reach unvisited squares on the other side of the board. On the other hand, visiting the squares with the fewest available moves first pushes the knight to visit the squares around the edges of the board first.

This ensures that the knight will visit the hard-to-reach corners early and can use the middle squares to **hop across the board only when necessary**. Utilizing this kind of knowledge to speed up an algorithm is called a heuristic. This particular heuristic is called Warnsdorff's algorithm.

```
In [24]: def knight_tour(n, path, u, limit):
          u.color = "gray"
          path.append(u)
          if n < limit:
              neighbors = order_by_avail(u)
              i = 0
              done = False
              while i < len(neighbors) and not done:
                  if neighbors[i].color == "white":
                      done = knight_tour(n + 1, path, neighbors[i], limit)
                      i = i + 1
              if not done: # prepare to backtrack
                  path.pop()
                  u.color = "white"
          else:
              done = True
          return done
```

```

In [25]: board_size = 8
kt_graph = knight_graph(board_size)
path = []
start = kt_graph.get_vertex(0)
finished = knight_tour(1, path, start, board_size * board_size)
if finished:
    solution = [(v.get_key() // board_size, v.get_key() % board_size)
                for v in path]
    print("One possible path:")
    print(solution)
else:
    print("No path found.")

```

One possible path:

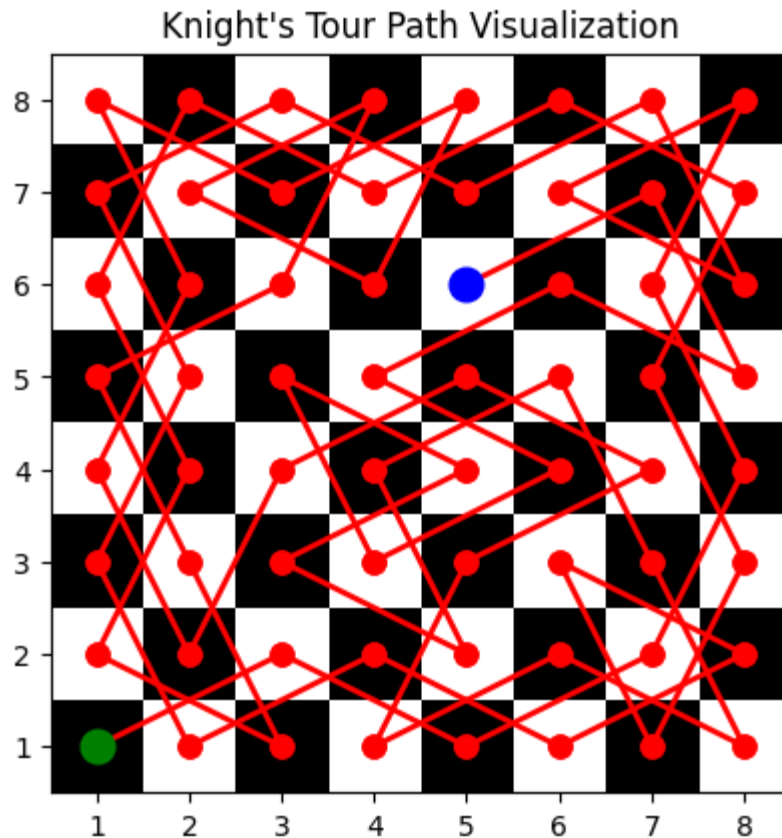
```

[(0, 0), (1, 2), (0, 4), (1, 6), (3, 7), (5, 6), (7, 7), (6, 5), (5,
7), (7, 6), (6, 4), (7, 2), (6, 0), (4, 1), (2, 0), (0, 1), (1, 3),
(0, 5), (1, 7), (2, 5), (0, 6), (2, 7), (4, 6), (6, 7), (7, 5), (6,
3), (7, 1), (5, 0), (3, 1), (1, 0), (0, 2), (2, 1), (4, 0), (5, 2),
(7, 3), (6, 1), (5, 3), (7, 4), (6, 2), (7, 0), (5, 1), (3, 0), (1,
1), (3, 2), (4, 4), (3, 6), (2, 4), (0, 3), (1, 5), (0, 7), (2, 6),
(4, 5), (3, 3), (1, 4), (2, 2), (3, 4), (4, 2), (2, 3), (3, 5), (4,
3), (5, 5), (4, 7), (6, 6), (5, 4)]

```



```
In [26]: visualize_knight_path(board_size, solution)
```



7.15. General Depth-First Search

The knight's tour is a special case of a depth-first search where the goal is to create the deepest depth-first tree **without any branches**.

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The more general depth-first search is actually easier. Its goal is to search as deeply as possible, connecting as many nodes in the graph as possible and branching where necessary.

The knight's tour is a special case of a depth-first search where the goal is to create the deepest depth-first tree **without any branches**.

The more general depth-first search is actually easier. Its goal is to search as deeply as possible, connecting as many nodes in the graph as possible and branching where necessary.

It is even possible that a depth-first search will create more than one tree. When the depth-first search algorithm creates a group of trees we call this a **depth-first forest**.

As with the breadth-first search, our depth-first search makes use of predecessor links to construct the tree. In addition, the depth-first search will make use of two additional instance variables in the `Vertex` class. The new instance variables are the discovery and closing times.

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- The discovery time tracks the number of steps in the algorithm before a vertex is first encountered.
- The closing time is the number of steps in the algorithm before a vertex is colored black.

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- The discovery time tracks the number of steps in the algorithm before a vertex is first encountered.
- The closing time is the number of steps in the algorithm before a vertex is colored black.

As we will see after looking at the algorithm, the discovery and closing times of the nodes provide some interesting properties we can use in later algorithms.


```
In [27]: class DFSGraph(Graph):
    def __init__(self):
        super().__init__()
        self.time = 0

    def dfs(self):
        for vertex in self:
            vertex.color = "white"
            vertex.previous = -1
        for vertex in self:
            if vertex.color == "white":
                self.dfs_visit(vertex)

    def dfs_visit(self, start):
        start.color = "gray"
        self._time = self._time + 1
        start.discovery_time = self._time
        for next_vertex in start.get_neighbors():
            if next_vertex.color == "white":
                next_vertex.set_previous(start)
                self.dfs_visit(next_vertex)
        start.color = "black"
        self._time = self._time + 1
        start.closing_time = self._time
```

You will notice that the `dfs()` method iterates over all of the vertices in the graph calling `dfs_visit()` on the nodes that are white.

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The reason we iterate over all the nodes, rather than simply searching from a chosen starting node, is to make sure that all nodes in the graph are considered and that no vertices are left out of the depth-first forest.

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The reason we iterate over all the nodes, rather than simply searching from a chosen starting node, is to make sure that all nodes in the graph are considered and that no vertices are left out of the depth-first forest.

It may look unusual to see the statement `for vertex in self`, but remember that in this case `self` is an instance of the `DFSGraph` class, and iterating over all the vertices in an instance of a graph is a natural thing to do.

The `dfs_visit()` method starts with a single vertex called `start_vertex` and explores all of the neighboring white vertices as deeply as possible.

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If you look carefully at the code for `dfs_visit()` and compare it to breadth-first search, what you should notice is that the algorithm is almost identical to `bfs()` except that on the last line of the inner for loop, `dfs_visit()` calls itself recursively to continue the search at a deeper level, whereas `bfs()` adds the node to a queue for later exploration!

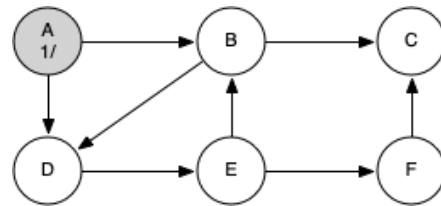
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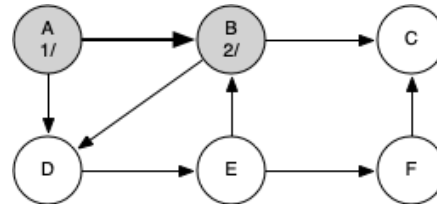
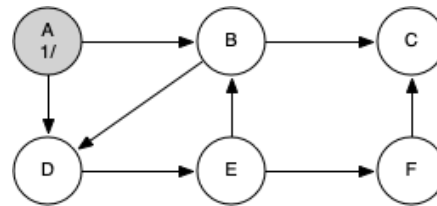
It is interesting to note that where `bfs()` uses a queue, `dfs_visit()` uses a stack. You don't see a stack in the code, but it is implicit in the recursive call.

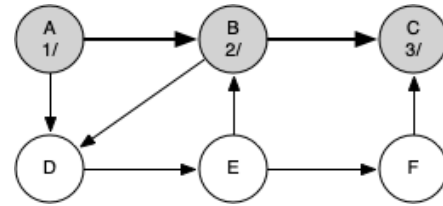
The following sequence of figures illustrates the depth-first search algorithm in action for a small graph. In these figures, the dotted lines indicate edges that are checked, but the node at the other end of the edge has already been added to the depth-first tree.

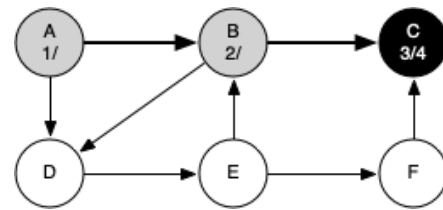
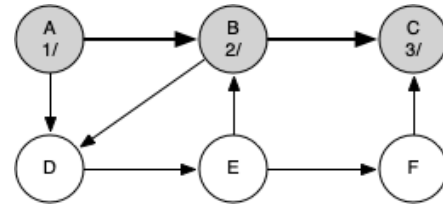
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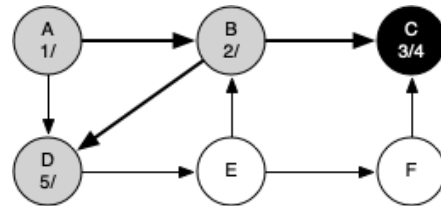
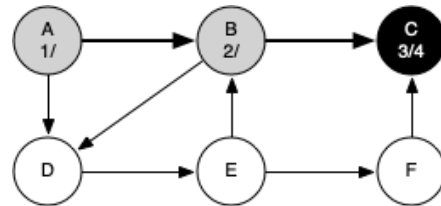
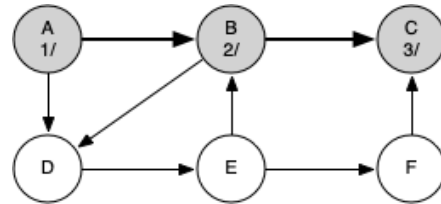


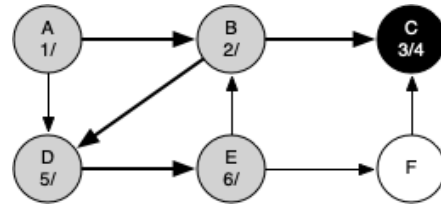
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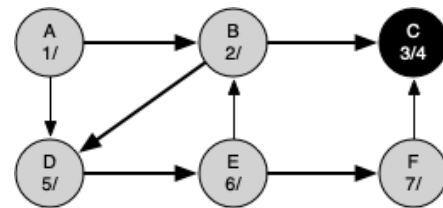
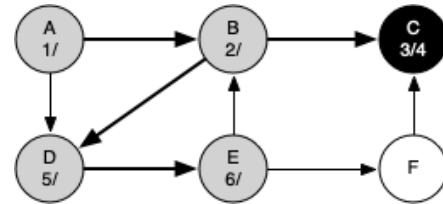


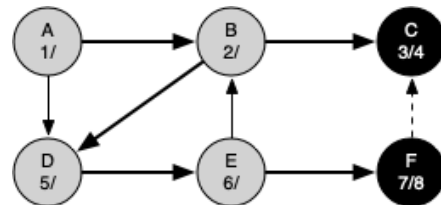
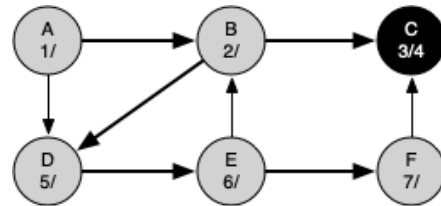
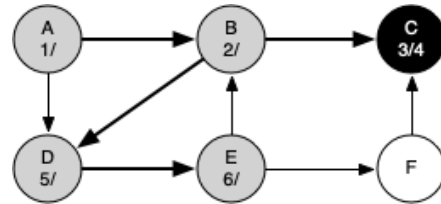


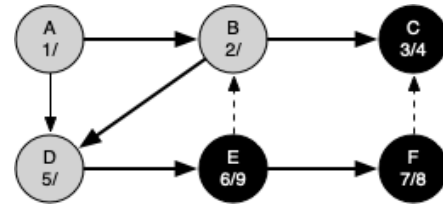


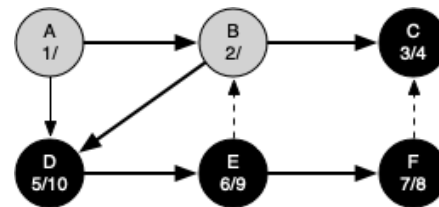
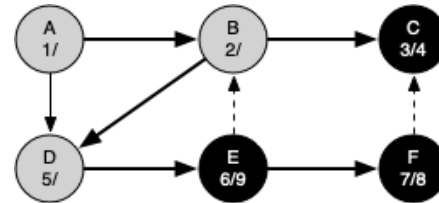


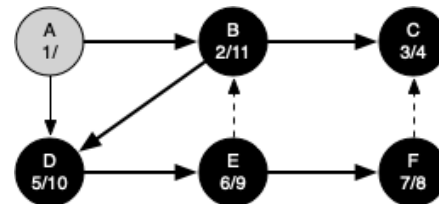
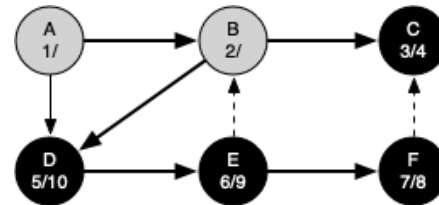
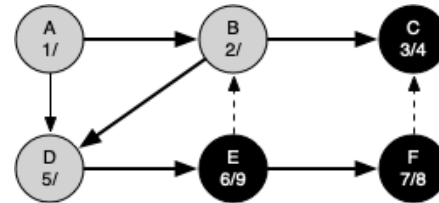


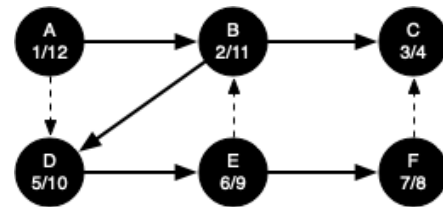


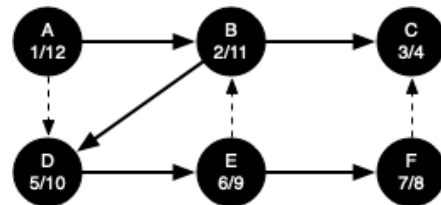












The discovery and closing times for each node display a property called the parenthesis property. This property means that all the children of a particular node in the depth-first tree have a later discovery time and an earlier closing time than their parent.

In [28]:

```
g = DFSGraph()
g.add_edge('A', 'B')
g.add_edge('B', 'C')
g.add_edge('A', 'D')
g.add_edge('B', 'D')
g.add_edge('D', 'E')
g.add_edge('E', 'B')
g.add_edge('E', 'F')
g.add_edge('F', 'C')

# Run DFS
g.dfs()

header = f"{'Key':^8}|{'Color':^8}|{'Distance':^8}|{'Discover':^4}|\\
{'Closing':^8}|{'Previous'}"
print(header)
# Print vertex states post-DFS
for key in g.get_vertices():
    vertex = g.get_vertex(key)
    print(vertex)
```

Key	Color	Distance	Discover	Closing	Previous
A	black	inf	1	12	None
B	black	inf	2	11	A
C	black	inf	3	4	B
D	black	inf	5	10	B
E	black	inf	6	9	D
F	black	inf	7	8	E

7.16. Depth-First Search Analysis

The general running time for depth-first search is as follows. The loops in `dfs()` run in $\mathcal{O}(V)$, not counting what happens in `dfs_visit()`, since it is executed once for each vertex in the graph.

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In `dfs_visit()` the loop is executed once for each edge in the adjacency list of the current vertex. Since it is only called recursively if the vertex is white, the loop will execute a maximum of once for every edge in the graph, or $\mathcal{O}(E)$. Therefore, the total time for depth-first search is $\mathcal{O}(V + E)$.

In [32]: `display_quiz(quiz_path+"bfs_dfs.json", max_width=800)`

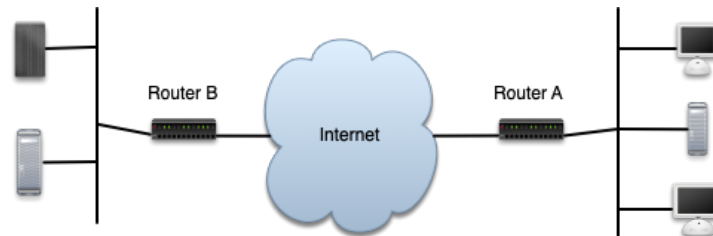
What is the primary structural difference in how Breadth-First Search (BFS) and Depth-First Search (DFS) track the next vertex to explore?

- ☐ BFS tracks vertices using a dictionary, while DFS only uses a simple list.
- ☐ Both algorithms must use a priority queue to ensure the correct search order.
- ☐ BFS uses a queue to explore layer by layer, while DFS uses a stack (often via recursion) to explore as deep as possible.
- ☐ BFS uses a stack while DFS uses a queue.

7.19. Shortest Path Problems

When you surf the web, send an email, or log in to a laboratory computer from another location on campus, a lot of work is going on behind the scenes to get the information on your computer transferred to another computer. The in-depth study of how information flows from one computer to another over the internet is the primary topic for a class in computer networking.

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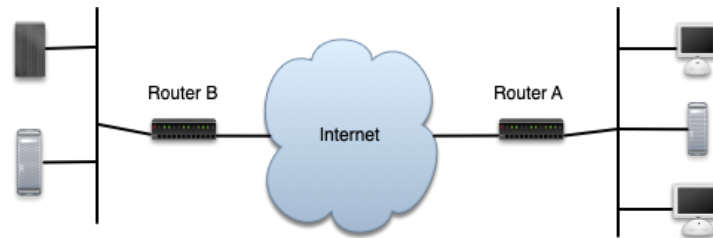


Figure above shows you a high-level overview of how communication on the internet works. When you use your browser to request a web page from a server, the request must travel over your local area network and out onto the internet through a router.

Each router on the internet is connected to one or more other routers. If you run the `tracert` command at different times of the day, you are likely to see that your information flows through different routers at different times. This is because there is a cost associated with each connection between a pair of routers that depends on the volume of traffic, the time of day, and many other factors.

Each router on the internet is connected to one or more other routers. If you run the `traceroute` command at different times of the day, you are likely to see that your information flows through different routers at different times. This is because there is a cost associated with each connection between a pair of routers that depends on the volume of traffic, the time of day, and many other factors.

By this time it will not surprise you to learn that we can represent the network of routers as a graph with weighted edges.

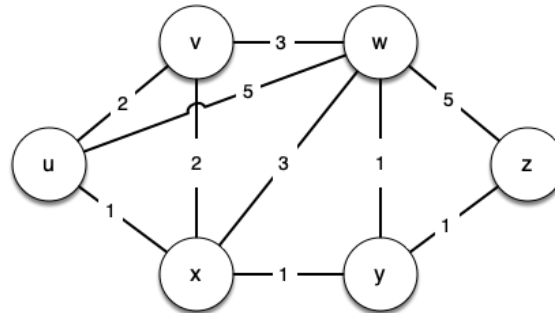
Each router on the internet is connected to one or more other routers. If you run the `traceroute` command at different times of the day, you are likely to see that your information flows through different routers at different times. This is because there is a cost associated with each connection between a pair of routers that depends on the volume of traffic, the time of day, and many other factors.

By this time it will not surprise you to learn that we can represent the network of routers as a graph with weighted edges.

Figure below shows a small example of a weighted graph that represents the interconnection of routers in the internet. The problem that we want to solve is to find the shortest path, one with the smallest total weight along which to route any given message.

This problem should sound familiar because it is similar to the problem we solved using a breadth-first search, except that here we are concerned with the total weight of the path rather than the number of hops in the path. It should be noted that if all the weights are equal, the problem is the same!

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```
In [34]: display_quiz(quiz_path+"bfs.json", max_width=800)
```

Why is Breadth-First Search (BFS) guaranteed to find the shortest path in an unweighted graph?

- ☐ It explores all vertices at distance k from the starting vertex before moving to vertices at distance $k+1$.
- ☐ It marks vertices as black only when the shortest path has been confirmed.
- ☐ It uses a greedy approach to prune paths that are too long.
- ☐ It always chooses the edge with the smallest numerical weight first.

7.20. Dijkstra's Algorithm

Dijkstra's algorithm is an iterative algorithm that provides us with the shortest path from one particular starting node to all other nodes in the graph. Again this is similar to the results of a breadth-first search.

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The algorithm iterates once for every vertex in the graph; however, the order that it iterates over the vertices is controlled by a priority queue. The value that is used to determine the order of the objects in the priority queue is **distance**.

A priority queue acts like a queue in that you dequeue an item by removing it from the front. However, in a priority queue the logical order of items inside a queue is determined by their priority. The highest priority items are at the front of the queue and the lowest priority items are at the back.

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Thus when you enqueue an item on a priority queue, the new item may move all the way to the front.

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Thus when you enqueue an item on a priority queue, the new item may move all the way to the front.

When a vertex is first created, distance is set to a very large number. Theoretically you would set distance to infinity, but in practice we just set it to a number that is larger than any real distance we would have in the problem we are trying to solve.

```

In [37]: from pythonds3.trees.priority_queue import PriorityQueue #heapq
# If checking the visited node, the algorithm can terminate
# when there is negative edge
def dijkstra(graph, start):
    pq = PriorityQueue()
    start.distance = 0
    pq.insert((start.distance, start))
    visited = set() # Set to track visited vertices

    while not pq.is_empty():
        print(pq)
        distance, current_v = pq.delete()
        # Mark the vertex as visited once it's pulled from the queue
        visited.add(current_v)

        # Process each adjacent vertex
        for next_v in current_v.get_neighbors():
            if next_v in visited:
                continue # Skip if next vertex is already visited
            new_distance = distance + current_v.get_neighbor(next_v)
            if new_distance < next_v.distance:
                next_v.distance = new_distance
                next_v.previous = current_v
                if next_v in pq:
                    pq.change_priority(next_v, new_distance)
                else:
                    pq.insert((new_distance, next_v))

```

The `PriorityQueue` class stores tuples of `(priority, key)` pairs. This is an important point, because Dijkstra's algorithm requires the key in the priority queue to match the key of the vertex in the graph.

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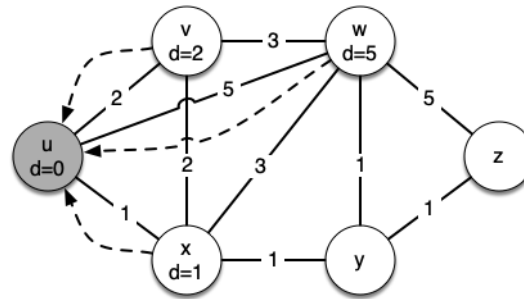
Note that there is a `change_priority()` method. This method is used when the distance to a vertex that is already in the queue is reduced, and thus the vertex is moved toward the front of the queue!

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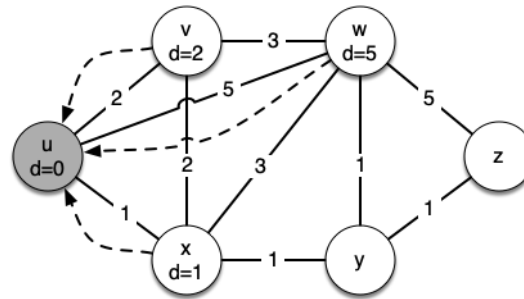
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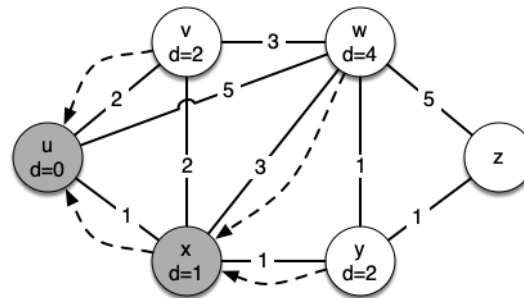
Let's walk through it. We begin with the vertex `s`. The three vertices adjacent to it are `a`, `b`, and `c`. Since the initial distances to them are all initialized to `sys.maxsize`, the new costs to get to them through the start node are all their direct costs. So we update the costs to each of these three nodes. We also set the predecessor for each node to `s` and we add each node to the priority queue. We use the distance as the key for the priority queue.



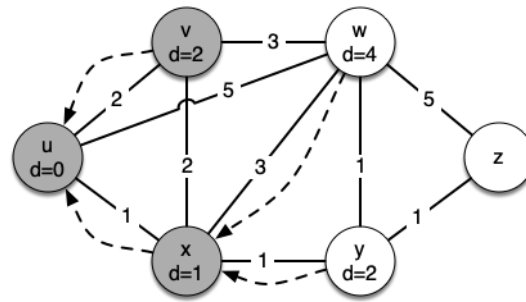
PQ = x, v, w



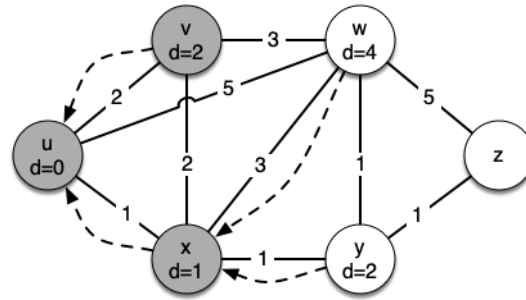
PQ = x, v, w



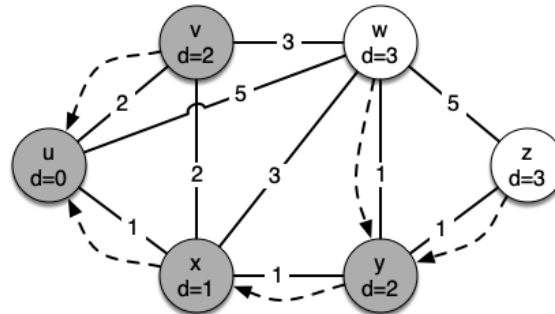
PQ = v, y, w



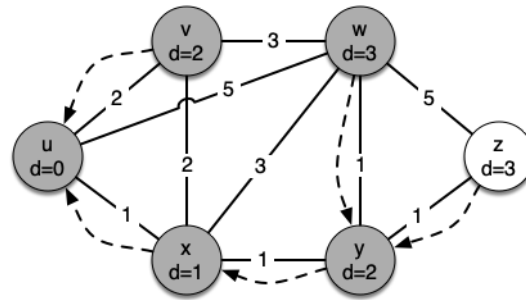
PQ = y, w



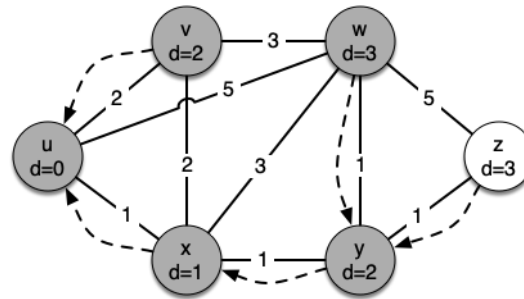
PQ = y, w



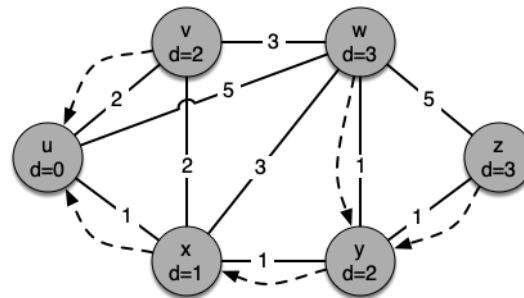
PQ = w, z



PQ = z



PQ = z



PQ = None

In [38]:

```
g = Graph()
g.add_edge('u', 'v', 2)
g.add_edge('v', 'u', 2)
g.add_edge('v', 'w', 3)
g.add_edge('w', 'v', 3)
g.add_edge('w', 'z', 5)
g.add_edge('z', 'w', 5)
g.add_edge('u', 'x', 1)
g.add_edge('x', 'u', 1)
g.add_edge('u', 'w', 5)
g.add_edge('w', 'u', 5)
g.add_edge('x', 'v', 2)
g.add_edge('v', 'x', 2)
g.add_edge('x', 'w', 3)
g.add_edge('w', 'x', 3)
g.add_edge('x', 'y', 1)
g.add_edge('y', 'x', 1)
g.add_edge('y', 'w', 1)
g.add_edge('w', 'y', 1)
g.add_edge('y', 'z', 1)
g.add_edge('z', 'y', 1)
```

In [39]:

```
# Run
dijkstra(g, g.get_vertex('u'))

header = f"{'Key':^8}|{'Color':^8}|{'Distance':^8}|{'Discover':^8}|\\
{'Closing':^8}|{'Previous'}"
print(header)
# Print vertex states
for key in g.get_vertices():
    vertex = g.get_vertex(key)
    print(vertex)
```

```
[(0, Vertex(u))]
```

```
[(1, Vertex(x)), (2, Vertex(v)), (5, Vertex(w))]
```

```
[(2, Vertex(v)), (4, Vertex(w)), (2, Vertex(y))]
```

```
[(2, Vertex(y)), (4, Vertex(w))]
```

```
[(3, Vertex(w)), (3, Vertex(z))]
```

```
[(3, Vertex(z))]
```

Key	Color	Distance	Discover	Closing	Previous
u	white	0	0	0	None
v	white	2	0	0	u
w	white	3	0	0	y
z	white	3	0	0	y
x	white	1	0	0	u
y	white	2	0	0	x


```

In [40]: def find_path(graph, start_key, end_key):
    start_vertex = graph.get_vertex(start_key)
    end_vertex = graph.get_vertex(end_key)
    path = []
    current_vertex = end_vertex

    # Backtrack from the end vertex to the start
    # vertex using the 'previous' pointers
    while current_vertex is not None and current_vertex != start_vertex:
        path.append(current_vertex._key)
        current_vertex = current_vertex.previous

    # Check if a path exists, i.e., we are back at the start vertex
    if current_vertex is None:
        return "No path"
    else: # Include the start vertex in the path
        path.append(start_vertex._key)
        path.reverse() # Reverse the path to show it from start to end
        return path

# Example of how to use the find_path function
dijkstra(g, g.get_vertex('u')) # Run Dijkstra from 'u'
path = find_path(g, 'u', 'z')
print("Path from 'u' to 'z':", " -> ".join(path))

```

```

[(0, Vertex(u))]
Path from 'u' to 'z': u -> x -> y -> z

```

It is important to note that Dijkstra's algorithm works only when the weights are all **positive**. You should convince yourself that if you introduced a negative weight on one of the edges of the graph, the algorithm would never exit.

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We will note that to route messages through the internet, other algorithms are used for finding the shortest path. One of the problems with using Dijkstra's algorithm on the internet is that you must have a complete representation of the graph in order for the algorithm to run. The implication of this is that every router has a complete map of all the routers in the internet!

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In practice this is not the case and other variations of the algorithm allow each router to discover the graph as they go.

7.21. Analysis of Dijkstra's Algorithm

Let's look at the running time of Dijkstra's algorithm. We first note that building the priority queue takes time since we initially add every vertex in the graph to the priority queue. Once the queue is constructed, the `while` loop is executed once for every vertex since vertices are all added at the beginning and only removed after that.

Let's look at the running time of Dijkstra's algorithm. We first note that building the priority queue takes $\Theta(V^2)$ time since we initially add every vertex in the graph to the priority queue. Once the queue is constructed, the `while` loop is executed once for every vertex since vertices are all added at the beginning and only removed after that.

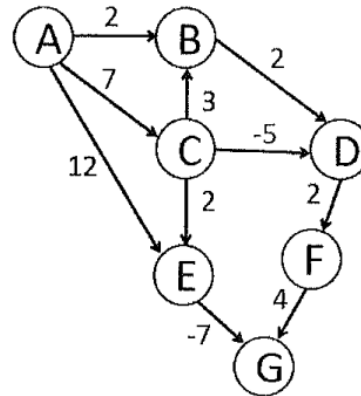
Within that loop each call to `delete()` takes $\Theta(V)$ time. Taken together, that part of the loop and the calls to `delete()` take $\Theta(V^2)$. The `for` loop is executed once for each edge in the graph, and within the for loop the call to `change_priority()` takes $\Theta(V)$ time. So the combined running time is $\Theta(V^2)$.

```
In [42]: display_quiz(quiz_path+"dijk.json", max_width=800)
```

Which of the following is a known limitation or characteristic of Dijkstra's algorithm?

- ☐ It is an iterative algorithm that provides the shortest path from a single source to all other reachable vertices.
- ☐ The time complexity is independent of the number of edges in the graph.
- ☐ It does not require a complete map of the graph to function in a routing environment.
- ☐ It can accurately find shortest paths in graphs with negative edge weights.

Exercise 1: Given a weighted directed graph with negative edges as follows, what is the resulting shortest path from vertex A to D after performing Dijkstra's algorithm if we do check the visited node?



(A) A -> B -> D

(B) A -> C -> D

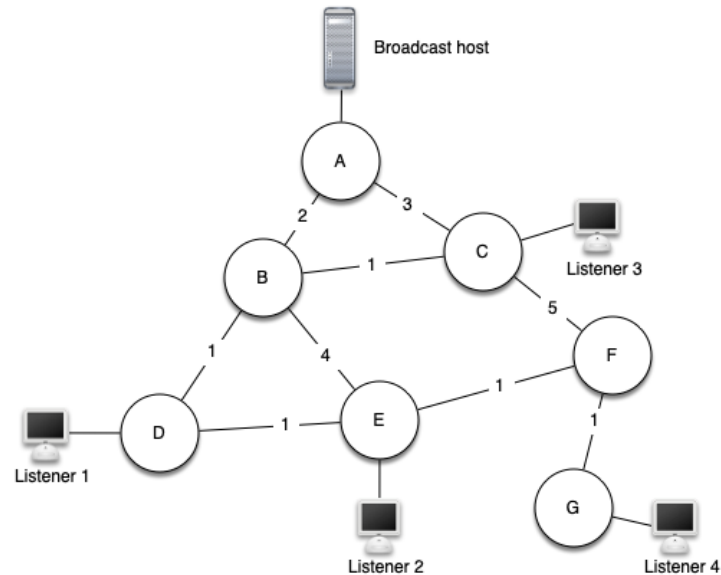
(C) A -> E -> G -> D

(D) A -> C -> E -> G -> D

7.22. Prim's Spanning Tree Algorithm

For our last graph algorithm let's consider a problem that online game designers. The problem is that they want to efficiently transfer a piece of information to anyone and everyone who may be listening. This is important in gaming so that all the players know the very latest position of every other player!

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In the above figure we show a small network with a broadcaster and some listeners. Using this first approach, four copies of every message would be sent. Assuming that the **least cost path** is used, let's see how many times each router would handle the same message.

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In the above figure we show a small network with a broadcaster and some listeners. Using this first approach, four copies of every message would be sent. Assuming that the **least cost path** is used, let's see how many times each router would handle the same message.

All messages from the broadcaster go through router A, so A sees all four copies of every message. Router C sees only one copy of each message for its listener. However, routers B and D would see three copies of every message since routers B and D are on the **cheapest path** for listeners 1, 2, and 4! When you consider that the broadcast host must send hundreds of messages each second for a radio broadcast, that is a lot of extra traffic.

Another brute force solution is for the broadcast host to send a single copy of the broadcast message and let the routers sort things out. In this case, the easiest solution is a strategy called uncontrolled flooding:

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Each message starts with a time to live (TTL) value set to some number greater than or equal to the number of edges between the broadcast host and its most distant listener. Each router gets a copy of the message and passes the message on to all of its neighboring routers. When the message is passed on the TTL is decreased. Because each router continues to send copies of the message to all its neighbors until the TTL value reaches 0, it is easy to convince yourself that uncontrolled flooding generates many more unnecessary messages than our first strategy.

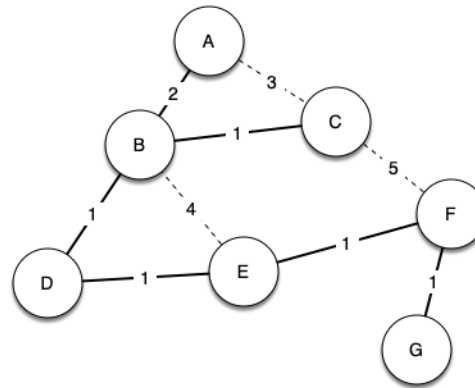
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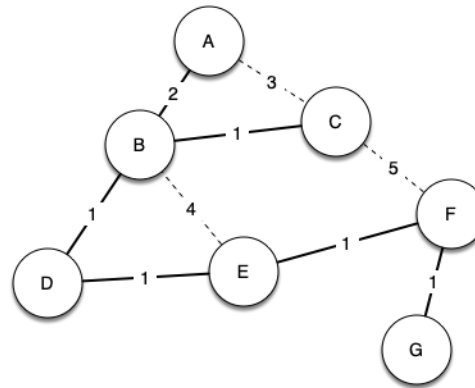
The solution to this problem lies in the construction of a minimum weight spanning tree. Formally we define the minimum spanning tree for a graph as follows. is an acyclic subset of that connects all the vertices in . The sum of the weights of the edges in is minimized.

Below shows a simplified version of the broadcast graph and highlights the edges that form a minimum spanning tree for the graph.

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In this example A forwards the message to B. B forwards the message to D and C. D forwards the message to E, which forwards it to F, which forwards it to G. No router sees more than one copy of any message, and all the listeners that are interested see a copy of the message.

The algorithm we will use to solve this problem is called Prim's algorithm. Prim's algorithm belongs to a family of algorithms called the greedy algorithms that applies to **undirected graph** because at each step it will choose the cheapest next step. In this case the cheapest next step is to follow the edge with the lowest weight.

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```
While T is not yet a spanning tree
  Find an edge that is safe to add to the tree
  Add the new edge to T
```

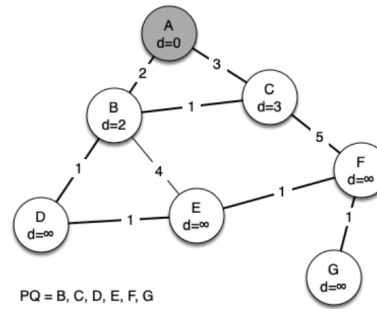
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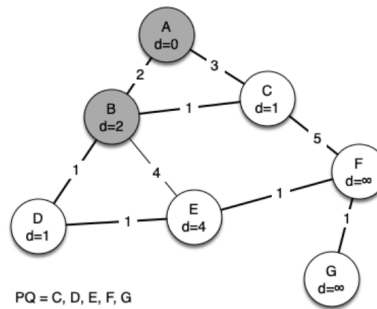
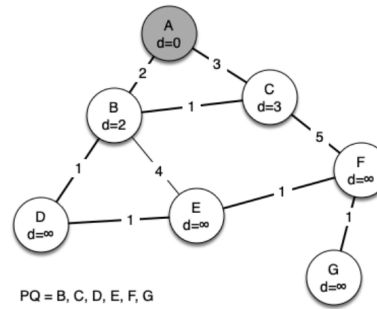
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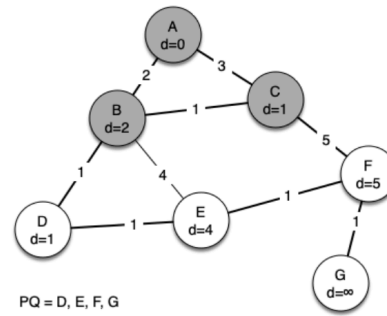
The trick is in the step that directs us to "find an edge that is safe." We define a safe edge as any edge that connects a vertex that is in the spanning tree to a vertex that is not in the spanning tree. This ensures that the tree will always remain a tree and therefore have no cycles.

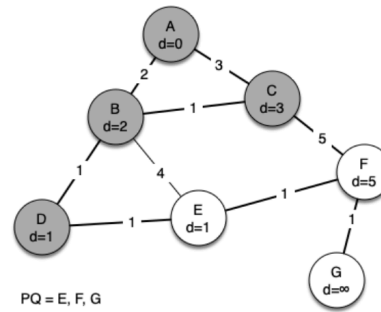
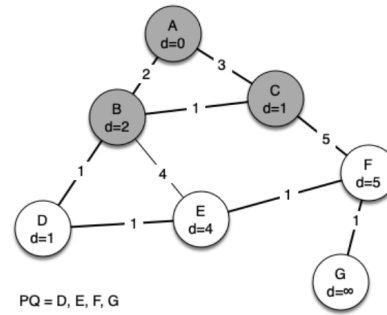

```
In [43]: import sys
from pythonds3.trees.priority_queue import PriorityQueue #heapq

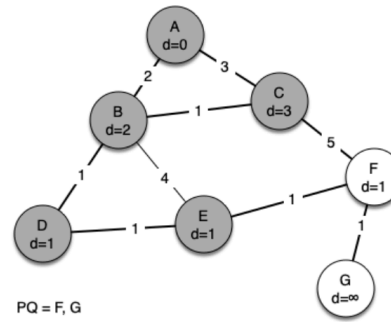
def prim(graph, start):
    pq = PriorityQueue()
    for vertex in graph:
        vertex.distance = sys.maxsize
        vertex.previous = None
    start.distance = 0
    pq.heapify([(vertex.distance, vertex) for vertex in graph])
    while not pq.is_empty():
        print(pq)
        distance, current_v = pq.delete()
        for next_v in current_v.get_neighbors():
            new_distance = current_v.get_neighbor(next_v)
            if next_v in pq and new_distance < next_v.distance:
                next_v.previous = current_v
                next_v.distance = new_distance
                pq.change_priority(next_v, new_distance)
```

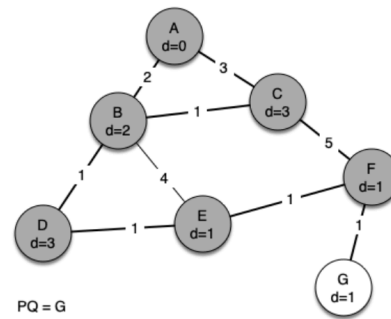
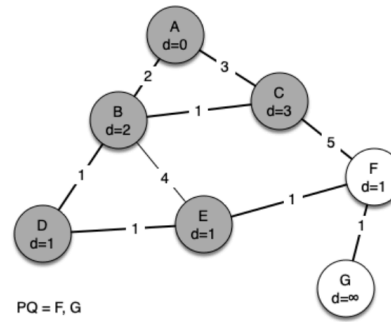


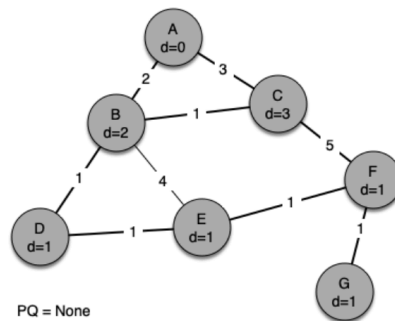












PQ = None

In [45]:

```
g = Graph()
g.add_edge('A', 'B', 2)
g.add_edge('B', 'A', 2)
g.add_edge('A', 'C', 3)
g.add_edge('C', 'A', 3)
g.add_edge('B', 'D', 1)
g.add_edge('D', 'B', 1)
g.add_edge('B', 'C', 1)
g.add_edge('C', 'B', 1)
g.add_edge('B', 'E', 4)
g.add_edge('E', 'B', 4)
g.add_edge('D', 'E', 1)
g.add_edge('E', 'D', 1)
g.add_edge('C', 'F', 5)
g.add_edge('F', 'C', 5)
g.add_edge('E', 'F', 1)
g.add_edge('F', 'E', 1)
g.add_edge('F', 'G', 1)
g.add_edge('G', 'F', 1)
```

```
In [46]: # Example of how to use the find_path function
def print_spanning_tree(graph, start_vertex):
    """Prints the edges in the minimum spanning tree."""
    print("Edges in the Minimum Spanning Tree:")
    for vertex in graph:
        if vertex.previous is not None:
            print(f"({vertex.previous.key}, {vertex.key}) \
                with weight {vertex.get_neighbor(vertex.previous)}")

prim(g, g.get_vertex('A')) # Run prim from 'A'
print_spanning_tree(g, g.get_vertex('A'))
```

[(0, Vertex(A)), (9223372036854775807, Vertex(B)), (9223372036854775807, Vertex(C)), (9223372036854775807, Vertex(D)), (9223372036854775807, Vertex(E)), (9223372036854775807, Vertex(F)), (9223372036854775807, Vertex(G))]

[(2, Vertex(B)), (9223372036854775807, Vertex(D)), (3, Vertex(C)), (9223372036854775807, Vertex(G)), (9223372036854775807, Vertex(E)), (9223372036854775807, Vertex(F))]

[(1, Vertex(C)), (1, Vertex(D)), (9223372036854775807, Vertex(F)), (9223372036854775807, Vertex(G)), (4, Vertex(E))]

[(1, Vertex(D)), (4, Vertex(E)), (5, Vertex(F)), (9223372036854775807, Vertex(G))]

[(1, Vertex(E)), (9223372036854775807, Vertex(G)), (5, Vertex(F))]

[(1, Vertex(F)), (9223372036854775807, Vertex(G))]

[(1, Vertex(G))]

Edges in the Minimum Spanning Tree:

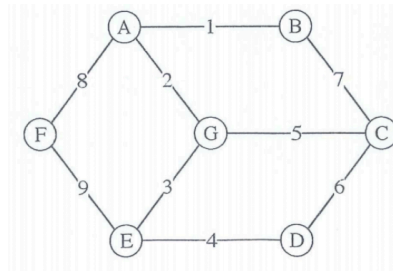
(A, B)	with weight 2
(B, C)	with weight 1
(B, D)	with weight 1
(D, E)	with weight 1
(E, F)	with weight 1
(F, G)	with weight 1

In [47]: `display_quiz(quiz_path+"prim.json", max_width=800)`

In Prim's algorithm for finding a Minimum Spanning Tree (MST), what defines a safe edge?

- ☐ Any edge that does not create a cycle when added to the current tree.
- ☐ An edge that has a weight lower than the average of all edges in the graph.
- ☐ The first edge encountered during a depth-first search of the graph.
- ☐ The edge with the minimum weight that connects a vertex in the tree to a vertex outside the tree.

Exercise 2: Considering the following graph, when Prim's algorithm is used and starts from node F, what is the weight of the last edge to be added into the minimum spanning tree?



```
In [48]: from jupytercards import display_flashcards  
fpath= "https://raw.githubusercontent.com/phonchi/nsysu-math208/refs/heads/main/flashcards/"  
#fpath= "flashcards/"  
display_flashcards(fpath + 'ch8.json')
```

Graph



Next

>

References

1. Textbook CH7

