# Problem Set 5

This problem set is divided into two parts: Part A problems are theory questions, and Part B problems are programming tasks.

Part A questions are due Tuesday, November 16th at 11:59PM.

#### Part B questions are due Friday, November 19th at 11:59PM.

Solutions should be turned in through the course website. Your solution to Part A should be in PDF form using LATEX or scanned handwritten solutions. Your solution to Part B should be a valid Python file, together with one PDF file containing your solutions to the two theoretical questions in Part B.

Templates for writing up solutions in LATEX are available on the course website.

Remember, your goal is to communicate. Full credit will be given only to the correct solution which is described clearly. Convoluted and obtuse descriptions might receive low marks, even when they are correct. Also, aim for concise solutions, as it will save you time spent on write-ups, and also help you conceptualize the key idea of the problem.

# Part A: Due Tuesday, November 16th

## 1. (15 points) No Odd Cycles

Given an undirected graph G = (V, E), design an algorithm that decides whether or not G contains a cycle of odd length. Prove its correctness. Your algorithm should run in time O(|V| + |E|).

## 2. (15 points) Maximum Bandwidth Path

Suppose we have a telephone network N with n nodes, m links (undirected edges) between them, and a distinguished source node s. Each link has an associated *bandwidth* which is a non-negative integer that indicates the maximum number of requests that link can handle. In a path of links, the link in the path with the smallest bandwidth constrains the number of requests that can be passed along the path. Hence, we define the bandwidth of a path Pconsisting of links  $\ell_1, \ldots, \ell_k$  to be min(bandwidth( $\ell_1$ ), ..., bandwidth( $\ell_k$ )). The maximum bandwidth path to a node v is the path from s to v with the maximum bandwidth.

Design an algorithm that computes the maximum bandwidth path to every node v in the network, and prove its correctness. Your algorithm should have the same asymptotic running time as Dijkstra's.

## 3. (20 points) Shortest Roundtrip Path

Suppose G is a strongly connected directed graph, meaning that for every two vertices u and v in G, there is a directed path from u to v and a directed path from v to u. Each edge has

a weight, not necessarily non-negative. Let s be a distinguished node in G. For every vertex v in the graph, we want to calculate a path from s to v and a path from v to s such that the sum of the weights of edges on the two paths is minimized. Give an efficient algorithm to perform this task, and show correctness. If your algorithm encounters a negative-weight cycle, it should output false. Otherwise, it should output the pair of paths from s to v and from v to s that minimizes the roundtrip cost.

#### Part B: Due Friday, November 19th

1. (50 points) Speeding up Dijkstra.

The Howe & Ser Moving Company is transporting the Caltech Cannon from Caltech's campus to MIT's and wants to do so most efficiently. Fortunately, you have at your disposal the National Highway Planning Network (NHPN), packaged for you in ps5\_dijkstra.zip. You can learn more about the NHPN at

http://www.fhwa.dot.gov/planning/nhpn/

This data includes node and link text files from the NHPN. Open nhpn.nod and nhpn.lnk in a text editor to get a sense of how the data is stored (datadict.txt has a more precise description of the data fields and their meanings). To save you the trouble of parsing these structures from a file, we have provided you with a Python module nhpn.py containing code to load the text files into Node and Link objects. Read nhpn.py to understand the format of the Node and Link objects you will be given.

Your goal in this problem is to implement and test several techniques for speeding-up Dijkstra's algorithm in order to compute shortest paths between various pairs of locations.

Implementation of Dijkstra's algorithm is already provided. Function

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dijkstra(nodes, edges, weight, source)
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is given a graph with non-negative edges (represented as a list of Node objects and a list of undirected Edge objects), a function weight (node1, node2) that returns the weight of any edge between node1 and node2, and a source node source. This function updates the node.visited field for all nodes, which indicates whether a shortest path to node has been found, as well as node.distance and node.parent for visited nodes, which are the length of the shortest path from the source node to node and the previous node on that path respectively. The function returns the number of nodes visited during the execution of the algorithm.

The links you are given do not include weights, so instead we use the geographical distance between their endpoints. Function distance(node1, node2) returns the distance between two NHPN nodes. Nodes come with latitude and longitude (in millionths of degrees). For simplicity, we treat these as (x, y) coordinates on a flat surface, where the distance between two points can be calculated using the Pythagorean Theorem.

Dijkstra's algorithm uses a priority queue, but this priority queue has one subtle requirement. Dijkstra's algorithm calls decrease\_key, but decrease\_key requires the index of an item in the heap, and Dijkstra's algorithm would have no way of knowing the current index corresponding to a particular Node. To solve this problem, the course staff have written an augmented heap object, heap\_id, with the following extra features:

- insert (key) returns a unique ID.
- decrease\_key\_using\_id(ID, key) takes an ID instead of an index.
- extract\_min\_with\_id() extracts the minimum element and returns a pair (key, ID).

Additionally, we have provided some tools to help you visualize the output from your algorithms. You can use the Visualizer class to produce a KML (Google Earth) file. To view such a file on Google Maps, place it in a web-accessible location, such as your Athena Public directory, and then search for its URL on Google Maps.

For this problem, you will modify the file dijkstra.py. As you solve each part of the problem, check your work by running the appropriate test functions. We have provided several test functions that test each part separately or perform comparison tests of several methods. You should follow the instructions for each part of the problem, perform appropriate tests and draw conclusions. Please submit the modified dijkstra.py file with the code and dijkstra.pdf file with proofs and short answers. Keep them short.

- (a) (3 points) Examine the code provided in nhpn.py, heap.py and dijkstra.py to learn the structure of the Node and Link classes and the implementation of Dijkstra's algorithm. Run test\_a(). Is there a significant difference in the execution time for different pairs of nodes? Explain your observation.
- (b) (7 points) One way to speed up Dijkstra's algorithm is to terminate the algorithm early once a shortest path to the destination has been found.Implement function

dijkstra\_early\_stop(nodes, edges, weight, source, dest)

that performs this optimization. As with the function dijsktra(), this function should update the node.visited, node.distance and node.parent fields, and return the number of nodes visited during its execution. Run test\_b(). What characterizes pairs of nodes for which there is a significant speed-up using this optimized version of Dijkstra's algorithm?

HINT: Reuse the implementation of Dijkstra's algorithm provided, making the required changes to allow for early termination.

(c) (20 points) We will apply the potentials method with a landmark node to obtain a faster shortest path algorithm. For a given landmark node l, we denote the potential of a node u with respect to a destination node t by  $\lambda_t^l(u)$ . The potential is defined as  $\lambda_t^l(u) = \delta(u, l) - \delta(t, l)$  if there exists a path from u to t through l, and  $\lambda_t^l(u) = C$ 

where C is some fixed constant if no such path exists. (Here,  $\delta(u, v)$  denotes the length of a shortest path from u to v).

- i. Prove that this potential function is feasible, i.e. the modified weight of every edge is non-negative.
- ii. Implement function

compute\_landmark\_distances (nodes, edges, weight, landmark) that computes shortest paths from all nodes to the given landmark node landmark. For each node node, node.land\_distance should be set to the value of the shortest path distance from node to landmark or to some constant C if no such path exists (e.g.,  $C = 10^9$ ). Why is it more useful to precompute distances to the landmark node than precomputing the potentials themselves?

iii. Implement function

dijkstra\_with\_potentials(nodes, edges, weight, source, dest) that performs Dijkstra's algorithm using edge weights modified according to the potentials method (i.e.  $w'(u, v) = w(u, v) - \lambda_t^l(u) + \lambda_t^l(v)$  for a landmark l and destination t) and terminates as soon as a shortest path to the destination node dest has been found. This function assumes that node.land\_distance is already set to the proper value (no need to call compute\_landmark\_distances() from it). As before, this function should update node.visited, node.distance and node.parent fields, and return the number of nodes visited during its execution. Run test\_d(). In which scenarios is the speed-up most significant (compare to both Dijkstra's algorithm and Dijkstra's algorithm with early termination)?

Hint: Reuse the implementation of Dijkstra's algorithm provided, making the necessary changes.

- (d) (20 points) We will now describe a potentials method where multiple landmarks are used. For a given set of landmarks L, the potential of a node u with respect to a destination node t is  $\lambda_t^L(u) = \max_{l \in L} \lambda_t^l(u)$ , where  $\lambda_t^l(u)$  is defined as before.
  - i. Prove that this potential function is also valid, i.e. the modified weight of every edge is non-negative.
  - ii. How does the potential function  $\lambda_t^L(u)$  compare to a potential function  $\lambda_t^l(u)$  for any single landmark vertex  $l \in L$  in terms of the number of visited nodes when used in Dijkstra's algorithm with early termination? Explain which is better and why.
  - iii. Implement function

that computes shortest paths from all nodes to all given landmark nodes in landmarks. For each node node, node.land\_distances should be set to a list of values, such that node.land\_distances[i] is the shortest path length from node to landmarks[i] or some constant C if no such path exists (e.g.,  $C = 10^9$ ).

iv. Implement function

that performs Dijkstra's algorithm using edge weights modified according to the potentials method with multiple landmarks (i.e.  $w'(u,v) = w(u,v) - \lambda_t^L(u) + \lambda_t^L(v)$  for a set of landmarks L and destination t), and terminates as soon as a shortest path to the destination node dest is found. This function assumes that node.land\_distances is already set to the proper list of values. As before, this function should update node.visited, node.distance and node.parent fields, and return the number of nodes visited during its execution. Run test\_f(). Does the performance of your algorithm match your assertions from part ii.?

(e) (Optional) Included in nhpn.py is a method to convert a list of nodes to a .kml file.
.kml files can be viewed using Google Maps, by putting the file in a web-accessible location (like your Athena Public directory), going to

http://maps.google.com and putting the URL in the search box.

Run visualize\_path.py. This will create two files, path\_flat.kml and path\_curved.kml. Both should be paths from Pasadena CA to Cambridge MA. path\_flat.kml was created using the distance function you wrote in part (b), and path\_curved.kml was created using a distance function that does not assume that the Earth is flat. Can you explain the differences? Also, try asking Google Maps for driving directions from Caltech to MIT to get a sense of how similar their answer is.