Shortest Path Trees and Reach in Road Networks

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We provide the code along with several comments and implementation notes in the src/ folder.

Special implementations notes

For all the questions, our implementation gives the correct answer for multiple shortest paths, aka when there are several paths of optimal length between two points.

When we output point coordinates between two vertices, we make a mean of their coordinates according to the distance between them (we suppose the edge is a straight line and keep the same ratio).

First part

For (1.1) and (1.2), using Dijkstra is obvious:

- It is very efficient for shortest paths with single source (fastest algorithm we have heard of if implemented with Fibonacci heaps)
- Because of the min-heap structure used, and the scanning of the vertices in the increasing order of distance, it is very simple to stop the execution as soon as we go further than the target distance, thus providing a faster algorithm.

For (1.3), we can extend the algorithm by retaining the vertices at distance t_1 and propagating them until we find a vertex at distance t_2 , or stop as soon as we go further.

If we consider multiple shortest paths, it leads to an inefficient memory usage because we must propagate sets.

The elegant method was to first compute shortest paths using Dijkstra, then select the vertices at distance greater than t_2 and backtrack until we are at distance less than t_1 .

Another advantage of this method is that it allows for

$$t_2 < t_1$$

And thus one can solve the original problem by setting

$$t_2 = 0$$

Optimisation tricks

Dijkstra stop

If our goal is the points at distance t_2 , we can stop the Dijkstra for the vertices at distance more than

$$t_2 + \max_{e \in E} e.length$$

but we must be careful, if we allow $t_2 < t_1$, the condition becomes

$$\max(t_1, t_2) + \max_{e \in E} e.length$$

Dijkstra memoisation

Since Dijkstra represent a huge part of our computations, and it is basically the same for all functions when we set one starting vertex in a graph, we can compute the Dijkstra over the whole graph (about the same running time as reading the input), and use it for other computations.

This part of the code has been replaced with the Online Dijkstra optimisation.

Distance sorting

Because of memoisation, if we use small values of t_1 and t_2 (or close ones), we actually lose time because we must test each vertex of the Dijkstra to ensure it is at distance at least t_2 , even if in the end we will never use vertices whose distance to v is not between approximately between t_1 and t_2 .

The solution is to return, along with the Dijkstra output, a list of all computed vertices sorted by their distance.

Thus with binary search, we can find the first and last starting vertices that are necessary for the backtrack, and iterate through the list.

This is a major optimisation for the part B for example, because we have to answer numerous requests like (1.2).

Online Dijkstra

This optimisation is by far the most interesting, because a similar idea can be used for the B part.

The problem we have is that two major optimizations are incompatible: the Dijkstra stop and the Dijkstra memoisation. If we stop the Dijkstra and then use it with a greater stopping value, then it will be incorrect. Conversely, to memoise the Dijkstra, we have to compute it over the whole graph, which is really stupid for the country-sized one for example.

The answer to this is simple: return a value while not throwing away the current computation step.

We implemented this in a very elegant fashion using Python's generators. We can yield the current result from the generator, and conversely we can send the new parameter value to the generator which is a less known feature of generators.

Option B

We note dist the distance function.

(B.1)

Suppose there is v with r(v) > D/2. Then there are s and t such that a fastest path $s \rightarrow t$ visits v, and dist(s, v) >= r(v) and dist(v, t) >= r(v). Then dist(s, v) > D which is a contradiction.

(B.2)

Suppose there exists a point $s \in S_{in}$ and a point $t \in T_{out}$, such that some quickest path from s to t passes through v. Let us show that $\mathtt{dist}(s, v) = 1$ hour and $\mathtt{dist}(v, t) = 1$ hour. By symmetry of the definitions, we only need to show it for $\mathtt{dist}(s, v)$. There exist a shortest path from s to a vertex such that v is 1 hour away of s. But any subpath of a shortest path is a shortest path (or else the main path is not the shortest). Thus $\mathtt{dist}(s, v) = 1$ hour and $\mathtt{dist}(v, t) = 1$ hour and $\mathtt{r}(v) >= 1$ hour.

(B.3)

If the reach of v is at least 2 hours, there are s and t such that a fastest path s \rightarrow t visits v, and dist(s, v) >= r(v) >= 2 hours and dist(v,t) >= r(v) >= 2 hours. Choose $t' \in T_{out}$ on the shortest path v \rightarrow textracted from s \rightarrow t and $s' \in S_{in}$ on the shortest path s \rightarrow v extracted from s \rightarrow t. But any subpath of a shortest path is a shortest path (or else the main path is not the shortest). Thus the path s' \rightarrow t' extracted from s \rightarrow t is a shortest path that contains v.

(B.4)

If we take the contrapositions of those propositions, if there are $s \in S_{in}$ and $t \in T_{out}$ such that dist(s, t) = dist(s, v) + dist(v, t), then r(v) is more than 1 hour. If those s and t do not exist, then r(v) is less than 2 hours.

We can model this with an oracle O(r) such that : if O(r) then r(v) >= r else $r(v) < 2 \times r$, where r = 1 hour in the previous example.

Conversely, the oracle has a defined behaviour only outside the interval [r(v)/2, r(v)] (left gives True and right gives False).

1. We find a factor 4 approximation

If we know a value r such that O(r/2) = True and O(r) = False, then we know that x is in $[r/2, 2 \times r[$.

And for any starting point y, while O(y) = True we multiply y by 2 until we have O(y) = False, and conversely for O(y) = False to find such a value.

By the properties of the oracle, this terminates. We have a logarithmic complexity. But when evaluating $O(y \times 2^k)$ for an integer k, we know that there is an index that separates the True values from the False because of the properties of the oracle. Thus we can apply binary search on this logarithmic scale to have a $\log(\log(r(v)))$ number of evaluations of the oracle.

2. We transform a $2^{1+\frac{1}{k}}$ approximation into a $2^{1+\frac{1}{2k}}$ approximation

Suppose you know $r(v) \in [a, b[$ with $\frac{b}{a} = 2^{1 + \frac{1}{k}}$.

Take $c = a \times 2^{\frac{1}{2k}}$.

We know that if $r(v) \in [a, c[$, then O(c) is false, and that if $r(v) \in [2c, b[$, then O(c) is true.

Thus we can eliminate one interval of ratio $2^{\frac{1}{2k}}$, and the remaining interval is of ratio $2^{1+\frac{1}{k}-\frac{1}{2k}}=2^{1+\frac{1}{2k}}$.

3. Complexity

We make $\log(\log(r(v)))$ calls to the oracle in the first stage, and then i to get a $2^{1+\frac{1}{2^i}}$ approximation factor.

Since we work with integers, the stopping condition is $r(v) \times (2^{1+\frac{1}{2^i}} - 2) < 1$. A Taylor expansion gives us $i > \log(r(v)) + 1$ with the first term, so $i > \log(r(v)) + 2$ should be sufficient.

4. Other strategies

Like shown below, more evaluations per step could get us immediately better approximation factors than 2.

However, it requires too much evaluations, with no improvement in the worst case, that is when the oracle is a threshold function.

Let $x_1, x_2 = \frac{r}{2}, x_3, x_4 = b, x_5, x_6 = 2r$ be numbers in geometric progression of ratio $\sqrt{2}$ with r found after the first stage.

We know $O(x_2) = T$, $O(x_4) = F$, $O(x_6) = F$, and we know $x \in [x_2, x_6]$. We split this interval in 4 and look at the values of $O(x_1)$, $O(x_3)$, $O(x_5)$. We note? when a value can be True or False.

For $r(v) \in [x_2, x_3[$, the possible values are ?, F, F.

For $r(v) \in [x_3, x_4]$, the possible values are T, ?, F.

For $r(v) \in [x_4, x_5[$, the possible values are T, ?, F.

For $r(v) \in [x_5, x_6]$, the possible values are T, T, ?.

If we get $O(x_1) = F$ or $O(x_5) = T$, we know a $\sqrt{2}$ factor approximation of r(v). Else, the value of $x_3 = c$ allows us to eliminate either the leftmost or the rightmost interval, which is exactly the first step of the second stage of our general strategy to transform a $4 = 2^{1+\frac{1}{4}}$ approximation into a $2\sqrt{2} = 2^{1+\frac{1}{2}}$ approximation.

5. Computation time of the oracle

Let's determine the computation time of the oracle :

We can memoize two Dijkstra results starting from \mathbf{v} and following the normal edges and the reversed edges. Thus we can compute quickly the two sets S_{in} and T_{out} .

As we saw in the examples, the size of S_{in} is not huge. Thus for each point of S_{in} we compute a Dijkstra, stopping when we know the distances to all points of T_{out} . Finally we test dist(s, t) = dist(s, v) + dist(v, t) for each (s, t)

t) pair (dist(s, v) and dist(v, t) are known from the memoized Dijkstra results).

6. Practical implementation and optimisations

As explained in reach.py:

A dichotomy for the 4-factor approximation reduces the number of calls to the oracle to $O(\log(\log(r)))$ for this stage.

But it would also call it with bigger inputs and more inputs that evaluate to False thus taking a bit more time.

If we are not lucky, it could call oracle with a huge value, so it is not a good idea in practice because oracle is optimised for small values.

Thus we did not implement the optimal 4-factor approximation.

For the Dijkstra between S_{in} and T_{out} , we used the same method of online algorithm with a specific function.

It leads to significant improvement because during the final steps of the algorithm, there are a lot of elements in common between the S_{in} sets.

We also clean the memory of those small Dijkstra at the end of each reach computation (since they are only used by the oracle).

(B.5)

We could notice a huge variance between the computation time of the **reach** function for different vertices, positively correlated (but not proportional) with the answer: for a bigger answer, we made more evaluations of the oracle, and over bigger subgraphs.

Our program compute_reach.py can be launched from the command line. The bonus is that multiple instances can be running at the same time, while updating the same file of results and not wasting resources on the same starting vertex.

We also provide a small launch.sh bash script that launches multiple instances of compute_reach.py in separate screen sessions and also logs the running time.

(B.6)

We used launch.sh to compute the reach of $2^{14} = 16384$ vertices of the france.in graph.

The results are available in CSV format in the reach.log file.

We computed two histograms, one as requested by the assignment, and one using a logarithmic scale with a constant bin width chosen by the Freedman–Diaconis rule¹.

Performances

Those are performances for france.in.

As we use Python 3, one could expect low performances, but they are quite good thanks to all the optimisations.

We used PyPy3² to speed up the computations.

The mean computation time of reach on a MacBook Pro Unibody 2012 is about 20-30 seconds per vertex, but many last less that 1 second, while some can last over 10 minutes.

The drawback from the memoisation is the memory usage (and Python is particularly not efficient unless you use array.array). After the graph is loaded, Python uses 2.5GB of memory, and after the converse graph is generated, it uses almost 4GB. If we delete the coordinates, we save about 200MB. But when it computes a huge value of reach, the memory usage can go up to 10GB (which is cleaned when we change of vertex). We could have saved some memory by cleaning more the Dijkstra with targets (a simple heuristic would clean the instances that have not been used by the last k calls to the oracle, where k depends on the mean edge length on the graph and the current interval in the reach function).

 $^{^1 \}rm https://en.wikipedia.org/wiki/Freedman\%E2\%80\%93Diaconis_rule$

²http://pypy.org/

Figures

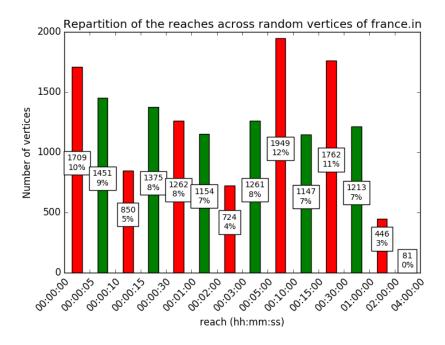


Figure 1:

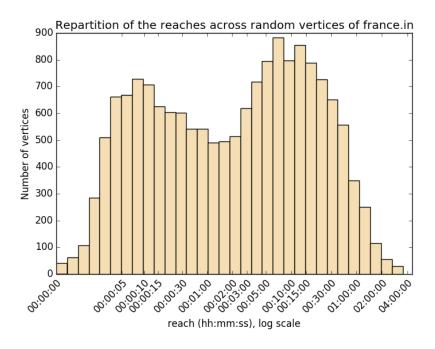


Figure 2: