

MAXimal

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Algorithm for finding the shortest paths in Leviticus from a given vertex to all other vertices

Suppose we are given a graph with N vertices and M edges, each of which indicated its weight L_i . Also, given the starting vertex V_0 . Required to find the shortest path from vertex V_0 to all other vertices.

Leviticus algorithm solves this problem very efficiently (about the asymptotic behavior and the speed of the cm. Below).

Description

Let array $D[1..N]$ will contain the current shortest path lengths, ie D_i - this is the current length of the shortest path from vertex V_0 to vertex i . Initially D array is filled with values "infinity", except $D_{V_0} = 0$. At the end of the algorithm, this array will contain the final shortest distance.

Let array $P[1..N]$ contains the current ancestors, ie P_i - is the peak preceding the vertex i in the shortest path from vertex V_0 to i . As well as an array D , array P is changed gradually during the algorithm and at its end receives the final values.

Now actually the algorithm Leviticus. At each step, supported by three sets of vertices:

- M_0 - vertex distance which has already been calculated (but perhaps not entirely);
- M_1 - vertex distance are calculated;
- M_2 - vertex distance is yet to be calculated.

Vertices in the set M_1 is stored in the form of bi-directional queue (deque).

Initially all nodes are placed in a plurality of M_2 , apart from the vertex V_0 , which is placed in a plurality of M_1 .

At each step of the algorithm, we take the top of the set M_1 (We reach the top element of the queue). Let V - is the selected vertex. Translate this vertex in the set M_0 . Then review all the edges emanating from this vertex. Let T - this is the second end of this rib (i.e., not equal to V), and L - the length of this edge.

- If T belongs to M_2 , then T is transferred to a set of M_1 to the end of the queue. D_T is set equal to $D_V + L$.
- If T belongs to M_1 , then try to improve the value of D_T : $D_T = \min(D_T, D_V + L)$. The very top of T is never moved in the queue.
- If T belongs to M_0 , and if D_T can be improved ($D_T > D_V + L$), then improve D_T , and T return to the top of the set M_1 , placing it in the top of the queue.

Of course, every time you update the array D should be updated and the value in the array P .

Implementation Details

Create an array $ID[1..N]$, in which each vertex will be stored, which set it belongs: 0 - if M_2 (ie, the distance is infinite), 1 - if M_1 (ie, the vertex is queue) and 2 - when M_0 (a path has been found, the distance is less than infinity).

Queue processing can be realized by a standard data structure deque. However, there is a more efficient way. Firstly, it is obvious in the queue at any one time will be stored a maximum of N elements. But secondly, we can add elements and beginning and end of the queue. Therefore, we can arrange a place on the array size N , but you have to loop it. Ie make an array $Q[1..N]$, pointers (int) to the first element QH and the element after the last QT . The queue is empty when $QH == QT$. Appending - a record in the $Q[QT]$ and increase QT 1; if QT then went beyond the line ($QT == N$), then do $QT = 0$. Adding the queue - reduce the QH -1, if it has moved beyond the stage of ($QH == -1$), then do $QH = N - 1$.

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Implement the algorithm itself is exactly the description above.

Asymptotics

I do not know more or less good asymptotic estimate of this algorithm. I have seen only the estimate $O(NM)$ of the similar algorithm.

However, in practice, the algorithm has proven itself very well: while it is running I rate as **$O(M \log N)$** , although, again, this is only an **experimental** evaluation.

Implementation

```
typedef pair <int, int> rib;
typedef vector <vector <rib>> graph;

const int inf = 1000 * 1000 * 1000;

int main ()
{
    int n, v1, v2;
    graph g (n);

    Graph reading ... ..

    vector <int> d (n, inf);
    d [v1] = 0;
    vector <int> id (n);
    deque <int> q;
    q.push_back (v1);
    vector <int> p (n, -1);

    while (! q.empty ())
    {
        int v = q.front (), q.pop_front ();
        id [v] = 1;
        for (size_t i = 0; i < g [v] .size (); ++ i)
        {
            int to = g [v] [i] .first, len = g [v] [i] .second;
            if (d [to] > d [v] + len)
            {
                d [to] = d [v] + len;
                if (id [to] == 0)
                    q.push_back (to);
                else if (id [to] == 1)
                    q.push_front (to);
                p [to] = v;
                id [to] = 1;
            }
        }
    }

    Conclusion ... the result ...

}
```

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according to the description in the inner cycle there must be "else if (id[to] == 2) q.push_front(to);" instead of "else if (id[to] == 1) q.push_front(to);" (vertex "to" is in M_0 - set). Besides, where did you set vertex V as M_0 ? Some misunderstanding of mine here, i guess...

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**Tornike** • 2 года назад

is it working for negative edges ?

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Yes, it is, and that's the main purpose of this algorithm - because standard algorithms like Dijkstra or breadth-first search don't work with negative-cost edges.

On the other hand, you should take into account that this is a heuristical algorithm - it's even non-polynomial, though it works very good in practise (especially on random graphs).

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**RiaD** ➔ e_maxx • 10 месяцев назадI'd add that it works about 2^n on some case right in Асимптотика section.

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А тут точно всё правильно с номерами множеств (id[n])? А то 2 ни разу не встречается, да и вершины из M_1 не выходят.

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**Jarekczek** • год назад

I can't leave a direct link, cause the exact article is hardly available on the net. But I saw in several places, that in pessimistic case this algorithm (Pape-Levit's, correct?) performs in exponential (2^n) time. Although in most cases it showed to be faster than Dijkstra.

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