Steiner tree problem

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1 Problem description

We are going to study the Steiner tree problem, which is defined as follows: Given a graph G=(V,E), a weight function $w:e\mapsto\mathbb{R}$ assigning a weight w(e) for every $e\in V$ and a set of terminal nodes $T\subset V$. The goal is to find a subset of edges $M\subset E$ of minimal weight that connects all terminals in G.

This problem is NP-COMPLETE. Here we are going to be presenting and comparing different heuristic approaches to the problem.

2 Definition

2.1 Checking admissibility

Definition 2.1. Since a solution needs to connect all terminal nodes, I will define a solution to be admissible when all terminal nodes $t \in T$ are in the same connected component.

2.2 Gain function

Definition 2.2. My gain function will simply be the sum of all edges plus the diameter of the graph.

The diameter being the maximum eccentricity over all vertices. The eccentricity of a vertex being the distance (shortest path over the weighted edges) from that vertex to the furthest away vertex.

Remark 2.3. Note that an optimal solution will always have its terminal nodes as leaves.

3 Overview

The basic framework I will use is the one seen in class.

- First, I will initialise a solution from a known approximation algorithm.
- Then I will generate new solutions by considering variations of my current solutions.
- Finally I will select a few solutions from which I will iterate, optimizing the gain function.
- I will stop after a fixed amount of iterations.
- λ being the number of variations I am considering.
- μ being the number of solutions I am selecting.

Algorithm 1: GENERIC_FRAMEWORK $(\lambda, \mu, \text{ VARIATION}, \text{ SELECTION}, k, G, T)$

- VARIATION being a function used to consider variations of the current solutions.
- \bullet SELECTION being a function used to select μ solutions from the current solutions.
- k is a simple integer, G our graph and T our terminal nodes.

4 Initialisation

The problem will be initialised by the following 2-apprximation algorithm.

For all pair of terminals, compute the shortest path between all pair of terminal vertices. Then return the minimum spanning tree of this subgraph.

This algorithm is clearly correct as terminal nodes are never disconnected from the connected component.

Remark 4.1. I will admit that it is a 2-approximation as I don't want my report to be 20 pages and all that really matters is that it yields an admissible solution.

5 Variation

Here I will present the different ways of generating a new solution based on already computed solutions that I studied.

5.1 Mutation

I will refer as mutation all variations that modify a solution by only changing it a little bit.

5.1.1 Adding an edge

A straight-forward mutation that could be applied to an admissible solution would be adding an edge. The edge should have one end in the current solution as we have no interest in creating another connected component. The solution will obviously stay admissible as no terminal nodes can be disconnected from that operation.

This operation is quite limited as it doesn't change the solution much. We observe that our algorithm converges rather quickly to a not so good solution. Therefore, we might want to modify the solution more to bypass those potential local optimums.

5.1.2 Adding a path

Another mutation that would be worth considering is adding a whole path to our current solution. In order to do so, we would randomly select two nodes from out current solution then add a shortest path based on the graph G we started with

This mutation is interesting as it changes the solution significantly. Solutions looke better than the previous solution.

5.1.3 Clean the solution

After either of the previous two mutations, it is interesting to remove as many unnecessary edges as possible. An unnecessary edge is an edge that when removed doesn't make the solution unfeasible.

This operation is more time consuming as it would entail checking if the solution is still feasible.

5.1.4 Multiple mutation

Definition 5.1 (Multiple mutation). With a probability p add an edge, with probability 1-p add a path then clean the solution.

5.2 Crossover

I will refer as crossover a variation that merges from two previous solutions.

The crossover mutation I will consider is merging two solutions $S_1 = (V_1, E_1)$ and $S_2 = (V_2, E_2)$ into $S = (V_1 \cup V_2, E_1 \cup E_2)$ then cleaning it as explained above.

5.3 Combining both

Another variation that I will consider is a mix of both a mutation and a crossover. Depending on how many offsprings we request, I will generate a multiple mutation then a crossover then a crossover on a multiple mutation.

This yields good variations of our current set of solutions.

6 Selection

In this section I will be presenting different ways of selecting solutions.

6.1 Elitist selection

The elitist selection will sample the μ best solutions out of the $\lambda + \mu$ solutions.

6.2 Elitist selection on offsprings only

The elitist selection on offsprings only will sample the μ best solutions out of the λ offpsrings.

It could be a problem when $\lambda < \mu$. If this happens, it will fall back to the previous Elitist selection.

6.3 Fitness proportional selection

The fitness proportional selection samples μ solutions with probability proportional to their gains such that the solution closest to the optimum has the highest probability of being chosen.

From my experience, this doesn't work very well as our solutions tend to have a relatively high gain (a few thousands) and aren't very far from each other (a few hundreds). Therefore the probability vector that I compute is close to uniform.

A workaround could be to compute the probability vector based only on the distance to the current best solution so that the probabilities aren't so close to uniform.

6.4 Boltzmann selection

The Boltzmann selection samples μ solutions out of the λ offsprings. It selects an offspring x if it is better than the best previous solutions y. If it is worse, it selects it with probability $e^{\frac{gain(x)-gain(y)}{T}}$. For some value of T.

6.5 Threshold selection

The Threshold selection samples μ solutions out of the λ offsprings. It selects an offspring x if it is better than the best previous solutions y. If it is worse, it selects it with probability gain(x) > gain(y) + T. For some value of T.

7 Comparison

Here I will be comparing the variation and selection strategies.

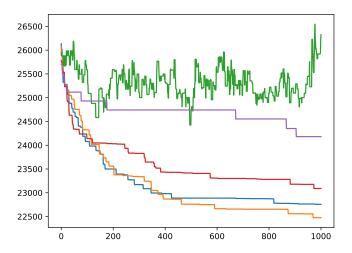


Figure 1: Comparaison for $\lambda=5$, $\mu=2$ and mutation variation of classic elitist selection (in blue), elitist selection on offsprings (in orange), fitness proportional (in green), Boltzmann with constant T=1000 (in red) and Threshold selection with constant parameter T=-150

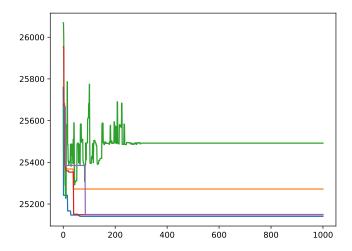


Figure 2: Comparaison for $\lambda=5$, $\mu=2$ and crossover variation of classic elitist selection (in blue), elitist selection on offsprings (in orange), fitness proportional (in green), Boltzmann with constant T=1000 (in red) and Threshold selection with constant parameter T=-150

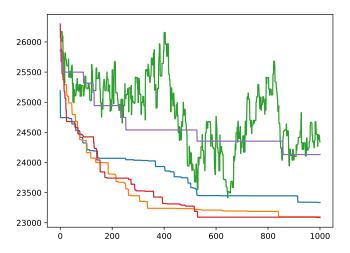


Figure 3: Comparaison for $\lambda=5$, $\mu=2$ and multiple variation of classic elitist selection (in blue), elitist selection on offsprings (in orange), fitness proportional (in green), Boltzmann with constant T=1000 (in red) and Threshold selection with constant parameter T=-150

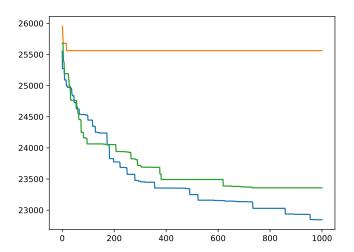


Figure 4: Comparaison for $\lambda=5,\,\mu=2$ and classic elitist selection of mutation variation (in blue), crossover variation (in orange) and another variation consisting of a mix of both (in green).

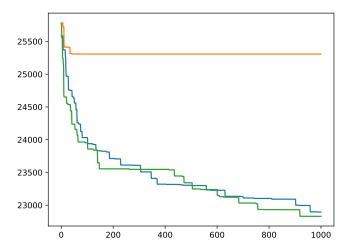


Figure 5: Comparaison for $\lambda = 5$, $\mu = 2$ and offsprings elitist selection of mutation variation (in blue), crossover variation (in orange) and another variation consisting of a mix of both (in green).

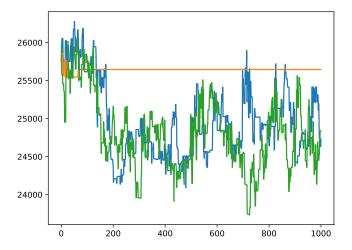


Figure 6: Comparaison for $\lambda=5,~\mu=2$ and Fitness proportional selection of mutation variation (in blue), crossover variation (in orange) and another variation consisting of a mix of both (in green).

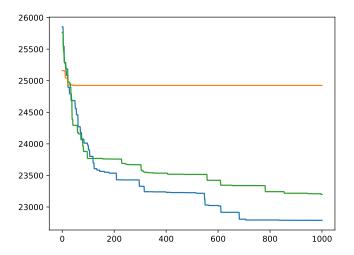


Figure 7: Comparaison for $\lambda = 5$, $\mu = 2$ and Boltzmann selection with constant parameter T = 1000 of mutation variation (in blue), crossover variation (in orange) and another variation consisting of a mix of both (in green).

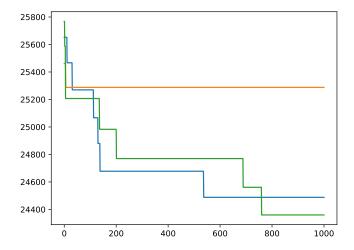


Figure 8: Comparaison for $\lambda=5,\,\mu=2$ and threshold selection with constant parameter T=-150 of mutation variation (in blue), crossover variation (in orange) and another variation consisting of a mix of both (in green).

- 7.1 Variations
- 7.2 Selections
- 8 Conclusion

For this problem,