

Exercises – Algorithms for Genome Rearrangement

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<http://wiki.techfak.uni-bielefeld.de/gi/Teaching/2015summer/gr>

Exercise List 3 — 27.04.2015

Discussion of exercises on: 04.05.2015

Exercise 1

(2 Points)

Given the signed permutation (genome)

$$\pi = [0 \ 2 \ 1 \ -3 \ 4]$$

- Draw the breakpoint graph $BP(\pi)$ and calculate the reversal distance of π to the identity permutation.
- Find a sorting scenario using the overlap graph $O(\pi)$ to find safe reversals, at each step redrawing both the BP and Overlap graphs.

Exercise 2

(2 Points)

Given the signed permutation (genome)

$$\pi = [0 \ -3 \ 1 \ 2 \ 4 \ 6 \ 5 \ 7 \ -15 \ -13 \ -14 \ -12 \ -10 \ -11 \ -9 \ 8 \ 16]$$

- Draw the breakpoint graph $BP(\pi)$. You may use the software from the Wiki if you already did the Exercise 1 by hand, and you are feeling a little lazy... :-)
- What is the lower bound on the reversal distance of π to the identity? Is this bound tight? Why?

Exercise 3

(3 Points)

Consider the special case of Sorting By Reversals where only reversals of length two are allowed (swap of consecutive elements), called SB2R.

- Give an algorithm for SB2R of an *unsigned* permutation. Can you show if this algorithm is optimal, that is, that it sorts the permutation in the minimum possible number of swaps?
- The same as the previous item, but now for *signed* permutations.

Exercise 4

(3 Points)

In the overlap graph $O(\pi)$, applying a reversal induced by a vertex v will have two effects:

- The subgraph induced by v and its neighbours is complemented.
- All neighbours of v have their orientation inverted.

Using only the fact that v is an oriented vertex (odd degree) and the item (i), prove item (ii), that is, prove that all neighbours of v have their orientation inverted.

Tip: for a neighbour u of v , check how many edges it loses and how many it gains after the reversal, and check what is the parity of the total change.