

## Algorithms for Bioinformatics (Autumn 2015)

### Exercise 5 (Tue 6.10., 10-12, B222)

If you cannot make it to the exercise session, please e-mail your solutions and the reason why you cannot attend to [daniel.valenzuela@cs.helsinki.fi](mailto:daniel.valenzuela@cs.helsinki.fi) before the exercise session to get credit.

Some of the problems below are programming exercises on the Rosalind platform at <http://rosalind.info/problems/list-view/?location=bioinformatics-textbook-track>

1. Solve the Rosalind problem BA6A: *Implement GreedySorting to Sort a Permutation by Reversals.*
2. Solve the Rosalind problem BA6B: *Compute the Number of Breakpoints in a Permutation.*
3. Prove that if a signed permutation  $P$  contains negative signs, there is always a reversal that decreases  $\text{Breakpoints}(P)$ .
4. Perform the breakpoint reversal sort algorithm with  $P = (+3 +4 -6 -5 +7 +1 -8 -2)$  as the input and show all intermediate permutations. Is this the optimal solution to this instance of reversal sorting problem?
5. Compute the 2-break distance  $d(P, Q)$  for  $P = (+a +b +c +d +e +f +g +h)$  and  $Q = (+c +d -f -e +g +a -h -b)$ . Show the breakpoint graph  $\text{Breakpoint}(P, Q)$ .