## Exercises – Algorithms for Genome Research

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http://wiki.techfak.uni-bielefeld.de/gi/Teaching/2014 winter/AlgoGR

## Exercise List 3 — 14.11.2014

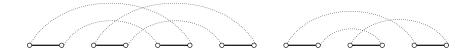
Discussion of exercises on: 21.11.2014

**Exercise 1** For the following permutations, answer the questions:

 $\begin{aligned} \pi &= (0 + 3 + 5 + 4 + 6 + 2 + 1 + 7 - 8 + 9) \\ \pi &= (0 + 2 + 4 + 3 + 5 + 1 + 6 + 8 + 10 + 9 + 11 + 7 + 12 + 14 + 16 + 15 + 17 + 13 + 18) \\ \pi &= (0 + 2 + 3 - 4 + 5 + 7 + 6 + 8 + 1 + 9 + 11 + 10 + 12) \\ \pi &= (0 + 7 + 1 - 2 + 3 + 5 + 4 + 6 + 8 + 10 + 9 + 11) \\ \pi &= (0 + 1 - 2 + 3 + 7 + 5 + 6 + 4 + 8 + 10 + 13 + 11 + 12 + 9 + 13) \end{aligned}$ 

- (a) Find the components in the permutation, and the type of each component.
- (b) Compare the components that you found with the breakpoint graph components. *Tip: use the available software*
- (c) Draw the component tree  $T_P$ .
- (d) What is the reversal distance?

**Exercise 2** Consider the following breakpoint graph.



- (a) What is the reversal distance for this breakpoint graph?
- (b) Find a permutation  $\pi$  that has this breakpoint graph.

**Exercise 3** After a *cycle merge* reversal, that is, a reversal defined by black edges in two different cycles, the two cycles are merged into one. Prove that this new cycle is always oriented. *Tip: What happens when we apply the same reversal again?*