

# Exercises – Algorithms for Genome Research

Universität Bielefeld, WS 2014, Dr. Pedro Feijao

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## Exercise List 11 — 23.01.2015

Discussion of exercises on: 30.01.2015

**Exercise 1** Find all common intervals of the two permutations below:

$$\sigma_A = (1, 2, 3, 4, 5, 6, 7, 8)$$

$$\sigma_B = (5, 3, 1, 4, 2, 8, 7, 6)$$

### Exercise 2

Read the following paper:

Uno, T., & Yagiura, M. (2000). **Fast Algorithms to Enumerate All Common Intervals of Two Permutations**. *Algorithmica*, 26(2), 290-309.

On section 4. **An Algorithm with  $O(n + K)$  Worst-Case Running Time.**, the authors describe the algorithm seen in the lecture. About this algorithm, answer the following questions:

- (a) Describe the data structure used to store the values of  $u(x, y)$ ,  $l(x, u)$ , and the list of  $y$ .
- (b) How does the data structure change when the next  $x - 1$  is included?

### Exercise 3

Write an algorithm that finds all the common intervals between  $k$  permutations, with  $k > 2$ . You can use a common intervals algorithm for 2 permutations as an internal step.