

# Algorithms in Comparative Genomics

Universität Bielefeld, WS 2021/2022

Dr. Marília D. V. Braga · Leonard Bohnenkämper

<https://gi.cebitec.uni-bielefeld.de/teaching/2021winter/cg>

## Exercise sheet 7, 09.12.2021

### Exercise 1 (Bounds for the SCJ distance)

(6 pts)

Theoretical bounds for the SCJ distance with respect to the DCJ and the inversion distances are

$$d_{\text{DCJ}}(\mathbb{A}_{\triangleright}^f, \mathbb{B}_{\triangleright}^f) \leq d_{\text{SCJ}}(\mathbb{A}_{\triangleright}^f, \mathbb{B}_{\triangleright}^f) \leq 4 d_{\text{DCJ}}(\mathbb{A}_{\triangleright}^f, \mathbb{B}_{\triangleright}^f)$$

$$2 d_{\text{INV}}(\mathbb{A}_{\triangleright}^f, \mathbb{B}_{\triangleright}^f) \leq d_{\text{SCJ}}(\mathbb{A}_{\triangleright}^f, \mathbb{B}_{\triangleright}^f) \leq 4 d_{\text{INV}}(\mathbb{A}_{\triangleright}^f, \mathbb{B}_{\triangleright}^f)$$

For each one of these four bounds:

Show that it is tight by giving an example of pairs of mutually distinct genomes that fulfill it.

### Exercise 2 (Singular DCJ-indel model)

(12 pts)

Consider the following singular genomes:

$$\mathbb{A} = [a_1 \ 4 \ 1] \ [a_2 \ 6 \ a_3 \ 3 \ a_4 \ 2 \ a_5 \ 5] \ [a_6 \ a_7 \ 7 \ 8 \ a_8 \ \bar{9}] \ [a_9] \ \text{and}$$

$$\mathbb{B} = [b_1 \ 1 \ b_2 \ 2 \ b_3 \ 3 \ b_4 \ 4 \ b_5] \ [5 \ b_6 \ 6] \ [b_7 \ 7 \ 8 \ b_8 \ 9] \ (b_9 \ b_{10})$$

1. Give the sets of genes  $\mathcal{G}_*$ ,  $\mathcal{A}$  and  $\mathcal{B}$ .
2. Construct the relational graph  $RG(\mathbb{A}, \mathbb{B})$ .
3. For each component  $C$  of the relational graph  $RG(\mathbb{A}, \mathbb{B})$ :
  - (a) Give the type of  $C$  (cycle, singleton,  $\mathbb{A}\mathbb{B}$ -path,  $\mathbb{A}\mathbb{A}$ -path or  $\mathbb{B}\mathbb{B}$ -path).
  - (b) Give the number of runs  $\Lambda(C)$ ;
  - (c) Give the run-type of  $C$  ( $\varepsilon$ ,  $\mathcal{A}$ ,  $\mathcal{B}$ ,  $\mathcal{AB}$  or  $\mathcal{BA}$ );
  - (d) Compute the minimum number of indels  $\lambda(C)$  that are necessary for sorting  $C$  separately.
4. Find all chains of deducing recombinations.
5. Compute the DCJ-indel distance  $d_{\text{DCJ}}^{\text{ID}}(\mathbb{A}, \mathbb{B})$ .