## Algorithms in Comparative Genomics

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## Exercise sheet 11, 26.01.2024

**Exercise 1 (Calculating Distances)** 

Given two linear, unichromosomal genomes

$$A = [12 - 4 - 35 - 6]$$

and

B = [1 - 3 - 2456]

- 1. What is the breakpoint distance between A and B?
- 2. What is the SCJ distance between A and B?
- 3. Give an SCJ sorting scenario from A to B.
- 4. Draw the adjacency graph of A and B.
- 5. What is the DCJ distance between A and B?
- 6. Give a DCJ sorting scenario from A to B.

**Exercise 2 (Cooking Recipe)** The explode chromosome operation  $(ECO)^1$  transforms a chromosome  $[g_1...g_k]$  or  $(g_1...g_k)$  of arbitrary length k into k linear chromosomes  $[g_1], ..., [g_k]$  with one marker  $g_i$  each. The assemble chromosome operation  $(ACO)^2$  transforms a set of linear chromosomes  $\{[C_1], ..., [C_l]\}$  into a linear chromosome [S] or circular chromosome (S) with  $S = C_1 C_2 ... C_l$  for any orientation and order of  $C_1$  to  $C_l$ .

For example:  $[1 2 3], [4 5] \xrightarrow{ECO} [1], [2], [3], [45] \equiv [1], [-5 - 4], [3], [-2] \xrightarrow{ACO} [1 - 5 - 4 3 - 2].$ 

- 1. Familiarize yourself with the operations. Give optimal sorting scenarios for the following:
  - (a) Sort [1 2 3], (4 5) into [1 3 2], (4 5).
  - (b) Sort [1 2 3] into [1], [2], [3].
  - (c) Sort [1], (234), [5] into [1234], [5].
- 2. Use the "cooking recipe" from the lecture to derive a general formula for the minimum number of ECOs and ACOs needed to transform one genome into the other. You do not need to invent a data structure for this. The "quantity" you need for the cooking recipe can be derived directly from the genomes themselves.

## Exercise 3 (Feather Median)

Given 2k + 3 genomes  $G_1, G_2, G_3, \dots G_{2k+3}$  and an algorithm to compute the median  $M_d(A, B, C)$  of three genomes (A, B, C) under a distance model d. The Feather Median<sup>3</sup> is defined as

$$M_f(G_1, G_2, G_3, \dots G_{2k+3-2}, G_{2k+3-1}, G_{2k+3}) = M_d(G_{2k+3-1}, M_f(G_1, G_2, G_3, \dots G_{2k+3-2}), G_{2k+3})$$
(1)

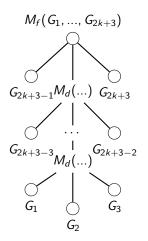
with recursion base

$$M_f(G_1, G_2, G_3) = M_d(G_1, G_2, G_3)$$
<sup>(2)</sup>

<sup>&</sup>lt;sup>1</sup>which I made up.

<sup>&</sup>lt;sup>2</sup>which I also made up :)

<sup>&</sup>lt;sup>3</sup>which I also made up ;)



- 1. Disprove (e.g. via counter example): The Feather Median under the breakpoint distance is always a breakpoint median<sup>4</sup>.
- 2. Prove or disprove: No metric d on a set with two or more distinct elements exists, under which the Feather Median is always a true Median. (Spoiler 1, 2, 3, 4, 5, 6).

<sup>&</sup>lt;sup>4</sup>The true median of a set  $K \subseteq S$  under metric d on space S being the element  $M_d \in S$  that minimizes  $m(M_d) = \sum_{k \in K} d(M_d, k)$ .