

**Algorithms in Comparative Genomics**  
**Summer 2018**

**Exercises**

**Number 6, return 2018 June 22**

1. Consider the linear genomes  $A = [1 \ a \ -2 \ b \ 3]$  and  $B = [1 \ c \ 2 \ d \ 3]$ .
  - (a) Draw the adjacency graph  $AG(A, B)$ . What is the DCJ-indel distance between  $A$  and  $B$ ?
  - (b) Find a scenario with 3 indels, applying only optimal DCJs. Redraw the adjacency graph after the first DCJ.
  - (c) Find a scenario with 2 indels. Redraw the adjacency graph after the first DCJ.  
*(Hint: Find a neutral DCJ that reduces the number of runs.)*
2. Prove the formula for the DCJ-indel potential  $\lambda$  of a cycle  $C$  of the adjacency graph  $AG(A, B)$ ,

$$\lambda(C) = \left\lceil \frac{\Lambda(C) + 1}{2} \right\rceil,$$

where  $\Lambda(C) > 0$  is the number of runs of indels in  $C$ .

How does the corresponding formula for a path  $P$  with  $\Lambda(P)$  runs look like?

3. Consider the following two genomes  $A = [1 \ 2 \ 3 \ 4 \ 5 \ 6]$  and  $B = [-3 \ 1 \ 2 \ -6 \ 5]$ .
  - (a) Find a third genome  $C$  such that the triangle inequality for the uncorrected DCJ-indel distance does not hold, i.e.

$$d_{\text{DCJ-indel}}(A, B) > d_{\text{DCJ-indel}}(A, C) + d_{\text{DCJ-indel}}(C, B).$$

- (b) Show how the distance can be corrected such that the triangle inequality holds again.