TME2 - PHYG

Witold PODLEJSKI & Jérémie PERRIN

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- 1 Exercise 1 : Genome evolution simulation
- 1.1 Question 1
- 1.2 Question 2
- 1.3 Question 3
- 2 Exercise 2: Implementation of simple events

Cf. folder "code".

- 3 Exercise 3: Small dataset simulation
- 3.1 Question 1

Cf. folder "code/results/[1-10]".

3.2 Question 2

| | Sim 1 | Sim 2 | Sim 3 | Sim 4 | Sim 5 | Sim 6 | Sim 7 | Sim 8 | Sim 9 | Sim 10 |
|---------------|-------|-------|-------|-------|-------|-------|-------|-------|-------|--------|
| Inversion | 7 | 10 | 11 | 11 | 12 | 10 | 7 | 8 | 7 | 10 |
| Translocation | 8 | 1 | 5 | 1 | 4 | 7 | 6 | 2 | 1 | 6 |
| Duplication | 0 | 1 | 0 | 1 | 2 | 0 | 1 | 3 | 3 | 0 |
| Deletions | 0 | 2 | 1 | 1 | 2 | 0 | 3 | 2 | 4 | 0 |
| Fusions | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Fissions | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| WGD | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |

3.3 Question 3

3.4 Question 4-5



Figure 1: Real tree for simulation 1



Figure 2: The resulting tree of Maximum of likelihood algorithm for simulation $\mathbf{1}$



Figure 3: Real tree for simulation 6

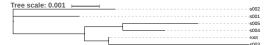


Figure 4: The resulting tree of Maximum of likelihood algorithm for simulation 6