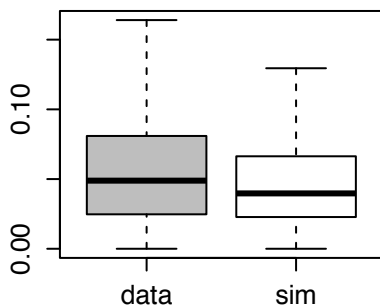
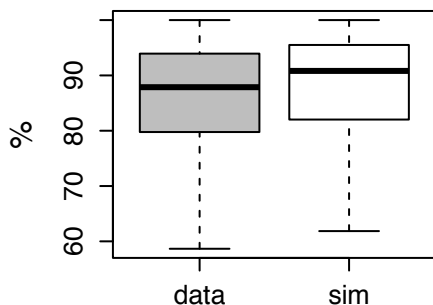


relative edit distance

**average distance within alignment**

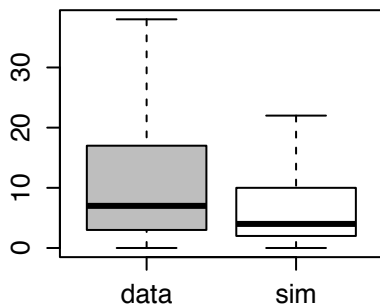


**% of invariant sites per alignment**

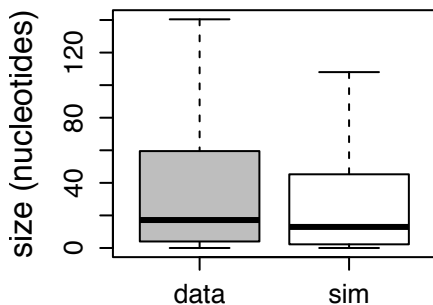


count

**Number of indels per alignment**

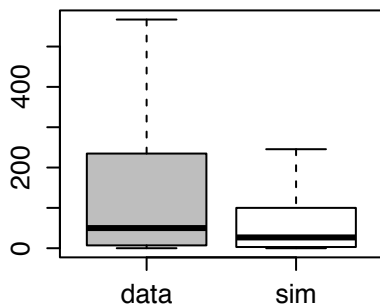


**Average size of indels per alignment**



gap count

**Number of gaps per sequence**



**% gaps per sequence**

