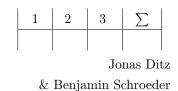
Bioinformatics I

WS 15/16

Tutor: Alexander Seitz



Assignment 3

(Abgabe am 2. November 2015)

Exercise 1 - Equivalence of distance and similarity alignments

Exercise 1 - Using BLAT to align 454 reads to the Helicobacter pylori genome

Exercise 1 - Bonus: Use SSAHA2 to align 535 reads to the Helicobacter pylori genome