

Bioinformatics I

WS 15/16

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## 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***