

deepBlockAlign v1.0

Block group alignment: cluster-483-chr9-His-GTG with cluster-199-chr16-Gly-CCC (Score: 0.755482)

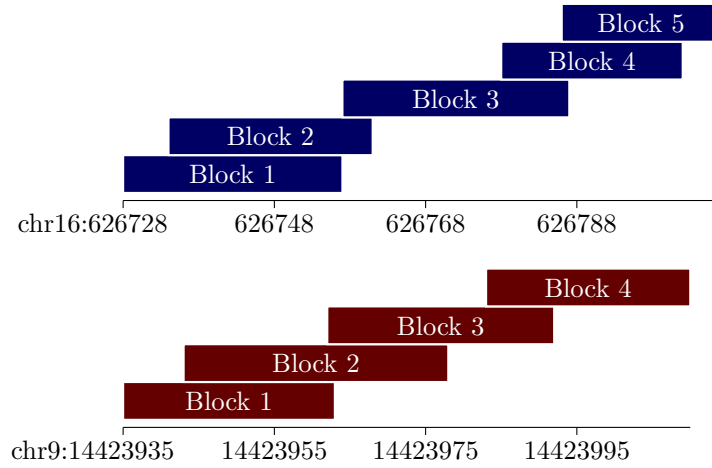


Table 1: Aligned Blocks

tRNA	tRNA
Block 1	Block 1
Block 2	Block 2
Block 3	Block 3
Block 4	Block 4

Block alignment:

