

deepBlockAlign v1.0

Block group alignment: cluster-201-chr16-Lys-CTT with cluster-736-chr16-Pro-TGG (Score: 0.817387)

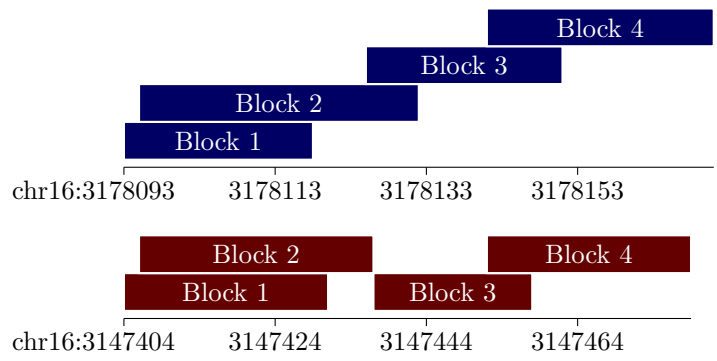


Table 1: Aligned Blocks

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

Block alignment:

