

deepBlockAlign v1.0

Block group alignment: cluster-638-chr11-Lys-TTT with cluster-1003-chr6-Gln-CTG (Score: 0.571359)

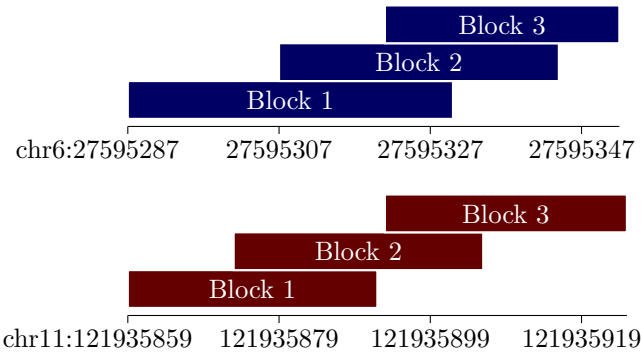


Table 1: Aligned Blocks

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

Block alignment:

