

deepBlockAlign v1.0

Block group alignment: cluster-271-chr2-Gly-CCC with cluster-1007-chr6-Ala-TGC (Score: 0.832154)

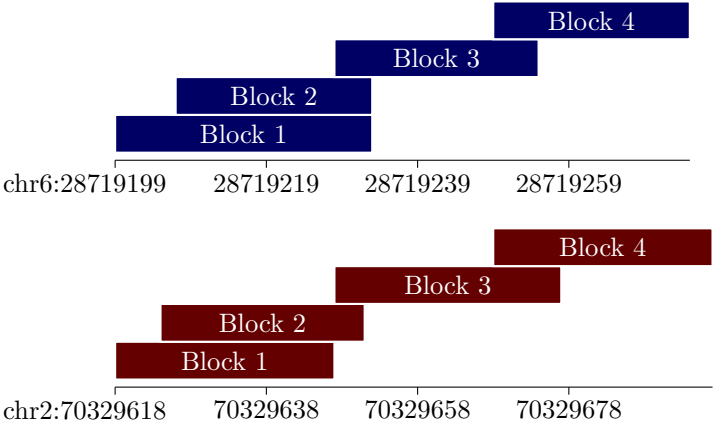


Table 1: Aligned Blocks

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

Block alignment:

