

deepBlockAlign v1.0

Block group alignment: cluster-413-chr6-Leu-CAA with cluster-994-chr6-Tyr-GTA (Score: 0.737246)

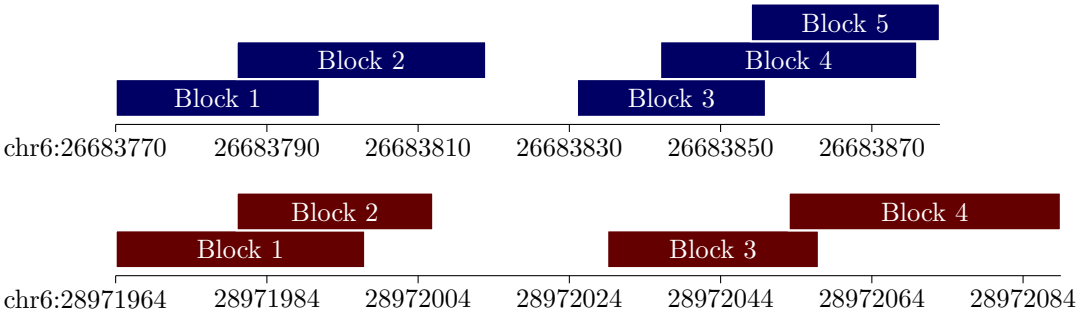


Table 1: Aligned Blocks

tRNA	tRNA
Block 1	Block 1
Block 2	Block 2
Block 3	Block 3
Block 4	Block 5

Block alignment:

