

deepBlockAlign v1.0

Block group alignment: *cluster-991-chr6-Ala-CGC* with *cluster-674-chr14-Pro-TGG* (Score: 0.815323)

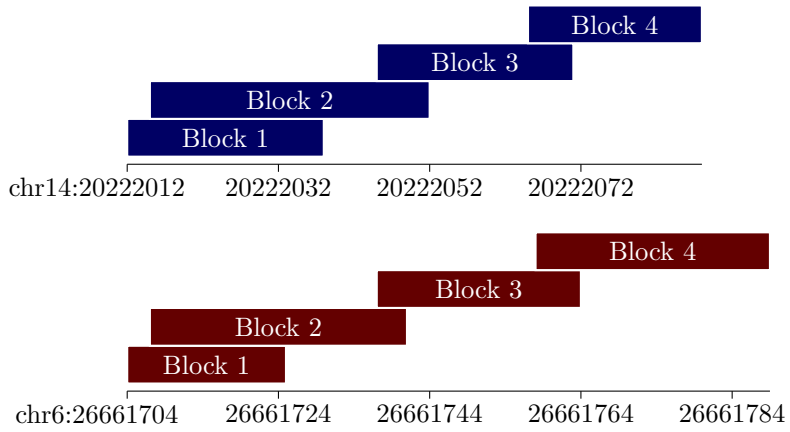


Table 1: Aligned Blocks

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

Block alignment:

