

deepBlockAlign v1.0

Block group alignment: cluster-228-chr17-Cys-GCA with cluster-184-chr15-His-GTG (Score: 0.709088)

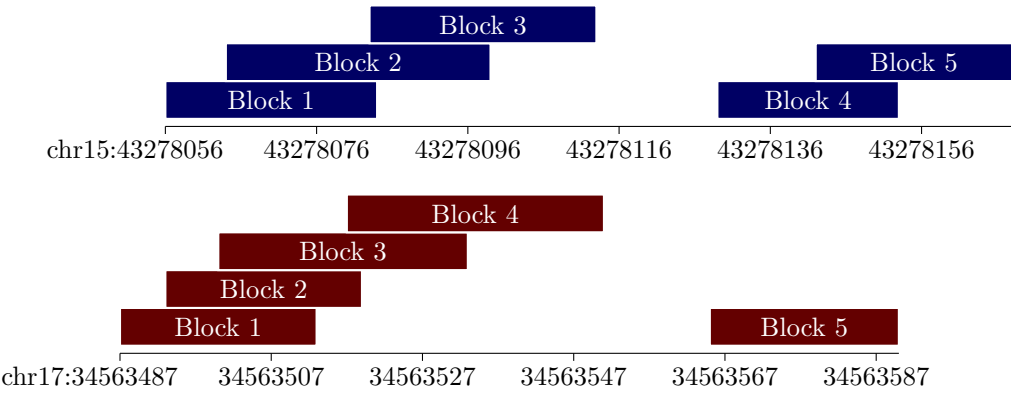


Table 1: Aligned Blocks

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

Block alignment:

