

deepBlockAlign v1.0

Block group alignment: *cluster-853-chr2-Ala-AGC* with *cluster-767-U49A* (Score: 0.813908)

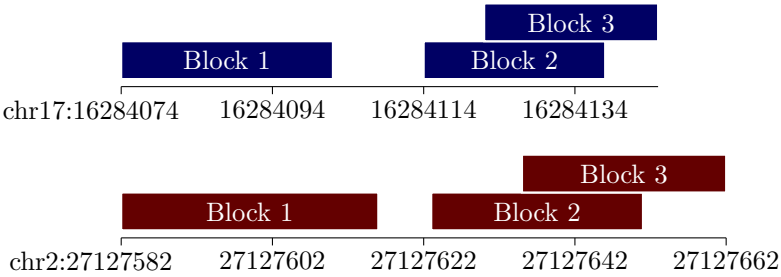


Table 1: Aligned Blocks

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

Block alignment:

