

deepBlockAlign v1.0

Block group alignment: cluster-1070-chr8-Ala-AGC with cluster-163-hsa-mir-15a (Score: 0.759703)

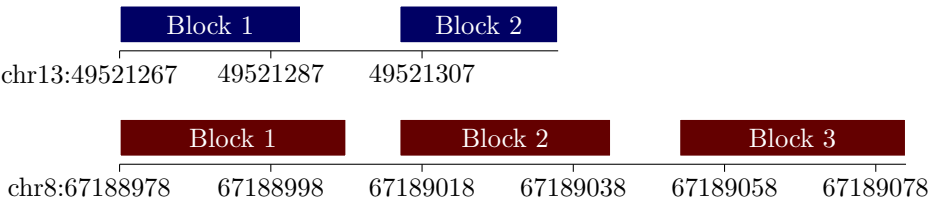


Table 1: Aligned Blocks

tRNA	miRNA
Block 1	Block 1
Block 2	Block 2

Block alignment:

