

deepBlockAlign v1.0

Block group alignment: cluster-218-chr17-Ser-AGA with cluster-841-hsa-mir-516b-1 (Score: 0.784787)

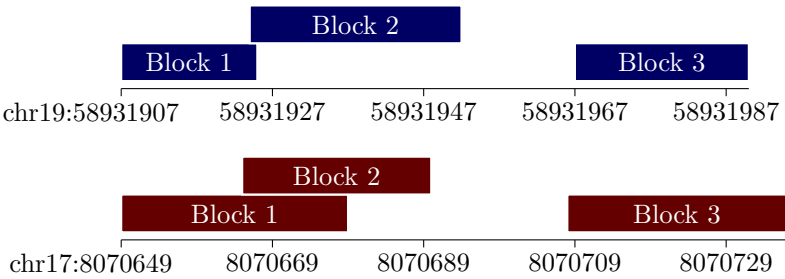


Table 1: Aligned Blocks

tRNA	miRNA
Block 1	Block 1
Block 2	Block 2
Block 3	Block 3

Block alignment:

