

deepBlockAlign v1.0

Block group alignment: cluster-115-U27 with cluster-49-hsa-mir-181b-1 (Score: 0.872690)

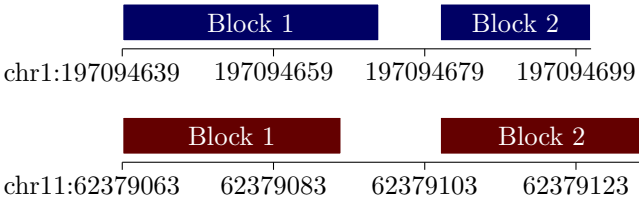


Table 1: Aligned Blocks

snoRNA_CD	miRNA
Block 1	Block 1
Block 2	Block 2

Block alignment:

