

deepBlockAlign v1.0

Block group alignment: cluster-114-U28 with cluster-501-hsa-mir-199b (Score: 0.940100)

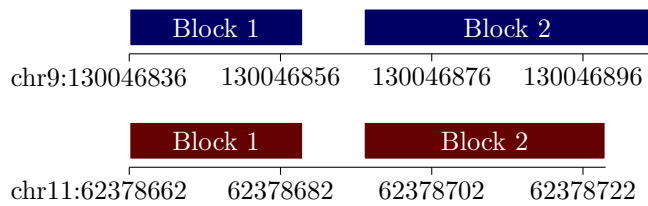


Table 1: Aligned Blocks

snoRNA_CD	miRNA
Block 1	Block 1
Block 2	Block 2

Block alignment:

