

deepBlockAlign v1.0

Block group alignment: cluster-557-U38B with cluster-822-hsa-mir-519c (Score: 0.669391)

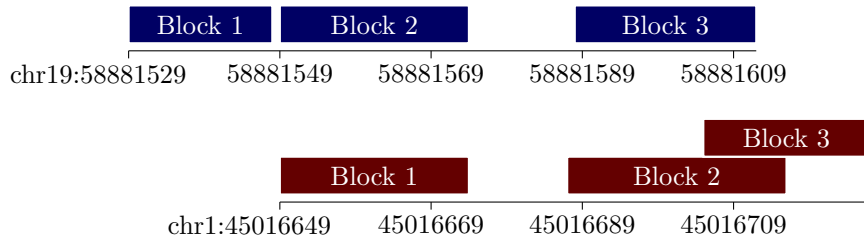


Table 1: Aligned Blocks

snoRNA_CD	miRNA
Block 1	Block 2
Block 2	Block 3

Block alignment:

