

deepBlockAlign v1.0

Block group alignment: cluster-711-HBII-85-29 with cluster-669-hsa-mir-20a (Score: 0.692862)

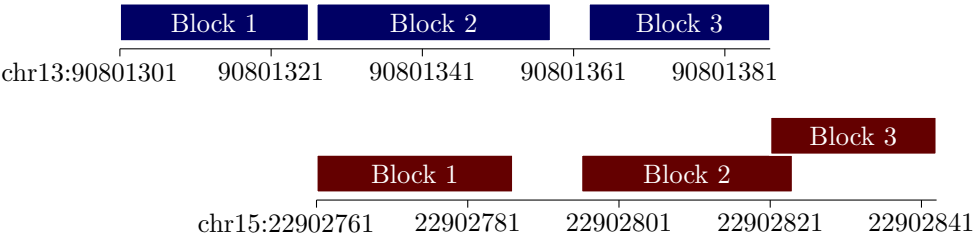


Table 1: Aligned Blocks

snoRNA_CD	miRNA
Block 1	Block 2
Block 2	Block 3

Block alignment:

