

deepBlockAlign v1.0

Block group alignment: cluster-320-U43 with cluster-815-hsa-mir-125a (Score: 0.930653)

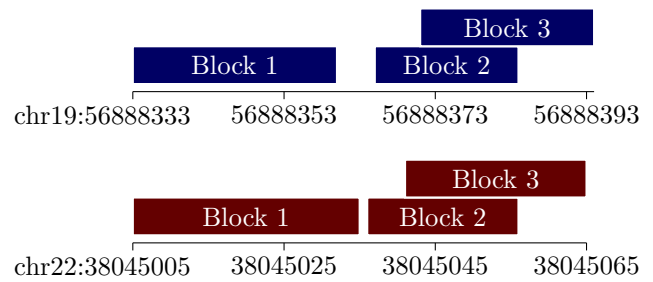


Table 1: Aligned Blocks

snoRNA_CD	miRNA
Block 1	Block 1
Block 2	Block 2
Block 3	Block 3

Block alignment:

