

deepBlockAlign v1.0

Block group alignment: *cluster-303-U20* with *cluster-843-hsa-mir-371* (Score: 0.759263)

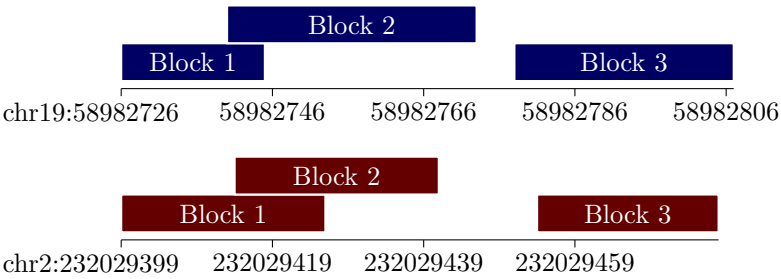


Table 1: Aligned Blocks

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

Block alignment:

