

deepBlockAlign v1.0

Block group alignment: cluster-117-U25 with cluster-519-hsa-mir-374a (Score: 0.755549)

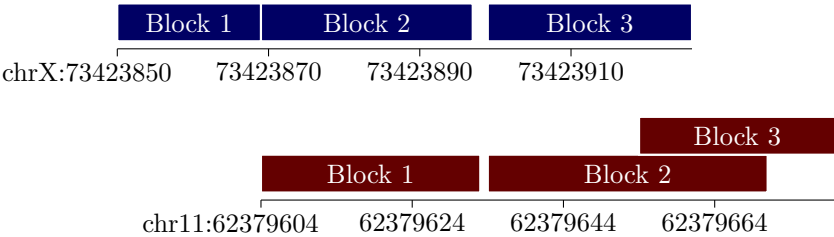


Table 1: Aligned Blocks

snoRNA_CD	miRNA
Block 1	Block 2
Block 2	Block 3

Block alignment:

