

deepBlockAlign v1.0

Block group alignment: *cluster-9-ACA61* with *cluster-517-hsa-mir-421* (Score: 0.819691)

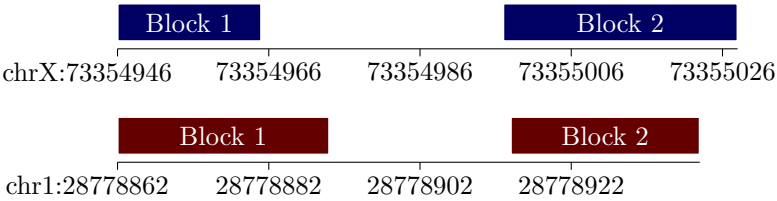


Table 1: Aligned Blocks

snoRNA_HACA	miRNA
Block 1	Block 1
Block 2	Block 2

Block alignment:

