

deepBlockAlign v1.0

Block group alignment: cluster-45-HBI-100 with cluster-52-hsa-mir-135b (Score: 0.927850)

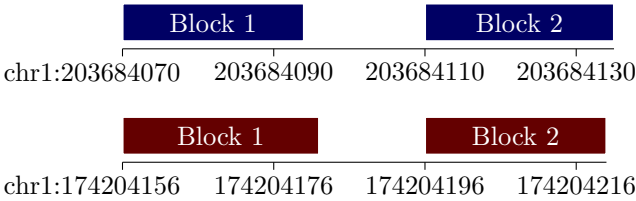


Table 1: Aligned Blocks

snoRNA_HACA	miRNA
Block 1	Block 1
Block 2	Block 2

Block alignment:

