

deepBlockAlign v1.0

Block group alignment: *cluster-929-E3* with *cluster-30-hsa-mir-9-1* (Score: 0.937599)

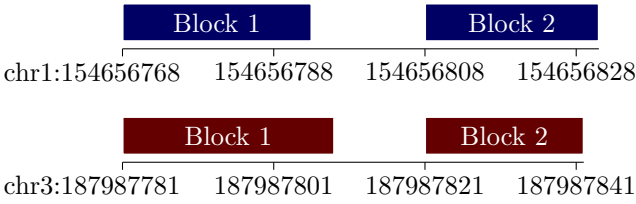


Table 1: Aligned Blocks

snoRNA_HACA	miRNA
Block 1	Block 1
Block 2	Block 2

Block alignment:

