

deepBlockAlign v1.0

Block group alignment: *cluster-58-ACA36B* with *cluster-482-hsa-mir-151* (Score: 0.858776)

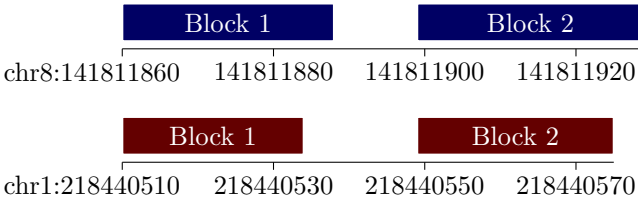


Table 1: Aligned Blocks

snoRNA_HACA	miRNA
Block 1	Block 1
Block 2	Block 2

Block alignment:

