

deepBlockAlign v1.0

Block group alignment: *cluster-10-HBII-251* with *cluster-1046-hsa-mir-129-1* (Score: 0.866384)

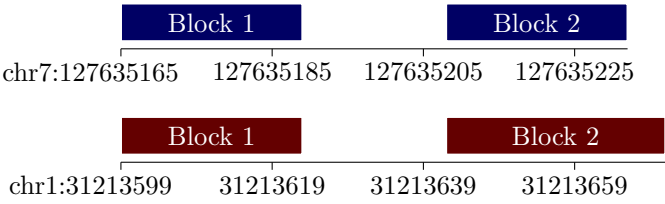


Table 1: Aligned Blocks

snoRNA_CD	miRNA
Block 1	Block 1
Block 2	Block 2

Block alignment:

