

deepBlockAlign v1.0

Block group alignment: cluster-428-n-a with cluster-620-hsa-mir-129-2 (Score: 0.852714)

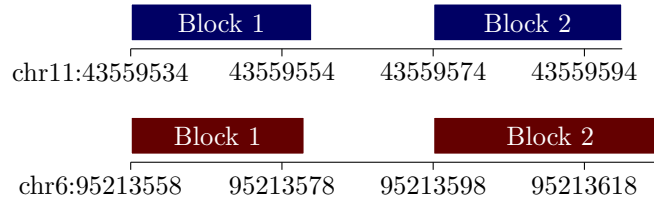


Table 1: Aligned Blocks

n/a	miRNA
Block 1	Block 1
Block 2	Block 2

Block alignment:

