

deepBlockAlign v1.0

Block group alignment: cluster-723-ACA45 with cluster-461-hsa-mir-486 (Score: 0.923554)

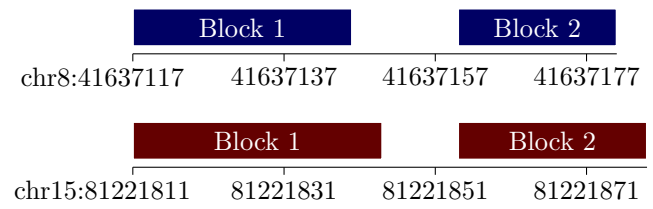


Table 1: Aligned Blocks

snoRNA_scaRna	miRNA
Block 1	Block 1
Block 2	Block 2

Block alignment:

