

deepBlockAlign v1.0

Block group alignment: *cluster-216-U8* with *cluster-923-hsa-mir-16-2* (Score: 0.883556)

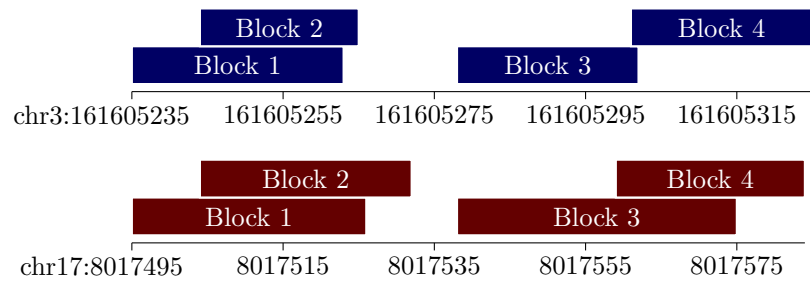


Table 1: Aligned Blocks

snoRNA_CD	miRNA
Block 1	Block 1
Block 2	Block 2
Block 3	Block 3
Block 4	Block 4

Block alignment:

