

deepBlockAlign v1.0

Block group alignment: *cluster-989-chr6-Ser-AGA* with *cluster-494-hsa-mir-7-1* (Score: 0.780132)

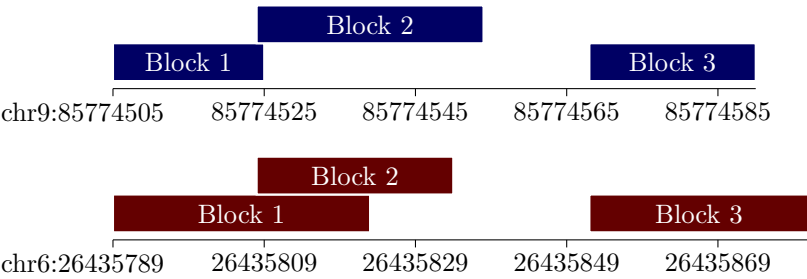


Table 1: Aligned Blocks

tRNA	miRNA
Block 1	Block 1
Block 2	Block 2
Block 3	Block 3

Block alignment:

