

deepBlockAlign v1.0

Block group alignment: *cluster-305-SNORD119* with *cluster-1002-chr6-Ser-GCT* (Score: 0.870516)

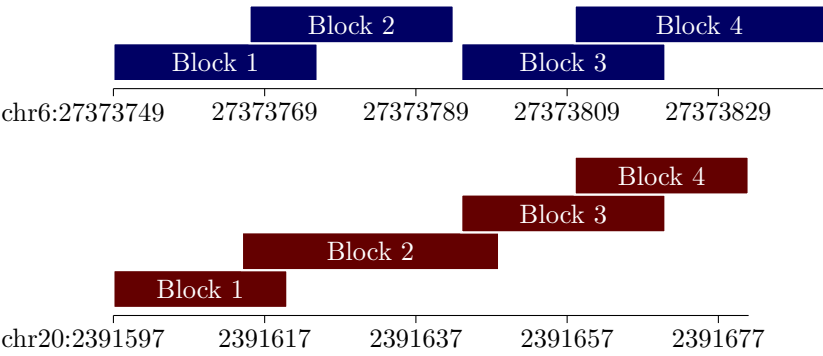


Table 1: Aligned Blocks

snoRNA_CD	tRNA
Block 1	Block 1
Block 2	Block 2
Block 3	Block 3
Block 4	Block 4

Block alignment:

