

deepBlockAlign v1.0

Block group alignment: cluster-710-HBII-85-27 with cluster-163-hsa-mir-15a (Score: 0.754321)

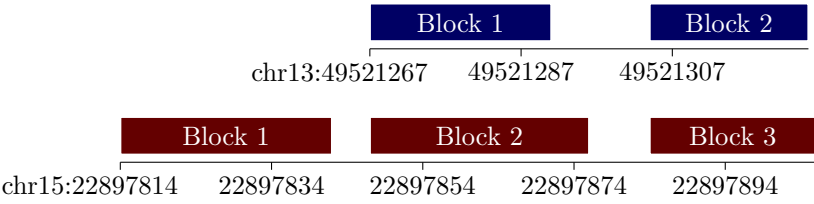


Table 1: Aligned Blocks

| snoRNA_CD | miRNA   |
|-----------|---------|
| Block 2   | Block 1 |
| Block 3   | Block 2 |

Block alignment:

