

deepBlockAlign v1.0

Block group alignment: *cluster-1013-U52* with *cluster-93-hsa-mir-483* (Score: 0.832938)

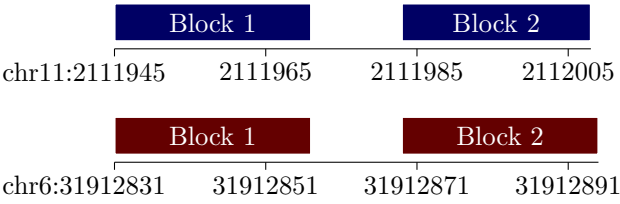


Table 1: Aligned Blocks

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

Block alignment:

