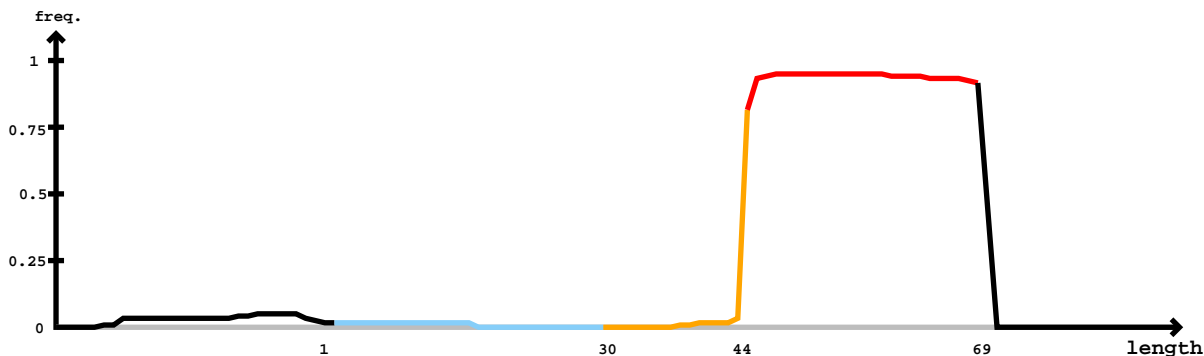
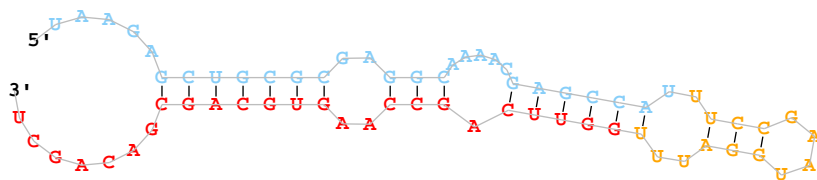


Provisional ID : 3R\_dna:chromosome\_chromosome:BDGP6:3R:1:32079331:1\_REF\_21941  
 Score total : 1.6  
 Score for star read(s) : -1.3  
 Score for read counts : 0  
 Score for mfe : 1.9  
 Score for randfold : 1.6  
 Score for cons. seed : -0.6  
 Total read count : 111  
 Mature read count : 111  
 Loop read count : 0  
 Star read count : 0



Star

Mature

| 5' -  | reads | mm | sample |
|---|-------|----|--------|
| aguuuccuuaauuaagggugccaaa <u>uaagagcugcgagggcaaaacagacc</u> au <u>uuccgaauggauuugguucagccaagugcagcgacagcu</u> uugcuuuuaugaaacuguu | 1     | 0  | seq    |
| .....uuccuuaauuaagggugccaaa.....  | 2     | 0  | seq    |
| .....ccuuaauuaagggugccca.....   | 1     | 0  | seq    |
| .....ccuuaauuaagggugccaa.....   | 1     | 0  | seq    |
| .....guggccaaaguaagagcugcgcgag.....   | 1     | 0  | seq    |
| .....ggccaaaguaagagcugcgcgag.....   | 1     | 0  | seq    |
| .....uggauuugguucagccaagugcagcg.....  | 1     | 0  | seq    |
| .....gauuugguucagccaagugc.....  | 1     | 0  | seq    |
| .....Agguucagccaagugcagcgacagcu.....  | 1     | 1  | seq    |
| .....ugguucagccaagugcagcgacagcu.....  | 1     | 0  | seq    |
| .....gguucagccaagugcagcgacag.....   | 1     | 0  | seq    |
| .....gguucagccaagugcagcgacagc.....  | 1     | 0  | seq    |
| .....gguucagccaagugcagcgacaAcu.....   | 1     | 1  | seq    |
| .....gguucagccaagCgcagcgacagcu.....   | 1     | 1  | seq    |
| .....gguucagccaagugcagcgacagAu.....   | 1     | 1  | seq    |
| .....gguucagccaagugcagcgacaNcu.....   | 1     | 1  | seq    |
| .....ggGuucagccaagugcagcgacagcu.....  | 1     | 1  | seq    |
| .....gguucagccaagUAcagcgacagcu.....   | 1     | 1  | seq    |
| .....Cguucagccaagugcagcgacagcu.....   | 1     | 1  | seq    |
| .....gguucagccaagugcagcgacGgcu.....   | 1     | 1  | seq    |
| .....gguucagccaagugcagcgacagcu.....   | 82    | 0  | seq    |
| .....gguucagccaagugcagcgacagcuu.....  | 1     | 0  | seq    |
| .....guucagccaagugcagcgacagcu.....  | 8     | 0  | seq    |
| .....guucagccaagCgcagcgacagcu.....  | 1     | 1  | seq    |
| .....guucagccaagugcagcgacagcu.....  | 1     | 1  | seq    |
| .....guucagUcaagugcagcgacagcuu.....   | 1     | 1  | seq    |
| .....guucagccaagugcagcgacagcuu.....   | 3     | 0  | seq    |
| .....uucagccaagugcagcgacagcu.....   | 1     | 0  | seq    |
| .....ucaAccaagugcagcgacagcu.....  | 1     | 1  | seq    |