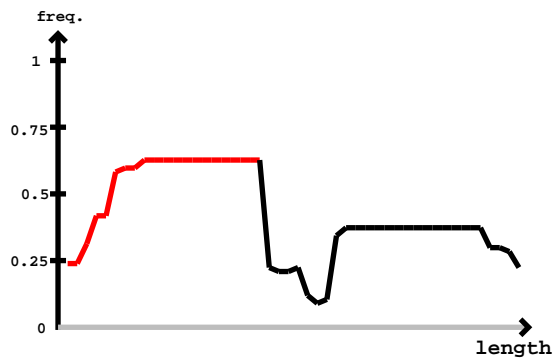
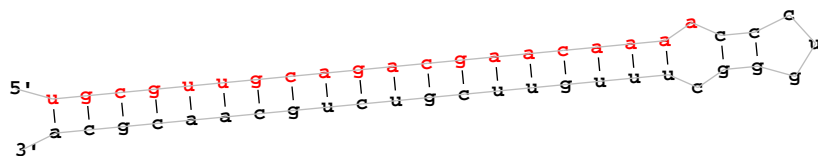


miRBase precursor : chromosome\_5\_19070  
Total read count : 67  
chromosome\_5\_19070 read count : 67  
remaining reads : 26



chromosome_5_19070			
5'-ugcguugcagacgaacaaaaaccugggcuuuguucgucugcaacgca-3'	exp		
((((((((((((((((((((.....))))))))))))))))))))))	reads	mm	sample
ugcguugcagacgaacaaaa.....	6	0	seq
ugcguuAcagacgaacaaaa.....	1	1	seq
ugcguugcagacgaalaaaa.....	1	1	seq
Agcguugcagacgaacaaaa.....	2	1	seq
ugcguugcagacgaacaaaG.....	5	1	seq
ugcguugcagacgaGcaaaa.....	1	1	seq
.gcuugcagacgaacaaaa.....	4	0	seq
.Acguugcagacgaacaaaa.....	1	1	seq
.Gguugcagacgaacaaaa.....	1	1	seq
.Aguugcagacgaacaaaa.....	5	1	seq
.cguugcagacgaacaaaac.....	1	0	seq
...uugcagacgaacaaaacc.....	1	0	seq
...uugcagacgaacUaaacc.....	2	1	seq
...uugcagacgaacaaaaccA.....	6	1	seq
...uugcagacgaacaaaaccu.....	1	0	seq
...uugcagacgaacaaaaccCggg.....	1	1	seq
...ugcagacgaacaaaacc.....	1	0	seq
.....cagaUgaacaaaaccug.....	1	1	seq
.....cagacgaacaaaaccug.....	1	0	seq
.....ugggUuuuguucgucugcaa.....	5	1	seq
.....gUuuuguucgucugcaacg.....	1	1	seq
.....Uuuuguucgucugcaacgc.....	4	1	seq
.....Uuuuguucgucugcaacgca.....	8	1	seq
.....cuuuguucgucugcaacgca.....	5	0	seq
.....uuuguucgucugUaacgca.....	2	1	seq