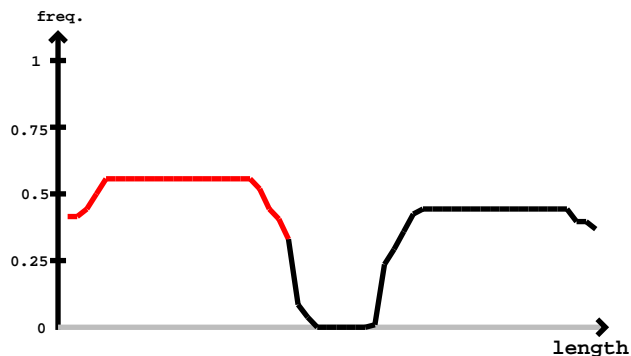
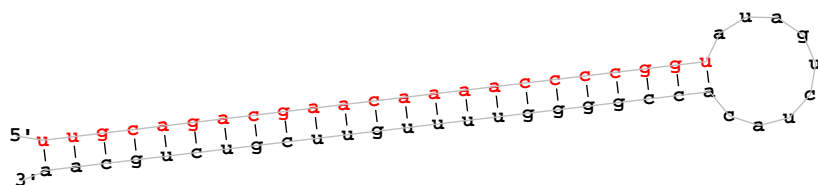


miRBase precursor : chromosome_11_52764
 Total read count : 106
 chromosome_11_52764 read count : 50
 remaining reads : 47



chromosome_11_52764

5'-	reads	mm	sample
uugcagacgaacUaaaccc.....	2	1	seq
uugcagacgaacaaaaccc.....	1	0	seq
uugcagacgaacaaaacccA.....	6	1	seq
uugcagacgaacaaaaccc.....	1	0	seq
uugcagacgaacaaaaccccg.....	3	0	seq
uugcagacgaacaaaacccU.....	1	1	seq
uugcagacgaacaaaaccccg.....	3	0	seq
uugcagacgaacaaaaccccgC.....	3	1	seq
uugcagacgaacaaaaccccgG.....	1	1	seq
uugcagacgaacaaaaccccggu.....	21	0	seq
uugcagacgaacaaaaccccgguA.....	2	0	seq
.ugcagacgaacaaaaccc.....	1	0	seq
.ugcagacgaacaaaaccc.....	1	0	seq
.ugcagacgaacaaaaccccgguU.....	1	1	seq
.gcagacgGacaaaaccccgguA.....	2	1	seq
.gGagacgaacaaaaccccgguau.....	3	1	seq
.gcagacgaacaaaacccUcgguau.....	1	1	seq
..cagacgaacaaaaccccgA.....	1	1	seq
..cagacgaacaaaaccccg.....	3	0	seq
..cagacgaacaaaaccccgC.....	1	1	seq
..cagacgaacaaaaccccggu.....	1	0	seq
.....Aaccgggguuuuuguucgucgca.....	1	1	seq
.....accgggguuuuuguucgucg.....	1	0	seq
.....accgggguuuuuguucgAucgca.....	1	1	seq
.....accgggguuuuuguucgucgcaa.....	17	0	seq
.....Cccgggguuuuuguucgucgcaa.....	2	1	seq
.....Gccgggguuuuuguucgucgcaa.....	3	1	seq
.....ccgggguuuuuguucgucg.....	2	0	seq
.....Ucgggguuuuuguucgucg.....	1	1	seq
.....Gcgggguuuuuguucgucg.....	1	1	seq
.....ccgggguuuuuguucgucgcaa.....	1	0	seq
.....ccgggguuuuuguucguAagcaa.....	1	1	seq
.....cggggguuuuuguucgucgcaa.....	3	0	seq
.....Ugggguuuuuguucgucgcaa.....	3	1	seq
.....Aggggguuuuuguucgucgcaa.....	1	1	seq

uugcagacgaacaaaaccccgguauagucuaacacccgggguuuugnuucgucugcaa			
.....Ugggguuuugnuucgucugcaa	5	1	seq
.....ggggguuuugnuucgucugcaa	1	0	seq
.....Agggguuuugnuucgucugcaa	1	1	seq
.....ggggguuuugnuucgucugca.	1	0	seq
.....ggggguuuugnuucAucugcaa	1	1	seq