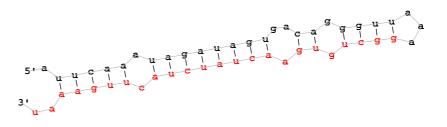
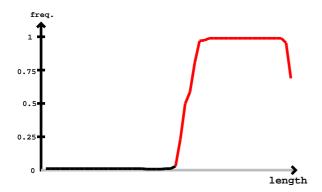
miRBase precursor : chromosome_5_16995

Total read count : 263 chromosome_5_16995 read x26f7t remaining reads : 6





chromosome_5_16995

- auucaaauagauagugacaggguuaaaggcugugaacuaucuacuugaaau	-3' exp		
.(((((.(((((((()))).)).))))))))	reads	mm	sample
auucaaauagauagugacag	3	0	seq
Uguuaaaggcugugaacuaucuacuugaaa.	2	1	seq
aaaggcugugaacuaucuacuugaaa.	1	0	seq
aggcugugaacuaucuacuugaaa.	1	0	seq
Uggcugugaacuaucuacuugaaa.	1	1	seq
Uggcugugaacuaucuacuugaaau	3	1	seq
ggcugugaacuaucuacuugaCa.	1	1	seq
ggcugugaacuaucuacuugaaa.	2	0	seq
Agcugugaacuaucuacuugaaa.	13	1	seq
ggcugugGacuaucuacuugaaau	8	1	seq
Agcugugaacuaucuacuugaaau	7	1	seq
ggcugugaacuaucCacuugaaau	1	1	seq
ggcugugaacuaucuacuugaaau	20	0	seq
Ugcugugaacuaucuacuugaaau	1	1	seq
gcugugGacuaucuacuugaa	1	1	seq
Ucugugaacuaucuacuugaaa.	2	1	seq
gcugugaacuaucuacuugaaa.	16	0	seq
gcugugaacuaucCacuugaaa.	1	1	seq
Acugugaacuaucuacuugaaa.	5	1	seq
Acugugaacuaucuacuugaaau	11	1	seq
gcugugaacuaucuacuugaaau	17	0	seq
Ucugugaacuaucuacuugaaau	3	1	seq
gcugugGacuaucuacuugaaau	12	1	seq
gcugugaacCaucuacuugaaau	1	1	seq
gcugugaacuaucuacuAgaaau	1	1	seq
cugugaacuaucuacuugaa	2	0	seq
cugugaacuaucuacuugaaa.	4	0	seq
Augugaacuaucuacuugaaa.	2	1	seq
cugugaacuaucuGcuugaaau	1	1	seq
cugugGacuaucuacuugaaau	9	1	seq
cugugaGcuaucuacuugaaau	1	1	seq
cugugaacuaucuacuugaaau	4	0	seq
ugugaacuaucuacuugaa	1	0	seq
ugugaacuaucuacuugaaa.	13	0	seq
ugugaacuaucuacuCgaaa.	1	1	seq

chromosome_5_16995

2	1	seq
24	1	seq
1	1	seq
13	0	seq
1	1	seq
1	1	seq
1	1	seq
2	1	seq
3	0	seq
3	0	seq
24	1	seq
8	0	seq
2	1	seq
1	1	seq
1	1	seq
3	1	seq
1	0	seq
	24 1 13 1 1 1 2 3 3 24 8 2 1 1 3	24 1 1 1 13 0 1 1 1 1 1 1 1 1 2 1 3 0 3 0 24 1 8 0 2 1 1 1 1 1 1 1 3 1