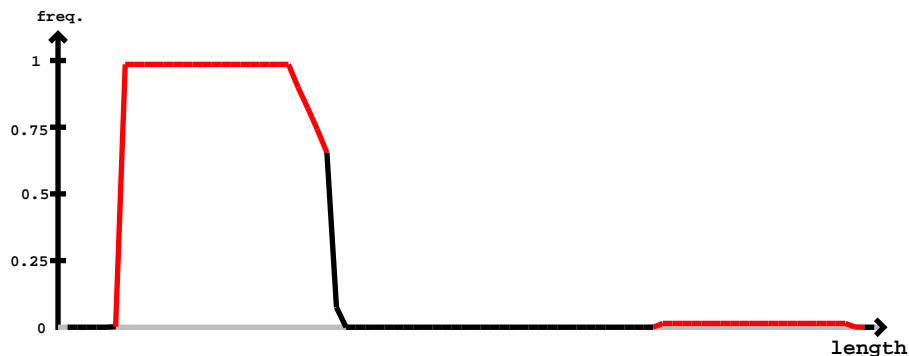
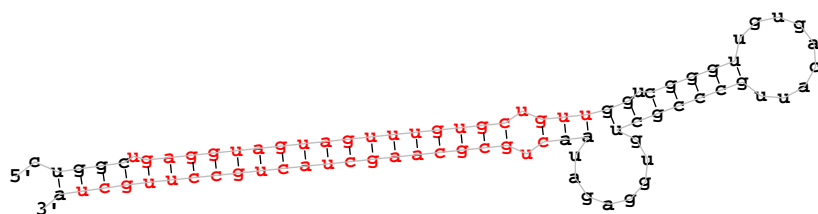


miRBase precursor : hsa-let-7i
Total read count : 2768
hsa-let-7i-5p read count : 2728
hsa-let-7i-3p read count : 40
remaining reads : 0



hsa-let-7i-5p

hsa-let-7i-3p

5'	reads	mm	sample
cuggcugagguaguaguuugugcuguuggucggguugugacauugcccgcugaggagauaacugcgcaagcuacugccuugcua	2768	0	seq
.....cugagguaguaguuugugcugu.....	1	0	seq
.....Uugagguaguaguuugugcugu.....	2	1	seq
.....Uugagguaguaguuugugcuguuu.....	2	1	seq
.....Augagguaguaguuugugcuguuu.....	1	1	seq
.....ugagguaguaguuugugG.....	3	1	seq
.....ugaggCaguaguuugugc.....	1	1	seq
.....ugagguaguaguuugugc.....	234	0	seq
.....ugagguaguaguuugugcu.....	209	0	seq
.....ugagguaguaguuuAugcu.....	2	1	seq
.....ugagguaguaguuugugcG.....	1	1	seq
.....ugagguuACuaguuugugcu.....	1	1	seq
.....Agagguaguaguuugugcu.....	1	1	seq
.....ugaggGaguaguuugugcug.....	1	1	seq
.....ugagguaguaguuugCgucug.....	1	1	seq
.....ugagguaguaguuuCGugcug.....	1	1	seq
.....ugagguaguaguCuugucug.....	1	1	seq
.....ugagguuACuaguuugugcug.....	1	1	seq
.....ugagguaguaguuugugcug.....	217	0	seq
.....ugagguaguagAuugugcugu.....	1	1	seq
.....ugagguaguaguuugugcugC.....	2	1	seq
.....ugagguaguaguuugugcugu.....	234	0	seq
.....ugGgguaguaguuugugcugu.....	1	1	seq
.....ugagguaguaguuugugcuguA.....	6	1	seq
.....ugagguaguaguuugGuguu.....	3	1	seq
.....ugagguagCaguugugcuguuu.....	1	1	seq
.....ugagguaguaguCuugucuguuu.....	2	1	seq
.....ugagguaguCuugugugcuguuu.....	1	1	seq
.....ugagguaguaguuugugcuAuu.....	1	1	seq
.....ugaggGaguaguuugugcuguuu.....	1	1	seq
.....ugagguaguaguuugCgucuguuu.....	1	1	seq
.....ugagguaguaguuuUugcuguuu.....	1	1	seq
.....ugagguaguaguCuugucuguuu.....	1	1	seq
.....ugagguaguuaUuuugugcuguuu.....	1	1	seq
.....ugagguaguuaAuuugugcuguuu.....	1	1	seq

hsa-let-7i-5p

hsa-let-7i-3p

cuggcugagguaguaguuugugcuguugggucggguugugacauugcccgcuguggagauaacugcgcaagcuacugccuugcua

....ugagguacuaguuugugcuguu.....	1	1	seq
....ugagguaguaguuugugcuguu.....	1578	0	seq
....Cgagguaguaguuugugcuguu.....	2	1	seq
....Agagguaguaguuugugcuguu.....	2	1	seq
....ugagguaguGguuuugugcuguu.....	1	1	seq
....ugagguaguaguuugugcuguuU.....	30	1	seq
....ugagguaguaguuugugcuguuA.....	173	1	seq
....ugagguaguaguuugugcuguug.....	3	0	seq
.....Ccugcgcaagcuacugccuug...	1	1	seq
.....cugcgcaagcuacugccuug...	36	0	seq
.....cugcgcaagcuacugccuugc..	3	0	seq