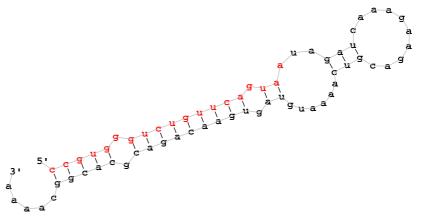
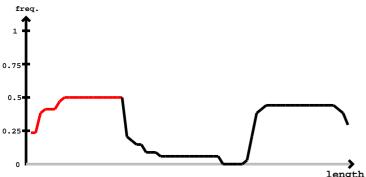
miRBase precursor : chromosome\_14\_72515

Total read count : 34 chromosome\_14\_72515 read **&6**unt remaining reads : 18





## chromosome\_14\_72515

5'-	ccgugggucuguucaguaauagaucaaagaagacgucaaaauguagugaacagacgcacggcaaaa	-3'	exp	
	(((((((((((((((((((((((((((((((((((((((	reads	s mm	sample
	ccgugggucuguucaguaa	1	0	seq
	ccgugggucuguucaguUa	1	1	seq
	ccgugCgucuguucaguaa	5	1	seq
	ccgugggucuguucaguaau	1	0	seq
	.cgugCgucuguucaguaa	5	1	seq
	gugCgucuguucaguaaua	1	1	seq
	Cgucuguucaguaauaga	2	1	seq
	gucuguucaguaauagauca	1	0	seq
	uagaucaCagaagacgucaa	2	1	seq
	aCugaacagacgcacggca	1	1	seq
	Cugaacagacgcacggcaa	1	1	seq
	Cugaacagacgcacggcaaa.	3	1	seq
	gugaacagacgcacggcaaaa	2	0	seq
	ugaacagacgcacggcaaaa	5	0	seq
	ugaacagacAcacggcaaaa	1	1	seq
	gaacagacgcacggcaaaG	1	1	seq
	Hacagacgcaaaa	1	1	sea