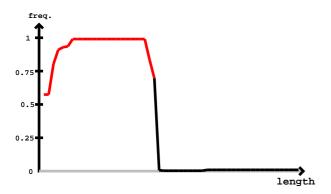
miRBase precursor : chromosome\_6\_24347
Total read count : 310

Total read count : 310 chromosome\_6\_24347 read x306t remaining reads : 4





## chromosome\_6\_24347

-	cgaucacgucuagaacacccaaagauugaaucuuucaguguucuagacguaau	-3' exp		
	(((((((((((((((((((((((((((((((((((	reads	mm	sample
	cgaucacguUuagaacaccca	1	1	seq
	Ugaucacgucuagaacaccca	20	1	seq
	cgaucacgucuagaacaccca	6	0	seq
	Ugaucacgucuagaacacccaa	22	1	seq
	Ggaucacgucuagaacacccaa	3	1	seq
	cUaucacgucuagaacacccaa	1	1	seq
	cgaucacgucuagaacacccaa	8	0	seq
	cgaucacgucuGgaacacccaaa	1	1	seq
	cgaGcacgucuagaacacccaaa	1	1	seq
	Ugaucacgucuagaacacccaaa	50	1	seq
	cgaucacgucuagaacacccaaa	53	0	seq
	cgaucacgGcuagaacacccaaa	1	1	seq
	Agaucacgucuagaacacccaaa	5	1	seq
	cgaucacgucGagaacacccaaa	1	1	seq
	Ggaucacgucuagaacacccaaa	3	1	seq
	Agaucacgucuagaacacccaaag	1	1	seq
	cgaucacgucuagaacacccaaagauugaa	1	0	seq
	.Aaucacgucuagaacaccca	3	1	seq
	.Uaucacgucuagaacaccca	2	1	seq
	.gaucacgucuagaacaccca	8	0	seq
	.gaucacgCcuagaacacccaa	1	1	seq
	.gaucacgucuagaacacccaa	4	0	seq
	.gaucacgCcuagaacacccaaa	3	1	seq
	.gaCcacgucuagaacacccaaa	2	1	seq
	.gaucacgucuagaacacccaaa	39	0	seq
	.gaucacgucCagaacacccaaa	1	1	seq
	.Uaucacgucuagaacacccaaa	1	1	seq
	.gaucacgucuagaacacccUaa	1	1	seq
	.gaucacgucuagaacacUcaaa	1	1	seq
	.gaucacgucuagaacacccaGa	2	1	seq
	.Aaucacgucuagaacacccaaa	3	1	seq
	Uucacgucuagaacaccca	1	1	seq
	Gucacgucuagaacaccca	1	1	seq
	aucacgucuagaacaccca	5	0	seq
	aucacgucuagaacacccaa	2	0	seq

## cgaucacgucuagaacacccaaagauugaaucuuucaguguucuagacguaau

aucacgCcuagaacacccaaa	1	1	seq
Gucacgucuagaacacccaaa	4	1	seq
aAcacgucuagaacacccaaa	1	1	seq
aucacgucuagaacacccaaa	18	0	seq
ucacgucuagaacaccca	1	0	seq
ucacgucuagaacGcccaa	1	1	seq
ucacgucuCgaacacccaaa	1	1	seq
ucacgucuagaacacccaaa	3	0	seq
Uacgucuagaacacccaa	1	1	seq
Uacgucuagaacacccaaa	1	1	seq
Gcgucuagaacacccaaa	1	1	seq
Ucgucuagaacacccaaa	1	1	seq
acgucuagaacacccaaa	15	0	seq
ucuuucaguguucuagacguaau	1	0	seq
uucaguguucuagacguaau	2	0	seq