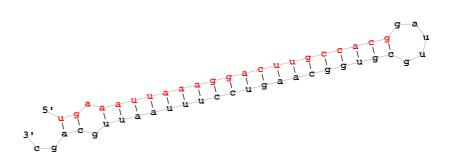
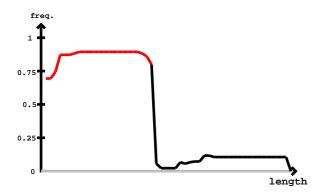
miRBase precursor : chromosome_2_3806

Total read count : 180 chromosome_2_3806 read co**\u00ednf7** remaining reads : 23





chromosome_2_3806

-	ugaaauuaaaggacuugccacggauugcguggcaaguccuuuaauugcagc	-3'	exp	
	((.(((((((((((((((((())))))))))))))	reads	mm	sample
	ugaaauuaaaggacuugcc	2	0	seq
	ugaaauuaaaggacuugcca	4	0	seq
	ugaaauuaaaggacuugccac	8	0	seq
	Cgaaauuaaaggacuugccacg	1	1	seq
	Agaaauuaaaggacuugccacg	1	1	seq
	ugaaauuaaaggacuugccUcg	2	1	seq
	ugaaauuaaagUacuugccacg	1	1	seq
	ugaaauuaaaggacuugccaAg	1	1	seq
	ugaaauCaaaggacuugccacg	1	1	seq
	ugaaauuaaaggacuugccacA	29	1	seq
	ugaaauuaGaggacuugccacg	2	1	seq
	uAaaauuaaaggacuugccacg	1	1	seq
	ugaaauuaaaggacuCgccacg	1	1	seq
	ugaaauuaaaggacuGgccacg	1	1	seq
	ugaaauuaaaggacuugccacg	61	0	seq
	ugaaauuaaaggacuugccacU	2	1	seq
	ugaaauGaaaggacuugccacg	1	1	seq
	ugaaauuaaaggacuugccacgA	2	1	seq
	ugaaauuaaaggacuugccacgU	4	1	seq
	.gaaauuaaaggacuugccac	2	0	seq
	.gaaauuaaaggacuugccacg	5	0	seq
	.gaaauuaaaggacuugccacA	2	1	seq
	aaauuaaaggacuugccac	1	0	seq
	aaauuaaaggacuugccacg	19	0	seq
	aaauuaaaggacuugccacU	1	1	seq
	aaauuaaaggacuugccacA	2	1	seq
	uuaaaggacuugccacggauugU	2	1	seq
	uaaaggacuugccacggauugcguggca	2	0	seq
	cguggcaaguccuuuaauuUcag.	8	1	seq
	uggcaaguccuuuaauuUcag.	2	1	seq
	ggcaaguccuuuaauuUcag.	1	1	seq
	caaguccuuuaauuUcag.	8	1	seq