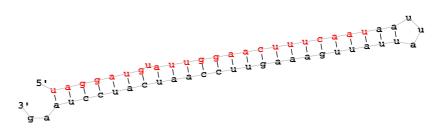
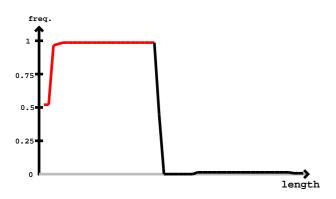
miRBase precursor : chromosome_12_57748

Total read count : 292 chromosome_12_57748 read **288**nt remaining reads : 4





chromosome_12_57748

	CHI OMOSOME_12_5//40				
5'	- uaggauguauuggaacuuucaauaauuauugaaaguuccaaucauccuaag	-3'	exp		
	(((((((((((((((((((((((((((((((((((((((reads	s m	m.	sample
	uaggauguauuggaacuuucaGu	1	1		seq
	uaggauguauuggaacuuucaau	70	0		seq
	uaggauguauuggaGcuuucaau	3	1		seq
	uaggauguaCuggaacuuucaau	2	1		seq
	uaggauguauuggCacuuucaau	1	1		seq
	uaggauguauuggaacuuucaaC	7	1		seq
	uaggauguauuggaacCuucaau	1	1		seq
	uaggauguauuggaacGuucaau	1	1		seq
	uGggauguauuggaacuuucaaua	2	1		seq
	uaggauguauuggaacuuucaaCa	1	1		seq
	uaggauguauuggGacuuucaaua	1	1		seq
	uaggGuguauuggaacuuucaaua	1	1		seq
	uaggauguauugAaacuuucaaua	1	1		seq
	uaggauguauuggaacuuucaaua	57	0		seq
	uaggauguauuggaacuuucaaGa	1	1		seq
	uaggauguauuggaacuuucaUua	1	1		seq
	uaggauguauuggaacAuucaaua	1	1		seq
	.aggauguauuggaacCuucaau	1	1		seq
	.aggauguauuggaacuuucaaC	4	1		seq
	.aggauguauuggaacuuucaau	60	0		seq
	.aggauguauuggaacuuucaGu	1	1		seq
	.aggauguauuggaGcuuucaaua	1	1		seq
	.aggauguauuggaacuuucaaua	58	0		seq
	.aggauguauuggaacuuucGaua	1	1		seq
	.aggauguauuggaacuuucaauC	1	1		seq
	.aggauguauuggaacuuucaaGa	1	1		seq
	.aggauguauCggaacuuucaaua	1	1		seq
	.aAgauguauuggaacuuucaaua	1	1		seq
	ggauguauuggaacuuucaau	3	0		seq
	gauguauuggaacuuucaau	3	0		seq
	uugaaaguuccaauccC	1	1		seq
	uugaaaguuccaaucauccu	1	0		seq
	uugaaaguuccaaucauccuaag	2	0		seq