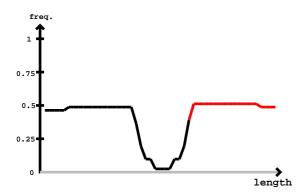
miRBase precursor : chromosome\_1\_1120

Total read count : 41 chromosome\_1\_1120 read colifit remaining reads : 24





## chromosome\_1\_1120

5'- uugccuaugguggauucuggcaauauugccagaauccaccauaggcaa	-3'	exp	
.(((((((((((((((())))))))))))))))))	reads	mm	sample
uugccuaugguggauucu	4	0	seq
uugccuaugguggaGucu	1	1	seq
uugccuaugguggauucuA	2	1	seq
uugccuauggugAauucug	1	1	seq
uugccuaugguggauucug	4	0	seq
uugccuaugguggauucugg	4	0	seq
uugccuaugguggauucuggca	3	0	seq
cuaugguggauucuggcaau	1	0	seq
auugccagaauccaccauag	1	0	seq
ugccagaauccaccauaggcaa	3	0	seq
ccagaauccaccauaggcaa	3	0	seq
ccagaauccaccauaggcaC	1	1	seq
cagaauccaccauaggcaa	4	0	seq
	3	1	seq
cagaauUcaccauaggcaa	1	1	seq
agaauccaccauaggcaa	5	0	seq