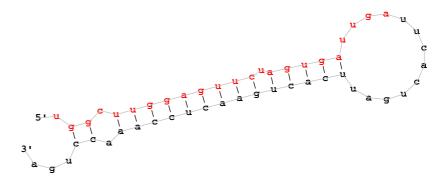
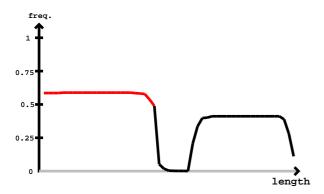
miRBase precursor : chromosome\_4\_15207

Total read count : 1269 chromosome\_4\_15207 read v 46t remaining reads : 523





## chromosome\_4\_15207

٠-	uggcuuggaguucuagugauugauucacugauucacugaacuccaaaccuga	-3' e	хp	
	.((.(((((((((((())))))))))))))	reads	mm	sample
	uggcuuggaguucuagug	4	0	seq
	uggcuuggaguucuaguU	1	1	seq
	uggcuuggaguucuaguga	1	0	seq
	uggcuuggaguucuagGga	2	1	seq
	uggcuuggaguucuagugau	5	0	seq
	uggcuuggaguucuagugauC	2	1	seq
	uggcuuggaguuAuagugauu	1	1	seq
	uggcuuggaguucuagugCuu	2	1	seq
	uggcuuggaguucuagugauu	46	0	seq
	Gggcuuggaguucuagugauu	1	1	seq
	Aggcuuggaguucuagugauug	1	1	seq
	Gggcuuggaguucuagugauug	1	1	seq
	ugCcuuggaguucuagugauug	1	1	seq
	uggcuuggaguucuaguCauug	1	1	seq
	uggcuuggaguucuagugauuU	1	1	seq
	uggcuuggaguucuagugauuA	7	1	seq
	uggcuuggaguucuagugauug	50	0	seq
	uggcuuggaguuAuagugauuga	1	1	seq
	uggcAuggaguucuagugauuga	1	1	seq
	ugAcuuggaguucuagugauuga	1	1	seq
	uggUuuggaguucuagugauuga	1	1	seq
	uggcuuggaguucuaguAauuga	1	1	seq
	uggcuuggaguucuagugauCga	1	1	seq
	uggcuCggaguucuagugauuga	2	1	seq
	uggcuuggaguCcuagugauuga	3	1	seq
	uggcuuggaguucuagugauugU	17	1	seq
	uggcuuAgaguucuagugauuga	1	1	seq
	uggcuuggaguucuagugaCuga	1	1	seq
	Aggcuuggaguucuagugauuga	3	1	seq
	uggcuuggGguucuagugauuga	2	1	seq
	uUgcuuggaguucuagugauuga	1	1	seq
	uggcCuggaguucuagugauuga	2	1	seq
	uggcuuggaguucuagugauGga	11	1	seq
	uggcuuggCguucuagugauuga	1	1	seq
	ugUcuuggaguucuagugauuga	1	1	seq

uggcuuggaguucuagugauugauucacugauucacugaacuccaaaccuga			
uggcuuggaguucuagugGuuga	2	1	seq
uggcuuggaguucuagugauuga	481	0	seq
uggcuuggaAuucuagugauuga	2	1	seq
Cggcuuggaguucuagugauuga	1	1	seq
uggcuuggaguAcuagugauuga	1	1	seq
uggcuuggaguucuGgugauuga	5	1	seq
uggcuuggaguucuagAgauuga	2	1	seq
uggcGuggaguucuagugauuga	1	1	seq
uggcuuggaguucuagugauugG	9	1	seq
uggcuuggGguucuagugauugau	1	1	seq
uggcuuggaguucuagugau	38	0	seq
uggcuuggaguucuagugauugaC	3	1	seq
uggcuuggaguucuagugauugaA	1	1	seq
uggcuuggaguucuagugauugauA	2	1	seq
uggcuuggaguucuagugaAu	3	1	seq
uggcuuggaguucuagugauu	8	0	seq
uggcuuggaguucuaCugauugauu	1	1	seq
			_
uggcuuggaguucuagugauugaAuc	2	1	seq
ugAcuuggaguucuagugauuca	1	1	seq
uggcuuggaguucuagugauuUauucacu	1	1	seq
cuuggaguucuagugauugauu	4	0	seq
uucacugauucacugaacuccaaaccuga	1	0	seq
auucacugaacuccaaacc	4	0	seq
auucacugaacuccaaacU	3	1	seq
auucacugaacuccaaacA	14	1	seq
auucacugaacuccaaaccA	11	1	seq
auucacugaacuccaaaccu	51	0	seq
auucacugaacuccaaaccC	9	1	seq
aCucacugaacuccaaaccu	1	1	seq
auucacCgaacuccaaaccu	1	1	seq
auucacugaacuccaaaccug.	53	0	seq
Guucacugaacuccaaaccug.	3	1	seq
auucacugaacuccaaaccuU.	2	1	seq
auucacCgaacuccaaaccug.	3	1	seq
	2		_
auucacugaacuccaaaccuC.		1	seq
auucacugaacuccaaaccuA.	7	1	seq
auCcacugaacuccaaaccug.	2	1	seq
auucGcugaacuccaaaccuga	1	1	seq
auucacugaacuccaaaccGga	2	1	seq
auucacCgaacuccaaaccuga	1	1	seq
auucacugaacuccaaaccuga	80	0	seq
auucacugaacuccaaaccugU	7	1	seq
auucacGgaacuccaaaccuga	1	1	seq
auucacugaacuccaaaccugG	2	1	seq
auucacugaacuccaaaccugC	1	1	seq
auucacugaacuccaaaccuUa	1	1	seq
auucacugGacuccaaaccuga	1	1	seq
uucacugaacuccaaacA	1	1	seq
uucacugaacuccaaacU	4	1	seq
uucacugaacuccaaacG	3	1	seq
uucacugaacuccaaacUu	2	1	seq
	5	1	
uucacugaacuccaaaccC			seq
uGcacugaacuccaaaccu	1	1	seq
uucacugaacuccaaaccu	38	0	seq
uucacugaacuccaaaccA	5	1	seq
uucacAgaacuccaaaccug.	1	1	seq
uAcacugaacuccaaaccug.	1	1	seq
uucacuCaacuccaaaccug.	1	1	seq
uucacugaacuccaaaccug.	70	0	seq
uucacugaacuccaaaccuU.	2	1	seq
Cucacugaacuccaaaccug.	1	1	seq
uucacugaGcuccaaaccug.	1	1	seq
uucacugaacuccaaaccuA.	8	1	seq
uucacugaacuccaaaccuga	16	0	seq
uucacugaacuccaaGccuga	2	1	seq
uucacugaacuccaaaccugU	1	1	seq
ucacugaacuccaaaccA	1	1	seq
ucacugaacuccaaaccu	12	0	seq
ucGcugaacuccaaaccug.	1	1	seq
ucacugaacuccaaaccug.	35	0	seq
		•	204

uggcuuggaguucuagugauugauucacugauucacugaacuccaaaccuga			
ucacugaUcuccaaaccug.	1	1	seq
ucacCgaacuccaaaccug.	1	1	seq
ucacugGacuccaaaccug.	1	1	seq
ucaUugaacuccaaaccug.	2	1	seq
ucacugaacuccaaaccuA.	11	1	seq
ucacugaacuccaaaccuU.	2	1	seq
ucacugaacuccaaaccugU	4	1	seq
ucacugaacuccaaaccuga	7	0	seq
cacugaacuccaaaccug.	2	0	seq
	_	_	

seq

seq

seq

seq

1

.....cacugaacuccaaaccug. 2 .....cacugaacuccaaaccuA.

.....acugaacuccaaaccugC 3

.....acugaacuccaaaccuga 9

.....acugaacuccaaaccGga 1 1