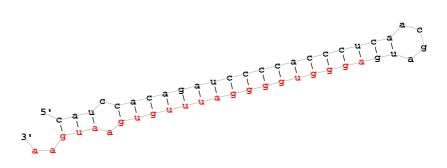
miRBase precursor : chromosome\_12\_59393

Total read count : 25 chromosome\_12\_59393 read **25**unt remaining reads : 0





## chromosome\_12\_59393

5'- cauccacagauccccacccucaacgaugagggugggggauuugugaaugaa	-3' ex	p	
(((.(((((((((((((((((((((((((((((((((((	reads	mm	sample
Agaggguggggauuugugaau	2	1	seq
ugaggguggggauuugugaaug	2	0	seq
	1	1	seq
ugaggguggggauuugugaauga.	2	0	seq
	1	1	seq
ugaggguggggauuugugaaugaa	5	0	seq
gaggguggggauuugugaauga.	1	0	seq
gaggguggggauuugugaaugaa	2	0	seq
Uggguggggauuugugaaug	2	1	seq
aggguggggauuugugaaugaa	6	0	seq
	1	1	sea