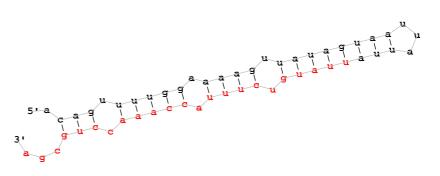
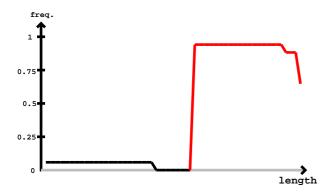
miRBase precursor : chromosome\_12\_58629

Total read count : 17 chromosome\_12\_58629 read **t6**unt remaining reads : 1





## chromosome\_12\_58629

5'- acaguuuuggaaaaguuauaguaauuauuaugucuuuaccaaaccugcga	-3'	exp	
.(((.((((((((((((((((())))))))))))))	reads	mm.	sample
acaguuuuggaaaaguuauagu	1	0	seq
uuaugucuuuaccaaacGu	1	1	seq
uuaugucuuuaccaaaccugcg.	4	0	seq
uuaugucuuuaccaaaccugcga	11	0	seq