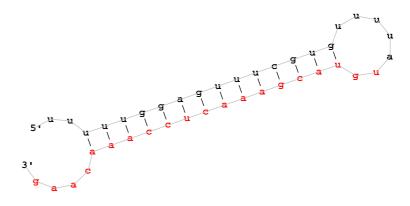
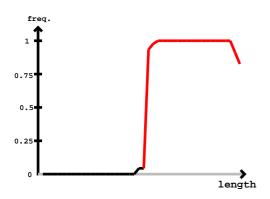
miRBase precursor : chromosome\_4\_12905

Total read count : 46 chromosome\_4\_12905 read x\(\phi\)6nt remaining reads : 0





## chromosome\_4\_12905

5'- uuuuuggaguuucguguuuu <mark>auguacgaaacuccaaacaag</mark>	-3' exp		
(((((((((((())))))))))))))	reads	mm	sample
uauguacgaaacuccaaaca	2	0	seq
uguacgaaGcuccaaaca	1	1	seq
uguacgaaaAuccaaaca	1	1	seq
uguacgaaacuccaaacaa.	4	0	seq
uguacgaGacuccaaacaag	1	1	seq
uguacgaaacuccaaacaGg	1	1	seq
uguacgaaacuccaaacaaA	1	1	seq
uguacgaaacuccaaacaag	29	0	seq
uguacgGaacuccaaacaag	1	1	seq
uguacgaaGcuccaaacaag	1	1	seq
uguacgaaacuccaaacaaC	1	1	seq
guacgaaacuccaaacaag	2	0	seq
uacgaaacuccaaacaag	1	0	seq