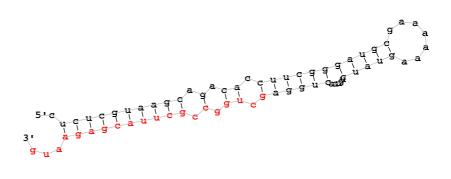
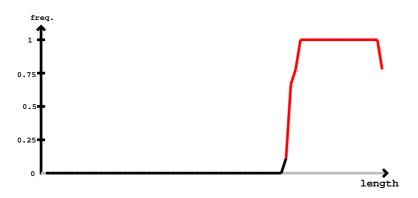
miRBase precursor : chromosome_2_4922

Total read count : 9 chromosome_2_4922 read co\(\text{9nt} \) remaining reads : 0





chromosome_2_4922

	exp	-3'	5'- cucucguaagcagacaccuucgggaugcgaaaaaaguaucuguaacugga <mark>gcuggccgcuuacgagaaug</mark>	5
sample	mm	reads	.((((((((((((((((((((((((((((((((((((((
seq	1	1	Ugcuggccgcuuacgagaaug	
seq	1	1	Acuggccgcuuacgagaau.	
seq	1	4	Acuggccgcuuacgagaaug	
seq	1	1	Auggccgcuuacgagaau.	
seq	0	1	uggccgcuuacgagaaug	
seq	1	1	uggccgcuuacgagaauA	