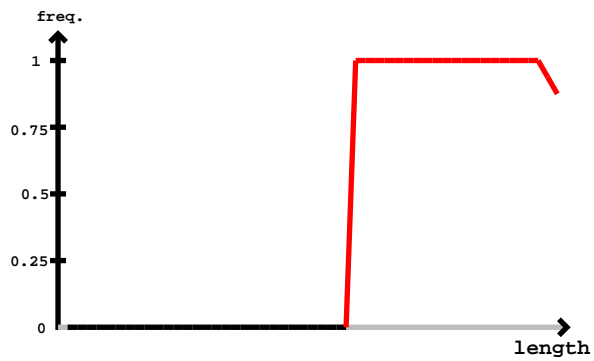
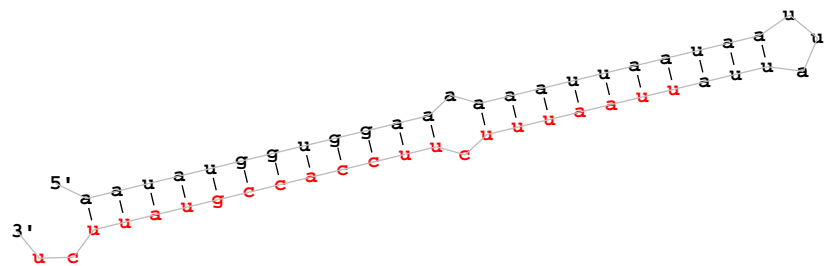


miRBase precursor : chromosome_1_2986
 Total read count : 16
 chromosome_1_2986 read count : 16
 remaining reads : 0



chromosome_1_2986			
5'-	aaauagguggaaaaaaauaauuuuuuuuaauuuuuccaccguauuc	-3'	exp
	((((((((.....(((.....)))))))))).)))))))).)..	reads	mm
uuuuuuuuccaccguauC..	1	1
uuuuuuuuccaccguauuc.	1	0
uuuuuuuuccaccguauucA	5	1
uuuuuuuuccaccguauucC	2	1
uuuuuuuuccaccguauuc	7	0
		seq	seq