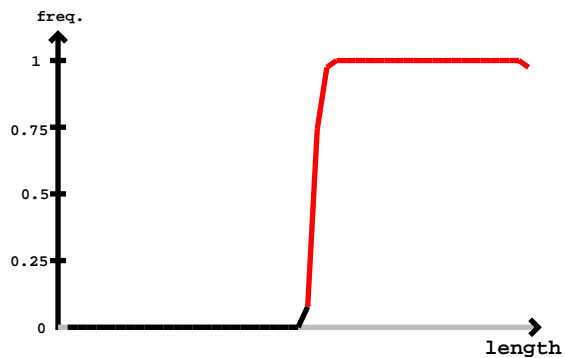
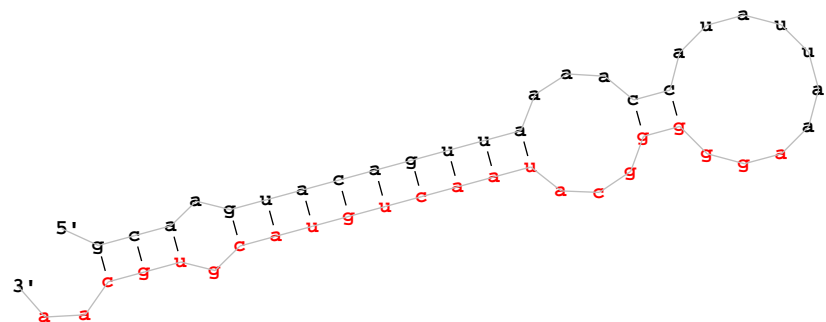


miRBase precursor : chromosome_11_54784
 Total read count : 39
 chromosome_11_54784 read count : 39
 remaining reads : 0



chromosome_11_54784			
5'-	gcaaguacaguuaaaaccauauuaaagggggc <u>auaacuguacgugcaa</u>	-3'	exp
	(((((((((((((((.....)).....)))))))))))).))..	reads	mm sample
.....	aaUggggc <u>auaacuguacgugcaa</u>	1	1 seq
.....	aagggggc <u>auaacuguacgugcaa</u>	1	0 seq
.....	Gagggggc <u>auaacuguacgugcaa</u>	1	1 seq
.....	Uggggggc <u>auaacuguacgugcaa</u>	1	1 seq
.....	Gggggggc <u>auaacuguacgugcaa</u>	4	1 seq
.....	aggggggc <u>auaacuguacgugcaa</u>	20	0 seq
.....	Cggggggc <u>auaacuguacgugcaa</u>	1	1 seq
.....	gggggggc <u>auaacuguacgugca</u> .	1	0 seq
.....	Ugggggc <u>auaacuguacgugcaa</u>	3	1 seq
.....	ggggggc <u>auaacuguacgugcaa</u>	3	0 seq
.....	Agggggc <u>auaacuguacgugcaa</u>	2	1 seq
.....	Uggggc <u>auaacuguacgugcaa</u>	1	1 seq