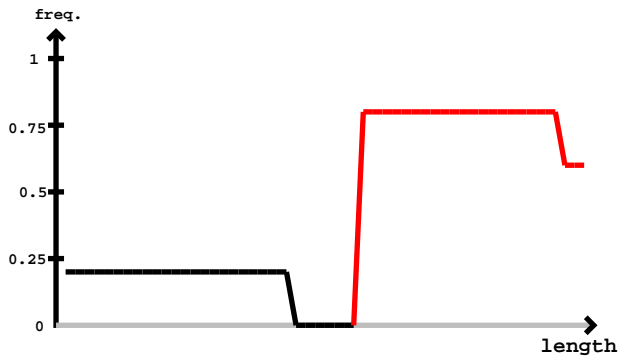
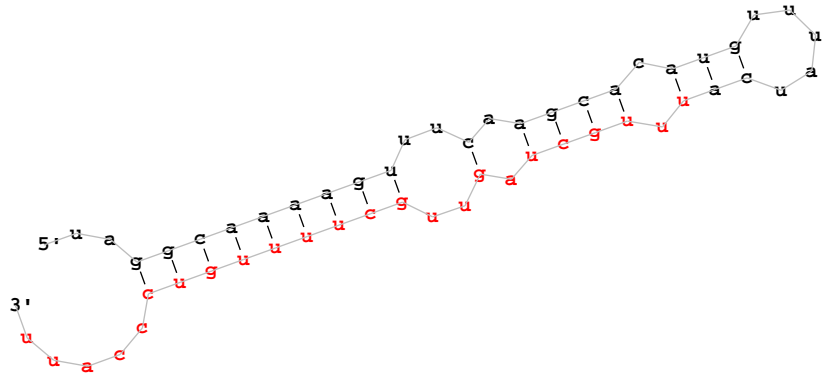


miRBase precursor : chromosome_12_56385

Total read count : 5

chromosome_12_56385 read count

remaining reads : 1



chromosome_12_56385			
5' -		-3'	exp
uaggcaaaaguuucaagcacauguuuaucauuugcuaguugcuuuuugucccau	reads	mm	sample
..((((((((((..((((((((.....)))))).)))).)))).)).....	1	0	seq
uaggcaaaaguuucaagcacaug.....uuugcuaguugcuuuuuguccc...	1	0	seq
.....uuugcuaguugcuuuuugucccau	1	1	seq
.....uuugcuaguugcuuuuugucccau	2	0	seq