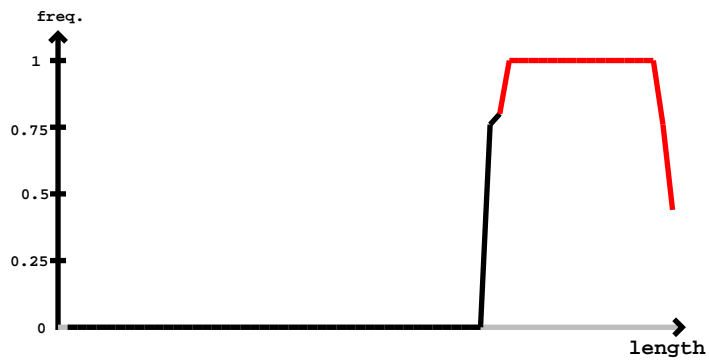
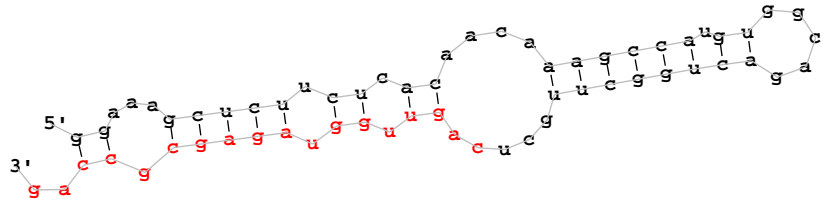


miRBase precursor : chromosome_12_57483
 Total read count : 25
 chromosome_12_57483 read 25
 remaining reads : 0



chromosome_12_57483			
5'-		-3'	exp
ggaaagcucucucacacaacaagccaugggcagacuggcugcucag <u>uugguagagcgccag</u>	reads	mm	sample
((...(((((((((...(((((((...)))))))))...)).)).))))))..	1	1	seq
.....cucaguugguagagcgGc..	1	1	seq
.....cucaguuggGagagcgcc..	1	1	seq
.....cucaguugguagagcgC..	1	1	seq
.....cucaguugguagagcgUc..	2	1	seq
.....cucaguCgguagagcgcc..	1	1	seq
.....cucaguugguagagcgGca..	1	1	seq
.....cucaguugguagagcgccU..	3	1	seq
.....cucaguugguagagcgUca..	1	1	seq
.....cucaguugguagagcgcca..	2	0	seq
.....cucaguugguagagcgccUg	6	1	seq
.....ucaguugguagagcgUca..	1	1	seq
.....caguugguagagcgGcag	3	1	seq
.....caguugguagagcgUcag	2	1	seq