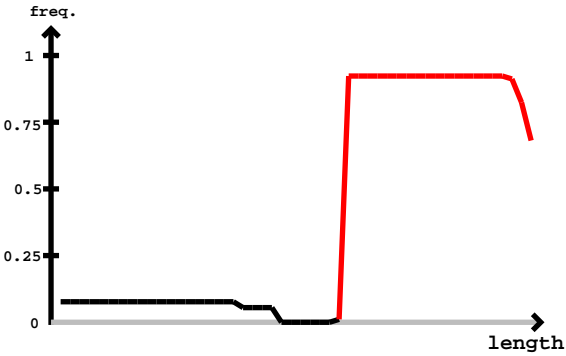
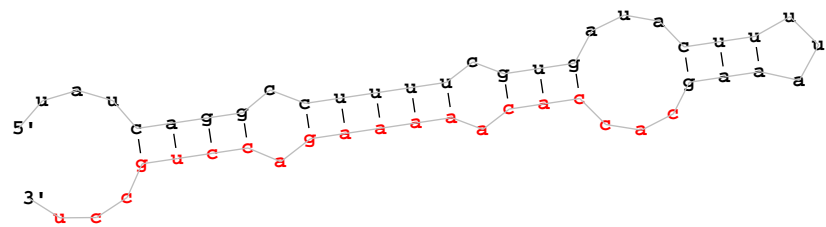


miRBase precursor : chromosome\_11\_50389  
Total read count : 91  
chromosome\_11\_50389 read count : 86  
remaining reads : 7



chromosome_11_50389				
5'-	u-a-u-c-a-g-g-c-u-u-u-u-c-g-a-u-a-c-u-u-a-a-a-g-c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	reads	exp	sample
...	(((((((((((.....)))))).....)))))).....	1	0	seq
u-a-u-c-a-g-g-c-u-u-u-u-c-g-a-u-a-c-u-u-a-a-a-g-c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	u-a-u-c-a-g-g-c-u-u-u-u-c-g-a-u-a-c-u-u-a-a-a-g-c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	1	1	seq
u-a-u-c-a-g-g-c-u-u-u-u-c-g-a-u-a-c-u-u-a-a-a-g-c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	u-a-u-c-a-g-g-c-u-u-u-u-c-g-a-u-a-c-u-u-a-a-a-g-c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	3	0	seq
u-a-u-c-a-g-g-c-u-u-u-u-c-g-a-u-a-c-u-u-a-a-a-g-c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	u-a-u-c-a-g-g-c-u-u-u-u-c-g-a-u-a-c-u-u-a-a-a-g-c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	2	1	seq
.....g-c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....g-c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	1	0	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	1	1	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	1	1	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	2	1	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	4	0	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	1	1	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	1	1	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	1	1	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	3	1	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	4	0	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	4	1	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	6	1	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	1	1	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	7	1	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	1	1	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	1	1	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	45	0	seq
.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	.....c-a-c-a-c-a-a-a-a-g-a-c-c-u-g-c-c-3'	1	1	seq