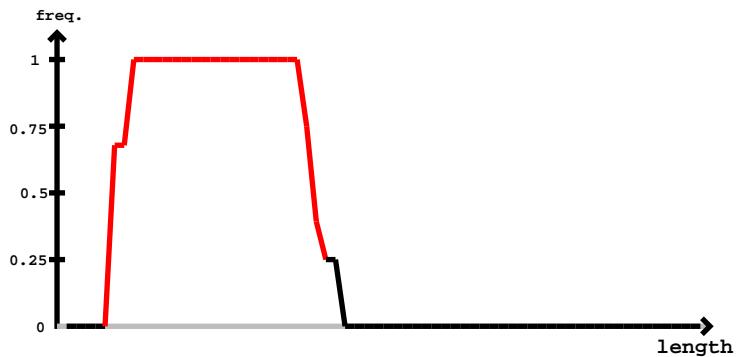
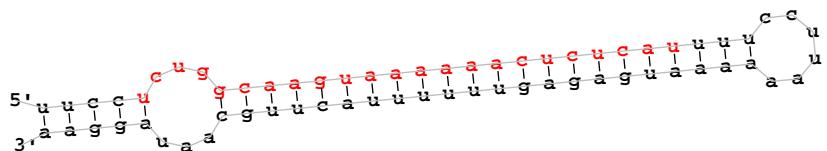


miRBase precursor : hsa-mir-3128

Total read count : 28

hsa-miR-3128 read count : 28

remaining reads : 0



hsa-miR-3128

5'	-3'	exp	reads	mm	sample
5'- uucc <u>c</u> ugg <u>g</u> caagu <u>aaaaaa</u> acuc <u>u</u> cauu	uuu <u>cc</u> uu <u>aaaaaa</u> aug <u>ag</u> ag <u>uuuuuu</u> ac <u>ug</u> ca <u>au</u> agg <u>aa</u>	.	1	0	Z3_-
....ucugg <u>g</u> caagu <u>aaaaaa</u> acuc <u>u</u> cu.....	9	0	Z1_-	
....ucugg <u>g</u> caagu <u>aaaaaa</u> acuc <u>u</u> cu.....	2	0	Z1_-	
....ucugg <u>g</u> caagu <u>aaaaaa</u> acuc <u>u</u>	5	0	GDM	
....ucugg <u>g</u> caagu <u>aaaaaa</u> acuc <u>u</u>	5	0	GDM	
....ucugg <u>g</u> caagu <u>aaaaaa</u> acuc <u>u</u>	5	0	GDM	
....ugg <u>g</u> caagu <u>aaaaaa</u> acuc <u>u</u> ca.....	2	0	GDM	
....ucugg <u>g</u> caagu <u>aaaaaa</u> acuc <u>u</u>	1	0	Z2_-	
....ucugg <u>g</u> caagu <u>aaaaaa</u> acuc <u>u</u> cu.....	1	0	Z2_-	
....ugg <u>g</u> caagu <u>aaaaaa</u> acuc <u>u</u> cauu.....	7	0	Z2_-	