

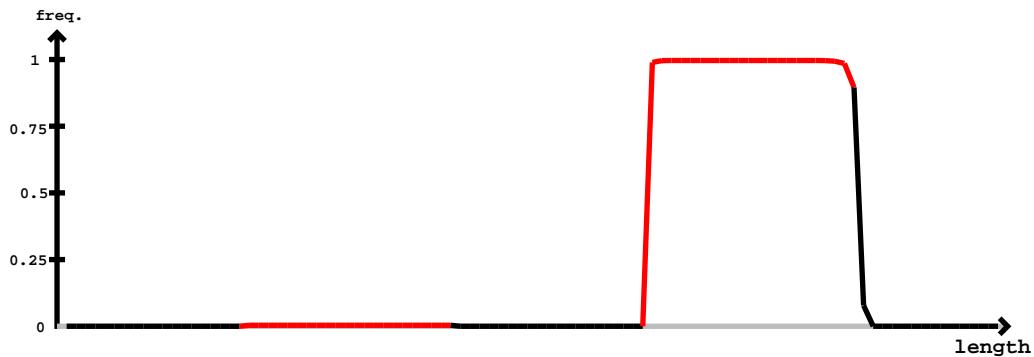
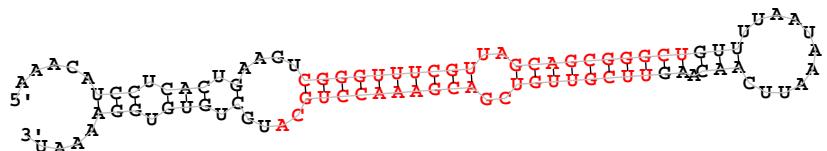
miRBase precursor : dme-mir-981_MI0005838

Total read count : 2191

dme-miR-981-3p_MIMAT0005497 read count

dme-miR-981-5p_MIMAT00020878 read count

remaining reads : 0



dme-miR-981-5p_MIMAT00020878

dme-miR-981-3p_MIMAT0005497

5' - aaacaucacugaagu	cggguuucguuagcagcgccu	guuuuaauaaacaag	uucguugucgacgaaaccugca	ugcuguguggaaaaau	-3'	exp	reads	mm	sample
.....(((((.....(((((((((.....((.....))))))))))))....))))....	0	seq
.....cggguuucguuagcagcgccu	7	0	seq
.....cggguuucguuagcagcgccu	1	1	seq
.....uucguugucgacgaaacc	1	0	seq
.....uucguugucgacgaaaccu	5	0	seq
.....uucguugucgacgaaaccug	18	0	seq
.....uucguugucgacgaaacGuc	1	1	seq
.....uucguugucgacgaaaccUG	1	1	seq
.....uucGUuugucgacgaaaccugc	1	1	seq
.....uucGUuugucgacgaaaccugc	1	1	seq
.....uucGUuugucgacgaaaccugc	1	1	seq
.....uucGUuugucgacgaaaccugc	189	0	seq
.....uucGUuugucgacgaaaccCgc	1	1	seq
.....uucGUuugGcgacgaaaccugc	1	1	seq
.....Cucguugucgacgaaaccugc	1	1	seq
.....uucGUuugucgacgaaaccugc	1	1	seq
.....uucGUuugucgacgaaaccugc	6	1	seq
.....uucGUuugucCcgaaaccugca	1	1	seq
.....uucGUuugucgacgaaaAcugca	1	1	seq
.....uucGUuugucgacgaaaACugca	1	1	seq
.....uucGUuugucgacgaaaccuUca	1	1	seq
.....Cucguugucgacgaaaccugca	4	1	seq
.....uuGUuugucgacgaaaccugca	1	1	seq
.....uucGUuugucgacgaaaccuCca	3	1	seq
.....uucGUuugucgacgaaaccugca	1	1	seq
.....uucGUuugucgacgUaacccugca	1	1	seq
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.....uucGUuUucgacgaaaccugca	2	1	seq
.....Aucguugucgacgaaaccugca	17	1	seq
.....uucGUuugucgacAAaccugca	4	1	seq
.....uucGUuugucgacaaacUugca	1	1	seq
.....uucGUuugucgacgaaaccugca	1662	0	seq
.....uucGUuugucgacgaaaccugcU	2	1	seq
.....uucGUuugucAacgaaaccugca	2	1	seq

dme-miR-981-5p_MIMAT0020878

dme-miR-981-3p_MIMAT0005497

aaacaucuccacugaagu**cggguuuucguuagcagcgccu**guuuuuauaaaauaacaaguucguugucgacgaaaccugcaugcuguguggaaaaau

.....uu <u>cgguugucgacgaaaccugAa</u>	1	1	seq
.....uu <u>cgguugucgacgaGaccugca</u>	3	1	seq
.....uu <u>cgguugucgacgaGccugca</u>	5	1	seq
.....uu <u>cgguugucgacgaaaccuAca</u>	1	1	seq
.....uu <u>cgguuAucgacgaaaccugca</u>	3	1	seq
.....uu <u>cgguugucgUcgaaaccugca</u>	1	1	seq
.....uu <u>cgguAgucgacgaaaccugca</u>	1	1	seq
.....uu <u>cgguugucgacgaaaccGgca</u>	4	1	seq
.....uu <u>cgguugucgacgCaaccugca</u>	4	1	seq
.....uu <u>cgguugucJucgaaaccugca</u>	2	1	seq
.....uu <u>cgguCgucgacgaaaccugca</u>	1	1	seq
.....uu <u>cgguugucgacgaaaccugca</u>	1	1	seq
.....uu <u>cgguugucgacgaaaccugUa</u>	4	1	seq
.....u <u>Acgguugucgacgaaaccugca</u>	1	1	seq
.....uu <u>cgguuCucgacgaaaccugca</u>	1	1	seq
.....uu <u>cgguugucgacgaaaccugC</u>	8	1	seq
.....uu <u>cgguugucgacgaaUccugca</u>	1	1	seq
.....uu <u>cgguugucgacgCaaccugca</u>	2	1	seq
.....uu <u>cgguugucgacgaaaccCgca</u>	2	1	seq
.....uu <u>cgguugucgCgcgaaaccugca</u>	5	1	seq
.....uu <u>cgguuugucgacgaaaccugca</u>	3	1	seq
.....uu <u>cgguGgucgacgaaaccugca</u>	1	1	seq
.....uu <u>cgguCugucgacgaaaccugca</u>	1	1	seq
.....uu <u>cgguugucgacgaaCcugca</u>	2	1	seq
.....uu <u>cgguugCcgacgaaaccugca</u>	1	1	seq
.....uu <u>cgguugucgacgaaaccAga</u>	1	1	seq
.....uu <u>cgguugucgacCaaaccugca</u>	3	1	seq
.....uu <u>cgguuguUgacgaaaccugca</u>	4	1	seq
.....uu <u>cgguugucgacgaaaccugcaA</u>	138	1	seq
.....uu <u>cgguugucgacgaaaccugcaC</u>	7	1	seq
.....uu <u>cgguugucgacgaaaccugcaG</u>	1	1	seq
.....uu <u>cgguugucgacgaaaccugcau</u>	17	0	seq
.....u <u>cgguugucgacgaaaccCgca</u>	1	1	seq
.....u <u>cgguugucgacgaaaccugca</u>	6	0	seq
.....A <u>cgguugucgacgaaaccugca</u>	1	1	seq
.....u <u>cgguugucgacgaaaccugcaA</u>	7	1	seq
.....c <u>guugucgacgaaaccugcaA</u>	1	1	seq