

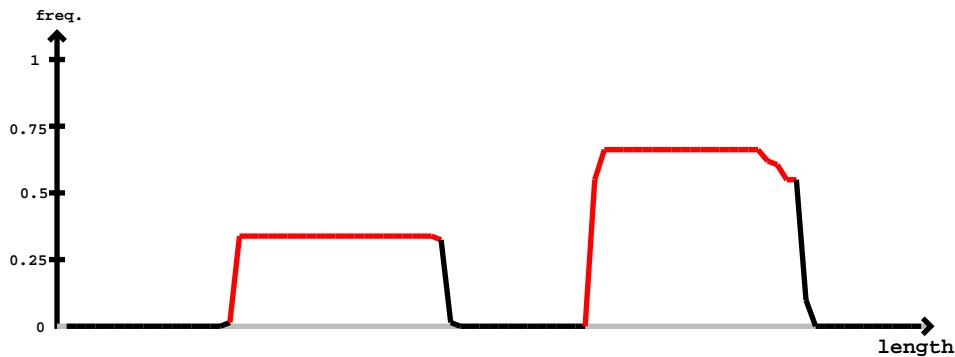
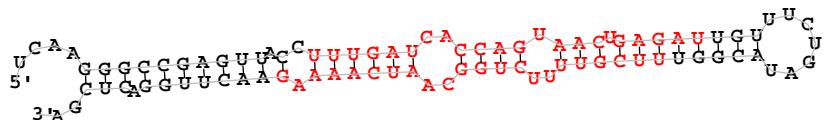
miRBase precursor : dme-mir-961_MI0005816

Total read count : 71

dme-miR-961-3p MIMAT0000856 read count

dme-miR-961-5p MIMAT00025859 read count
dme-miR-961-5p MIMAT00025475 read count

remaining reads : 0



dme-miR-961-3p_MIMAT0020856

dme-miR-961-5p_MIMAT0005475

5'- ucaaggggcagaguaccuuugaucaccaguacugagauuuucugauacgguuucguuuucuggcaa <u>aaaag</u> acuuggacu	-3'	exp	
	reads	mm	sample
.....((((((.(((.((((.((((((.....))))))))))))....))))....))))....))))..	1	1	seq
.....cuuugaucaccaguacugagauA.....	1	0	seq
.....uuugaucaccaguacugaga.....	1	1	seq
.....uuugaucaccaguaaGugagau.....	1	1	seq
.....uCugaucaccaguacugagau.....	20	0	seq
.....uuugaucaccaguacugagau.....	3	0	seq
.....uuucguuuuucuggcaa <u>u</u> ca.....	1	0	seq
.....uuucguuuuucuggcaa <u>u</u> caa.....	4	0	seq
.....uuucguuuuucuggc <u>G</u> aucaa <u>a</u> ag.....	1	1	seq
.....uuucguuuuucuggcaa <u>u</u> caa <u>a</u> ag.....	27	0	seq
.....uuucguuuuucuggcaa <u>u</u> caa <u>a</u> ag.....	1	1	seq
.....uuucguuuuucuggc <u>G</u> ucaa <u>a</u> ag.....	1	1	seq
.....uuucguuuuucuggcaa <u>u</u> caa <u>a</u> aga.....	1	0	seq
.....u <u>c</u> guuuuucuggcaa <u>u</u> caa <u>a</u> ag.....	2	0	seq
.....u <u>c</u> guuuuucuggcaa <u>u</u> caa <u>a</u> aga.....	2	0	seq
.....u <u>c</u> guuuuucuggcaa <u>u</u> caa <u>a</u> ag <u>C</u>	1	1	seq
.....u <u>c</u> quuuuucu <u>q</u> caa <u>u</u> caa <u>a</u> ag <u>U</u>	3	1	seq