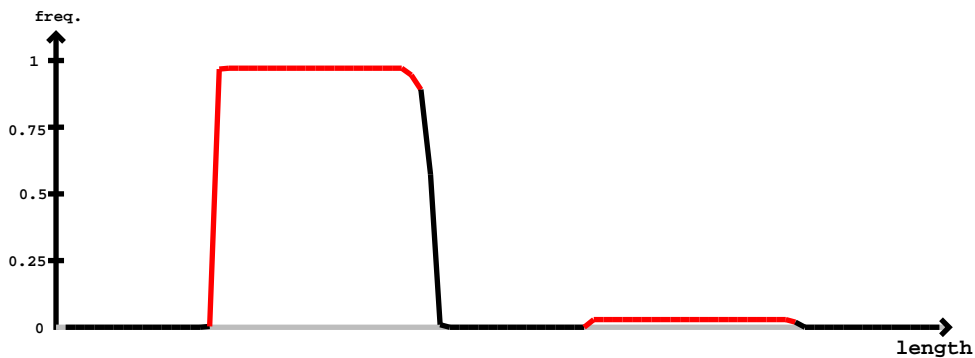
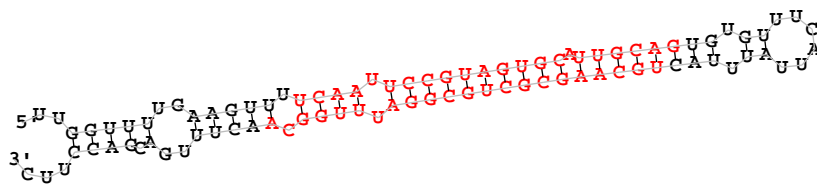


miRBase precursor : dme-mir-932_MI0005820
 Total read count : 312
 dme-miR-932-3p_MIMAT0020860 read count : 164
 dme-miR-932-5p_MIMAT0005479 read count : 148
 remaining reads : 0



dme-miR-932-3p_MIMAT0020860

dme-miR-932-5p_MIMAT0005479

5'	-3'	exp	reads	mm	sample
uugguuugaaguuu <u>ucaauuccguagugcauugc</u> aguguucauuuuuacugcaagcgcugcggaauuggcaacuugagaccuuc	..(((((((.....)))))))))				
.....Aucaauuccguagugcauugcag.....			1	1	seq
.....ucaauuccguaguAcauugc.....			1	1	seq
.....ucaauuccguagugcauugc.....			7	0	seq
.....ucaauuccguagugcauugca.....			17	0	seq
.....ucaauuccguagugcauugcaA.....			1	1	seq
.....ucaauuccguagugcauugcag.....			1	1	seq
.....ucaauuccgGagugcauugcag.....			1	1	seq
.....ucaauuccguagugcauugcag.....			90	0	seq
.....ucaauAccguagugcauugcag.....			2	1	seq
.....ucaGuuccguagugcauugcag.....			2	1	seq
.....ucaauuccguagugAauugcagu.....			1	1	seq
.....ucaauuccguagCgcauugcagu.....			2	1	seq
.....ucaauuccguagugcaAugcagu.....			1	1	seq
.....ucaauuccguagugcauugcaCu.....			1	1	seq
.....ucaauuccguagugcUuugcagu.....			1	1	seq
.....ucaauuccguagugcGuugcagu.....			1	1	seq
.....ucaauuccguaAugcauugcagu.....			1	1	seq
.....ucaauuccguagugcauugcagC.....			1	1	seq
.....ucaauuccguagugcauugcag.....			1	1	seq
.....ucaauuccguagugcauugcGgu.....			1	1	seq
.....ucaauuccguagugcauugcag.....			1	1	seq
.....ucaauuccguagugcauugcagu.....			164	0	seq
.....ucaauuccguagugcauugcaguU.....			3	1	seq
.....caauuccguagugcauugcag.....			1	0	seq
.....ugcaagcgcugcggaauuggc.....			3	0	seq
.....ugcaagcgcugcggaauuggca.....			5	0	seq
.....uAcaagcgcugcggaauuggca.....			1	1	seq