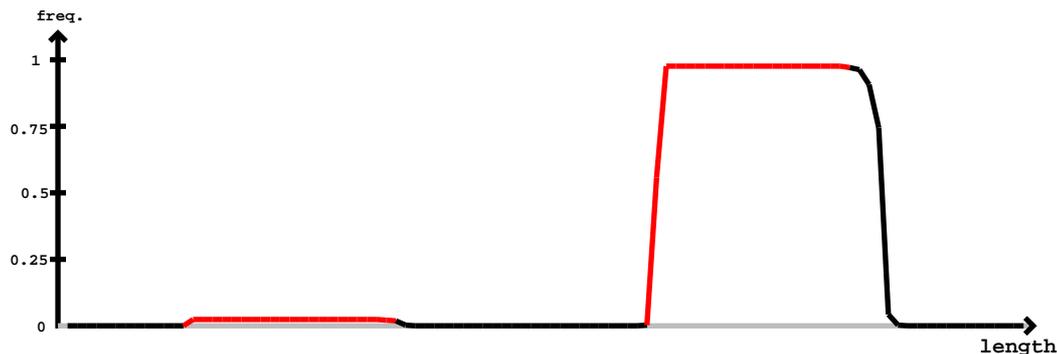
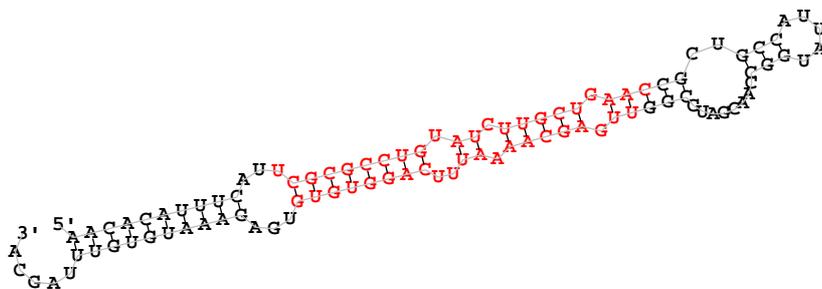


miRBase precursor : dme-mir-87_MI0000382
 Total read count : 376
 dme-miR-87-5p_MIMAT0020821 read count : 20821
 dme-miR-87-3p_MIMAT0000361 read count : 361
 remaining reads : 0



dme-miR-87-3p_MIMAT0000361

dme-miR-87-5p_MIMAT0020821

5'	3'	exp	reads	mm	sample
aacacauuucauucgcgccuguaucucugcugaaccgugcgaauauggccaacgaucggugagcaaaaauucaggugugugagaaauguuagca					
.....ucgcgccuguaucucugcuga.....			1	0	seq
.....ucgcgccugCaucuugcugaa.....			1	1	seq
.....ucgcgccuguaucucugcugaac.....			6	0	seq
.....ucgcgccuguaucucugcugaCcc.....			1	1	seq
.....guugagcaaaaauucagguguguU.....			1	1	seq
.....uugagcaaaaauucaggugugu.....			2	0	seq
.....uugagcaaaaauucaggugugA.....			1	1	seq
.....uugagcaaaaauucaggugug.....			2	0	seq
.....uugagcaaaaauucaggCgugugu.....			1	1	seq
.....uugagcaaaaauucagAugugu.....			1	1	seq
.....uugagcaaaaauucaggugugu.....			15	0	seq
.....uugagcaaaaauucaggugugug.....			1	1	seq
.....uugagcaaaaauUaggugugug.....			1	1	seq
.....Augagcaaaaauucaggugugug.....			1	1	seq
.....uugagcaaaaauucaggugugAug.....			2	1	seq
.....uugagcaaaaauucagguguguU.....			2	1	seq
.....uugagcaaaaauucaggAgugug.....			2	1	seq
.....uugagcaaaaauucaAgugugug.....			1	1	seq
.....uugagcaaaaauucaggugugCug.....			1	1	seq
.....uugagcaaaaauucaggugugug.....			2	1	seq
.....uugagcaaaaauucagCugugug.....			1	1	seq
.....uugagcaaaaauucaggugugug.....			25	0	seq
.....uugagcaaaaauucaggCguguga.....			1	1	seq
.....uugaAcaaaaauucagguguguga.....			1	1	seq
.....uugagcaaaaauucagguguguga.....			118	0	seq
.....Augagcaaaaauucagguguguga.....			2	1	seq
.....uugagcaaaaauucaggugugCga.....			2	1	seq
.....uugagcaaaaauucagCuguguga.....			1	1	seq
.....uugagcaaaaauucUgguguguga.....			1	1	seq
.....uugagcaaaaauucaggAguguga.....			4	1	seq
.....uugagAaaaaauucagguguguga.....			1	1	seq
.....uugagcaaaaauucaggugugAga.....			2	1	seq
.....uugagcaaaaauucagguguguUa.....			1	1	seq
.....uugagcaaaaauucagguUuguga.....			1	1	seq

aacacauuucauucgcgccgugaucucugcugaaccgcugccaauauggccaacgauccggugugagcaaaaauucaggugugugagaaauguguuagca

.....uugagcaaaaAuucagguguguga.....	3	1	seq
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.....Cugagcaaaaauucagguguguga.....	1	1	seq
.....uugagcaaaaCuucagguguguga.....	1	1	seq
.....uugagcaaaaauucagguguguga.....	2	1	seq
.....uugagcaaaaauucaggugugugU.....	2	1	seq
.....uugagcaaaaauuLagguguguga.....	1	1	seq
.....uugagcaaaaauucaggugugugaA.....	2	1	seq
.....ugagcaaaaauucagguguga.....	1	1	seq
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.....ugagcaaaaauucaggugugugaA.....	6	1	seq
.....ugagcaaaaauucaggugugugaC.....	3	1	seq
.....ugagcaaaaauucaggugugugaAa.....	1	1	seq