

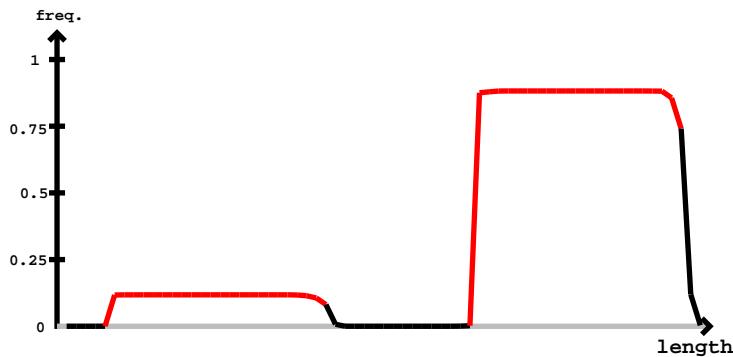
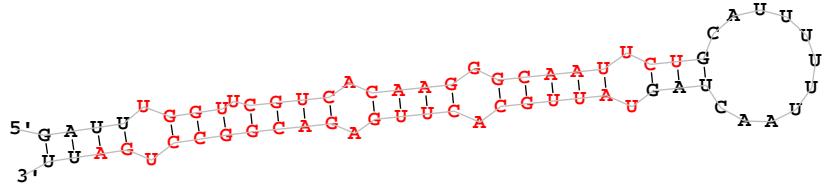
miRBase precursor : dme-mir-312_MI0000424

Total read count : 11387

dme-miR-312-3p_MIMAT0000404 read count

dme-miR-312-5p_MIMAT0020838 read count

remaining reads : 0



dme-miR-312-5p_MIMAT0020838

dme-miR-312-3p_MIMAT0000404

5' - sequence	- 3'	exp	reads	mm	sample
(((((.(((.(((.((((.(((((.....))))....))))....))))....))))....)))					
...Augguucguacacaaggggcaauucu.....			3	1	seq
...ugguucguacacaaggggcaaa.....			8	0	seq
...ugguucguacacaaggggcaau.....			30	0	seq
...ugguucguacGagggcaau.....			1	1	seq
...ugguucguacacaaggggcaaA.....			1	1	seq
...Gguuucguacacaaggggcaauu.....			1	1	seq
...ugguucguacacaaggggcaGuu.....			1	1	seq
...ugguucguacacaaggggcaauu.....			86	0	seq
...ugguucguacacaagAgcaauu.....			1	1	seq
...ugguucguacacaagCgcaauu.....			1	1	seq
...ugguucguacacaaggggcaauA.....			3	1	seq
...ugguucguacacaaggCcuaau.....			1	1	seq
...ugguucguacacaaggggAaaa.....			1	1	seq
...ugguucguacacaaggggcaauuU.....			1	1	seq
...Cgguucguacacaaggggcaauuc.....			1	1	seq
...ugguucguacacaaggggcaCuuc.....			1	1	seq
...ugguucguAacaaggggcaauuc.....			1	1	seq
...ugguucguacacaaggggcaauuc.....			258	0	seq
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...ugguucguacacaAGggcaauuc.....			1	1	seq
...ugUuucguacacaaggggcaauuc.....			1	1	seq
...ugguucguacacaagUgcuaauuc.....			1	1	seq
...ugguucguacacaaggggcaauG.....			1	1	seq
...ugguucguacacaaggggcaaGd.....			1	1	seq
...ugguucgucaUaggggcaauuc.....			1	1	seq
...ugguucguacacaaggggGaaauuc.....			1	1	seq
...ugguucguacacaaggggcaauuG.....			1	1	seq
...ugguucguacacaaggggcaaCuc.....			1	1	seq
...ugguucguacacaaggggcaauuA.....			4	1	seq
...ugguucguCcacaaggggcaauuc.....			1	1	seq
...ugguucguacUaggggcaauuc.....			1	1	seq
...Agguucguacacaaggggcaauuc.....			1	1	seq
...ugguuAguucacaaggggcaauuc.....			2	1	seq
...ugguGcgucacaaggggcaauuc.....			1	1	seq
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gauuuugguucguacacaaggggcaauuucugcauuuuuuuacuauguaauugcacuugagacggccugauu

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....ugguucAucacaaggggcaauuucu.....	2	1	seq
....ugguucguacGagggcaauuucu.....	1	1	seq
....ugguucguacacaaggggcaauuU.....	2	1	seq
....ugguucguacacaaggggcaCuucu.....	4	1	seq
....ugguucguacacaaggggcaauucA.....	52	1	seq
....ugguucguacacaaggUgcuaauucu.....	1	1	seq
....ugguucguacacaaggggcaUuuu.....	2	1	seq
....ugguucguacacaaggggcaauuuc.....	699	0	seq
....ugguucguacacaaggggcaauuAcu.....	1	1	seq
....Agguucguacacaaggggcaauuucu.....	23	1	seq
....ugguucguacacaaggAcaauuucu.....	1	1	seq
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....ugguuGGuacacaaggggcaauuucu.....	1	1	seq
....ugguucguacacaaggggUaaauucu.....	3	1	seq
....ugguucguacacaaggggCauuucu.....	2	1	seq
....ugguucguacacaaggggAaaauucu.....	2	1	seq
....uggCucguacacaaggggcaauuucu.....	3	1	seq
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....ugguucguacacaGgggcaauuucu.....	3	1	seq
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....ugguucguacacaaggggcaauuucU.....	2	1	seq
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....ugguucguacacaaggggcaauuucugc.....	1	0	seq
....ugguucguacacaaggggcaauuucuAc.....	1	1	seq
....ggucguacacaaggggcaauuucu.....	7	0	seq
....uucguacacaaggggcaauuucu.....	1	0	seq
....Aucguacacaaggggcaauuucu.....	1	1	seq
.....aAuauugcacuugagacggccuga...	5	1	seq
.....Uuauugcacuugagacggccug...	1	1	seq
.....Auauugcacuugagacggccug...	2	1	seq
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.....uauuugcacuugagacggccCg...	3	1	seq
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.....uauuugcacuugagGggccug...	3	1	seq
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.....uauuugcacuugagacggccuA...	15	1	seq

dme-miR-312-5p_MIMAT0020838

dme-miR-312-3p_MIMAT0000404

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.....Aauugcacuugagacggccug...	14	1	seq
.....uaauugcacuugagacggGcug...	1	1	seq
.....uaauugcacuugagacggccug...	1201	0	seq
.....uaauugcacu <u>Cgagacggccug...</u>	2	1	seq
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.....uaauu <u>Ucacuugagacggccug...</u>	1	1	seq
.....ua <u>Augcacuugagacggccug...</u>	1	1	seq
.....uaauug <u>cacuugagac<u>Gccug...</u></u>	3	1	seq
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.....uaauug <u>cacuui<u>Uagacggccug...</u></u>	1	1	seq
.....uaauug <u>cacuuga<u>Aacggccug...</u></u>	1	1	seq
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.....ua <u>Ugcacuugagacggccug...</u>	2	1	seq
.....u <u>Guugcacuugagacggccug...</u>	3	1	seq
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.....uaauug <u>cacuugagacggccu<u>A...</u></u>	8	1	seq
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.....ua <u>Cugcacuugagacggccuga...</u>	14	1	seq
.....uaauug <u>cacuugaga<u>Uggccuga...</u></u>	6	1	seq
.....uaauug <u>cacuugagac<u>cgccAuga...</u></u>	11	1	seq
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.....uaauug <u>cacuugagac<u>Gccuga...</u></u>	12	1	seq
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.....uaauug <u>cacuugagac<u>Cccuga...</u></u>	1	1	seq
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.....uaauug <u>cacuugagac<u>Uggccuga...</u></u>	12	1	seq
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.....uaauug <u>cacu<u>Ggagacggccuga...</u></u>	10	1	seq
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.....uaauug <u>cacuugagacgg<u>ccuA...</u></u>	5	1	seq
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.....uaauug <u>cacuugagacgg<u>Gcuga...</u></u>	1	1	seq
.....uaauug <u>cacuuu<u>Cagacggccuga...</u></u>	3	1	seq
.....uaauug <u>cacuugagac<u>Agccuga...</u></u>	8	1	seq
.....uaauu <u>Ucacuugagacggccuga...</u>	5	1	seq

dme-miR-312-5p_MIMAT0020838

dme-miR-312-3p_MIMAT0000404

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.....	.uauugcacuugagacggccGga..	11	1	seq
.....	.uauugcacuugagacggccugG..	20	1	seq
.....	.uaGugcacuugagacggccuga..	6	1	seq
.....	.uauugcacuugUgacggccuga..	1	1	seq
.....	.uauugcacuuUagacggccuga..	2	1	seq
.....	.uauugcGcuuugagacggccuga..	4	1	seq
.....	.uauugcacuugagacGgccuga..	3	1	seq
.....	.uauugcacuugagaaacggccuga..	10	1	seq
.....	.uauugcacuugagacggccugU..	3	1	seq
.....	.uauugcaUuugagacggccugau..	1	1	seq
.....	.uauugcacuugagacggccugau..	93	0	seq
.....	.uauugcacuugagaAggccugau..	1	1	seq
.....	.uauugcacuugagacggccugaC..	12	1	seq
.....	.Aauugcacuugagacggccugau..	1	1	seq
.....	.uauugcacuugagacggAcugau..	1	1	seq
.....	.uauugcacuugagaAacggccugau..	1	1	seq
.....	.uauugcacuugagacggccugCu..	2	1	seq
.....	.uauugcacuugagacggccugaA..	1180	1	seq
.....	.uauugcacuugagacggccugaG..	8	1	seq
.....	.uauugcacuugagacggccugaAu	18	1	seq
.....	.uauugcacuugagacggccugaA	15	1	seq
.....	.uauugcacuugagacggccugaC	1	1	seq
.....	.uauugcacuugagacggccugaGu	1	1	seq
.....	.uauugcacuugagacggccugauu	5	0	seq
.....	.auugcacuugagacggccu....	1	0	seq
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.....	.auugcacuugagacggccugau..	5	0	seq
.....	.auugcacuugagacggccugaA..	4	1	seq
.....	.uugcacuugagacggccug....	6	0	seq
.....	.uugcacuugagacggccugU..	1	1	seq
.....	.uugcacuugagacggccuga..	21	0	seq
.....	.AugcacuugagacggccugaC..	1	1	seq
.....	.uugcacuugagacggccugac..	1	1	seq
.....	.uugcacuugagacggccugaA..	4	1	seq
.....	.ugcacuugagacggccuga..	3	0	seq