

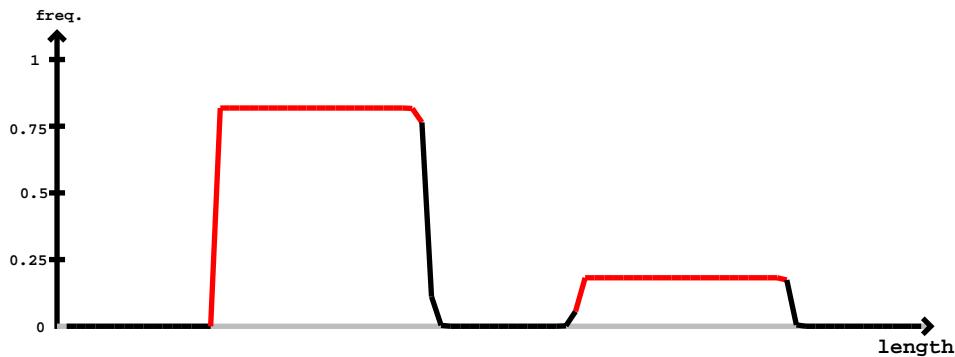
miRBase precursor : dme-mir-316_MI0000428

Total read count : 594

dme-miR-316-3p_MIMAT00020842 read count

dme-miR-316-5p_MIMAT0000408 read count

remaining reads : 0



dme-miR-316-3p_MIMAT00020842

5' - seq	-3' seq	exp	sample
.....aaaaucuagucgauuuugucuuuuuuccgcuuacuggcguuucaauuuccacaacg	acaggaaaaggaaaaaggcguauuuacuaugaguuu
.....ugucuuuuuuccgcuuacugg.....	1	0
.....ugucuuuuuuccgAuuuacuggc.....	1	1
.....ugucuuuuuuccgcuuacuggc.....	29	0
.....ugucuuuuuuccgUuuuacuggc.....	1	1
.....ugucuuuuuuccgcuuuacuggcg.....	1	1
.....ugucuuuuuuccgcuuuacuggcg.....	1	1
.....ugucuuuuuuccgcuuuacuggcg.....	1	1
.....ugucuuuuuuccgcuuuacuggcg.....	1	1
.....ugucuuuuuuccgcuuuacuggcg.....	1	1
.....ugucuuuuuuccgcuuuacuggcg.....	2	1
.....uAucuuuuuuccgcuuuacuggcg.....	2	1
.....ugucuuuuuuccgcuuuacuggcC.....	1	1
.....ugucuuuuuuccgcuuuacuggcg.....	1	1
.....ugucuuuuuuccgcuuuacuggcg.....	1	1
.....ugucuuuuuuccgcuuuacuggcg.....	3	1
.....ugucuuuuuuccgAuuuacuggcg.....	1	1
.....ugucuuuuuuccgcuuuacuggcA.....	3	1
.....ugucuuuuuuccgcuuuacuUgcg.....	1	1
.....ugucuuuuuuccgcuuuacuggG.....	1	1
.....ugucuuuuuuccgcuuuacugUcg.....	1	1
.....Cgucuuuuuuccgcuuuacuggcg.....	1	1
.....Agucuuuuuuccgcuuuacuggcg.....	9	1
.....ugucuuuuuuccgcuuuGcuggcg.....	1	1
.....ugucuuuuuuccgcuuuacAggcg.....	1	1
.....ugucuCuuuuccgcuuuacuggcg.....	1	1
.....uguUuuuuuuccgcuuuacuggcg.....	1	1
.....ugucuuuuuuccgcuuuUcuggcg.....	1	1
.....ugucuuuuuuccAcuuuacuggcg.....	2	1
.....ugucuuuuuuccgcuuuacuggcg.....	352	0
.....ugucuuuuuuccgcuuuacuCgcg.....	1	1
.....ugucuuuuuuccgcuuuacugAcg.....	1	1
.....ugucuuuuuuccgcuuuCcuggcg.....	1	1
.....Agucuuuuuuccgcuuuacuggcg.....	2	1
.....ugucuuuuuuccAcuuuacuggcg.....	1	1
.....ugucuuuuuuccgcuuuacuggcg.....	54	0
.....ugucuuuuuuccgcuuuacuUgcg.....	1	1
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dme-miR-316-5p_MIMAT0000408

aaaaucuagugucguuuugucuuuuuuccgcuuucuacuacgacgaaagggaaaagggguauuuuacuaugaguuu

.....ugucuuuuuccgcuuacuggcgA.....	1	1	seq
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.....ugucuuuuuccgcuuacuggcguu.....	1	0	seq
.....ugucuuuuuccgcuuacuggcg <u>A</u>	1	1	seq
.....cgacaggaaaggaaaaaggcg.....	1	0	seq
.....gacaggaaagg <u>G</u> aaaaggcgua.....	1	1	seq
.....gacaggaaag <u>A</u> gaaaaggcgua.....	1	1	seq
.....gacaggaaaggaaaaAgcgua.....	1	1	seq
.....gacaggaaaggaaaaaggcgua.....	27	0	seq
.....gacaggaaaggaaaaaggcg <u>C</u> a.....	2	1	seq
.....acaggaaaggaaaaaggcg.....	4	0	seq
.....acaggaaaggGaaaaggcgua.....	1	1	seq
.....aAaggaaaggaaaaaggcgua.....	2	1	seq
.....acaggaaagUaaaaaggcgua.....	2	1	seq
.....acaggaaaggaaaaaggc <u>A</u> ua.....	1	1	seq
.....acaggaaaggaaaaaggcgua.....	57	0	seq
.....acaggCaaggaaaaaggcgua.....	1	1	seq
.....acaggaaagg <u>A</u> aaaaggcgua.....	1	1	seq
.....acaggaaaggaaaaGggcgua.....	1	1	seq
.....acaggaaaggaaaaCgcgu.....	1	1	seq
.....acaggaGaggaaaaaggcgua.....	1	1	seq
.....acaggaaaggaaaaaggcg <u>u</u> au.....	1	0	seq
.....acaggaaaggaaaaaggcg <u>A</u>	2	1	seq