

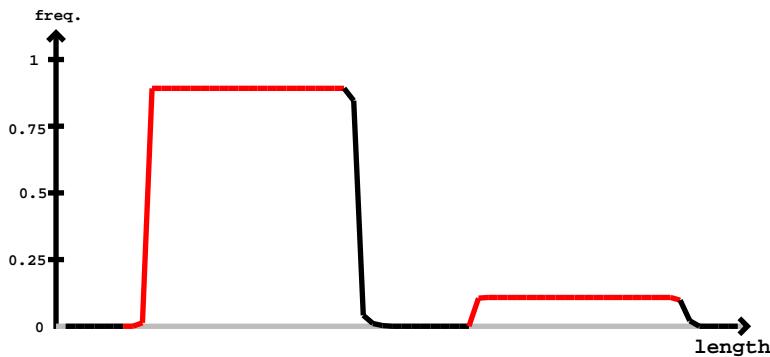
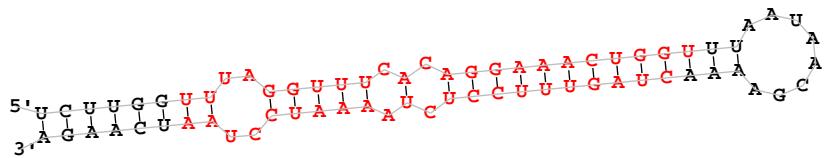
miRBase precursor : dme-mir-303\_MI0000409

Total read count : 389

dme-miR-303-5p\_MIMAT0000388 read count

dme-miR-303-3p\_MIMAT0020824 read count

remaining reads : 0



dme-miR-303-3p\_MIMAT0020824

5' -	-3'	exp	
reads	mm	sample	
(((((.....((((((.....)))))))))))).....	0	seq	
.....uaaggguuucacaggaaacugguu.....	2	0	seq
.....uaaggguuucacaggaaacugguuA.....	1	1	seq
.....uaaggguuucacaggaaacugguu.....	1	0	seq
.....Auaggguuucacaggaaacugguu.....	1	1	seq
.....uaggguuucacaggaaacugguu.....	13	0	seq
.....uaggguuucacaggaaacugguu.....	1	1	seq
.....uaggguuucacaggaaacugguu.....	1	1	seq
.....uagggGuacaggaaacugguu.....	1	1	seq
.....uaggguuucacaggaaGcuggu.....	1	1	seq
.....uaggguuucacaggaaacugguuA.....	6	1	seq
.....uaggguuucacaggaaAcAgguu.....	1	1	seq
.....uaggguuucacaggGaacugguu.....	2	1	seq
.....uaggguuucacaGaaaacugguu.....	3	1	seq
.....uagggUuuucacaggaaacugguu.....	1	1	seq
.....uaggguuucacaggaaGacugguu.....	1	1	seq
.....uaggAuuucacaggaaacugguu.....	1	1	seq
.....uaggGuuacaggaaacugguu.....	1	1	seq
.....uaggguuucacaggaaaUugguu.....	1	1	seq
.....uaggguuucacaggaaacCgguu.....	1	1	seq
.....uaggguuucacaggaaacugguUC.....	1	1	seq
.....uaggguuucacaggaaCcgguu.....	1	1	seq
.....uaggguuucacaggaaacugguUu.....	1	1	seq
.....uaggguuucacaggaaacugggGu.....	3	1	seq
.....uaggguuucacaggCaacugguu.....	1	1	seq
.....uaggguuucacaggaaacugguu.....	285	0	seq
.....uaggguuucacaggaaacugguu.....	8	0	seq
.....uaggguuucacaggaaacugguUC.....	1	1	seq
.....uaggguuucacaggaaacugguA.....	3	1	seq
.....uaggguuucacaggaaacugguAA.....	3	1	seq
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.....cuaggguuuccucuaaaauccua.....	3	0	seq
.....cAaggguuuccucuaaaauccua.....	1	1	seq
.....cAaggguuuccucuaaaauccua.....	1	1	seq
.....cuaggguuuccucuaaaauccua.....	27	0	seq
.....cuaggguuuccucuuaGauccua.....	1	1	seq

dme-miR-303-3p\_MIMAT0000388

ucuugguuuagguuucacaggaaacugguuuuaauaacgaaaacuaguuuccucuaaaaauccuaaucaaga

.....cAaguuuccucuaaaaauccuaau.....	2	1	seq
.....cuaguuuccucuaaaaauccuaau.....	6	0	seq
.....uaguuuccucuaaaaauccuaa.....	1	0	seq