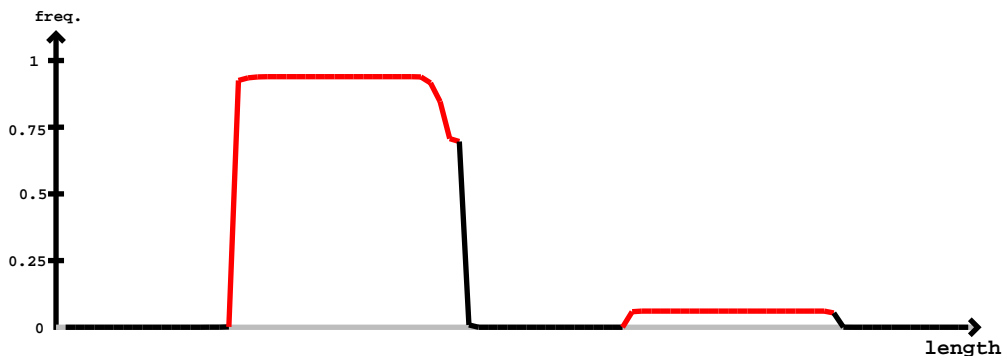
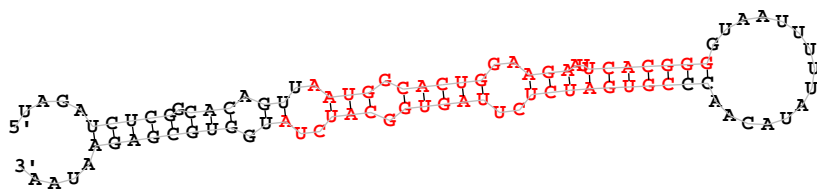


miRBase precursor : dme-mir-263a\_MI0000343  
 Total read count : 3415  
 dme-miR-263a-5p\_MIMAT0000319 read count  
 dme-miR-263a-3p\_MIMAT0000799 read count  
 remaining reads : 1



dme-miR-263a-3p\_MIMAT0020799

dme-miR-263a-5p\_MIMAT0000319

5' -	reads	mm	sample
uagaucucggcacaguu...aauggcacuggaagaauuacacggg...gaaauuuuuuauacaacccgugaucucuaguggcaucauuggugcgagaauaa	195	0	seq
...guuaauggcacuggaagaauuacac...	1	0	seq
...Aaauggcacuggaagaauuacac...	1	1	seq
...Aaauggcacuggaagaauuacacg...	1	1	seq
...Aaauggcacuggaagaauuacacggg...	2	1	seq
...aauggcacuggaagaauuc...	3	0	seq
...aauggcacuggaagGauuca...	1	1	seq
...aauggcacuggaagaAaLuca...	1	1	seq
...aauggcacuggaagaauuca...	64	0	seq
...aauggcacugAaagaauuca...	2	1	seq
...aauggcacuggaagaauucU...	1	1	seq
...aaUgAcacuggaagaauuca...	1	1	seq
...aauggcacuggaagaauucG...	1	1	seq
...aauggcacuggaagaauAca...	2	1	seq
...aauggcacuggaAaauuca...	1	1	seq
...aGuggcacuggaagaauuca...	1	1	seq
...aauggcacuAgaagaauuacac...	1	1	seq
...aauggcacuGgaagaauuacac...	1	1	seq
...aauggcacuggaagaauuacac...	195	0	seq
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...aauggcacuggaCgaauuacac...	2	1	seq
...aauggcacuggaagaCuucac...	1	1	seq
...aauggcacuggaagUauuacac...	3	1	seq
...aauggcacuggaagaauucaA...	1	1	seq
...aauggcacuggaagaauucaU...	1	1	seq
...aauggcacCggaagaauuacac...	1	1	seq
...aauggcacuggaAaauuacac...	6	1	seq
...aaUgCacuggaagaauuacac...	1	1	seq
...aauggcacugAaagaauuacac...	5	1	seq
...Uauggcacuggaagaauuacac...	1	1	seq
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.....aauggcacugCaagaauucacg.....	1	1	seq
.....aauggcacuggaaga <u>A</u> ucacg.....	6	1	seq
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.....aauggcacug <u>A</u> aagaauucacg.....	3	1	seq
.....aauggcacuggaagaauucacg.....	404	0	seq
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.....aauggcacuggaagaauGc <u>ac</u> g.....	1	1	seq
.....aauggcacuggaGgaauucacg.....	1	1	seq
.....aauggcacuggaagaCuc <u>ac</u> g.....	2	1	seq
.....aauggcacCggaagaauucacg.....	1	1	seq
.....aauggcacugga <u>A</u> auucacg.....	2	1	seq
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.....aauggcacuggaagaauuc <u>U</u> cg.....	1	1	seq
.....aauggcGcuggaagaauucacg.....	1	1	seq
.....aauggcacuggaagaauU <u>ac</u> g.....	2	1	seq
.....aauggcacuggaCaauucacg.....	1	1	seq
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.....aaUgacacuggaagaauucacg.....	1	1	seq
.....aauggcacuggaagaauucac <u>A</u> .....	6	1	seq
.....aauggcacuggaagaau <u>A</u> cacg.....	7	1	seq
.....aauggcacuggaagaauucacg <u>g</u> .....	31	0	seq
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.....aauggcacuggaagaauucacg <u>A</u> .....	1	1	seq
.....aauggcacCggaagaauucacg <u>gg</u> .....	1	1	seq
.....aauggcGcuggaagaauucacg <u>gg</u> .....	2	1	seq
.....aauggcacuggaagaau <u>A</u> cg <u>gg</u> .....	3	1	seq
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.....aaGggcacuggaagaauucacg <u>gg</u> .....	7	1	seq
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