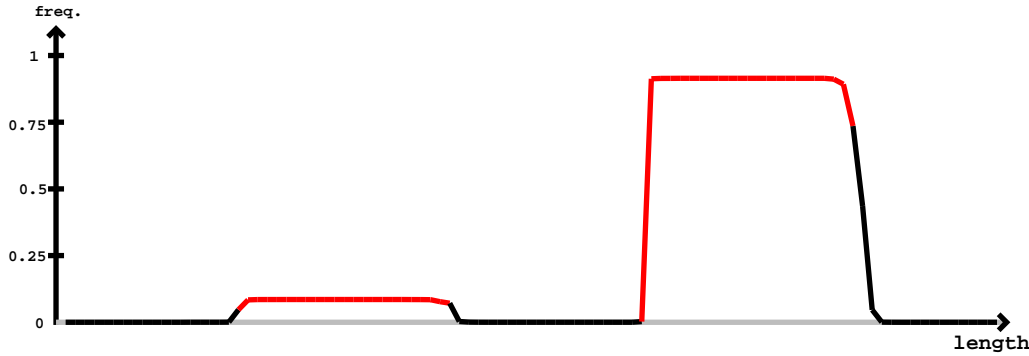
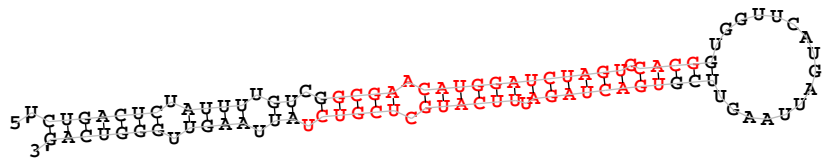


miRBase precursor : dme-mir-996\_MI000587  
 Total read count : 32021  
 dme-miR-996-5p\_MIMAT0020895 read count : 27895  
 dme-miR-996-3p\_MIMAT0005515 read count : 4126  
 remaining reads : 9



dme-miR-996-3p\_MIMAT0005515

dme-miR-996-5p\_MIMAT0020895

5' -	reads	mm	sample
ucugagacucuaauuuugucgcggaacauggaucuagugcaccgugguucaugauuaaguucgugacuagauuucaugcucgucuaauaaguuggucag			
.....uugcggcgaacauggaucuagu.....	4	0	seq
.....Augcggcgaacauggaucuagu.....	1	1	seq
.....ucggcgaacauggaucuagugca.....	1	0	seq
.....Aggcgaacauggaucuagugcacg.....	4	1	seq
.....cggcgaacauggaucuagugcacg.....	6	0	seq
.....ggcgaacauggaucuagug.....	1	0	seq
.....Agcgaacauggaucuagugc.....	1	1	seq
.....ggcgaacauggaucuaAugc.....	1	1	seq
.....ggcgaacauggaucuagugc.....	10	0	seq
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.....ggcgaacauggaCcuagugca.....	1	1	seq
.....ggcAaacauggaucuagugca.....	1	1	seq
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.....ggcUaacauggaucuagugca.....	1	1	seq
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.....ggcgaacauggaucuagugcaU.....	1	1	seq
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.....ggcgaacauggaucuagugcCcg.....	1	1	seq
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.....ggcgaUcauggauc <u>uagugcacg</u> .....	1	1	seq
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.....gUcgaacauggaucuagugcacg.....	6	1	seq
.....ggcgaacauggaucuagAgcacg.....	2	1	seq
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.....ggGgaacauggaucuagugcacg.....	1	1	seq
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. . . . . . acauggaucuagugcaccg . . . . .	1	0	seq
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. . . . . . . . gCgacuagauuucaugcucguc . . . . .	1	1	seq
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. . . . . . . . Augacuagauuucaugcucguc . . . . .	3	1	seq
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. . . . . . . . Uugacuagauuucaugcucgucua . . . . .	1	1	seq
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. . . . . . . . gugacuagauuucaugcucgucua . . . . .	1	0	seq
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. . . . . . . . ugacuagauuuCugcucgu . . . . .	1	1	seq
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. . . . . . . . ugacuagauuucaugcuUgu . . . . .	1	1	seq
. . . . . . . . ugacuagaCucaugcucgu . . . . .	1	1	seq
. . . . . . . . ugacuagauuucauAcucgu . . . . .	2	1	seq

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.....uUacuagauuucaugcucguc.....	7	1	seq
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.....ugacuUgauuucaugcucgucu.....	3	1	seq
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.....ugacuagauuucaugcucgucuG.....	16	1	seq
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.....ugacuagauuucauCcucgucua.....	6	1	seq
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.....ugacuagauuucaugcucAucua.....	9	1	seq
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.....ugacuCgauuucaugcucgucua.....	3	1	seq
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.....ugacuagauuucaugGucgucua.....	2	1	seq
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.....gacuagauuucaugcucgucU.....	18	0	seq
.....UacuagauuucaugcucgucU.....	1	1	seq
.....gacuagauuucaugcucgCcu.....	1	1	seq

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.....gacuagauuucaugcucgucuaA.....	1	1	seq
.....acuagauuucaugcucgucua.....	2	0	seq
.....cuagauuucaugcucguc.....	2	0	seq
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