

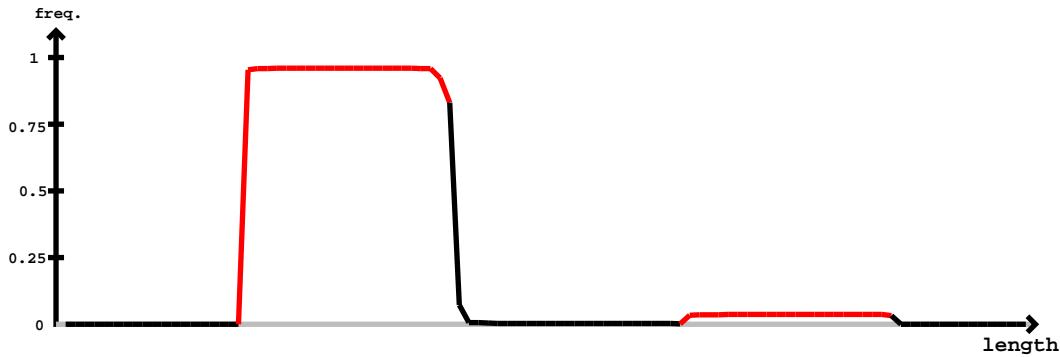
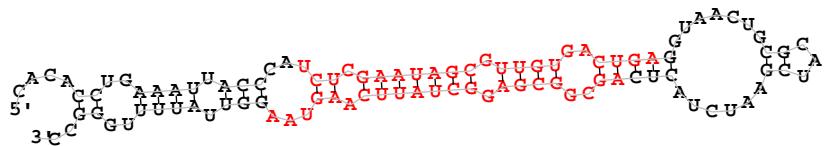
miRBase precursor : dme-mir-986_MI0005845

Total read count : 650

dme-miR-986-5p_MIMAT0005503 read count

dme-miR-986-3p_MIMAT0020884 read count

remaining reads : 2



dme-miR-986-3p_MIMAT0020884

| 5' - | -3' | exp | reads | mm | sample |
|--|-------|-----|-------|-----|--------|
|(((((.....(((((.....((((((.....))))....))))....))))....)))).... | | | | | |
|ucucgaaauagcgguugugacuga..... | | 1 | 0 | seq | |
|ucucgaaauagUguugugacu..... | | 18 | 1 | seq | |
|ucucgaaauagcgguugugacu..... | | 4 | 0 | seq | |
|ucucgaaauagcgguugugCcu..... | | 1 | 1 | seq | |
|ucucgaaauagAguugugacug..... | | 1 | 1 | seq | |
|ucucgaaauagUguugugacug..... | | 49 | 1 | seq | |
|ucucgAAauagcgguugugacug..... | | 1 | 1 | seq | |
|ucucgAAauagcgguugAgacug..... | | 1 | 1 | seq | |
|ucucgAAauagcgguugugacug..... | | 8 | 0 | seq | |
|ucucgAAauagcgguuguguCacuga..... | | 1 | 1 | seq | |
|ucucgAAauagAguugugacuga..... | | 2 | 1 | seq | |
|ucucgAAauagcgguugAgacuga..... | | 3 | 1 | seq | |
|ucucgAAauagCagcgguugugacuga..... | | 1 | 1 | seq | |
|ucucgAAauagcgguugCgacuga..... | | 2 | 1 | seq | |
|ucuUgaaauagcgguugugacuga..... | | 1 | 1 | seq | |
|ucucgAAauagcgguugugacugU..... | | 1 | 1 | seq | |
|ucucgAAauagcgguugugGcuga..... | | 1 | 1 | seq | |
|ucucgAAauagcgguugugacuga..... | | 103 | 0 | seq | |
|AcucgAAauagcgguugugacuga..... | | 2 | 1 | seq | |
|ucucgAAauagUguugugacuga..... | | 377 | 1 | seq | |
|ucucgAAauagcgguugugacugA..... | | 15 | 1 | seq | |
|ucucgAAauagcgguugugacugaU..... | | 3 | 1 | seq | |
|ucucgAAauagUguugugacugag..... | | 19 | 1 | seq | |
|ucucgAAauagcgguugugacugag..... | | 4 | 0 | seq | |
|ucucgAAauagUguugugacugaggu..... | | 1 | 1 | seq | |
|cucgAAauagUguugugacuga..... | | 1 | 1 | seq | |
|cucgAAauagcgguugugacugag..... | | 1 | 0 | seq | |
|cucgAAauagUguugugacugag..... | | 1 | 1 | seq | |
|cgaaauagcgguugugacugag..... | | 1 | 0 | seq | |
|gguaacugcgcaaucuacu..... | | 1 | 0 | seq | |
|gguaacugcgcaaucuacu..... | | 1 | 0 | seq | |
|agcggcgaggcuaucaagu..... | | 3 | 0 | seq | |
|agcggcgaggcuaucaagu..... | | 1 | 1 | seq | |
|agcggcgaggcuauUaagu..... | | 5 | 1 | seq | |

dme-miR-986-3p_MIMAT0005503

cacaccugaaauuacccaucucgaaauagcguugugacugagguaacuacuacuacagcgccgaggcuauucaaguaagguuauuuuggccc

| | | | | |
|-------|-------------------------|----|---|-----|
| | agcgccgaggcuauucaaguaa. | 12 | 0 | seq |
| | agcgccgaggcuauucaaguaa. | 1 | 1 | seq |
| | gccccgaggcuauucaaguaa. | 1 | 0 | seq |
| | gcgaggcuauucaaguaa. | 1 | 0 | seq |