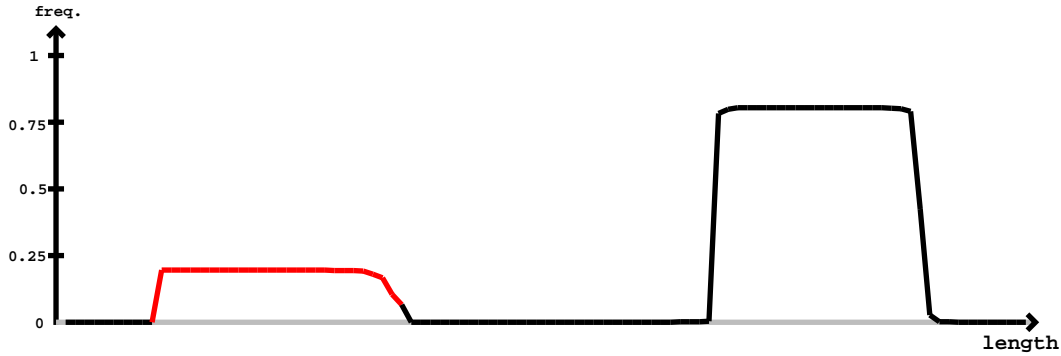
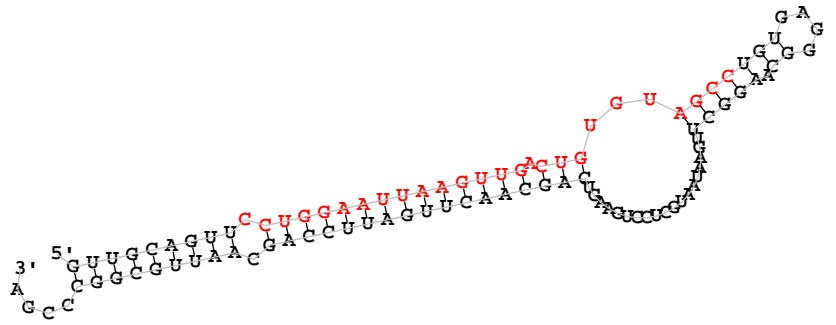


miRBase precursor : dme-mir-284_MI000369
 Total read count : 516
 dme-miR-284-5p_MIMAT0020811 read count : 20811
 remaining reads : 415



dme-miR-284-5p_MIMAT0020811

| 5' | 3' | exp | reads | mm | sample |
|-----------|--|-----|-------|----|--------|
| guugcaguu | ccuggaauaaguugacugugagccugugagggcaaggcuugaauauguccugaagucagcaacuugauccagcaauugcgccga | | | | |
| | | | 1 | 0 | seq |
| | | | 1 | 0 | seq |
| | | | 1 | 1 | seq |
| | | | 5 | 0 | seq |
| | | | 1 | 1 | seq |
| | | | 1 | 1 | seq |
| | | | 5 | 0 | seq |
| | | | 1 | 1 | seq |
| | | | 1 | 1 | seq |
| | | | 30 | 0 | seq |
| | | | 1 | 1 | seq |
| | | | 1 | 1 | seq |
| | | | 18 | 0 | seq |
| | | | 1 | 1 | seq |
| | | | 1 | 1 | seq |
| | | | 1 | 1 | seq |
| | | | 1 | 1 | seq |
| | | | 28 | 0 | seq |
| | | | 1 | 1 | seq |
| | | | 1 | 1 | seq |
| | | | 1 | 0 | seq |
| | | | 1 | 0 | seq |
| | | | 1 | 0 | seq |
| | | | 5 | 0 | seq |
| | | | 4 | 1 | seq |
| | | | 2 | 1 | seq |
| | | | 1 | 1 | seq |
| | | | 1 | 1 | seq |
| | | | 2 | 1 | seq |
| | | | 2 | 1 | seq |
| | | | 1 | 1 | seq |
| | | | 3 | 1 | seq |
| | | | 1 | 1 | seq |

| | | | |
|--|-----|---|-----|
| guugcaguuccuggaauaaguugacuguguagccugugagggcaaggcuugaauaugcuccugaagucagcaacuugauuccagcaauugcgcccgga | | | |
|gucagcaacuugGuuccagca..... | 1 | 1 | seq |
|gucagcaacuGgauuccagca..... | 1 | 1 | seq |
|gucagcaacuugauuccagca..... | 165 | 0 | seq |
|gucGcaacuugauuccagca..... | 1 | 1 | seq |
|gucagcaacuugauuccaAcaa..... | 2 | 1 | seq |
|gucagcaacuugauuccagcaa..... | 2 | 1 | seq |
|gucagcaacuugaGuccagcaa..... | 1 | 1 | seq |
|gGcagcaacuugauuccagcaa..... | 1 | 1 | seq |
|gucagcaacuugauCccagcaa..... | 1 | 1 | seq |
|gucagcaacuugauuccagUaa..... | 1 | 1 | seq |
|gucAcaacuugauuccagcaa..... | 1 | 1 | seq |
|gucagcaacuAuuuccagcaa..... | 1 | 1 | seq |
|gucagcaacuugauuccagcaa..... | 174 | 0 | seq |
|gucagcaacuugauuccagcaC..... | 3 | 1 | seq |
|gucagUaacuugauuccagcaa..... | 1 | 1 | seq |
|gucagcaacuugauuccagcCa..... | 1 | 1 | seq |
|gucagcaacuAgauccagcaa..... | 2 | 1 | seq |
|gucagGaacuugauuccagcaa..... | 1 | 1 | seq |
|gucagAaacuugauuccagcaa..... | 1 | 1 | seq |
|gucagcaacuugauuccagcGa..... | 1 | 1 | seq |
|gucagcaacuugauuccGgcaa..... | 1 | 1 | seq |
|gucagcaacuugauuccaCcaa..... | 1 | 1 | seq |
|gucagcaacuugauucUagcaa..... | 1 | 1 | seq |
|gucagcaacuugauAccagcaa..... | 3 | 1 | seq |
|gucagcaacuugauuccagcUa..... | 1 | 1 | seq |
|gucagcaacuugauuccagcaau..... | 2 | 0 | seq |
|gucagcaacuugauuccagcaaA..... | 6 | 1 | seq |
|ucagcaacuugauuccagca..... | 1 | 0 | seq |
|ucagcaacuugauuccagcaa..... | 2 | 0 | seq |
|ucagcaacuugauuccagcaau..... | 4 | 0 | seq |
|ucagcaacuugaAuccagcaau..... | 1 | 1 | seq |
|cagcaacuugauuccagca..... | 1 | 0 | seq |
|cagcaacuugauuccagcaa..... | 2 | 0 | seq |