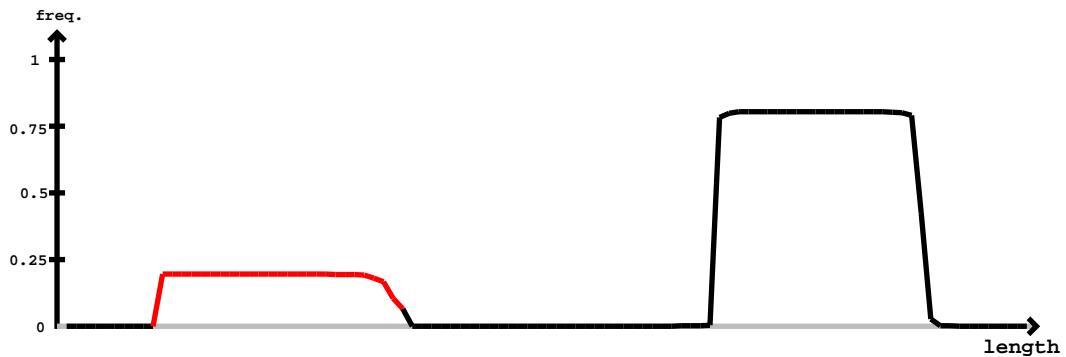
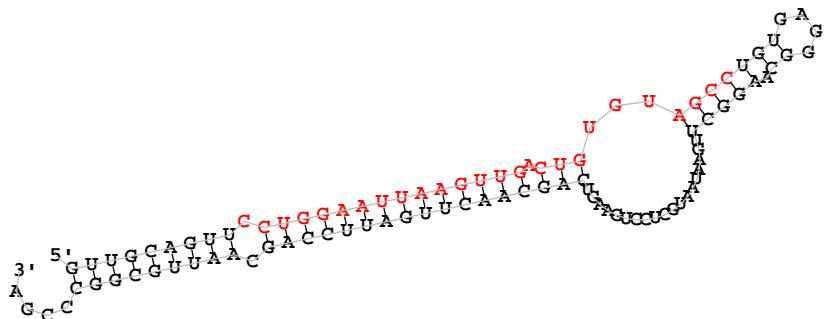


miRBase precursor : dme-mir-284\_MI0000369  
 Total read count : 516  
 dme-miR-284-5p\_MIMAT0020811 read count  
 remaining reads : 415



#### dme-miR-284-5p\_MIMAT0020811

5' - sequence	-3' sequence	exp	reads	mm	sample
.....ccuggaaauuaaguuugacugugacccuugaggcaaggcuuugaauuaugcuccugaagucagcaacuuugauuccagaauugcgccccga	-3'				
.....ccuggaaauuaaguuugacu.....			1	0	seq
.....ccuggaaauuaaguuugacugug.....			1	0	seq
.....ccuggaaauuaagGugacugugu.....			1	1	seq
.....ccuggaaauuaaguugacugugu.....			5	0	seq
.....ccuggaaauuaaguugacAgugua.....			1	1	seq
.....ccuggaaauuaaguugacuCugua.....			1	1	seq
.....ccuggaaauuaaguugacugugua.....			5	0	seq
.....cAuggaaauuaaguugacuguguag.....			1	1	seq
.....ccuggaaauuaaguugUcuguguag.....			1	1	seq
.....ccuggaaauuaaguugacuguguag.....			30	0	seq
.....ccuggaaCuaaguugacuguguagc.....			1	1	seq
.....ccuggaaauuaaguuAacuguguagc.....			1	1	seq
.....ccuggaaauuaaguugacuguguagc.....			18	0	seq
.....ccuggaaauuaaguugacuguguUgcc.....			1	1	seq
.....ccuggaaauuaaguugacuguguagcG.....			1	1	seq
.....Gcuggaaauuaaguugacuguguagcc.....			1	1	seq
.....ccuggaaauuaaguugacuguguagUc.....			1	1	seq
.....ccuggaaauuaaguugacuguguagcc.....			28	0	seq
.....ccuggaaauuaaguugacuguguagGc.....			1	1	seq
.....ccuUGaaauuaaguugacuguguagcc.....			1	1	seq
.....ugaagucagcaacuuugauuccagcaauug.....			1	0	seq
.....agucagcaacuuugauuccagca.....			1	0	seq
.....gucagcaacuuugauucca.....			1	0	seq
.....gucagcaacuuugauuccag.....			1	0	seq
.....gucagcaacuuugauuccagc.....			5	0	seq
.....gucagcaacuuugauucUagca.....			4	1	seq
.....gucagcaacuuCaucuccagca.....			2	1	seq
.....gucagcaaAuugauuccagca.....			1	1	seq
.....gucagcaacCugauuccagca.....			1	1	seq
.....gucagAAacuuugauuccagca.....			1	1	seq
.....gucagcaacuuugauuccGgca.....			2	1	seq
.....gucagcaacuuugauuccUgca.....			2	1	seq
.....gucagcaacuuugaAuccagca.....			1	1	seq
.....gucagcaacuAgauuccagca.....			3	1	seq
.....gucagcaacuuugacuccagca.....			1	1	seq

guugcaguuccuggaauaaguugacugguagccugugaggggcaaggcuugaaauaugcuccugaaggcaaucugaauccucccaaauugcgggcccga  
.....gucagcaacuugGuuccagca..... 1 1 seq  
.....gucagcaacuCgauuccagca..... 1 1 seq  
.....gucagcaacuugauuccagca..... 165 0 seq  
.....gucGgcaacuugauuccagca..... 1 1 seq  
.....gucagcaacuugauuccaAcaa..... 2 1 seq  
.....gucagcaacAugauuccagcaa..... 2 1 seq  
.....gucagcaacuugauuccagcaa..... 1 1 seq  
.....gGcagcaacuugauuccagcaa..... 1 1 seq  
.....gucagcaacuuugauCccagcaa..... 1 1 seq  
.....gucagcaacuugauuccagUa..... 1 1 seq  
.....guAcacuugauuccagcaa..... 1 1 seq  
.....gucagcaacuuauuccagcaa..... 1 1 seq  
.....gucagcaacuuugauuccagcaa..... 174 0 seq  
.....gucagcaacuugauuccagca..... 3 1 seq  
.....gucagUaacuugauuccagcaa..... 1 1 seq  
.....gucagcaacuugauuccAgCa..... 1 1 seq  
.....gucagcaacuuAgauuccagcaa..... 2 1 seq  
.....gucagGaacuugauuccagcaa..... 1 1 seq  
.....gucagAaacuugauuccagcaa..... 1 1 seq  
.....gucagcaacuugauuccGa..... 1 1 seq  
.....gucagcaacuuugauuccGgaa..... 1 1 seq  
.....gucagcaacuugauuccCaaa..... 1 1 seq  
.....gucagcaacuugauuccUagcaa..... 1 1 seq  
.....gucagcaacuugauAccagcaa..... 3 1 seq  
.....gucagcaacuugauuccAgUa..... 1 1 seq  
.....gucagcaacuugauuccAgcaa..... 2 0 seq  
.....gucagcaacuugauuccAgcaaA..... 6 1 seq  
.....ucagcaacuugauuccagca..... 1 0 seq  
.....ucagcaacuugauuccAgcaa..... 2 0 seq  
.....ucagcaacuugauuccAgcaa..... 4 0 seq  
.....ucagcaacuugauuccAgcaa..... 1 1 seq  
.....cagcaacuugauuccAgcaa..... 1 0 seq  
.....cagcaacuugauuccAgcaa..... 2 0 seq