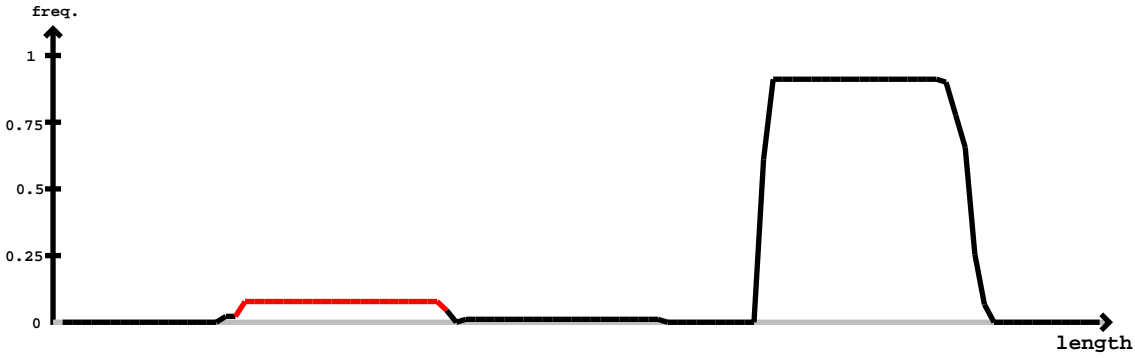
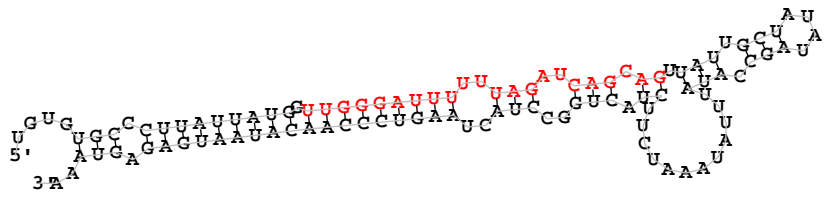


miRBase precursor : dme-mir-193\_MI0005809  
Total read count : 90  
dme-miR-193-5p\_MIMAT0020849 read count  
remaining reads : 83



dme-miR-193-5p\_MIMAT0020849

5'	-3'	exp	reads	mm	sample
ugugugcccuauuuauugguugggauuuuuuagaucagcaguuauugcuauuagccauuuuuuaaauucucuaucuggccuacuaagucccaacaauaagagagauaaa					
(((.....(((((((.....(((.....(((.....)))))).....)))))).....)))))).....)))))).....)))))).....))))..					
.....gguugggauuuuuuagaucagca.....			1	0	seq
.....gguugggauuuuuuagaucagcaA.....			1	1	seq
.....uugAgauuuuuuagaucagca.....			1	1	seq
.....uugggauuuuuuagaucagca.....			1	0	seq
.....uugggauuuuuuagaucagcag.....			3	0	seq
.....uuugcuauuagccauuuu.....			1	0	seq
.....uacuggccuacuaaguccc.....			1	0	seq
.....uacugCccuacuaaguccca.....			1	1	seq
.....uacuggccuacuaaguccca.....			7	0	seq
.....uacuggccuacuaagCccca.....			1	1	seq
.....uacuggccuacuaagucccaa.....			10	0	seq
.....uGcuggccuacuaagucccaac.....			1	1	seq
.....uacGggccuacuaagucccaac.....			1	1	seq
.....uacuggccuacuaagucccaaA.....			2	1	seq
.....uacugCccuacuaagucccaac.....			1	1	seq
.....uacuggccuacuaagAcccaac.....			1	1	seq
.....uacuggccuacuaagucccaac.....			20	0	seq
.....uCcuggccuacuaagucccaac.....			1	1	seq
.....uacuggccuacuaaguUccaac.....			1	1	seq
.....Aacuggccuacuaagucccaac.....			7	1	seq
.....acuggccuacuaaguccca.....			3	0	seq
.....acuggccuacuaagucccaac.....			1	0	seq
.....acuggccuacuaagucccUaca.....			1	1	seq
.....acuggccuacuaagucccGaca.....			1	1	seq
.....acuggccuacuaagucccaacU.....			1	1	seq
.....acuggccuacuaagucccaaca.....			14	0	seq
.....acuggccuacuaagucccaacau.....			6	0	seq