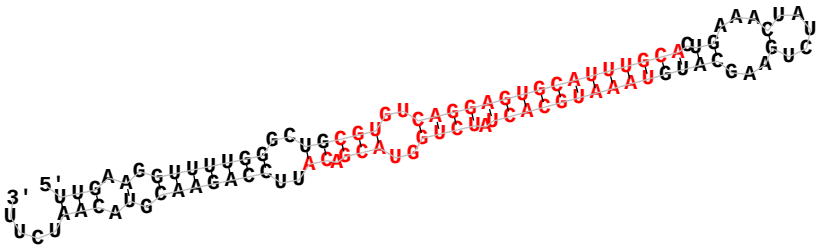
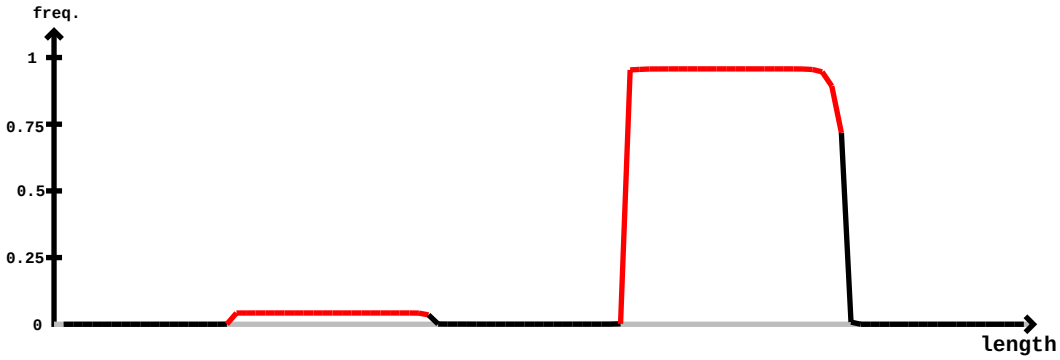
miRBase precursor : dme-mir-277_MI0000360
Total read count : 15576
dme-miR-277-5p_MIMAT0020804 read count
dme-miR-277-3p_MIMAT0000338 read count
remaining reads : 6





dme-miR-277-3p_MIMAT0000338

dme-miR-277-5p_MIMAT0020804

5'- uugaagguuuugggcugcgugucaggagugcauuugcacugaaacuaucugaagcaugcaaaugcacuaucugguacgacauuccagaacguacaaucuu -3' exp

(((.(.(((((((..(((((..((((((((((((((((.((...(.....)...)))))))))))))).))))..))).))..)))))))).).))).... reads mm sample

.....gcgugAcaggagugcauuugc.....	1	1 seq
.....gcgugucaggagugcauuugc.....	3	0 seq
.....gcgugucaggagugcauuugca.....	1	0 seq
.....gcgugucaggagugcauuugcaA.....	1	1 seq
.....cgugucaggagugcauuCg.....	1	1 seq
.....cgugucaggagGgcauuug.....	1	1 seq
.....cgugucaggagugcauuug.....	6	0 seq
.....cgugucaggaAugcauuugc.....	2	1 seq
.....cgugucaggagugcauuugc.....	80	0 seq
.....cgugucaAgagugcauuugc.....	1	1 seq
.....cguUucaggagugcauuugc.....	1	1 seq
.....cgugucaggagugcaAuugc.....	1	1 seq
.....cgugucaggagugcauuugA.....	1	1 seq
.....cgugucaggagugcauuuUc.....	1	1 seq
.....cgugucaggagugcauuAgc.....	2	1 seq
.....Agugucaggagugcauuugc.....	1	1 seq
.....cgugucaggagugcauAugc.....	3	1 seq
.....cgugucaggagAgcauuugc.....	1	1 seq
.....cgugucaggagugcauuugU.....	2	1 seq
.....cgugucaggagCgcauuugca.....	3	1 seq
.....cgugucaggagugcauuugcU.....	2	1 seq
.....cgugucaggagugcGuuugca.....	1	1 seq
.....cgugucaggagugcaGuugca.....	1	1 seq
.....cgugucaggagugcUuuugca.....	2	1 seq
.....cgugucaggaAugcauuugca.....	7	1 seq
.....cgugucagCagugcauuugca.....	1	1 seq
.....cgugucaggagugcauuCgca.....	1	1 seq
.....cguguGaggagugcauuugca.....	1	1 seq
.....cgugAcaggagugcauuugca.....	2	1 seq
.....cgugucGggagugcauuugca.....	1	1 seq
.....cgugucaggagAgcauuugca.....	5	1 seq
.....cgugucaggagugcauuAgca.....	9	1 seq
.....cgugucaggagugcaAuugca.....	8	1 seq
.....cgugucaggagugcauCugca.....	5	1 seq

dme-miR-277-3p_MIMAT0000338

dme-miR-277-5p_MIMAT0020804

uugaaggguuuugggcugcgugucaggagugcauuugcacugaaacuaucugaagcauguaaaugcacuaucugguacgacauuccagaacguacaaucuu

.....cgugCcaggagugcauuugca.....	3	1	seq
.....cgugucaggagugcauuugca.....	459	0	seq
.....cgugucaggagugcauAugca.....	6	1	seq
.....cgugucaAgagugcauuugca.....	1	1	seq
.....cguCucaggagugcauuugca.....	1	1	seq
.....cgugucaggagugcaCuugca.....	5	1	seq
.....cgugucaggagugcauuugcC.....	1	1	seq
.....cguguAaggagugcauuugca.....	1	1	seq
.....cgugucaggagugGauuugca.....	3	1	seq
.....cgugucaggaguAcauuugca.....	1	1	seq
.....cgGgucaggagugcauuugca.....	1	1	seq
.....cgugucaggagugcauuugUa.....	1	1	seq
.....Agugucaggagugcauuugca.....	4	1	seq
.....cgugucaggaCugcauuugca.....	1	1	seq
.....cAugucaggagugcauuugca.....	1	1	seq
.....cgugucaggagugcauuugcac.....	1	0	seq
.....cgugucaggagugcauuugcaA.....	3	1	seq
.....cgugucaggagugcauuugcacug.....	2	0	seq
.....cgugucaggagugcauuugcacuga.....	2	0	seq
.....cgugucaggagugcauuugcacugaa.....	1	0	seq
.....gugucaggagugcauuugc.....	1	0	seq
.....ugucaggagugcauuugc.....	1	0	seq
.....cugUaacuaucugaagcaug.....	1	1	seq
.....cugaaacuaucugaagcaug.....	4	0	seq
.....agcauguaaaugcacuaucugguac.....	1	0	seq
.....uguaaaugcacuaucugguacgaca.....	4	0	seq
.....Auaaaugcacuaucugguacga.....	2	1	seq
.....Auaaaugcacuaucugguacgac.....	5	1	seq
.....Auaaaugcacuaucugguacgaca.....	9	1	seq
.....uaaaugcacuaucuggua.....	7	0	seq
.....uaaaugcacuaucugguaU.....	1	1	seq
.....uaaaugcacuaucuggCac.....	1	1	seq
.....uaaaugcacuaucugguac.....	25	0	seq
.....uaaaugcacuaucugguaA.....	1	1	seq
.....uaaaugcacuaucuggCacg.....	2	1	seq
.....uaaaugcacuaCcugguacg.....	1	1	seq
.....uaaaugcacCaucugguacg.....	1	1	seq
.....uaaaugcacuaucuggAacg.....	3	1	seq
.....uaaauAcacuaucugguacg.....	1	1	seq
.....uaaaugcacuauGugguacg.....	1	1	seq
.....uaaaugcacuaucugUuacg.....	1	1	seq
.....uaaaugcacuaucugguacA.....	2	1	seq
.....uaaaugcacuaucugguacg.....	123	0	seq
.....uaaaugcacuaucugAuacg.....	3	1	seq
.....uaaaugcacuaucugguaGg.....	1	1	seq
.....uaaaugcacuaucuAguacga.....	1	1	seq
.....uaaaugcacuaucugguacgU.....	3	1	seq
.....uaaaugcacuaucugguGcga.....	1	1	seq
.....uaaaugcacuaGcugguacga.....	1	1	seq
.....uaaaugcacuaucugguacga.....	719	0	seq
.....uaaaugcaUuaucugguacga.....	1	1	seq
.....uaaaugUacuaucugguacga.....	1	1	seq
.....uaaaugcacuaucuCguacga.....	6	1	seq
.....uaaaugcacuaucCgguacga.....	4	1	seq
.....uaaauAcacuaucugguacga.....	6	1	seq
.....uaaaugcacuaucugCuacga.....	6	1	seq
.....Caaaugcacuaucugguacga.....	2	1	seq
.....uaaaugcacuaucuUguacga.....	2	1	seq
.....uaaauCcacuaucugguacga.....	2	1	seq
.....uaaaugcacuaucuggAacga.....	9	1	seq
.....uaaaugcacuaucuggCacga.....	3	1	seq
.....uaaaugcacuaucugAuacga.....	18	1	seq
.....uaaaCgcacuaucugguacga.....	1	1	seq
.....uaaaugcacuaucugUuacga.....	1	1	seq
.....uaaaugcacuaucugguaUga.....	1	1	seq
.....uaaaugcacuaucugguaGga.....	2	1	seq
.....uaaaugcacAaucugguacga.....	2	1	seq
.....uCaaugcacuaucugguacga.....	1	1	seq
.....uaaaugcacGaucugguacga.....	1	1	seq

dme-miR-277-3p_MIMAT0000338

dme-miR-277-5p_MIMAT0020804

uugaaggguuuugggcugcgugucaggagugcauuugcacugaaacuaucugaagcaguaaaugcacuaucugguacgacauuccagaacguacaaucuu

.....Gaaaugcacuaucugguacga.....	1	1	seq
.....uaaaugcacuauUugguacga.....	1	1	seq
.....uaaaugcacuaucugguUcga.....	1	1	seq
.....uGaaugcacuaucugguacga.....	1	1	seq
.....uaaaugcacuaucugguacAa.....	8	1	seq
.....uaaaugcacuaCcugguacga.....	1	1	seq
.....uaaaugcacuaucAgguacga.....	5	1	seq
.....uaaaugcacuauAugguacga.....	1	1	seq
.....uaaaugcacuauGugguacga.....	1	1	seq
.....uaaaugcacCaucugguacga.....	2	1	seq
.....uaaaugcacuaucugguacgG.....	2	1	seq
.....uaUaugcacuaucugguacga.....	1	1	seq
.....uaaaugcacuaucugguacUa.....	1	1	seq
.....uaaaAgcacuaucugguacgac.....	6	1	seq
.....uaaaugcacuaucuggAacgac.....	43	1	seq
.....uaaaugcacuaucuAguacgac.....	9	1	seq
.....Aaaaugcacuaucugguacgac.....	2	1	seq
.....uaaaugcaGuaucugguacgac.....	2	1	seq
.....uaaaugcacuaucugguacgaU.....	55	1	seq
.....uaaaugcacuaucugguCcgac.....	1	1	seq
.....uaUaugcacuaucugguacgac.....	1	1	seq
.....uaaaugcacuauGugguacgac.....	3	1	seq
.....uaaaugcacuaucCgguacgac.....	5	1	seq
.....uaaaugcacuaAcugguacgac.....	8	1	seq
.....uaaaugcacuaucugguGcgac.....	1	1	seq
.....uaaauCcacuaucugguacgac.....	3	1	seq
.....uaaauUcacuaucugguacgac.....	3	1	seq
.....uaaaugcacuaucugguacgCc.....	2	1	seq
.....uaaaugcacAaucugguacgac.....	7	1	seq
.....uaaaugcaUuaucugguacgac.....	4	1	seq
.....uaaaugcGcuaucugguacgac.....	5	1	seq
.....uaaauAcacuaucugguacgac.....	14	1	seq
.....uaaaugcacuaucugguacgaA.....	57	1	seq
.....uaaaugcacuaucugguacAac.....	16	1	seq
.....uaaaugcacuaCcugguacgac.....	8	1	seq
.....uaaaugcacuaucugguacgGc.....	6	1	seq
.....uaaaugcacuaucugguUcgac.....	4	1	seq
.....uaaaugcacCaucugguacgac.....	8	1	seq
.....uaaaugcacuaucuCguacgac.....	11	1	seq
.....uaaaugcacuaucugguacCac.....	7	1	seq
.....uGaaugcacuaucugguacgac.....	1	1	seq
.....uaaaugcacuaucugguAgac.....	5	1	seq
.....uaaGugcacuaucugguacgac.....	7	1	seq
.....uaaaugcacuaucuUguacgac.....	4	1	seq
.....uaGaugcacuaucugguacgac.....	3	1	seq
.....uaaaugcacuaucAgguacgac.....	14	1	seq
.....uaaUugcacuaucugguacgac.....	1	1	seq
.....uaaaCgcacuaucugguacgac.....	3	1	seq
.....uaaaugcacuaGcugguacgac.....	1	1	seq
.....uaaaugcacuGucugguacgac.....	1	1	seq
.....uaaaugcacuauUugguacgac.....	11	1	seq
.....uaaaugcacuaucuggGacgac.....	2	1	seq
.....uaaaugcacuaucugguaGgac.....	1	1	seq
.....uaaaugUacuaucugguacgac.....	2	1	seq
.....uaaaugcacuaucuggCacgac.....	29	1	seq
.....uaaaugcacuaucugCuacgac.....	12	1	seq
.....uaaaugcacuaucugguacgUc.....	9	1	seq
.....uaaaugcacuaucugAuacgac.....	44	1	seq
.....uaaaugcacuaucugUuacgac.....	1	1	seq
.....uaaaugcacuaucugguacgac.....	2243	0	seq
.....uaaaugcacuauAugguacgac.....	4	1	seq
.....uaaaugcacuaucugguacgaG.....	22	1	seq
.....Caaaugcacuaucugguacgac.....	2	1	seq
.....uaaauCcacuaucugguacgaca.....	9	1	seq
.....uaaaugcacuaAcugguacgaca.....	74	1	seq
.....uaaaugcacuaucugUuacgaca.....	8	1	seq
.....uGaaugcacuaucugguacgaca.....	8	1	seq
.....uaaaugcacuUucugguacgaca.....	4	1	seq
.....uaaaugcacuaucugAuacgaca.....	178	1	seq

dme-miR-277-3p_MIMAT0000338

dme-miR-277-5p_MIMAT0020804

uugaaggguuuugggcugcgugucaggagugcauuugcacugaaacuaucugaagcauguaaaugcacuaucugguacgacauuccagaacguacaaucuu

.....uaaaugcacuaucugguacgUca.....	41	1	seq
.....uaaaugUacuaucugguacgaca.....	2	1	seq
.....uaaaugcacGaucugguacgaca.....	2	1	seq
.....uaaauAcacuaucugguacgaca.....	85	1	seq
.....uaaaugcacuGucugguacgaca.....	10	1	seq
.....uCaaugcacuaucugguacgaca.....	6	1	seq
.....uaaaugcacuaucugguacgaUa.....	90	1	seq
.....uaaaugcacuaucugguCcgaca.....	3	1	seq
.....uaaaugcacuaucCgguacgaca.....	70	1	seq
.....uaaaugcacuaucuCguacgaca.....	52	1	seq
.....uUaaugcacuaucugguacgaca.....	2	1	seq
.....uaaaugcacuauAugguacgaca.....	13	1	seq
.....uaaaugcacuaucugguacgCca.....	15	1	seq
.....uaaaugcacCaucugguacgaca.....	22	1	seq
.....uaaaugcGcuaucugguacgaca.....	28	1	seq
.....uaaCugcacuaucugguacgaca.....	2	1	seq
.....uaaaugcacuaucAgguacgaca.....	46	1	seq
.....uaaGugcacuaucugguacgaca.....	18	1	seq
.....uaaaugcCcuaucugguacgaca.....	2	1	seq
.....uaaaugcacuaucugguacUaca.....	4	1	seq
.....uaaauUcacuaucugguacgaca.....	5	1	seq
.....uaaaugcacuaucugguacgacC.....	32	1	seq
.....uaaaugcacuaucugguaUgaca.....	34	1	seq
.....uaaaugcacuCucugguacgaca.....	2	1	seq
.....uaaaugcacuaucuggAacgaca.....	203	1	seq
.....uaaaCgcacuaucugguacgaca.....	19	1	seq
.....uaaaugcacuaucugguacgaAa.....	10	1	seq
.....uaaaugcacuaucugguacgacU.....	14	1	seq
.....uaUaugcacuaucugguacgaca.....	1	1	seq
.....uaaaugcacuaucugguGcgaca.....	12	1	seq
.....uaaaugcacuaucugguaGgaca.....	4	1	seq
.....uaaaugGacuaucugguacgaca.....	2	1	seq
.....uaaaugcacuaucugguacgaGa.....	13	1	seq
.....uaaaugcacuaucuggCacgaca.....	105	1	seq
.....uaaaugcacuaucugguacAaca.....	140	1	seq
.....uaaUugcacuaucugguacgaca.....	10	1	seq
.....uaaaugcacuaucugguaAgaca.....	3	1	seq
.....uaaaugcacAaucugguacgaca.....	17	1	seq
.....uaCaugcacuaucugguacgaca.....	2	1	seq
.....uaaaugcacuaucuggGacgaca.....	12	1	seq
.....uaaaugAacuaucugguacgaca.....	4	1	seq
.....uaaaugcacuauGugguacgaca.....	17	1	seq
.....uaaaugcacuaucugguUcgaca.....	58	1	seq
.....uaaaugcacuaucugguacgacG.....	24	1	seq
.....uaaaugcUcuaucugguacgaca.....	4	1	seq
.....uaaaugcaUuaucugguacgaca.....	3	1	seq
.....uaaaugcacuaucuAguacgaca.....	50	1	seq
.....uaaaugcacuauUugguacgaca.....	29	1	seq
.....Caaaugcacuaucugguacgaca.....	8	1	seq
.....uaaaugcacuaucugCuacgaca.....	58	1	seq
.....uaaaugcacuaucuUguacgaca.....	8	1	seq
.....uaaaugcacuaucugguacgaca.....	9129	0	seq
.....Aaaaugcacuaucugguacgaca.....	21	1	seq
.....uaaaugcacuaucugguacgGca.....	39	1	seq
.....uaaaugcacuaucugguaCaca.....	58	1	seq
.....uaGaugcacuaucugguacgaca.....	9	1	seq
.....uaaaAgcacuaucugguacgaca.....	12	1	seq
.....uaaaugcacuaucGgguacgaca.....	7	1	seq
.....uaaaugcacuaCcugguacgaca.....	28	1	seq
.....uaaaugcacuaucugguacgacaA.....	85	1	seq
.....uaaaugcacuaucugguacgacaG.....	2	1	seq
.....uaaaugcacuaucugguacgacaC.....	2	1	seq
.....uaaaugcacuaucuggAacgacau.....	1	1	seq
.....uaaaugcacuaucugguUcgacau.....	1	1	seq
.....uaaaugcacuaucugguacgacau.....	27	0	seq
.....uaaaugcacuaAcugguacgacau.....	1	1	seq
.....uaaaugcacuaucugguacgaUau.....	1	1	seq
.....uaaaugcacuaucugguacgUcau.....	1	1	seq
.....uaaaugcacuaucugguacgacauu.....	1	0	seq

dme-miR-277-3p_MIMAT0000338

dme-miR-277-5p_MIMAT0020804

uugaaggguuuugggcugcgugucaggagugcauuugcacugaaacuaucugaagcaguguaaaugcacuaucugguacgacauuccagaacguacaaucuu

.....uaaaugcacuaucugguacgacaAu.....	1	1	seq
.....aaaugcacuaucugguacga.....	3	0	seq
.....Uaaugcacuaucugguacga.....	2	1	seq
.....aaaugcacuaucugguacgac.....	2	0	seq
.....aaaugcacuaAcugguacgaca.....	1	1	seq
.....Uaaugcacuaucugguacgaca.....	2	1	seq
.....aaaugcacuaucugguacgaca.....	19	0	seq
.....aaaugcacuaucugguacgacau.....	3	0	seq
.....aaugcacuaucugguacg.....	1	0	seq
.....aaugcacuaucugguacga.....	1	0	seq
.....aaugcacuaucugguacgac.....	5	0	seq
.....aaugcacuaucugguacgaA.....	1	1	seq
.....Uaugcacuaucugguacgac.....	1	1	seq
.....aaCgcacuaucugguacgaca.....	1	1	seq
.....aaugcacuaucugguacgaca.....	13	0	seq
.....augcacuaucugguacgac.....	1	0	seq
.....augcacuaucugguacgaca.....	1	0	seq
.....ugcacuaucugguacgUca.....	1	1	seq
.....ugcacuaucugguacgaca.....	2	0	seq
.....gcacuaucugguacgacaA.....	1	1	seq