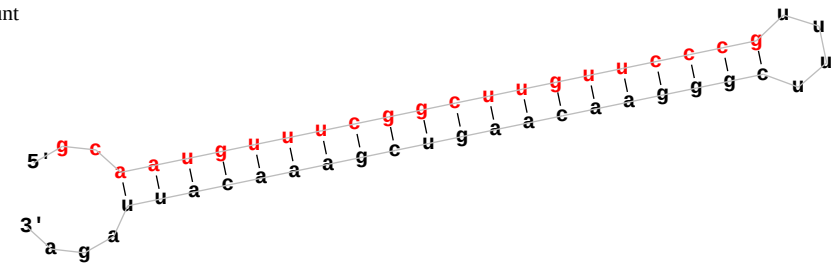
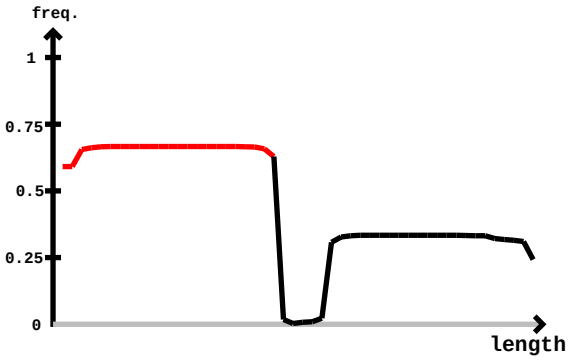
miRBase precursor : Montipora_capitata_HIv3___Scaffold_2_96328
Total read count : 1064
Montipora_capitata_HIv3___Scaffold_2_96328 read count : 709
remaining reads : 355





Montipora_capitata_HIv3___Scaffold_2_96328

5'- gcaauguuucggcuuguucccguuuucgggaacaagucgaaacauuaga -3' exp

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

Acaauguuucggcuuguuccc.....	4	1	s60
AcaauguuucggcuuguucccU.....	4	2	s60
AcCCuguuucggcuuguucccg.....	1	3	s60
AcaauguCucggcuuguucccg.....	1	2	s60
Acaauguuucggcuuguucccg.....	138	1	s60
AcCauguuucggcuuguucccg.....	2	2	s60
AcaCuguuucggcuuguucccg.....	1	2	s60
AcaauguuucggcuuguucccC.....	2	2	s60
AUaauguuucggcuuguucccg.....	2	2	s60
Acaauguuucggcuuguucccgu.....	1	1	s60
.Uaauguuucggcuuguucccg.....	1	1	s60
.caauguuucggcuuguucccg.....	16	0	s60
.cCauguuucggcuuguucccU.....	1	2	s60
..aauguuucggcuuguucccg.....	2	0	s60
...auguuucggcuuguucccg.....	1	0	s60
.....Aucgggaacaagucgaaacauu...	1	1	s60
.....Aucgggaacaagucgaaacauuag.	1	1	s60
.....AggAaaGaagucgaaaca.....	1	3	s60
.....Agggaacaagucgaaacau....	1	1	s60
.....Agggaacaagucgaaacauuaga	1	1	s60
.....Aggaacaagucgaaacauuag.	10	1	s60
.....AggaacaagucgaaacauuaC.	1	2	s60
.....AggaacCagucgaaCcauuaga	1	3	s60
.....AggaCcaagucgaaacauuaga	3	2	s60
.....Aggaacaagucgaaacauuaga	50	1	s60
.....AggaacaagucgaaacauuagU	3	2	s60
.....ggaacaagucgaaacauuag.	1	0	s60
.....ggaacaagucgaaacauuaCa	1	1	s60
.....ggaacaagucgaaacauuaga	4	0	s60
.....gaacaagucgaaacauuaga	2	0	s60
Acaauguuucggcuuguuccc.....	1	1	s13
Acaauguuucggcuuguucccg.....	2	1	s13
...auCuuucggGuuguuccc.....	1	2	s13
.....AuucgggaaUaagGcgaa.....	1	3	s13

Montipora_capitata_HIv3___Scaffold_2_96328

gcaauguuucggcuuguucccguuuucgggaacaagucgaaacauuaga

.....AggAaaGaagucgaaaca.....	1	3	s13
.....Aggaacaagucgaaacauuag.	4	1	s13
.....ggaCcaagucgaaacauuag.	1	1	s13
Acaauguuucggcuuguuc.....	1	1	s72
Acaauguuucggcuuguucc.....	2	1	s72
Acaauguuucggcuuguuccc.....	5	1	s72
AcCauguuucggcuuguucccg.....	1	2	s72
AcaCuguuucggcuuguucccg.....	2	2	s72
AcaauguuucggcuuguucccU.....	4	2	s72
Acaauguuucggcuuguucccg.....	135	1	s72
AUaauguuucggcuuguucccg.....	6	2	s72
AcaauguuucggcuugCucccg.....	1	2	s72
AcaauguuucggcuuguucccC.....	2	2	s72
Acaauguuucggcuuguucccgu.....	5	1	s72
.caauguuucggcuuguucccU.....	2	1	s72
.caauguuucggcuuguucccg.....	16	0	s72
.caCuguuucggcuuguucccg.....	1	1	s72
.Aaauguuucggcuuguucccg.....	1	1	s72
.caauguuucggcuuguucccgu.....	1	0	s72
..aauguuucggcuuguucccg.....	1	0	s72
..aauguuucggcuuguucccgu.....	1	0	s72
.....Acgggaacaagucgaaaca.....	1	1	s72
.....AggAaaGaagucgaaaca.....	1	3	s72
.....Aggaacaagucgaaacauuag.	6	1	s72
.....AggCacaagucgaaacauuaga	1	2	s72
.....AggaacaagucgaaCcauuaga	1	2	s72
.....Aggaacaagucgaaacauuaga	28	1	s72
.....ggaacaagucgaaacauuaga	2	0	s72
Acaauguuucggcuuguuccc.....	2	1	s09
.....AucgggaacaagucgaaacauuaC.	1	2	s09
.....AggAaaGaagucgaaaca.....	1	3	s09
.....AggaacaagucgaaacauuaC.	1	2	s09
.....AggCacaagucgaaacauuaC.	1	3	s09
.....Aggaacaagucgaaacauuag.	7	1	s09
Acaauguuucggcuuguucc.....	2	1	s85
Acaauguuucggcuuguuccc.....	5	1	s85
AUaauguuucggcuuguuccc.....	1	2	s85
Acaauguuucggcuuguucccg.....	74	1	s85
AcaauguuucggcuuguucccU.....	1	2	s85
AcCauguuucggcuuguucccg.....	1	2	s85
AUaauguuucggcuuguucccg.....	4	2	s85
Acaauguuucggcuuguucccgu.....	2	1	s85
.caauguuucggcuuguucccg.....	4	0	s85
.....Agggaacaagucgaaaca.....	1	1	s85
.....AggAaaGaagucgaaaca.....	2	3	s85
.....Aggaacaagucgaaacau.....	2	1	s85
.....Aggaacaagucgaaacauuag.	7	1	s85
.....AggCacaagucgaaacauuag.	1	2	s85
.....AggaacaagucgaaacauuaC.	1	2	s85
.....AggaacaCgucgaaacauuaga	1	2	s85
.....AggaacaagucgaaaAauuaga	1	2	s85
.....AggaacaagucgaaacaGuaga	1	2	s85
.....Aggaacaagucgaaacauuaga	39	1	s85
.....ggaacaagucgaaacauuag.	1	0	s85
Acaauguuucggcuuguuccc.....	2	1	s23
AcaauguuucggcuuguucccA.....	1	2	s23
Acaauguuucggcuuguucccg.....	2	1	s23
.....AgggaacaagucgaaacauuaCC	1	3	s23
.....Aggaacaagucgaaacauuag.	2	1	s23
.....Aggaacaagucgaaacauuaga	8	1	s23
Acaauguuucggcuuguu.....	1	1	s35
Acaauguuucggcuuguuccc.....	4	1	s35
AcaauguuucggcuuguucccC.....	1	2	s35
Acaauguuucggcuuguuccag.....	1	2	s35
Acaauguuucggcuuguucccg.....	77	1	s35

Montipora_capitata_HIv3___Scaffold_2_96328

gcaauguuucggcuuguucccguuuucgggaacaagucgaaacauuaga

AcaauguuCcggcuuguucccg.....	1	2	s35
gcaauguuucggcuuguucccgu.....	1	0	s35
Acaauguuucggcuuguucccgu.....	2	1	s35
.caauguuucggcuuguucccg.....	9	0	s35
.caauguuucggcuuguucccgu.....	1	0	s35
..aGuCuuucggGuuguucc.....	1	3	s35
..aauguuucggcuuguucccg.....	1	0	s35
...auguuucggcuuguucccg.....	1	0	s35
.....Auuuucgggaacaagucgaaacauuaga	1	1	s35
.....uucgggaacaagucgaaa.....	1	0	s35
.....Aucgggaacaagucgaaacauua..	1	1	s35
.....ucgggaacaagucgaaacauuaC.	1	1	s35
.....AggAaaGaagucgaaaca.....	1	3	s35
.....cgggaacaagucgaaacauuaC.	1	1	s35
.....Aggaacaagucgaaacauu...	1	1	s35
.....Aggaacaagucgaaacauua..	2	1	s35
.....AggaacaagucgaaacauuG..	1	2	s35
.....Aggaacaagucgaaacauuag.	8	1	s35
.....AggaCcaagucgaaacauuaga	1	2	s35
.....Aggaacaagucgaaacauuaga	53	1	s35
.....AggaacaagucgaaacauuaCU	1	3	s35
.....AggaacaagucgaaacauuaCC	1	3	s35
.....AggaacaagucgaaacaCuaga	1	2	s35
.....AggaacaagucgaaacauuaCa	3	2	s35
.....ggaacaagucgCCCcauuag.	1	3	s35
.....Agaacaagucgaaacauuag.	1	1	s35
.....ggaacaagucgaaacauuaCa	1	1	s35
.....ggaacaagucgaaacauuaga	5	0	s35
.....gaacaagucgaaacauuaga	1	0	s35

Acaauguuucggcuuguucc.....	2	1	s52
Acaauguuucggcuuguuccc.....	7	1	s52
AUaauguuucggcuuguucccg.....	7	2	s52
AcaCuguuucggcuuguucccg.....	1	2	s52
Acaauguuucggcuuguucccg.....	96	1	s52
AcaauguuucggcuuguucccC.....	2	2	s52
AcaauguuucggcuuguucccU.....	3	2	s52
Acaauguuucggcuuguucccgu.....	2	1	s52
AcaauguuucggcuuguucccCu.....	1	2	s52
.caauguuucggcuuguuccc.....	1	0	s52
.caauguuucggcuuguucccg.....	13	0	s52
.caCuguuucggcuuguucccg.....	1	1	s52
..aauguuucggcuuguucccg.....	1	0	s52
...auguuucggcuuguucccg.....	1	0	s52
....uguuucggcuuguucccg.....	1	0	s52
.....Auuuucgggaacaagucgaaacauuaga	1	1	s52
.....AggAaaGaagucgaaaca.....	2	3	s52
.....Aggaacaagucgaaacau....	1	1	s52
.....Aggaacaagucgaaacauu...	1	1	s52
.....Aggaacaagucgaaacauua..	1	1	s52
.....Aggaacaagucgaaacauuag.	12	1	s52
.....Aggaacaagucgaaacauuaga	36	1	s52
.....ggaacaagucgaaacauuag.	2	0	s52
.....ggaacaagucgaaacauuaga	1	0	s52
.....gaacaagucgaaacauuaga	2	0	s52
.....aacaagucgaaacauuaga	2	0	s52