Provisional ID	:	Pocillopora_meandrina_HIv1___Sc0000002_135259
Score total	:	4.6
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	1.3
Score for randfold	:	1.6
Score for cons. seed	:	3
Total read count	:	258
Mature read count	:	258
Loop read count	:	0
Star read count	:	0

	reads	mm	sample
5'- ucagagcgauggacgaagggcuaacgcccugacgaagggcaaacgcucgaaacgucagcuuuuuuacccuuuaccguggcuaauuuacguuuucaacuuaguuguuaacac -3'			exp
....(((.(((((..(((((((.(((....(((((((.(((((....)))).....))))))).....)))))))).)))).))).......(((..(((...)))..)))..			
ucagagcgauggacgaagggcG.	1	1	seq
.cagagcgauggacgaagggcuaacgcccG.	3	1	seq
........................cugacgaagggcaaacgcucgaaacg.	2	0	seq
..................................Ccucgaaacgucagcuuuuu.	1	1	seq
..................................gcucgaaacgucagcuuuuuu.	2	0	seq
..................................cucgaaacgucagcuuuuu.	2	0	seq
..................................cucgaaacgucagcuuuuuu.	8	0	seq
...................................ucgaaacgucagcuuuuu.	1	0	seq
...................................ucgaaacgucagcuuuuuuacA.	1	1	seq
...................................ucgaaacgucagcuuuuuuacccu.	1	0	seq
......................................gaaacgucagcuuuuuuacccu.	1	0	seq
....aaacgucagcuuuuuuac.	2	0	seq
....aaacgucagcuuuuuuacA.	1	1	seq
....aaacgucagcuuuuuuacc.	3	0	seq
....aaacgucagcuuuCuuaccc.	1	1	seq
....aaacgucagcuuuuuuaccG.	3	1	seq
....aaacgucagcuuuuuuacAc.	2	1	seq
....aCacgucagcuuuuuuaccc.	1	1	seq
....aaacgucagcuuuuuuaccc.	55	0	seq
....aaacgucagcuuuCuaccc.	1	1	seq
....aaacgucagcuuuuuuacAcu.	3	1	seq
....aaaAgucagcuuuuuuacccu.	1	1	seq
....aaacgucagcuuuuuuacccG.	5	1	seq
....aaacgucagcuuuuuuacccu.	39	0	seq
....aaacgucagcuuuuuuacAcuu.	2	1	seq
....aaacgucagcuuuuuuacccGu.	1	1	seq
....aaacgucagcuuuuuuacccuA.	1	1	seq
....aGacgucagcuuuuuuacccuu.	1	1	seq
....aaacgucagcuuuuuuacccuG.	1	1	seq
....aaacgucagcuuuuuuacccuu.	3	0	seq
....aaacgucagcuuuuuuacccuuu.	1	0	seq
....aaacgucagcuuuuuuacccuuuac.	1	0	seq
....aacgucagcuuuuuuacA.	1	1	seq
....aacgucagcuuuuuuacc.	1	0	seq

ucagagcgauggacgaagggcuaacgcccgacgaagggcaaacgcucgaaacgucagcuuuuuuacccuuuaccguggcuaauuuacguuuucaacuuaguuguuaacac

....aacgucCgcuuuuuuaccc..	1	1	seq
....aacgucagcuuuuuuaccc..	22	0	seq
....aGcgucagcuuuuuuaccc...	1	1	seq
....aacgucagcuuuuuuaccG..	2	1	seq
....aacgucaCcuuuuuuacccu...	1	1	seq
....Cacgucagcuuuuuuacccu...	3	1	seq
....aacguUagcuuuuuuacccu...	2	1	seq
....aacgucagcuuuuuuacccu...	36	0	seq
....aacgucagcuuuuuuacccG...	6	1	seq
....aacgucagcuuuuCuacccu...	1	1	seq
....aacgucagcuuuuuuacccuA..	1	1	seq
....aacguUagcuuuuuuacccuu..	1	1	seq
....aacgucagcuuuuuuacccuu..	9	0	seq
....aGcgucagcuuuuuuacccuu..	1	1	seq
....aaUgucagcuuuuuuacccuuuac.....................................	2	1	seq
....acgucagcuuuuuuaccc..	5	0	seq
....acgucagcuuuuuuacccG...	2	1	seq
....acguUagcuuuuuuacccu...	1	1	seq
....acgucagcuuuuuuacAcu...	1	1	seq
....acgucagcuuuuuuacccu...	5	0	seq
....acgucagcuuuuuuacccuu..	8	0	seq
....acgucagcuuuuuuacccuA..	3	1	seq
....Gcgucagcuuuuuuacccuu.......................................	1	1	seq
....acgucagcuuuuuuacccuG..	1	1	seq
....acgucagcuuuuuuacccuuu.......................................	3	0	seq
....cgucagcuuuuuuacccuu.......................................	2	0	seq
....gucagcuuuuuuacccuuuac....................................	1	0	seq
....ucagcuuuuuuacccuuu.....................................	2	0	seq
....ucagcuuuuuuacccuuua....................................	2	0	seq
....ucagcuuuuuuacccuuuac...................................	1	0	seq
....uuaccguggcuaauuuacguu..........................	1	0	seq
....ccguggcuGauuuacguuuucaacuua..................	1	1	seq