Provisional ID	: Pocillopora_meandrina_HIv1___Sc0000033_892145
Score total	: 623.5
Score for star read(s)	: 3.9
Score for read counts	: 615
Score for mfe	: 0.1
Score for randfold	: 1.6
Score for cons. seed	: 3
Total read count	: 1218
Mature read count	: 1212
Loop read count	: 0
Star read count	: 6


```
5'- uuaguuaguuuagaguaaaaggcuucuuccagcguaugaacggagaaaaguuucaagcacauguuuaucauuugcuaguugcuuuugucccguuuguucauuggauuu  -3'    obs
    uuaguuaguuuagaguaaaaggcuucuuccagcguaugaacggagaaaaguuucaagcacauguuuaucauuugcuaguugcuuuugucccguuuguucauuggauuu         exp
    ..........(((((.....))))..(((((.(.(..(((((((.(((((((..(.(((((.(((.....))).)))))).)..))))))).).)))))..).).)))))...   reads   mm    sample
```

	reads	mm	sample
....ggagaaaaguuucaagcacaug...	5	0	seq
....ggagaaaaguuucaagcacauA...	1	1	seq
....Cuuugcuaguugcuuuugucccgu...............................	1	1	seq
....uuugcuaguugcuuuugu.................................	1	0	seq
....uuugcuaguugcuuuuguc................................	1	0	seq
....uuugcuaguugcuuuuguU................................	2	1	seq
....uuugcuaguugcGuuuguc................................	2	1	seq
....uuugcuaguugcuuuugucc...............................	13	0	seq
....uuugcuaguugcCuuugucc...............................	1	1	seq
....uuugcuaguugcuuuugucA...............................	6	1	seq
....uuugcuaguugcuuuugucG...............................	1	1	seq
....uuugcuaguugcuuuugucU...............................	7	1	seq
....uuugcuaguugcuuuuguccG..............................	4	1	seq
....uuugcuaguugcuuuuguccU..............................	7	1	seq
....uuugcuaguugcuuuuguccc..............................	13	0	seq
....uuugcuaguugcuuuuguccA..............................	10	1	seq
....uuugcuaguugcCuuuguccc..............................	1	1	seq
....uuugcuaguugcuuuugucccA.............................	10	1	seq
....uuugcuaguugcuuuuguccAg.............................	2	1	seq
....uuugcuaguugcuuuuguccccg............................	7	0	seq
....uuugcuaguugcuuuuguccccU............................	1	1	seq
....uuugcuaguugcuuuCgucccgu...........................	1	1	seq
....uuugcuaguugcuuuugucAcgu...........................	1	1	seq
....uuugcuaguugcCuuugucccgu...........................	1	1	seq
....uuugcuaguugcuuuuguccccAu..........................	10	1	seq
....uuugcuaguugcuuuAgucccgu...........................	1	1	seq
....uAugcuaguugcuuuuguccccgu..........................	1	1	seq
....uuugcCaguugcuuuuguccccgu..........................	1	1	seq
....uuugcuaguugcuuuuguccccgG..........................	18	1	seq
....uuugcuaguCgcuuuuguccccgu..........................	1	1	seq
....uuugcuaguugcuuuugucccUgu..........................	8	1	seq
....uuugcuaguugcuuuuguccccgC..........................	74	1	seq
....uuugcuaguugcuuuuguccccgu..........................	279	0	seq

uuaguuaguuuagaguaaaaggcuucuuccagcguaugaacggagaaaaguuucaagcacauguuuaucauuugcuaguugcuuuugucccguuuguucauuggauuu

....uuuAcuaguugcuuuugucccgu..............	1	1	seq
....uuugcuGguugcuuuugucccgu..............	1	1	seq
....uuugcuaguugcuuuugucccCu..............	3	1	seq
....Cuugcuaguugcuuuugucccgu..............	7	1	seq
....uuugcuagCugcuuuugucccgu..............	2	1	seq
....uuugcuaguugcuuuugucccgA..............	30	1	seq
....uuugcuaguugcuuuugucccgAu..............	16	1	seq
....uuugcuaguugcuuuugucccgGu..............	26	1	seq
....uuugcuaguugcuuuuguccAguu..............	1	1	seq
....uuugcuaguugcuuuCgucccguu..............	1	1	seq
....uuugcuaguugcCuuugucccguu..............	5	1	seq
....uuugcuaguugcuuuugucccguG..............	28	1	seq
....uCugcuaguugcuuuugucccguu..............	1	1	seq
....uuugcuaguugcuuuugucccguC..............	99	1	seq
....uuugcuaguugcuuuugCcccguu..............	2	1	seq
....uuugcuaguugcuuuugucccguA..............	61	1	seq
....uuugcuaguuAcuuuugucccguu..............	1	1	seq
....uuugcuaguugcuuuugucccguu..............	374	0	seq
....Cuugcuaguugcuuuugucccguu..............	6	1	seq
....uuugcuaguugcuuuugucccgCu..............	5	1	seq
....uuugcuaguugcuCuugucccguu..............	3	1	seq
....uuugcuaguugcuuuugucccguuu..............	25	0	seq
....uuugcuaguugcuuuCgucccguuu..............	2	1	seq
....uuugcuGguugcuuuugucccguuu..............	1	1	seq
....uuugcuaguugAuuuugucccguuu..............	1	1	seq
....uuugUuaguugcuuuugucccguuu..............	2	1	seq
....uuugcuaguugcuuuugucccguuC..............	5	1	seq
....uuugcuaguugcuuuugucccguuA..............	4	1	seq
....uuugcuaguugcuuuugucccgCuu..............	1	1	seq
....uuugcuaguugcuuuugucccguuU..............	5	1	seq
....uuugcuaguugcuuuugucccguuuA..............	2	1	seq
....uuugcuaguugcuuuugucccguuuUu..............	1	1	seq
....uugcuaguugcuuuugucccguu..............	1	0	seq
....ugcuaguugcuuuugucccAu..............	1	1	seq
....cuaguugcuuuugucccguuA..............	1	1	seq
....uaguugcuuuugucccguu..............	1	0	seq