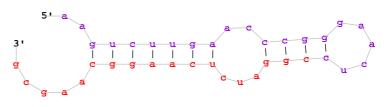
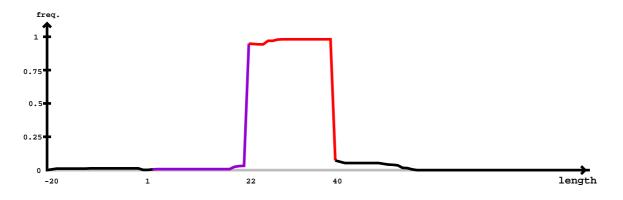
Provisional ID : Montipora_capitata_HIv3___Scaffold_11_652458

242.1 Score total Score for star read(s) 3.9 239.2 Score for read counts Score for mfe 1.1 Score for randfold -2.2 Score for cons. seed Total read count 481 478 Mature read count Loop read count 0 Star read count 3





5'- auucaacgaguggucagaacaagucuugaacccgggaacuccggaucucaaggcaagcguccuaaccacugggccgcgcacugccuccuacaggcccaauuauagguga

Star Mature

auucaacgaguggucagaacaagucuugaacccggga <mark>acuccggaucucaaggcaagc</mark> guccuaaccacugggccgccgcacugccuccuacaggcccaauuauagguga	exp		
((((((((((((((((((((((((((((((((((reads	mm	sample
.Acggaucucaaggcaagc.	99	1	s52
.Acggaucucaaggcaagcg.	1	1	s52
.Acggaucucaaggcaagcguccuaacca	2	1	s52
aucucaaggcaagcguccuaaccacugg	5	0	s52
Aucaaggcaagcguccuaaccacugggc	2	1	s52
Aucaaggcaagcguccuaaccacugggcc	2	1	s52
Acaaggcaagcguccuaaccacugggc	1	1	s52
Acggaucucaaggcaagc	35	1	s72
Acggaucucaaggcaagcguccuaaccacug	1	1	s72
aucucaaggcaagcguccuaaccacugg	1	0	s72
Aucaaggcaagcguccuaaccacugggcc	1	1	s72
aagucuugaacccgggaacuG	2	1	s23
cuccggaucucaaggcCagc.	1	1	s23
.Acggaucucaaggcaagc.	79	1	s23
Acggaucucaaggcaagcg	2	1	s23
auucaacgaguggucaga	2	0	s13
agucuugaacccgggaaGucagucuugaacccgggaaGuc	1	1	s13
.acuceggaucucaaggcaagc.	3	0	s13
	55	1	s13
	2	1	s13
.Acggaucucaaggcaagcguccuaaccacu	1	1	s13
aucucaaggcaagcguccuaaccacugg	1	0	s13
auucaacgaguggucaga	1	0	s60
.acuceggaucucaaggcaagc.	1	0	s 60
cuccggaucucaaggcaagc	1	0	s 60
Acggaucucaaggcaagc	51	1	s60
aucucaaggcaagcguccuaaccac	1	0	s 60
aucucaaggcaagcguccuaaccacugg	2	0	s 60
acuccggaucucaaggcaagc	6	0	s09

Star	Mature
olai	mature

auucaacgaguggucagaacaagucuugaacccgggaacuc <mark>cggaucucaaggcaagcg</mark> uccuaaccacugggccgccgcacugccuccuacaggcccaauuauagguga			
Acggaucucaaggcaagc	51	1	s09
Acggaucucaaggcaagcg	1	1	s09
Acggaucucaaggcaagcgu	1	1	s09
Acggaucucaaggcaagcguccuaaccac	1	1	s09
.aucucaaggcaagcguccuaaccacugggc	2	0	s09
auucaacgaguggucaga	2	0	s35
Aaguggucagaacaagucuugaacccggga	1	1	s 35
.cuccggaucucaaggcaagc.	1	0	s 35
Accggaucucaaggcaagcgu.	1	1	s35
Acggaucucaaggcaagc	78	1	s 35
ceggaucucaaggcaagc.	2	0	s 35
	2	1	s 35
Acggaucucaaggcaagcguccuaacca	1	1	s 35
Acggaucucaaggcaagcguccuaaccac	1	1	s 35
Acggaucucaaggcaagcguccuaaccacug	1	1	s 35
.aucucaaggcaagcguccuaaccacugg	1	0	s 35
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auucaacgaguggucaga	1	0	s85
Acggaucucaaggcaagc	15	1	s 85