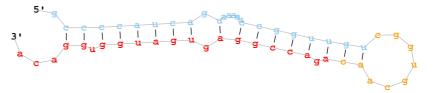
Provisional ID : Montipora_capitata_HIv3___Scaffold_13_836552

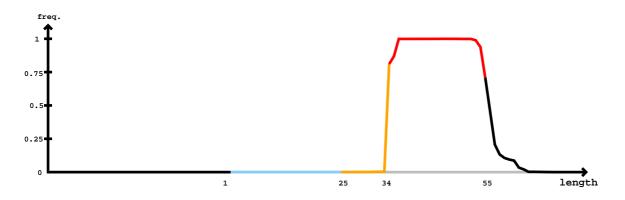
Score total : 1.7
Score for star read(s) : -1.3
Score for read counts : 0
Score for mfe : 1.4
Score for randfold : 1.6
Score for cons. seed :



cagaccggagugaugguggacagagacgguagggggacaccc

Total read count : 27747
Mature read count : 27747
Loop read count : 0
Star read count : 0

5'- aggaaggugacccuugcacuuagaugagauuuuacu



(((((((((((((((((((((((((((((((((((reads	mm	sample
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.Augcaacagaacggaguga	1	1	s13
	1	1	s13
	1	1	s13
aacGgacoggagugauggugaca	3	1	s13
acGgaccggagugaugguca	2	1	s13
agacoggagugauggugg	9	0	s13
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Star Mature			
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Star Mature			
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Star Mature			
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. Aaccggagugaugga.	20	1	s 35
. Aaccggagugaugguc	23	1	s 35
accggagugaugguc	78	0	s 35
accgg/Ggugaugguggac	1	1	s 35
augguggacGgagacgguagg	1	1	s 35
.augguggacGgagacgguagggg.	1	1	s 35
	_		
accggagugaugguggac	2	0	s 86
_	_	_	
Aucggugcaacagaccggagugauggugga	1	1	s52
Agcaacagaccggagugaugg	1	1	s52
Agcaacagaccggagugauggugga	1	1	s52
acGgaccggagugaugguggaca	2	1	s52
agaccggagugauggugg	12	0	s52
agaccggagugauggugga	77	0	s52
	1	1	s52
agCccggagugauggugga	1	1	s52
agCccggagugaugguggacagCccggagugaugguggac	1	1	s52
agaccggagugaugguggac	138	0	s52
agaccggagugauggucc.	1	1	s52
agaccggagugaugguggacU	1	1	s52
agaccggagugaugguggacG.	945	1	s52
agaccggagugaugguggacGgagaccggagugaugguggacGg	360	1	s52
agaccggagugaugguggacGga	183	1	s52
agaccggagugaugguggacGgag	70	1	s52
agaccggagugauggucggacUgaga.	1	1	s52
agaccggagugaugguggacGgaga	29	1	s52
agaccggagugauggucggacGgagac	20	1	s52
agaccggagugaugguggacGgagacg	112	1	s52
agaccggagugaugguggacGgagacggagaccggagugaugguggacGgagacgg	42	1	s52
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agaccggagugguggacGgagacgguag	2	1	s52
agaccggagugguggacGgagacgguaggagaccggagugaugguggacGgagacgguagg	2	1	s52
agaccggagugguggacGgagacgguagggg	1	1	s52
Aaccggagugauggugga	34	1	s52
Aaccggagugaugguggac	98	1	s52
Aaccggagugaugguggacagagacg	1	1	s52
accggagugCugguggac	2	1	s52
accggagugaugguggac	274	0	s52

aggaaggugacccuugcacuuagaugagauuuuacugecccaucaguaaaaucegguuguc <mark>ggugcaacagaccggagugaugguggaca</mark> gagacgguaggggacacec			
accggagugauggaA	3	1	s52
accggagugauggukgac	1	1	s52
accggCgugaugguggac	1	1	s52
	1	1	s52
accgAagugaugguggac	1	1	s52
accggagugauggacaaccggagugauggaca.	2	0	s52
accggagugauggucgacU	1	1	s52
accggagugauggucGgagacgguag	1	1	s52
accggagugauggucggacggagagg	3	1	s52
Augaugguggacagagacgg	1	1	s52
augguggacGgagacgguagggg	1	1	s52