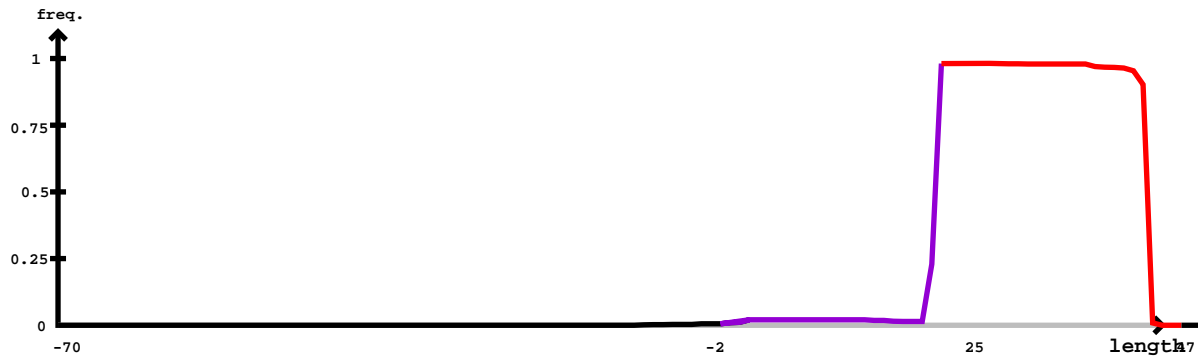
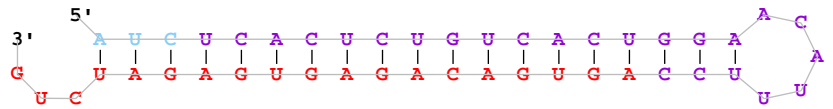


Provisional ID : NC_058070.1_Acropora_millepora_isolate_JS-1_chromosome_5_Amil_v2.1_whole_genome_shotgun_sequence_204549
Score total : 0.1
Score for star read(s) : -1.3
Score for read counts : 0
Score for mfe : 2
Score for randfold : 0
Score for cons. seed : -0.6
Total read count : 1254
Mature read count : 1235
Loop read count : 0
Star read count : 19



		Star		Mature					
5'						-3'	obs		
caacaaucacugccgauuuuguacagcgccaaaucaacacagaguuagacaccaccuuguacccagauucacucucugucacuggaacauuuccagugacagagugagauucug							exp		
caacaaucacugccgauuuuguacagcgccaaaucaacacagaguuagacaccaccuuguacccagauucacucacucugucacuggaacauuuccagugacagagugagauucug									
.....(((.....))).....(((.....))).....((((((((((((((((((((.....))))))))))))))))))						reads	mm	sample	
.....guaccagaucucacucugCcacu.....						2	1	seq	
.....uaccagaucucacucugucacu.....						1	0	seq	
.....cccagaucucacucugucacugU.....						1	1	seq	
.....agaucucacucugucacugg.....						3	0	seq	
.....ucucacucugucacugga.....						1	0	seq	
.....ucucacucugucacuggaacauuucc.....						1	0	seq	
.....ucucacucugucacuggaacauuucU.....						2	1	seq	
.....ucucacucugucacuggaacauuuccaguga.....						1	0	seq	
.....cucacucugucacuggaacauuucU.....						1	1	seq	
.....cucacucugucacuggaacauuuccaC.....						1	1	seq	
.....ucacucugucacuggaacauuucc.....						10	0	seq	
.....ucacucugucacuggaacauuucU.....						1	1	seq	
.....ucacucugucacuggaacauuuccagu.....						1	0	seq	
.....Cuuccagugacagagug.....						1	1	seq	
.....uuuccagugacagagug.....						11	0	seq	
.....uuuccagugacagagugU.....						2	1	seq	
.....uuuccagugacagagugaA.....						1	1	seq	
.....uuuccagugacagagugaga.....						1	0	seq	
.....uuuccagugacagagugagU.....						1	1	seq	
.....uuuccagugacagagugagU.....						2	1	seq	
.....uuuccagugacagagugagau.....						3	0	seq	
.....uuuccagugacagagugagaG.....						2	1	seq	
.....uuuccagugacagagugagaA.....						1	1	seq	
.....uuuccagugacagagugaAauc.....						2	1	seq	
.....uuuccagugacagagugagaAc.....						1	1	seq	
.....uuuccagugacagagugagauG.....						1	1	seq	
.....uuuccagugacagagugagauA.....						4	1	seq	
.....uuuccagugacagagugagauU.....						44	1	seq	
.....uuuccagugacagagugagauC.....						1	1	seq	
.....uuuccagugacagagugagauC.....						19	1	seq	
.....uuuccagugacagagugagauC.....						14	1	seq	
.....uuuccagugacagagugagauC.....						2	1	seq	
.....uuuccagugacagagugagauU.....						2	1	seq	

Star

Mature

caacaaucaugccgauuuuguaacagcgccaaaucaacacgaguuagacaccaccuuguaccagaucu	caacucugucacuggaacaauuccagugacagagugagau	cug
.....uuuccagugacagagugagGucu.	2	1seq
.....uuuccagugacagagugagaCcu.	2	1seq
.....uuuccagugacagagugagau.	151	0seq
.....uuuccagugacagagugagauU	1	1seq
.....uuccagugacagagugG.....	1	1seq
.....uuccagugacagagugaga....	1	0seq
.....uuccagugacagagugagau..	4	0seq
.....uuccagugacagagugagUu...	1	1seq
.....uuccagugacagagugGgau.	2	1seq
.....uuccagugacagagugagauA..	2	1seq
.....uuccagugacagagugagauU..	9	1seq
.....uuccagugacagagugagauUu.	6	1seq
.....uuccUgugacagagugagau.	1	1seq
.....uuccagugacagagugagaGcu.	1	1seq
.....uuccagugacagagugagauC.	109	1seq
.....uuccagugacagagCgagau.	2	1seq
.....uuccagugacagagugagGucu.	2	1seq
.....uuccagugacUgagugagau.	1	1seq
.....uuccagugacagagugagaCcu.	4	1seq
.....uuccagugacagagugagau.	643	0seq
.....uuccagugacagagugagauC.	144	1seq
.....uuccagugacagagugGgau.	5	1seq
.....uuccagugacagagugagauC	1	1seq
.....uuccagugacagagugagauCg	1	1seq
.....uuccagugacagagugagauU	4	1seq
.....uuccagugacagagugagauA	4	1seq
.....uccagugacagagugagauC.	1	1seq
.....uccagugacagagugagau.	11	0seq
.....ccagugacagagugagau.	3	0seq
.....cagugacagagugagauC.	1	1seq