

Provisional ID : NC\_058067.1\_Acropora\_millepora\_isolate\_JS-1\_chromosome\_2\_Amil\_v2.1\_whole\_genome\_shotgun\_sequence\_92488

Score total : 0.2

Score for star read(s) : -1.3

Score for read counts : 0

Score for mfe : 2.1

Score for randfold : 0

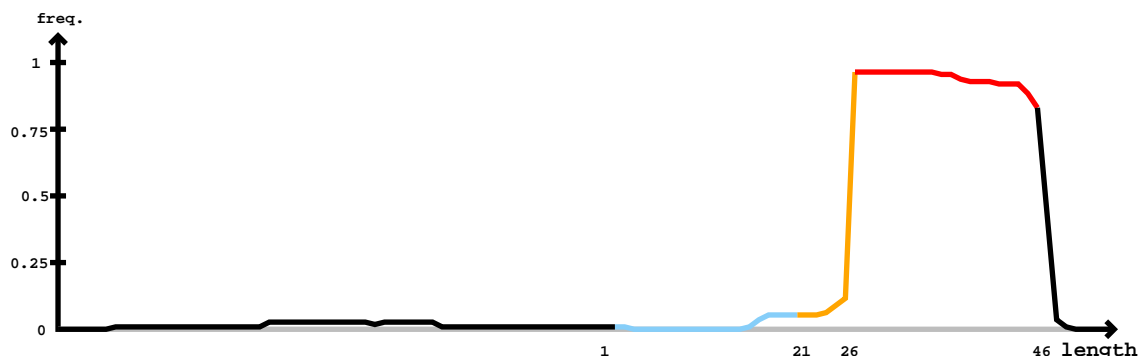
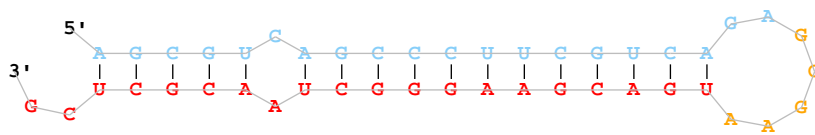
Score for cons. seed : -0.6

Total read count : 101

Mature read count : 101

Loop read count : 0

Star read count : 0



		Star		Mature					
5'						-3'	exp		
acgaguugauaaagguggaauuaccaccgugaaagaacuggaagcugacguuccgagcugacagccuucgucagagcgaaugacgaagggcuaacgcucgaaauucg						reads	mm	sample	
.....(((((((.....)))))).....(((((((.....)))))).....(((((((.....)))))).....(((((((.....)))))).....(((((((.....)))))).....						1	1	seq	
.....guugauaaagguAgaauuaccaccgug.....						2	0	seq	
.....uuaccaccgugaaagaac.....						1	1	seq	
.....aagaacuggaagcugacguGccgag.....						1	1	seq	
.....gucagagcgaaGgacgaagggcuaacgcu.....						1	1	seq	
.....ucagagcgaaugacgaagA.....						1	1	seq	
.....ucagagcgaaugacgaagggc.....						2	0	seq	
.....cagagcAaaugacgaagggcu.....						1	1	seq	
.....cagagcgaaGgacgaagggcuaac.....						1	1	seq	
.....gaaugacgaagggcuaacgcucgU.....						1	1	seq	
.....aaugacAaagggcuaacgcu.....						1	1	seq	
.....aaugacgaagggcuaacgcucg.....						1	0	seq	
.....aaGgacgaagggcuaacgcucgaaa.....						1	1	seq	
.....aGgacgaagggcuaacgcucga.....						1	1	seq	
.....aGgacgaagggcuaacgcucgaa.....						2	1	seq	
.....ugacgaagggcuaacgcA.....						1	1	seq	
.....ugacgaagggcuaacgcu.....						1	0	seq	
.....ugacgaagggcuaacgcuc.....						5	0	seq	
.....ugacgaagggcuaacgcCc.....						1	1	seq	
.....ugacgaagggcuaacgcucg.....						68	0	seq	
.....ugacgaGgggcuaacgcucg.....						2	1	seq	
.....ugacgaagggcuaacgcCc.....						1	1	seq	
.....ugacgaagggcGacgcucg.....						5	1	seq	
.....ugacgaagggcuaacgcucga.....						2	0	seq	
.....ugacgaagggcuaacgcucgU.....						8	1	seq	
.....ugacgaagggcuaacgcucgaU.....						1	1	seq	