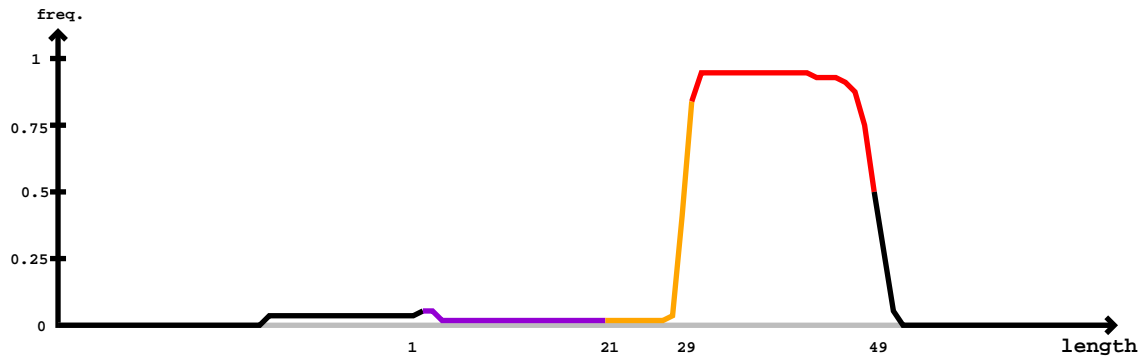
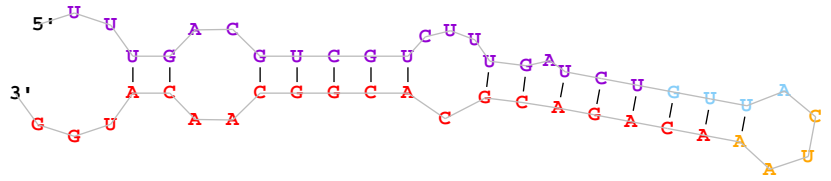


Provisional ID : NC\_058076.1\_Acropora\_millepora\_isolate\_JS-1\_chromosome\_11\_Amil\_v2.1\_whole\_genome\_shotgun\_sequence\_433115  
Score total : 21.8  
Score for star read(s) : 3.9  
Score for read counts : 21  
Score for mfe : -2.4  
Score for randfold : 0  
Score for cons. seed : -0.6  
Total read count : 53  
Mature read count : 52  
Loop read count : 0  
Star read count : 1



### Star Mature

5' -	obs	exp	reads	mm	sample
aacgccucgugggcacuaauuugcucuuaacauuuuugacgucgucuuuugaucuguuuacuaaaacagacgcacggcaacauggaaccuaauuugguaaaauuugagcc	-3'				
aacgccucgugggcacuaauuugcucuuaacauuuuugacgucgucuuuugaucuguuuacuaaaacagacgcacggcaacauggaaccuaauuugguaaaauuugagcc					
...(((.....))).....((((.....(((.....(((.....(((.....)))))))).....(((.....))).....)))).....					
.....uuugcucuuaacauuu.....uuugacgucgucuGugaucu.....			2	0	seq
.....uuugacgucgucuGugaucu.....uuacuaaaacagacgcacggc.....			1	1	seq
.....uuacuaaaacagacgcacggc.....Uaacagacgcacggcaac.....			1	0	seq
.....Uaacagacgcacggcaac.....aacagacgcacggcaaca.....			1	1	seq
.....aacagacgcacggcaaca.....aGcagacgcacggcaaca.....			1	1	seq
.....aGcagacgcacggcaaca.....aacagacgcacggcaacau.....			2	0	seq
.....aacagacgcacggcaacau.....aacagacgcacggcaacauC.....			1	1	seq
.....aacagacgcacggcaacauC.....aacagacUcacggcaacaug.....			1	1	seq
.....aacagacUcacggcaacaug.....aacagacgcacggcaacauU.....			2	1	seq
.....aacagacgcacggcaacauU.....aacagacgcacggcaacauA.....			2	1	seq
.....aacagacgcacggcaacauA.....aacagacgcacggcaacaug.....			7	0	seq
.....aacagacgcacggcaacaug.....aacagacgcacggcaacaugg.....			2	0	seq
.....aacagacgcacggcaacaugg.....aGcagacgcacggcaacaugg.....			1	1	seq
.....aGcagacgcacggcaacaugg.....aacagacgcacggcaacaugga.....			1	0	seq
.....aacagacgcacggcaacaugga.....acagacgcacggcaacau.....			4	0	seq
.....acagacgcacggcaacau.....acagacgcacggcaacaC.....			1	1	seq
.....acagacgcacggcaacaC.....acagacgcacggcaacaugg.....			15	0	seq
.....acagacgcacggcaacaugg.....acagacgUacggcaacaugga.....			1	1	seq
.....acagacgUacggcaacaugga.....acagacgcacggcaacauggaU.....			1	1	seq
.....acagacgcacggcaacauggaU.....acagacgcacAgcaacauggaa.....			2	1	seq
.....acagacgcacAgcaacauggaa.....cagacCcacggcaacaug.....			1	1	seq
.....cagacCcacggcaacaug.....cagacgcAaggcaacaugg.....			1	1	seq
.....cagacgcAaggcaacaugg.....cagacgcacggcaacauggU.....			1	1	seq
.....cagacgcacggcaacauggU.....cagacgcacggcaacaugga.....			1	0	seq
.....cagacgcacggcaacaugga.....cagacUcacggcaacaugga.....			2	1	seq