

Provisional ID : NW\_025322672.1\_Acropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_Amil\_v2.1\_Sc0000178\_whole\_genome\_shotgun\_sequenc

Score total : 2.5

Score for star read(s) : -1.3

Score for read counts : 0

Score for mfe : 0.8

Score for randfold : 0

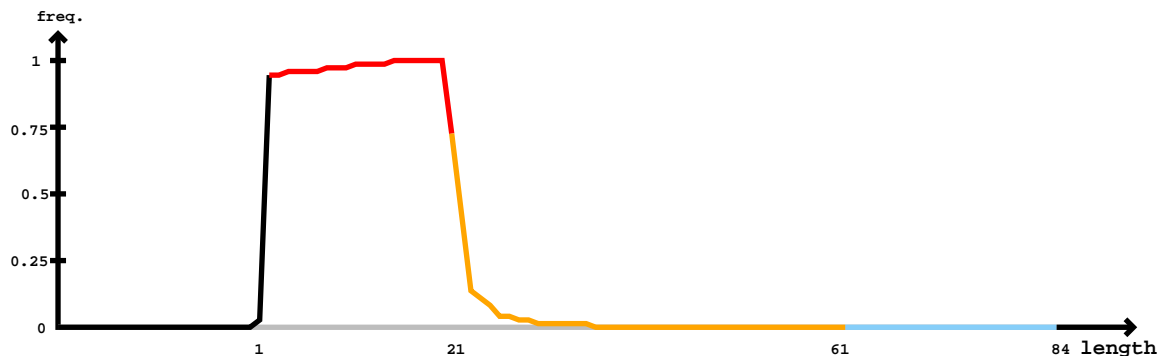
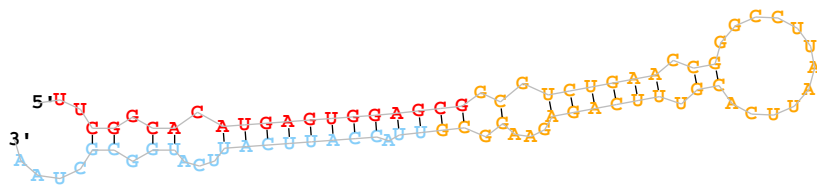
Score for cons. seed : 3

Total read count : 70

Mature read count : 70

Loop read count : 0

Star read count : 0



## Mature

## Star

5'	ucgaaccuaauugaauuaga <u>uucggcacaugaguggagcg</u> gucugaaccgggcccuaauucacguuucagagagggcg <u>uuaccauuc<u>auucauggcgcua</u>auuacaac</u>	-3'	exp	
	.....((((((((.....(((.....((((((((((((((((.....)))))).....)))))).....)))))).....)))))).....))))..	reads	mm	sample
	.....auucggcacaugaguggagc.....	1	0	seq
	.....auucggcacaugaguggagcA.....	1	1	seq
	.....uucggcacaugaguggagA.....	7	1	seq
	.....uucggcacaugaguggagc.....	12	0	seq
	.....uucggcacaugaguggagcA.....	13	1	seq
	.....uucggcacaugaguggagcg.....	13	0	seq
	.....uucggcGcaugaguggagcg.....	1	1	seq
	.....uucggcacaugaguggagcU.....	7	1	seq
	.....uucggcacaugaguggagcgg.....	8	0	seq
	.....uucggcacaugaguggagcggU.....	1	1	seq
	.....uucggcacaugaguggagcggA.....	1	1	seq
	.....uucggcacaugaguggagcggcg.....	1	0	seq
	.....uucggcacaugaguggagcggcA.....	1	1	seq
	.....uucggcGcaugaguggagcggcgu.....	1	1	seq
	.....uAcggcacaugaguggagcggcgu.....	1	1	seq
	.....cggcGcaugaguggagcggcgu.....	1	1	seq
	.....acaugaguggagcggcgucA.....	1	1	seq
	.....ugaguggagcggcgucuga.....	1	0	seq
	.....uggagcggcgucugaaccggg.....	1	0	seq