

Provisional ID : NC\_058075.1\_Acropora\_millepora\_isolate\_JS-1\_chromosome\_10\_Amil\_v2.1\_whole\_genome\_shotgun\_sequence\_394516

Score total : 5.5

Score for star read(s) : 3.9

Score for read counts : 4.7

Score for mfe : -6

Score for randfold : 0

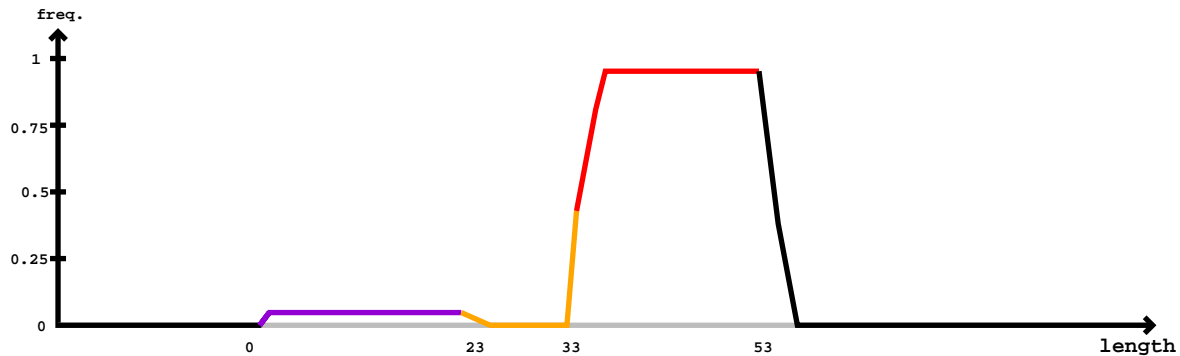
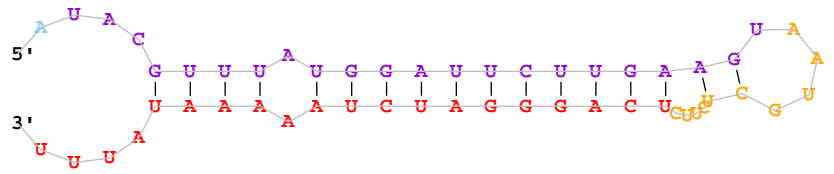
Score for cons. seed : 3

Total read count : 21

Mature read count : 20

Loop read count : 0

Star read count : 1



Star

Mature

5' -		-3'	obs		
	aacuccaaaauagacgaaaa <u>uacguuuu</u> augga <u>uuc</u> uuga <u>agu</u> <u>aaugc</u> u <u>uuc</u> ucagggau <u>cu</u> aaaa <u>uuuuu</u> uuggu <u>uuuuuuuuu</u> agauuugaagaaaaaggcccguaaag				
	aacuccaaaauagacgaaa <u>uacguuuu</u> augga <u>uuc</u> uuga <u>ag</u> <u>aaugc</u> u <u>uuc</u> ucagggau <u>cu</u> aaaa <u>uuuuu</u> uuggu <u>uuuuuuuuu</u> agauuugaagaaaaaggcccguaaag		exp		
	.....(((.....(((((((.....)))))))))))).....(((((((.....)))))))).....	reads	mm	sample	
	.....uacguuuu <u>augga</u> u <u>uuc</u> uuga.....	1	0	seq	
	.....ucagggau <u>cu</u> aaaa <u>uuuu</u> C.....	1	1	seq	
	.....ucagggau <u>cu</u> aaaa <u>uuuu</u> .....	1	1	seq	
	.....ucagggau <u>cu</u> aaaa <u>uuuuu</u> .....	5	0	seq	
	.....ucagggau <u>cu</u> aaaa <u>uuuuuuu</u> .....	2	0	seq	
	.....cagggau <u>cu</u> aaaa <u>uuuu</u> C.....	1	1	seq	
	.....cagggau <u>cu</u> aaaa <u>uuuuuuu</u> .....	2	0	seq	
	.....cagggau <u>cu</u> aaaa <u>uuuuuuu</u> .....	1	0	seq	
	.....agggau <u>cu</u> aaaa <u>uuuuuuu</u> .....	2	0	seq	
	.....agggau <u>cu</u> aaaa <u>uuuuuuu</u> .....	1	0	seq	
	.....agggau <u>cu</u> aaaa <u>uuuuuuu</u> g.....	1	1	seq	
	.....gggau <u>cu</u> aaaa <u>uuuuuuu</u> g.....	3	1	seq	