

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

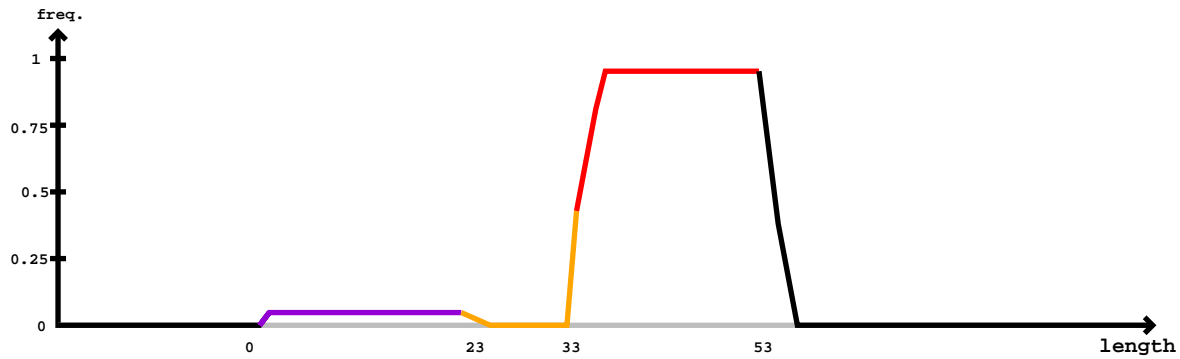
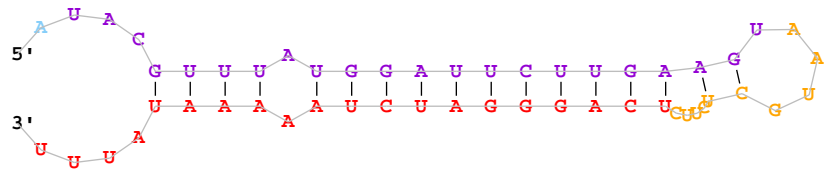
Score for cons. seed : 3

Total read count : 21

Mature read count : 20

Loop read count : 0

Star read count : 1



Star

Mature

5' -		-3'	obs		
	aacuccaaaauagacgaaaa <u>uacg</u> uuuauaggauuc <u>uugaagu</u> <u>aaugc</u> uc <u>uuc</u> cagggau <u>cuaaaaa</u> uuuuuuggu <u>aaaauuuuuu</u> agauuugaagaaaaaggcccguaaag				
	aacuccaaaauagacgaaa <u>uacg</u> uuuauaggauuc <u>uugaag</u> <u>uaa</u> ugc <u>uc</u> uuc <u>c</u> cagggau <u>cuaaaaa</u> uuuuuuggu <u>aaaauuuuuu</u> agauuugaagaaaaaggcccguaaag				
	.....(((.....(((((((.....)))))))))))))).....(((((((.....)))))))))).....	reads	mm	sample	
	.....uacguuuauaggauuc <u>uugaagu</u> .....	1	0	seq	
	.....ucagggau <u>cuaaaaa</u> uuuu.....	1	1	seq	
	.....ucagggau <u>cuaaaaa</u> uuuu.....	5	0	seq	
	.....ucagggau <u>cuaaaaa</u> uuu <u>C</u> .....	1	1	seq	
	.....ucagggau <u>cuaaaaa</u> uuuuu.....	2	0	seq	
	.....cagggau <u>cuaaaaa</u> uuu <u>C</u> .....	1	1	seq	
	.....cagggau <u>cuaaaaa</u> uuuuu.....	2	0	seq	
	.....cagggau <u>cuaaaaa</u> uuuuu.....	1	0	seq	
	.....agggau <u>cuaaaaa</u> uuuuu.....	2	0	seq	
	.....agggau <u>cuaaaaa</u> uuuuu.....	1	0	seq	
	.....agggau <u>cuU</u> aaaauuuuuug.....	1	1	seq	
	.....gggau <u>cuU</u> aaaauuuuuug.....	3	1	seq	