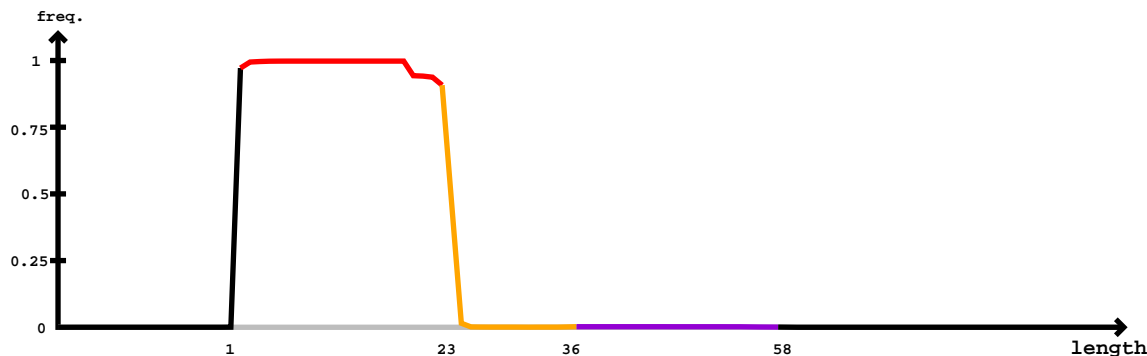
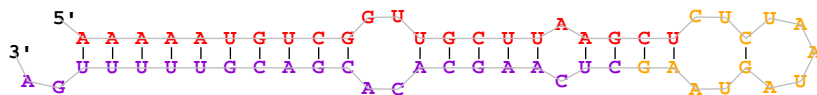


Provisional ID : NC\_058067.1\_Acropora\_millepora\_isolate\_JS-1\_chromosome\_2\_Amil\_v2.1\_whole\_genome\_shotgun\_sequence\_107317  
 Score total : 4346.5  
 Score for star read(s) : 3.9  
 Score for read counts : 4337.2  
 Score for mfe : 2.4  
 Score for randfold : 0  
 Score for cons. seed : 3  
 Total read count : 8519  
 Mature read count : 8506  
 Loop read count : 0  
 Star read count : 13



**Mature**

**Star**

5' -	obs	reads	mm	sample
guuucuguccguggcucaaaaaugucgguugcuuaagcucucuaauaguaagcucaagcacacgacgguuuuugagccacgggcgggcaaaagcaagugagcugcauuuuu	exp			
guuucuguccguggcucaaaaaugucgguugcuuaagcucucuaauaguaagcucaagcacacgacgguuuuugagccacgggcgggcaaaagcaagugagcugcauuuuu		1	0	seq
guuucuguccguggcucaaaaaugucgguugcuuaagcucucuaauaguaagcucaagcacacgacgguuuuugagccacgggcgggcaaaagcaagugagcugcauuuuu		2	1	seq
guuucuguccguggcucaaaaaugucgguugcuuaagcucucuaauaguaagcucaagcacacgacgguuuuugagccacgggcgggcaaaagcaagugagcugcauuuuu		1	1	seq
guuucuguccguggcucaaaaaugucgguugcuuaagcucucuaauaguaagcucaagcacacgacgguuuuugagccacgggcgggcaaaagcaagugagcugcauuuuu		464	1	seq
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guuucuguccguggcucaaaaaugucgguugcuuaagcucucuaauaguaagcucaagcacacgacgguuuuugagccacgggcgggcaaaagcaagugagcugcauuuuu		35	1	seq
guuucuguccguggcucaaaaaugucgguugcuuaagcucucuaauaguaagcucaagcacacgacgguuuuugagccacgggcgggcaaaagcaagugagcugcauuuuu		2	1	seq
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guuucuguccguggcucaaaaaugucgguugcuuaagcucucuaauaguaagcucaagcacacgacgguuuuugagccacgggcgggcaaaagcaagugagcugcauuuuu		13	1	seq

## Mature

## Star

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.....	aaaaugucggguugcuuaagc	Au.....			2	1	seq
.....	aaaaugucggguugcuuGagc	cu.....			25	1	seq
.....	aaaaugucggguugcuuaagc	Ccu.....			2	1	seq
.....	aaaaugucggguugcuuaagc	Uu.....			1	1	seq
.....	aaaaugucggguugcuuaagc	cC.....			15	1	seq
.....	aaaaugucggguugcuuaagc	cu.....			34	0	seq
.....	aaaaugucggguugcuuaagc	cG.....			2	1	seq
.....	aaaaugucggguugcuuaagc	cuA.....			1	1	seq
.....	aaaaugucggguugcuuaagc	cuAuc.....			1	1	seq
.....	aaaaugucggguugcuuaagc	cuU.....			1	1	seq
.....	aaaugucggguugcuuaagc	cu.....			4	0	seq
.....	Uaaugucggguugcuuaagc	cu.....			2	1	seq
.....	aaaugucggguugcuuaagc	C.....			4	1	seq
.....	aaaugucggguugcuuaagc	A.....			1	1	seq
.....	aaaugucggguugUuaagc	cu.....			1	1	seq
.....	aaaugucggguugcuuaagc	cu.....			3	0	seq
.....	aaaugucggguugcuuaagc	cuA.....			1	1	seq
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.....	aaugucggguugcuuGagc	cu.....			3	1	seq
.....	aaugucggguugcuuaagc	cu.....			2	0	seq
.....	augucggguugcuuaagc	cu.....			1	0	seq
.....	augucggguugcuuaagc	cC.....			1	1	seq
.....	ugucggguugcuuaagc	cucu.....			1	0	seq
.....	cuauaguaagcucaagcacacgacg	G.....			1	1	seq
.....	cuauaguaagcucaagcacacgacg	uuuu.....			1	0	seq
.....	aauguaagcucaagcaca	.....			1	0	seq
.....	uaagcucaagcacacgacg	uuuC.....			1	1	seq
.....	uaagcucaagcacacgacg	uuuu.....			1	0	seq
.....	gcucaagcacacgacg	uuuu.....			2	0	seq
.....	gcucaagcacacgacg	uuuUC.....			1	1	seq
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.....	cucaagcacacgacg	uuuu.....			1	0	seq
.....	cucaagcacacgacg	uuuuGga.....			6	1	seq