Provisional ID: NC\_058070.1\_Acropora\_millepora\_isolate\_JS-1\_chromosome\_5\_Amil\_v2.1\_whole\_genome\_shotgun\_sequence\_204550

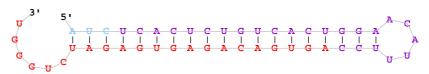
0.8 Score total Score for star read(s) -1.3 Score for read counts 0 2.7 Score for mfe Score for randfold 0 Score for cons. seed -0.6 Total read count 1254 Mature read count 1235

0

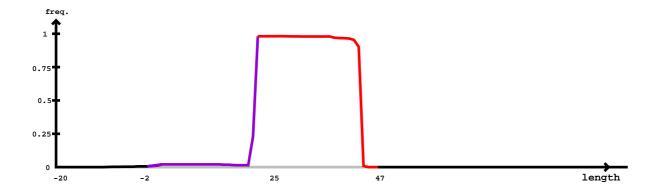
19

Loop read count

Star read count



sample



	Star	Mature
5'- ca	ccaccuuguacccagaucucacucugucacuggaacauuuc	cagugacagagugagaucuggguacgaaggugauuccucagaaaaaaaguucccgaugaauuuuuuugaa
ca	ccaccuuguacccagaucucacucugucacuggaacauuuc	cagugacagagugagaucuggguacgaaggugauuccucagaaaaaaaguucccgaugaauuuuuuugaa

.....uuuccagugacagagugagaucC.....

guacccagaucucacucugCcacuguacccagaucucacucucucucucucucucucucucucu	2	1	seq
uacccagaucucacucugucacu	1	0	seq
ccagaucucacucugucacugU	1	1	seq
agaucucacucugucacugg	3	0	seq
ucucacucugucacugga	1	0	seq
ucucacucugucacuggaacauuucc	1	0	seq
ucucacucugucacuggaacauuucU	2	1	seq
ucucacucugucacuggaacauuuccaguga	1	0	seq
cucacucugucacuggaacauuucU	1	1	seq
cucacucugucacuggaacauuuccaC	1	1	seq
.ucacucugucacuggaacauuucc	10	0	seq
.ucacucugucacuggaacauuucU	1	1	seq
.ucacucugucacuggaacauuuccagu	1	0	seq
.uuuccagugacagagug	11	0	seq
	1	1	seq
.uuuccagugacagagugU	2	1	seq
.uuuccagugacagagugaA.	1	1	seq
.uuuccagugacagagugagU	1	1	seq
.uuuccagugacagagugaga	1	0	seq
.uuuccagugacagagugagaA	1	1	seq
.uuuccagugacagagugagUu	2	1	seq
.uuuccagugacagagugagaG	2	1	seq
.uuuccagugacagagugagau	3	0	seq
.uuuccagugacagaggauG	1	1	seq
.uuuccagugacagaggauA	4	1	seq
.uuuccagugacagagugagaAc	1	1	seq
.uuuccagugacagagugagauU	44	1	seq
.uuuccagugacagagugaAauc	2	1	seq
.uuuccagugacagagugUgaucu	1	1	seq
.uuuccagugacagagugagaucA	19	1	seq
.uuuccagugacagagugagauUu.	2	1	seq
uuuccagugacagagugagaucu	151	0	seq

Star Mature

2002			
caccaccuuguacccagaucucacucugucacuggaacauuucc <mark>agugacagagugagaucugggua</mark> cgaaggugauuccucagaaaaaaguucccgaugaauuuuuuugaa			
	2	1	seq
.uuuccagugacagagugGgaucu	2	1	seq
	2	1	seq
	1	1	seq
.uuccagugacagagugG	1	1	seq
.uuccagugacagagugaga	1	0	seq
.uuccagugacagagugagUu	1	1	seq
.uuccagugacagagugagau	4	0	seq
.uuccagugacagagugGgauc	2	1	seq
.uuccagugacagagugagauU	9	1	seq
.uuccagugacagagugagauA.	2	1	seq
.uuccagugacagagugagaucC	144	1	seq
.uuccagugacagagCgagaucu	2	1	seq
.uuccagugacagagugagaCcu	4	1	seq
.uuccagugacagagugagaucA	109	1	seq
.uuccagugacagagugagauUu.	6	1	seq
.uuccagugacagagugagaGcu	1	1	seq
.uuccagugacUgagugaucu.	1	1	seq
.uuccagugacagagugagGucu.	2	1	seq
.uuccUgugacagagugagaucu	1	1	seq
.uuccagugacagagugGgaucu	5	1	seq
.uuccagugacagagugagaucu	643	0	seq
.uuccagugacagagugagaucuC	1	1	seq
.uuccagugacagagugagaucuU	4	1	seq
.uuccagugacagagugagaucuA	4	1	seq
.uuccagugacagagugagaucAg	1	1	seq
uccagugacagagugagaucu	11	0	seq
uccagugacagagugagaucC	1	1	seq
ccagugacagagugagaucu.	3	0	seq
.cagugacagagugagaucC	1	1	seq