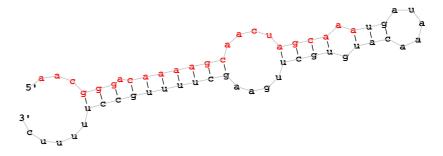
miRBase precursor : chromosome_12_56380

Total read count : 40 chromosome_12_56380 read **40**unt remaining reads : 0





chromosome_12_56380

5'- aacgggacaaaagcaacuagcaaaugauaaacaugugcuugaagcuuuuugccuuuuuc	-3'	exp		
(((.(((((((())).)))))))))))))	read	s	mm	sample
aacgggacaaaagcaaGuagcaaa	1		1	seq
aacgggacaaaagcaacuagcaaa	11		0	seq
Gacgggacaaaagcaacuagcaaa	1		1	seq
aaUgggacaaaagcaacuagcaaa	2		1	seq
Cacgggacaaaagcaacuagcaaa	1		1	seq
aacgggacaaaGgcaacuagcaaa	1		1	seq
aacgggacaaaagcaacuagcaaG	2		1	seq
aacgggacaaaagcaacuagcaGa	3		1	seq
aacgggacaaaagcaacuagcaaaG	1		1	seq
.Gcgggacaaaagcaacuagcaaa	2		1	seq
.acgggacaaaagcaacuagcaGa	3		1	seq
.acgggacaaaagcaacuagcaaa	10		0	seq
.Ucgggacaaaagcaacuagcaaa	1		1	seq
gggacaaaagcaacuagcaaa	1		0	seq