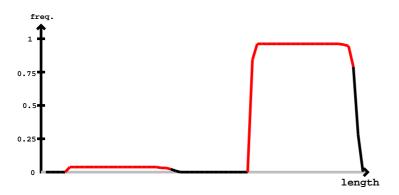
miRBase precursor : hsa-mir-378a

Total read count : 315 hsa-miR-378a-5p read count 12 hsa-miR-378a-3p read count 303 remaining reads : 0





hsa-miR-378a-5p

hsa-miR-378a-3p

inda min 370d 3p			
5'- agggcuccugacuccagguccuguguguuaccuagaaauagcacuggacuuggagucagaaggccu	-3'	exp	
(((.((.(((((((((((()))))).))))))	reads	s mm	sample
cuccugacuccagguccug	2	0	seq
cuccugacuccagguccugug	3	0	seq
cuccugacuccagguccugugA	1	1	seq
cuccugacuccagguccugugu	3	0	seq
cuccugacuccagguccuguguU	2	1	seq
cuccugacuccagguccugugug	1	0	seq
acuggacuuggagucagaa	1	0	seq
	1	1	seq
acuggacuuggagucagaag	3	0	seq
	4	1	seq
acuggacuuggagucagaagg	44	0	seq
acuggacuuggagucUgaagg	1	1	seq
acuggacuuggagucagaaggU	3	1	seq
acuggacuuggagucagaaggc	96	0	seq
acuggacuuggaguUagaaggc	1	1	seq
acuggacuuggagucaAaaggc	1	1	seq
	6	1	seq
acuggacuuggagucagaaggA	28	1	seq
acAggacuuggagucagaaggc	1	1	seq
acuggacuuggagucagaaggcG.	2	1	seq
acuggacuuggagucagaaggAc.	1	1	seq
acuggacuuggagucagaaggcU.	31	1	seq
acuggacuuggagucagaaggcA.	40	1	seq
acuggacuuggagucagaaggcAu	1	1	seq
cuggacuuggagucagaaggA	2	1	seq
cuggCcuuggagucagaaggc	1	1	seq
cuggacuuggagucagaaggc	22	0	seq
cuggacuuggagucagaaggcA.	4	1	seq
cuggacuuggagucagaaggcU.	9	1	seq