



hsa-miR-378a-3p

5'	aggggcuccugagacuccaggguccuguguguuaccuagaaaagcacuggacuuggagucagagaaggccu	-3'	exp
	(((((((( ((((((((((((((((((((((.(. ....).).))))).)))))))))))))).).)))	reads	mm sample
...	cuccugagacuccaggguccug...	2	0 seq
...	cuccugagacuccaggguccugug...	3	0 seq
...	cuccugagacuccaggguccugugA...	1	1 seq
...	cuccugagacuccaggguccugugug...	3	0 seq
...	cuccugagacuccaggguccugugugU...	2	1 seq
...	cuccugagacuccaggguccugugug...	1	0 seq
...	...acuggacuuggagucagaaa...	1	0 seq
...	...Ccuggacuuggagucagaaa...	1	1 seq
...	...acuggacuuggagucagaaag...	3	0 seq
...	...Ccuggacuuggagucagaaagg...	4	1 seq
...	...acuggacuuggagucagaaagg...	44	0 seq
...	...acuggacuuggagucUgaagg...	1	1 seq
...	...acuggacuuggagucagaaaggU...	3	1 seq
...	...acuggacuuggagucagaaaggc...	96	0 seq
...	...acuggacuuggagucUgaaggc...	1	1 seq
...	...acuggacuuggagucAaaaggc...	1	1 seq
...	...Ccuggacuuggagucagaaaggc...	6	1 seq
...	...acuggacuuggagucagaaaggA...	28	1 seq
...	...acAaggacuuggagucagaaaggc...	1	1 seq
...	...acuggacuuggagucagaaaggcG...	2	1 seq
...	...acuggacuuggagucagaaaggAc...	1	1 seq
...	...acuggacuuggagucagaaaggcU...	31	1 seq
...	...acuggacuuggagucagaaaggcA...	40	1 seq
...	...acuggacuuggagucagaaaggcAu...	1	1 seq
...	...cuggacuuggagucagaaaggA...	2	1 seq
...	...cuggCcuuggagucagaaaggc...	1	1 seq
...	...cuggacuuggagucagaaaggc...	22	0 seq
...	...cuggacuuggagucagaaaggcA...	4	1 seq
...	...cuggacuuggagucagaaaggcU...	9	1 seq