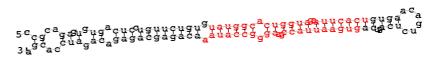
miRBase precursor : hsa-mir-183

Total read count : 173 hsa-miR-183-5p read count : 162 hsa-miR-183-3p read count : 10 remaining reads : 1





hsa-miR-183-3p

hsa-miR-183-5p

5'- ccgcagagugugacuccuguucuguguauggcacugguagaauucacugugaacagucucagucag	-3'	exp	
.((((.(.((.((((((((((((((((((((((((((read	ls mm	sample
uauggcacugguagaauu	5	0	seq
uauggcacugguagaauuc	2	0	seq
uauggcacugguagaauuca	2	0	seq
uauggcacugguagaauucac	9	0	seq
uauggcacugguagaauucacu.	38	0	seq
uauggcacugAuagaauucacu	1	1	seq
uauggcacugguagaauucacug	12	0	seq
uauggcacugguagaauucacugA.	4	1	seq
.auggcacugguaAaauuca	1	1	seq
.auggcacugguagaauuca	1	0	seq
.auggcacugguagaauucac	4	0	seq
.auggcacCgguagaauucacu	1	1	seq
.augAcacugguagaauucacu	1	1	seq
.auggcacugguagaauucacu	37	0	seq
.Uuggcacugguagaauucacu	1	1	seq
.Cuggcacugguagaauucacug	1	1	seq
auggcacugguagaauucacug	16	0	seq
.Cuggcacugguagaauucacugu	1	1	seq
auggcacugguagaauucacugG	1	1	seq
auggcacugguagaauucacugA	17	1	seq
.auggcacugguagaauucacugu	7	0	seq
ucagugaauuUccgaagggcc	1	1	seq
gugaauuaccgaagggcca	3	0	seq
gugaauuaccgaagggccauU	1	1	seq
	1	0	seq
	1	0	seq
ugaauuaccgaagggccauaa	3	0	seq
ugaauuaccgaagggccauaaa	1	0	seq