

5' c g c a g g u a g u a g u u c u g a u a u g g c a c u g g u a u u c a u g u g a c a
g g c a c u a g a g a g a g u u c u g a u a u g g c a c u g g u a u u c a u g u g a c a



5'	hsa-miR-183-5p	3'	exp	reads	mm	sample
ccgcagagugugacuccuguuucugug	uauggcacugguagaaauucacug	gugaacagucucagucagug	aaauaccgaagggccauaa	acagagcagagacagauccacga		
(((.....)))						
.....uauggcacugguagaaau				5	0	seq
.....uauggcacugguagaaauuc				2	0	seq
.....uauggcacugguagaaauuca				2	0	seq
.....uauggcacugguagaaauucac				9	0	seq
.....uauggcacugguagaaauucacu				38	0	seq
.....uauggcacugAuaagaaauucacu				1	1	seq
.....uauggcacugguagaaauucacug				12	0	seq
.....uauggcacugguagaaauucacugA				4	1	seq
.....auggcacugguAaauuca				1	1	seq
.....auggcacugguagaaauuca				1	0	seq
.....auggcacugguagaaauucac				4	0	seq
.....auggcacCgguagaaauucacu				1	1	seq
.....augAacugguagaaauucacu				1	1	seq
.....auggcacugguagaaauucacu				37	0	seq
.....Uuggcacugguagaaauucacu				1	1	seq
.....Cuggcacugguagaaauucacug				1	1	seq
.....auggcacugguagaaauucacug				16	0	seq
.....Cuggcacugguagaaauucacugu				1	1	seq
.....auggcacugguagaaauucacugG				1	1	seq
.....auggcacugguagaaauucacugA				17	1	seq
.....auggcacugguagaaauucacugu				7	0	seq
.....ucagugaaauUccgaagggcc				1	1	seq
.....gugaaauuaccgaagggccca				3	0	seq
.....gugaauuaccgaagggccauU				1	1	seq
.....ugaauuaccgaagggccca				1	0	seq
.....ugaauuaccgaagggcccau				1	0	seq
.....ugaauuaccgaagggcccauaa				3	0	seq
.....ugaauuaccgaagggcccauaaa				1	0	seq