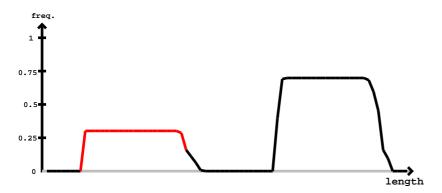
miRBase precursor : hsa-mir-2110

Total read count : 126 hsa-miR-2110 read count : 38 remaining reads : 88





hsa-miR-2110

IISA-MIR-ZIIV			
5'- cagggguuuggggaaacggccgcugagugaggcgucggcuguguuucucaccgcggucuuuuccucccacucuug	-3'	exp	
(((((((((((((((((((((((((((((((((((((reads	mm	sample
uuggggaaacggccgcugag	2	0	seq
uuggggaaacggccgcugagu	16	0	seq
uuggggaaacggccgcugagug	6	0	seq
uuggggaaacggccgcugaguga	5	0	seq
uuggggaaCcggccgcugaguga	1	1	seq
uuggggaaacggccgcugagugag	6	0	seq
uuggggaaacggccgcugagugaU	1	1	seq
uuggggaaacggccgcugagugagg	1	0	seq
ucaccgcggucuuuuccuc	2	0	seq
Ccaccgcggucuuuuccucc	1	1	seq
ucaccgcggucuuuuccucc	9	0	seq
ucaccgcggucuuuuccuccU	3	1	seq
ucaccgcggucuuuuccuccc	8	0	seq
ucaccgcggucuuuuccuccA	1	1	seq
ucaccgcggucuuuuccucccU	14	1	seq
ucaccgcggucuuuuccuccca	8	0	seq
ucaccgcggucuuuuccucccaU	1	1	seq
ucaccgcggucuuuuccucccaA	1	1	seq
ucaccgcggucuuuuccucccaUu	1	1	seq
ucaccgcggucuuuuccucccacA	1	1	seq
caccgcggucCuuuccucc	1	1	seq
caccgcggucuuuuccuccc	3	0	seq
caccgcggucuuuuccuccU	3	1	seq
caccgcggucuuuuccucccU	11	1	seq
caccgcggucuuuuccuccca	4	0	seq
caccgcggucuuuuccucccaU	2	1	seq
caccgcggucuuuuccucccac	3	0	seq
cCccgcggucuuuuccucccac	1	1	seq
Gaccgcggucuuuuccucccacu	1	1	seq
caccgcggucuuuuccucccacu	5	0	seq
caccgcggucuuuuccucccaAu	1	1	seq
caccgcggucuuuuccucccaUu	2	1	seq
accgcggucuuuuccucccacu	1	0	seq