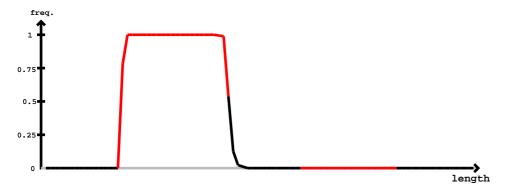
miRBase precursor : hsa-mir-181b-2

Total read count : 175 hsa-miR-181b-5p read count 175 hsa-miR-181b-2-3p read count remaining reads : 0





hsa-miR-181b-5p

hsa-miR-181b-2-3p

inda min 1010 2 5p			
5'- cugauggcugcacucaacauucauugcugucgguggguuugagucugaaucaacucacugaucaaugaaug	-3'	exp	
$\dots (((((((((((((((((((((((((((((((((((($	reads	s mm	sample
aacauucauugcugucggug	1	0	seq
aacauucauugcugucggugA	1	1	seq
aacauucauuAcugucgguggg	1	1	seq
aacauucauugcugucgguggg	60	0	seq
aacauucCuugcugucgguggg	1	1	seq
Cacauucauugcugucgguggg	4	1	seq
aacauucauugcugucgguggA	3	1	seq
aacauucauugcugucggugggA	21	1	seq
aacauucauugcugucggugggG	2	1	seq
aacauucauugcugucggugggu	25	0	seq
Cacauucauugcugucggugggu	3	1	seq
aacauucauugcugucgguggUu	1	1	seq
aacauucauugcugucgguggguG	1	1	seq
aacauucauugcugucgguggguu	10	0	seq
aacauucauugcugucgguggguuu	1	0	seq
aacauucauugcugucgguggguuuU	2	1	seq
acauucauugcugucgguggg	9	0	seq
Ccauucauugcugucgguggg	1	1	seq
Ccauucauugcugucggugggu	4	1	seq
acauucauugcugucggugggu	15	0	seq
Ucauucauugcugucggugggu	1	1	seq
acauucauugcugucgguggguG	1	1	seq
acauucauugcugucgguggguu	2	0	seq
acauucauugcugGcgguggguu	1	1	seq
acauucauugcugucgguggguA	2	1	seq
acauucauugcugucggugggAu	1	1	seq
acauucauugcugucgguggguuu	1	0	seq