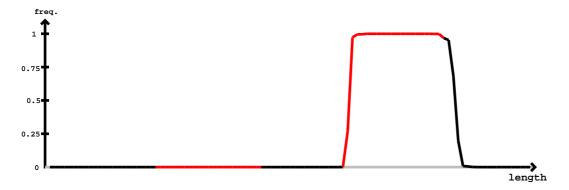
miRBase precursor : hsa-mir-140
Total read count : 2901
hsa-miR-140-5p read count : 1
hsa-miR-140-3p read count : 2897
remaining reads : 3



hsa-miR-140-3p

5'- ugugucucucucuguguccugccagugguuuuacccuaugguagguuacgucaugcuguucuaccacaggguagaaccacggacaggauaccggggcacc					
$\cdot ((((((((((((((((((((((((((((((((($					

hsa-miR-140-5p

	-		
	reads	mm	sample
cagugguuuuacccuauggu	1	0	seq
	2	1	seq
uaccacaggguagaacca	3	0	seq
uaccacaggguagaaccacg	33	0	seq
uaccCcaggguagaaccacg	2	1	seq
uaccacagggCagaaccacg	1	1	seq
uaccCcaggguagaaccacgg	1	1	seq
uaccacaggguagaaccacgg	14	0	seq
.uaccacaggguagaaccacgA	1	1	seq
uaccacaggguagaaUcacgga	1	1	seq
uaccacGggguagaaccacgga	1	1	seq
uaccacagggCagaaccacgga	2	1	seq
uaccacaUgguagaaccacgga	1	1	seq
.uaccCcaggguagaaccacgga	24	1	seq
uaccUcaggguagaaccacgga	1	1	seq
uaccacaggguagaaccacggG	2	1	seq
uaccacaggguagaaccacgga	300	0	seq
uaccacaggguagCaccacgga	1	1	seq
.uaccacaggguahaaccacgga	1	1	seq
.uaccCcaggguagaaccacggac	7	1	seq
uCccacaggguagaaccacggac	1	1	seq
uaccacagggAagaaccacggac	1	1	seq
uaccacagCguagaaccacggac	1	1	seq
uaccacaggguagaaccacggaA	234	1	seq
.uaccacaAgguagaaccacggac	1	1	seq
.uaccacaggguagaaccacggaU	1	1	seq
	81	0	seq
uaccacaggguagaaccacggaGa	5	1	seq
uaccacaggguagaaccacggacG	3	1	seq
uaccCcaggguagaaccacggaca	1	1	seq
uaccacaggguagaaccacggaAa	6	1	seq
uaccacaggguagaaccacggaca	4	0	seq
uaccacaggguagaaccacggacU	47	1	seq
Cccacaggguagaaccacg	2	1	seq

ugugucucucuguguccugccagugguuuuacccuaugguagguuacgucaugcuguucuaccacaggguagaaccacggacaggauaccggggcacc

.accacaggguagaaccacg	49	0	seq
accacaggAuagaaccacg	1	1	seq
	2	1	seq
accacaggguagaaccacgA	4	1	seq
.accacaggguagaaccacgg	21	0	seq
.accacaggguagaaccacgU	1	1	seq
accacaggguagaaccaUgga	1	1	seq
accacaggguagaaccacgAa	3	1	seq
accacagggGagaaccacgga	1	1	seq
accacaggguagaaAcacgga	3	1	seq
accacaggguagaCccacgga	1	1	seq
accacaggguagaaccacggG	1	1	seq
	2	1	_
accacaggguagaGccacgga			seq
Cccacaggguagaaccacgga	27	1	seq
accacaAgguagaaccacgga	1	1	seq
Gccacaggguagaaccacgga	1	1	seq
.accacaggguagaaccacgga	395	0	seq
.accacaggguagaaccGcgga	1	1	seq
accacaggguagaaccacggaU	13	1	seq
accacaggguagaaccacgAac	1	1	seq
accacaggguagaaccacggac	701	0	seq
accCcaggguagaaccacggac	1	1	seq
accacaggguagaaccacggaG	14	1	seq
accacaCgguagaaccacggac	1	1	seq
accacCggguagaaccacggac	1	1	seq
.accacaggguagaaAcacggac	1	1	seq
.accacaggguagaaccacggaA.	287	1	seq
accacaggguagaCccacggac	2	1	seq
accacaggguaUaaccacggac	1	1	seq
accacaggAuagaaccacggac	1	1	seq
	31	1	seq
accacaggguagaUccacggac	1	1	seq
accacaggguagaaccacggacU	230	1	seq
	3	1	
	11	1	seq
accacaggguagaaccacggaUa	1	1	seq
	75	1	
accacaggguagaaccacggaAa	4	1	seq
accacaggguagaaccacggacG	37	1	seq
	87	0	
accacaggguagaaccacggaca	1	1	seq
accacaggguagaaccacggacag	3	0	seq
accacaggguagaaccacggaAag	3	1	seq
accacaggguagaaccacggacaAg	5	1	seq
accacaggguagaaccacggacagU	1	1	seq
accacaggguagaaccacggacagA	4	1	seq
accacaggguagaaccacggacaAga	1	1	seq
ccacaggguagaaccacg	1	0	seq
ccacaggguagaaccacgga	5	0	seq
	4	1	seq
cacaggguagaaccacggaU	1	1	seq
ccacaggguagaaccacggac	11	0	seq
ccacaggguagaaccacggaG	1	1	seq
ccacagggCagaaccacggac	1	1	seq
ccacaggguagaaccacggacG	2	1	seq
ccacaggguagaaccacggaAa	3	1	seq
ccacaggguagaaccacggacU	33	1	seq
ccacaggguagaaccacggacUg	1	1	seq
ccacaggguagaaccacggacaAga	1	1	seq
cacaggguagaaccacgga	1	0	seq
cacaggguagaaccacggacU	2	1	seq
acaggguagaaccacgga	5	0	seq
acaggguagaaccacggac	1	0	seq
acaggguagaaccacggacagg	3	0	seq
acaggguagaaccacggacagga	1	0	seq
		-	