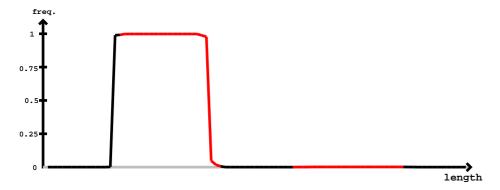
miRBase precursor : hsa-mir-142
Total read count : 17943
hsa-miR-142-5p read count : 17881
hsa-miR-142-3p read count : 30
remaining reads : 32





hsa-miR-142-3p

		_			
ısa	-m;	.ĸ-	14	2-	5p

118a-mir-142-5p			
5'- gacagugcagucacccauaaaguagaaagcacuacuaacagcacuggaggguguaguguuuccuacuuuauggaugaguguacugug	-3'	exp	
.((((((((((((((((((((((((((((((((((((reads	mm	sample
Ccccauaaaguagaaagca	1	1	seq
acccauaaaguagaaagcac	1	0	seq
Ccccauaaaguagaaagcac	1	1	seq
Ccccauaaaguagaaagcacu	29	1	seq
cccauaaaguagaaagca	169	0	seq
cccCuaaaguagaaagca	1	1	seq
cccauaaaguGgaaagca	1	1	seq
Accauaaaguagaaagca	1	1	seq
cccauaaCguagaaagca	1	1	seq
cccauaaaguagaGagca	1	1	seq
cccauaaaguagCaagca	1	1	seq
cccauCaaguagaaagcac	4	1	seq
cccauUaaguagaaagcac	1	1	seq
cccauaaaguGgaaagcac	1	1	seq
cccCuaaaguagaaagcac	2	1	seq
ccUauaaaguagaaagcac	1	1	seq
cccauaaaguagaaagcac	179	0	seq
cccauaaaguagaaagUac	1	1	seq
cccauaaaguagaaagcaA	1	1	seq
cccauaaaguCgaaagcacu	3	1	seq
cccauaaUguagaaagcacu	2	1	seq
cccauaaaguagaaagcacu	16052	0	seq
cccauUaaguagaaagcacu	6	1	seq
cccauaaaguaCaaagcacu	4	1	seq
cccauaaGguagaaagcacu	3	1	seq
cccauaaagGagaaagcacu	6	1	seq
cccauaaaguagUaagcacu	3	1	seq
cccauaaaguagGaagcacu	3	1	seq
cccauGaaguagaaagcacu	3	1	seq
cccauaaaCuagaaagcacu	3	1	seq
cccauaaaguagaUagcacu	8	1	seq
cUcauaaaguagaaagcacu	12	1	seq
cccauaaaguagaaagcGcu	2	1	seq
cccaCaaaguagaaagcacu	4	1	seq

cccauaaCguagaaagcacu	15	1	seq
ccUauaaaguagaaagcacu	7	1	seq
cccaAaaaguagaaagcacu	5	1	seq
cccauaaaguagaaUgcacu	1	1	seq
cccauaaaguaUaaagcacu	1	1	seq
cccauaaaguagaaagcacCccAauaaaguagaaagcacu	10 1	1	seq
cccauaaaguagaaCgcacu	8	1	_
cccauaCaguagaacgcacu	2	1	seq
	2	1	seq
cccauaaaguagaGagcacu	5	1	seq
Uccauaaaguagaaagcacu	20	1	seq
cccauaaaAuagaaagcacu	6	1	seq
cccauCaaguagaaagcacu	193	1	seq
cccauaaaguGgaaagcacu	3	1	seq
Accauaaaguagaaagcacu	8	1	seq
cccauaaaguagaaaCcacu	1	1	seq
	3	1	seq
	2	1	seq
cccauaaaguagaCagcacu	6	1	seq
cccauaaaguagaaagUacu	3	1	seq
cccauaaaguagaaagcCcu	6	1	seq
cccauaaaguagaaagcacG	27	1	seq
cccauaaaguagaaagcacA	6	1	seq
cccauaaagCagaaagcacu	5	1	seq
cccCuaaaguagaaagcacu	71	1	seq
cccauaaaguagaaagcaUu	2	1	seq
Gccauaaaguagaaagcacu	3	1	seq
cccUuaaaguagaaagcacu	12	1	seq
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cccGuaaaguagaaagcacu	6	1	seq
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cccauaaaguagaaCgcacua	1	1	seq
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	1 2	1	seq
cccauaaaguagaaagcacuG	1	1	seq
Uccauaaaguagaaagcacua	1	1	seq
cccauCaaguagaaagcacua	8	1	seq
cccauaaaguagaaagcGcua	1	1	seq
cccauaaaguagaaagcacua	516	0	seq
cccauaaaguagaaagcacuaG	9	1	seq
cccauCaaguagaaagcacuac	1	1	seq
cccauaaaguagaaagcacuaA	58	1	seq
	15	1	seq
cccauaaaguagaaagcacuac	111	0	seq
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cccauaaaguagaaagcacuacu	9	0	seq
cccauaaaguagaaagcacuacG	8	1	seq
cccauaaaguagaaagcacuacA	34	1	seq
cccauaaaguagaaagcacuacC	10	1	seq
cccauaCaguagaaagcacuacu	1	1	seq
cccauaaaguagaaagcacuaAu	2	1	seq
ccauaaaguagaaagcacu	31	0	seq
ccauCaaguagaaagcacu	2	1	seq
ccauCaaguagaaagcacua	1	1	seq
ccauaaaguagaaagcacua	2	0	seq
cauaaaguagaaagcacu	61	0	seq
cauaCaguagaaagcacu	2	1	seq
cauaaaguagaaaCcacu	1	1	seq
cauaaaguagaaagcacua	12	0	seq
cauaaaguagaaagcacuac	4	0	seq
cauaCaguagaaagcacuac	1 37	1 0	seq
cauaaguagaaagcacuacucauaCaguagaaagcacuacu	37	0 1	seq
cauaCaguagaaagcacuacu	2	0	seq
uguaguguuuccuacCuuaugga	1	1	seq
uguaguguauccaaccaaagga	_	-	204

hsa-miR-142-3p

hsa-miR-142-5p

gacagugcagucacccauaaaguagaaagcacuacuaacagcacuggaggguguaguguuuccuacuuuauggaugaguguacugug

uguaguguuuccuacuuuaugga	11	0	seq
uguaguguuuccuacuuuauggU	1	1	seq
guaguguuuccuacuuuaugga	1	0	seq
uaguguuuccuacuuuaugga	1	0	seq
Cguguuuccuacuuuauggaug	2	1	seq
aguguuuccuacuuuauggaug	13	0	seq