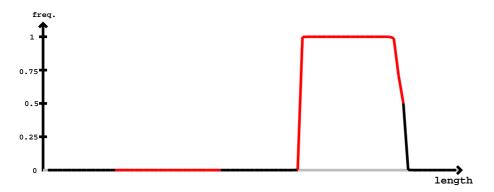
miRBase precursor : hsa-mir-22
Total read count : 31713
hsa-miR-22-5p read count : 2
hsa-miR-22-3p read count : 31711
remaining reads : 0





hsa-miR-22-3p

hsa-miR-22-5p

118a-mir-22-3p			
5'- ggcugagccgcaguaguucuucaguggcaagcuuuauguccugacccagcuaaagcugccaguugaagaacuguugcccucugcc	-3'	exp	
(((.(((((((((((((((((((((((((((((((((read	s mm	sample
aguucuucaguggcaagcu	1	0	seq
Cguucuucaguggcaagcu	1	1	seq
aaagcugccaguugaagaacu	14	0	seq
Caagcugccaguugaagaacu	3	1	seq
aaagcugccGguugaagaacu	1	1	seq
	2	1	seq
aaagcugccaguugaagaacug	9	0	seq
	5	1	seq
aaagcugccaguugaagaacugu	15	0	seq
aagcugccaguugaagaa	3	0	seq
	1	1	seq
Cagcugccaguugaagaac	12	1	seq
aagcugccGguugaagaac	1	1	seq
aCgcugccaguugaagaac	1	1	seq
aagcugccaguugaagaac	286	0	seq
aagcugccagAugaagaac	1	1	seq
aagcugccaguugaCgaac	1	1	seq
aagcugccaguugaagCac	1	1	seq
aagcugccaguugaaCaacu	1	1	seq
aGgcugccaguugaagaacu	2	1	seq
aagcugccaguugaUgaacu	1	1	seq
aagcugccaguugCagaacu	2	1	seq
aagcugcAaguugaagaacu	1	1	seq
aagcugccaguAgaagaacu	4	1	seq
aagcugccagGugaagaacu	1	1	seq
aagcugccaguCgaagaacu	9	1	seq
aUgcugccaguugaagaacu	4	1	seq
Gagcugccaguugaagaacu	4	1	seq
aagcugUcaguugaagaacu	7	1	seq
aagcugccaguugaagaaGu	1	1	seq
aagGugccaguugaagaacu	1	1	seq
	567	1	seq
aagAugccaguugaagaacu	1	1	seq
aagcugccCguugaagaacu	8	1	seq

ggcugagccgcaguaguucuucaguggcaagcuuuauguccugacccagcuaaagcugccaguugaagaacuguugcccucugcc

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aagcugccaguugaagCacu	1	1	seq
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aagcugccaguugaagaacG	32	1	seq
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aagcugccagAugaagaacu	2	1	seq
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aagcugccaguugaagaCcu	2	1	seq
aagcuAccaguuqaagaacu	7	1	seq
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aagcugccaCuugaagaacu	3	1	seq
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aagcuCccaguugaagaacu	1	1	seq
aagcugAcaguugaagaacu	1	1	seq
aCgcugccaguugaagaacu	10	1	seq
aagcugccaguugaagaUcu	1	1	seq
aagcugccaguuAaagaacu	4	1	seq
aagUugccaguugaagaacu	2	1	seq
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aagcugccGguugaagaacu	4	1	seq
aagcCgccaguugaagaacu	2	1	seq
aagcugccaguuCaagaacu	1	1	seq
aagcGgccaguugaagaacu	1	1	seq
aagcugccaguuUaagaacu	2	1	seq
aagcugccaguugUagaacu	2	1	seq
aagcugccaguugGagaacu	7	1	seq
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aagcugccaguugUagaacug	4	1	seq
aagcCgccaguugaagaacug	1	1	seq
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aagcugccaguugaagJacug	1	1	_
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aagcugccaguugaGgaacug	1	1	seq
aagcugccaguugCagaacug	1	1	seq
aagcugccCguugaagaacug	8	1	seq
aagcugccagAugaagaacug	8	1	seq
aagcGgccaguugaagaacug	2	1	seq
aagcugccaguugaagaacug	6121	0	seq
aagcugccaguugaagaacCg	4	1	seq
aaCcugccaguugaagaacug	2	1	seq
aUgcugccaguugaagaacug	3	1	seq
aagcugccaguAgaagaacug	1	1	seq
aagcugccaguugaCgaacug	1	1	seq
aagcugccaguugaagaacGg	27	1	peq
aagcugGcaguugaagaacug	4	1	seq
aagcugccaguugaaCaacug	1	1	seq
	3	1	seq
aagcugccaguuCaagaacug	3	1	seq
aagcAgccaguugaagaacug	1	1	seq
aagcugccagGugaagaacug	1	1	seq
aagUugccaguugaagaacug	3	1	seq
aagcugccaguugaaAaacug	2	1	seq
aagcugccaguugaagaacuA	2	1	seq
aagcugccaguugaagaacuU	2	1	seq
aagcugccUguugaagaacug	4	1	seq
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aagcugccagCugaagaacug	3	1	seq
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aagcugccaguCgaagaacug	3		seq
aCgcugccaguugaagaacug	3	1	seq
aagcuCccaguugaagaacug	2	1	seq

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Uagcugccaguugaagaacug	4	1	pea
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aagcugccaguugaGgaacugu	2	1	seq
Gagcugccaguugaagaacugu	8	1	seq
aagcugccaguugaagaGcugu	1	1	seq
aagcugccaguugaagaacCgu	2	1	seq
aagcGgccaguugaagaacugu	1	1	seq
aagGugccaguugaagaacugu	1	1	seq
aagUugccaguugaagaacugu	3	1	seq
aagcugccaguugaagaaUugu	3	1	seq
aagcugccaguugGagaacugu	9	1	seq
aagcugccaguugaagaacugA	27	1	seq
aagcugccagAugaagaacugu	7	1	seq
aagcugccaguugaagaUcugu	1	1	seq
aagcugccagGugaagaacugu	2	1	seq
aagcugcAaguugaagaacugu	2	1	seq
aagAugecaguugaagaacugu	2	1	seq
aUgcugccaguugaagaacugu	4	1	seq
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aagcuQccaguuqaagaacugu	2	1	seq
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aagcCgccaguugaagaacugu	2	1	seq
aaCcugccaguugaagaacugu	1	1	seq
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aagcugccaguugaagaCcugu	2	1	seq
aagcugccaguugCagaacugu	3	1	seq
aagcugccaguugUagaacugu	10	1	seq
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Uagcugccaguugaagaacuguu	1	1	seq
agcugccaguugaagaacu	13	0	seq
agcugccaguCgaagaacu	1	1	seq

hsa-miR-22-3p

hsa-miR-22-5p

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Cgcugccaguugaagaacug	4	1	seq
Cgcugccaguugaagaacugu	3	1	seq
agcugccaguugaagaacCgu	1	1	seq
agcugccaguugaagaacugu	13	0	seq
gcugccaguugaagaacu	1	0	seq