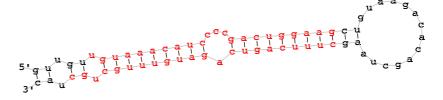
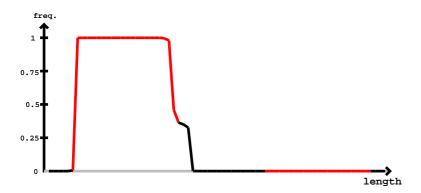
miRBase precursor : hsa-mir-30d
Total read count : 7383
hsa-miR-30d-5p read count : 7381
hsa-miR-30d-3p read count : 2
remaining reads : 0





hsa-miR-30d-3p

hsa-miR-30d-5p

iisa-mir-30d-3p			
5'- guuguuguaaacauccccgacuggaagcuguaagacacagcuaagcuuucagucag	-3'	exp	
((.(((((((((((((((((((((((((((((((((reads	mm	sample
uuguaaacauccccgacugga	24	0	seq
Guguaaacauccccgacugga	2	1	seq
uCguaaacauccccgacuggaa	1	1	seq
uuguaaacauccccgacuggaa	5	0	seq
uuguaaacauccccgacuggaaA	1	1	seq
uuguaaacauccccgacuggaagc	2	0	seq
uuguaaacauccccgacuggaagcu	12	0	seq
Cuguaaacauccccgacuggaagcu	1	1	seq
uuguaaaAauccccgacuggaagcu	1	1	seq
uguaaacauccccAacug	1	1	seq
uguaaacauccccgacug	24	0	seq
uguaCacauccccgacug	1	1	seq
uguaCacauccccgacugg	5	1	seq
uguaaacauccccgacugg	110	0	seq
Cguaaacauccccgacugg	1	1	seq
uguaaCcauccccgacugg	1	1	seq
uguaaacauccccgacugA	4	1	seq
uguaaacCuccccgacugg	1	1	seq
uguaaGcauccccgacugga	1	1	seq
uguaaacauccccgaUugga	1	1	seq
uguaaacauccccgacCgga	2	1	seq
uguaaacauccccgacGgga	1	1	seq
uCuaaacauccccgacugga	2	1	seq
uguUaacauccccgacugga	2	1	seq
uguaaacCuccccgacugga	1	1	seq
uguaaacauAcccgacugga	1	1	seq
uguaaCcauccccgacugga	2	1	seq
uguaaacaucUccgacugga	12	1	seq
uguaaacauccccgacuCga	2	1	seq
uguaaacaAccccgacugga	2	1	seq
uguaaacaucGccgacugga	1	1	seq
uguaaacGuccccgacugga	2	1	seq
uguaaUcauccccgacugga	2	1	seq
uguaaacauccccgGcugga	1	1	seq

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uUuaaacauccccgacugga	5	1	seq
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uguaaacauccccgacuAga	1	1	seq
uguaaacauccAcgacugga	1	1	seq
uguaaacauccccgacuggC	2	1	seq
uguaaacauccccgaAugga	2	1	seq
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uguaaacauccccAacugga	1	1	seq
uguaaacauccccgacugUa	1	1	seq
uguaaacaucccGgacugga	1	1	seq
uguaaacauccccCacugga	1	1	seq
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uguaaacauccGcgacugga	1	1	seq
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uguaaacauccccgacuggaagU	5	1	seq
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cuuucagucagauguuugcu	2	0	seq