miRBase precursor : hsa-mir-451a Total read count : 37495 hsa-miR-451a read count : 37458 remaining reads : 37





hsa-miR-451a

1010			
- cuugggaauggcaagg <mark>aaaccguuaccauuacugaguuu</mark> aguaaugguaaugguucucuugcuauacc	caga -3'	exp	
.(((((.(((((((((((((((((((((((((((((((())). read	ds mm	sample
Aaaaccguuaccauuacug	2	1	seq
gaaaccguuaccauuacug	1	0	seq
Aaaaccguuaccauuacuga	2	1	seq
Aaaaccguuaccauuacugag	5	1	seq
Aaaaccguuaccauuacugagu	45	1	seq
Caaaccguuaccauuacugagu	2	1	seq
Aaaaccguuaccauuacugaguu	13	1	seq
Aaaaccguuaccauuacugaguuu	1	1	seq
aaaccguuUccauuacug	1	1	seq
aCaccguuaccauuacug	4	1	seq
Gaaccguuaccauuacug	1	1	seq
aaaccguuGccauuacug	1	1	seq
	1	1	seq
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aaaccguuaccauuacuC	1	1	seq
aaaccguuaccauuaGug	1	1	seq
aUaccguuaccauuacug	1	1	seq
aaaccguuacGauuacug	1	1	seq
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aaaccguuaccauuacuA	2	1	seq
aaaccguuaccauuacCg	3	1	seq
aaaccguuaccaCuacug	1	1	seq
aaaccguuaccauuacGg	7	1	seq
aaaccguuaccauuacug	219	1 0	seq
aaaccgAuaccauuacug	3	1	seq
aaaccguuCccauuacug	5	1	seq
	151	1	seq
aaaccguuacUauuacug	1	1	seq
aaaccguuaccauuaUug	1	1	seq
	1	1	seq
aaaccguuaccauuacuga	1009	9 0	seq
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aaaccguuacUauuacuga	2	1	seq
aGaccguuaccauuacuga	1	1	seq
aaaccguuaccauuacugG	6	1	seq

cuugggaauggcaagg <mark>aaaccguuaccauuacugaguu</mark> uaguaaugguaaugguucucuugcuauacccaga			
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aaaccguuaccauCacuga	1	1	seq
aUaccguuaccauuacuga	1	1	seq
aaaccguuacAauuacuga	1	1	seq
aCaccguuaccauuacuga	3	1	seq
Caaccguuaccauuacuga	45	1	seq
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aaaccguuacGauuacugag	1	1	seq seq
aaaccguuaccauuCcugag	2	1	seq
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aaCccguuaccauuacugag	2	1	seq
aaaccguuaccauuacugaC	3	1	seq
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aaaccguuaccauuacugCgu	9	1	seq
aaaccguuaccauuGcugagu	3	1	seq
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Caaccguuaccauuacugagu	1311	1	seq
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aaaccguuacAauuacugagu	9	1	seq
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	2	1	seq
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ccguuaccauuacugagu	2	0	seq
ccguuaccauuacugaguu	1	0	seq
	1	0	seq
	3	0	seq
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guuaccauuacugaguuu	3	0	seq
uuaccauuacugaguuuagu	1	0	seq
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	-	-	