

[illegible]

hsa-miR-376b

caguccuucuuugguauuuaaaacguggauauuccuucuauguuuacgugauuccugguua**aucauagagggaaauccauguuuucaguaucaaaugcug**

Sequence	Count	Count	Species
.....aUCAUAGAGGAAAAACCAUGUUU.....	1	1	hsa
.....aUCAUAGAGGAAAAUCUUGUUUU.....	1	1	hsa
.....aUCAUAGAGGAAAAUCCAUGUUUA.....	3	1	hsa
.....aUCGUAGAGGAAAAUCCAUGUUU.....	1	1	hsa
.....aUCAUAGAGGAAAAUCCAUGUUC.....	1	1	hsa
.....aUCAUAGAGGAAAAUCCAUGUUU.....	9	1	hsa
.....aUCAUAGAGGAAAAUCGUGUUUU.....	1	1	hsa
.....aUCAUAGAGGAAAAUCCAUGUUUU.....	6	0	hsa
.....aUCAUAGAGGAAAAUCCAUGUUUC.....	1	1	hsa
.....UCAUAGAGGAAAAUCCAUGU.....	2	0	hsa
.....UCAUAGAGGAAAAUCCAUGU.....	1	1	hsa
.....UCAUAGAGGAAAAUCCAUGUU.....	1	0	hsa
.....UCAUAGAGGAAAAUCCAUGUUU.....	10	0	hsa
.....AUAGAGGAAAAUCCAUGU.....	1	0	hsa
.....AUAGAGGAAAAUCCAUGU.....	1	1	hsa
.....AUGGAGGAAAAUCCAUGU.....	1	1	hsa
.....AUAGAGGAAAAUCCAUGUUU.....	3	0	hsa
.....AUAGAGGAAAAUCCAUGUUUU.....	1	1	hsa