

☐ miRDeep

min score -10

submit

progress

miRNA ID	Score	expr	loci	sequence
hsa-mir-21	1,450.72	2,818	chr17+55273394~55273498	UAGCUUUAUCAGACUGAUGUUGAC
hsa-mir-221	1,396.14	2,709	chrX-45490531~45490633	AGCUACAUUGUCUGCGUGGUUU
hsa-mir-29a	1,390.85	2,728	chr7-130212025~130212127	UAGCACCAUCUGAAACGGGUUA
hsa-mir-29b	1,377.12	1,825	chr7-25056045~25056147	UACAGUCCAGUACAGAAUUCU

[click for details](#)

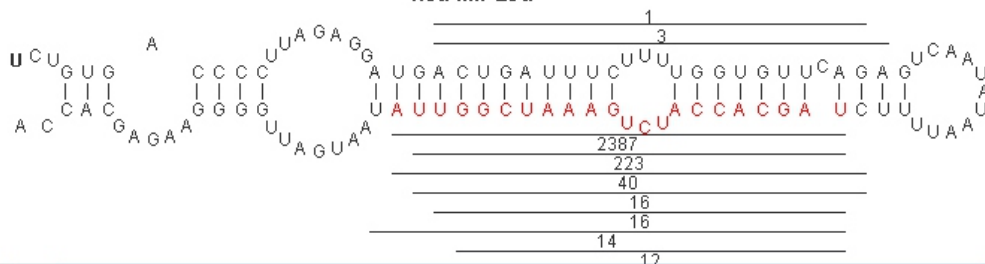


Target genes

ucugugaccccuagagggaugacugauuucuuuuggguguucagagucacaaauuuuucUAGCACAUCUGAAAUCGGUUAuaaugauuggggaaga
gcacca

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))))..

hsa-mir-29a



target Genes

targetGene	RefSeq	contextScore	PCT_score
PI15	NM_015886	-0.346	3.612
COL4A5	NM_000495	-0.358	1.881