

Scale
chr1:

36131200| 36131300| 36131400| 36131500| 36131600| 36131700|

200 bases

EvoFold Predictions of RNA Secondary Structure

Perfect Matches to Short Sequence (AUAAAGCUAGAUAAACCGAAAGU)

UCSC Genes Based on RefSeq, UniProt, GenBank, CCDS and Comparative Genomics

EIF2C1
EIF2C1



Human mRNAs from GenBank



Non-Human RefSeq Genes

Other RefSeq

Ensembl Gene Predictions - archive Ensembl 54 - may2009

ENST00000373206
ENST00000373204



TargetScan miRNA Regulatory Sites

Simple Nucleotide Polymorphisms (dbSNP build 128)

SNPs (128)

C/D and H/ACA Box snoRNAs, scaRNAs, and microRNAs from snoRNABase and miRBase

Reported Poly(A) Sites from PolyA_DB

Predicted Poly(A) Sites Using an SVM