

Diagram illustrating the secondary structure of a single-stranded RNA molecule. The sequence is 5'-G C U C U U U U C A C A U U G U G C U A C U G U C U G A C A C U G A C A C-3'. The structure shows several base pairs forming stems and loops. The 3' end is labeled '3' U A'.

[illegible]

Star

Mature

agggccggcgaugccucugcugcugggccagagcucuuuucacauugugcuacugucugcaccugucacuaagcagugcaauguuuaaagggcgauuggccguguagugcuacc

.....cagugcaauAuuaaaagggcgau.....	2	1	hsa
.....cGgugcaauguuuaaagggcgau.....	3	1	hsa
.....cGgugcaauguuuaaagggcgau.....	3	1	hsa
.....cagCgcaauguuuaaagggcgau.....	3	1	hsa
.....cagugcGauguuuaaagggcgau.....	2	1	hsa
.....cagugcaauguuuaaagggcgau.....	1	1	hsa
.....cagugcaauUuuuaaagggcgau.....	5	1	hsa
.....cagugcaauguuuaaaggAcgau.....	3	1	hsa
.....Aagugcaauguuuaaagggcgau.....	1	1	hsa
.....cagugcaauguuuaaagggcgau.....	2937	0	hsa
.....cagugcaauguuuaaaggUcgau.....	7	1	hsa
.....cagugcaauguuUaaagggcgau.....	2	1	hsa
.....cagGgcaauguuuaaagggcgau.....	3	1	hsa
.....caAugcaauguuuaaagggcgau.....	2	1	hsa
.....cagugcaauguuuaaaggCcgau.....	2	1	hsa
.....cagugcaauguuuaaagggcgau.....	1	1	hsa
.....caUugcaauguuuaaagggcgau.....	3	1	hsa
.....cagugcaauguuuaaagggcCu.....	1	1	hsa
.....cagugcaauguuuaaagggcCaC.....	293	1	hsa
.....cagugcaauguuuaaagggcgau.....	7	1	hsa
.....caguUcaauguuuaaagggcgau.....	5	1	hsa
.....cagugcaauguuCaaagggcgau.....	1	1	hsa
.....cagugcaauguuuaaagggcCaA.....	7	1	hsa
.....cagugcaUuguuuaaagggcgau.....	1	1	hsa
.....cagugcaauguAaaaagggcgau.....	2	1	hsa
.....cagugcaauguuuaaagggcGau.....	1	1	hsa
.....cagugAaauguuuaaagggcgau.....	1	1	hsa
.....cagugcaaugAuuaaagggcgau.....	4	1	hsa
.....cagugcaaugCuuaaagggcgaUU.....	1	1	hsa
.....cagugcaauguCaagggcgaUU.....	1	1	hsa
.....caguUcaauguuuaaagggcgaUU.....	2	1	hsa
.....cagugcaauguuuaaagggcgaUU.....	1	1	hsa
.....cagugcaauguuuaaagggcgaUG.....	14	1	hsa
.....cagugcaaugAuuaaagggcgaUU.....	2	1	hsa
.....cagugcaauguuuaaagggcgaUA.....	196	1	hsa
.....cagugcaauguuuaaagggcCaCu.....	25	1	hsa
.....cagugcaauguuUagggcgaUU.....	1	1	hsa
.....caCugcaauguuuaaagggcgaUU.....	1	1	hsa
.....cagugcaGuguuuaaagggcgaUU.....	1	1	hsa
.....cagugcaauguuuaaagggcgaUC.....	14	1	hsa
.....cagugcaauguuuaaagggcgaUU.....	354	0	hsa
.....NagugcaauguuuaaagggcgaUU.....	1	1	hsa
.....cagugcaauUuuuaaagggcgaUU.....	1	1	hsa
.....UagugcaauguuuaaagggcgaUU.....	1	1	hsa
.....cagugcaauguuuaaagggcgaUUA.....	28	1	hsa
.....cagugcaauguuuaaagggcgaUUU.....	72	1	hsa
.....cagugcaauguuuaaagggcgaUAg.....	1	1	hsa
.....cagugcaauguuuaaagggcgaUAgg.....	1	1	hsa
.....agugcaauguuuaaagggc.....	1	0	hsa
.....agugcaauguuuaaagggcA.....	1	0	hsa
.....agugcaauguuuaaagggcAC.....	1	1	hsa
.....Ngugcaauguuuaaagggcgau.....	1	1	hsa
.....agugcaauguuuaaagggcgau.....	13	0	hsa
.....agugcaauguuuaaagggcgaUU.....	2	0	hsa
.....Cugcaauguuuaaagggcgau.....	1	1	hsa
.....Ccaauguuuaaagggcgau.....	1	1	hsa