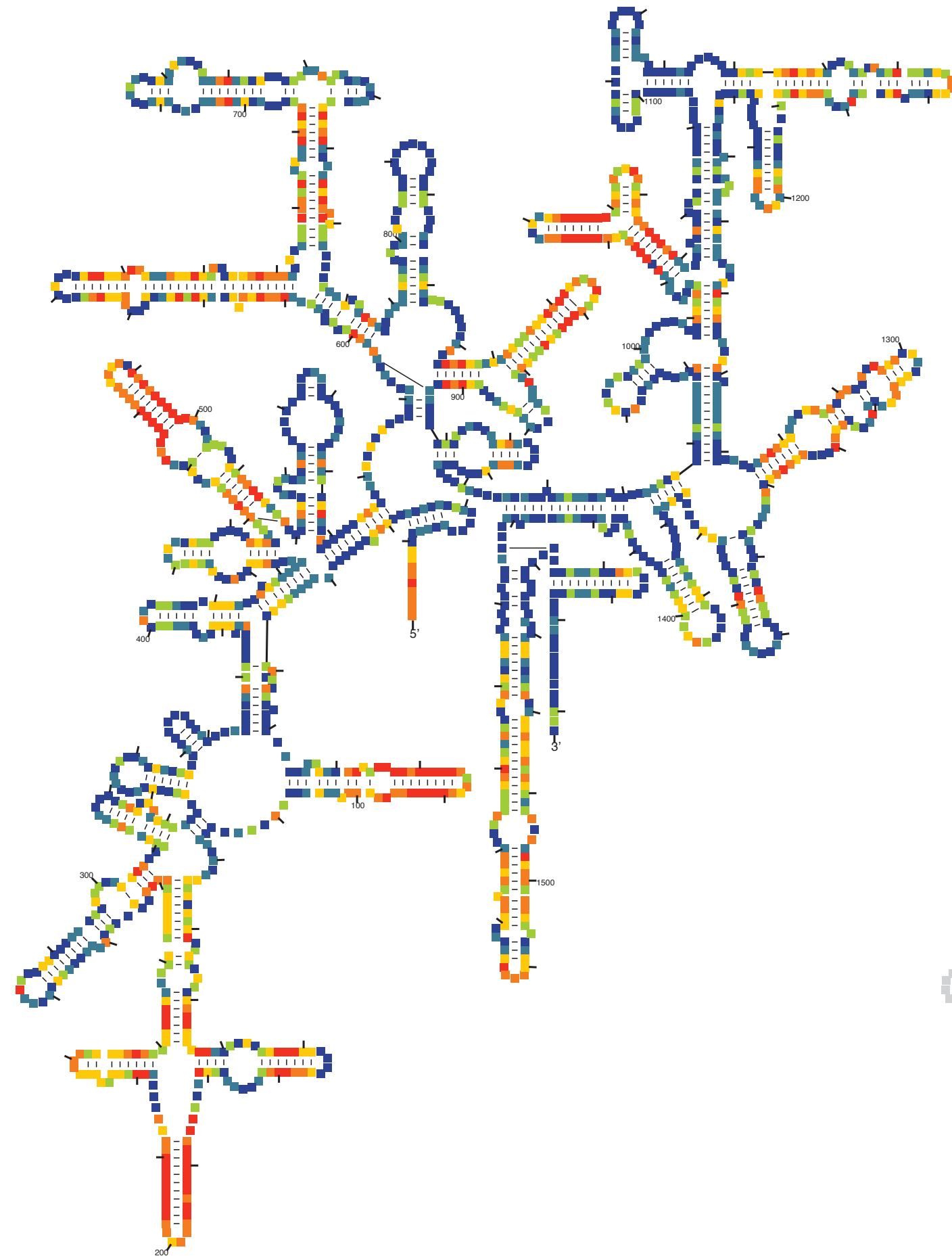


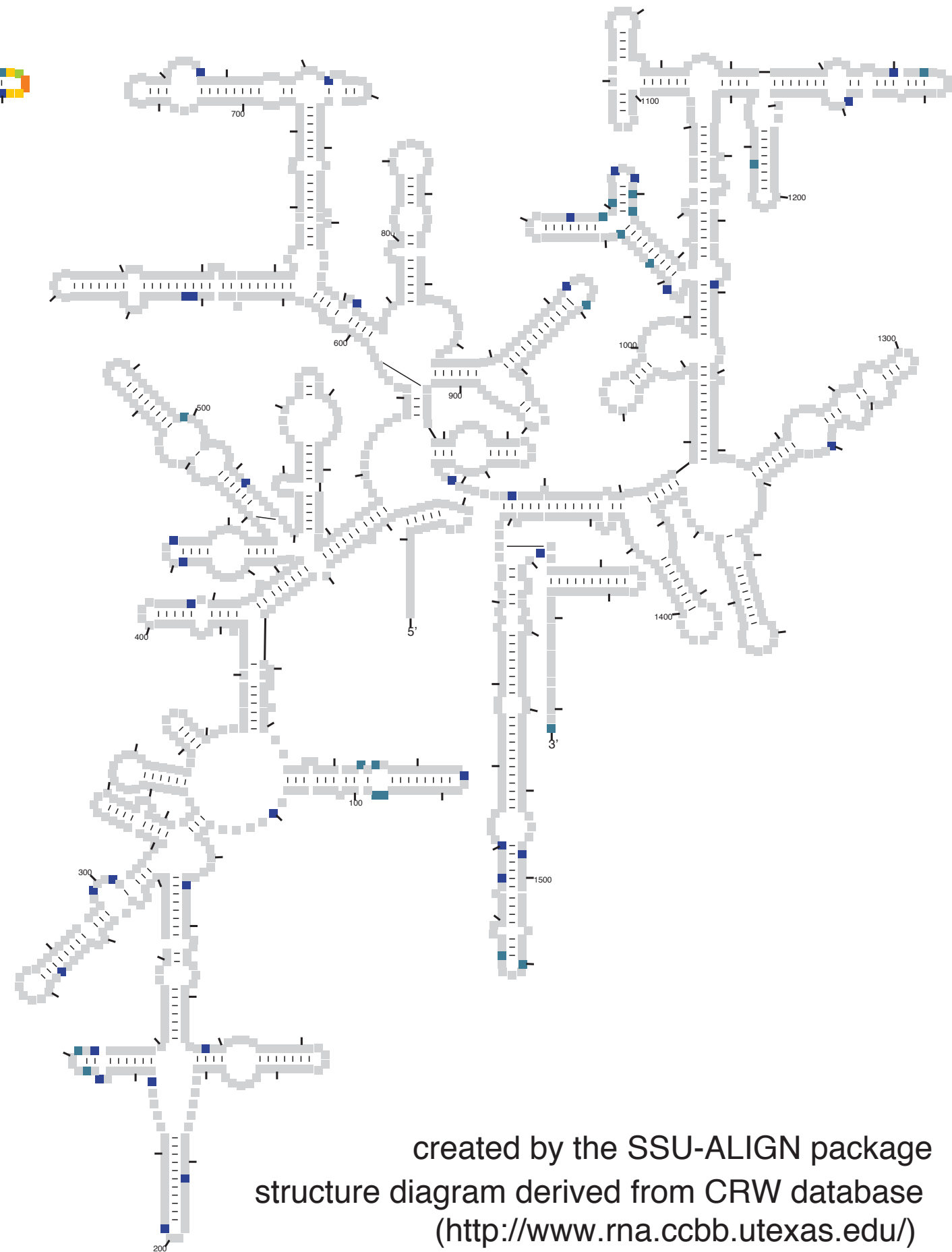
Sequence conservation per position

blue: highly conserved red: highly variable



Frequency of insertions after each position

grey: zero to very few inserts teal: 1-2% inserts



created by the SSU-ALIGN package
structure diagram derived from CRW database
(<http://www.rna.cccb.utexas.edu/>)