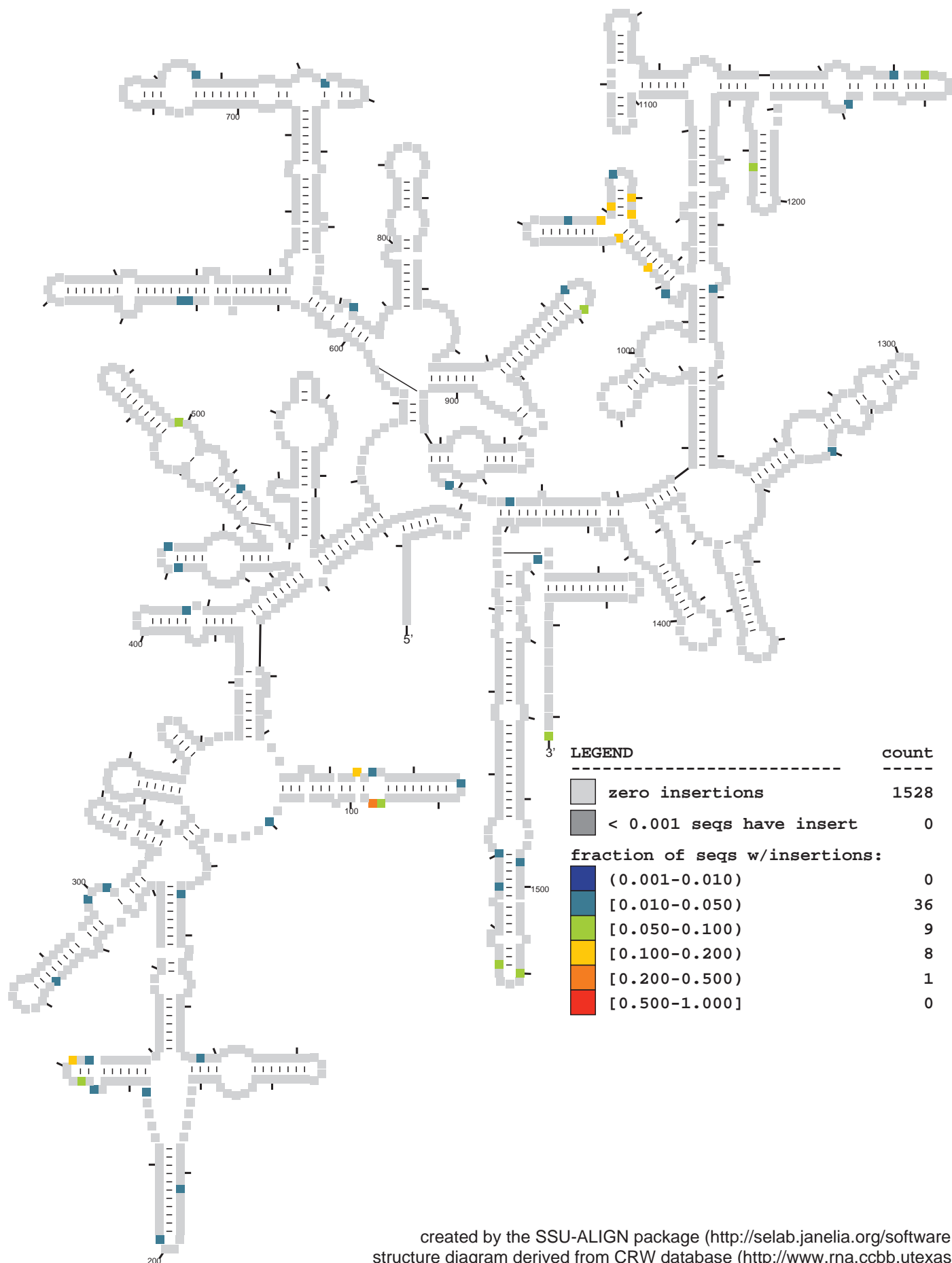


model	#pos	#bps	#seqs	description
bacteria	1582	480	93	frequency of insertions after each position



created by the SSU-ALIGN package (<http://selab.janelia.org/software.html>)
 structure diagram derived from CRW database (<http://www.rna.ccbb.utexas.edu/>)