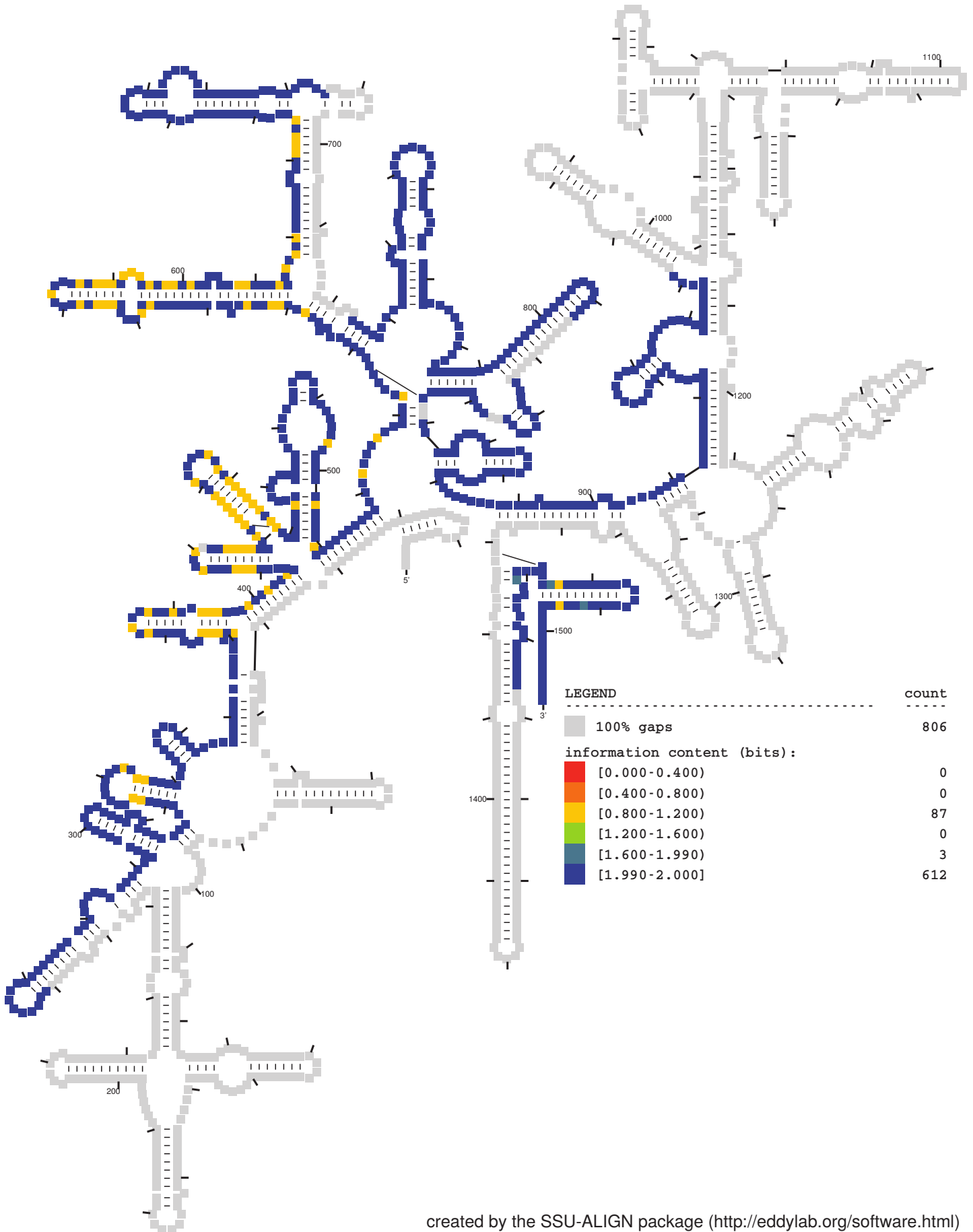
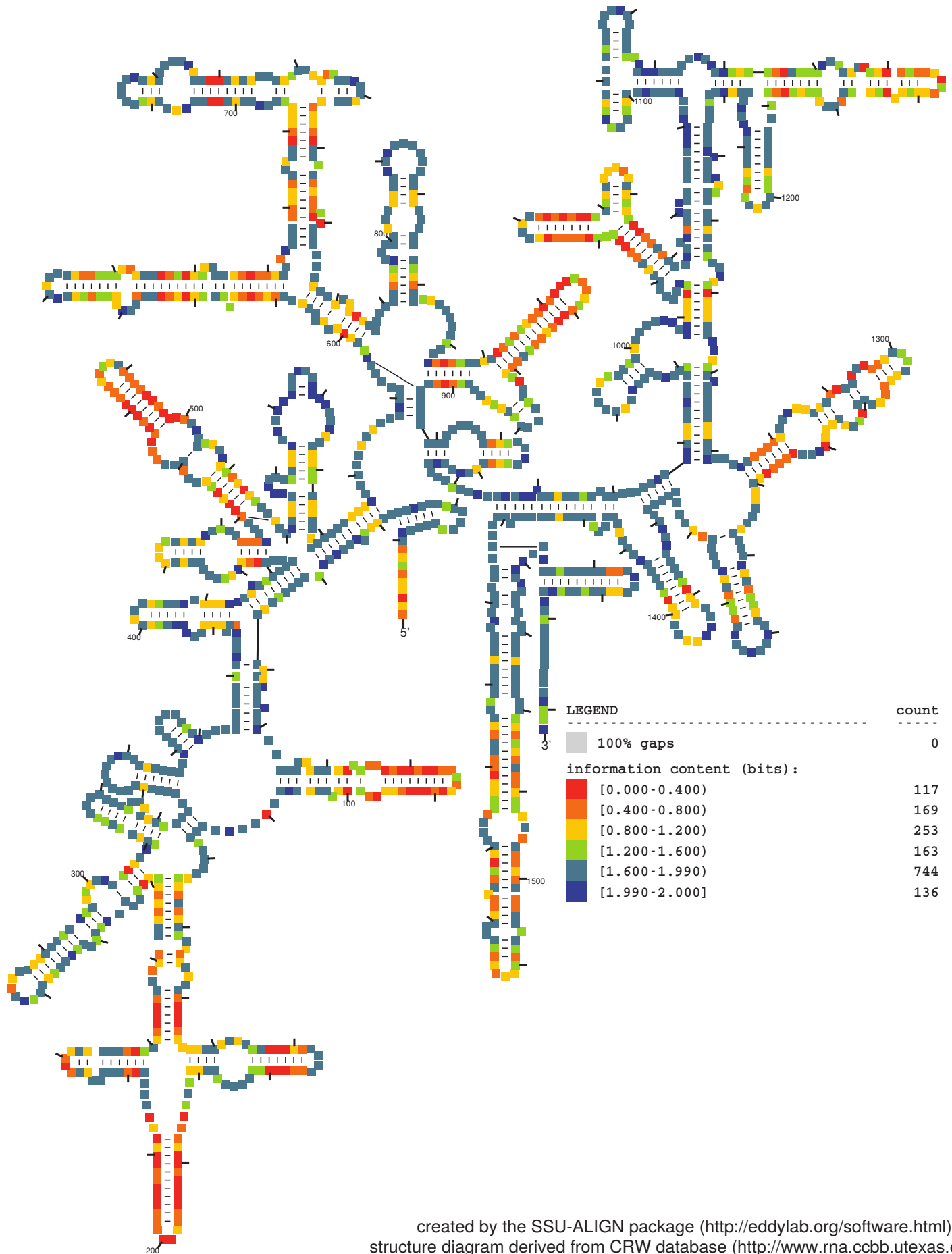


model	#pos	#bps	#seqs	description
archaea	1508	471	23	information content per position



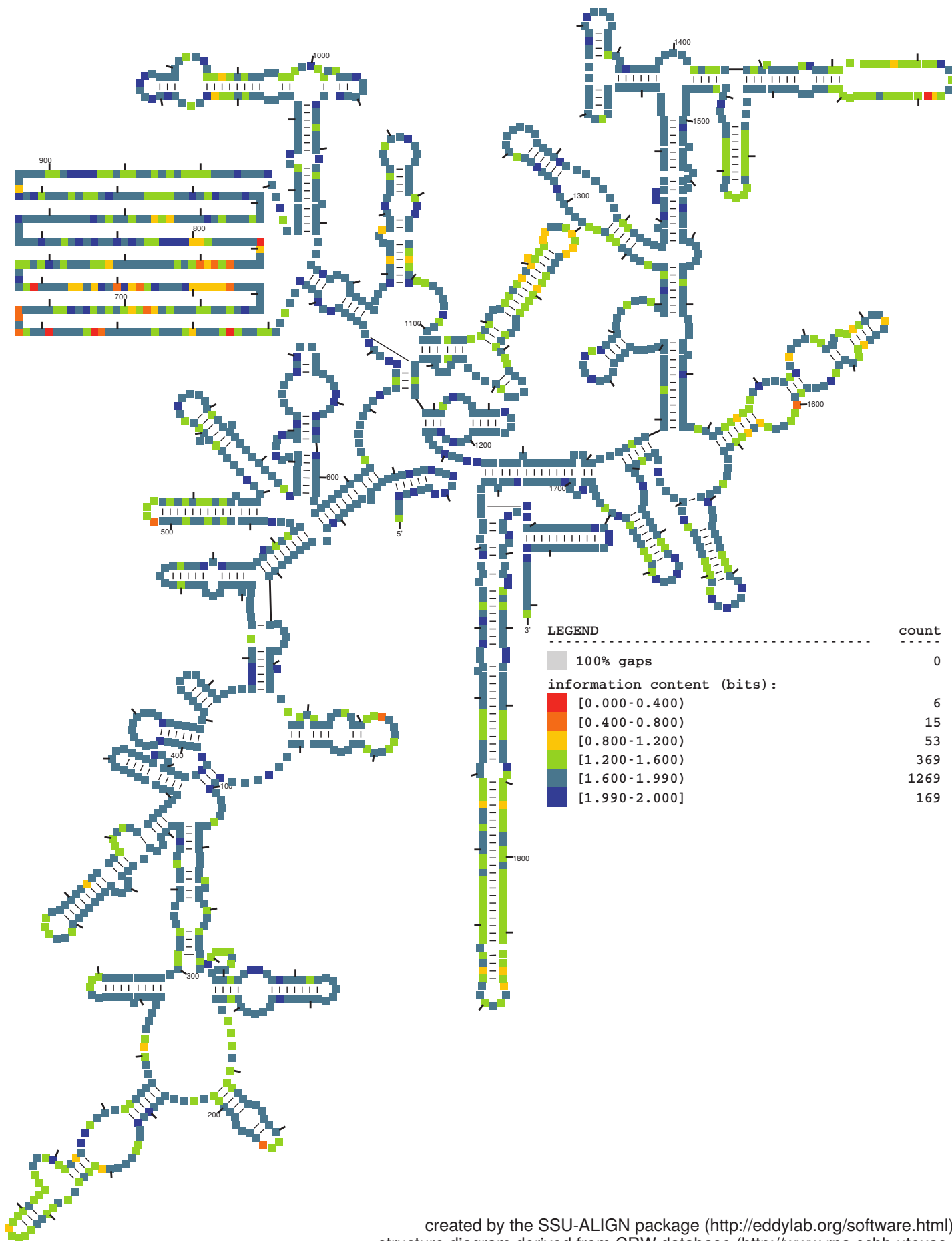
created by the SSU-ALIGN package (<http://eddylib.org/software.html>)  
structure diagram derived from CRW database (<http://www.rna.ccbb.utexas.edu/>)

model	#pos	#bps	#seqs	description
bacteria	1582	480	6313	information content per position



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

model	#pos	#bps	#seqs	description
eukarya	1881	448	6328	information content per position



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