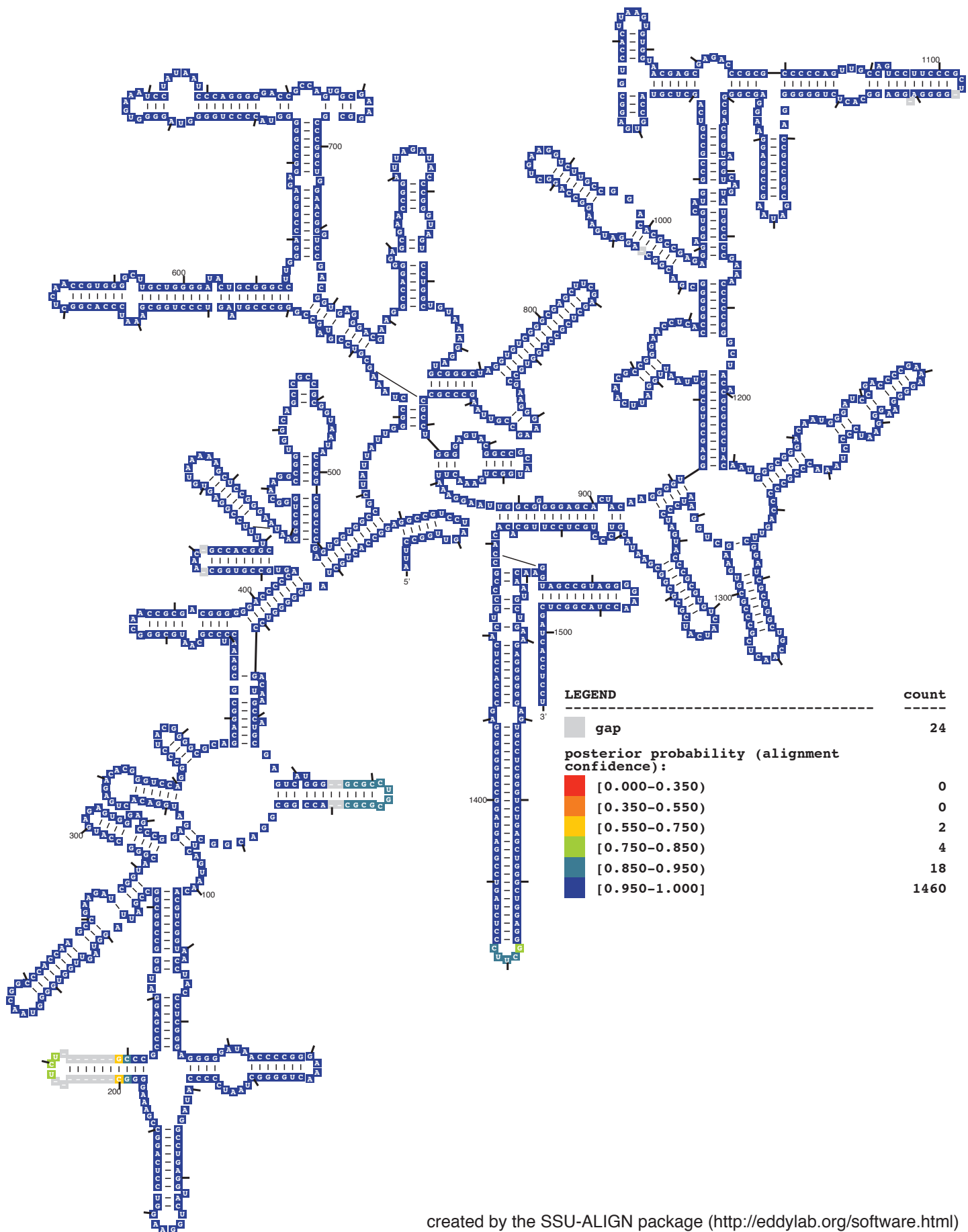


model	#pos	#bps	seqlen	sequence name
archaea	1508	471	1486	gi 176199 gb M21529.1 THCRRD



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