model	#pos	#bps	seqlen	sequence name
archaea	1508	471	1486	00121::Thermococcus_celer::M21529

LEGEND count
--- -----

gap 24

posterior probability (alignment confidence):

	count
[0.000–0.350)	0
[0.350–0.550)	0
[0.550–0.750)	2
[0.750–0.850)	4
[0.850–0.950)	18
[0.950–1.000]	1460
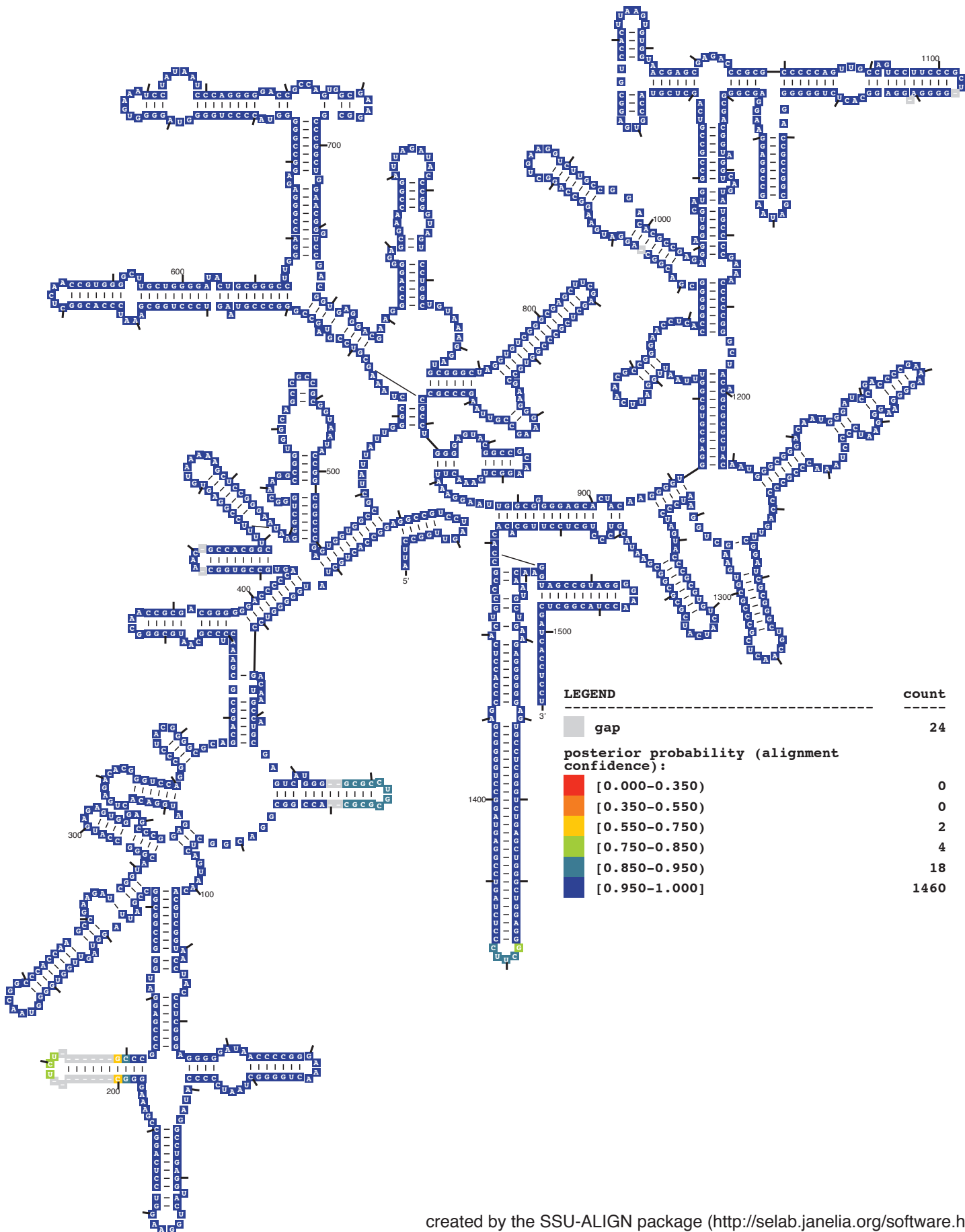
created by the SSU-ALIGN package (http://selab.janelia.org/software.html)
structure diagram derived from CRW database (http://www.rna.ccbb.utexas.edu/)