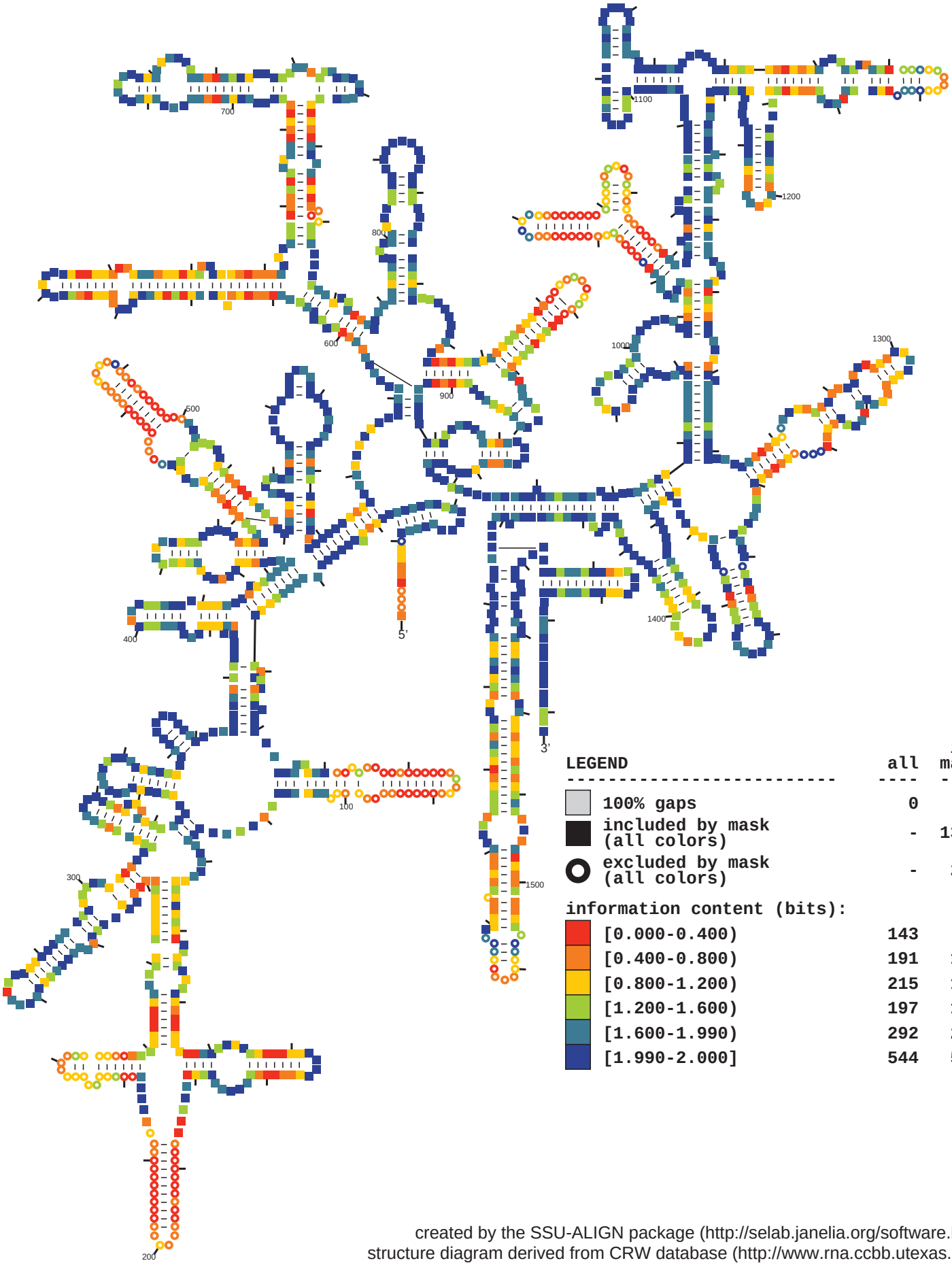
model	#pos	#bps	#seqs	description
bacteria	1582	480	93	information content per position



LEGEND	all	in mask
100% gaps	0	0
included by mask (all colors)	-	1376
excluded by mask (all colors)	-	206
information content (bits):		
[0.000-0.400)	143	72
[0.400-0.800)	191	132
[0.800-1.200)	215	178
[1.200-1.600)	197	181
[1.600-1.990)	292	282
[1.990-2.000]	544	531

created by the SSU-ALIGN package (http://selab.janelia.org/software.html)
structure diagram derived from CRW database (http://www.rna.ccbb.utexas.edu/)
alignment file: ../../seeds/bacteria-0p1.p.stk; mask file: ../../seeds/masks/bacteria-0p1.mask;