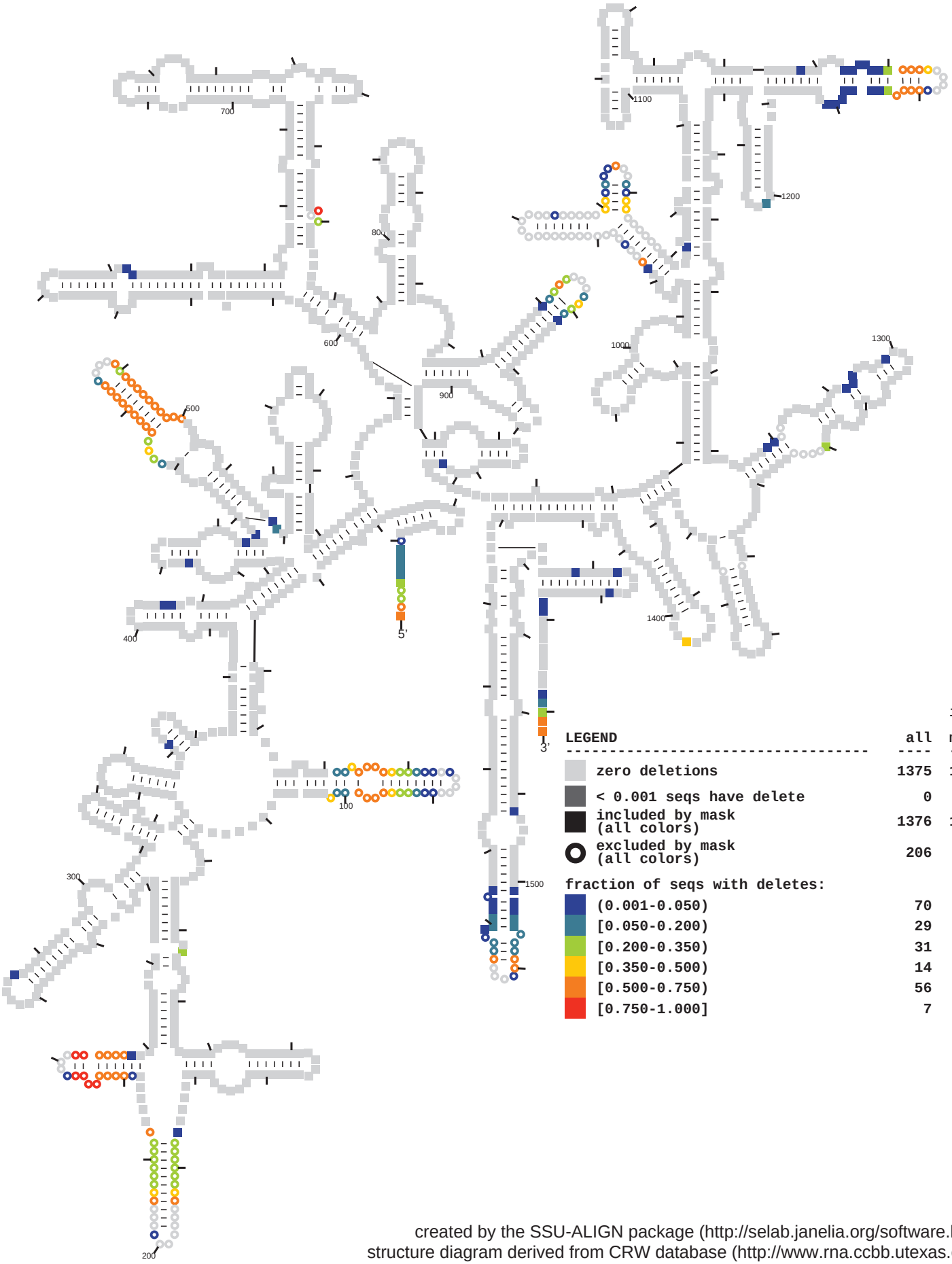
model	#pos	#bps	#seqs	description
bacteria	1582	480	93	frequency of deletions at each position



LEGEND	all	incl by mask
zero deletions	1375	1304
< 0.001 seqs have delete	0	0
included by mask (all colors)	1376	1376
excluded by mask (all colors)	206	0
fraction of seqs with deletes:		
(0.001-0.050)	70	51
[0.050-0.200)	29	11
[0.200-0.350)	31	6
[0.350-0.500)	14	1
[0.500-0.750)	56	3
[0.750-1.000]	7	0

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