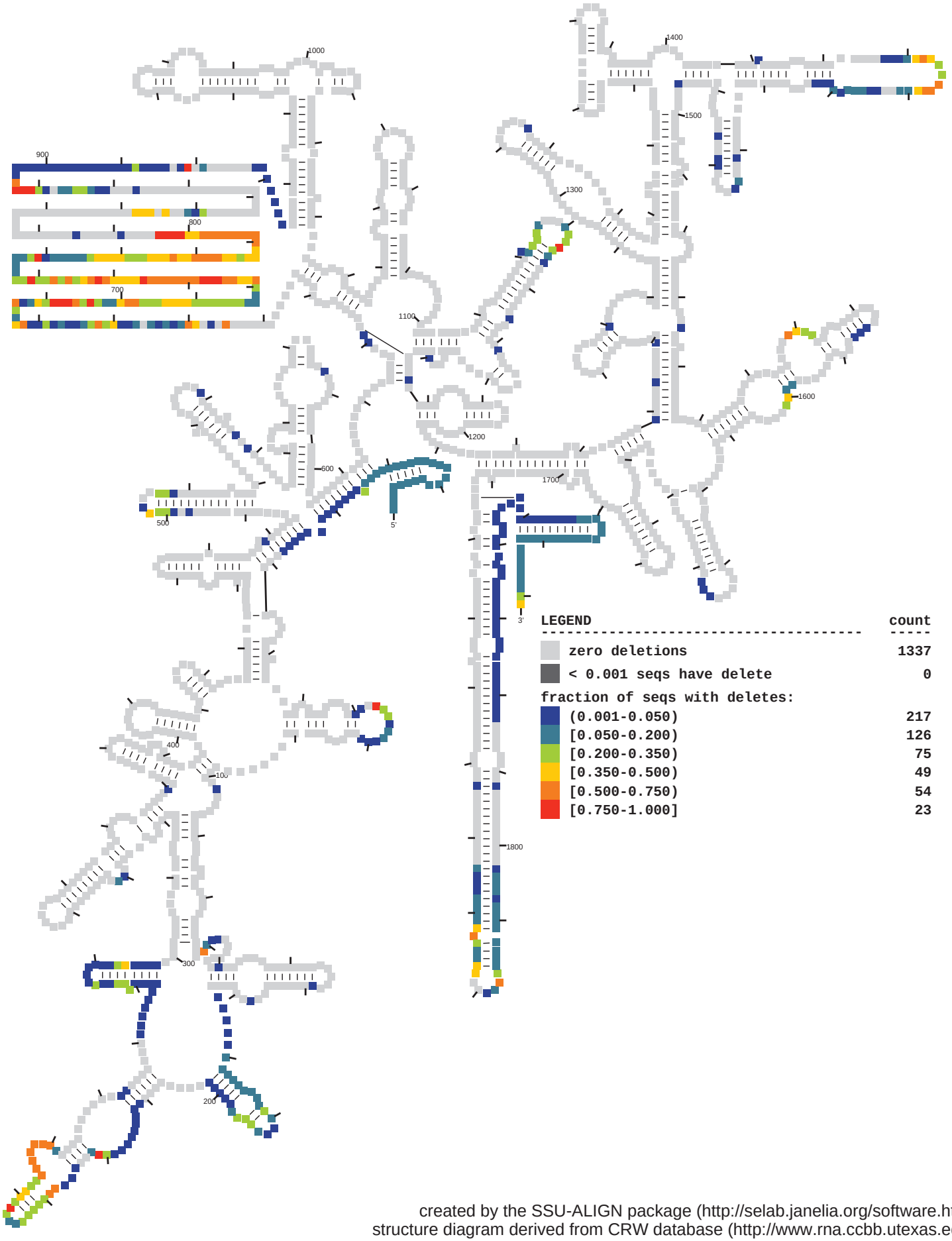
model	#pos	#bps	#seqs	description
eukarya	1881	448	89	frequency of deletions at each position



LEGEND	count
zero deletions	1337
< 0.001 seqs have delete	0
fraction of seqs with deletes:	
(0.001-0.050)	217
[0.050-0.200)	126
[0.200-0.350)	75
[0.350-0.500)	49
[0.500-0.750)	54
[0.750-1.000]	23

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