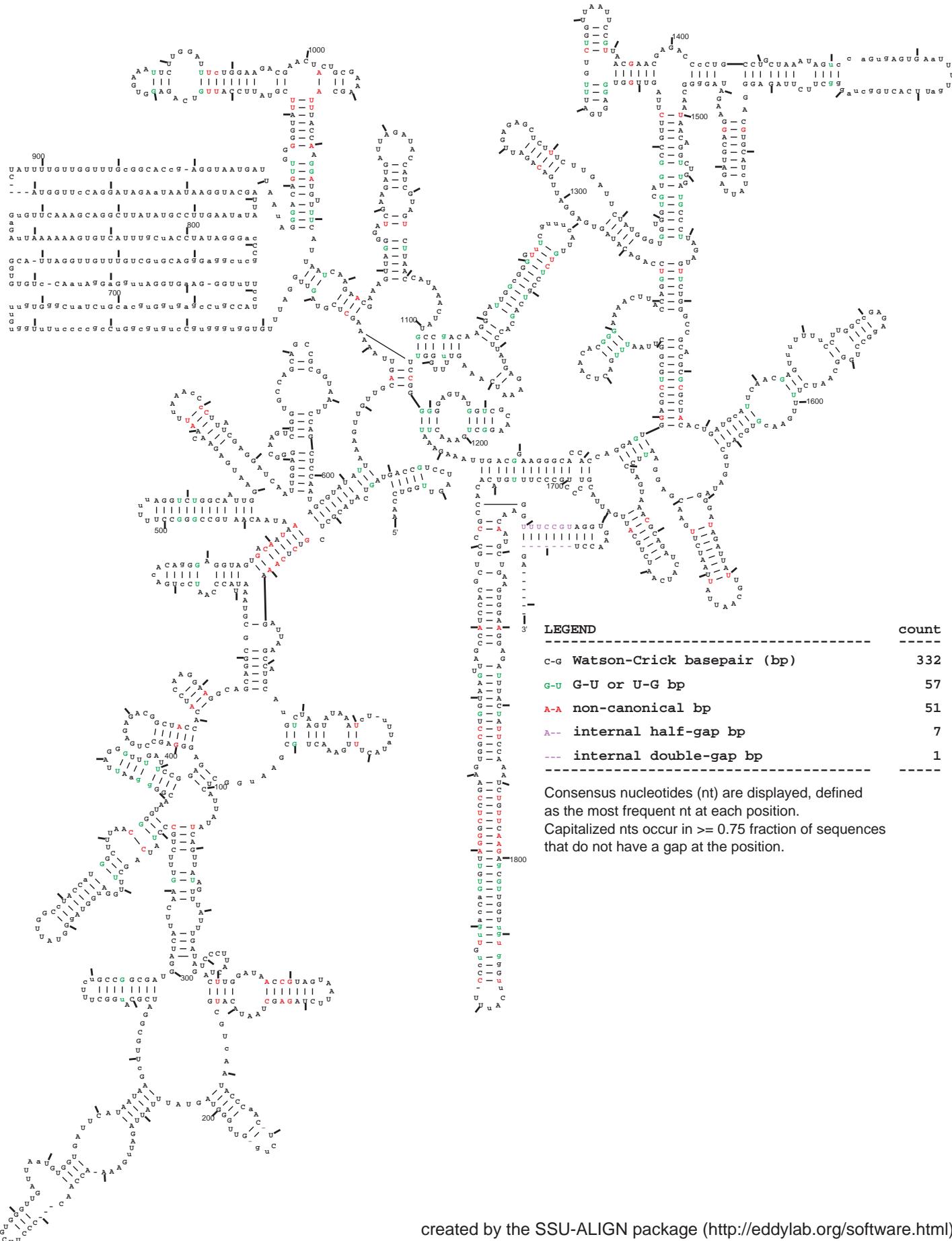


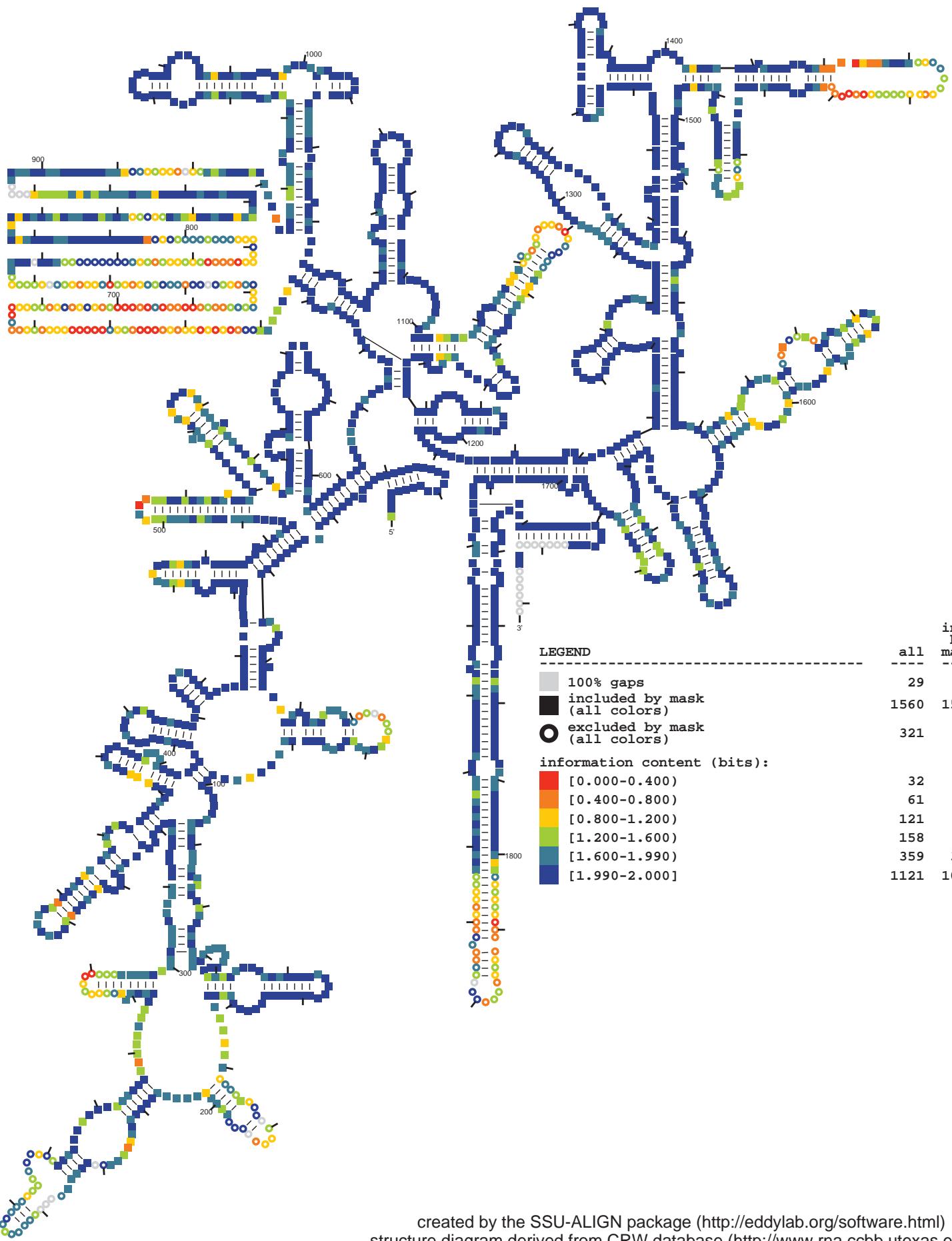
model #pos #bps
 eukarya 1881 448

sequence name
 alignment consensus sequence



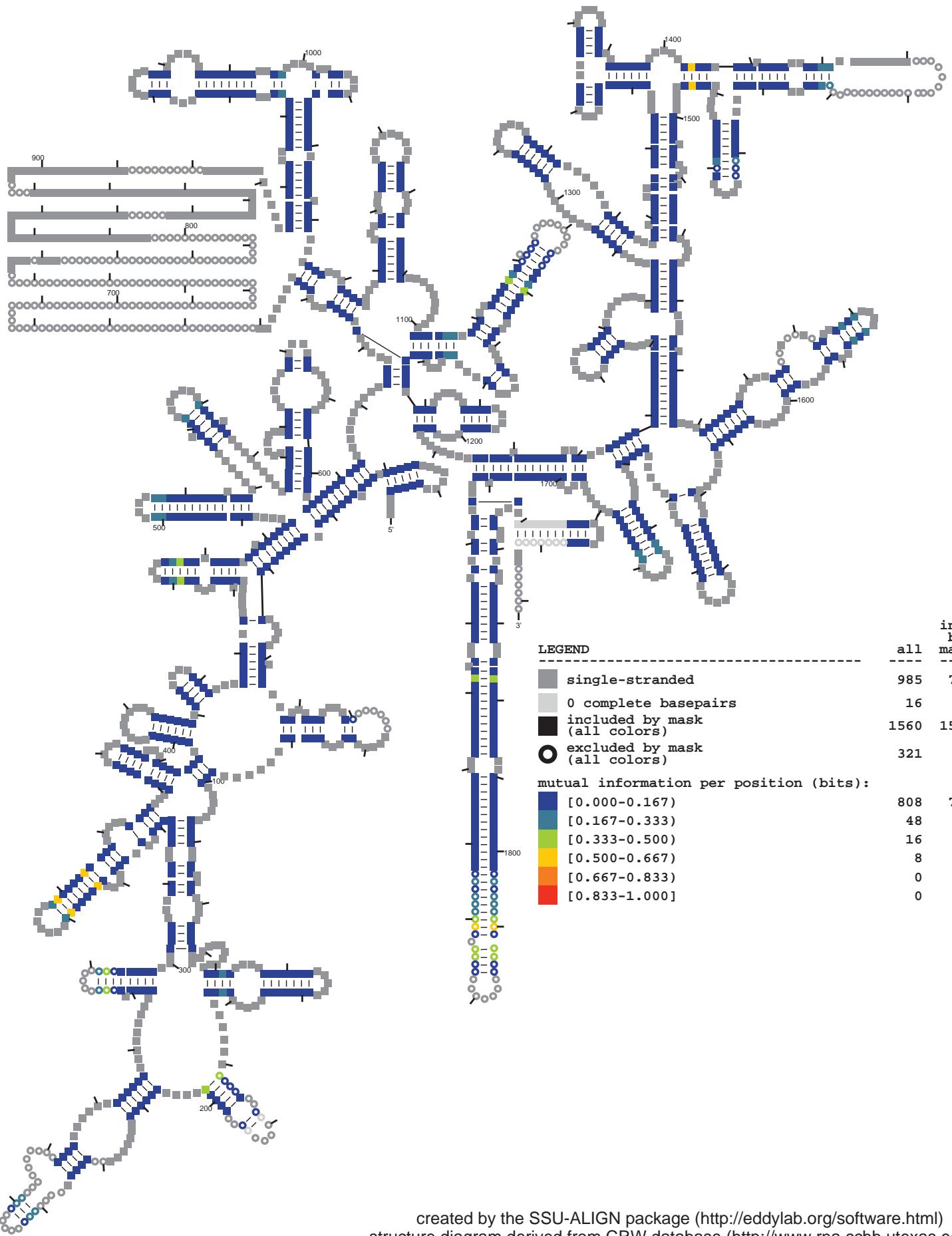
created by the SSU-ALIGN package (<http://eddylab.org/software.html>)
 structure diagram derived from CRW database (<http://www.rna.ccbb.utexas.edu/>)

model	#pos	#bps	#seqs	description
eukarya	1881	448	94	information content per position



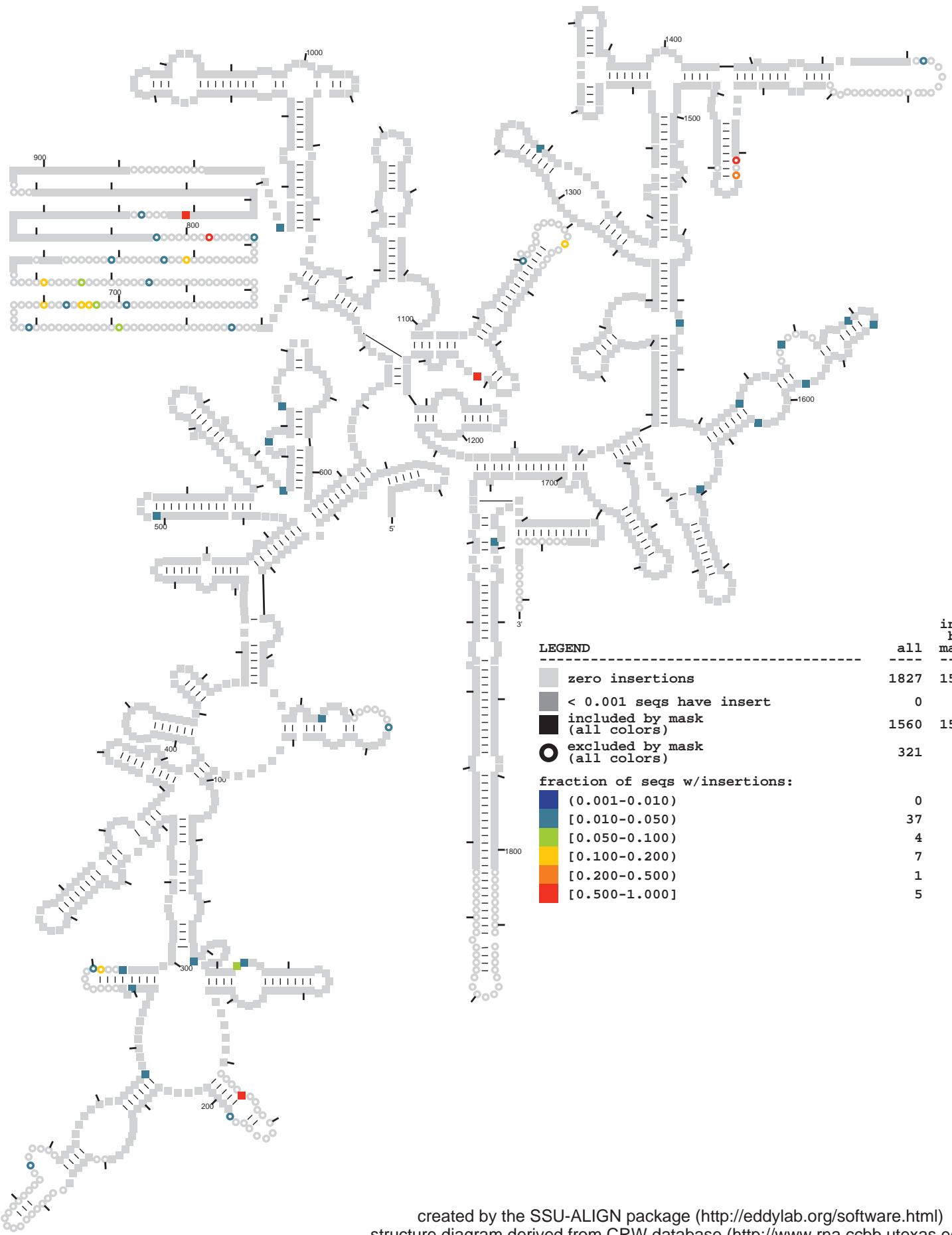
created by the SSU-ALIGN package (<http://eddylab.org/software.html>)
structure diagram derived from CRW database (<http://www.rna.ccbb.utexas.edu/>)

model	#pos	#bps	#seqs	description
eukarya	1881	448	94	mutual information per basepaired position



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

model	#pos	#bps	#seqs	description
eukarya	1881	448	94	frequency of insertions after each position

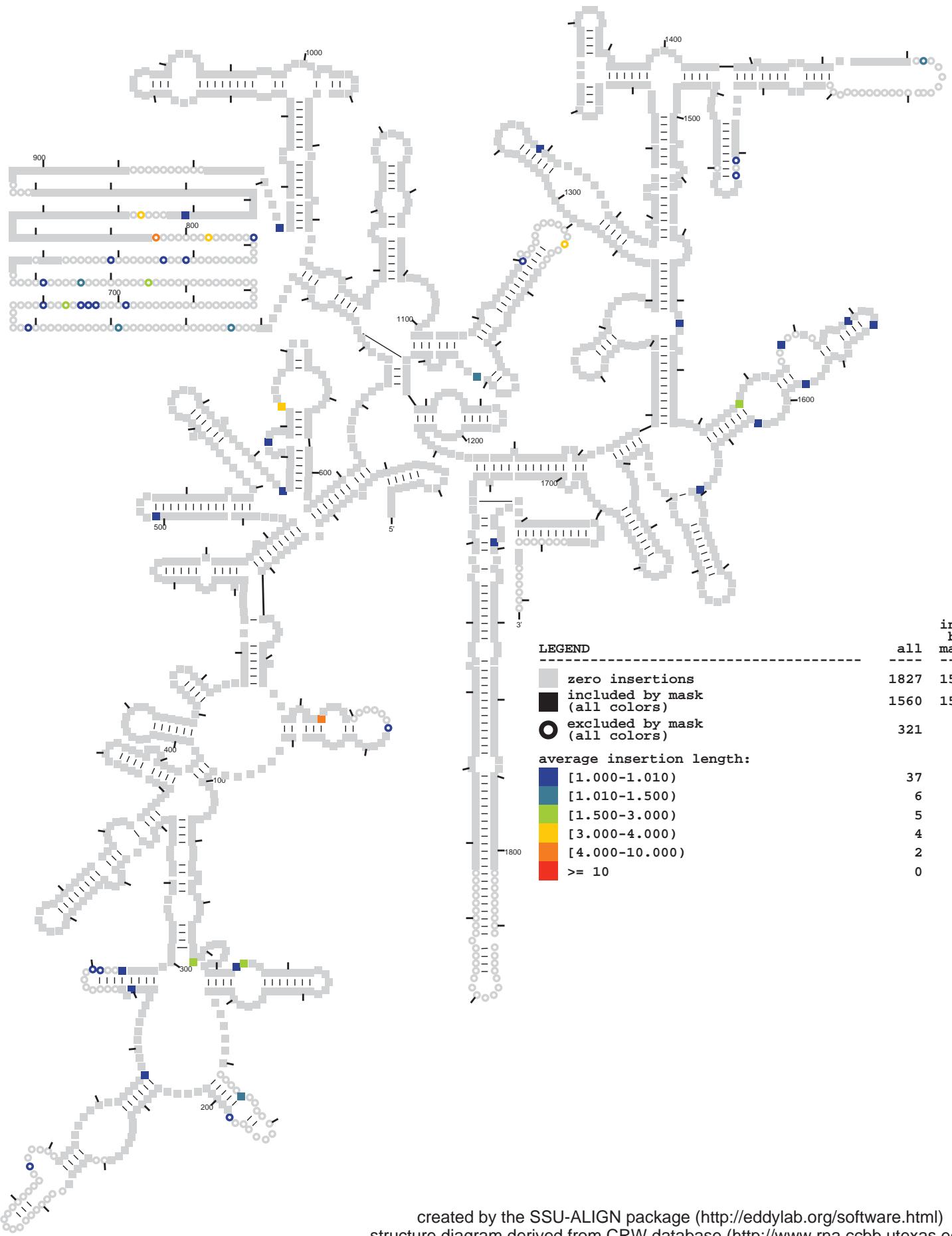


created by the SSU-ALIGN package (<http://eddylab.org/software.html>)
structure diagram derived from CRW database (<http://www.rna.ccbb.utexas.edu/>)

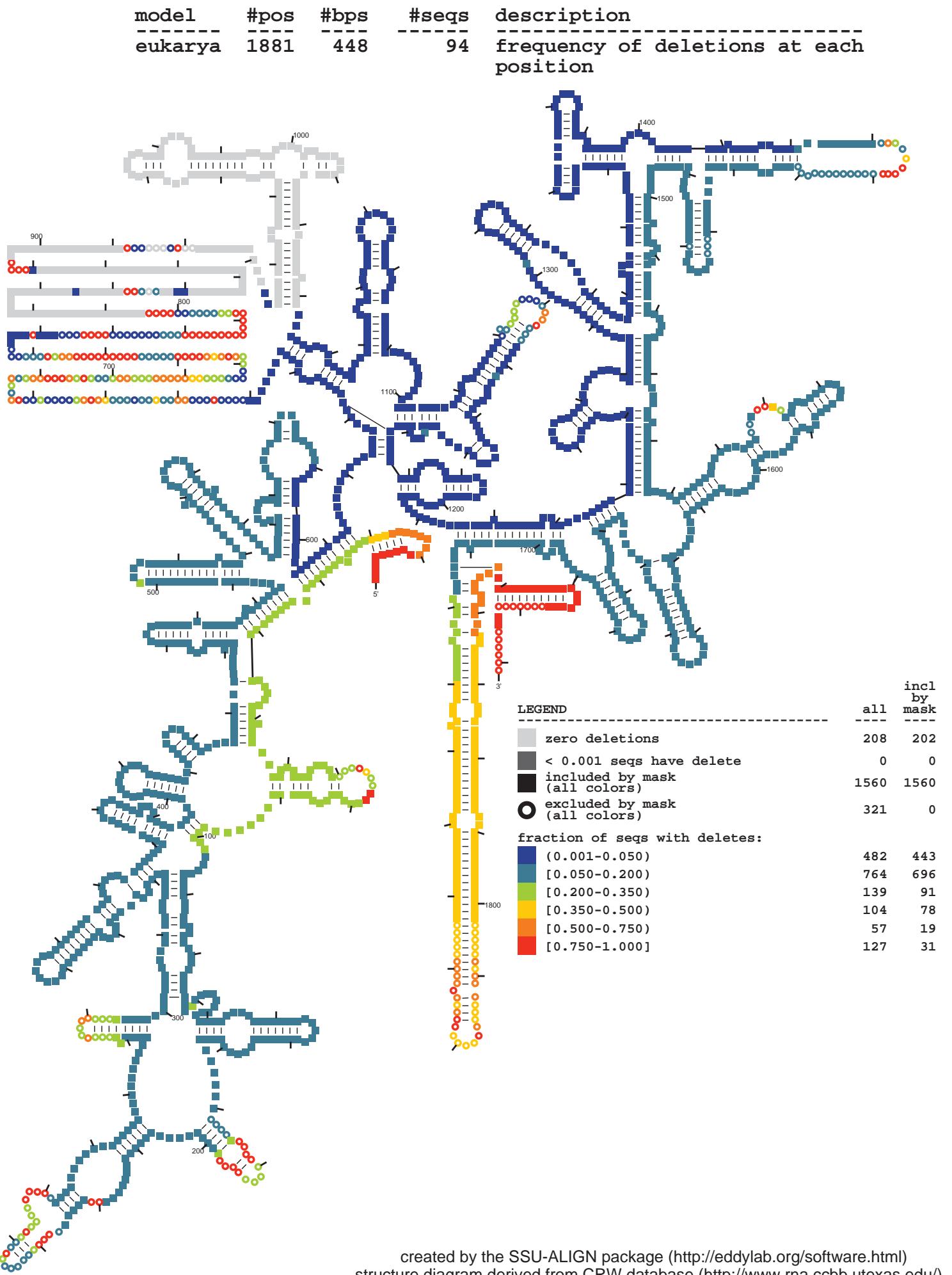
alignment file: raphids_ssu/raphids_ssu.eukarya.stk; mask file: raphids_ssu/raphids_ssu.eukarya.mask;

page 4

model	#pos	#bps	#seqs	description
eukarya	1881	448	94	average insertion length after each position

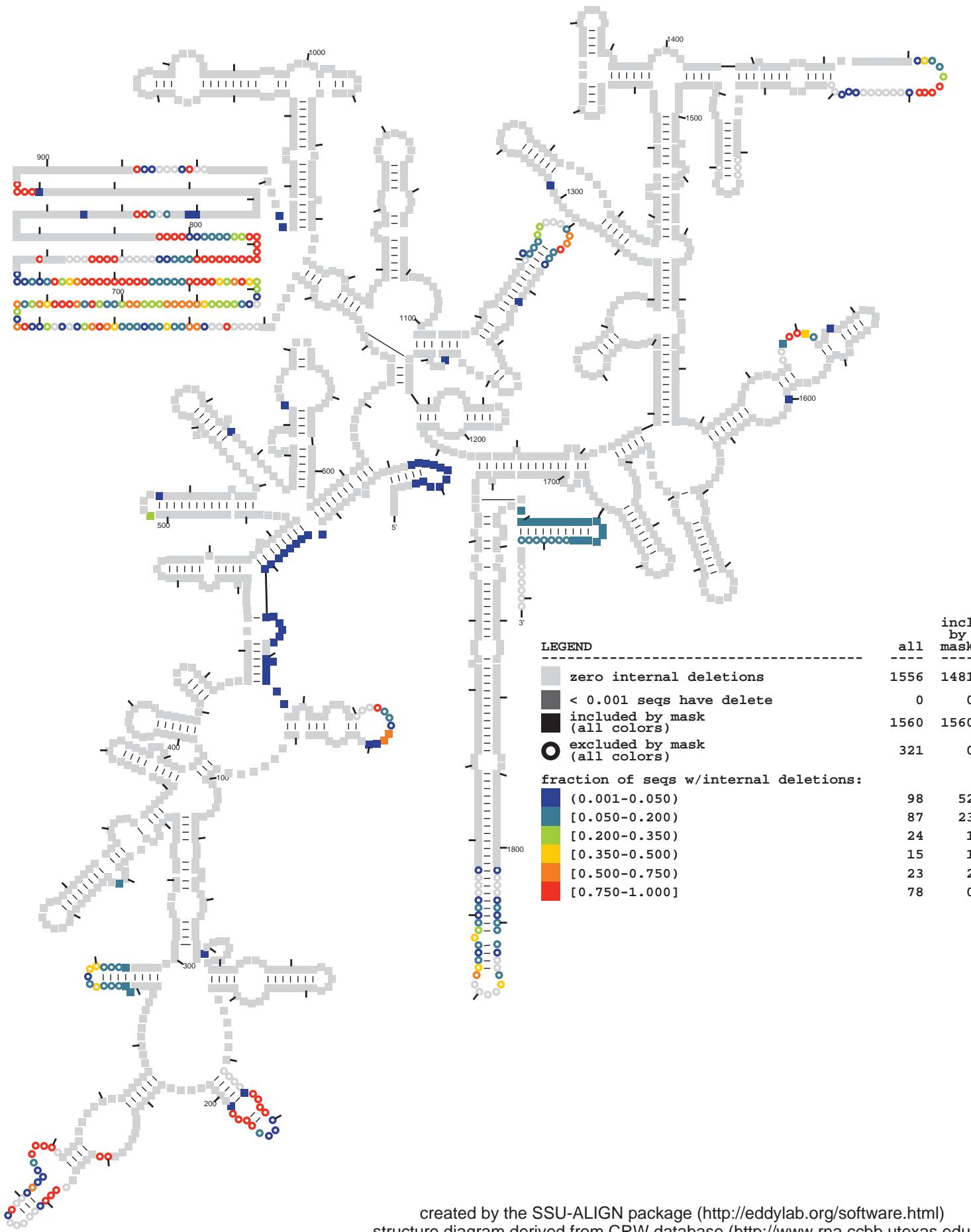


created by the SSU-ALIGN package (<http://eddylab.org/software.html>)
structure diagram derived from CRW database (<http://www.rna.ccbb.utexas.edu/>)



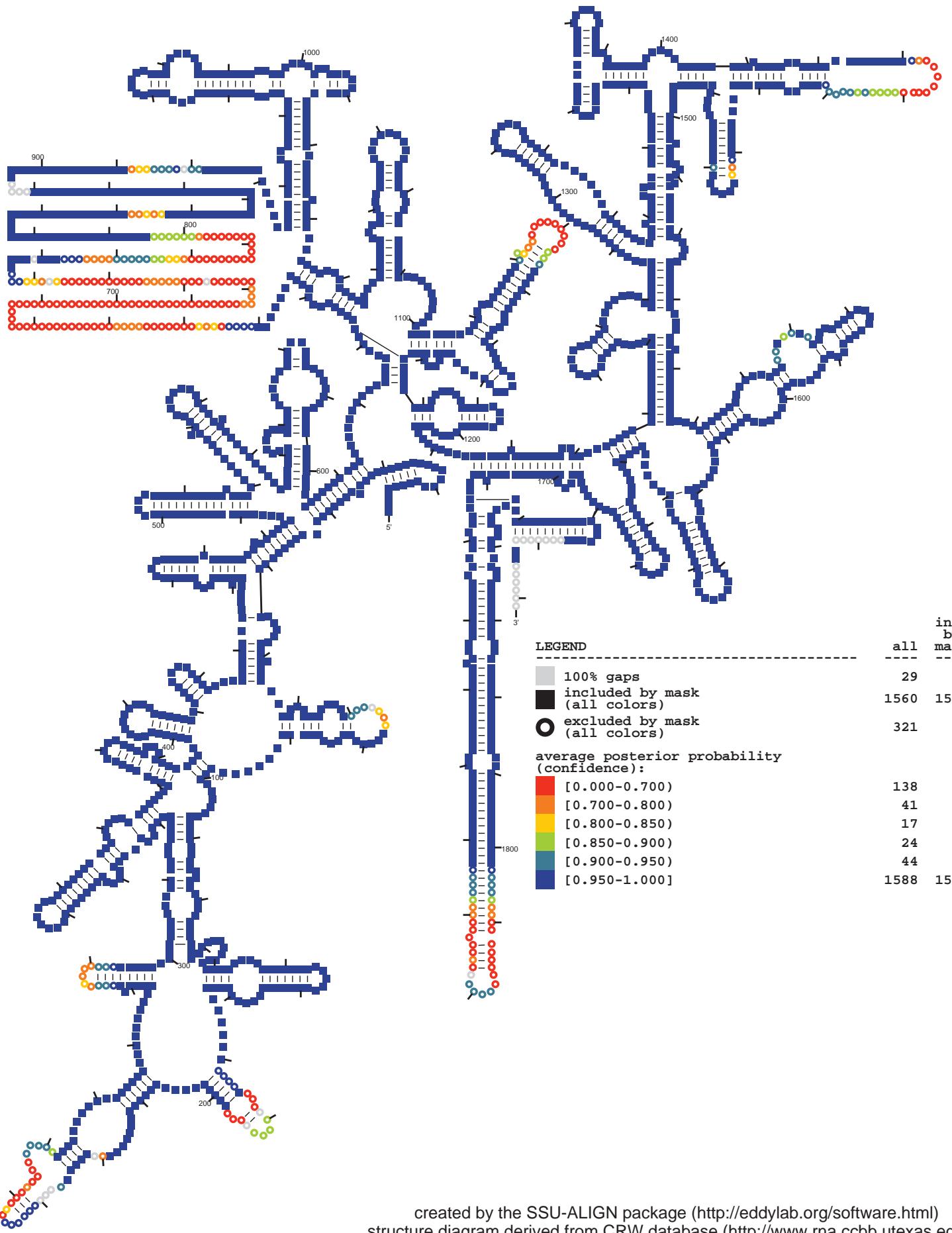
created by the SSU-ALIGN package (<http://eddylab.org/software.html>)
structure diagram derived from CRW database (<http://www.rna.ccbb.utexas.edu/>)

model	#pos	#bps	#seqs	description
eukarya	1881	448	94	frequency of internal deletions in each position

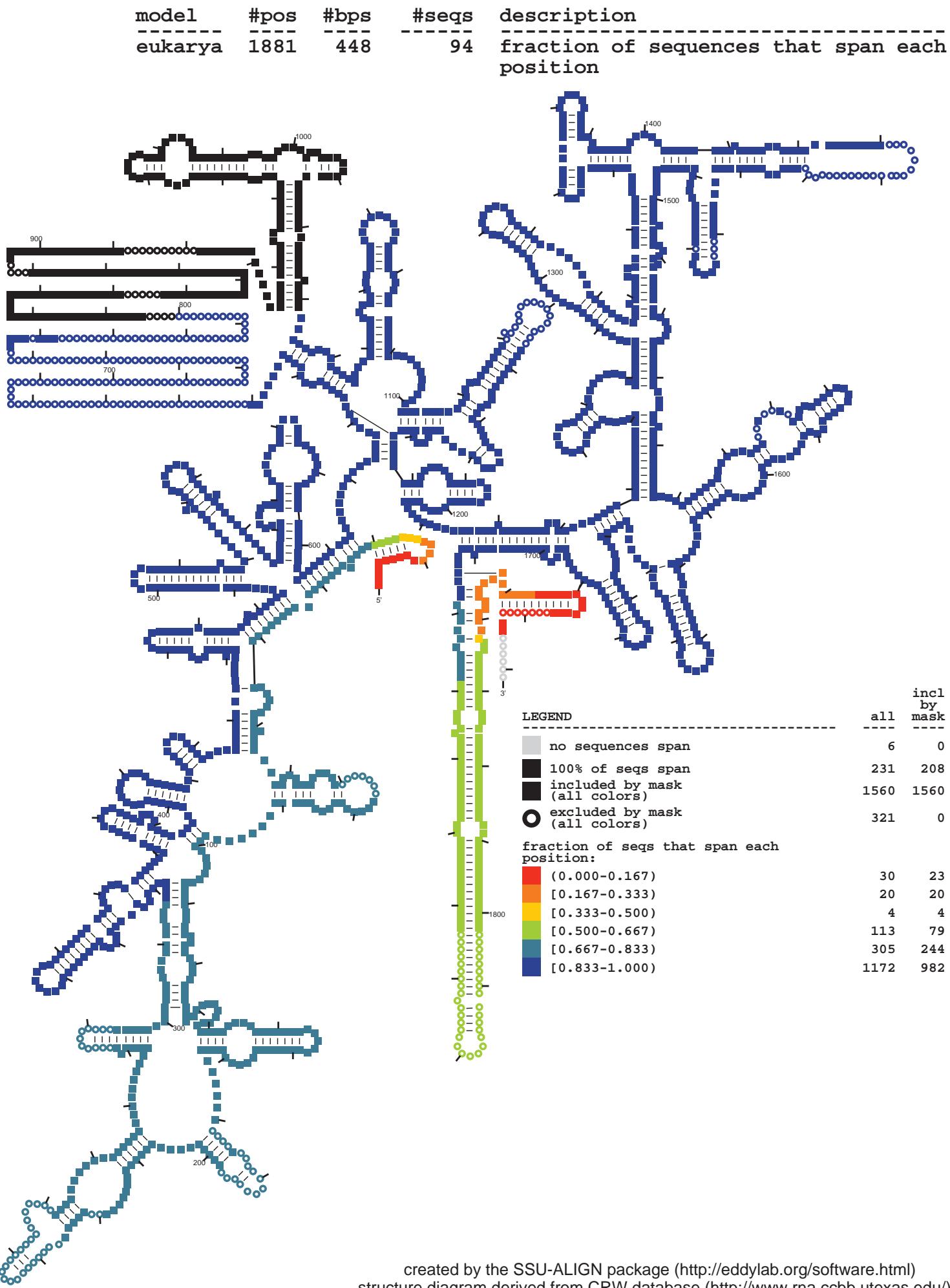


created by the SSU-ALIGN package (<http://eddylab.org/software.html>)
structure diagram derived from CRW database (<http://www.rna.ccbb.utexas.edu/>)

model	#pos	#bps	#seqs	description
eukarya	1881	448	94	average posterior probability per position



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structure diagram derived from CRW database (<http://www.rna.ccbb.utexas.edu/>)



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structure diagram derived from CRW database (<http://www.rna.ccbb.utexas.edu/>)