model	#res	#bps	#seqs
bacteria	1582	480	93

LEGEND

		all	in mask
	100% gaps	0	0
	included by mask (all colors)	-	1471
O	excluded by mask (all colors)	-	111

average posterior
probability (alnment confidence):

		all	in mask
	[0.000-0.800)	4	0
	[0.800-0.900)	16	0
	[0.900-0.925)	13	0
	[0.925-0.950)	12	0
	[0.950-0.975)	33	1
	[0.975-1.000]	1504	1470