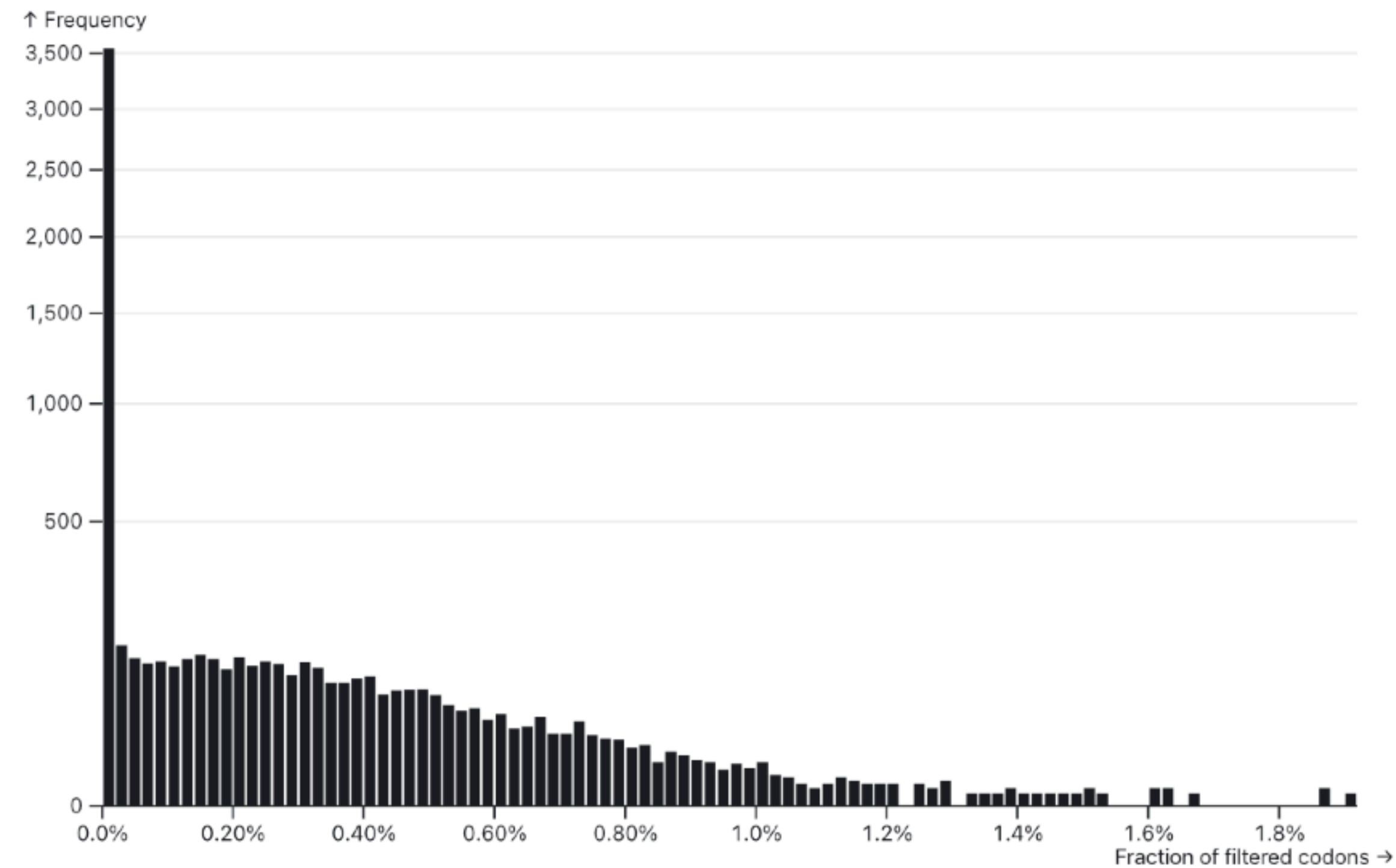
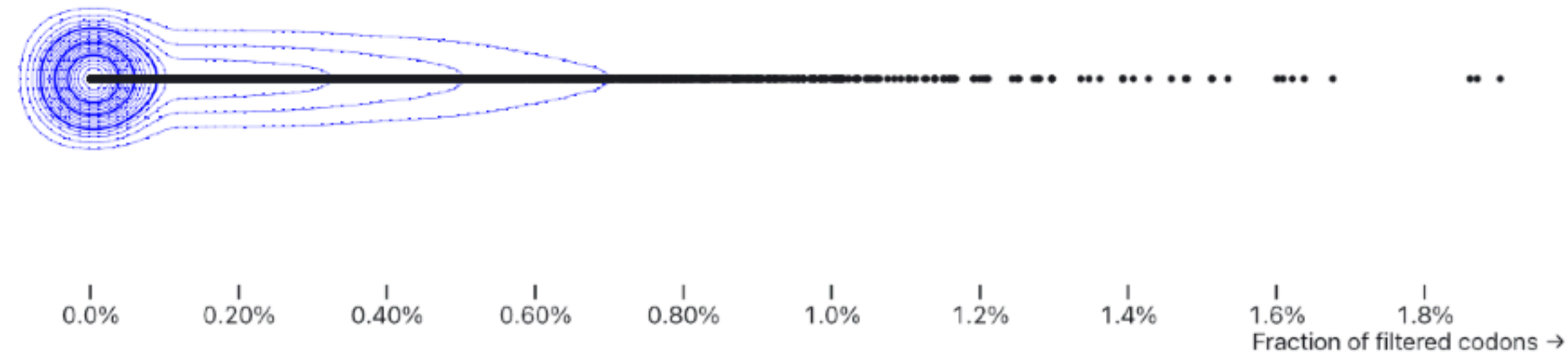


- The distribution of per MSA filtered codon counts

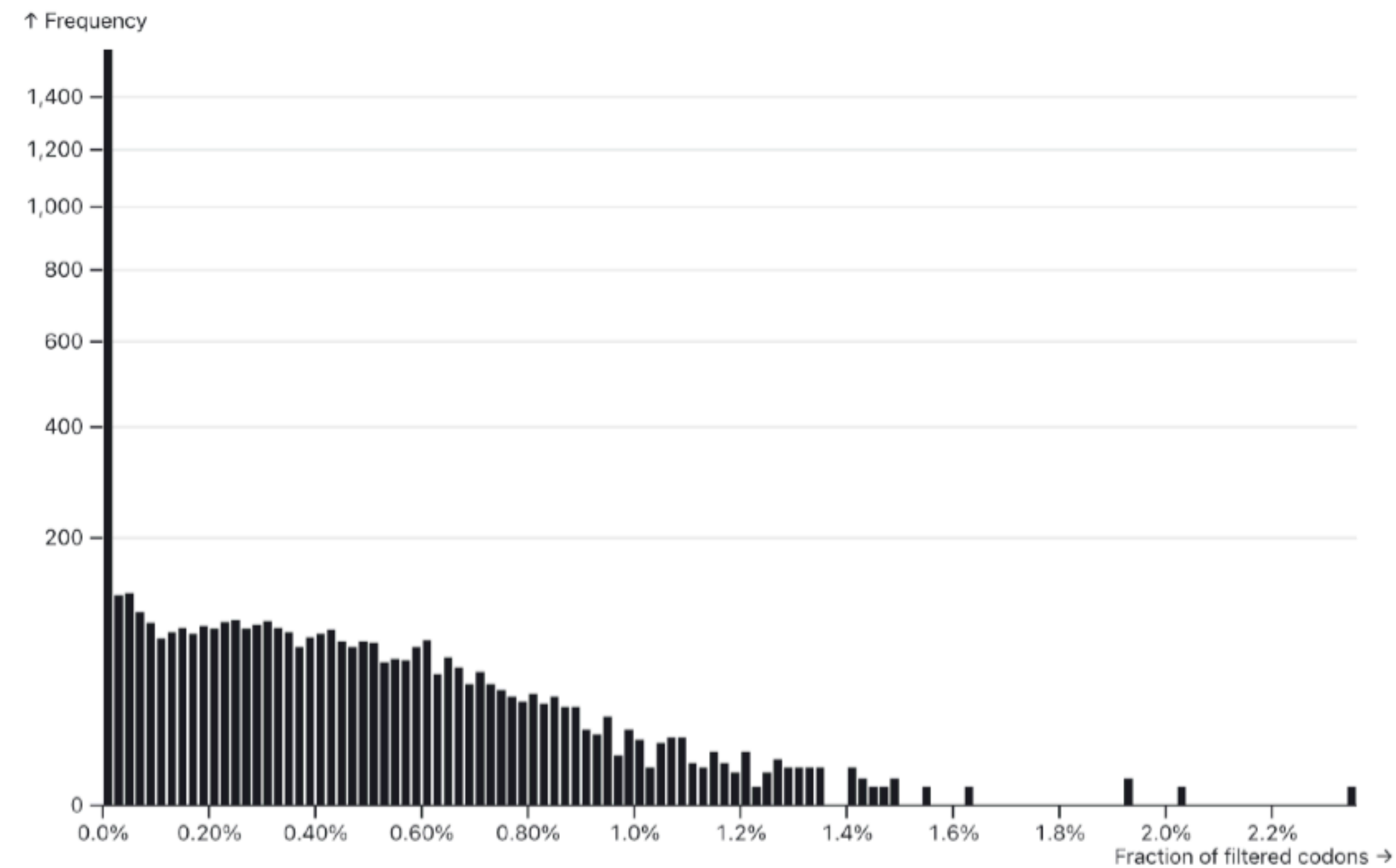
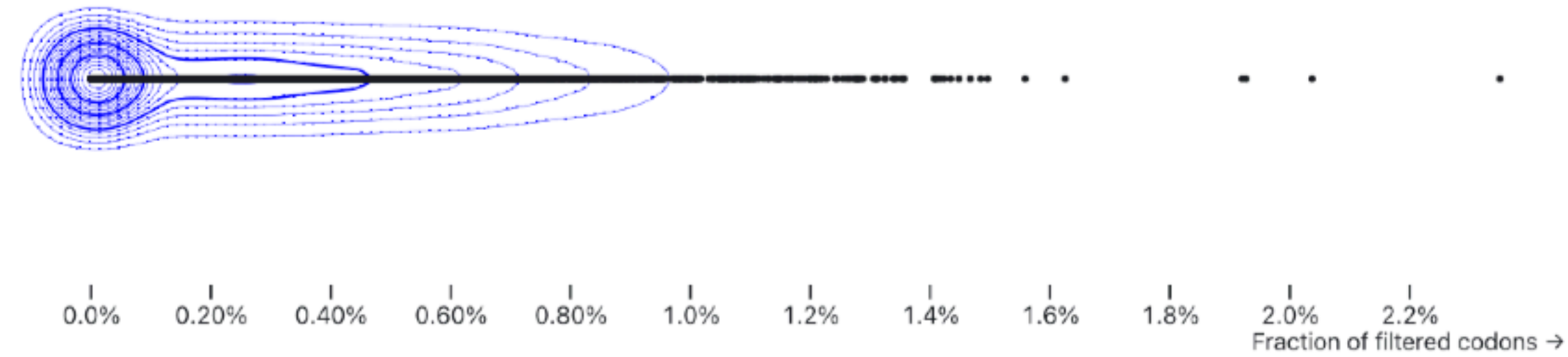
Fractions of filtered sites (of the entire alignment character count, seqs x codons)



32-taxon Zoonomia tree
Codons filtered / 1000 codons

- The distribution of per MSA filtered codon counts

Fractions of filtered sites (of the entire alignment character count, seqs x codons)



64-taxon Zoonomia tree
Codons filtered / 1000 codons