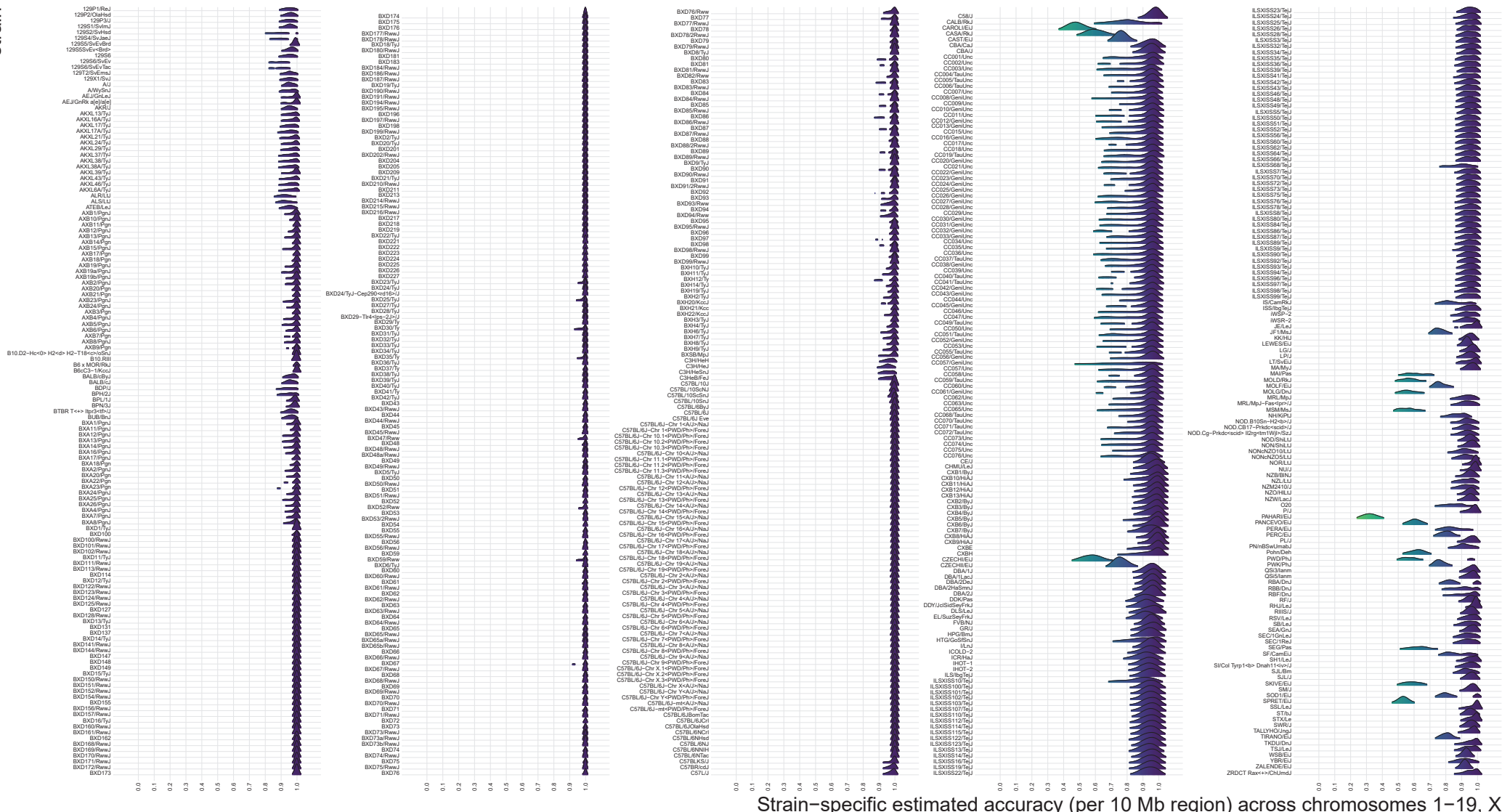


strain



Supplemental Fig. S1. Distributions of strain-specific estimated accuracy across chromosomes 1-19 and X. Strain-specific accuracy was estimated in each 10Mb region using a ‘held out’ test set that was not used to impute allelic states for the strain in the 10Mb region. Ridgeline plots illustrate the distribution of estimated accuracies.