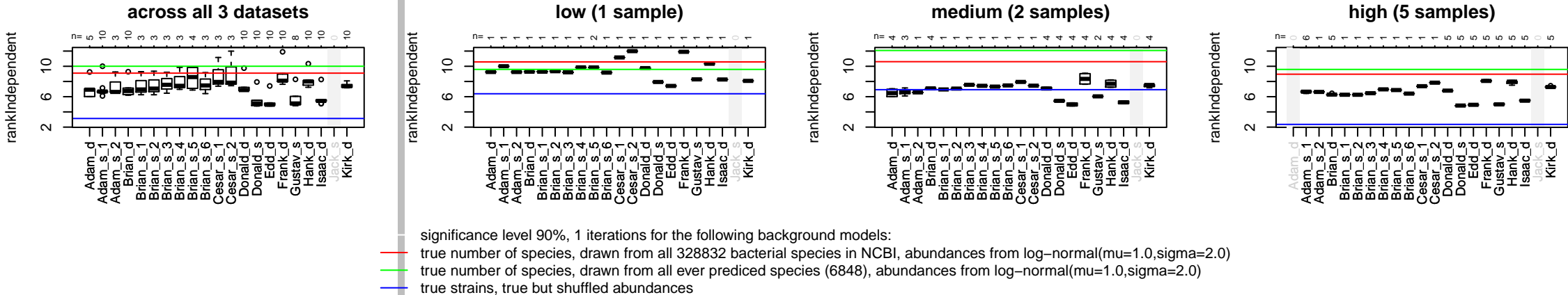
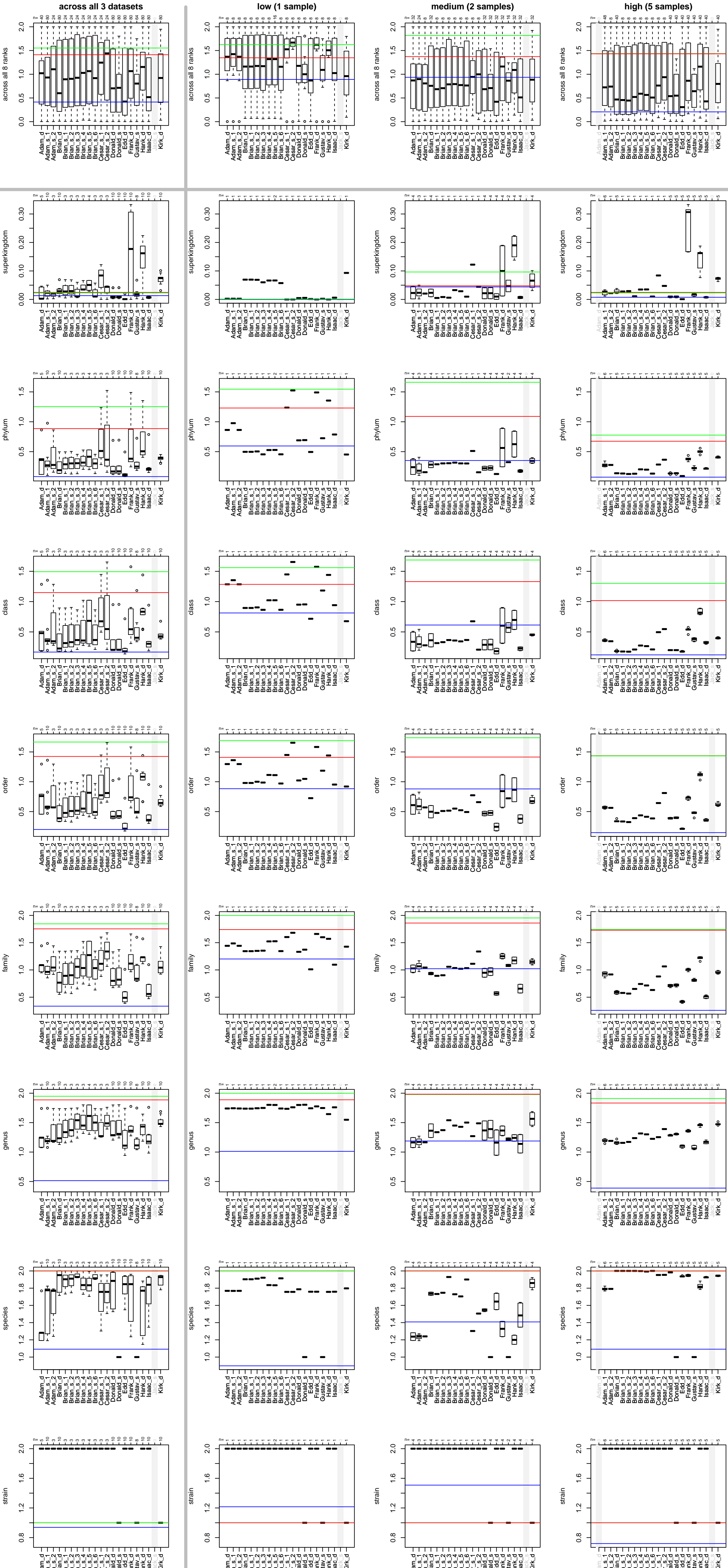


CAMI competition, metric: Unifrac



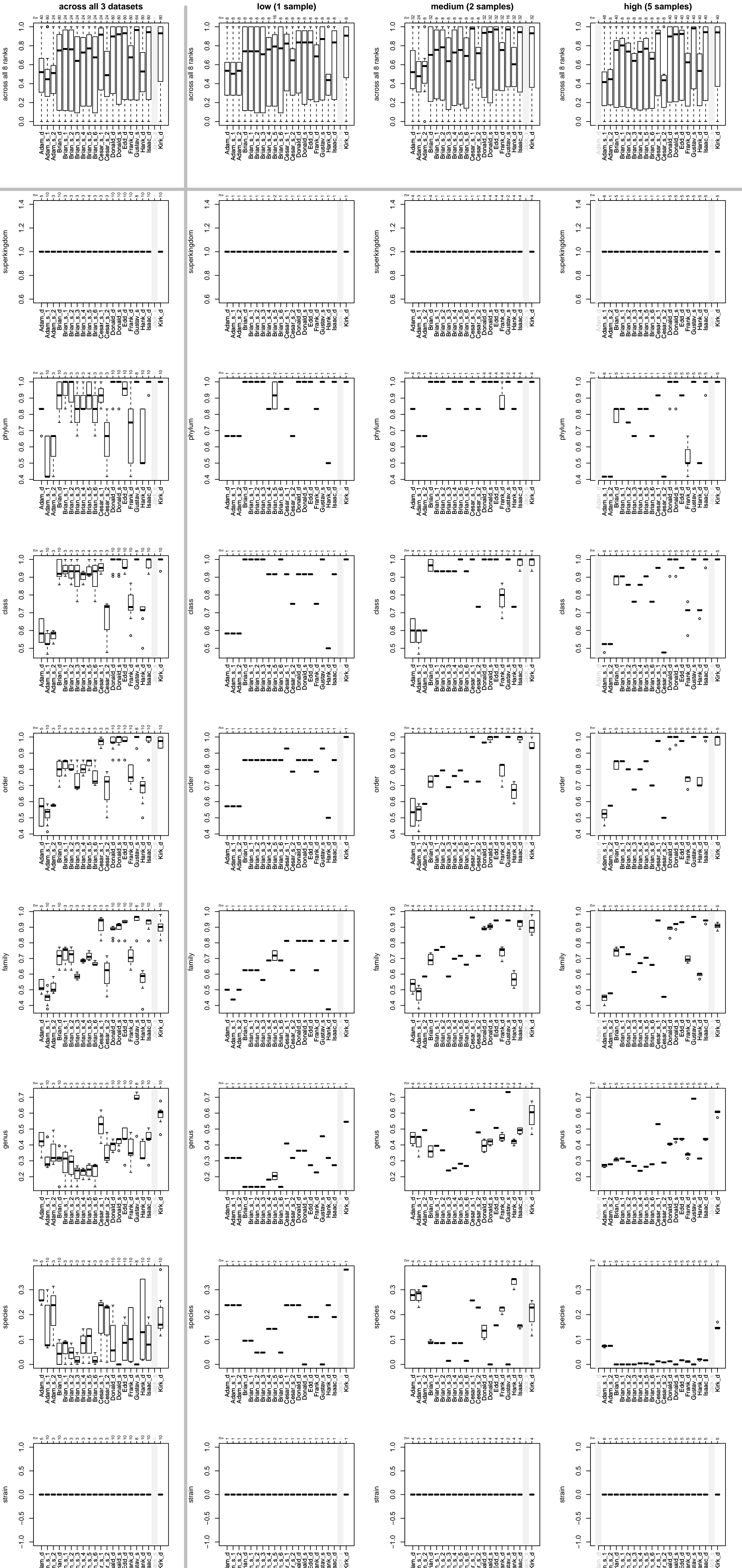
CAMI competition, metric: L1norm



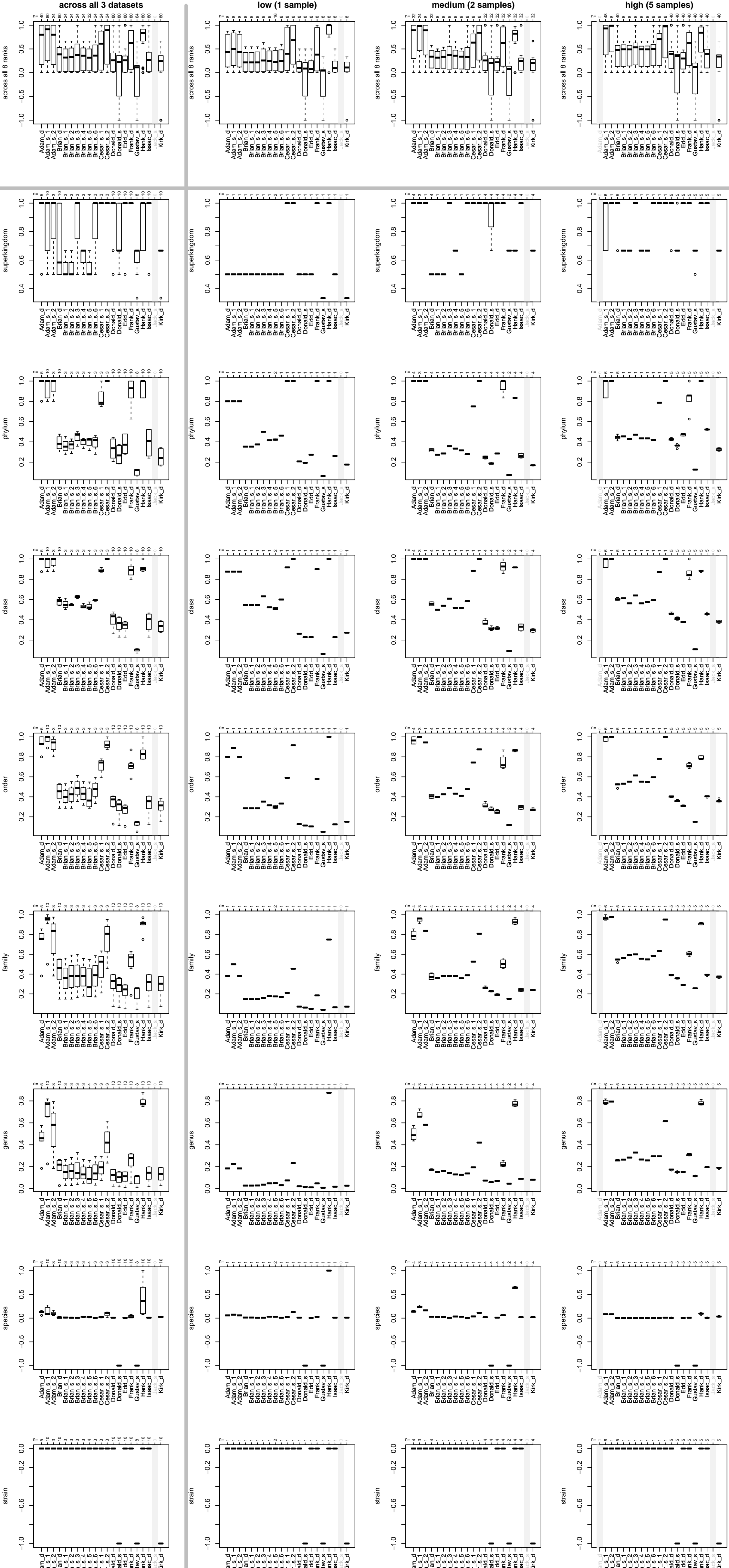
significance level 90%, 1 iterations for the following background models:

- grey true number of species, drawn from all 328832 bacterial species in NCBI, abundances from log-normal($\mu=1.0, \sigma=2.0$)
- red true number of species, drawn from all ever predicted species (6848), abundances from log-normal($\mu=1.0, \sigma=2.0$)
- green true strains, true but shuffled abundances
- blue true strains, true but shuffled abundances

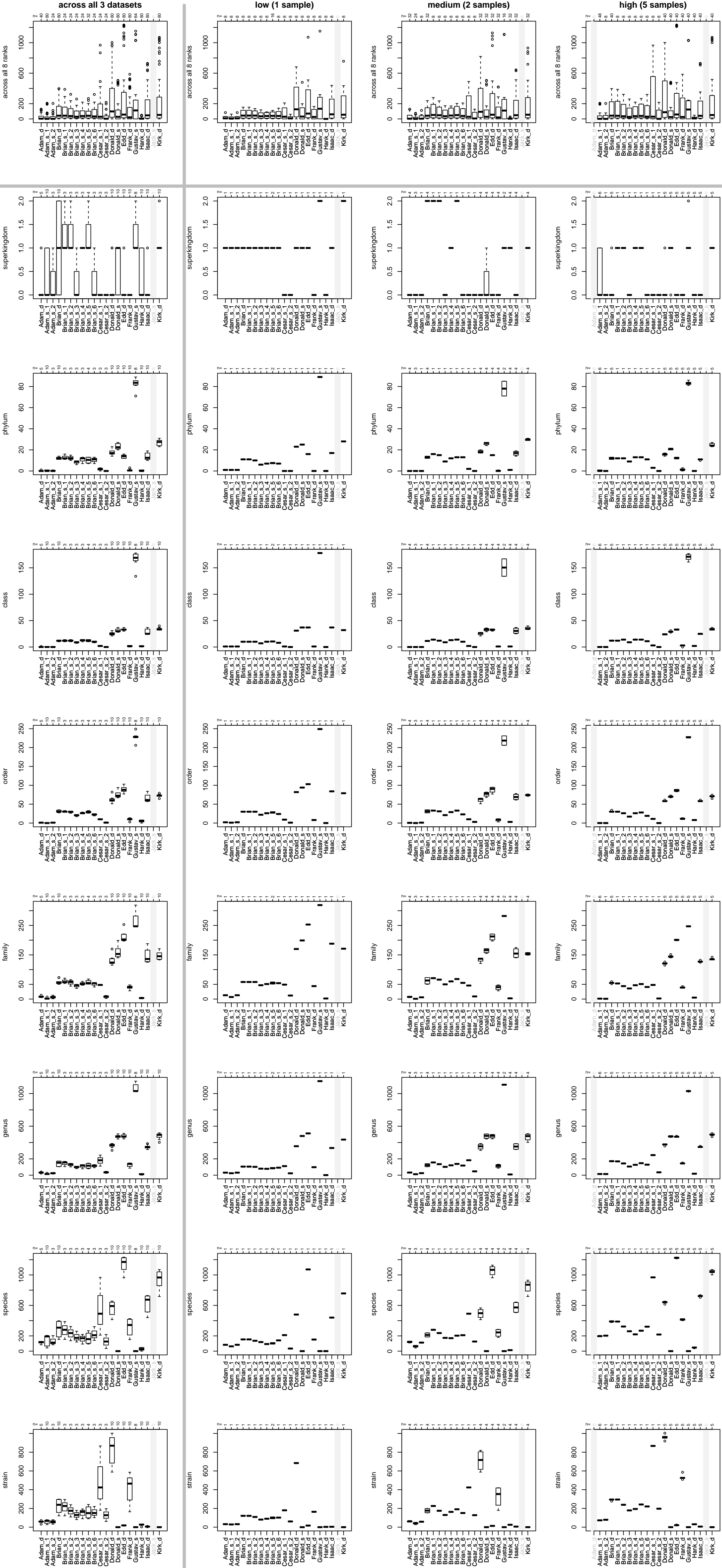
CAMI competition, metric: Sensitivity



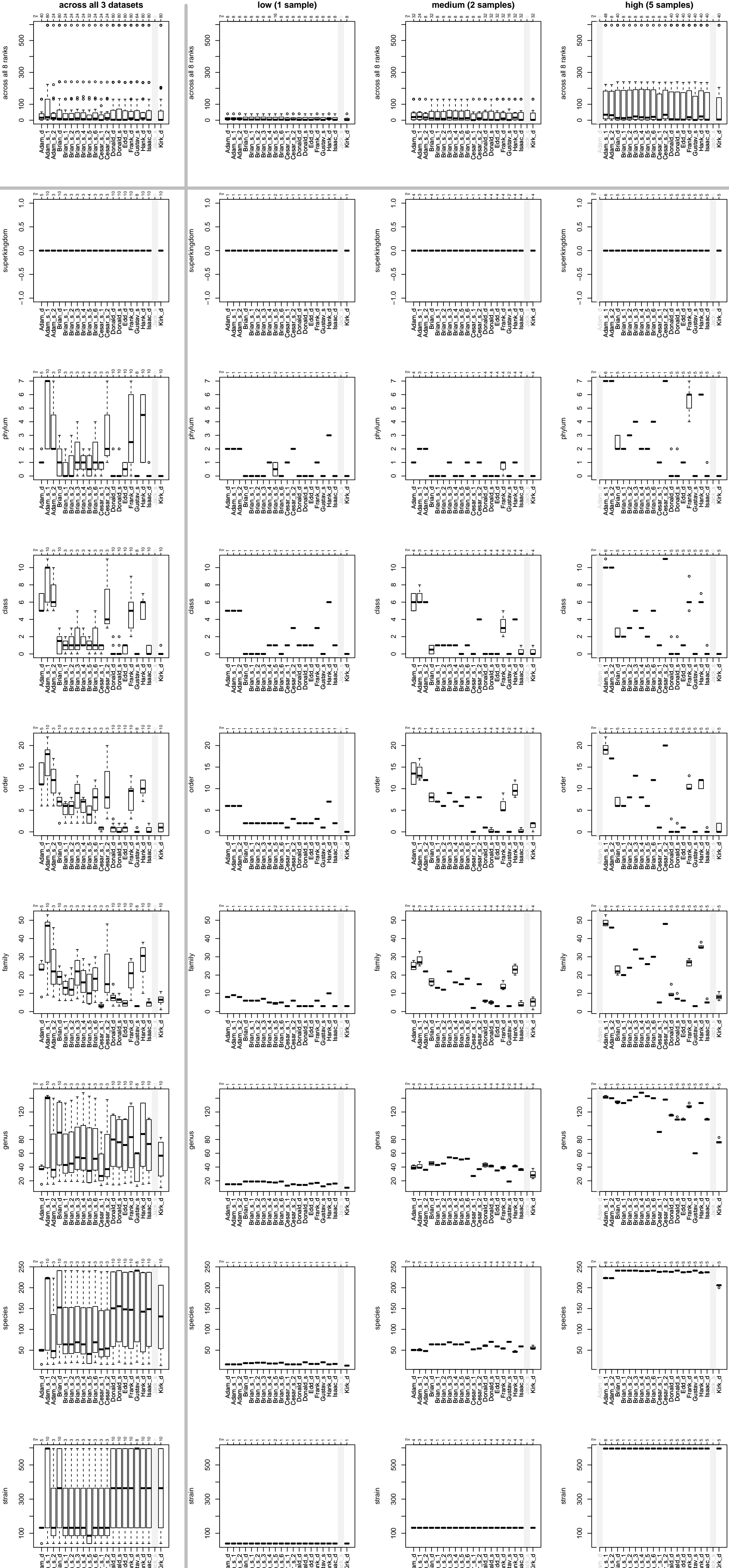
CAMI competition, metric: Precision



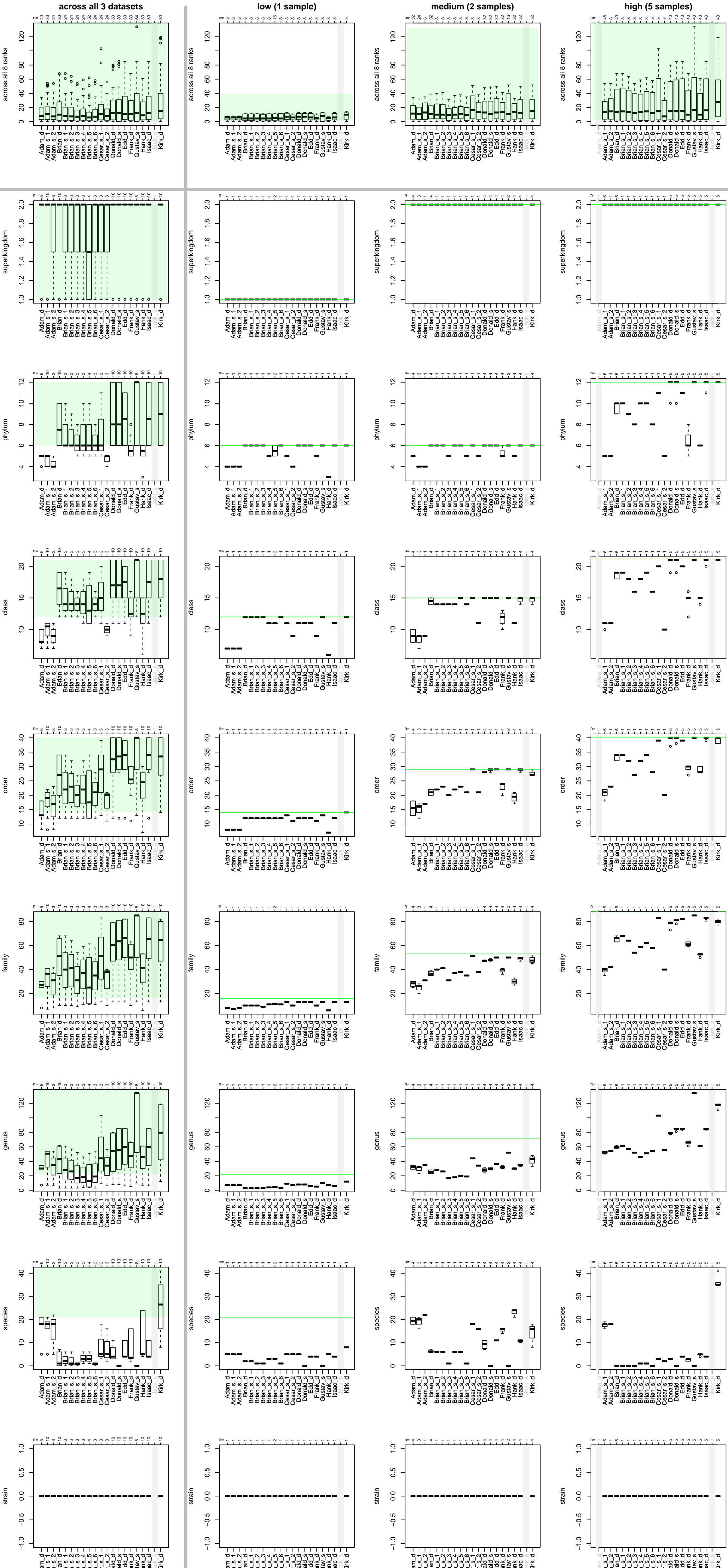
CAMI competition, metric: False Positives



CAMI competition, metric: False Negatives



CAMI competition, metric: True Positives



— number of taxonomic classes in gold standard profile(s)