Figure: Box plots comparing methods across datasets. Top row (accuracy): rmp, camp, cirrhosis, genomemed. Bottom row (negative MSE): centralparksoil, americangut, $\times 10^6$ hiv, tara.

Legend: Baseline, SVM-RBF, Lin/Log-L1, RF, LogCont-L1, KernelBiome, KB-Aitchison