CNS−GBM (paired, all sigs)

Legend (top to bottom): fullREDMSL, HMP, permutation, HMP2, ILR, perturbation, ttest, Harris_chi, diagREDM, fullREM, diagREDMSL

X-axis: Percentage of mixture — 0, 6.1e-06, 4.5e-05, 0.00034, 0.018, 0.12, 0.27, 0.5, 0.98, 1

Y-axis: Accuracy — 0.00, 0.25, 0.50, 0.75, 1.00