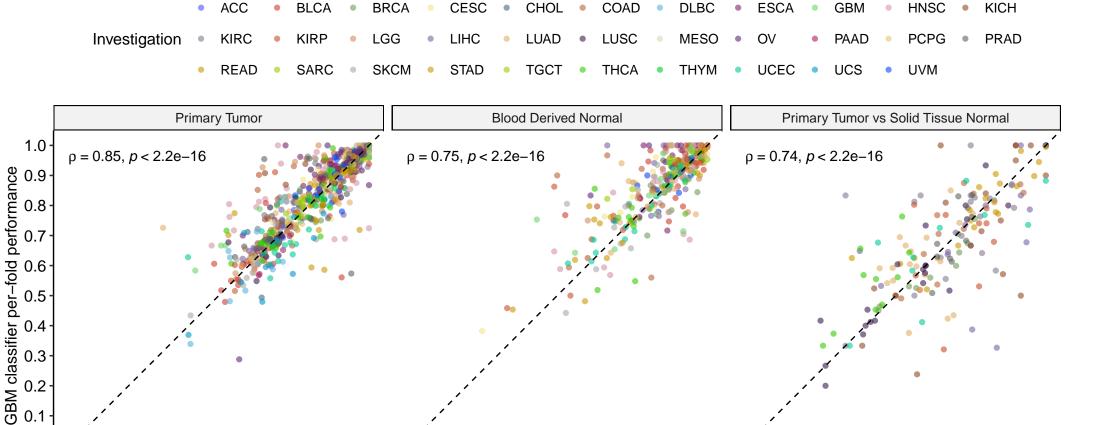
Matched GBM vs. RF Per–Fold CV Performance | AUROC | TCGA Raw Decontaminated Data (All Sequencing Centers)



0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

RF classifier per-fold performance

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

0.0

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0