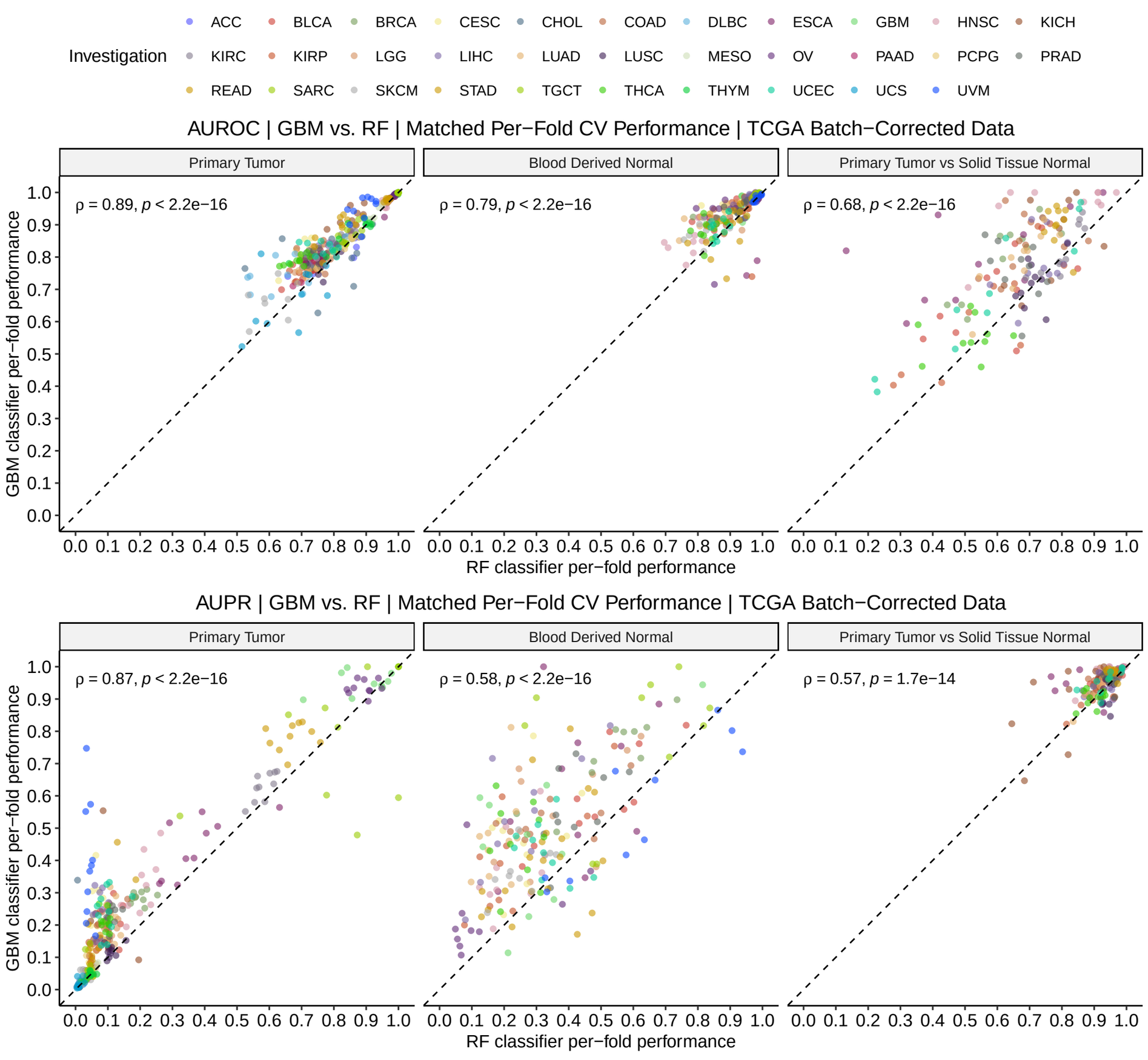
Investigation
ACC
BLCA
BRCA
CESC
CHOL
COAD
DLBC
ESCA
GBM
HNSC
KICH
KIRC
KIRP
LGG
LIHC
LUAD
LUSC
MESO
OV
PAAD
PCPG
PRAD
READ
SARC
SKCM
STAD
TGCT
THCA
THYM
UCEC
UCS
UVM
AUROC | GBM vs. RF | Matched Per−Fold CV Performance | TCGA Batch−Corrected Data
Primary Tumor
Blood Derived Normal
Primary Tumor vs Solid Tissue Normal
ρ = 0.89, p < 2.2e−16
ρ = 0.79, p < 2.2e−16
ρ = 0.68, p < 2.2e−16
GBM classifier per−fold performance
RF classifier per−fold performance
AUPR | GBM vs. RF | Matched Per−Fold CV Performance | TCGA Batch−Corrected Data
Primary Tumor
Blood Derived Normal
Primary Tumor vs Solid Tissue Normal
ρ = 0.87, p < 2.2e−16
ρ = 0.58, p < 2.2e−16
ρ = 0.57, p = 1.7e−14
GBM classifier per−fold performance
RF classifier per−fold performance