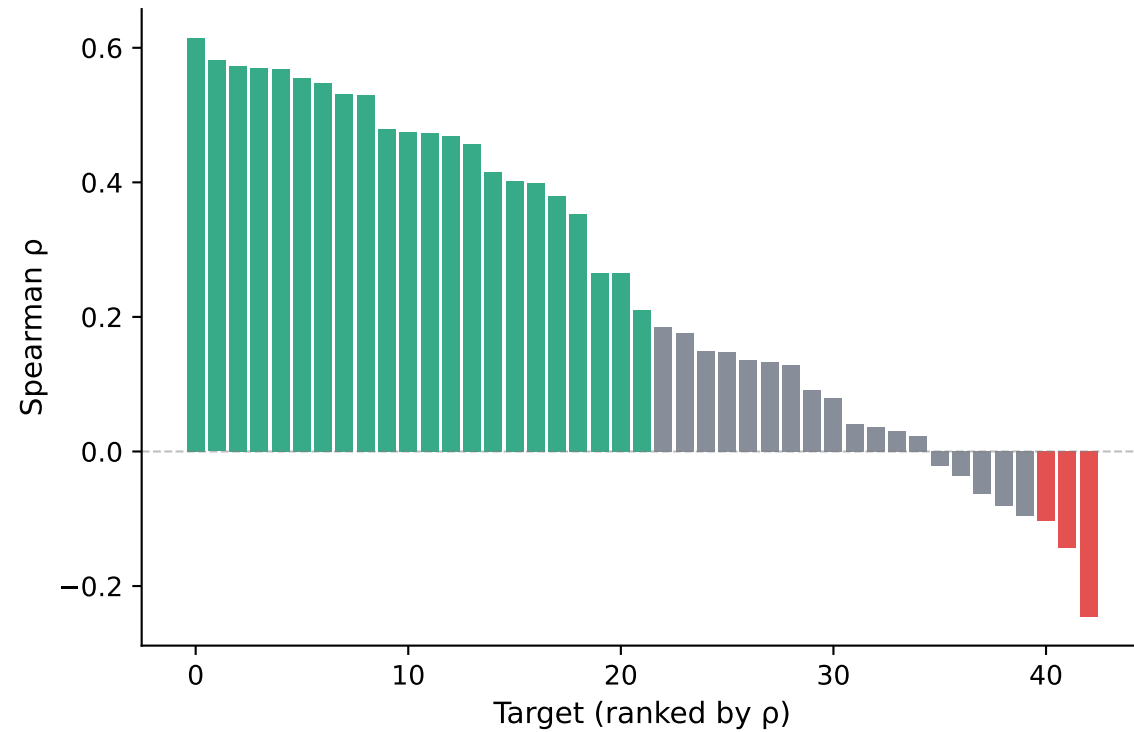
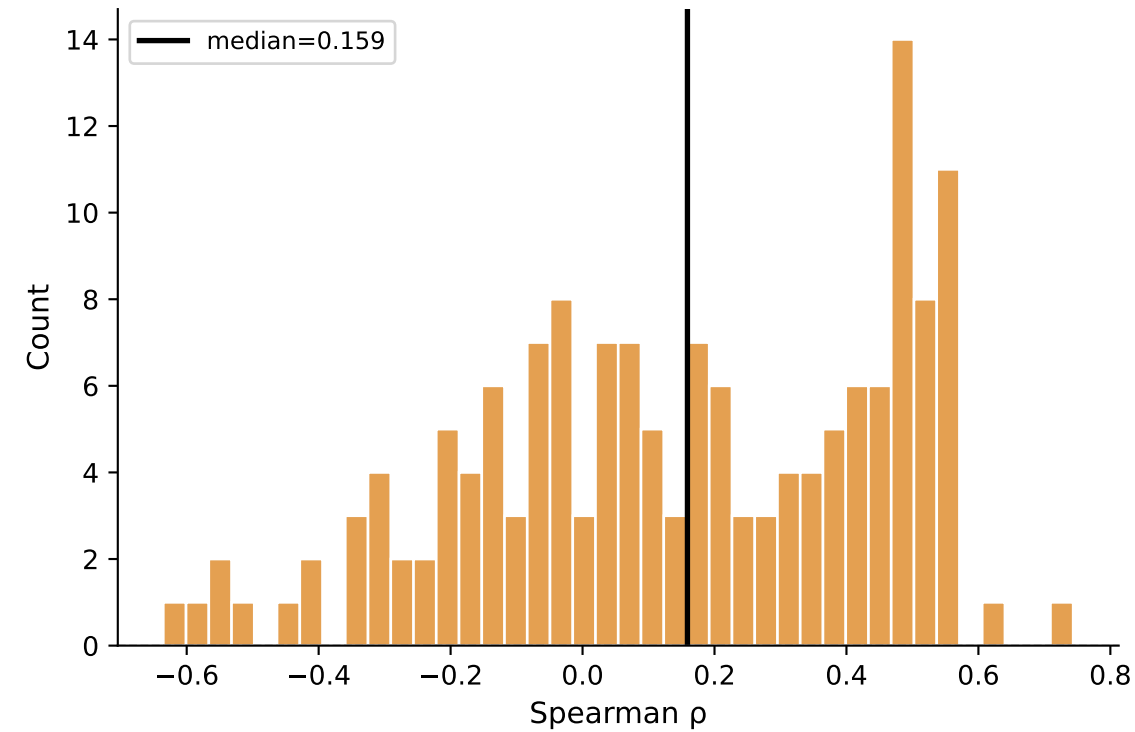


Bulk RNA-seq Validation: Donor-Level Spearman Correlation (CytoSig/LinCytoSig/SecAct predicted activity vs target gene expression)

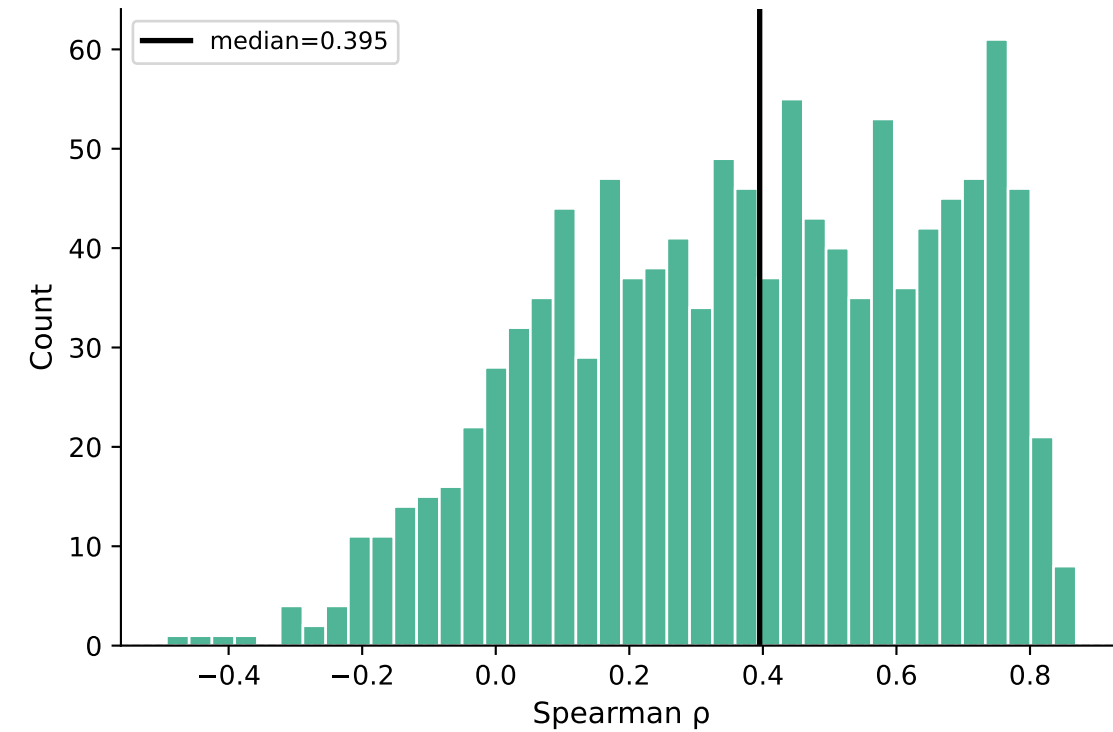
GTEx — CYTOSIG
(n=43, median $\rho=0.211$)



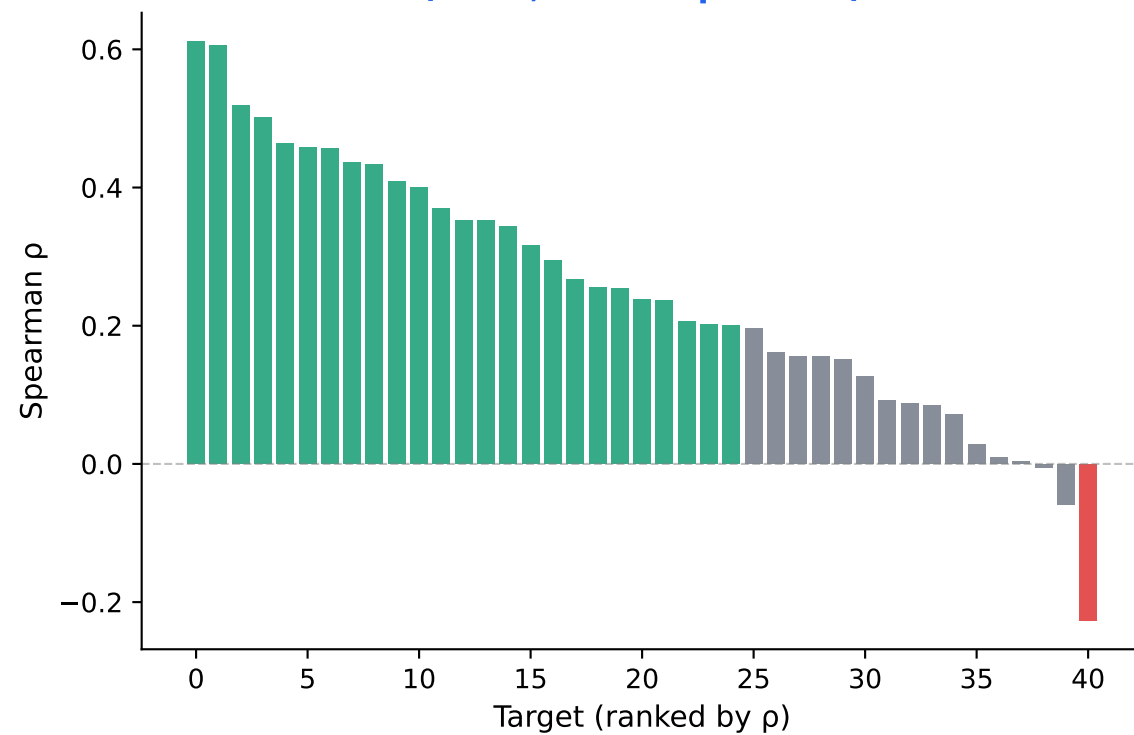
GTEx — LINCYTOSIG
(n=156, median $\rho=0.159$)



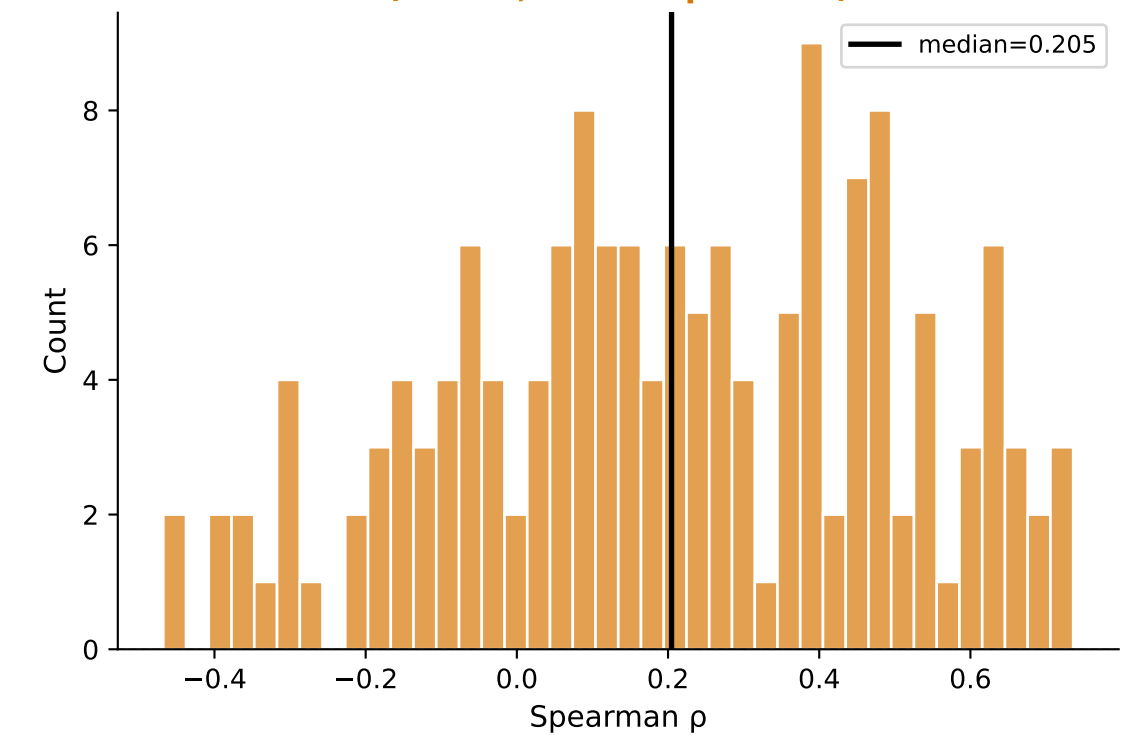
GTEx — SECACT
(n=1132, median $\rho=0.395$)



TCGA — CYTOSIG
(n=41, median $\rho=0.238$)



TCGA — LINCYTOSIG
(n=152, median $\rho=0.205$)



TCGA — SECACT
(n=1085, median $\rho=0.415$)

