GO MF Dot Plot: IC2_vs_Normal_wilcox.csv GTPase regulator activity phospholipid binding ubiquitin-like protein ligase binding structural constituent of ribosome p.adjust GTPase binding 2e-04 cadherin binding 4e-04 kinase regulator activity-6e-04 histone binding Count phosphatase binding ubiquitin-like protein binding rRNA binding 100 virus receptor activity-MHC protein complex binding peptide antigen binding-K63-linked polyubiquitin modification-dependent-0 protein binding 0.01 0.02 0.03 0.04 GeneRatio