GO MF Bar Plot: IC2\_vs\_Normal\_wilcox.csv structural constituent of ribosome ubiquitin-like protein ligase binding MHC protein complex binding GTPase binding GTPase regulator activity phosphatase binding p.adjust rRNA binding 2e-04 cadherin binding 4e-04 ubiquitin-like protein binding 6e-04 peptide antigen binding kinase regulator activity phospholipid binding virus receptor activity-K63-linked polyubiquitin modification-dependent protein binding histone binding 30 60 90 0 120 Count