GO MF Bar Plot: IIB\_vs\_Normal\_wilcox.csv structural constituent of ribosome GTPase regulator activity cadherin binding ubiquitin protein ligase binding kinase regulator activity actin binding p.adjust GTPase binding 1e-05 MHC class II protein complex 2e-05 binding 3e-05 4e-05 rRNA binding 5e-05 protein serine/threonine kinase activity translation factor activity, RNA binding peptide bindingtransmembrane transporter binding phospholipid binding SMAD binding 30 60 90 0 Count