GO MF Dot Plot: IIB_vs_Normal_wilcox.csv GTPase regulator activity actin binding phospholipid binding structural constituent of ribosome p.adjust cadherin binding 1e-05 protein serine/threonine 2e-05 kinase activity 3e-05 ubiquitin protein ligase 4e-05 binding 5e-05 GTPase binding Count 20 kinase regulator activitypeptide binding 60 transmembrane transporter 100 binding translation factor activity, RNA binding SMAD binding rRNA binding MHC class II protein complex 0 binding 0.01 0.02 0.03 0.04 0.05 GeneRatio