GO MF Dot Plot: IIIB_vs_Normal_wilcox.csv GTPase regulator activity actin binding DNA-binding transcription factor binding cadherin bindingstructural constituent of ribosome p.adjust GTPase binding-1e-05 ubiquitin protein ligase binding 2e-05 proton transmembrane 3e-05 transporter activity ribonucleoprotein complex binding Count electron transfer activity 100 unfolded protein bindingtranslation factor activity, RNA binding oxidoreduction-driven active transmembrane transporter activity NADH dehydrogenase (ubiquinone) activity endopeptidase activator activity 0.01 0.02 0.03 0.04 GeneRatio