GO MF Dot Plot: IC2_vs_Normal_wilcox.csv GTPase regulator activity actin binding **DNA**-binding transcription factor binding cadherin binding ubiquitin-like protein ligase binding p.adjust GTPase binding 2e-06 structural constituent of 4e-06 ribosome 6e-06 phosphatidylinositol binding transcription corepressor Count activity 50 phosphatase binding 100 150 integrin binding extracellular matrix structural constituent growth factor binding collagen binding MHC protein complex binding-0 0.01 0.02 0.03 0.04 0.05 GeneRatio