GO MF Dot Plot: IIIC\_vs\_Normal\_wilcox.csv DNA-binding transcription factor binding GTPase regulator activity catalytic activity, acting on RNA cadherin binding p.adjust ubiquitin-like protein ligase binding 2.50e-06 GTPase binding 5.00e-06 7.50e-06 transcription coactivator 1.00e-05 activity 1.25e-05 histone binding Count transcription corepressor 100 activity 150 structural constituent of 200 ribosome 250 ribonucleoprotein complex 300 binding helicase activity nuclear receptor binding transcription coregulator binding SMAD binding 0.01 0.02 0.03 GeneRatio