GO MF Bar Plot: IIIB_vs_Normal_wilcox.csv structural constituent of ribosome ubiquitin-like protein ligase binding cadherin binding DNA-binding transcription factor binding GTPase regulator activity transcription corepressor activity p.adjust GTPase binding 2e - 06nuclear receptor binding basal RNA polymerase II 4e-06 transcription machinery binding 6e-06 ribonucleoprotein complex binding histone binding actin binding translation factor activity, RNA binding histone modifying activity phosphatase binding-50 100 150 0 200 Count