GO MF Bar Plot: IC2_vs_Normal_wilcox.csv GTPase regulator activity cadherin binding-GTPase bindingubiquitin-like protein ligase binding DNA-binding transcription factor binding structural constituent of ribosome p.adjust protein serine/threonine kinase activity transcription coactivator 2e-06 activity 4e-06 histone modifying activity-6e-06 transcription corepressor activity actin bindingmagnesium ion bindingmodification-dependent protein binding histone H2A kinase activitycatalytic activity, acting on DNA 100 200 0 300 Count