GO Enrichment Analysis: Molecular function S-acyltransferase activity ubiquitin protein ligase _ binding protein phosphatase regulator activity GeneRatio phosphatase activator activity -0.01 0.02 0.03 ubiquitin-like protein ligase binding p.adjust 0.020 catalytic activity, acting on DNA 0.015 0.010 single-stranded DNA binding -0.005 ATP-dependent activity, acting on DNA microtubule motor activity -ATP hydrolysis activity down up (2327)(2102)Cluster