Patient AM 16 Metabolic Function ubiquitin-like protein ligase binding ubiquitin protein ligase binding GTP binding p.adjust 5.0e-07 phosphatidylinositol binding -1.0e-06 Count structural constituent of 40 ribosome 80 120 160 signaling adaptor activity double-stranded RNA binding single-stranded DNA helicase \bigcirc activity 0.01 0.03 0.02 GeneRatio