Patient AM 08 Metabolic Function actin binding guanyl nucleotide binding guanyl ribonucleotide binding GTP binding GTPase activity extracellular matrix structural constituent actin filament binding p.adjust primary active transmembrane transporter activity 0.005 0.010 growth factor binding-0.015 protein-folding chaperone 0.020 binding SMAD binding Count oxidoreduction-driven active 20 transmembrane transporter activity extracellular matrix structural constituent conferring tensile strength oxidoreductase activity, acting on NAD(P)H, quinone or similar compound as acceptor GDP binding NAD(P)H dehydrogenase (quinone) activity transmembrane receptor protein serine/threonine kinase activity calcium channel inhibitor activity transforming growth factor 0 beta receptor activity 0 titin binding 0.01 0.02 0.04 0.03 GeneRatio