Patient AM 06 Metabolic Function structural constituent of ribosome cadherin binding catalytic activity, acting on RNA ribonucleoprotein complex binding ligase activity · unfolded protein binding ribosome binding p.adjust catalytic activity, acting 0.003 on a tRNA 0.006 electron transfer activity · 0.009 growth factor binding 0.012 translation regulator activity, nucleic acid binding Count oxidoreduction-driven active transmembrane transporter activity 40 rRNA binding oxidoreductase activity, acting on NAD(P)H, quinone or similar compound as acceptor translation initiation factor activity disulfide oxidoreductase activity mRNA 5'-UTR binding voltage-gated monoatomic anion channel activity oxidoreductase activity, acting on metal ions ubiquitin-protein transferase inhibitor activity 0.02 0.03 0.01 GeneRatio