Patient CMP 03 Metabolic Function protein serine/threonine kinase activity extracellular matrix structural constituent sulfur compound binding primary active transmembrane transporter activity growth factor binding integrin binding extracellular matrix structural constituent conferring tensile strength p.adjust cytokine receptor activity -0.01 oxidoreduction-driven active transmembrane transporter 0.02 activity 0.03 flavin adenine dinucleotide binding 0.04 oxidoreductase activity, acting on the aldehyde or-Count oxo group of donors phosphatidylinositol-3-phosphate binding 20 proteoglycan binding phosphatidylinositol 3-kinase binding platelet-derived growth factor binding aldehyde dehydrogenase (NAD+) activity aldehyde dehydrogenase [NAD(P)+] activity extracellular matrix constituent conferring 0 elasticity RAGE receptor binding structural molecule activity 0 conferring elasticity

0.02

GeneRatio

0.01

0.03