## Patient AM 07 Metabolic Function actin binding · ubiquitin-like protein ligase binding cadherin binding ubiquitin protein ligase binding GTP binding -DNA-binding transcription repressor activity p.adjust extracellular matrix structural constituent **DNA**-binding transcription repressor activity, RNA 2e-04 polymerase II-specific ribonucleoprotein complex 4e-04 binding structural constituent of 6e-04 ribosome Count growth factor binding -20 isomerase activity -60 integrin binding ribosome binding · 100 transcription coregulator binding SMAD binding extracellular matrix binding oxidoreductase activity, acting on a sulfur group of donors extracellular matrix structural constituent conferring tensile strength disulfide oxidoreductase activity 0.02 0.03 0.01 GeneRatio