

term	p-value	q-value
Neutrophil Degranulation	0,000116	0,012716
Innate Immune System	0,000443	0,017599
Post-translational Modification Synthesis of GPI-anchored Proteins	0,00048	0,017599
Metabolism of Amino Acids and Derivatives	0,002741	0,075389
Immune System	0,006621	0,129227
Alternative Complement Activation	0,008224	0,129227
Sodium-coupled Phosphate Cotransporters	0,008224	0,129227
Phenylalanine Metabolism	0,00986	0,13558
Thyroxine Biosynthesis	0,016382	0,169963
Diseases Associated With Surfactant Metabolism	0,016382	0,169963
Phenylalanine and Tyrosine Metabolism	0,018005	0,169963
Lysine Catabolism	0,019626	0,169963
Glutamate and Glutamine Metabolism	0,022861	0,169963
RHOA GTPase Cycle	0,025031	0,169963
Formation of the Nephric Duct	0,029299	0,169963
Metabolism of Amine-Derived Hormones	0,029299	0,169963
Initiation of Nuclear Envelope (NE) Reformation	0,030902	0,169963
Competing Endogenous RNAs (ceRNAs) Regulate PTEN Translation	0,030902	0,169963
Defective GALNT12 Causes CRCS1	0,030902	0,169963
Defective GALNT3 Causes HFTC	0,030902	0,169963
Defective C1GALT1C1 Causes TNPS	0,032503	0,170253
Insertion of Tail-Anchored Proteins Into the Endoplasmic Reticulum Membrane	0,035696	0,170721
Common Pathway of Fibrin Clot Formation	0,035696	0,170721
RHO GTPase Cycle	0,037495	0,171851
Termination of O-glycan Biosynthesis	0,042052	0,182779
Cholesterol Biosynthesis	0,043635	0,182779
Regulation of PTEN mRNA Translation	0,045215	0,182779
Surfactant Metabolism	0,048368	0,182779
Dectin-2 Family	0,048368	0,182779