proton motive force—driven ATP synthesis regulation of protein localization regulation of catalytic activity lipid metabolic process—ubiquitin—dependent protein catabolic process—regulation of protein stability—into some biogenesis—mitochondrion organization—vesicle—mediated transport—positive regulation of transcription by RNA polymerase II—organic acid metabolic process—intracellular calcium ion homeostasis—chondroitin sulfate biosynthetic process—attivacid metabolic process—fatty acid metabolic process—fatty acid metabolic process—fatty acid metabolic process—fatty acid metabolic process—anatomical structure morphogenesis—negative regulation of transcription by RNA polymerase II—anatomical structure development—anatomical structure de morphogenesis of an epithelium cell division response to salt response to heat mRNA export from nucleus mitochondrial respiratory chain complex I assembly response to starvation regulation of muscle contraction regulation of muscle contraction regulation of chemotaxis metabolic process actin cytoskeleton organization regulation of multicellular organismal process Arp2/3 complex—mediated actin nucleation mRNA processing engulfment of apoptotic cell central nervous system development acyl—CoA metabolic process regulation of signal transduction regulation of cellular component organization peptidyl—serine phosphorylation modulation of chemical synaptic transmission animal organ morphogenesis regulation of cell migration potassium ion transmembrane transport positive regulation of cell population proliferation cell activation translation regulation of oviposition muscle organ development cell differentiation cell activation translation regulation of oviposition muscle organ development chemical synaptic transmission tricarboxylic acid cycle RNA methylation regulation of cell adhesion proton transmembrane transport proteolysis establishment of mitotic spindle orientation defense response to bacterium B cell differentiation rRNA processing response to stimulus Golgi organization DNA topological change response to stimulus Golgi organization DNA topological change response to wounding regulation of translation neuron remodeling intracellular transport glycolytic process epigenetic regulation of gene expression cellular response to light stimulus protein phosphorylation protein glycosylation protein glycosylation protein glycosylation protein folding positive regulation of axon regeneration pigment accumulation protein colactor biosynthetic process epithelial cell migration dopamine biosynthetic process lamin depolymerization defense response to fungus chloride transmembrane transport blood coagulation tetrahydrofolate biosynthetic process regulation of stem cell population maintenance regulation of cellular process regulation of stem cell population maintenance regulation of cellular process regulation of stem cell population maintenance regulation of cellular process tetrahydrofolate biosynthetic process regulation of stem cell population maintenance regulation of cellular process.

L—ascorbic acid transmembrane transport acid transmembrane transport acid transmembrane transport. G protein—coupled receptor signaling pathway cytochrome c—hemie linkage xenobiotic metabolic process transcription initiation at RNA polymerase II promoter stem cell fate determination. RNA polymerase II promoter del fate determination protein polyubiquitination pidment metabolic process muclear pore organization—multicellular organismal process methylation—sinsulin processing—histone H3—k4 demethylation—collagen fibril organization—collagen fibril organization—amyloid precursor protein catabolic process—tRNA threonylcarbamoyladenosine metabolic process—regulation of amine metabolic process—regulation of amine metabolic process—protein poly—ADP—ribosylation—protein deglutathionylation—protein degluta 0 20 40 60