GO:0000226: microtubule cytoskeleton organization GO:0006259: DNA metabolic process GO:0032774: RNA biosynthetic process GO:0080135: regulation of cellular response to stress GO:0009314: response to radiation GO:1990778: protein localization to cell periphery GO:0098609: cell-cell adhesion GO:0030705: cytoskeleton-dependent intracellular transport GO:1901987: regulation of cell cycle phase transition GO:0010562: positive regulation of phosphorus metabolic process GO:0034330: cell junction organization GO:1902414: protein localization to cell junction GO:0006790: sulfur compound metabolic process GO:1901615: organic hydroxy compound metabolic process GO:0032648: regulation of interferon-beta production GO:0006298: mismatch repair GO:0010810: regulation of cell-substrate adhesion GO:0044089: positive regulation of cellular component biogenesis GO:1905605: positive regulation of blood-brain barrier permeability GO:0097435: supramolecular fiber organization GO:0009611: response to wounding GO:0031175: neuron projection development GO:0032956: regulation of actin cytoskeleton organization GO:0007163: establishment or maintenance of cell polarity GO:1901699: cellular response to nitrogen compound GO:1903530: regulation of secretion by cell GO:0006029: proteoglycan metabolic process GO:0031344: regulation of cell projection organization GO:1902074: response to salt GO:0046777: protein autophosphorylation GO:0035860: glial cell-derived neurotrophic factor receptor signaling pathway GO:0048511: rhythmic process GO:0043087: regulation of GTPase activity GO:0009612: response to mechanical stimulus GO:0002700: regulation of production of molecular mediator of immune response GO:0003018: vascular process in circulatory system GO:1902692: regulation of neuroblast proliferation GO:0016236: macroautophagy GO:0072001: renal system development GO:0048871: multicellular organismal-level homeostasis GO:0051896: regulation of phosphatidylinositol 3-kinase/protein kinase B signal transduction GO:0006260: DNA replication GO:0046545: development of primary female sexual characteristics GO:0033365: protein localization to organelle GO:0043410: positive regulation of MAPK cascade GO:0032965: regulation of collagen biosynthetic process GO:0045216: cell-cell junction organization GO:0007169: transmembrane receptor protein tyrosine kinase signaling pathway GO:0045445: myoblast differentiation GO:0001666: response to hypoxia GO:0048255: mRNA stabilization GO:0042063: gliogenesis GO:0090100: positive regulation of transmembrane receptor protein serine/threonine kinase signaling pathway GO:0006368: transcription elongation by RNA polymerase II GO:0051056: regulation of small GTPase mediated signal transduction GO:0042045: epithelial fluid transport GO:0046530: photoreceptor cell differentiation GO:0006913: nucleocytoplasmic transport GO:0030111: regulation of Wnt signaling pathway GO:0010639: negative regulation of organelle organization GO:0010457: centriole-centriole cohesion GO:0043604: amide biosynthetic process GO:0042391: regulation of membrane potential GO:0006641: triglyceride metabolic process GO:0001667: ameboidal-type cell migration GO:0045785: positive regulation of cell adhesion GO:0031396: regulation of protein ubiquitination GO:1900180: regulation of protein localization to nucleus GO:0017156: calcium-ion regulated exocytosis GO:0009066: aspartate family amino acid metabolic process GO:0040007: growth GO:0051091: positive regulation of DNA-binding transcription factor activity GO:1902017: regulation of cilium assembly GO:0048638: regulation of developmental growth GO:0007158: neuron cell-cell adhesion GO:1900034: regulation of cellular response to heat GO:0045787: positive regulation of cell cycle GO:0006338: chromatin remodeling GO:0034284: response to monosaccharide GO:0006595: polyamine metabolic process GO:0032784: regulation of DNA-templated transcription elongation GO:0007632: visual behavior GO:0043122: regulation of canonical NF-kappaB signal transduction GO:0031929: TOR signaling GO:0051968: positive regulation of synaptic transmission, glutamatergic GO:0002693: positive regulation of cellular extravasation GO:0043086: negative regulation of catalytic activity GO:0038061: non-canonical NF-kappaB signal transduction GO:0060348: bone development GO:0061024: membrane organization GO:2000146: negative regulation of cell motility GO:0006662: glycerol ether metabolic process GO:0036297: interstrand cross-link repair GO:0042743: hydrogen peroxide metabolic process GO:0098810: neurotransmitter reuptake GO:0032069: regulation of nuclease activity GO:0007030: Golgi organization 2 8 4 6 -log10(P)