

**dephosphorylation, GO:0016311**

- phosphoprotein phosphatase activity, GO:0004721
- phosphatase activity, GO:0016791
- protein tyrosine phosphatase activity, GO:0004725
- peptidyl-tyrosine dephosphorylation, GO:0035335
- sphingolipid biosynthetic process, GO:0030148
- inositol phosphate metabolic process, GO:0043647
- gluconeogenesis, GO:0006094
- hematopoietic progenitor cell differentiation, GO:0002244
- response to insulin, GO:0032868
- phospholipid metabolic process, GO:0006644
- phosphatidylinositol-3-phosphatase activity, GO:0004438
- platelet-derived growth factor receptor binding, GO:0005161
- anchored component of membrane, GO:0031225
- neuron development, GO:0048666
- drug metabolic process, GO:0017144
- phosphatidylethanolamine biosynthetic process, GO:0006646
- lymphangiogenesis, GO:0001946
- transmembrane receptor protein tyrosine phosphatase activity, GO:0005001
- positive regulation of glycolytic process, GO:0045821
- regulation of innate immune response, GO:0045088
- endochondral ossification, GO:0001958
- regulation of insulin secretion, GO:0050796
- negative regulation of T cell receptor signaling pathway, GO:0050860
- regulation of gluconeogenesis, GO:0006111
- endothelial cell migration, GO:0043542
- negative regulation of T cell activation, GO:0050868
- hormone metabolic process, GO:0042445
- negative regulation of collateral sprouting, GO:0048671
- negative regulation of protein kinase activity by regulation of protein phosphorylation, GO:0044387
- phosphate-containing compound metabolic process, GO:0006796
- fatty acid catabolic process, GO:0009062
- pyrimidine nucleoside salvage, GO:0043097
- fructose metabolic process, GO:0006000
- negative regulation of hormone secretion, GO:0046888
- atrioventricular canal development, GO:0036302
- MAP kinase tyrosine/serine/threonine phosphatase activity, GO:0017017
- homotypic cell-cell adhesion, GO:0034109
- L-methionine salvage from methylthioadenosine, GO:0019509
- purine nucleotide catabolic process, GO:0006195
- positive regulation of regulatory T cell differentiation, GO:0045591
- myelin assembly, GO:0032288
- negative regulation of production of miRNAs involved in gene silencing by miRNA, GO:1903799