

**protein dephosphorylation, GO:0006470**

- phosphoprotein phosphatase activity, GO:0004721
- dephosphorylation, GO:0016311
- phosphatase activity, GO:0016791
- calmodulin binding, GO:0005516
- protein tyrosine phosphatase activity, GO:0004725
- peptidyl-tyrosine dephosphorylation, GO:0035335
- protein serine/threonine phosphatase activity, GO:0004722
- phosphotyrosine residue binding, GO:0001784
- sphingolipid biosynthetic process, GO:0030148
- cytoplasmic side of plasma membrane, GO:0009898
- protein tyrosine/serine/threonine phosphatase activity, GO:0008138
- T cell activation, GO:0042110
- glycogen metabolic process, GO:0005977
- cell morphogenesis, GO:0000902
- face morphogenesis, GO:0060325
- entrainment of circadian clock by photoperiod, GO:0043153
- negative regulation of MAP kinase activity, GO:0043407
- positive regulation of protein export from nucleus, GO:0046827
- peptidyl-threonine dephosphorylation, GO:0035970
- sarcomere organization, GO:0045214
- B cell receptor signaling pathway, GO:0050853
- embryonic skeletal system morphogenesis, GO:0048704
- endothelial cell migration, GO:0043542
- regulation of axonogenesis, GO:0050770
- cation binding, GO:0043169
- mitogen-activated protein kinase kinase kinase binding, GO:0031435
- regulation of acetyl-CoA biosynthetic process from pyruvate, GO:0010510
- negative regulation of T cell activation, GO:0050868
- phosphatidylinositol-3-phosphatase activity, GO:0004438
- focal adhesion assembly, GO:0048041
- non-membrane spanning protein tyrosine phosphatase activity, GO:0004726
- cellular response to ATP, GO:0071318
- negative regulation of DNA biosynthetic process, GO:2000279
- phospholipid metabolic process, GO:0006644
- negative regulation of insulin secretion, GO:0046676
- JUN kinase binding, GO:0008432
- regulation of epidermal growth factor receptor signaling pathway, GO:0042058
- negative regulation of epithelial cell migration, GO:0010633
- phosphate-containing compound metabolic process, GO:0006796
- lymphangiogenesis, GO:0001946
- positive regulation of actin filament depolymerization, GO:0030836
- positive regulation of toll-like receptor 3 signaling pathway, GO:0034141
- cardiac muscle fiber development, GO:0048739
- negative regulation of MAPK cascade, GO:0043409
- negative regulation of T cell receptor signaling pathway, GO:0050860
- positive regulation of insulin secretion involved in cellular response to glucose stimulus, GO:0035774
- nucleotide phosphatase activity, GO:0098519
- MAP kinase tyrosine/serine/threonine phosphatase activity, GO:0017017
- negative regulation of protein tyrosine kinase activity, GO:0061099
- positive regulation of regulatory T cell differentiation, GO:0045591
- positive regulation of toll-like receptor 4 signaling pathway, GO:0034145
- negative regulation of peptidyl-tyrosine phosphorylation, GO:0050732