

phosphoprotein phosphatase activity, GO:0004721

- dephosphorylation, GO:0016311
- phosphatidylinositol biosynthetic process, GO:0006661
- protein tyrosine phosphatase activity, GO:0004725
- peptidyl-tyrosine dephosphorylation, GO:0035335
- phosphatase activity, GO:0016791
- T cell differentiation, GO:0030217
- glycogen metabolic process, GO:0005977
- protein tyrosine/serine/threonine phosphatase activity, GO:0008138
- cellular response to drug, GO:0035690
- peptidyl-threonine dephosphorylation, GO:0035970
- phosphatidylinositol dephosphorylation, GO:0046856
- cytoplasmic side of plasma membrane, GO:0009898
- phosphatidylinositol-3-phosphatase activity, GO:0004438
- negative regulation of MAPK cascade, GO:0043409
- phosphatidylinositol-3, GO:0052629
- negative regulation of BMP signaling pathway, GO:0030514
- regulation of acetyl-CoA biosynthetic process from pyruvate, GO:0010510
- negative regulation of T cell receptor signaling pathway, GO:0050860
- regulation of ventricular cardiac muscle cell action potential, GO:0098911
- phosphate-containing compound metabolic process, GO:0006796
- lymphangiogenesis, GO:0001946
- response to denervation involved in regulation of muscle adaptation, GO:0014894
- negative regulation of T cell proliferation, GO:0042130
- regulation of innate immune response, GO:0045088
- megakaryocyte development, GO:0035855
- positive regulation of hormone secretion, GO:0046887
- entrainment of circadian clock by photoperiod, GO:0043153
- negative regulation of insulin receptor signaling pathway, GO:0046627
- calcineurin complex, GO:0005955
- negative regulation of epithelial cell migration, GO:0010633
- negative regulation of cell adhesion mediated by integrin, GO:0033629
- positive regulation of triglyceride biosynthetic process, GO:0010867
- signaling, GO:0023052
- lamin binding, GO:0005521
- semicircular canal morphogenesis, GO:0048752
- locomotion involved in locomotory behavior, GO:0031987
- non-membrane spanning protein tyrosine phosphatase activity, GO:0004726
- embryonic skeletal system morphogenesis, GO:0048704
- positive regulation of insulin secretion involved in cellular response to glucose stimulus, GO:0035774
- anatomical structure development, GO:0048856
- presynaptic membrane assembly, GO:0097105
- positive regulation of regulatory T cell differentiation, GO:0045591
- phosphatidate phosphatase activity, GO:0008195
- negative regulation of lipid storage, GO:0010888
- MAP kinase tyrosine/serine/threonine phosphatase activity, GO:0017017
- leading edge membrane, GO:0031256
- heparan sulfate proteoglycan binding, GO:0043395
- negative regulation of hormone secretion, GO:0046888
- maternal behavior, GO:0042711
- negative regulation of protein kinase activity by regulation of protein phosphorylation, GO:0044387
- calcineurin-NFAT signaling cascade, GO:0033173