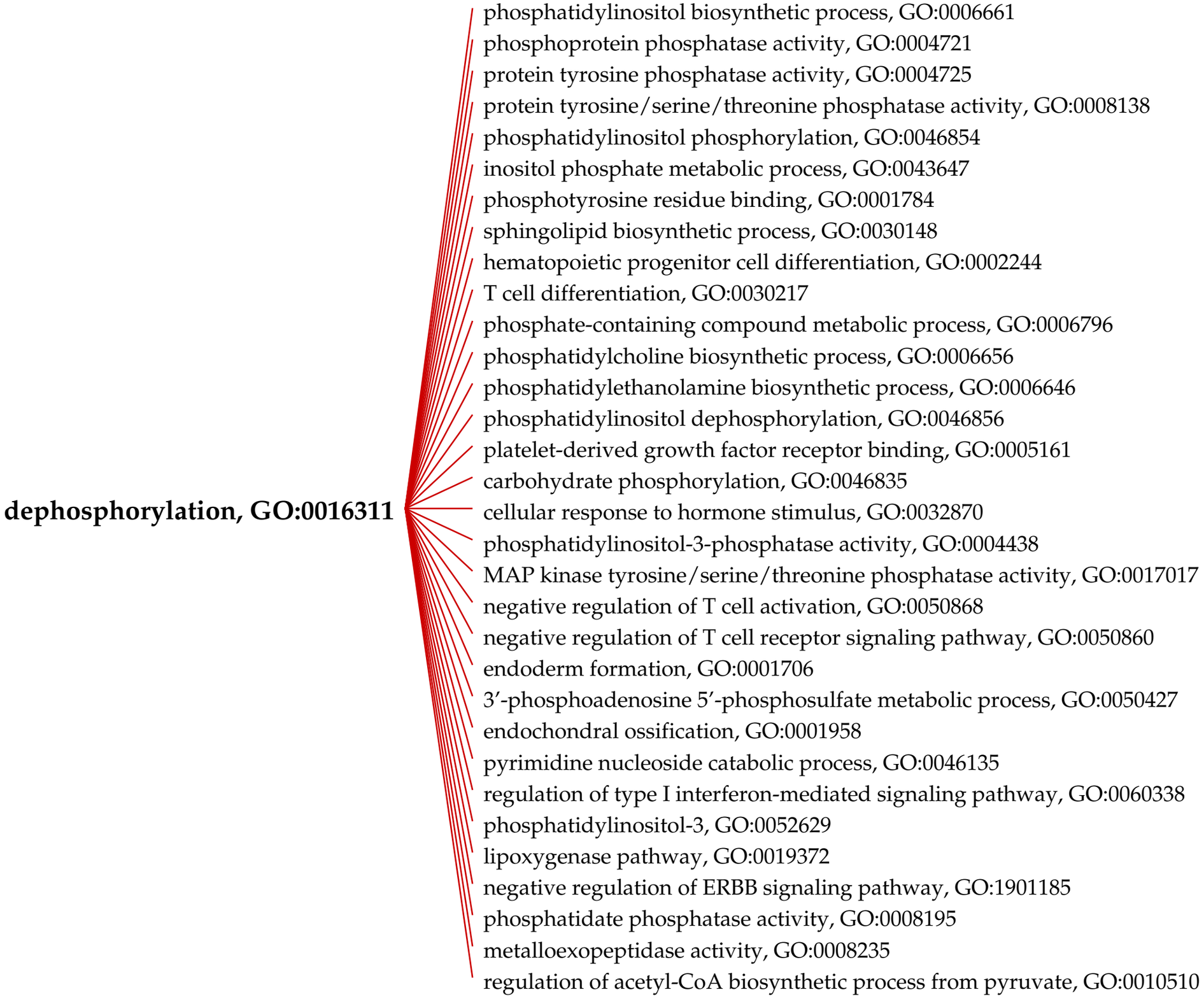
dephosphorylation, GO:0016311
phosphatidylinositol biosynthetic process, GO:0006661
phosphoprotein phosphatase activity, GO:0004721
protein tyrosine phosphatase activity, GO:0004725
protein tyrosine/serine/threonine phosphatase activity, GO:0008138
phosphatidylinositol phosphorylation, GO:0046854
inositol phosphate metabolic process, GO:0043647
phosphotyrosine residue binding, GO:0001784
sphingolipid biosynthetic process, GO:0030148
hematopoietic progenitor cell differentiation, GO:0002244
T cell differentiation, GO:0030217
phosphate-containing compound metabolic process, GO:0006796
phosphatidylcholine biosynthetic process, GO:0006656
phosphatidylethanolamine biosynthetic process, GO:0006646
phosphatidylinositol dephosphorylation, GO:0046856
platelet-derived growth factor receptor binding, GO:0005161
carbohydrate phosphorylation, GO:0046835
cellular response to hormone stimulus, GO:0032870
phosphatidylinositol-3-phosphatase activity, GO:0004438
MAP kinase tyrosine/serine/threonine phosphatase activity, GO:0017017
negative regulation of T cell activation, GO:0050868
negative regulation of T cell receptor signaling pathway, GO:0050860
endoderm formation, GO:0001706
3′-phosphoadenosine 5′-phosphosulfate metabolic process, GO:0050427
endochondral ossification, GO:0001958
pyrimidine nucleoside catabolic process, GO:0046135
regulation of type I interferon-mediated signaling pathway, GO:0060338
phosphatidylinositol-3, GO:0052629
lipoxygenase pathway, GO:0019372
negative regulation of ERBB signaling pathway, GO:1901185
phosphatidate phosphatase activity, GO:0008195
metalloexopeptidase activity, GO:0008235
regulation of acetyl-CoA biosynthetic process from pyruvate, GO:0010510