calmodulin binding, GO:0005516

interferon-gamma-mediated signaling pathway, GO:0060333 calcium ion transport, GO:0006816 sarcoplasmic reticulum, GO:0016529 calcium ion transmembrane transport, GO:0070588 phosphoric diester hydrolase activity, GO:0008081 phosphatidylinositol-3, GO:0005547 generation of precursor metabolites and energy, GO:0006091 potassium ion transmembrane transport, GO:0071805 muscle contraction, GO:0006936 cellular response to calcium ion, GO:0071277 calcium channel activity, GO:0005262 inner ear development, GO:0048839 negative regulation of signal transduction, GO:0009968 sodium ion transport, GO:0006814 glycogen catabolic process, GO:0005980 plasma membrane raft, GO:0044853 sarcoplasmic reticulum membrane, GO:0033017 potassium channel activity, GO:0005267 sodium ion transmembrane transport, GO:0035725 intrinsic component of plasma membrane, GO:0031226 lipid catabolic process, GO:0016042 phosphatidylserine binding, GO:0001786 sodium ion import across plasma membrane, GO:0098719 actomyosin structure organization, GO:0031032 muscle filament sliding, GO:0030049 phosphatidylinositol phospholipase C activity, GO:0004435 neuronal action potential, GO:0019228 striated muscle contraction, GO:0006941 positive regulation of calcium ion-dependent exocytosis, GO:0045956 positive regulation of blood vessel diameter, GO:0097755 calcium ion-regulated exocytosis of neurotransmitter, GO:0048791 release of sequestered calcium ion into cytosol by sarcoplasmic reticulum, GO:0014808 regulation of calcium ion-dependent exocytosis, GO:0017158 exogenous drug catabolic process, GO:0042738 calcium ion homeostasis, GO:0055074 voltage-gated potassium channel activity, GO:0005249 cAMP catabolic process, GO:0006198 membrane depolarization during action potential, GO:0086010 positive regulation of peptidyl-tyrosine autophosphorylation, GO:1900086 3', GO:0004114 negative regulation of blood pressure, GO:0045776 regulation of postsynaptic membrane potential, GO:0060078