

**positive regulation of protein targeting to mitochondrion, GO:1903955**

- phospholipid biosynthetic process, GO:0008654
- protein K11-linked ubiquitination, GO:0070979
- ubiquitin conjugating enzyme activity, GO:0061631
- protein palmitoylation, GO:0018345
- regulation of fat cell differentiation, GO:0045598
- piRNA metabolic process, GO:0034587
- cardiac left ventricle morphogenesis, GO:0003214
- labyrinthine layer blood vessel development, GO:0060716
- granulocyte differentiation, GO:0030851
- phosphotransferase activity, GO:0016773
- startle response, GO:0001964
- neuronal stem cell population maintenance, GO:0097150
- activation of transmembrane receptor protein tyrosine kinase activity, GO:0007171
- muscle alpha-actinin binding, GO:0051371
- cytochrome-b5 reductase activity, GO:0004128
- embryonic camera-type eye development, GO:0031076
- positive regulation of peptidyl-threonine phosphorylation, GO:0010800
- pancreas development, GO:0031016
- inner ear receptor stereocilium organization, GO:0060122
- peripheral nervous system development, GO:0007422
- structural molecule activity conferring elasticity, GO:0097493
- regulation of systemic arterial blood pressure, GO:0003073
- pituitary gland development, GO:0021983
- proteoglycan binding, GO:0043394
- muscle myosin complex, GO:0005859