

post-embryonic development, GO:0009791

- transcriptional repressor activity, GO:0001078
- hemopoiesis, GO:0030097
- multicellular organism growth, GO:0035264
- palate development, GO:0060021
- skeletal system development, GO:0001501
- transcriptional activator activity, GO:0001228
- cellular response to transforming growth factor beta stimulus, GO:0071560
- regulation of synaptic plasticity, GO:0048167
- ureteric bud development, GO:0001657
- heart morphogenesis, GO:0003007
- bone development, GO:0060348
- signal transduction by protein phosphorylation, GO:0023014
- skeletal system morphogenesis, GO:0048705
- pattern specification process, GO:0007389
- embryonic cranial skeleton morphogenesis, GO:0048701
- memory, GO:0007613
- retina development in camera-type eye, GO:0060041
- SMAD protein signal transduction, GO:0060395
- positive regulation of pathway-restricted SMAD protein phosphorylation, GO:0010862
- anterior/posterior pattern specification, GO:0009952
- face morphogenesis, GO:0060325
- negative regulation of TORC1 signaling, GO:1904262
- GATOR2 complex, GO:0061700
- embryonic limb morphogenesis, GO:0030326
- pancreas development, GO:0031016
- transforming growth factor beta receptor binding, GO:0005160
- odontogenesis, GO:0042476
- embryonic pattern specification, GO:0009880
- chondrocyte development, GO:0002063
- low-density lipoprotein particle, GO:0034362
- smooth muscle tissue development, GO:0048745
- bone mineralization, GO:0030282
- heterotrimeric G-protein complex, GO:0005834
- voltage-gated sodium channel complex, GO:0001518
- positive regulation of mesenchymal cell proliferation, GO:0002053
- positive regulation of bone mineralization, GO:0030501
- definitive hemopoiesis, GO:0060216
- enhancer binding, GO:0035326
- response to light stimulus, GO:0009416
- eye photoreceptor cell development, GO:0042462
- positive regulation of smooth muscle cell migration, GO:0014911
- hair follicle morphogenesis, GO:0031069
- artery morphogenesis, GO:0048844
- decidualization, GO:0046697
- regulation of pH, GO:0006885
- positive regulation of smoothened signaling pathway, GO:0045880
- developmental growth, GO:0048589
- long-term memory, GO:0007616
- exogenous drug catabolic process, GO:0042738
- cardiac right ventricle morphogenesis, GO:0003215
- maternal behavior, GO:0042711
- neuron fate commitment, GO:0048663
- nuclear outer membrane, GO:0005640
- anion transmembrane transporter activity, GO:0008509
- embryonic retina morphogenesis in camera-type eye, GO:0060059
- positive regulation of cartilage development, GO:0061036
- head development, GO:0060322
- mesenchymal cell differentiation, GO:0048762
- anatomical structure formation involved in morphogenesis, GO:0048646
- dopaminergic neuron differentiation, GO:0071542
- female pregnancy, GO:0007565
- mammary gland development, GO:0030879
- cardiac atrium morphogenesis, GO:0003209
- ear development, GO:0043583
- aromatase activity, GO:0070330
- cellular response to BMP stimulus, GO:0071773