multicellular organism growth, GO:0035264 smoothened signaling pathway, GO:0007224 axoneme, GO:0005930 chondrocyte differentiation, GO:0002062 BMP signaling pathway, GO:0030509 bone development, GO:0060348 determination of left/right symmetry, GO:0007368 palate development, GO:0060021 response to progesterone, GO:0032570 negative regulation of smoothened signaling pathway, GO:0045879 mammary gland development, GO:0030879 odontogenesis, GO:0042476 positive regulation of bone mineralization, GO:0030501 semaphorin-plexin signaling pathway, GO:0071526 response to mechanical stimulus, GO:0009612 regulation of smoothened signaling pathway, GO:0008589 motile cilium assembly, GO:0044458 cullin-RING ubiquitin ligase complex, GO:0031461 heart looping, GO:0001947 negative chemotaxis, GO:0050919 positive regulation of cell cycle arrest, GO:0071158 face morphogenesis, GO:0060325 SMAD protein signal transduction, GO:0060395 positive regulation of protein dephosphorylation, GO:0035307 cytoplasmic sequestering of transcription factor, GO:0042994 positive regulation of pri-miRNA transcription from RNA polymerase II promoter, GO:1902895

## neural tube closure, GO:0001843

female pregnancy, GO:0007565 camera-type eye morphogenesis, GO:0048593 smooth muscle tissue development, GO:0048745 negative regulation of actin filament polymerization, GO:0030837 head development, GO:0060322 positive regulation of receptor recycling, GO:0001921 plasma membrane raft, GO:0044853 negative regulation of fibroblast proliferation, GO:0048147 cell junction assembly, GO:0034329 embryonic skeletal joint morphogenesis, GO:0060272 lamellipodium membrane, GO:0031258 positive regulation of smoothened signaling pathway, GO:0045880 positive regulation of fatty acid beta-oxidation, GO:0032000 maintenance of epithelial cell apical/basal polarity, GO:0045199 cell-cell junction organization, GO:0045216 pulmonary valve morphogenesis, GO:0003184 receptor agonist activity, GO:0048018 BMP receptor binding, GO:0070700 actin polymerization or depolymerization, GO:0008154 positive regulation of T-helper 2 cell differentiation, GO:0045630 outflow tract septum morphogenesis, GO:0003148 growth factor activity, GO:0008083 cardiac muscle cell proliferation, GO:0060038 sperm principal piece, GO:0097228 BBSome, GO:0034464 cartilage condensation, GO:0001502 central nervous system neuron differentiation, GO:0021953 response to cholesterol, GO:0070723

spinal cord motor neuron differentiation, GO:0021522

cochlea morphogenesis, GO:0090103

epithelial cell morphogenesis, GO:0003382

frizzled binding, GO:0005109

cellular response to BMP stimulus, GO:0071773