positive regulation of osteoblast differentiation, GO:0045669

cytokine receptor activity, GO:0004896
positive regulation of T cell proliferation, GO:0042102
regulation of proteasomal ubiquitin-dependent protein catabolic process, GO:0032434
regulation of protein localization to cell surface, GO:2000008
intrinsic component of membrane, GO:0031224
embryonic skeletal system development, GO:0048706

ventricular septum development, GO:0003281
armadillo repeat domain binding, GO:0070016
proton-transporting V-type ATPase, GO:0033180
glomerular visceral epithelial cell development, GO:0072015
T-helper 17 cell lineage commitment, GO:0072540

multicellular organismal iron ion homeostasis, GO:0060586