ubiquitin ligase complex, GO:0000151

ligand-dependent nuclear receptor transcription coactivator activity, GO:0030374 mediator complex, GO:0016592 ubiquitin conjugating enzyme activity, GO:0061631 protein K11-linked ubiquitination, GO:0070979 ubiquitin conjugating enzyme binding, GO:0031624 RNA polymerase II transcription cofactor activity, GO:0001104 K63-linked polyubiquitin binding, GO:0070530 regulation of androgen receptor signaling pathway, GO:0060765 positive regulation of ubiquitin-protein transferase activity, GO:0051443 regulation of reactive oxygen species metabolic process, GO:2000377 PRC1 complex, GO:0035102 Cul4A-RING E3 ubiquitin ligase complex, GO:0031464 monocyte differentiation, GO:0030224 cellular response to steroid hormone stimulus, GO:0071383 histone ubiquitination, GO:0016574 IkappaB kinase complex, GO:0008385 positive regulation of mammary gland epithelial cell proliferation, GO:0033601 activation of MAPKKK activity, GO:0000185 cellular protein catabolic process, GO:0044257 regulation of dendrite morphogenesis, GO:0048814 regulation of transforming growth factor beta receptor signaling pathway, GO:0017015 ventricular trabecula myocardium morphogenesis, GO:0003222 thyroid hormone generation, GO:0006590 sex chromatin, GO:0001739