ubiquitin ligase complex, GO:0000151

viral entry into host cell, GO:0046718 RNA polymerase II transcription cofactor activity, GO:0001104 mediator complex, GO:0016592 protein K11-linked ubiquitination, GO:0070979 ubiquitin conjugating enzyme binding, GO:0031624 ubiquitin conjugating enzyme activity, GO:0061631 ventricular trabecula myocardium morphogenesis, GO:0003222 IkappaB kinase complex, GO:0008385 positive regulation of ubiquitin-protein transferase activity, GO:0051443 protein maturation, GO:0051604 positive regulation of cellular protein catabolic process, GO:1903364 limb development, GO:0060173 ubiquitin-dependent SMAD protein catabolic process, GO:0030579 thyroid hormone generation, GO:0006590 histone ubiquitination, GO:0016574 suckling behavior, GO:0001967 positive regulation of cell cycle arrest, GO:0071158 photoreceptor cell outer segment organization, GO:0035845