regulation of MHC class II biosynthetic process (GO:0045346) regulation of intrinsic apoptotic signaling pathway in response to DNA damage by p53 class mediator (GO:1902165) 8.0 regulation of DNA methylation (GO:0044030) negative regulation of protein acetylation (GO:1901984) negative regulation of peptidyl-lysine acetylation (GO:2000757) 0.6 negative regulation of myoblast differentiation (GO:0045662) negative regulation of intrinsic apoptotic signaling pathway in response to DNA damage by p53 class mediator (GO:1902166) 0.4negative regulation of intrinsic apoptotic signaling pathway by p53 class mediator (GO:1902254) negative regulation of histone acetylation (GO:0035067) macrophage differentiation (GO:0030225) 0.2 lymphoid progenitor cell differentiation (GO:0002320) histone H2A ubiquitination (GO:0033522) anatomical structure regression (GO:0060033) granulocyte differentiation (GO:0030851) negative regulation of peptidase activity (GO:0010466) DNA recombination (GO:0006310) transmembrane receptor protein serine/threonine kinase signaling pathway (GO:0007178) cellular response to transforming growth factor beta stimulus (GO:0071560) response to transforming growth factor beta (GO:0071559) regulation of cellular response to growth factor stimulus (GO:0090287) regulation of transmembrane receptor protein serine/threonine kinase signaling pathway (GO:0090092) midbrain development (GO:0030901) forebrain morphogenesis (GO:0048853) lateral ventricle development (GO:0021670) protein deubiquitination (GO:0016579) protein modification by small protein removal (GO:0070646) proteolysis involved in cellular protein catabolic process (GO:0051603) ubiquitin-dependent protein catabolic process (GO:0006511) modification-dependent macromolecule catabolic process (GO:0043632) modification-dependent protein catabolic process (GO:0019941) negative regulation of transforming growth factor beta receptor signaling pathway (GO:0030512) regulation of transforming growth factor beta receptor signaling pathway (GO:0017015) negative regulation of cellular response to growth factor stimulus (GO:0090288) negative regulation of transmembrane receptor protein serine/threonine kinase signaling pathway (GO:0090101) regulation of proteasomal protein catabolic process (GO:0061136) transforming growth factor beta receptor signaling pathway (GO:0007179) regulation of cellular protein catabolic process (GO:1903362) regulation of proteolysis involved in cellular protein catabolic process (GO:1903050)