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regulation of MHC class II biosynthetic process (GO:0045346)

regulation of DNA methylation (GO:0044030)

phosphatidylinositol dephosphorylation (GO:0046856)

parental behavior (GO:0060746)

oocyte development (GO:0048599)

negative regulation of protein acetylation (GO:1901984)

negative regulation of peptidyl–lysine acetylation (GO:2000757)

negative regulation of histone acetylation (GO:0035067)

multicellular organismal reproductive behavior (GO:0033057)

macrophage differentiation (GO:0030225)

lymphoid progenitor cell differentiation (GO:0002320)

apoptotic process involved in morphogenesis (GO:0060561)

granulocyte differentiation (GO:0030851)

magnesium ion transport (GO:0015693)

Fc receptor signaling pathway (GO:0038093)

Fc–epsilon receptor signaling pathway (GO:0038095)

regulation of protein depolymerization (GO:1901879)

regulation of fatty acid metabolic process (GO:0019217)

regulation of protein complex disassembly (GO:0043244)

negative regulation of protein complex disassembly (GO:0043242)

negative regulation of protein depolymerization (GO:1901880)

regulation of microtubule polymerization or depolymerization (GO:0031110)

regulation of microtubule depolymerization (GO:0031114)

regulation of fatty acid biosynthetic process (GO:0042304)

positive regulation of fatty acid metabolic process (GO:0045923)

positive regulation of fatty acid biosynthetic process (GO:0045723)

nucleotide–binding oligomerization domain containing signaling pathway (GO:0070423)

negative regulation of microtubule polymerization or depolymerization (GO:0031111)

negative regulation of microtubule depolymerization (GO:0007026)

midbrain development (GO:0030901)

lateral ventricle development (GO:0021670)

cytoplasmic pattern recognition receptor signaling pathway (GO:0002753)

forebrain morphogenesis (GO:0048853)

I–kappaB kinase/NF–kappaB signaling (GO:0007249)

nucleotide–binding domain, leucine rich repeat containing receptor signaling pathway (GO:0035872)

