8 regulation of MHC class II biosynthetic process (GO:0045346) regulation of DNA methylation (GO:0044030) phosphatidylinositol dephosphorylation (GO:0046856) 6 parental behavior (GO:0060746) oocyte development (GO:0048599) negative regulation of protein acetylation (GO:1901984) 4 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) ls10ppm IsLHet10ppm