

Is10ppm

IsCdHet

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- pigmentation (GO:0043473)
- chromatin modification (GO:0016568)
- carbohydrate derivative biosynthetic process (GO:1901137)
- regulation of Rho GTPase activity (GO:0032319)
- histone ubiquitination (GO:0016574)
- positive regulation of Rho GTPase activity (GO:0032321)
- positive regulation of Ras GTPase activity (GO:0032320)
- positive regulation of GTP catabolic process (GO:0033126)
- regulation of Wnt signaling pathway (GO:0030111)
- regulation of sulfur metabolic process (GO:0042762)
- regulation of microtubule depolymerization (GO:0031114)
- regulation of cofactor metabolic process (GO:0051193)
- regulation of coenzyme metabolic process (GO:0051196)
- regulation of acyl-CoA biosynthetic process (GO:0050812)
- regulation of acetyl-CoA biosynthetic process from pyruvate (GO:0010510)
- positive regulation of potassium ion transport (GO:0043268)
- positive regulation of dendrite morphogenesis (GO:0050775)
- polyamine metabolic process (GO:0006595)
- polyamine biosynthetic process (GO:0006596)
- peptide hormone processing (GO:0016486)
- organ morphogenesis (GO:0009887)
- neurotrophin TRK receptor signaling pathway (GO:0048011)
- negative regulation of protein depolymerization (GO:1901880)
- negative regulation of organelle organization (GO:0010639)
- negative regulation of microtubule polymerization or depolymerization (GO:0031111)
- negative regulation of microtubule depolymerization (GO:0007026)
- negative regulation of cell-matrix adhesion (GO:0001953)
- histone H2A ubiquitination (GO:0033522)
- cerebral cortex cell migration (GO:0021795)
- cellular biogenic amine biosynthetic process (GO:0042401)
- amine biosynthetic process (GO:0009309)
- body morphogenesis (GO:0010171)
- nucleotide-binding oligomerization domain containing signaling pathway (GO:0070423)
- cellular response to fatty acid (GO:0071398)
- neurotrophin signaling pathway (GO:0038179)
- replacement ossification (GO:0036075)
- regulation of vascular endothelial growth factor receptor signaling pathway (GO:0030947)
- regulation of ryanodine-sensitive calcium-release channel activity (GO:0060314)
- regulation of membrane repolarization (GO:0060306)
- negative regulation of potassium ion transport (GO:0043267)
- endochondral ossification (GO:0001958)
- lung alveolus development (GO:0048286)
- glycolytic process (GO:0006096)
- negative regulation of signal transduction by p53 class mediator (GO:1901797)
- tricarboxylic acid cycle (GO:0006099)

