pigmentation (GO:0043473) DNA packaging (GO:0006323) chromatin modification (GO:0016568) positive regulation of Ras GTPase activity (GO:0032320) regulation of Wnt signaling pathway (GO:0030111) regulation of sulfur metabolic process (GO:0042762) regulation of potassium ion transmembrane transport (GO:1901379) 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 GTP catabolic process (GO:0033126) 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) nucleotide-binding oligomerization domain containing signaling pathway (GO:0070423) neurotrophin TRK receptor signaling pathway (GO:0048011) neurotrophin signaling pathway (GO:0038179) 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 ubiquitination (GO:0016574) histone H2A ubiquitination (GO:0033522) cerebral cortex cell migration (GO:0021795) cellular response to fatty acid (GO:0071398) cellular biogenic amine biosynthetic process (GO:0042401) carbohydrate derivative biosynthetic process (GO:1901137) amine biosynthetic process (GO:0009309) body morphogenesis (GO:0010171) 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) endochondral ossification (GO:0001958) negative regulation of potassium ion transport (GO:0043267) glycolytic process (GO:0006096) negative regulation of signal transduction by p53 class mediator (GO:1901797) tricarboxylic acid cycle (GO:0006099) IsCdHet10ppm

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