positive regulation of heterotypic cell-cell adhesion positive regulation of cell-substrate adhesion positive regulation of cell adhesion cell adhesion negative regulation of cell-matrix adhesion monocyte aggregation cartilage condensation spindle assembly regulation of skeletal muscle cell proliferation positive regulation of skeletal muscle cell proliferation cell cycle and proliferation positive regulation of G2/M transition of mitotic cell cycle positive regulation of epithelial cell proliferation positive regulation of endothelial cell proliferation mitotic spindle organization lymphocyte proliferation transcription initiation from RNA polyme spermatogenesis, exchange of chromatogenesis, cell organization and biogenesis Spemann organizer formation regulation of neurotransmitter uptake positive regulation of synaptic transmission, GABAergic cell-cell signaling positive regulation of synaptic transmission, dopaminergic negative regulation of growth hormone secretion modulation of chemical synaptic transmission long-term synaptic potentiation suppression by virus of host apoptotic process regulation of smooth muscle cell apoptotic process negative regulation of cell death apoptotic process anterior ed weight near batte in teach transcription-coupled nucleotide-excision repair pyrimidine dimer repair by nucleotide-excision repair positive regulation of mitotic recombination nucleotide-excision repair, DNA incisior nucleotide-excision repair, DNA duplex unwinding mitotic recombinati mitochondrial DNA replication mitochondrial DNA repair double-strand break repair via single-strand annealing over_represented_pvalue DNA replication, synthesis of RNA primer DNA replication, removal of RNA primer DNA replication, Okazaki fragment processing white fat cell differentiation positive regulation et positive regulation et psy positive regulation of humoral immune respense positive regulation of the of the second other biological processes macrophage attration in the comment of the comment detection of temperature stimulus develveship activation of Infiliation regulation of generation of precurs positive regulation of biosynthetic process of antibacterial peditides ac positive regulation of biosynthetic process of antibacterial peditides ac positive regulation of biosynthetic process of antibacterial peditides ac other metabolic processes negative regulation of prostaglandin negative regulation — arabino aerobic electron transport oran regulation of receptor internalization regulation of proteptory internalization regulation of protein ublquitination regulation of protein ublquitination protein transport to vacuole involved in ubiquitin—dependent protein catabolic process via the multivesicular body sorting pathway protein by sulfation protein by protein p tRNA methylation transcription initiation from RNA polymerase II promoter transcription elongation from RNA polymerase I promoter spliceosomal complex assembly snoRNA metabolic process rDNA heterochromatin assembly production of siRNA involved in RNA interference positive regulation of transcription by RNA polymerase II negative regulation of NF-kappaB transcription factor activity mRNA splicing, via spliceosome metabolism mRNA 3'-end processing by stem-loop binding and cleavage maturation of LSU-rRNA from tricistronic rRNA transcript (SSU-rRNA, 5.8S rRNA, LSU-rRNA) glycyl-tRNA aminoacylation DNA replication, synthesis of RNA primer DNA replication, removal of RNA primer -DNA methylation-dependent heterochromatin assembly carbon catabolite repression of transcription by glucose -7-methylguanosine RNA capping -Vmt signaling pathway transmembrane receptor protein tyrosine kinase signaling pathway transmembrane receptor protein serine/threonine kinase signaling pathway regulation of Rho protein signal transduction regulation of MyD88—dependent toll—like receptor signaling pathway regulation of MAPK cascade regulation of BMP signaling pathway receptor recycling Rap protein 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