positive regulation of cell adhesion
negative regulation of cell-matrix adhesion
positive regulation of skeletal muscle cell proliferation
positive regulation of epithelial cell proliferation
positive regulation of endothelial cell proliferation
transcription initiation from RNA polymerase
positive regulation of
DNA methylation-dependent heterochromatin assembly
regulation of neurotransmitter uptake
negative regulation of growth hormone secretion
modulation of chemical synaptic transmission
long-term synaptic potentiation

HAUS complex
cornified envelope

regulation of smooth muscle cell apoptotic process
negative regulation of cell death
apoptotic process

progression of morphogenetic furrow involved in
embryonic
translesion synthesis
pyrimidine dimer repair by nucleotide-excision repair
DNA replication, synthesis of RNA primer

enzyme regulator activity
acetyltransferase activator activity

Golgi apparatus

transforming growth factor beta receptor activity, type I
activin receptor activity, type I

mitochondrial membrane

low-density lipoprotein particle
host cell

unmethylated CpG binding
RNA binding
double-stranded DNA binding
transcription factor TFIIH core complex
transcription factor
core TFIIH complex portion of holo TFIIH complex
alpha DNA polymerase:primase complex

positive regulation of
positive regulation of
detection of mechanical stimulus
detection of
RNA
RNase
preribosome, large subunit precursor
molybdopterin synthase complex
condensed chromosome
late endosome
ESCRT II complex
mitochondrial membrane
ESCRT II complex
heterotrimeric
activin receptor complex

positive regulation of biosynthetic process of
positive regulation of phosphatidyl
negative regulation of
diadenosine
RNA
NAD-dependent histone
heparosan-N
alpha-L

heterotrimeric G-protein complex
activin receptor complex

protein transport to vacuole involved in ubiquitin-dependent protein catabolic process via the multivesicular
regulation of proteolysis
positive
negative regulation of
M6 regulation of
transcription initiation from
maturation of LSU-rRNA from tricistronic
carbon catabolite repression of
regulation of MyD88-dependent toll-like receptor signaling pathway
nucleotide-binding
negative regulation of DNA damage response
interferon-gamma-mediated signaling pathway
transmembrane receptor protein tyrosine phosphatase activity
platelet-derived growth factor receptor binding
activin receptor activity, type I

positive regulation of biosynthetic process of
negative regulation of
negative regulation of DNA damage response, signal
cell

behavioral response to starvation

RNA polymerase II general transcription initiation factor activity
ribosomal large subunit export from nucleus

protein transport to vacuole involved in ubiquitin-dependent protein catabolic process via the
positive regulation of
negative regulation of
negative regulation of growth
endocytic recycling

iron ion transmembrane transporter activity

cell adhesion

cycle and

organization and

cell-cell signaling

cytoskeleton

death

developmental processes

DNA metabolism

enzyme regulator activity

ER/Golgi

kinase activity

mitochondrion

non-structural extracellular

acid binding

nucleus

other biological processes

other cellular component

other cytoplasmic organelle

other membranes

other metabolic processes

other molecular function

plasma membrane

protein metabolism

RNA metabolism

signal transduction

signal transduction activity

stress response

transcription regulatory activity

transport

transporter activity

over_represented_pvalue

ontology