

iron, 2 sulfur cluster binding	●	cysteinyl–tRNA aminoacylation	●	protein neddylation
–hydroxyphenylpyruvate dioxygenase activity	●	dipeptidyl–peptidase activity	●	protein quality control for misfolded or incompletely synthesized proteins
–nucleotidase activity	●	hydroxymethylglutaryl–CoA synthase activity	●	protein tyrosine kinase activity
midphosphoribosyltransferase activity	●	intrinsic apoptotic signaling pathway	●	protein–L–isoaspartate (D–aspartate) O–methyltransferase activity
P–1 adaptor complex	●	isoprenoid biosynthetic process	●	purine nucleobase biosynthetic process
apoptotic process	●	large ribosomal subunit	●	purine–rich negative regulatory element binding
beta–tubulin binding	●	Las1 complex	●	pyridoxamine–phosphate oxidase activity
ation:chloride symporter activity	●	lipid binding	●	pyridoxine biosynthetic process
cell communication	●	membrane coat	●	respiratory chain complex IV assembly
cellular copper ion homeostasis	●	NAD+ ADP–ribosyltransferase activity	●	RNA–dependent DNA biosynthetic process
cellular developmental process	●	NEDD8 activating enzyme activity	●	sphingolipid biosynthetic process
copper ion transport	●	oxidoreductase activity, acting on single donors with incorporation of molecular oxygen	●	sphingolipid delta–4 desaturase activity
ubiquitin–RING E3 ubiquitin ligase complex	●	peptide–methionine (S)–S–oxide reductase activity	●	thiopurine S–methyltransferase activity
cystathionine beta–synthase activity	●	peptidyl–amino acid modification	●	transferase activity, transferring amino–acyl groups
cysteine biosynthetic process from serine	●	peptidyl–diphthamide biosynthetic process from peptidyl–histidine	●	tricarboxylic acid cycle
cysteine biosynthetic process via cystathionine	●	phosphorelay signal transduction system	●	tubulin complex assembly
cysteine–tRNA ligase activity	●	post–chaperonin tubulin folding pathway		