_	 proteolysis		
<u> </u>	 transmembrane transport		
<u> </u>	 ATP synthesis coupled proton transport		
-	 immune response		
H	 xenobiotic transport		
H	 nitrogen compound transport		
<b>⊢</b>	 oxidation–reduction process		
<b>⊢</b>	 protein folding		
<b>⊢</b>	 response to oxidative stress		
<b>⊢</b>	 protein N–linked glycosylation		
<b>⊢</b>	 phospholipid metabolic process		
<b>⊢</b>	 arachidonic acid secretion		
<b>⊢</b>	 microtubule-based process		
-	 chromosome segregation		
-	 cilium assembly		
<b>-</b>	CTP biosynthetic process		
H	 DNA-templated transcription, initiation	as.	factor(cluster)
F	 kinetochore assembly		,
H	vacuolar transport	<b>A</b>	1
	superoxide metabolic process	_	2
<u> </u>	phosphatidylinositol metabolic process	_	_
	toll–like receptor signaling pathway	<b>A</b>	3
	mitochondrial electron transport, ubiquinol to cytochrome c		4
-	negative regulation of transcription by RNA polymerase II		4
	regulation of microtubule polymerization or depolymerization	<b>A</b>	5
	CENP–A containing nucleosome assembly		•
	cilium movement involved in cell motility	_	6
	tRNA 3'-trailer cleavage	<b>A</b>	7
	nucleosome assembly		
	regulation of DNA replication		
	chaperone-mediated protein transport		
	A dTMP biosynthetic process		
	mitotic cytokinesis		
	mitotic sister chromatid segregation		
	phospholipid catabolic process ribosomal large subunit assembly		
	A spindle organization		
	deoxyribonucleotide biosynthetic process		
	establishment of mitotic spindle localization		
L	witotic recombination—dependent replication fork processing		
	negative regulation of biosynthetic process		
	negative regulation of nucleic acid-templated transcription		
ł	nucleoside diphosphate phosphorylation		
	GTP biosynthetic process		
	UTP biosynthetic process		
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