

UP	Day 1	Day 3	Day 6	GO term	Description
				GO:0000080	G1 phase of mitotic cell cycle
				GO:0043648	dicarboxylic acid metabolic process
				GO:0015850	organic alcohol transport
				GO:0000422	mitochondrion degradation
				GO:0075306	regulation of conidium formation
				GO:0006468	protein amino acid phosphorylation
				GO:0034727	piecemeal microautophagy of nucleus
				GO:0070887	cellular response to chemical stimulus
				GO:0019395	fatty acid oxidation
				GO:0000160	two-component signal transduction system (phosphorelay)
				GO:0005993	trehalose catabolic process
				GO:0043603	cellular amide metabolic process
				GO:0019751	polyol metabolic process
				GO:0032258	CVT pathway
				GO:0034498	early endosome to Golgi transport
				GO:0007231	osmosensory signaling pathway
				GO:0009063	cellular amino acid catabolic process
				GO:0007039	vacuolar protein catabolic process
				GO:0000045	autophagic vacuole assembly
				GO:0019662	non-glycolytic fermentation
				GO:0012501	programmed cell death
				GO:0006081	cellular aldehyde metabolic process
				GO:0015718	monocarboxylic acid transport
				GO:0007047	cellular cell wall organization
				GO:0051173	positive regulation of nitrogen compound metabolic process
				GO:0032569	gene-specific transcription from RNA polymerase II promoter
				GO:0045860	positive regulation of protein kinase activity
				GO:0000165	MAPKKK cascade
				GO:0048468	cell development
				GO:0007186	G-protein coupled receptor protein signaling pathway
				GO:0006800	oxygen and reactive oxygen species metabolic process
				GO:0000747	conjugation with cellular fusion
				GO:0001403	invasive growth in response to glucose limitation
				GO:0000122	negative regulation of transcription from RNA polymerase II promoter
				GO:0007124	pseudohyphal growth

DOWN	Day 1	Day 3	Day 6	GO term	Description
				GO:0006360	transcription from RNA polymerase I promoter
				GO:0045039	protein import into mitochondrial inner membrane
				GO:0000027	ribosomal large subunit assembly
				GO:0006555	methionine metabolic process
				GO:0009108	coenzyme biosynthetic process
				GO:0006418	tRNA aminoacylation for protein translation
				GO:0000463	maturation of LSU-rRNA from tricistronic rRNA transcript
				GO:0030150	protein import into mitochondrial matrix
				GO:0000472	endonucleolytic cleavage from (SSU-rRNA, 5.8S rRNA, LSU-rRNA)
				GO:0006733	oxidoreduction coenzyme metabolic process
				GO:0000447	endonucleolytic cleavage in ITS1 to separate tricistronic rRNA transcript
				GO:0006407	rRNA export from nucleus
				GO:0051258	protein polymerization
				GO:0015986	ATP synthesis coupled proton transport
				GO:0033108	mitochondrial respiratory chain complex assembly
				GO:0042026	protein refolding
				GO:0000154	rRNA modification
				GO:0031204	posttranslational protein targeting to membrane, translocation
				GO:0071038	nuclear polyadenylation-dependent tRNA catabolic process
				GO:0006547	histidine metabolic process
				GO:0006614	SRP-dependent cotranslational protein targeting to membrane
				GO:0009067	aspartate family amino acid biosynthetic process
				GO:0042274	ribosomal small subunit biogenesis
				GO:0006450	regulation of translational fidelity
				GO:0006007	glucose catabolic process
				GO:0009060	aerobic respiration