

value

regulation of (1→3)–beta–D–glucan biosynthetic process			inositol lipid–mediated signaling			regulation of vesicle–mediated transport			negative regulation of phosphorylation			N–glycan processing		histidyl–tRNA aminoacylation		branched–chain amino acid biosynthetic process		pollen exine formation		radial axis specification		embryonic meristem development												
																polyamine biosynthetic process																		
hyperosmotic response			negative regulation of developmental growth		brassinosteroid homeostasis		regulation of defense response to insect		regulation of transcription from RNA polymerase II promoter in response to oxidative stress		negative regulation of transcription by RNA polymerase II		response to light intensity		mitochondrial translation		base conversion or substitution editing		DNA catabolic process		tetrapyrrole metabolic process		CUT catabolic process		anther wall tapetum cell differentiation		protein–containing complex assembly		maturation of 5.8S rRNA					
			cellular response to nitrogen starvation		response to flooding		cellular response to UV–C		photoperiodism, flowering		positive regulation of cellular response to oxidative stress		regulation of cellular component organization		negative regulation of DNA–templated transcription, initiation		pectin biosynthetic process		tRNA N2–guanine methylation		cytokinin catabolic process		D–gluconate catabolic process		protein deglycosylation		socket cell differentiation		male–female gamete recognition during double fertilization forming a zygote and endosperm		embryo development		dehiscence	
positive regulation of hydrogen peroxide metabolic process		positive regulation of catalytic activity		regulation of developmental process		regulation of double–strand break repair		regulation of seedling development		positive regulation of flavonoid biosynthetic process		regulation of ARF protein signal transduction		regulation of gibberellic acid mediated signaling pathway		indole–containing compound metabolic process		N–terminal protein myristoylation		snoRNA splicing		protein desumoylation		protein phosphorylation		plant–type primary cell wall biogenesis		abscission		endoplasmic reticulum tubular network organization		shoot system morphogenesis		
regulation of mitotic cell cycle		response to aluminum ion		regulation of carbohydrate metabolic process		defense response by callose deposition		signal transduction		double–strand break repair		response to zinc ion		negative regulation of vernalization response		phosphatidic acid biosynthetic process		galactolipid biosynthetic process		acylglycerol transport		neutral amino acid transport		iron ion transmembrane transport		lignin metabolic process								
negative regulation of response to salt stress		response to organonitrogen compound		regulation of cellular response to hypoxia		calcium–mediated signaling		response to toxic substance		interstrand cross–link repair		innate immune response		negative regulation of transcription by RNA polymerase III		tRNA export from nucleus		clathrin–coated vesicle cargo loading, AP–1–mediated		urea transport		chlorophyll cycle												
positive regulation of cell death		detection of brassinosteroid stimulus		regulation of tryptophan metabolic process		regulation of response to osmotic stress		regulation of histone H3–K9 dimethylation		cellular response to mechanical stimulus		positive regulation of metabolic process		regulation of cell death		phospholipid catabolic process		lipid biosynthetic process		sphingolipid biosynthetic process		ER to chloroplast lipid transport		oxygen transport		calcium ion transport		glyoxal metabolic process						