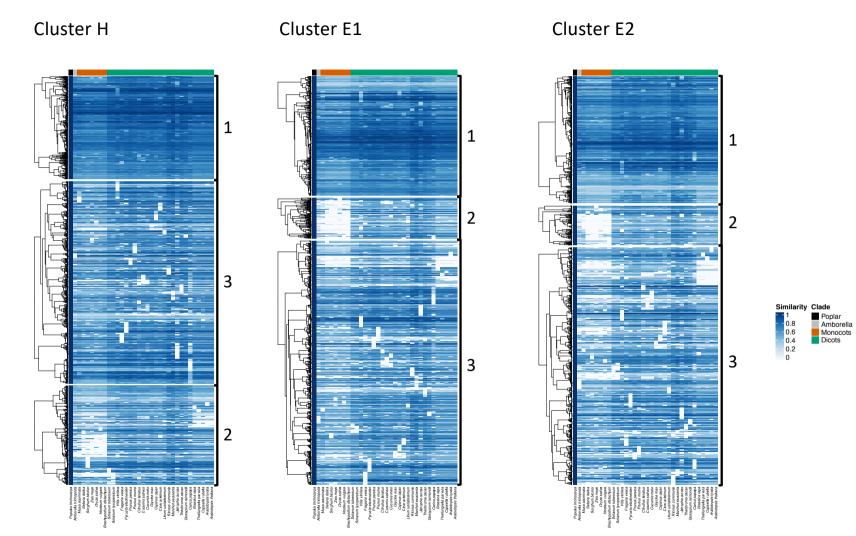
O category	GO description	Enrich	nmen	
OLOGICAL_PROCESS	1,2-dichloroethane catabolic process abscisic acid-activated signaling pathway			
	amino acid transmembrane transport			
	anther dehiscence			
	arabinan catabolic process			
	aromatic amino acid family metabolic process			
	aspartate family amino acid metabolic process			
	auxin efflux	•		
	auxin polar transport	•		
	auxin transport			
	callose deposition in cell wall	•		
	carbohydrate derivative catabolic process			
	carboxylic acid catabolic process			
	carboxylic acid transport			
	cell morphogenesis			
	cell morphogenesis involved in differentiation	•		
	cell proliferation			
	cellular amino acid biosynthetic process			
	cellular biogenic amine metabolic process			
	cellular metal ion homeostasis	•		
	cellular response to acid chemical		•	
	cellular response to oxygen-containing compound			
	cellular response to phosphate starvation	•		
	cellular response to starvation			
	cellulose biosynthetic process			
	cellulose microfibril organization			
	chemical homeostasis	Ĭ		
	coenzyme biosynthetic process		ı,	
	cotyledon vascular tissue pattern formation			
	cotyledon vascular tissue pattern formation cytoskeleton organization			
	defense response to bacterium defense response to bacterium, incompatible interaction			
	developmental cell growth divalent metal ion transport	•		
	DNA replication		•	
	embryonic morphogenesis			
	ethylene biosynthetic process	•		
	fatty acid biosynthetic process		•	
	fatty acid metabolic process			
	floral whorl development			
	fruit dehiscence			
	galactose metabolic process	•	•	
	gametophyte development			
	generation of precursor metabolites and energy			
	glycerolipid metabolic process	•		
	glycogen metabolic process			
	glycosaminoglycan catabolic process	•		
	glycosyl compound biosynthetic process			
	glycosyl compound metabolic process	•		
	hemicellulose metabolic process	•		
	histidine metabolic process	•		
	homogalacturonan biosynthetic process			
	hormone-mediated signaling pathway		•	
	hydrogen ion transmembrane transport			
	import across plasma membrane			
	indole-containing compound metabolic process	•		
	induced systemic resistance, jasmonic acid mediated signaling pathway			
	intracellular signal transduction			
	isoleucine catabolic process			
	jasmonic acid and ethylene-dependent systemic resistance			
	leaf development	•		
	leucine catabolic process	•		
	lignin biosynthetic process			
	lignin catabolic process			
	limonene catabolic process	•		
	lipid biosynthetic process	•		
	microtubule-based process	•		
	mitochondrial fission	•	•	
	monocarboxylic acid biosynthetic process	•		
	mRNA processing	•		
	negative regulation of ethylene-activated signaling pathway	•		
	negative regulation of seed germination	•		
	nitrogen compound transport		•	
	nucleobase metabolic process			
	nucleotide-sugar biosynthetic process			
	one-carbon metabolic process			
	peptidyl-histidine phosphorylation			
	peptidyl-serine phosphorylation	•		
	pigment catabolic process	Ĭ		
	plant organ formation			
	plant-type primary cell wall biogenesis			
	plant-type primary cell wall biogenesis plant-type secondary cell wall biogenesis			
			l .	
	polarity specification of adaxial/abaxial axis			
	pollen development	•		
	pollination	•		
	porphyrin-containing compound catabolic process	•		
	positive gravitropism		•	
	positive regulation of developmental process	•		
	positive regulation of reproductive process	•		
	post-embryonic plant morphogenesis	•		
	potassium ion transmembrane transport			
	primary shoot apical meristem specification			
	protein autophosphorylation		• .	
	protein catabolic process			
	purine nucleoside metabolic process			
	purine ribonucleotide biosynthetic process			
	purine ribonucleotide biosynthetic process purine ribonucleotide metabolic process			
	partine injuriacientine metaponic process			
	regulation of root development			

	GO description	Enrich	ment
	regulation of stomatal movement		•
	response to cadmium ion		•
	response to cold		•
	response to fungus response to heat		
	response to hydrogen peroxide		
	response to light stimulus		
	response to lipid		•
	response to organic cyclic compound		
	response to red or far red light		1
	response to salt stress	•	•
	response to sucrose response to water deprivation		
	response to wounding		Ĭ
	ribonucleoside metabolic process		
	ribonucleoside monophosphate metabolic process		
	ribonucleoside triphosphate metabolic process		
	root hair cell differentiation		
	root hair elongation		
	S-adenosylmethionine biosynthetic process serine family amino acid metabolic process		
	starch metabolic process	ă	•
	sterol biosynthetic process		
	sucrose metabolic process		•
	sugar mediated signaling pathway	•	
	sulfur compound metabolic process		•
	trehalose metabolism in response to stress		
	UDP-glucose metabolic process unidimensional cell growth		
	vacuolar acidification		
	vacuolar protein processing	•	
	valine catabolic process	•	
	xylem and phloem pattern formation	•	
CELLULAR_COMPONENT	apoplast		•
	cellulose synthase complex	•	
	chloroplast part		:
	cytosol endocytic vesicle		•
	endocytic vesicie endomembrane system		
	Golgi membrane		
	integral component of plasma membrane		
	kinesin complex	•	
	microtubule	•	
	nucleolus		
	plant-type cell wall	•	•
	plant-type vacuole		•
	plasma membrane part plasmodesma		
	protein histidine kinase complex		
	protein phosphatase type 1 complex		
	tubulin complex	•	
	vacuolar membrane		•
	vacuolar proton-transporting V-type ATPase, V0 domain	٠	
MOLECULAR_FUNCTION	alpha,alpha-trehalose-phosphate synthase (UDP-forming) activity		
	amino acid transmembrane transporter activity  ATP binding	ě	
	ATPase activity, coupled to transmembrane movement of ions, phosphorylative mechanism		•
	ATPase coupled ion transmembrane transporter activity		
	auxin efflux transmembrane transporter activity	•	
	auxin influx transmembrane transporter activity	•	
	calcium ion binding		
	calcium-dependent protein serine/threonine kinase activity	•	
	calcium-dependent protein serine/threonine kinase activity calmodulin binding	•	•
	calcium-dependent protein serine/threonine kinase activity calmodulin binding calmodulin-dependent protein kinase activity	•	
	calcium-dependent protein serine/threonine kinase activity calmodulin binding	•	
	calcium-dependent protein serine/threonine kinase activity calmodulin binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity	• • • • • • • • • • • • • • • • • • • •	
	calcium-dependent protein serine/threonine kinase activity calmodulin-dipendent protein kinase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity	• • • • • • • • • • • • • • • • • • • •	
	calcium-dependent protein serine/threonine kinase activity calmodulin-dipendent protein kinase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene binding ethylene binding	•	•
	calcium-dependent protein serine/threonine kinase activity calmodulin-binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding	•	
	calcium-dependent protein serine/threonine kinase activity calmodulin-binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylar glucuronosyltransferase activity	•	
	calcium-dependent protein serine/threonine kinase activity calmodulin-dependent protein kinase activity cation-transporting ATPase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding	•	•
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	calcium-dependent protein serine/threonine kinase activity calmodulin-binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTPase activity hydrolase activity, hydrolyzing O-glycosyl compounds	•	
	calcium-dependent protein serine/threonine kinase activity calmodulin-dependent protein kinase activity cation-transporting ATPase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTPase activity	•	
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	calcium-dependent protein serine/threonine kinase activity calmodulin-dependent protein kinase activity cation-transporting ATPase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTP binding GTP activity fydrolase activity, hydrolyzing O-glycosyl compounds hydroquinonexoxygen oxidoreductase activity ligase activity, forming carbon-sulfur bonds	•	•
	calcium-dependent protein serine/threonine kinase activity calmodulin-dependent protein kinase activity cation-transporting ATPase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTP binding GTPase activity hydrolase activity, hydrolyzing O-glycosyl compounds hydroquinonexygen oxidoreductase activity ligase activity, forming carbon-sulfur bonds metal lon transmembrane transporter activity methionine adenosyltransferase activity microtubule binding	•	•
	calcium-dependent protein serine/threonine kinase activity calmodulin-binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTPase activity hydrolase activity, hydrolyzing O-glycosyl compounds hydroquinonexoxygen oxidoreductase activity ligase activity, forming carbon-sulfur bonds metal ion transmembrane transporter activity methionine adenosyltransferase activity microtubule binding monovalent inorganic cation transmembrane transporter activity	•	•
	calcium-dependent protein serine/threonine kinase activity calmodulin-binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTPase activity Hydrolase activity, hydrolyzing O-glycosyl compounds hydroquinone:oxygen oxidoreductase activity ligase activity, forming carbon-sulfur bonds metal ion transmembrane transporter activity microtubule binding monovalent inorganic cation transmembrane transporter activity N-acylphosphatidylethanolamine-specific phospholipase D activity		•
	calcium-dependent protein serine/threonine kinase activity calimodulin-binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTPase activity hydrolase activity, hydrolyzing O-glycosyl compounds hydroquinone:oxygen oxidoreductase activity ligase activity, forming carbon-sulfur bonds metal ion transmembrane transporter activity methionine adenosyltransferase activity microtubule binding monovalent inorganic cation transmembrane transporter activity NaCylphosphatidylethanolamine-specific phospholipase D activity NAO binding		•
	calcium-dependent protein serine/threonine kinase activity calmodulin-binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTPase activity hydrolase activity, hydrolyzing O-glycosyl compounds hydroquinonexoxygen oxidoreductase activity ligase activity, rorming carbon-sulfur bonds metal ion transmembrane transporter activity methionine adenosyltransferase activity microtubule binding monovalent inorganic cation transmembrane transporter activity N-acylphosphatidylethanolamine-specific phospholipase D activity NAD binding oligospetide transmembrane transporter activity		•
	calcium-dependent protein serine/threonine kinase activity calimodulin-binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTPase activity hydrolase activity, hydrolyzing O-glycosyl compounds hydroquinone:oxygen oxidoreductase activity ligase activity, forming carbon-sulfur bonds metal ion transmembrane transporter activity methionine adenosyltransferase activity microtubule binding monovalent inorganic cation transmembrane transporter activity NaCylphosphatidylethanolamine-specific phospholipase D activity NAO binding		•
	calcium-dependent protein serine/threonine kinase activity calmodulin-dependent protein kinase activity cation-transporting ATPase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTPase activity hydrolase activity, hydrolyzing O-glycosyl compounds hydroquinonexoxyen oxidoreductase activity ligase activity, forming carbon-sulfur bonds metal ion transmembrane transporter activity microtubule binding monovalent inorganic cation transmembrane transporter activity NAO binding oligopeptide transmembrane transporter activity NAO binding oligopeptide transmembrane transporter activity oligopeptide transmembrane transporter activity oligopeptide transmembrane transporter activity oligopeptide transmembrane transporter activity		•
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	calcium-dependent protein serine/threonine kinase activity calmodulin-binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTPase activity hydrolase activity, hydrolyzing O-glycosyl compounds hydroquinone:oxygen oxidoreductase activity hydrolase activity, forming carbon-sulfur bonds metal ion transmembrane transporter activity methionine adenoxyltransferase activity methionine adenoxyltransferase activity microtubule binding monovalent inorganic cation transmembrane transporter activity NAO binding iliqopeptide transmembrane transporter activity NAO binding iliqopeptide transmembrane transporter activity voxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxyger oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxyger oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxyger oxidoreductase activity, acting on the aldehydo or oxo group of donors, NAD or NADP as acceptor oxidoreductase activity, oxidizing metal ions		•
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	calcium-dependent protein serine/threonine kinase activity calmodulin-dependent protein kinase activity cation-transporting ATPase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding glucuronoxylan glucuronosyltransferase activity Hydrolase activity, hydrolyzing O-glycosyl compounds hydroquinonexoxyen oxidoreductase activity ligase activity, forming carbon-sulfur bonds metal ion transmembrane transporter activity microtubule binding monovalent inorganic cation transmembrane transporter activity NAD binding dispopetide transmembrane transporter activity NAD binding oligopeptide transmembrane transporter activity oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxyger oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxyger oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxyger oxidoreductase activity, acting on the aldehyde or oxo group of donors, NAD or NADP as acceptor oxidoreductase activity, oxidizing metal ions oxygen binding phospholpase D activity phosphorelay sensor kinase activity		
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	calcium-dependent protein serine/threonine kinase activity calmodulin-binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTPase activity hydrolase activity, hydrolyzing O-glycosyl compounds hydroquinone.oxygen oxidoreductase activity ligase activity, forming carbon-sulfur bonds metal ion transmembrane transporter activity methionine adenosyltransferase activity microtubule binding monovalent inorganic cation transmembrane transporter activity N-acylphosphatidylethanolamine-specific phospholipase D activity NAD binding oligopeptide transmembrane transporter activity oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxygen oxidoreductase activity, acting on the aldehyde or oxo group of donors, NAD or NADP as acceptor oxidoreductase activity, oxidizing metal ions oxygen binding phospholipase D activity phosphorelay sensor kinase activity potassium channel activity RNA polymerase II regulatory region sequence-specific DNA binding secondary active transmembrane transporter activity		
	calcium-dependent protein serine/threonine kinase activity calmodulin-binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTPase activity hydrolase activity, hydrolyzing O-glycosyl compounds hydroquinone:oxygen oxidoreductase activity hydrolase activity, forming carbon-sulfur bonds metal ion transmembrane transporter activity methionine adenosyltransferase activity microtubule binding monovalent inorganic cation transmembrane transporter activity N-acylphosphatidylethanolamine-specific phospholipase D activity NAD binding oliqopeptide transmembrane transporter activity oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxyger oxidoreductase activity, oxidizing metal ions oxygen binding phospholipase D activity phosphorelay sensor kinase activity phosphorelay sensor kinase activity phosphorelay sensor kinase activity protassium channel activity RNA polymerase il regulatory region sequence-specific DNA binding		
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	calcium-dependent protein serine/threonine kinase activity calmodulin-binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTPase activity hydrolase activity, hydrolyzing O-glycosyl compounds hydroquinone:oxygen oxidoreductase activity hydrolase activity, forming carbon-sulfur bonds metal ion transmembrane transporter activity methionine adenosyltransferase activity microtubule binding monovalent inorganic cation transmembrane transporter activity N-acylphosphatidylethanolamine-specific phospholipase D activity NAD binding oliqopeptide transmembrane transporter activity oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxyger oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxyger oxidoreductase activity, acting on the aldehyde or ox ox group of donors, NAD or NADP as acceptor oxidoreductase activity, oxidizing metal ions oxygen binding phospholipase D activity phosphorelay sensor kinase activity phosphorelay sensor kinase activity NAP polymerase il regulatory region sequence-specific DNA binding secondary active transmembrane transporter activity		
	calcium-dependent protein serine/threonine kinase activity calmodulin-binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding glucuronoxylan glucuronosyltransferase activity GTP binding GTPase activity, hydrolyzing O-glycosyl compounds hydroquinone:oxygen oxidoreductase activity ligiase activity, forming carbon-sulfur bonds metal ion transmembrane transporter activity methionine adenosyltransferase activity microtubule binding monovalent inorganic cation transmembrane transporter activity N-acylphosphatidylethanolamine-specific phospholipase D activity NAD binding oliqopeptide transmembrane transporter activity oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxyger oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxyger oxidoreductase activity, oxidizing metal ions oxygen binding hospholipase D activity phosphorelay sensor kinase activity phosphorelay sensor kinase activity NAP polymerase il regulator yegion sequence-specific DNA binding secondary active transmembrane transporter activity structural constituent of cytoskeleton symporter activity transferase activity, transferring hexosyl groups transmembrane receptor activity trehalose-phosphatase activity trehalose-phosphatase activity		
	calcium-dependent protein serine/threonine kinase activity calmodulin-binding calmodulin-dependent protein kinase activity cation-transporting ATPase activity cellulose synthase (UDP-forming) activity copper ion binding ethylene binding ethylene receptor activity FAD binding glucuronoxylan glucuronosyltransferase activity GTP binding GTPsase activity, hydrolyzing O-glycosyl compounds hydroquinoneoxygen oxidoreductase activity ligase activity, forming carbon-sulfur bonds metal ion transmembrane transporter activity microtubule binding monovalent inorganic cation transmembrane transporter activity NAD binding oliopopetide transmembrane bransporter activity NAD binding oliopopetide transmembrane transporter activity vaidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxyger oxidoreductase activity, acting on paired donors, with incorporation or molecular oxyger oxidoreductase activity, acting on the aldehyde or oxo group of donors, NAD or NADP as acceptor oxidoreductase activity, acting on paired donors, with incorporation or paired on the oxygen binding phospholpase D activity phosphorelay sensor kinase activity phosphorelay sensor kinase activity RNA polymerase II regulatory region sequence-specific DNA binding secondary active transmembrane transporter activity structural constituent of cytoskeleton symporter activity, transferring hexosyl groups transferase activity, transferring hexosyl groups transferase activity, transferring hexosyl groups		



	GO Description
Biological process	single-organism carbohydrate metabolism
	cell surface receptor signaling pathway
	plant organ development
	protein phosphorylation
	ion transmembrane transport
	organic substance catabolism
	cell wall organization
	nucleotide-sugar metabolism
	organic substance metabolism carbohydrate derivative metabolism
	stomatal movement
	stomatal movement single-organism process
	single-organism process single organism reproductive process
	single organism reproductive process
	response to stimulus
	reproduction
	nitrogen compound metabolism
	multicellular organismal process
	metabolism
	localization
	growth
	generation of precursor metabolites and energy
	developmental process
	cofactor metabolism
	coenzyme metabolism
	cellular process
	cellular component organization or biogenesis
	cell wall organization or biogenesis
	cell communication
	catabolism
	carbohydrate metabolism
Cellular component	cytosol
	external encapsulating structure
	bounding membrane of organelle
	membrane part
	endomembrane system
	catalytic complex
	whole membrane
	symplast
	side of membrane
	pollen tube membrane
	extracellular region
	cell-cell junction cell junction
	cell
	apoplast
Molecular function	phosphotransferase activity, alcohol group as acceptor
morecalar rarrectors	magnesium ion binding
	protein binding
	hydrolase activity, hydrolyzing 0-glycosyl compounds
	ion transmembrane transporter activity
	signaling receptor activity
	oxidoreductase activity, oxidizing metal ions
	ligase activity, forming carbon-sulfur bonds
	carbon-oxygen lyase activity, acting on polysaccharides
	transporter activity
	transferase activity
	structural constituent of cytoskeleton
	signal transducer activity
	oxidoreductase activity
	molecular transducer activity
	lyase activity
	lipid binding
	hydrolase activity
	coenzyme binding
	catalytic activity
	hinding

GO Category	GO Description	
Biological process	regulation of gene expression	•
	regulation of macromolecule metabolic process	•
	regulation of cellular macromolecule biosynthetic process	•
	regulation of macromolecule biosynthetic process	•
	regulation of primary metabolic process	•
	regulation of nitrogen compound metabolic process	•
	transcription, DNA-templated	•
	regulation of cellular metabolic process	•
	regulation of transcription, DNA-templated	•
	regulation of cellular biosynthetic process	•
	RNA biosynthetic process	•
	nucleic acid-templated transcription	•
	regulation of RNA biosynthetic process	•
	regulation of nucleic acid-templated transcription	•
	regulation of biosynthetic process	•
	regulation of metabolic process	•
	regulation of RNA metabolic process	•
	regulation of nucleobase-containing compound metabolic process	
	nucleic acid metabolic process	•
Cellular component	transcription factor complex	•
Molecular function	transcription factor activity, sequence-specific DNA binding	
	nucleic acid binding transcription factor activity	•

	Tier 3	SND1	NST1	NST2	VND6	VND7	
	Tier 2	SND3	MYB46	MYB61	MYB83	MYB103	
			MYB4	SND2	KNAT7	MYB20	
			MYB7	MYB52	MYB75	MYB58	
			MYB32	MYB54		MYB63	
				C3H14		MYB69	
	Tier 1					MYB79	
	Ξe					MYB85	
						ATHB18	
						At3g46080	
		Cellulose:	Hemicellulose:	Programmed cell death:	Signalling:	Lignin:	Other:
		CesA4	FRA8	XCP1	ROP1/2/4	PAL1	FLA11
		CesA7	IRX8	XCP2	CTL2	PAL4	FLA12
		CesA8	IRX9	MC8	LRR kinases	C4H	FLA13
		COBL4	IRX9-L	MC9	Rac GTPase	4CL1	FLA15
			IRX10	RNS3	Rho GTPase	4CL3	AGP4
	S		IRX14	BFN1	IQD10	HCT	AGP21
	au.		IRX14-L		RIC2	C3H1	Xyn3
	Structural genes		IRX15			CCoAOMT1	RIP4
	ra		IRX15-L			CCR1	ARAC2
	ਬੁੱ		CsIA9			F5H1	BGAL3
	ŝtr		CsIB1/2			COMT	TBL3
	0,		GXM1/2/3			CAD4/6	GLP10
			GUX1/2			CHS	CCoAOMT7
			PGSIP1			LAC4	
			PGSIP3			LAC10	
			UXS3			LAC17	
			PME				
			XTH10				
			PARVUS				