0 1 2 3 4 5 5 4 3 2 1 0 nextive regulation of mitotic		-log ₁	₀ <i>P</i> -va	lue	(8	a)			$-\log_{10} P$	-value	(b)	
DNA metabolic process SCC-depicator protoacomanti. mitotic sister chromatid segregation mitotic sister chromatid segregation metal ion transport phospholipid dephosphorylation metal ion transport proton transport mitotic metal ion metabolic process regulation of 5.85 rRNA from splicessomal if "arRAP. complex positive regulation of 5.85 rRNA from splicessomal if "arRAP. complex peophera bolic ion metal ion transport phosphatidylinositol-phosphate regulation of moved proton-transport consolid stack proton-transport such rotabolic process microtubule-based movement regulation of regulation of proton- regulation of splices- go consendiation complex cytosolic large rhoomal submit symptomenal complex cytosolic large rhoomal submit symptomenal submit APC-CG20 complex activity APC-CG20 complex activity alterion-streame regulation binding acyl-large-carrier protoin]. RNA polymerase II transcription phosphotic exter hydrolae activity withing e-gated chloride channel phosphotic exter hydrolae activity APC-CG20 complex activity withing e-gated chloride channel phosphotic exter hydrolae activity withing acgue activity APC-binding binding acyl-large-carrier protoin]. RNA polymerase II transcription phosphotic exter hydrolae activity withing e-gated chloride channel phosphotic exter hydrolae activity withing acgue activity APC binding binding acyl-large-carrier protoin]. RNA polymerase II transcription phosphotic exter hydrolae activity withing acgue activity APD binding binding acyl-large-carrier protoin]. RNA polymerase II transcription phosphotic exter hydrolae activity withing activity binositic transport activity binositic transport activity chinase acti		0 1	2	3	4	5	5	4	3	2	1	0
 cellular transition netal on attrainance of S.S. rRNA from phospholipid deptosphorylation prospholipid deptosphorylation actin filament deploymerization metal ion transport phosphatidid optionsito 3-phosphata. phosphatidid secondplat cellular macromolecule biosynthetic regulation of infloxes-ence meristem netalone complex regulation of complex condition metal ion binding defining definin	positive regulation of mitotic.											dhurrin biosynthetic process
mitotic sister chromatid segregation phospholigid dephosphotylation regulation of segressing of zinc in metal ion transport phosphatidylinositoltyhosphata. response to zinc ion procein ubiquitation involved in. response to zinc ion procein ubiquitation involved in. regulation of inforescence meristem celtara macromolecule biosynthetic regulation of inforescence meristem celtara macromolecule biosynthetic regulation of response to another organism maturation of LSU-RNA from celtara macromolecule biosynthetic regulation of inforescence meristem celtara macromolecule biosynthetic regulation of inforescence meristem celtara macromolecule biosynthetic regulation of response to another organism maturation of LSU-RNA from actin cytoskeleton smalt ribosome biogenesis actin cytoskeleton incohondria respiratory chain phosphore respiratory chain phosphore respiratory chain phosphore biogenes activity APC-Cd.20 comptex activity actin cytoskeleton incohondria respiratory chain phosphore set biogenes activity woltage-gated ebloride channel ADP binding acyltransferase activity voltage-gated ebloride channel cytoshore boghastas activity ADP binding acyltransferase activity woltage-gated ebloride channel cytoshore boghastas activity Protein kinase inhibitor activity childra boghastas activity childra boghastas activity childra cativity childra c	SCF-dependent proteasomal.											DNA metabolic process
phosphalipid dephospharylation splicecound trisnRNP complex regulation of zine ion positive regulation of miRNA metal ion transport mitochondrail deciron transport protein ubiquitination involved in regulation of regions to ion robosphality linssiol-3-phosphate. regulation of complex robosphality linssiol-3-phosphate. regulation of regions to ion nuclear SCF ubiquitin ligas complex regulation of response to ion contrasporting two-sector regulation of response to ion contrasporting two-sector regulation of freeponse to cosmotic stress defense response to inflorescence meristem regulation of foral meristem growth synaptonenal complex regulation of foral meristem growth synaptonenal complex regulation of foral meristem growth phosphoric seter hydrolase activity regulation of foral meristem growth phosphoride regulation binding regulation of foral meristem growth <td>cellular transition metal ion.</td> <td></td> <td>chromosome organization</td>	cellular transition metal ion.											chromosome organization
regulation of sequestering of zinc ion actin filament deolymerization medal ion transport phosphatidylinositol-3-phosphate. Proton-transport of zinc ion protein ubiquitination involved in. Proton-transport explaision of inflorescence meristem eclular macmolecule biosynthetic regulation of inflorescence meristem chiin catabolic process microthubi-based movement regulation of LSU-rRNA from actin cytoskeleton small ribosomal subuit phosphoric ester bydrolase activity ubiquitin conjugating enzyme activity biquitin conjugating enzyme activity agalaction-succes. Franking actin cytoskeleton sayle phosphoric ester bydrolase activity biquitin conjugating enzyme activity adaption-succes. Franking actin cytoskeleton sayle phosphoric ester bydrolase activity biquitin conjugating enzyme activity adaption-succes. Franking actin cytoskeleton actin cytoskeleton sayle phosphoric ester bydrolase activity biquitin conjugating enzyme activity adaption-succes. Franking actin cytoskeleton activity ADP binding avyl-[acyl-leartie-protein], ADP binding biosphatidae phosphatiase activity voltage-gated chloride channel. Franking biosphatidae phosphatiase activity protein himpophates activity protein kinase entivity voltage-gated chloride channel. Franking biosphatidae phosphatidae channel. Franking biosphatidae phosphatidae channel. Franking biosphatidae phosphatidae activity protein kinase entivity franking biosphatidae phosphate channel. Franking biosphatidae phosphate channel. Franking biosphatidae phosphate channel. Franking biosphatidae phosphate channel. Franking biosphatidae phosphate channel. Franking biosphate channel. Franking biosphate channel. Franking biosphate channel. Franking biosphate channel. Franking biosphate channel. Franking frank	mitotic sister chromatid segregation											maturation of 5.8S rRNA from
actin filament depolymerization metal ion transport phosphaticylinosiol-3-phosphate. response to zine ion protein ubiquitination involved in regulation of cyclins regulation of cyclins regulation of inflorescence meristem regulation of inflorescence meristem regulation of response to cosmotic stress Golgi stack proton-transporting two-sector core mediator complex cytosolic large ribosomal subunit synaptonemic complex synaptonemic complex issuing enzyme activity ubiquitin conjugating enzyme activity ubiquitin conjugating enzyme activity phosphoric ester hydrolase activity phosphoriage parted ebloride channel ADP binding acyt-[acyl-carrier-protein] Phosphatakae activity phosphatakae activity phosphatak	phospholipid dephosphorylation											spliceosomal tri-snRNP complex
metal ion transport circumitation phosphrid/jtionsitol-3-phosphate circumitation response to zinc ion mito-hondrial dector narmport protein ubiquitination involved in regulation of nortex-serve meristerm rubosome biogenesis collater accomplex mackear SCF ubiquitin ligase complex collater activity Colgi stack core mediator complex core mediator complex core mediator of seponse to osmotic stress core mediator complex actio cytoskeleton synaptonemal complex actio cytoskeleton actin cytoskeleton metalion of floar mercisterm growth phosphorie set hydrolase activity metalion of floar mercisterm growth ubiquitin conjugating enzyme activity mito-chondrial cector transport chain galaction-3uccreate. posphorie set hydrolase activity ubiquitin conjugating enzyme activity mito-chondrial cector transport chain galaction-3uccreate. posphorie set hydrolase activity APC-Cd20 complex activity mito-chondrial cector transport chain galaction-3uccreate. posphorie set hydrolase activity phosphatidae phosphates activity galaction-3uccreate. galactino-3uccreate. <td< td=""><td>regulation of sequestering of zinc ion</td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td>positive regulation of miRNA</td></td<>	regulation of sequestering of zinc ion											positive regulation of miRNA
phosphridylinositol-3-phosphate response to zino ion protein ubiquitiation involved in ribosome biogenesis Golgi stack Golgi stack Goldia Gol												circumnutation
picesphatidylinositol ->phosphate negative regulation of cyclin response to zhn cion negative regulation of cyclin protein ubiquifinition involved in regulation of findroscence meristem nuclear SCF ubiquifin ligase complex negative regulation of molorscence meristem Golgi stack netroubule-based movement regulation of norscence to commonic stress defense response to other organism actin cytoskeleton maturation of LSU-rRNA from astin cytoskeleton aerobic electron transport chain phosphoric ester hydrolase activity nuclear SCF ubiquifin ligging enzyme activity APC-Cdc20 complex activity nuclear stress gluestinol-sucrose poosphoric system respiratory chain complex set of find meristem growth proton-transport chain regulation of floral meristem growth proton-transport chain regulation of molar meristem growth proton-transport chain regulation system respiratory chain complex aerobic electron transport chain gluesorythosphatic poosphorelay signal transduction system respiratory chain respiratory chain complex IV ADP binding gluesorythosphate/lightsphate/lightsphate <td>-</td> <td></td> <td>mitochondrial electron transport,</td>	-											mitochondrial electron transport,
response to Zhe 100 cellular macromolecule biosynthetic protein ubiquitinitoni involved in regulation of inflorescence meristem nuclear SCF ubiquitin Tigse complex chifin catabolic process Core mediator complex cellular macromolecule biosynthetic core mediator complex cellular macromolecule biosynthetic cytosolic large ribosomal subunit cellular macromolecule biosynthetic synaptonemal complex cellular macromolecule biosynthetic actin cytoskeleton microtubale-based movement regulation of floral meristem growth regulation of floral meristem growth protein ubiquitine conjugatine enzyme activity aerobic electron transport chain actin cytoskeleton nucleous hinding galactionol-sucrose transition metal ion binding galactionol-sucrose acyl-[cyt-[cyt-[cyt-carrier-protein] protein bioding givesynthesphatidylinositol-N previbosome, large subunit precursor N-RNA polymerase II transcription previbosome, large subunit precursor N-acyltransferase activity 4hydroxyphenylacetaldehyde oxime ribonucleoprotein complex binding oxidoredutase activity voltage-gated chloride channel <t< td=""><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></t<>												
regulation inforcements medear SCF ubiquitin ligase complex Golgi stack proton-transporting two-sector core mediator complex actin cytoskeleton small rbosomal subunit phosphoric ester hydrolase activity ubiquitin conjugating enzyme activity ubiquitin conjugating enzyme activity actin cytoskeleton small rbosomal subunit phosphoric ester hydrolase activity ubiquitin conjugating enzyme activity actin cytoskeleton small rbosomal subunit phosphoric ester hydrolase activity ubiquitin conjugating enzyme activity actin cytoskeleton small rbosomal subunit phosphorica ester hydrolase activity ADP binding binding acyl-[acyl-carrier-protein] RNA polymerase II transcription phosphatidate phosphatese activity voltage-gated chloride channel binding actioredinates activity voltage-gated chloride channel binding actioredinates activity cytochrome-confake activity protein kinase activity RNA-directed DNA polymerase activity inositol ternkisphosphate J-kinase inositol ternkisphosphate J-kinase inositol 1,3,4-trisphosphate J-kinase inositol 1,3,4-trisphosphate J-kinase inositol ternkisphosphate J-kinase	-											
nuclear SCF ubiquiin ligas complex Golgi stack proton-transporting two-sector cytosofic large ribosomal subunit synaptonemal complex a chin cytoskeleton small ribosomal subunit phosphoric setter hydrolaes activity ubiquitin conjugating enzyme activity a APC-Cdc-20 complex activity a galactionl-sucrese transition metal ion binding binding a ceyl-[aeyl-carrier-protin] Phosphorical transcription RNA polymerase II transcription phosphoria excitivity voltage-gated chloride channel exploratione activity voltage-gated chloride channel exploratione activity rotoria metal ion binding a ceyl-[aeyl-carrier-protin] phosphorie complex activity woltage-gated chloride channel exploratione activity rotoria metal ion binding a ceyl-[aeyl-carrier-protin] phosphorie activity woltage-gated chloride channel exploratione activity rotoria metal ion binding a ceyl-[aeyl-carrier-protin] phosphatiate phosphatase activity woltage-gated chloride channel exploratione activity rotoria metal ion frage activity woltage-gated chloride channel exploratione activity protein kinase inhibitor activity chitimase activity inositol-1,3,4-trisphosphate 1-kinase inositol-1,3,4-trisphosphate 1-kinase inositol-1,3,4-trisphosphate 2-kinase activity chitimase activity inositol-1,3,4-trisphosphate 1-kinase inositol-1,3,4-trisphosphate 2-kinase activity chitimase activity inositol-1,3,4-trisphosphate 2-kinase activity chitimase activity inositol-1,3,4-trisphosphate 2-kinase activity inositol-1,3,4-trisphosphate 2-kinase activity inositol-1,3,4-trisphosphate 3-kinase activity inositol-1,3,4-trisphosphate 3-kinase activity inositol-1,3,4-trisphosphate 3-kinase activity inos												
Golgi stack microtubule-based movement proton-transporting two-sector microtubule-based movement core mediator complex maturation of LSU-rRNA from actin cytoskeleton actin cytoskeleton small ribosomal subunit aerobic electron transport chain phosphoric ester hydrolase activity aerobic electron transport chain actin cytoskeleton microtubule-based movement small ribosomal subunit aerobic electron transport chain phosphoric ester hydrolase activity aerobic electron transport chain actin cytoskeleton microtubule-based movement APC-Cdc20 complex activity aerobic electron transport chain galactinol-sucrose. microtubule-based movement ADP binding gycosylphosphatidylinositol-N gycosylphosphatidylinositol-N pretinormal complex IV gycosylphosphatidylinositol-N pretinor	-											
proton-transporting two-sector regulation of response to osmotic stress core mediator complex defense response to other organism actin cytoskelton actin cytoskelton small ribosomal subunit naturation of LSU-rRNA from phosphoric ester hydrolase activity naturation of foral meristem growth proton-function metal ion binding naturation of the protein inport into nucleus Leaf Vascular tissue pattern formation natice system respiratory chain phosphorie ster hydrolase activity galactinol-sucrose phosphorie ster hydrolase activity actin cytoskelton nitochondrial respiratory chain galactinol-sucrose phosphorelay signal transduction system respiratory chain phosphatidylinositol-N phosphatitate phosphatase activity glycosylphosphatidylinositol-N protosphatitate phosphatase activity storeductase activity voltage-gated chloride channel storeduc												
core mediator complex cytosolic large ribosomal subunit synaptonemal complex actin cytoskeleton small ribosomal subunit phosphoric ester hydrolase activity ubiquitin conjugating enzyme activity galactinol-success APC-Cdc20 complex activity galactinol-success ADP binding binding acyl-facyl-carrier-protein] RNA polymerase activity voltage-gated chloride channel explorability activity ubiquitin conjugatide phosphate activity insolute protein import into nucleus leaf vascular tissue pattern formation mitochondrial respiratory chain phosphorelay signal transduction system respiratory chain complex IV glycosylphosphatidylinositol-N preribosome, large subunit precursor N-acyltransferase activity voltage-gated chloride channel eytochrome-c oxidase activity inositol transiphosphate 1-kinase double-stranded RNA binding UTP-glucose-1-phosphate ADP binding	•											
cytosolic large ribosomal subunit synaptonemal complex actin cytoskeleton small ribosomal subunit phosphoric ester hydrolase activity ubiquitin conjugating enzyme activity alactinol-sucrose ADP binding acyl-[acyl-cardiereprotein] phosphorelay signal transduction system respiratory chain complex IV glacetinol-sucrose ADP binding acyl-[acyl-cardiereprotein] phosphorelay signal transduction system respiratory chain complex IV glycosylphosphatidylinositol-N preribosome, large subunit precursor Nacyltransferase activity voltage-gated chloride channel mitoschorter coxidase activity protein kinase inhibitor activity chitinase activity RNA_directed DNA polymerase activity inositol tertakisphosphate 1-kinase inositol-1,3,4-trisphosphate 5-kinase double-stranded RNA binding UTTP:glucose-1-phosphate ADP binding												regulation of response to osmotic stress
synaptonemal complex actin cytoskeleton small ribosomal subunit phosphoric ester hydrolase activity ubiquitin conjugating enzyme activity agateniol-sucrose APC-6d20 complex activity agateniol-sucrose https://dxi.org/lacyl.earlier.protein] ADP binding binding acyl-[acyl-carrier-protein] phosphorelasy signal transduction system respiratory chain complex IV glycosylphosphatidylinositol-N preribosome, large subunit precursor N-acyltransferase activity 4-hydroxyphenylacetaldehyde oxime ribonucleoprotein complex binding oxidoreductase activity voltage-gated chloride channel fiositol 1,3,4-trisphosphate 1-kinase inositol 1-1,3,4-trisphosphate 1-kinase inositol 1-1,3,4-trisphosphate 1-kinase inositol 1-1,3,4-trisphosphate 1-kinase inositol 1-1,3,4-trisphosphate 1-kinase inositol 1-1,3,4-trisphosphate 1-kinase inositol 1-1,3,4-trisphosphate 5-kinase double-stranded RNA binding UTP:glucose-1-phosphate ADP binding	*											defense response to other organism
actin cytoskeleton small ribosomal subunit phosphoric ester hydrolase activity ubiquitin conjugating enzyme activity galactinol-sucrose												maturation of LSU-rRNA from
small ribosomal subunit phosphoric ester hydrolase activity ubiquitin conjugating enzyme activity alactinol-sucrose ADP binding binding acyl-[acyl-carrie-protein] phosphatidate phosphatase activity voltage-gated chloride channel												aerobic electron transport chain
phosphoric ester hydrolase activity ubiquitin conjugating enzyme activity APC-Cdc20 complex activity galactinol-sucrose transition metal ion binding MADP binding ioning acyl-[acyl-carrier-protein] phosphatidate phosphatase activity woltage-gated chloride channel FINA polymerase II transcription phosphatidate phosphatase activity woltage-gated chloride channel FINA polymerase II transcription phosphatidate phosphatase activity woltage-gated chloride channel FINA polymerase II transcription FINA polymerase activity FINA polymerase activity FINA polymerase activity FINA polymerase II transcription FINA polymerase activity FINA polymerase ac												regulation of floral meristem growth
ubiquitin conjugating enzyme activity APC-Cdc20 complex activity galactinol-sucrose transition metal ion binding ADP binding binding acyl-[acyl-carrier-protein] RNA polymerase II transcription phosphatidate phosphatase activity voltage-gated chloride channel N-acyltransferase activity voltage-gated chloride channel												protein import into nucleus
APC-Cdc20 complex activity galactinol-sucrose https://www.complex.iv/ apple.pinding binding acyl-[acyl-carrier-protein] RNA polymerase II transcription phosphatidate phosphatase activity voltage-gated chloride channel												leaf vascular tissue pattern formation
transition metal ion binding ADP binding binding acyl-[acyl-carrier-protein] RNA polymerase II transcription phosphatidate phosphatase activity voltage-gated chloride channel cytochrome-c oxidase activity protein kinase inhibitor activity chitinase activity RNA-directed DNA polymerase activity inositol tetrakisphosphate 1-kinase inositol-1,3,4-trisphosphate 5-kinase double-stranded RNA binding UTP:glucose-1-phosphate ADP binding												mitochondrial respiratory chain
transition metal ion binding ADP binding binding acyl-[acyl-carrier-protein] RNA polymerase II transcription phosphatidate phosphatase activity voltage-gated chloride channel ribonucleoprotein complex binding oxidoreductase activity, acting on the cytochrome-c oxidase activity protein kinase inhibitor activity chitinase activity RNA-directed DNA polymerase activity inositol 1-1,3,4-trisphosphate 1-kinase inositol-1,3,4-trisphosphate 5-kinase double-stranded RNA binding UTP:glucose-1-phosphate ADP binding	galactinol-sucrose.											phosphorelay signal transduction system
ADP binding binding acyl-[acyl-carrier-protein] RNA polymerase II transcription phosphatidate phosphatase activity voltage-gated chloride channel voltage-gated chlorid	transition metal ion binding											
binding acyl-[acyl-carrier-protein] RNA polymerase II transcription phosphatidate phosphatase activity voltage-gated chloride channel ribonucleoprotein complex binding oxidoreductase activity, acting on the cytochrome-c oxidase activity protein kinase inhibitor activity protein kinase inhibitor activity chitinase activity RNA-directed DNA polymerase activity inositol tetrakisphosphate 1-kinase inositol-1,3,4-trisphosphate 5-kinase double-stranded RNA binding UTP:glucose-1-phosphate ADP binding	ADP binding											
acyl-[acyl-carrier-protein] N-acyltransferase activity RNA polymerase II transcription 4-hydroxyphenylacetaldehyde oxime phosphatidate phosphatase activity 4-hydroxyphenylacetaldehyde oxime voltage-gated chloride channel ribonucleoprotein complex binding oxidoreductase activity protein kinase inhibitor activity protein kinase inhibitor activity chitinase activity grotein kinase inhibitor activity chitinase activity inositol tetrakisphosphate 1-kinase inositol tetrakisphosphate 1-kinase double-stranded RNA binding UTP:glucose-1-phosphate ADP binding Disting	binding									-		
RNA polymerase II transcription 4-hydroxyphenylacetaldehyde oxime phosphatidate phosphatase activity ribonucleoprotein complex binding oxidoreductase activity, acting on the cytochrome-c oxidase activity rotein kinase inhibitor activity protein kinase inhibitor activity chitinase activity protein kinase inhibitor activity chitinase activity nositol tetrakisphosphate 1-kinase inositol tetrakisphosphate 5-kinase double-stranded RNA binding UTP:glucose-1-phosphate ADP binding	acyl-[acyl-carrier-protein].											
ribonucleoprotein complex binding oxidoreductase activity, acting on the cytochrome-c oxidase activity protein kinase inhibitor activity chitinase activity RNA-directed DNA polymerase activity inositol tetrakisphosphate 1-kinase inositol-1,3,4-trisphosphate 5-kinase double-stranded RNA binding UTP:glucose-1-phosphate ADP binding	RNA polymerase II transcription.											
oxidoreductase activity, acting on the cytochrome-c oxidase activity protein kinase inhibitor activity chitinase activity RNA-directed DNA polymerase activity inositol tetrakisphosphate 1-kinase inositol-1,3,4-trisphosphate 5-kinase double-stranded RNA binding UTP:glucose-1-phosphate ADP binding	phosphatidate phosphatase activity											
cytochrome-c oxidase activity protein kinase inhibitor activity chitinase activity RNA-directed DNA polymerase activity inositol tetrakisphosphate 1-kinase inositol-1,3,4-trisphosphate 5-kinase double-stranded RNA binding UTP:glucose-1-phosphate ADP binding	voltage-gated chloride channel.											
protein kinase inhibitor activity chitinase activity RNA-directed DNA polymerase activity inositol tetrakisphosphate 1-kinase inositol-1,3,4-trisphosphate 5-kinase double-stranded RNA binding UTP:glucose-1-phosphate ADP binding												
chitinase activity RNA-directed DNA polymerase activity inositol tetrakisphosphate 1-kinase inositol-1,3,4-trisphosphate 5-kinase double-stranded RNA binding UTP:glucose-1-phosphate ADP binding												cytochrome-c oxidase activity
RNA-directed DNA polymerase activity inositol tetrakisphosphate 1-kinase inositol-1,3,4-trisphosphate 5-kinase double-stranded RNA binding UTP:glucose-1-phosphate ADP binding										-		protein kinase inhibitor activity
inositol tetrakisphosphate 1-kinase inositol-1,3,4-trisphosphate 5-kinase double-stranded RNA binding UTP:glucose-1-phosphate ADP binding												chitinase activity
inositol-1,3,4-trisphosphate 5-kinase double-stranded RNA binding UTP:glucose-1-phosphate ADP binding										, i		RNA-directed DNA polymerase activity
double-stranded RNA binding UTP:glucose-1-phosphate ADP binding												inositol tetrakisphosphate 1-kinase
UTP:glucose-1-phosphate ADP binding												inositol-1,3,4-trisphosphate 5-kinase
ADP binding												double-stranded RNA binding
ADP binding												UTP:glucose-1-phosphate
												malate dehydrogenase