Treemap of significantly upregulated Gene Ontology terms in combined heat and drought

cellular response to unfolded protein	cellular response to heat		negative regulation of gibberellic acid mediated signaling pathway		ac res ^{ay} de	heat climat ponse siccat	ion e to ion	asparagine biosynthetic process -asparagine biosy	raffinose family nthessis	phosphatidylin b phosphatidylin b	
respons <mark>Çęljul</mark>	response to heat ar response	respo high se ^{inte}	nse to	o resp ded p	ponse to potelin	respor endop reticu stre	nse to lasmic ulum ess	cellular amino acio catabolic process		phosphatidylinosito	
herbivore	response to hydrogen	absei lic	nce of ght	hyperosm salinit <u>i</u> respons	to reticulum stress response to to to to to to to to to to	synthase					
cellular response to sucrose	peroxide cellular response	respo bacte resp	onse to terium ponse	respo suc	onse to rose	respor woun	nse to ding	transmembrane transporter activity	(glutamine-hy activi	amine–hydrolyzing) activity	
starvation	to hypoxia	to osmotic stress		respo abscis	onse to sic acid	response to salt stress		activity	linole	linoleate b	
induction of program induc cell death	regula ction of pr of p	tion ograr H	cellu potase n med ior homeos	ilar ^I sium si Cell stasis g	regulati reactive of pecies m death proce gene silo by R	ion of oxygen letabolic ess encing NA	DNA methylation	anaerobic respiration	13S-lipoxy linole 13S-lipoxy factiv 3'-monoox activ	vgenase /genase ityd f vgenase f	

plant-type cell wall loosening sitol-3-phosphate ding plant-type cell sitol-3-phosphate ding wall loosening modification chromatin modification involved in ol-3,5-bisphosphate abscission ding anchored component floral organ orphogenesis of plasma membrane anchored component polygalacturonase lipid activity of inding membrane oxygen binding endoplasmic rotein reticulum lumen olding wax biosynthesis