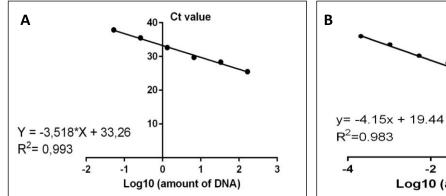
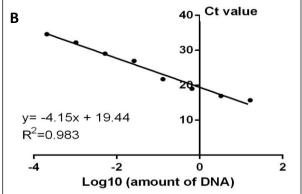
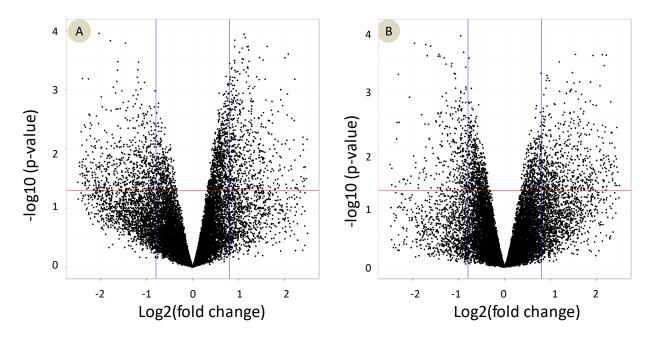
## Supplemental Figures





**Supplemental Figure 1.** Standard curves used for *Fragaria vesca* and *Botrytis cinerea* genomic DNA quantification. Curves were generated by amplifying the *EFTwo* (elongation factor 2-like) gene of *F. vesca* (**A**) and the *Bc3* (ribosomal IGS spacer) gene of *B. cinerea* (**B**), in a 5-fold serially diluted genomic DNA of *F. vesca* and *B. cinerea*, respectively. The standard curves show good linear relationship (R2 > 0.98) between the log5 value of the starting DNA concentration and the threshold cycle (Ct), both in the plant and in the fungus.



**Supplemental Figure 2.** Volcano plot of *Fragaria vesca* white fruit (**A**) and red fruit (**B**) gene expression values at 24 hpi. Genes in the left top rectangle [P. value > -log(10, 1.12) and fold change < log (2, |-0.80|) ] were selected as downregulated; whereas, genes in the right top rectangle were selected as upregulated [P. value > -log(10, 1.12) and fold change > log (2, 0.80)].

Gene II		N R	Description		Gene ID	W	R	Description
gene297	_		Glutamate receptor		gene 19766			Alcohol O-fatty-acyltransferase
gene173			Brefeldin A-inhibited GEP 2	wall	gene 24409			Callose synthase
gene193			Brefeldin A-inhibited GEP 2		gene 20547			Cellulose synthase Fold change
gene093			Calcium-binding protein	Cell	gene 24370			Postate lugge 1
gene17			Cysteine-rich RLK29		gene30789			Polygalacturonase (log2)
gene012 gene130 gene130			Disease resistance protein		gene02468	=		ABC transporter
gene130	57		Disease resistance protein		gene03853			ABC transporter
gene136	84		Disease resistance protein		gene 15834			ABC transporter
ø gene26€	50		Disease resistance protein					The state of the s
5 gene323	01		Disease resistance protein	l t	gene 16257			ABC transporter
gene323 gene343 gene249 gene240	24		Disease resistance protein	8	gene 30945			ABC transporter
gene249	62		Lectin-domain receptor kinase	Transport	gene01819			Aquaporin NIP5-1
gene240	85		MAP kinase	12	gene 19831			Oligopeptide transporter
gene230	86		PRA1 family protein		gene01912			Oligopeptide transporter
gene222	29		Strubbelig-receptor family 2		gene 30970			Potassium channel AKT1
gene030			TMV resistance protein N		gene05930			Protein translocase
gene165			TMV resistance protein N		gene03413			60S ribosomal export protein
gene319			TMV resistance protein N		gene06433			30S ribosomal protein
	_			=	gene08941			60S ribosome subunit biogenesis protein
gene193			AC9 transposase		-			
gene051		-	CAM transcription activator		gene 19176			Acetylglutamate kinase
gene133			Heat stress transcription factor B-2a		gene 29029			Adenylyl-sulfate kinase
gene082			Histone H1.2		gene01867	_		ADP,ATP carrier protein
gene191		1	Histone-lysine N-methyltransferase		gene06526			Ankyrin repeat-containing protein
gene175			RNA polymerase II transcription		gene 12868			Argininosuccinate lyase
gene132			Remorin		gene 34920			Aspartyl-tRNA synthetase
gene260			RPE-spondin		gene 10392			ATECP63
gene13: gene19: gene17: gene17: gene13: gene26: gene23: gene23: gene23: gene13:			S-acyltransferase		gene 27957			ATP-dependent RNA helicase
gene133	26		TraB domain-containing protein		gene32567			ATP-dependent RNA helicase
gene191	28		Trans. elongation factor B polypeptide3		gene 28599			B3 domain-containing protein
gene245	77		Transcription factor bHLH95		gene 23859			Coiled-coil domain-containing protein
gene250	55		Translation initiation factor IF-2		gene 20880			Cytokinin-N-glucosyltransferase
gene104			Two-component response regulator-like APRR2		gene00094			Cytokinin-O-glucosyltransferase
gene290			Acylamino-acid-releasing enzyme	=	_			
1000					gene31390			D-alanyl-D-alanine carboxypeptidase
gene236			Aspartyl aminopeptidase		gene 28423			Diphthamide biosynthesis protein
≌ gene305			Aspartyl aminopeptidase		gene 17572			Dolichol-phosphate mannosyltransferase
gene021			E3 ubiquitin-protein ligase MARCH6		gene 15570			F-box/FBD/LRR-repeat protein
gene305 gene021 gene169 gene292			Helicase		gene 28926			F-box/kelch-repeat protein
gene292			Protease Do-like 7		gene04407			F-box/LRR-repeat protein
gene296	71		PTS2 receptor		gene 24446			Formin-like protein
∝ gene119	69		Serine/threonine-protein kinase		gene 27685			Formin-like protein
gene218	93		Tim17		gene00561			Frizzled-3
gene074	98		Tim23		gene 14841			Gibberellin 3-beta-dioxygenase
	63		3-ketoacyl-CoA synthase		gene 15604			Glucose-6-phosphate 1-dehydrogenase
gene290 E gene188			Glycerol-3-phosphate acyltransferase 6	922	gene 24055			Glutamine amidotransferase
gene146			Palmitoyl-monogalactosyldiacylglycerol delta-7	J Miscellaneous			-	
gene146			desaturase	l Se	gene 10816			Golgi resident protein
gene043	1355			= ਵਿ	gene 17395			Histone deacetylase
			Anthocyanidin 5,3-O-glucosyltransferase	8	gene01761			Hypothetical protein
gene043			Anthocyanidin 5,3-O-glucosyltransferase	\second	gene03688			Hypothetical protein
gene281 gene213 gene233 gene268			Cationic peroxidase	-	gene 15082			Hypothetical protein
gene213		1111	Chal cone isomerase		gene 24705			Hypothetical protein
gene233			Chalcone isomerase 1		gene 25795			Hypothetical protein
gene268			Chalcone synthase		gene 12281			LIM domain-containing protein
≥ gene268			Chalcone synthase 2		gene 21051			Lysine-rich arabinogalactan protein
gene268 gene258 gene120 gene146			Flavonoid 3'-monooxygenase		gene01929			Lysine-specific demethylase
gene120	86		Laccase		gene00411			Modification methylase
gene14	11		Naringenin,2-oxoglutarate 3-dioxygenase					Monoglyceride lipase
gene097	53		Phenylalanine ammonia-lyase		gene 23304			
gene086			Vinorine synthase		gene 12793			Neuroblastoma-amplified sequence
gene065			Beta-galactosidase	i	gene31584			Oligogalacturonate-specific porin
gene060			Cryptochrome-1		gene 29012			Orotidine 5'-phosphate decarboxylase
gene008					gene 27497			PH and SEC7 domain-containing protein
1			Defensin-like protein 232		gene 14205			PHD and RING finger domain-containing protein
gene030			Heat shock protein		gene07883			Phospholipase C 4
gene091			Heat shock protein		gene 19207			Poly(A) RNA polymerase cid14
gene131		7/	Heat shock protein		gene 27828	_		Protein HIR1
gene154			Heat shock protein		gene 11038			Protein RIC1 homolog
gene208			Heat shock protein		gene07903			Pyridoxal biosynthesis protein
gene208	85		Heat shock protein					
gene322	58		Heat shock protein		gene 18342			Pyridoxal biosynthesis protein
gene154 gene208 gene208 gene322 gene070			Major allergen Pru ar 1		gene31671			Ribonuclease H protein
gene285			MLO-like protein 3		gene02490			Ribosomal protein L11 methyltransferase
gene043			Pentatricopeptide repeat-containing protein		gene01752			Sperm protamine-P1
gene091			Pentatricopeptide repeat-containing protein		gene02487			Subtilisin-like protease
gene136			Phenylalanyl-tRNA synthetase beta chain		gene 16495			Tektin-3
gene005		19	Polygalacturonase inhibitor		gene 18991			Transmembrane protein
geneous geneos					gene 26399			Transmembrane protein
geneus			Protease-activating factor 1 Syntaxin-24		gene 16934			U4/U6.U5 small nuclear ribonucleoprotein
gene188								

Supplemental Figure 3. Heat map showing the expression profile of the 149 common genes. Gene description and functional category, based on MapMan annotation, is provided. W, white fruit; R, red fruit. The scale bar displays fold change values (log2).