

Table 2 List of apoplastic proteins exclusively modulated in leaves of DR after inoculation with *T. deformans*.

Functional classification	Description	Acc N	Log2 FC	
			DR12-DR0	DR96-DR0
Cell wall	Alpha-L-fucosidase 1	M5X4F3	1.6	-1.6
	(1-4)-beta-mannan endohydrolase	M5WNY8		-2.1
	Endo-1,4-beta glucanase 3	M5W248		
	Glycosyl hydrolase family 3 protein	M5VW21		-1.9
	Pectinesterase family protein	M5VV51		6.8
	Pectinacetylerase	M5XMM4		5.2
	RD22; nutrient reservoir	M5XR24	4.1	-5.3
	Xyloglucan:xyloglucosyl transferase XTH33	M5XS44		
	Expansin-3	M5VZP0	1.3	
	Expansin B2	M5XKK6		7.1
Stress	PR2, Beta-1,3-endoglucanase	M5XWI7	1.6	
	PR2, Beta-1,3-endoglucanase	M5XD44	4.6	
	PR2, Beta-1,3-endoglucanase	M5VM18	-6.6	2.3
	PR-3, basic endochitinase	M5VQE4		1.5
	PR-3, Endochitinase 1	M5XMI9		-6.6
	PR4, Chitinase, ATEP3	M5XTV3		7.3
	PR5, Pathogenesis-related thaumatin family protein TLP1	M5WV03	-5.5	6.8
	PR5, Thaumatin-like protein 2	M5WA70		12.6
	PR9, Peroxidase, putative	M5W061		2.4
	PR9, Peroxidase, putative	M5WVGJ8		-7.1
	PR9, Peroxidase 12	M5W6X8		
	PR9, Peroxidase 27	M5XBP5		-2.5
	PR12, Knot1 domain-containing protein, defensin	M5Y0D4*		11.0

	PR14, Non-specific lipid-transfer protein 1-like	M5W0V2		10.1
	PR14, Lipid transfer protein 1	M5W0S9	9.0	
	PR14, Lipid transfer protein	M5W0V7		6.1
	PR14, Lipid Transfer protein3	M5WKL7		-1.6
	PR14, Lipid Transfer protein 2, putative	M5WW86*		9.2
	PR14, Lipid transfer protein 2 family	M5WX74		5.9
	PR14, Non specific lipid binding protein putative	M5Y0J5		-2.9
	PR14, Hydrophobic seed domain protein, belongs to the LTP2 family	M5WVP0	-11.7	1.6
	Disease resistance -responsive (dirigent-like protein) family protein	M5WZH0	5.6	
	PR17, Secretory protein, putative	M5VL90		5.6
	Trypsin and protease inhibitor family protein / Kunitz family protein	M5X024		-1.2
	Desiccation-like protein	M5XDI7		3.8
Protein degradation	Subtilase family protein	M5XAM8		-1.4
	Serine carboxypeptidase-like 25	M5WW69		-6.4
	Nucellin protein, putative Aspartic proteinase Asp1 precursor	M5XF88	1.3	
	Matrixin family protein Metalloendoproteinase 1 precursor	M5WU28		2.8
Signaling	Phospholipase C/ phosphoric diester hydrolase	M5WSV4	5.6	
	Disease resistance family protein / LRR family protein	M5XRG1		-5.8
	no original description	M5WB80		9.8
	Leucine-rich repeat transmembrane protein kinase	M5W7L1	-6.5	
Biodegradation of xenobiotics	Glyoxal oxidase-related	M5XBM0		8.8
	Glyoxalase I	M5WUG2		-2.0
Secondary metabolism	Laccase 14	M5XK01		-8.0
Lipid metabolism	Glycerophosphoryl diester phosphodiesterase family protein	M5W9K9		-1.5
	Phospholipase A2	M5XU00		5.3
	Gibberellin-regulated family protein, GASA, Snakin-2 like	M5XNP6	11.2	6.8

Hormone metabolism	Gibberellin-regulated family protein, GASA , Snakin-1	M5XFZ3*	7.4	10.3
	Auxin-responsive family AIR12	M5W7R0		7.2
	Auxin-responsive family protein AIR12	M5VKB3		-3.2
Miscellaneous	Beta Glucosidase 16	M5VK49	1.3	4.3
	Glycosyl hydrolase family 38 protein, alpha-mannosidase precursor	M5WXQ9		-2.3
	Glycosyl hydrolase family 38 protein, alpha-mannosidase precursor	L7RZF3		-3.3
	Alpha-glucosidase 1, maltase	M5WM42	-1.9	-2.0
	Sks4 (SKU5 Similar 4); copper ion binding/oxidoreductase	M5WN24		-6.5
	GDSL-motif lipase/hydrolase family protein	M5VGZ1		7.6
	GDSL-motif lipase/hydrolase family protein	M5XJN1	6.3	-1.3
	GDSL-motif lipase/hydrolase family protein	M5WSW8		-6.2
	(R)-mandelonitrile lyase, putative (berberine bridge enzyme)	M5XN40		7.1
	(R)-mandelonitrile lyase, putative (berberine bridge enzyme)	M5XQN8	6.3	-7.6
	(R)-mandelonitrile lyase, putative (berberine bridge enzyme)	M5WL V7		4.0
	Amidase family protein (berberine bridge enzyme)	M5VKZ0		6.3
	FAD-binding domain-containing protein (berberine bridge enzyme)	M5WGQ8	-5.8	-5.8
	FAD-binding domain-containing protein (berberine bridge enzyme)	M5WXM6		-5.8
Not assigned	Trichome birefringence-like 43	M5WTP1	1.0	
	Trichome birefringence-like 43	M5WK29	0.7	-5.5
	Glucuronidase 2	M5WWA5	1.1	
	Glucuronidase 2	M5WBD9	1.0	
	Exordium like 3	M5X0J0		6.3
	Phosphorylase family protein	M5X7J0	-5.8	-5.8

For each protein, functional classification based on MapMan, description, UniProtKB database accession number (Acc N) and log₂FC for proteins differentially abundant in DR12 and DR96 with respect to DR0 are provided. The log₂ Fold Change (FC) was calculated using Perseus software.* M5Y0D4, M5WW86 and M5XFZ3 are also repressed in FL12 with respect to FL0 (log₂FC=-10.1, log₂FC=-6.5, log₂FC=-11.2, respectively).