

Table 1 Twenty-two most increased apoplastic proteins in leaves of *P. persica* inoculated with *T. deformans*.

Functional classification	Description	Acc N	Max. increase		Log2 FC			
			Log2 FC	Sample	DR12-DR0	DR96-DR0	FL12-FL0	FL96-FL0
stress, biotic, PR5	Thaumatococcal protein 2 precursor (PpAZ8)	M5VQU4	12.92	DR96		12.92	-7.09	3.26
stress, biotic, PR5	Osmotin, Thaumatococcal-like	M5WR50	12.79	DR96		12.79		3.98
stress, biotic, PR5K	Thaumatococcal protein 2 (PpAZ8)	M5WA70	12.56	DR96		12.56		
cell wall, degradation	RD22; nutrient reservoir, BURP domain-containing protein	M5XEZ4	11.45	DR12	11.45	6.99	7.20	
hormone metabolism, gibberellin	GASA, Snakin-2 like	M5XNP6	11.21	DR12	11.21	6.82		
stress, biotic, PR12	Knot1 domain-containing protein, defensin	M5Y0D4	11.05	DR96		11.05	-10.11	
stress, biotic, PR5	Osmotin, Thaumatococcal-like	M5WT96	10.83	DR96		10.83		4.58
hormone metabolism, gibberellin	GASA, Snakin-1, Peamaclein	M5XFZ3	10.34	DR96		10.34	-11.22	
stress, biotic, PR14	Lipid-transfer protein 1-like	M5W0V2	10.13	DR96		10.13		
signalling, miscellaneous	No original description	M5WB80	9.78	DR96		9.78		
stress, biotic, PR5	Thaumatococcal protein 1 precursor (PpAZ44)	M5WTQ8	9.58	DR96		9.58	-2.44	
stress, biotic, PR9	Peroxidase, putative	M5X5U2	9.29	FL96		8.48	7.12	9.29
stress, biotic, PR14	Lipid Transfer protein 2	M5WW86	9.20	DR96		9.20	-6.50	
signalling, receptor kinases	Polygalacturonase inhibitor 1 precursor	M5W2E6	9.06	FL96		3.70		9.06
stress, biotic, PR14	Lipid transfer protein 1	M5W0S9	9.05	DR12	9.05			
biodegradation of xenobiotics	Glyoxal oxidase-related	M5XBM0	8.81	DR96		8.81		
stress, biotic, PR5	Thaumatococcal family protein	M5X697	8.66	DR96		8.66		3.52
signalling, misc	Rapid alkalization factor (RALF)	M5XT27	8.50	DR12	8.50	7.00	-3.40	
cell wall, modification	Xyloglucan endotransglycosylase	M5XCX9	8.48	DR12	8.48	7.95	6.38	5.57
cell wall, degradation	Putative beta-D-xylosidase 1 (PpAZ152)	M5XLA6	8.37	FL96		7.69		8.37
miscellaneous, GDSL-motif lipase	GDSL-motif lipase/hydrolase family protein	M5XRS5	7.55	DR96		7.55		

stress, biotic, PR6	Trypsin and protease inhibitor family protein/Kunitz family protein	M5WV82	7.53	DR96		7.53		3.61
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Relative amounts of proteins found in the apoplast of inoculated leaves were expressed as log₂ Fold Change (FC) in DR96 and DR12 with respect to DR0 (DR12-DR0 and DR96-DR0, respectively) in the case of the resistant genotype and in FL96 and FL12 in relation to FL0 (FL12-FL0 and FL96-FL0, respectively) in the case of the susceptible genotype. The 22 proteins exhibiting the greatest FC were selected from Supplementary Table 3. For each protein, functional classification based on MapMan, description, UniProtKB database accession number (Acc N) and FC for each comparison are provided. The comparison and the FC in which the maximum FC is detected are also shown.