

Supplementary Tables

Table S1. Identifiers of PR-like Genes from Filamentous Phytopathogens

Family	Pfam	Activity	<i>Magnaporthe oryzae</i>	<i>Botrytis cinerea</i>	<i>Puccinia striiformis</i>	<i>Fusarium oxysporum</i>	<i>Verticillium dahliae</i>	<i>Ustilago maydis</i>	<i>Phytophthora infestans</i>	<i>Hyaloperonospora arabidopsidis</i>	<i>Phytophthora ramorum</i>
PR1	PF00188	Immune signaling, Lipid-binding	MGG_03085	BCIN_08g02680	PSTG_01738	FOXG_00482	VDAG_09343	UMAG_01204	Phyinf117174	Hyaar151	Phyral_182067
			MGG_07807	BCIN_11g04730	PSTG_01739	FOXG_09795	VDAG_02733	UMAG_04343	Phyinf14883	Hyaar153	Phyral_182077
			MGG_06772	BCIN_01g09180	PSTG_13342	FOXG_10300	VDAG_03487		Phyinf14926	Hyaar18924	Phyral_182078
			MGG_13936	BCIN_14g00420	PSTG_11017	FOXG_12292	VDAG_04828		Phyinf14946	Hyaar1858	Phyral_182080
			MGG_03755		PSTG_15884	FOXG_12428			Phyinf14947	Hyaar1859	Phyral_182082
			MGG_05100		PSTG_06516	FOXG_14109			Phyinf14949	Hyaar110787	Phyral_173685
						FOXG_06245			Phyinf14951	Hyaar110788	Phyral_173684
									Phyinf14952	Hyaar110789	Phyral_173686
									Phyinf14953	Hyaar111490	Phyral_174540
									Phyinf14954	Hyaar112558	Phyral_178264
									Phyinf14955	Hyaar112559	Phyral_178267
									Phyinf14967	Hyaar1802	Phyral_178919
									Phyinf14970	Hyaar1803	Phyral_196220
									Phyinf16418		Phyral_196231
									Phyinf16419		Phyral_180241
									Phyinf16420		Phyral_180240
									Phyinf16421		Phyral_180242
									Phyinf18884		Phyral_180475
									Phyinf19703		Phyral_195732
									Phyinf111612		Phyral_196230
									Phyinf111613		Phyral_174818
									Phyinf111616		
									Phyinf112481		
									Phyinf112482		
									Phyinf112483		
									Phyinf14566		
									Phyinf14567		
									Phyinf14568		
									Phyinf14569		
PR2	PF00332	β -1,3-glucanase	MGG_04689	BCIN_03g03840	PSTG_16373	FOXG_15659	VDAG_00511	UMAG_06078	Phyinf1203	Hyaar18547	Phyral_185261
			MGG_06023	BCIN_08g00910	PSTG_16375	FOXG_03723	VDAG_07185	UMAG_06133	Phyinf113636	Hyaar1781	Phyral_172319
			MGG_10400	BCIN_01g11220		FOXG_16943	VDAG_09510		Phyinf113643	Hyaar114284	Phyral_172578
				BCIN_02g06910		FOXG_06041	VDAG_01689		Phyinf113647		Phyral_174267
							VDAG_02343		Phyinf113651		Phyral_180072
PR3	PF00182	Chitinase		/	/	/	/	/	Phyinf112400	Hyaar13729	Phyral_171328
									Phyinf115211		
									Phyinf115213		
PR4	PF00967	Chitinase	/	/	/	/	/	/	/	/	/
PR5	PF00314	Thaumatin	MGG_03044	BCIN_16g04140	PSTG_11165	FOXG_03360	VDAG_01910	UMAG_03807	Phyinf11803	/	/
				BCIN_09g04150	PSTG_11429				Phyinf18660		
PR6	PF00280	Protease inhibitor	/	/	/	/	/	/	/	/	/
			/	/	/	/	/	/	/	/	/
PR7	PF00082	Subtilisin-like endoprotease	MGG_10449	BCIN_10g02530	PSTG_10928	FOXG_01145	VDAG_06121	UMAG_04400	Phyinf116981	Hyaar11391	Phyral_184081
			MGG_08966	BCIN_08g02990	PSTG_14252	FOXG_01284	VDAG_05709	UMAG_02843	Phyinf1625	Hyaar12216	Phyral_184204
			MGG_07965	BCIN_05g03290	PSTG_04600	FOXG_08795	VDAG_05967	UMAG_12106	Phyinf11200	Hyaar15496	Phyral_138840
			MGG_03670	BCIN_06g01370	PSTG_05113	FOXG_09801	VDAG_07176	UMAG_03024	Phyinf117677	Hyaar15686	Phyral_173616
			MGG_02863	BCIN_16g04460	PSTG_00912	FOXG_12263	VDAG_02670	UMAG_06118	Phyinf117751	Hyaar16367	Phyral_179030
			MGG_15291	BCIN_05g05830	PSTG_00927	FOXG_13463	VDAG_01043		Phyinf17249	Hyaar18309	Phyral_179126
			MGG_09246	BCIN_15g03150	PSTG_08707	FOXG_03262	VDAG_05299		Phyinf17250		Phyral_195546
			MGG_04733	BCIN_03g02040	PSTG_09271	FOXG_17011	VDAG_06012		Phyinf19522		
			MGG_07358	BCIN_08g01020	PSTG_14941	FOXG_04749	VDAG_07131		Phyinf113152		
			MGG_16831	BCIN_06g01010	PSTG_04216	FOXG_04896	VDAG_02277		Phyinf113259		
			MGG_14860	BCIN_06g00620	PSTG_04259	FOXG_05775	VDAG_10474		Phyinf115494		
			MGG_09990	BCIN_06g00330	PSTG_04264	FOXG_09594	VDAG_07367		Phyinf113880		
			MGG_02649	BCIN_15g04670	PSTG_00696	FOXG_12879	VDAG_07373		Phyinf18629		
			MGG_10445		PSTG_00915	FOXG_13295	VDAG_08100				
			MGG_08415		PSTG_00930	FOXG_18343	VDAG_09626				
			MGG_08429		PSTG_00932	FOXG_14511	VDAG_03905				
			MGG_08436			FOXG_14793	VDAG_04865				
			MGG_03316			FOXG_02695	VDAG_03685				
			MGG_00282			FOXG_22424					
			MGG_03870			FOXG_16982					
			MGG_09352			FOXG_04700					
			MGG_09817			FOXG_05860					
			MGG_02531			FOXG_06062					
			MGG_09073			FOXG_19284					
			MGG_07404			FOXG_19376					
			MGG_07559			FOXG_19497					
			MGG_13977			FOXG_19498					
			MGG_00940			FOXG_08084					
			MGG_09322			FOXG_19740					
			MGG_03056			FOXG_02380					
						FOXG_14564					
						FOXG_13604					

PR8/11	PF00704	Chitinase	MGG_10333	BCIN_16g01590	PSTG_13182	FOXG_10827	VDAG_00322	UMAG_02758	Phyinf117427	Hyaar19540	Phyra1_181952
			MGG_04534	BCIN_07g00890	PSTG_14802	FOXG_14329	VDAG_05658	UMAG_10419	Phyinf113414	Hyaar114252	Phyra1_177376
			MGG_01336	BCIN_03g03220	PSTG_03810	FOXG_15329	VDAG_06206	UMAG_06190			
			MGG_03599	BCIN_05g02660	PSTG_03812	FOXG_15151	VDAG_06734				
			MGG_05533	BCIN_07g00160	PSTG_03813	FOXG_22639	VDAG_06825				
			MGG_04732	BCIN_02g00570	PSTG_03991	FOXG_22648	VDAG_00901				
			MGG_01247	BCIN_08g07050	PSTG_04016	FOXG_00277	VDAG_08741				
			MGG_11231		PSTG_15142	FOXG_00921	VDAG_09205				
			MGG_01876		PSTG_15161	FOXG_09583	VDAG_09560				
			MGG_08054		PSTG_16340	FOXG_10034	VDAG_02162				
			MGG_07927		PSTG_16360	FOXG_10748	VDAG_02356				
			MGG_08458		PSTG_00935	FOXG_11492	VDAG_02396				
			MGG_17153		PSTG_07122	FOXG_12653	VDAG_10493				
			MGG_00086		PSTG_07135	FOXG_12882	VDAG_04416				
			MGG_17552		PSTG_08478	FOXG_12492	VDAG_04782				
			MGG_04073			FOXG_14047	VDAG_04833				
						FOXG_14840	VDAG_04879				
						FOXG_15473	VDAG_07000				
						FOXG_03402					
						FOXG_16128					
						FOXG_16251					
						FOXG_16430					
						FOXG_17257					
						FOXG_17332					
						FOXG_06883					
						FOXG_08721					
						FOXG_15373					
						FOXG_17685					
PR9	PF00141	Heme-containing peroxidase	MGG_08200	BCIN_03g07850	PSTG_07052	FOXG_00142	VDAG_01449	UMAG_01947	Phyinf1450	Hyaar110151	Phyra1_151170
			MGG_04545	BCIN_07g05810	PSTG_07189	FOXG_10495	VDAG_00782	UMAG_02377	Phyinf11442	Hyaar14038	Phyra1_186067
			MGG_07790	BCIN_13g03680		FOXG_12260	VDAG_06204	UMAG_11067	Phyinf116039	Hyaar18166	Phyra1_138638
			MGG_00461	BCIN_01g09360		FOXG_13439	VDAG_10405		Phyinf116040	Hyaar18190	Phyra1_173141
			MGG_10877			FOXG_13788	VDAG_02834				Phyra1_173140
			MGG_10368			FOXG_14234	VDAG_03116				Phyra1_177036
			MGG_04337			FOXG_17106	VDAG_04826				
			MGG_14940			FOXG_17130					
			MGG_09398			FOXG_17180					
			MGG_09834			FOXG_17460					
PR10	PF00407	Ribonuclease-like	/	/	/	/	/	/	/	/	/
PR12	PF00304	Plant defensin	/	/	/	/	/	/	/	/	/
PR13	PF00321	Thionin	/	/	/	/	/	/	/	/	/
PR14	PF00234	Non-specific lipid-transfer protein	/	/	/	/	/	/	/	/	/
PR15/16	PF00190	Oxalate oxidase	MGG_07824	BCIN_16g03240	/	FOXG_08602	VDAG_00088	/	/	/	/
				BCIN_01g03150		FOXG_10358	VDAG_07656				
				BCIN_04g05650		FOXG_11962	VDAG_09056				
						FOXG_13375	VDAG_09019				
						FOXG_08770	VDAG_09251				
						FOXG_09843	VDAG_04026				
						FOXG_13201					
PR17	PF04450	Putative aminopeptidase	MGG_04184	BCIN_07g01400	/	FOXG_08519	VDAG_00452	/	/	/	/

The table gives gene identifiers of PR-like proteins that are present in fungal (left-hand part) and oomycetes (right-hand part) phytopathogens. /, indicates that there is no gene present that matches the annotation of the respective plant PR protein family. PR-like families that are absent from the genomes of fungal or oomycetes are shaded in light green, i.e., PR3-, PR4-, PR6-, PR10-, PR12-, PR13-, PR14-, PR17-like. Genes were identified by screening sequences of individual plant PR family members against the genome sequences of *Magnaporthe oryzae* (Phyicaria1_oryzae 70.15, v3.0), *B. cinerea* (B05.10), *Puccinia* spp. (*Puccinia striiformis* f.sp. *tritici* PST-78, v1.0), *Fusarium oxysporum* (*Fusarium oxysporum* f.sp. *lycopersici* 4287, v2), *Verticillium dahliae* (VdLs.17), *Ustilago maydis* (521, v2.0), *Phytophthora infestans* (T30-4), *Hyaloperonospora arabidopsidis* (Emoy2, v2.0), and *Phytophthora ramorum* (v1.1) in the PhytoPath database (<https://phytopathdb.org/>) (Pedro et al., 2016).