

Supplementary table S3: Spectra mapping to different sequences in the *R. solani* and wheat databases..

Treatment	Tissue	Time point	Rsolani Protein accession number	Rsolani Peptide sequence	Modifications identified by spectrum	Number of unique peptides	Number of total spectra	Spectrum name	SEQUEST XCorr score	Mascot ion score	Wheat entry found	Wheat Peptide sequence	SEQUEST XCorr score	Mascot ion score	Comment
Infection	Membrane	Early (3 dpi)	RSAG8_03810	NSPAIIIFIDEIDSIAPK		23	42	File10594 Spectrum31123 scans: 33809	4.61	74.5	Traes_4DL_D BA53B48C.1	NAPSIIIFIDEIDSIAPK	4.35	52.9	3AA shuffled (SPA vs APS)
Infection	Membrane	Early (3 dpi)	RSAG8_03810	NSPAIIIFIDEIDSIAPK		23	42	File10594 Spectrum31433 scans: 34140	4.34	60.4	Traes_4DL_D BA53B48C.1	NAPSIIIFIDEIDSIAPK	4.03	33.5	3AA shuffled (SPA vs APS)
Infection	Membrane	Early (3 dpi)	RSAG8_03810	NSPAIIIFIDEIDSIAPK		23	42	File10594 Spectrum31969 scans: 34726	3.68	34.2	Traes_4DL_D BA53B48C.1	NAPSIIIFIDEIDSIAPK	3.41	31	3AA shuffled (SPA vs APS)
Infection	Membrane	Late (7 dpi)	RSAG8_03810	NSPAIIIFIDEIDSIAPK		20	33	File10602 Spectrum30430 scans: 32990	4.86	93.4	Traes_4DL_D BA53B48C.1	NAPSIIIFIDEIDSIAPK	4.31	58.4	3AA shuffled (SPA vs APS)
Infection	Membrane	Early (3 dpi)	RSAG8_03810	VLNOQILTEMDGMNAK	Oxidation (+16)	23	42	File10594 Spectrum14006 scans: 15653	3.41	46.9	Traes_4DL_D BA53B48C.1	VLNOLLTEMDGMNAK	3.41	38.1	Ambiguous AA (I vs L)
Infection	Membrane	Early (3 dpi)	RSAG8_03810	VLNOQILTEMDGMNAK		23	42	File10594 Spectrum17448 scans: 19269	4.89	60	Traes_4DL_D BA53B48C.1	VLNOLLTEMDGMNAK	4.89	47.7	Ambiguous AA (I vs L)
Infection	Membrane	Late (7 dpi)	RSAG8_03810	VLNOQILTEMDGMNAK	Oxidation (+16)	20	33	File10602 Spectrum12703 scans: 14375	4.21	70.3	Traes_4DL_D BA53B48C.1	VLNOLLTEMDGMNAK	4.21	70.3	Ambiguous AA (I vs L)
Infection	Membrane	Late (7 dpi)	RSAG8_03810	VLNOQILTEMDGMNAK	Oxidation (+16)	20	33	File10602 Spectrum8925 scans: 10408	4.47	63	Traes_4DL_D BA53B48C.1	VLNOLLTEMDGMNAK	4.47	54.2	Ambiguous AA (I vs L)
Infection	Culture filtrate	Late (7 dpi)	RSAG8_05268	YPNSGDGPYAGHIMVVSEQK	Deamidated (+), Deamidated (+1)	3	73	File10590 Spectrum10668 scans: 11961	2.62	17.5	Traes_1AL_E9 6C0662D.2	YCSILGTATGGNLDCTYQR	4.26	37.1	
Infection	Membrane	Early (3 dpi)	RSAG8_07749	IPIFSAAGLPHNEIAAOICR	Carbamidome thyl (-57)	5	5	File10594 Spectrum19027 scans: 20927	4.31	61	tr W5DET9 WSDET9_WH EAT	IPIFSAAGLPHNEIAAOICR	4.31	45.4	Ambiguous AA (I vs L)
Infection	Membrane	Late (7 dpi)	RSAG8_07749	IPIFSAAGLPHNEIAAOICR	Carbamidome thyl (-57)	10	16	File10602 Spectrum17571 scans: 19487	4.2	46.4	tr W5DET9 W5DET9_WH EAT	IPIFSAAGLPHNEIAAOICR	4.2	38.2	Ambiguous AA (I vs L)