

Supplementary Materials: Interacting Environmental Stress Factors Affects Targeted Metabolomic Profiles in Stored Natural Wheat and That Inoculated with *F. graminearum*

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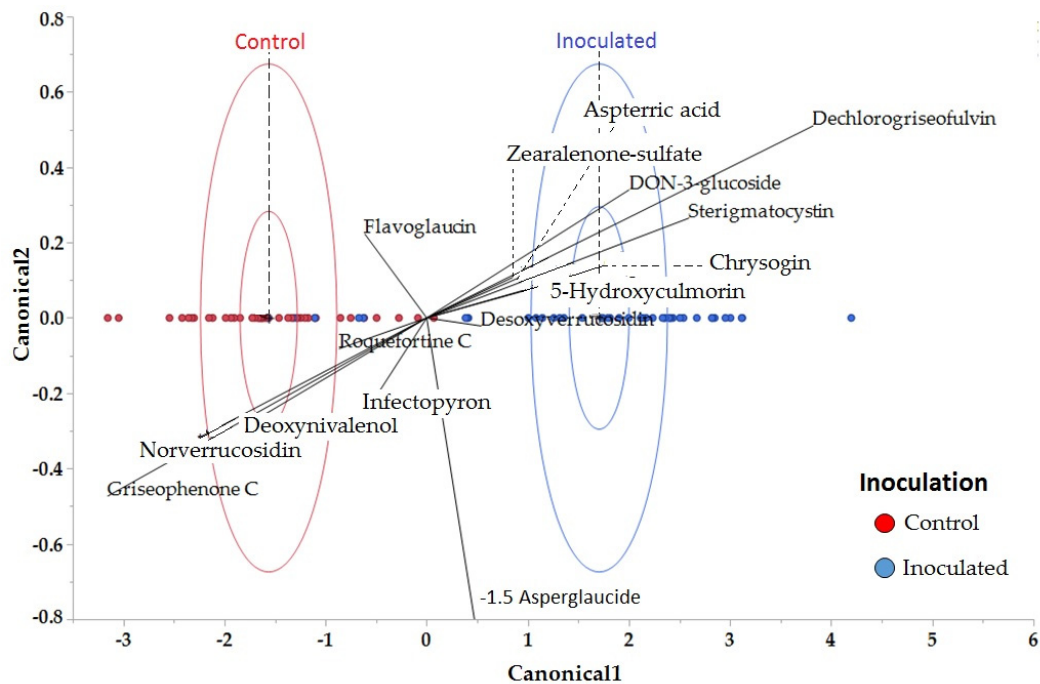
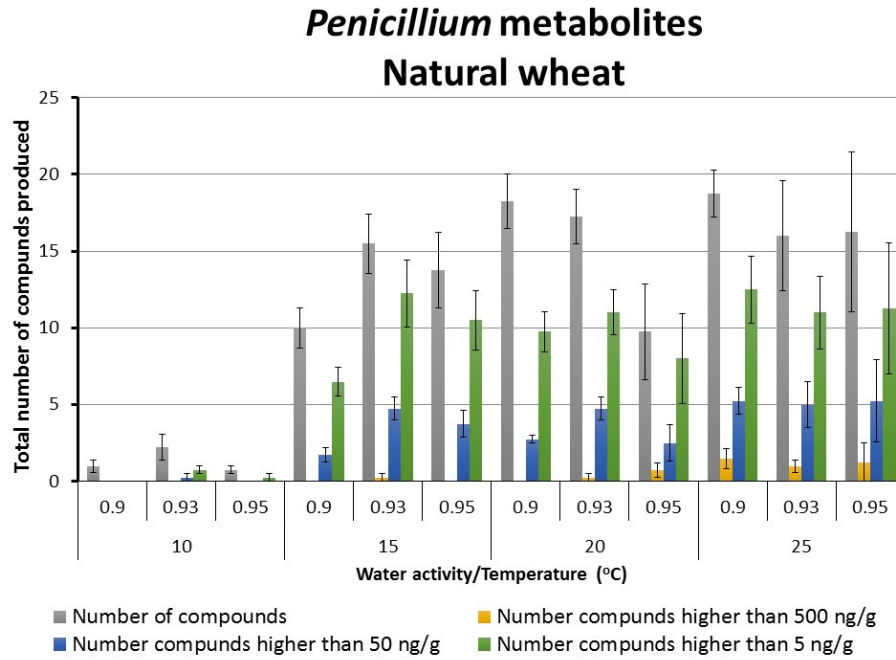
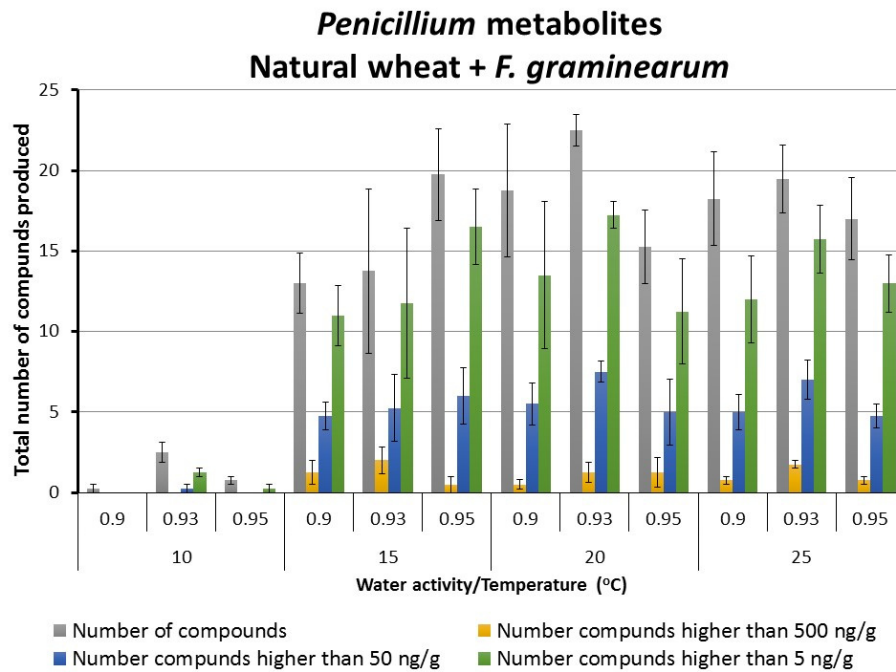


Figure S1. Canonical plot of the linear discriminant analysis between stored naturally contaminated wheat (control) and that inoculated with *F. graminearum*.



(a)



(b)

Figure S2. Effect of storage conditions on the *Penicillium* secondary metabolites (out of 45) produced in stored natural wheat grain (a) and in stored wheat grain + *F. graminearum* (b) under different interacting temperature x water activity conditions for 15 days. Data are for means + S.E.

Table S1. Secondary metabolites included in the linear discriminant analysis (LDA) grouped by fungal genera.

<i>Fusarium</i>	<i>Penicillium</i>	<i>Alternaria</i>	<i>Aspergillus</i>	Unspecific
Zearalenone-sulfate	Desoxyverrucosidin	Infectopyron	Sterigmatocystin	Asperglaucide
Deoxynivalenol	Norverrucosidin		Aspterric acid	
DON-3-glucoside	Roquefortine C			
5-Hydroxyculmorin	Griseophenone C			
Chrysogin	Dechlorogriseofulvin			
	Flavoglaucin			

Discriminant equation Canon1 = [1.7908027·10⁻⁷ x Zearalenone-sulfate + (-0.00173229067149818) x Deoxynivalenol + 0.0542366059471274 x DON-3-glucoside +0.00365544088741318 x 5-Hydroxyculmorin) + 0.0620752424956139 x Chrysogin + 0.0352002688455243 x Desoxyverrucosidin + -0.0370104256774014 x Norverrucosidin + -0.00184551544938296 x Roquefortine C + -0.791258703686439 x Griseophenone C + 0.231543704676479 x Dechlorogriseofulvin + (-0.00181899887872762) x Flavoglaucin + -0.0000759804488474338 x Infectopyron + 5.91681156878679 x Sterigmatocystin + 0.00574752284444148 x Aspterric acid + 0.128832906799609 x Asperglaucide] - 1.57772042376622.

Table S2. Canonical Details calculated from the overall pooled within-group covariance matrix Wilks' Lambda.

Eigenvalue	Percent	Cum Percent	Canonical Corr	Likelihood Ratio	Approx. F	NumDF	DenDF	Prob > F
2.73	100	100	0.86	0.268	13.81	15	76	<0.0001 *

* There are significant differences between groups.

Table S3. Discriminant scores.

Source	Count	Number Misclassified	Percent Misclassified	Entropy RSquare	-2LogLikelihood
Training	92	5	5.43	0.68	43.62

Table S4. Misclassified.

Actual	Predicted Count	
Inoculation	Control	Inoculated
Control	48	0
Inoculated	5	39

Table S5. Production of *Alternaria* secondary metabolites (altersolanol, altertoxin I, tentoxin, macrosporin, infectopyron, dihydroinfectopyron and total) in the two stored wheat grain treatments under different temperature x water activity conditions. Red indicates maximum production levels (ng/g) while green and shades of yellow/orange represents intermediate production levels. Key: <LOD, below limits of detection. Shading is per column.

T (°C)	aw	Natural Wheat							Natural Wheat + <i>F. graminearum</i>						
		AS ¹	ATX-I ²	TEN ³	Macros Porin	Infecto Pyron	Dihydro Infecto Pyron	Total Sum	AS ¹	ATX-I ²	TEN ³	Macros Porin	Infecto Pyron	Dihydro Infecto Pyron	Total Sum
10	0.9	<LOD	<LOD	1.8	0.6	642.5	<LOD	644.9	<LOD	<LOD	<LOD	<LOD	507.4	<LOD	507.4
	0.93	<LOD	2.7	<LOD	<LOD	778.3	<LOD	781.0	<LOD	<LOD	<LOD	<LOD	582.6	<LOD	582.6
	0.95	<LOD	<LOD	<LOD	<LOD	883.0	<LOD	883.0	<LOD	<LOD	0.1	<LOD	860.5	<LOD	860.7
15	0.9	<LOD	2.3	<LOD	<LOD	547.9	<LOD	550.1	<LOD	4.1	<LOD	<LOD	1021.0	<LOD	1025.0
	0.93	<LOD	2.7	<LOD	<LOD	1806.5	<LOD	1809.2	<LOD	<LOD	<LOD	0.4	2557.0	<LOD	2557.4
	0.95	<LOD	2.6	<LOD	<LOD	5078.4	<LOD	5081.1	<LOD	4.4	<LOD	<LOD	3769.6	<LOD	3773.9
20	0.9	<LOD	<LOD	<LOD	<LOD	1228.0	<LOD	1228.0	194.5	1.7	0.2	8.3	3316.3	<LOD	3521.1
	0.93	<LOD	<LOD	<LOD	<LOD	1291.1	<LOD	1291.1	938.7	<LOD	<LOD	23.5	2188.3	<LOD	3150.5
	0.95	<LOD	7.4	<LOD	0.3	8137.1	7.4	8152.2	<LOD	36.6	0.2	<LOD	5757.9	6.1	5800.8
25	0.9	<LOD	0.8	<LOD	<LOD	1900.8	<LOD	1901.6	<LOD	8.6	<LOD	<LOD	3803.0	<LOD	3811.7
	0.93	<LOD	11.1	0.3	0.2	5236.9	<LOD	5248.4	<LOD	36.5	0.1	0.4	11,564.5	<LOD	11,601.6
	0.95	<LOD	235.6	<LOD	<LOD	19,040.3	23.5	19,299.5	6784.1	87.0	0.2	79.7	18,530.2	23.7	25,505.1

¹ Altersolanol, ² Altertoxin-I, ³ Tentoxin (µg/kg).