

Mycotoxin producing potential of *Fusarium graminearum* isolates from Uruguayan barley

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Abstract

Twelve isolates of *Fusarium graminearum* were obtained from barley grains collected from different Uruguayan regions (harvest 1993–94). This was the predominant fungal species contaminating the crop due to a particular humid and warm season with cold nights conducive to toxin production. The isolates were grown on moist, sterile rice, extracted with aqueous methanol, and examined for mycotoxin production. Zearalenone (ZEA) and the trichothecenes deoxynivalenol (DON), 3- and 15-acetyl-DON (AcDON), nivalenol (NIV), fusarenon-X (FX) and T-2 toxin (T-2) were analyzed by thin-layer chromatography (TLC) and high-performance liquid chromatography (HPLC) and confirmed by gas chromatography–mass spectrometry (GC–MS).

Eleven of the 12 strains were DON and/or ZEA producers and 9 were AcDON positive. No NIV, or FX were detected. One strain produced T-2. The predominant acetyl-DON isomer was 15-AcDON. Mass-spectral analysis yielded detectable levels of other mycotoxins, 13-OH-apotrichothecenes, 11-epiapotrichothecenes, culmorin, sambucinol, and isotrichodermol being the most numerous.

From the metabolic profiles it is suggested that Uruguayan *F. graminearum* strains belong to the chemotype IB (DON/15-AcDON). The predominance of this chemotype is in accordance with data from Canada, United States, Mexico and Argentina which have similar climatic conditions that would favor *F. graminearum* growth and mycotoxin production.

Key words: Barley, Deoxynivalenol, *Fusarium graminearum*, Mycotoxins, Trichothecenes, Zearalenone

Introduction

Uruguay's main economic assets come from agriculture, with wheat, corn, barley and rice its most important crops. *Fusarium* disease affects all these grains and is caused mainly by *F. graminearum* [1,2]. Fungal development and mycotoxin formation are dependent on climatic conditions [3–6]. The fungal infection of grains by species of *Fusarium* is a common phenomenon in countries with temperate climates. In the River Plate region the 1993–94 period was particularly humid, warm and with cold nights, conducive to toxin production. *Fusarium* contamination caused severe losses to agriculture and animal productivity. Toxicogenic *Fusarium* species are now recognized to be a major agricultural problem with the production of a large number of toxic secondary metabolites. These

toxic metabolites include ZEA and the trichothecenes [2,7,8]. The toxigenic potential of monosporic *F. graminearum* strains isolated from barley grains collected from different regions of the country was determined by assaying for zearalenone and trichothecenes (DON, 3- and 15-AcDON, NIV, FX and T-2).

Material and methods

Isolation of Fusaria. Samples were collected from barley crops, harvest 1993–94 from 4 different regions of Uruguay. The code names indicate the location. One hundred grains from each location were surface disinfected with 1% sodium hypochlorite and 70% ethanol followed by three washes with sterile, distilled water and 5 grains/petri dish placed on dichloran–

chloramphenicol-peptone agar, DCPA [9] for 7 days at 25 °C. Selected *Fusaria* were subsequently subcultured onto potato dextrose agar, PDA [10] incubated for 7 days at 25 °C and on synthetic nutrient agar consisting of KH_2PO_4 , 1 g; KNO_3 , 1 g; $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$, 0.5 g; KCl, 0.5 g; glucose, 0.2 g; saccharose, 0.2 g; agar, 20 g and distilled water to 1 liter, with filter paper for 14 days at 25 °C. Cultures showing micro- and macroconidia in accordance to the *F. graminearum* species concept of Nelson *et al.* [11] were purified by single spore isolation and 12 of these isolates were chosen at random for determination of toxigenic potential.

Culture methods. Fungal cultures were prepared [12] in 250 erlenmeyer flasks containing 50 g of sterile, moist rice (50% moisture). They were inoculated with 2.5 ml of a spore suspension of each selected monosporic *F. graminearum* strain prepared by flooding the slant with 5 ml of sterile distilled water and vigorous agitation. The strains had been grown in PDA for 7 days at 25 °C. Incubation was carried out in the dark for one week at 25–28 °C and 2 weeks at 12 °C. At the end of the incubation period the moldy rice was extracted with 250 ml of methanol–water (45:55, v/v) containing 2.5% NaCl. Hexane (100 ml) was added and agitated for 30 min on a rotary shaker and filtered through Whatman No. 2 paper. The methanol–water phase was extracted 4 times with 25 ml chloroform each time and the combined extracts evaporated on a rotary evaporator at 45 °C. The residue obtained upon removal of the solvents was analyzed for ZEA and trichothecenes by TLC and HPLC.

Chemical analyses. Standards. Standards were from Agriculture Canada, Sigma Chemical Co. and Wako Chemicals. For HPLC they were dissolved in 4 ml of the solvent system to give concentrations of 2 µg ZEA, 1 µg NIV, 2 µg DON, 4 µg FX, 4 µg 3-AcDON and 4 µg 15-AcDON/ml. Injections of 20 µl of combined trichothecene standards and 20 µl of ZEA separately were used. For TLC analysis, standard concentrations were 50 µg ZEA, 100 µg NIV, 20 µg DON, 100 µg FX, 50 µg T-2, 100 µg, 3- and 100 µg 15-AcDON/ml. One, 3, 5 and 10 µl were spotted of each standard.

TLC. Extracts were dissolved in 1000 µl of chloroform and screened for ZEA, DON, 3- and 15-AcDON, NIV, FX and T-2 on precoated 0.20-mm thick silica gel plates without fluorescent indicator (E. Merck) using benzene–acetone (3:2, v/v), chloroform–acetone–isopropanol (8:1:1, v/v) as solvent systems and ethyl acetate–hexane (1:1, v/v) for the 3- and

15-AcDON separation. Five and 20 µl of the sample extracts were spotted in triplicate (250 mg and 1 g concentrations) with both internal and external standards used for positive mycotoxin identification. ZEA was observed at 254 nm and trichothecenes at 365 nm. ZEA was confirmed and Group B trichothecenes were detected by spraying with aluminum chloride (15% in ethanol–water, 15:85, v/v), heating at 120 °C for 2 min and observing blue fluorescence. Group A trichothecenes were sprayed after Group B detection with sulfuric acid (25% in water), heated at 120 °C for 2 min and observed as brownish–green fluorescent spots; ZEA turns orange and Group B trichothecenes disappear.

HPLC. Sample preparation. Equivalent to 50 g culture of each crude fungal extract was dissolved in 2 ml methanol to give 25 g/ml and 50 µl (1.25 g) of each were cleaned up on SPE-C18 columns (Supelco 0.5 g/6 ml). Preconditioning of the column was done with two 6-ml methanol washes followed by two 6-ml distilled water washes each with drainage under gravity. The vials containing the fungal extract were washed with three times 500-µl distilled water and added to the preconditioned columns. Columns were washed with 5.5 ml distilled water which was discarded. Trichothecenes and ZEA were eluted with 6 ml methanol–water (8:2, v/v) into a 50-ml round bottom flask. The eluted fraction was evaporated to dryness on a rotary evaporator at 45 °C, reconstituted by dissolving in 1 ml methanol to give 1.25 g/ml and kept in 4-ml screw cap vials with teflon liners. Before analysis the extracts were adjusted in the mobile phase to a concentration of 0.1 g/ml. Injections were 20 µl of each sample. HPLC was carried out using a Shimadzu LC10-AD pump, Rheodyne 7125 injector and a Waters model 440 UV/Vis detector (229 nm). A 5-µm Ultrachosphere C18 column (250 × 4.6 mm) was used with a flow rate of 1.5 ml of solvent per min. The mobile phase used for Type B trichothecenes was methanol–water (30:70, v/v) and acetonitrile–water (45:55, v/v) for ZEA. Retention times were 3.6 min for NIV, 5.3 min for DON, 7.6 min for FX, 15.9 min for AcDON and 10.8 min for ZEA.

HPLC was also carried out with a Hewlett Packard 1050 isocratic system with a variable wavelength UV/VIS detector (229 nm) and a 5-µm Hypersil C18 column (100 × 4.6 mm). The same mobile phases and flow rate were used as above.

Mass spectrometry. Toxins were confirmed at the Plant Research Centre, Agriculture and Agri-Food Canada, Ottawa. The fungal extracts prepared as above

Table 1. Mycotoxin production by *F. graminearum* isolates from barley

Isolate	DON ($\mu\text{g/g}$) ^a	15-AcDON ($\mu\text{g/g}$) ^a	ZEА ($\mu\text{g/g}$) ^a	Other mycotoxins identified by GC-MS	Diagnostic <i>m/z</i>
Y2	37.9	Neg.	41.4	7-deoxyDON 8-hydroxyisotrichodermol 15-OH-culmorin	280, 250, 249, 231, 203, 189 266, 251, 224, 218, 177, 123, 109 254, 236, 218, 206
Y3	20.3	15.0	80.5	15-OH-culmorin 15-OH-culmorone 3 β ,13-di-OH-11-epiapo	254, 236, 218, 206 252, 234, 219 252, 237, 224
Y4	25.0	11.0	55.0	culmorin 3 α ,11-oxy-13-OH-apo isotrichodermol 2,13-di-OH-3 α ,11-epoxyapo 7-deoxyDON sambucinol 3 α ,13-di-OH-11-epiapo 3 β ,13-di-OH-11-epiapo 15-OH-culmorin	238, 220, 205, 191, 177, 149 250, 235, 219, 161 202, 173, 124 266, 251, 177, 161 280, 250, 249, 231, 203, 189 266, 251, 235, 161, 124 252, 237, 224 252, 237, 224 254, 236, 218, 206
Y10	70.2	17.6	88.6	3 α ,11-oxy-13-OH-apo 3 α ,13-di-OH-11-epiapo 3 β ,13-di-OH-11-epiapo 15-OH-culmorin 7-deoxyDON 8-hydroxyisotrichodermol	250, 235, 219, 161 252, 237, 224 252, 237, 224 254, 236, 218, 206 280, 250, 249, 231, 203, 189 266, 251, 224, 218, 177, 123, 109
Y17	3.8	6.9	27.9	culmorin 3 α ,11-oxy-13-OH-apo 3 β ,13-di-OH-11-epiapo 3 α ,13-di-OH-11-epiapo 15-OH-culmorin 7-hydroxyisotrichodermol 7-hydroxyisotrichodermin 7-deoxyDON	238, 220, 205, 191, 177, 149 250, 235, 219, 161 252, 237, 224 252, 237, 224 254, 236, 218, 206 266, 218, 177, 159, 140, 139, 123, 109 308, 266, 218, 177, 159, 140, 123, 109 280, 250, 249, 231, 203, 189
S6	Neg.	Neg.	98.0	culmorin culmorone 3 α ,11-oxy-13-OH-apo isotrichodermol 3 β ,13-di-OH-11-epiapo 3 α ,13-di-OH-11-epiapo 12-OH-culmorin 15-OH-culmorin	238, 220, 205, 191, 177, 149 236, 218, 203 250, 235, 219, 161 202, 173, 124 252, 237, 224 252, 237, 224 236, 218, 205 254, 236, 218, 206
S13	2.7	1.7	9.0	culmorin culmorone 3 α ,11-oxy-13-OH-apo 3 β ,13-di-OH-11-epiapo 3 α ,13-di-OH-11-epiapo sambucinol 8-hydroxyisotrichodermol 8-hydroxyisotrichodermin 7-deoxyDON	238, 220, 205, 191, 177, 149 236, 218, 203 250, 235, 219, 161 252, 237, 224 252, 237, 224 266, 251, 235, 161, 124 266, 251, 224, 218, 177, 123, 109 308, 266, 224, 123, 109 280, 250, 249, 231, 203, 189

Table 1. Continued

Isolate	DON ($\mu\text{g/g}$) ^a	15-AcDON ($\mu\text{g/g}$) ^a	ZEA ($\mu\text{g/g}$) ^a	Other mycotoxins identified by GC-MS	Diagnostic <i>m/z</i>
S15	70.7	2.7	111	3 α ,11-oxy-13-OH- <i>apo</i>	250, 235, 219, 161
				isotrichodermol	202, 173, 124
				7-deoxyDON	280, 250, 249, 231, 203, 189
				7-hydroxyisotrichodermol	266, 218, 177, 159, 140, 139, 123, 109
				7-hydroxyisotrichodermin	308, 266, 218, 177, 159, 140, 123, 109
LE2	7.2	10.0	45.8	culmorin	238, 220, 205, 191, 177, 149
				3 α ,11-oxy-13-OH- <i>apo</i>	250, 235, 219, 161
				isotrichodermol	202, 173, 124
				15-OH-culmorin	254, 236, 218, 206
				3 α ,13-di-OH-11- <i>epiapo</i>	252, 237, 224
				3 β ,13-di-OH-11- <i>epiapo</i>	252, 237, 224
				sambucinol	266, 251, 235, 161, 124
LE19	18.0	9.0	60.0	3 α ,11-oxy-13-OH- <i>apo</i>	250, 235, 219, 161
				isotrichodermol	202, 173, 124
				3 β ,13-di-OH-11- <i>epiapo</i>	252, 237, 224
				3 α ,13-di-OH-11- <i>epiapo</i>	252, 237, 224
LE21	30.0	12.0	75.0	culmorin	238, 220, 205, 191, 177, 149
				15-OH-culmorin	254, 236, 218, 206
				3 α ,13-di-OH-11- <i>epiapo</i>	252, 237, 224
				3 β ,13-di-OH-11- <i>epiapo</i>	252, 237, 224
DC35P ^b	Neg.	Neg.	Neg.	No relevant toxins	

^aValues are average of TLC and LC determinations.

^bT-2 producer by TLC.

None of examined isolates produced NIV or FX by TLC, LC or MS.

Table 2. Deoxynivalenol and zearalenone recoveries

Sample	DON	ZEA	Recoveries	Average recoveries
	TLC/LC ($\mu\text{g/g}$)	TLC/LC ($\mu\text{g/g}$)	(% of added) TLC/LC	(% of added) TLC/LC
Isolate DC35P	100/99.2	100/99.2		
spiked with	98/97	—	98/97	97.6/96.1
100 $\mu\text{g/g}$ DON	95/92		95/92	
Isolate DC35P		50/37.5	100/75	
spiked with	—	49/30.0	98/60	97.3/71
50 $\mu\text{g/g}$ ZEA		47/38.8	94/78	

were taken to dryness under nitrogen, redissolved in ethyl acetate and analyzed using a Finnigan model 4500 gas chromatography/mass spectrometer system [13] with temperature program 100–280 °C at 15 °C/min. Mass spectra were determined in the elec-

tron impact mode and compared with the mycotoxin data base library of the Plant Research Centre, Agriculture and Agri-Food Canada.

Results and discussion

Mycotoxins detected in the *F. graminearum* extracts are reported in Table 1. All DON, AcDON and ZEA TLC and HPLC findings were confirmed by GC-MS. Diagnostic m/z values observed were 296, 278, 265 and 248 for DON; 278, 265, 249, 231 and 203 for 15-AcDON; and 318, 300, 249, 231, 204 and 188 for ZEA, although weaker ions were missing in some cases for DON and 15-AcDON. DON and/or ZEA were consistently produced by 11 of the 12 isolates. AcDON was produced by 9 out of 12 strains tested. The major isomer was 15-AcDON. AcDON separation was achieved by TLC and identity confirmed by GC-MS. The 3- and 15-AcDON were not separated by the HPLC method used. One strain (S6) produced only high yields of ZEA as the main mycotoxin. Isolate Y10 produced high levels of all three toxins, with yields of DON, AcDON and ZEA of 70.2, 17.6 and 88.6 $\mu\text{g/g}$ respectively. Isolate S15 also produced high concentrations while other isolates can be classified as moderate (e.g. Y3, Y4, LE21) and low (e.g. S13, Y17) producers.

Concentrations of major toxins varied. ZEA was produced by 91.7% of the strains (11/12) at an average concentration of 62.9 $\mu\text{g/g}$ (9–111), 83.3% (10/12) produced an average of 28.6 $\mu\text{g/g}$ (2.7–70.7) of DON and 75% (9/12) produced an average of 9.5 $\mu\text{g/g}$ (1.7–17.6) of 15-AcDON. DC35P was unique in that it produced unconfirmed T-2 by TLC but none of the other mycotoxins looked for were positively identified nor did it produce any relevant toxin by HPLC or GC-MS. No NIV or FX were detected in any of the 12 extracts.

Mass-spectral data revealed also a number of minor secondary metabolites [13,14] including derivatives of culmorin, sambucinol, isotrichodermin and isotrichodermol and of apotrichothecene (apo): 13-OH-apo and the epimers (3 α -OH and 3 β -OH) of 13-hydroxy-11-epiapo as the most commonly present (Table 1). Identifications were based on GC retention time and matching of mass spectra.

Control recoveries were determined by TLC for DON and ZEA (Table 2) with 97.6% and 97.3% as the average recoveries from control cultures spiked at 100 and 50 $\mu\text{g/g}$, respectively. Recoveries for DON and ZEA were also measured by HPLC with 96.1% and 71% average recoveries respectively.

The results of toxin production by the isolates tested suggest that there is a large potential for trichothecene, especially DON, and ZEA production in Uruguayan fields. The metabolic profiles of the *F. graminearum*

isolates analyzed were similar, having chemotype IB (DON, 15-AcDON) [15] in common.

Conclusions

This study indicates that *F. graminearum* strains isolated from barley are capable of producing a variety of toxins, constituting a potential human and animal health hazard. Barley is an important substrate for feed and the main raw material for malting and beer production, both processes which may not reduce mycotoxin levels significantly [16,17].

The results of this limited survey also indicate that our isolates would belong to the chemotype IB which is in agreement with reports from Canada, the United States, Mexico and Argentina [15,18–21] where this is the dominant chemotype of the species.

The data contribute to furthering our knowledge of chemotypes and geographic distribution and support the premise that there is a regional relationship between the origin of *F. graminearum* and the production of 3- or 15-AcDON as the major isomer. A greater percentage of the isolates producing 15-AcDON are found in North and South America analogous to DON and not NIV production in both continents [22,23].

Further studies involving a greater number of isolates from grain samples are currently under way.

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