CONFIRMATION OF DEOXYNIVALENOL PRESENCE IN CHILEAN WHEAT BY HIGH-PERFORMANCE THIN-LAYER CHROMATOGRAPHY-MASS SPECTROMETRY

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ABSTRACT

Deoxynivalenol (DON) is a mycotoxin that belongs to trichothecene subgroup B. Due its detrimental effects for human health, DON has been analyzed and detected by nearly all chromatographic methods. The present work reports for the first time the application of HPTLC/MS to confirm the presence of DON in wheat crops. Chromatography was performed on silica gel 60 F_{254} HPTLC plates using toluene - ethyl acetate - formic acid 6:3:1 (*V*/*V*/*V*) as mobile phase. After post-chromatographic derivatization with 10 % AlCl₃, fluorescence detection was carried out at UV 366/>400 nm. Bands identity and purity was confirmed by mass spectrometry acquiring analytes directly from the sample bands by means of TLC/MS elution head-based interface. Both, sample and standard bands showed clear mass signals at *m*/z 341 that corresponds to the formate adduct of DON [M+HCOO]. Thus, the presence of DON in the Chilean wheat sample was confirmed.

Keywords: Deoxynivalenol, mycotoxins, HPTLC/MS, Fusarium, TLC.

1. INTRODUCTION

Mycotoxins are secondary metabolites produced by several fungus genera, of which Aspergillus, Penicillium and Fusarium are the most important ones. Among other mycotoxins the genera Fusarium produces trichothecenes, which is one of the major class of mycotoxins composed of more than 200 toxins¹. They are chemically classified in four groups (type A, B, C and D) according to the type of substitution of the base structure (tricyclic 12,13-epoxytrichothec-9-ene). Deoxynivalenol (DON, Figure 1) is a type B trichothecene (containing a carbonyl group at C8) reported as one of the most prevalent food-associated mycotoxin, especially in cereal grains². US and European studies have reported that more than 50% of cereal and cereal-derived samples are positive for DON² and some of them with high concentrations3. Further, this mycotoxin is thermoresistant (up to 350°C) being therefore stable during processing, e.g. cooking² DON, also known as "vomitoxin" due to the emetic effect in pigs, produces relevant adverse (toxic) effects in humans. The alterations over the immune, gastrointestinal, endocrine and reproductive systems2, 3 led to the Joint FAO/ WHO Expert Committee on Food Additives (JECFA) to establish a provisional maximum tolerable daily intake (PMTDI) of 1 µg/kg body weight for DON and its acetylated derivatives (3-Ac-DON and 15-Ac-DON)4. Regarding carcinogenicity, the International Agency for Research on Cancer (IARC) classified DON into Group 35. In view of its toxic potential, the European Community6 established maximum levels for unprocessed and processed cereals (200 to 1750 μ g kg⁻¹), which has been used to routinely evaluate its presence in several countries finding samples that exceed the proposed limits7. DON determination can be done by several analytical methods, including almost all chromatographic techniques, e.g. high performance liquid chromatography (HPLC) coupled to ultraviolet (UV)8, fluorescence (FL)9 and mass spectrometry (MS)¹⁰ detectors; gas chromatography (GC) coupled to MS¹¹, flame ionization¹² and electron capture¹³ detectors and high performance thin layer chromatography (HPTLC) with FL detection (FLD)¹⁴. These methods use an extraction procedure that generally includes a clean-up step with QuEChERs15, immunoaffinity16 or multifunctional solid phase extraction (SPE) columns17.



Figure 1. Structure formula of 4-deoxynivalenol (DON) [12,13-epoxy-3,7,15-trihydroxytrichothec-9-en-8-one].

Many reports about cereal contamination with *Fusarium* toxins were published worldwide; whereas in Chile, located in the south-west of South America, there was no data about any outbreak attributable to *Fusarium* toxins present on domestic cereal production. First in February 2004, a fusariosis outbreak infecting a wheat crop was detected in the central part of the country. The wheat presented the classical symptoms of white head blight.

In this study, representative wheat crop samples were collected directly from the infected field as well as an overall sample was taken at the end of the harvest. A simple extraction procedure was used based on multifunctional SPE columns. These extracts were analyzed by a validated HPTLC/FLD method. Potential positive findings were confirmed by HPTLC/MS.

2. EXPERIMENTAL

2.1. Chemicals and reagents

Deoxynivalenol (DON) was purchased from Sigma (St. Louis, MO, USA). HPTLC plates silica gel 60 F_{254} (10 x 10 cm), aluminum chloride 6-hydrate, methanol, ethanol, toluene, ethyl acetate, formic acid and acetonitrile were obtained from Merck (Darmstadt, Germany). All solvents were of HPLC quality or distilled before its use. Ultra-pure water (18 M Ω cm) was produced by Synergy System (Millipore GmbH, Schwalbach, Germany). Romer Labs Mycosep DON multifunctional columns 225 were obtained from Coring System Diagnostix (Gernsheim, Germany). Before usage, the plates were prewashed with methanol and dried for 30 min at 120 °C.

2.2. Preparation of the standard solution

A methanolic stock solution was prepared from the crystalline DON (100 μ g mL⁻¹) and diluted 1:4 with methanol to obtain the standard solution (25 μ g mL⁻¹) stored at -20 °C.

2.3. Sample preparation

Due to the matrix complexity and the expected low levels of DON ($\mu g/$ kg to mg/kg), an exhaustive sample preparation was required to accomplish an adequate clean up and analyte concentration. DON was extracted applying the method reported by Radova et al.¹⁸ with slight modifications. Briefly, wheat crop samples were ground in a Romer mill (Romer Labs, Union, MO, USA) to obtain a coarse grained powder. The powder (25 g) was mixed with 100 mL acetonitrile and water, 21:4 (V/V) by an Ultra-turrax T 25 (Janke and Kunkel, IKA Labortechnik, Staufen, Germany) at high speed for 5 min. The solution obtained was filtered through a filter paper (MN-615, Macherey-Nagel, Düren, Germany). The filtrate (8 mL) was transferred to an assay tube and the Mycosep DON multifunctional column 225 was pushed over it. The purified extract (4 mL) was transferred to a vial and the solvent was evaporated to dryness under a gentle stream of nitrogen. The residue was dissolved in 500 μ L of ethyl acetate and methanol, 19:1 (V/V).

2.4. Chromatography

Sample extracts and standard solutions were applied with the Automatic TLC Sampler 4 (ATS 4, CAMAG, Muttenz, Switzerland) with the following settings: band length 6.0 mm, dosage speed 150 nL s⁻¹, distance from the left side 15.0 mm and from the lower side 8.0 mm, 10 tracks per plate. For MS recordings, the application volumes were 12 μ L standard solution (300 ng/band) and 8 μ L sample solution (resulting in a similar concentration). Chromatography was performed in a Horizontal Development Chamber, 10 x 10 cm (CAMAG), with toluene - ethyl acetate - formic acid 6:3:1 (*V*/*V*/*V*) as solvent system. After 2 min drying in a stream of warm air, the plate was immersed in a solution of 10 % AlCl₃ in ethanol - water 1:1 (*V*/*V*) using the TLC Immersion Device (CAMAG) and heated at 120 °C on the TLC Plate Heater (CAMAG) for 10 min. Fluorescence measurement of DON was performed at UV 366/>400 nm with the TLC Scanner 3 (CAMAG) using a slit dimension of 5.0 mm x 0.3 mm. All automated instruments were controlled via the software winCATS 1.4.1 Planar Chromatography Manager (CAMAG).

2.5. Mass spectrometry

For recording of mass spectra the underivatized plate was measured at 220 nm and the migration distance of each compound was marked with a soft pencil. Using an HPLC pump (HP 1100, Agilent Technologies, Palo Alto, USA) and the ChromeXtraktor interface (ChromAn, Holzhausen, Germany), the zone of interest on the HPTLC plate was directly eluted into the electrospray ionization (ESI) source of the VG platform II single-quadrupole mass spectrometer (Micromass, Manchester, United Kingdom). The zone of interest was sealed with the cutting edge of the elution head and eluted from the layer with a mixture of methanol and formate buffer (10 mmol L⁻¹, pH 4.0), 19:1 (V/V), at a flow rate of 0.1 mL min⁻¹. The MS system was operated in the full scan mode with the following parameters for ESI in the negative ionization mode: source temperature 120 °C; capillary voltage -3.5 kV; HV lens 0.5 kV; cone voltage -55 V; dwell time 0.5 s; inter channel delay 0.02; repeats 1; span 0.5. Data were processed with Mass Lynx 3.2 software (Micromass).

3. RESULTS AND DISCUSSION

3.1. Method validation

The validation of the method was carried out according to the guideline of the International Conference on Harmonization (ICH)¹⁹ (**Table 1**). The regression analysis was performed with six DON concentration levels in triplicate. The recovery was calculated via three separate blank samples, each spiked at the concentration level of 1 mg kg⁻¹. As indicator for the precision of the method, the repeatability of the sample analysis was determined. The limits of detection and quantification were calculated using a signal to noise ratio of 3 and 10, respectively.

Table 1.	Validation	data fo	r determination	of DON in	wheat by	HPTLC/
FLD.						

Linearity [ng band-1]	8-120		
Equation by peak area	y = 16.08x + 23.12		
r and R ²	0.9991 and 0.9982		
Recovery [%, $n = 3$, 1 mg kg ⁻¹]	90.1 ± 6.4		
Repeatability (%RSD) ^a	7.1		
Detection limit [mg kg ⁻¹] ^a	0.05		
Quantification limit [mg kg ⁻¹] ^a	0.19		

^aApplication of 20 µL sample extract solution

3.2. Sample analysis

Applying the validated method, representative wheat crop samples sampled directly from the field were analyzed. The overall wheat sample (sampled at the end of the harvest of 2004) was re-analyzed in 2005 and the mean value of DON (n = 3) was established to be 5.5 ± 0.3 mg kg⁻¹ (%*RSD* 5.1 %) via peak height and 5.4 ± 0.3 mg kg⁻¹ (%*RSD* 5.2 %) via peak area. These values were in good agreement with the previous results determined directly after the harvest of 2004. Using the described chromatographic system and the Mycosep DON multifunctional column 225, the analyte (DON) was completely separated from the matrix. The proof of this specificity in detection can be clearly seen in the chromatogram illuminated at UV 366 nm (**Figure 2**).

3.3. Confirmation by HPTLC/MS

The presence of DON in the wheat sample was confirmed by HPTLC/ ESI-MS. The elution head-based ChromeXtraktor interface²⁰, which reliability was already proven in trace analysis^{21; 22} and which was modified for elution from glass-backed HPTLC plates²³, was a highly targeted tool to combine HPTLC with MS. Only from the selected zone of interest, mass selective compound information was obtained. The targeted use of HPTLC/MS avoided the automatic transfer of background signals and matrix signals into the MS system, which is the routine case for column-based hyphenations. Full scan mass spectra were recorded in the negative ionization mode between m/z 200 and 500 (**Figure 3**). High amounts of DON (300 ng band⁻¹), which corresponded to the sample zone intensity, were selected for direct elution into the MS.



Figure 2. HPTLC chromatogram documented at UV 366 nm showing DON standard solutions (tracks 1-3 and 8-10: 75, 50 and 25 ng band⁻¹; twofold determination), the Chilean wheat sample containing DON (tracks 4-6) and the same sample extracted with a re-used Mycosep DON multifunctional column 225 (track 7).



Figure 3. Confirmation by HPTLC/MS via direct elution from the HPTLC plate silica gel 60, exemplarily showing full scan mass spectrum of the 300 ng band⁻¹ DON standard zone ionizing as formate adduct [M+HCOO]⁻ at m/z 341.

CONCLUSIONS

The outlined HPTLC/FLD/ESI-MS method for determination of DON in wheat offered a fast, low-cost alternative due to the coupling of an efficient screening method with a selective detector, if required. The confirmation of the results by HPTLC/MS avoided the assignment of false positives.

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