

Carry-over assessment of fumonisins and zearalenone to poultry tissues after exposure of chickens to a contaminated diet – A study implementing stable-isotope dilution assay and UHPLC-MS/MS

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- 1 Carry-over assessment of fumonisins and zearalenone to poultry tissues after
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- 5 Vincent Hort^a, Marina Nicolas^a, Angélique Travel^b, Catherine Jondreville^c, Corentin Maleix^a,
- 6 Elisabeth Baéza^c, Erwan Engel^d, Thierry Guérin^{a*}
- ^a Université Paris-Est, Anses, Laboratory for Food Safety, F-94701 Maisons-Alfort, France.
- 9 b ITAVI, INRA Centre Val de Loire, F-37380 Nouzilly, France.
- 10 ° INRA, URA83, F-37380 Nouzilly, France.
- ^d INRA, UR370 QuaPA, Microcontaminants, Aroma & Separation Science group (MASS), F-
- 12 63123 Saint-Genès-Champanelle, France.
- *Corresponding author. Tel.: +33 149 772 711; E-mail address: thierry.guerin@anses.fr

Abstract

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This study assessed the transfer of fumonisins B1 and B2 (FBs), zearalenone (ZEA) and 17 zearalenone metabolites α -zearalanol (α -ZAL) and α -zearalenol (α -ZEL) to poultry tissues. 18 Two experimental groups of 9 male chickens each were exposed for 8 weeks to a 19 contaminated diet (FB group: 12.7 mg FB1 + FB2 kg⁻¹, ZEA group: 0.40 mg ZEA kg⁻¹). To 20 measure the carry-over from feed to animal tissues (liver and muscle), a sensitive and accurate 21 SIDA-UHPLC-MS/MS method was developed and validated. For all mycotoxins, the limit of 22 detection (LOD) was 0.3 µg kg⁻¹ and the limit of quantification (LOQ) was 1.0 µg kg⁻¹. 23 Recoveries ranged from 92% to 107% and the intermediate precision coefficients of variation 24 (CV_{IP}) were between 4.3% and 13%. ZEA and α -ZAL were not detected in livers, whereas α -25 ZEL was detected in five out of eight samples at levels between LOD and LOQ. FBs were 26 detected and quantified in the livers of all animals exposed to the contaminated diet (mean 27 30.3 µg FB1 kg⁻¹ and 2.3 µg FB2 kg⁻¹). A significant correlation between the FB1 and the 28 FB2 contents in the liver was demonstrated and FB carry-over factors (CFs) from feed to liver 29 were determined (CF_{FB1}: 0.003 and CF_{FB2}: 0.001). Filet muscles from the same animals were 30 also analysed. FB1 was quantified at trace levels in eight samples out of nine (mean 2.0 µg 31 kg⁻¹) and was only detected in the remaining sample. FB2 was detected in only one muscle 32 33 sample.

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Keywords: SIDA-UHPLC-MS/MS, fumonisins, zearalenone, poultry tissues, contaminated diet, carry-over

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1. Introduction

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39 Fumonisins (FBs) and Zearalenone (ZEA) are mycotoxins produced by Fusarium species prior to cereal harvest or during poor storage conditions (Marin, Ramos, Cano-Sancho, & 40 Sanchis, 2013). FBs are hepatotoxic in all animal species studied, nephrotoxic for several of 41 them, and are also considered cytotoxic (Gelderblom et al., 2001; Völkel, Schröer-Merker, & 42 Czerny, 2011). FB1, the most abundant analogue of the fumonisins family, was classified by 43 the International Agency for Research on Cancer (IARC) in group 2B, compounds considered 44 carcinogenic to animals and possibly carcinogenic to humans (IARC, 1993). ZEA is an 45 endocrine disrupter with oestrogenic effects, thought to reduce male fertility in humans and 46 47 wildlife populations and possibly involved in cancer development (Stopper, Schmitt, & Kobras, 2005). 48 For human consumers, the main sources of exposure to FBs and ZEA are cereals and cereal-49 50 based products (AFSSA, 2006; Bailly & Guerre, 2009; Leblanc, Tard, Volatier, & Verger, 2005; SCOOP, 2003). Therefore, risk management is mainly based on the control of plant-51 52 based food and feed, as demonstrated by the regulations and recommendations adopted by the 53 European Union (EU) (European Union, 2006, 2007). Limits in feed and those in food are not always correlated. Usually, feed limits are defined on the basis of the contamination levels 54 commonly found, applying the principle of "as-low-as-reasonably achievable", whereas food 55 limits are based on tolerable daily intake levels for human consumption. This approach is 56 adopted mainly because of insufficient data for carry-over from feed to the target organ of the 57 animal or product (van Raamsdonk et al., 2009). Therefore, consumers may be exposed to the 58 toxic compounds indirectly due to the presence of residual contamination in foods from 59 animals that have been fed with contaminated feeds (Maragos, 2010). Although the 60 contribution of animal products is not considered significant, it should be taken into account 61 in risk assessments of compounds presenting chronic toxicity, such as mycotoxins (Meyer, 62

Mohr, Bauer, Horn, & Kovács, 2003). Moreover, depending on the metabolic pathways 63 involved, the passage through the animal may represent a detoxification process or, on the 64 contrary, lead to the appearance of new and sometimes more toxic compounds for the human 65 consumer (Bailly & Guerre, 2009). 66 Although FB1 is weakly bioavailable after oral ingestion and is mostly excreted in native 67 form in excreta, the bioavailable amount is extensively distributed and accumulates in tissues 68 (EFSA, 2018; Norred, Plattner, & Chamberlain, 1993; Prelusky, Trenhoim, & Savard, 1994). 69 FB1 toxicokinetic studies showed distribution in animal tissues with most of the absorbed 70 toxin found in the liver and kidneys (Martinez-Larranaga et al., 1999; Tardieu, Auby, Bluteau, 71 Bailly, & Guerre, 2008a; Tardieu, Bailly, Skiba, Grosjean, & Guerre, 2008b). One study in 72 weaned piglets also highlighted the presence of partially hydrolysed FB1 in tissues, and found 73 that unmetabolised FB1 was the most abundant form (Fodor et al., 2008). In France, the first 74 75 total diet study (TDS) revealed high FB1 levels in three poultry liver samples (50% of all the poultry livers analysed), with a content between 90 and 120 µg kg⁻¹, suggesting that human 76 77 exposure to FB1 by the ingestion of animal-derived food products should be considered more closely (Leblanc et al., 2005). 78 After oral ingestion, ZEA is quickly absorbed and biotransformed in the digestive tract. ZEA 79 and its associated metabolites are then mostly excreted in urine, faeces or bile, depending on 80 the animal species (Bailly & Guerre, 2009; EFSA, 2011). The principal metabolites known to 81 have affinities for oestrogenic receptors are in the following order: α -zearalanol (α -ZAL) $> \alpha$ -82 zearalenol (α -ZEL) > β -zearalenol (β -ZAL) > zearalenone (ZEA) > β -zearalenol (β -ZEL). 83 Because some of them have higher affinities for oestrogenic receptors than zearalenone, their 84 appearance during metabolic pathways can be considered as a bioactivation (Gaumy, Bailly, 85 & Guerre, 2001). However, only a few studies are available on the carry-over of ZEA and 86 metabolites to edible organs of different animal species (Dänicke et al., 2002; Goyarts, 87

Dänicke, Valenta, & Ueberschär, 2007; Mirocha, Robison, Pawlosky, & Allen, 1982). The 88 89 need for studies on the presence of ZEA and metabolites in animal-derived food was pointed out in a ZEA risk assessment conducted by the French Food Safety Agency, to assess their 90 91 transfer into food products of animal origin (AFSSA, 2006). Because mycotoxins are present at trace levels in animal products, sensitive and accurate 92 analytical methods are required. For ZEA and metabolites, enzyme-linked immunosorbent 93 assay (ELISA), liquid chromatography with fluorescence detection (LC-FLD) or single mass 94 spectrometry detection (LC-MS) have mostly been implemented (Gajecka et al., 2016; Iqbal, 95 Nisar, Asi, & Jinap, 2014; Pleadin et al., 2015; Zielonka et al., 2014, 2015). For analysis of 96 97 FBs in animal tissues, several liquid chromatography-tandem mass spectrometry (LC-MS/MS) methods have been developed in recent years. The combined implementation of LC-98 MS/MS methods with stable isotope dilution assays (SIDA) enabled optimal compensation of 99 100 FB losses at all analytical steps (Cao et al., 2018; Gazzotti et al., 2011; Schertz et al., 2018; Sørensen, Mogensen, & Nielsen, 2010). To our knowledge, this analytical strategy has not 101 102 been applied to the determination of ZEA and metabolites in animal products to date. 103 The aim of this work was to assess the carry-over of FBs, ZEA and metabolites from contaminated feed to chicken liver and muscle to meet the needs expressed by risk assessors. 104 105 This required beforehand the development and validation of a reliable and effective SIDA-LC-MS/MS method able to detect traces of FB1, FB2, ZEA, α-ZAL and α-ZEL in animal 106 tissues. 107

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2. Material and methods

- 110 *2.1. Chemicals and reagents*
- All solutions were prepared with analytical reagent-grade chemicals and ultrapure water (18.2)
- $M\Omega$ cm) obtained by purifying distilled water with a Milli-Q system associated with an Elix 5

pre-system (Millipore S.A., St Quentin-en-Yvelines, France). All standards (solutions, powder 113 and ¹³C-labelled standards) were purchased from Biopure (Tulln, Austria). A mixed stock 114 solution of FB1 and FB2 (500 ng mL⁻¹) was prepared in methanol/water 50/50 from a 50 µg 115 mL⁻¹ FB1 standard solution and a 50 µg mL⁻¹ FB2 standard solution. A ZEA stock solution 116 (100 µg mL⁻¹) was made by dissolving pure standard in acetonitrile. The ZEA concentration 117 was determined by absorption spectrometry ($\varepsilon = 12623 \text{ L mol}^{-1} \text{ cm}^{-1}$). A mixed ZEA, α -118 zearalanol, and α-zearalenol working standard solution (500 ng mL⁻¹ for each) was prepared 119 in methanol from the ZEA stock solution (100 μg mL⁻¹), α-ZAL (10 μg mL⁻¹), and α-ZEL (10 120 µg mL⁻¹) standard solutions. 121 The FB1 and FB2 (500 ng mL⁻¹) working standard solutions and the ZEA, α -ZAL, and α – 122 ZEL (500 ng mL⁻¹) working standard solution were combined to prepare a new mixed 123 working standard solution in water/methanol/formic acid 75/22/3 (v/v/v) (50 ng mL⁻¹ for 124 125 each). This solution was used for the preparation of calibration standards. A uniformly ¹³C-labelled FB1 and FB2 stock standard solution (1200 ng mL⁻¹) was prepared 126 in methanol/water 50/50 (v/v) from U-[$^{13}C_{34}$]-FB1 (25 μg mL $^{-1}$) and U-[$^{13}C_{34}$]-FB2 (10 μg 127 mL⁻¹) internal standard solutions. A uniformly ¹³C-labelled ZEA stock standard solution (880 128 ng mL⁻¹) was prepared in methanol from U-[¹³C₁₈]-ZEA internal standard solution (25 μg mL⁻¹ 129 ¹). ¹³C-labelled stock standard solutions of U-[13 C₃₄]-FB1 and U-[13 C₃₄]-FB2 (1200 ng mL⁻¹) 130 and ZEA (880 ng mL⁻¹) were combined to prepare a new mixed ¹³C-labelled working 131 standard solution in water/methanol/formic acid 75/22/3 (v/v/v) with a concentration of 120 132 ng mL⁻¹ for U-[$^{13}C_{34}$]-FB1 and U-[$^{13}C_{34}$]-FB2, and 88 ng mL⁻¹ for U-[$^{13}C_{18}$]-ZEA. This 133 solution was added to test samples, blank samples and calibration curves. 134 Calibration standards were prepared by diluting the combined mixed working standard 135 solution of FB1, FB2, ZEA, α-ZAL, and α-ZEL (50 ng mL⁻¹ for each) and the mixed ¹³C-136 labelled working standard solutions of U-[¹³C₃₄]-FB1, U-[¹³C₃₄]-FB2 (120 ng mL⁻¹ for both), 137

and U-[$^{13}C_{18}$]-ZEA (88 ng mL $^{-1}$) with water/methanol/formic acid 75/22/3 (v/v/v). The 138 following FB1, FB2, ZEA, α-ZAL and α-ZEL concentrations were obtained: 1.0, 2.0, 3.0 and 139 4.0 ng mL⁻¹. The ¹³C-labelled internal standard concentrations were the same for all the levels 140 $(3.0 \text{ ng mL}^{-1} \text{ for U-}[^{13}\text{C}_{34}]\text{-FB1} \text{ and U-}[^{13}\text{C}_{34}]\text{-FB2} \text{ and } 2.2 \text{ ng mL}^{-1} \text{ for U-}[^{13}\text{C}_{18}]\text{-ZEA}).$ 141 Because no stable isotope labelled standards are available for ZEA metabolites, the U-[¹³C₁₈]-142 ZEA was used as internal standard for the quantification of α -ZAL and α -ZEL. HPLC grade 143 acetonitrile, methanol and acetic acid were purchased from Fisher Scientific (Loughborough, 144 UK). Formic acid (Fisher Scientific) and ammonium formate (Alfa Aesar, Karlsruhe, 145 Germany) were of analytical grade. Phosphate buffered saline (PBS) tablets were purchased 146 from R-Biopharm AG (Darmstadt, Germany). PBS-buffer (pH 7.4) solution was prepared by 147 dissolving one tablet in 100 ml. In addition to calibration standard preparation, 148 water/methanol/formic acid 75/22/3 (v/v/v) was used to adjust the final volume before LC-149 150 MS/MS injection. Ultra-pure grade carrier argon (Ar, 99.9999% pure) and nitrogen (N₂, 99.999% pure) were purchased from Linde Gas (Montereau-Fault-Yonne, France). 151

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2.2. Preparation of feed

The composition and nutritional characteristics of diets distributed to chickens is indicated in 154 the Table SD1. First, uncontaminated starting feed was given over the period 0-27 days. Then 155 growing-finishing feed contaminated with raw materials provided by the National Veterinary 156 School of Toulouse (France) was given over the period 28-84 days. 157 The levels of the contamination of mycotoxins in feed were defined to approach the guidance 158 levels set by the European Union (2006/576/EC) relative to a feedingstuff with a moisture 159 content of 12% for calves, dairy cattle, sheep (including lambs) and goats (including kids) 160 (0.5 mg ZEA kg⁻¹) and for poultry, calves (<4 months), lambs and kids (20 mg FBs kg⁻¹) 161 (European Union, 2006). Contaminated raw materials of 44.62 g of rice flour (containing 650 162

mg ZEA kg⁻¹) and 138 g of corn flour (containing 6800 mg FB1 kg⁻¹ and 1600 mg FB2 kg⁻¹) 163 164 were mixed individually with uncontaminated feed (58 kg; Table SD1). The rice or corn flour were progressively mixed with the feed first in a mixer bowl (Hobart, Croissy Beaubourg, 165 France) containing 10 kg of feed then with a bigger feed mill mixer (Gondard, CTS, Le 166 Pontet, France) able to contain 400 kg feed. This mixture was then pelleted (diameter 2.5 mm) 167 with a feed mill pelletizer without using steam (CPM, CPM Europe B.V., Zaadam, 168 Netherlands). Special precautions had been taken to ensure the homogeneity of the 169 contaminated feed. The introduction of these small quantities of flour into the control feed 170 was not likely to significantly modify the feed value. The levels of contamination of the final 171 feed were measured by liquid chromatography with fluorescence detection (LC-FLD). 172

- For ZEA, the feed level was determined following the NF EN 15792 standard. The ZEA level
- measured was 0.40 ± 0.12 mg kg⁻¹, which represents 80% of the expected value.
- For FBs, levels were determined following the NF EN 16006 standard. The FB1 and FB2
- level measured was 12.7 ± 3.3 mg kg⁻¹, i.e. 64% of the expected value. FB1 represented 83%
- of the sum (10.5 mg kg $^{-1}$) and FB2, 17% (2.2 mg kg $^{-1}$).
- 178 Control feed was also analysed. ZEA was quantified at a level of 0.027 ± 0.009 mg kg⁻¹. This
- value is 15 times lower than the level measured in the ZEA contaminated feed. Neither FB1
- nor FB2 were detected (LOD: 0.02 mg kg⁻¹ for FB1 and 0.03 mg kg⁻¹ for FB2).

182 *2.3. Experimental design*

- 183 The experimental design involved 24 male chickens (genus Gallus slow growth rate strain
- JA 657) with a weight of 35 ± 3 g, purchased from Boyé Accouvage (La Boissière-en-Gatîne,
- France). All the chickens were vaccinated against Marek's disease and infectious bronchitis.
- The light duration was set to 23 hours daily and the temperature to 30-31 °C up to the age of 3
- days. These two parameters were then gradually decreased to reach 18 hours light daily and a

temperature of 19-20 °C from the age of 21 days. During the whole rearing period, water was always available. Until the age of 28 days, the chickens were reared together in the same room on wood shavings litter. Starter feed was distributed *ad libitum*. During the following 8 weeks of the exposure period (28 to 84 days of age), chickens of 28 days of age were distributed into 3 groups presenting the same average body weight after a fasting period of 12 hours. There were 15 chickens in the control group and 9 chickens in each mycotoxins treated groups. The first one was treated with ZEA and the second with FBs.

The animals were distributed into individual cages allowing monitoring of feed consumption quantities. The grower feed was used until slaughter. It was distributed daily, at 68, 80, 87, 98, 106, 111, 126 and 153 g per day, respectively during weeks 5, 6, 7, 8, 9, 10, 11 and 12, according to breeder recommendations. Feed refusals were collected individually and weighed weekly to calculate the feed intake.

None of the treatments induced systematic signs of toxicity. However, one chicken died during the 5th week in the ZEA and metabolites treated group, reducing the number of individuals to eight. At the age of 84 days, after a 12 hour fasting period, all chickens were weighed and slaughtered by electronarcosis, followed by bleeding.

2.4. Sample preparation

Since previous studies have shown that the liver is more strongly contaminated than muscle (Mirocha et al., 1982; Tardieu, Auby, et al., 2008a; Tardieu, Bailly, et al., 2008b; Völkel et al., 2011), the analytical strategy implemented consisted in first analysing the livers and if their levels were higher than the LOQ, the muscles belonging to the same animals were also analysed.

2.5. Extraction 213

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For the extraction process, 250 µL of ¹³C-labelled combined working standard solution were added to the test portion of 2.50 ± 0.01 g of muscle or liver sample placed into a 50 mL tube. The tube content was mixed using a vortex mixer. 9.5 mL methanol/water 80/20 (v/v) were 216 added and the samples were homogenised at 10 000 ± 500 rpm for 2 min using a Polytron® 217 (Kinematica AG, Luzern, Switzerland). After centrifugation at 9 000 g for 10 min at 3 °C, the 218 supernatant was placed in a volumetric flask of 20 mL. The extraction of mycotoxins from the 219 220 test portion with 9.5 mL methanol/water 80/20 (v/v), the centrifugation step, and the supernatant transfer were repeated once. The solution was adjusted to 20 mL and filtered 221 using a 0.20 µm polyester (PET) syringe filter (Chromafil®, Macherey-Nagel GmbH & Co. KG, Düren, Germany). 223

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- 2.6. Hexane washing of the extract 225
- Before FB clean-up, hexane washing of the extract was necessary. 5 mL of extract were added 226
- 227 to a 15 mL falcon tube and 5 mL of hexane were added. The tube was vortex mixed for 30 s.
- After centrifugation at 3 000 g for 3 min at 20 °C, the upper hexane phase was removed and 228
- the hexane washing was repeated once. 229

- 2.7. Clean-up 231
- Two parallel immunoaffinity clean-up steps were implemented. Fumoniprep[®] immunoaffinity 232
- clean-up (R-Biopharm AG, Darmstadt, Germany) was specific for FBs, and Easi-extract® 233
- Zearalenone immunoaffinity clean-up (R-Biopharm AG, Darmstadt, Germany) was specific 234
- for ZEA and metabolites. Both for FBs and for ZEA and metabolites, 8 mL of PBS were 235
- added to 2 mL of the hexane washed extract in a 15 mL Falcon tube. These PBS solutions 236
- were slowly (drop by drop) passed through their respective immunoaffinity columns, then 237

placed in a vacuum manifold. For FBs, the column was washed with 10 mL of PBS and for

ZEA and metabolites, the column was washed with 20 mL of PBS. Both columns were then

dried by pushing air through them with a syringe. Mycotoxins were finally eluted by applying

241 the backflushing technique with 3.0 mL methanol/acetic acid 98/2 (v/v).

The eluates were evaporated at 50 ± 5 °C under a gentle stream of nitrogen to 0.1 mL and

adjusted to 1 mL with water/methanol/formic acid 75/22/3 (v/v/v). The final extracts were

mixed using the vortex mixer, filtered (PET, 20 µm), and analysed by UHPLC-MS/MS.

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2.8. LC conditions

The LC system was an Accela 1250 (Thermo Fisher Scientific, San Jose, CA, USA). Ultrahigh performance liquid chromatographic (UHPLC) separation of mycotoxins was performed using a column with particles composed of a solid core and a porous shell. A Kinetex[®] C18 column (100 Å, 2.6 µm particle size, 50 x 2.1 mm) equipped with a Kinetex C18 security guard cartridge (Phenomenex, Torrance, CA, USA) was used. For the FBs analysis, eluent A was composed of water/formic acid 99.9/0.1 (v/v) and eluent B of methanol/water/formic acid 94.9/5/0.1 (v/v/v). Both eluents contained 0.5 mmol/L ammonium formate. For the ZEA and metabolites analysis, eluent A was composed of water and eluent B of methanol/water 95/5 (v/v). Both gradients were the same and programmed as follows: 25% B (initial), 25-100% B (6.4) min), 100% B (hold 1.1 min), 100-25% B (0.5 min), 25% B (hold 1 min). The column effluent was transferred via a divert valve (Rheodyne, USA) either to the mass spectrometer (between 1.5 and 5.5 min) or to waste. The total flow rate was 0.5 mL min⁻¹, while the injection volume was 10 µL. The column temperature was maintained at 30 °C. A chromatogram of a blank matrix spiked at the quantification limit for all mycotoxins (0.1 µg kg⁻¹) is presented in **Fig. 1.**

2.9. MS/MS conditions

Detection was performed with a TSQ Vantage triple quadrupole mass spectrometer (Thermo Fisher Scientific), equipped with an ElectroSpray Ionisation (ESI) source (HESI-II probe). The mass spectrometer was operated in Selected Reaction Monitoring (SRM). The spray voltage was -2625 V in negative mode and + 4000 V in positive. The source temperature was set at 500 °C and capillary temperature at 350 °C. Nitrogen was used as the nebulising gas with a sheath gas pressure of 50 (arbitrary unit) and an auxiliary gas pressure of 18 (arbitrary unit). The collision gas was argon, with a gas pressure of 1.5 mTorr. One transition was used for quantification (Q) and another as qualifier transition (q). The optimised SRM parameters by injection of each compound are listed in Table 1. A mass resolution of 0.7 Da full width at half maximum (FWHD) was set on the first (Q1) and the third (Q3) quadrupoles. Instrument control and data were handled by a computer equipped with TSQ Tune Master version 2.3.0, Xcalibur version 2.1.0 and TraceFinder version 1.0.1 (Thermo Fisher Scientific).

2.10. Method validation methodology

The linearity of the method was statistically evaluated by the Fisher-Snedecor test with a significance level $\alpha=0.01$. The other performance characteristics were validated according to the accuracy profile approach. This procedure summarises every validation element on a single plot, giving a graphical representation of the method's performance. The validity domain is defined between the lowest and the highest tested concentrations, with tolerance limits (β -expectation limits) between the acceptance limits (λ). Tolerance limits were calculated at each concentration level and take into account the bias, the repeatability, and the intermediate precision. The limit of quantification (LOQ) was defined as the lowest concentration level validated.

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2.11. Quality control

To ensure reliable results, samples were analysed in batches including several internal quality controls (IOCs). When acceptance criteria were not met, results were discarded and samples were re-analysed. A blank matrix was analysed in the same conditions as for all samples, to check the absence of any contamination. The concentration had to be below the detection limit. A blank matrix spiked at the quantification limit (1 µg kg⁻¹ for all the mycotoxins) was prepared and analysed in the same conditions as for all the samples. Recoveries had to be between 70 and 130%. All mycotoxins were quantified using bracketing calibration curves. The determination coefficient (r^2) of the calibration curve had to be ≥ 0.98 and the slope variation between two sets of bracketing calibration curves had to be below 15%. Variation of the retention time in samples had to be below 5% in comparison to the standard retention time. Before use, the capacity of each batch of immunoaffinity columns was checked. For fumonisins, 200 ng of FB1 and 200 ng of FB2 were deposited at the same time on the Fumoniprep® immunoaffinity columns, the capacity had to be higher than 100 ng for each mycotoxin. The same quantities of ZEA, α -ZAL and α -ZEL were deposited at the same time on the Easi-extract® Zearalenone immunoaffinity column. The capacity also had to be higher than 100 ng for each.

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2.12. Statistical data analysis

308 Statistical data analyses were performed using R studio software version 1.0.143 (R Studio,

309 Boston, USA).

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3. Results and discussion

3.1. Method validation

For each mycotoxin, an accuracy profile was built following the NF V03-110 Standard (AFNOR, 2010; Mermet & Granier, 2012). The main advantage of this approach is the robustness of the validation methodology. It included a range of 3 concentration levels (1.0, 4.0 and 10.0 µg kg⁻¹) and 6 series repeated on different days over a period of 1 month for ZEA and metabolites, and 4 months for FBs. For each series, 2 replicates of each of the 3 concentration levels were analysed in bracketing with five calibration standards to establish the response function. Spiked samples followed the whole analytical procedure, including extraction. The probability β was set to 80%, meaning that the risk of results falling outside β expectation tolerance intervals was below 20% on average. The acceptance limits (λ) were set at \pm 25%. For all the concentration levels, β -expectation tolerance intervals were within the acceptability limits. The accuracy profile and the performance criteria obtained for each mycotoxin are presented in Table 2, and as an example, the FB1 accuracy profile is presented in Fig. 2. The method performances in terms of trueness and precision were very satisfactory. Recoveries ranged from 92% to 107% for all toxins. The repeatability coefficient of variation, CVr, varied from 4.3% to 10.9% on the validity domain, and the intermediate precision coefficient of variation, CV_{IP}, was between 4.3% and 13.1%. Because mycotoxins are present at trace levels in animal matrices, sensitivity was the key parameter of the method. The combined use of immunoaffinity purification, SIDA and UHPLC allowed us to achieve an LOQ of 1.0 µg kg⁻¹ for all mycotoxins studied. To the best of our knowledge, the sensitivity of the FB method is the best observed among the existing methods for meat products (Fodor et al., 2008; Gazzotti et al., 2011; Guillamont et al., 2005; Meyer et al., 2003; Sørensen et al., 2010; Szabó-Fodor et al., 2015; Tardieu, Auby, et al., 2008a). For ZEA, method sensitivity is equal to that obtained with several other methods (Fang, Chen, & Guo, 2002; Goyarts et al., 2007; Yan et al., 2018), and is very close to the most sensitive ones (0.4 to 0.5 µg kg⁻¹) (Chen et al., 2013; Pleadin et al., 2015; Zöllner et al., 2002). Importantly, in this study, the LOQ was

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assessed by a very robust method and was defined as the lowest validated level. Finally, the validity domain of the analytical method ranged from 1.0 to 10.0 μ g kg⁻¹ for ZEA, α -ZAL and α -ZEL and from 1.0 to 16.0 μ g kg⁻¹ for FB1 and FB2.

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3.2. Carry-over of ZEA to the liver

ZEA, α -ZAL and α -ZEL levels were assessed in the 15 control samples and the 8 samples of the ZEA treated group. ZEA and α-ZAL were not detected in any samples from the control or treated groups (Table 3). α-ZEL was not detected in control samples, but levels between the limit of detection (0.3 µg kg⁻¹) and the limit of quantification (1.0 µg kg⁻¹) were found in 5 out of 8 samples in the ZEA treated group, with concentrations estimated to be in the range 0.4 – 0.8 µg kg⁻¹. The non-detection of zearalenone could be explained by the fact that ZEA is largely and rapidly eliminated in excreta (Mirocha et al., 1982). Therefore, in this study, excreta were collected during the last two days before animal slaughter, and ZEA was analysed by applying the NF EN 15792 Standard. A level of 0.270 ± 0.090 mg kg⁻¹ was quantified for the chickens exposed to the contaminated diet, whereas a level 23 times lower was measured for the chickens exposed to the control feed $(0.012 \pm 0.006 \text{ mg kg}^{-1})$. These levels confirm the considerable elimination of ZEA in excreta. This mycotoxin is also rapidly metabolised by three major routes in mammals: hydroxylation resulting, among others, in the formation of α-ZEL; glucuronidation in the small intestine and the liver; and cytochrome P450-mediated oxidation which produces catechol metabolites (EFSA, 2011). The detection of α -ZEL in several samples supports this metabolisation process. Dänicke et al. (2002) also studied the carry-over of ZEA to laying hen tissues following hen exposure to ZEA feed with 4 times higher contamination than in the current study (1.6 mg kg⁻¹ ¹). Without β-glucuronidase and aryl-sulfatase treatment of the samples, and when no Mycofix®Plus was added to the feed, similarly to our results, ZEA was not detected in breast

meat or in the liver, whether α -ZEL was quantified in livers. To quantify the extent of carry-over, a carry-over factor (CF) was calculated to measure the degree of release of the mycotoxin from the feed matrix in the digestive tract, and its absorption, distribution metabolisation and elimination (**Equation (1)**).

Carry-over factor (CF) = Toxin level in tissue ($\mu g kg^{-1}$) / Toxin level in diet ($\mu g kg^{-1}$) (1)

Since our results in tissues were below the limit of quantification, it was not possible to calculate a CF; consequently, we calculated this factor between α -ZEL concentrations in the liver and ZEA concentrations in feed from the Dänicke et al. (2002) study. This CF amounted to 0.0008 (1.3 µg kg⁻¹ in liver divided by 1580 µg kg⁻¹ in feed). When we applied this carry-over factor to the level of ZEA in the contaminated feed used in this study, we obtained an estimated α -ZEL level in the liver of 0.3 µg kg⁻¹ (400 µg kg⁻¹ x 0.0008). This value is equal to the LOD (0.3 µg kg⁻¹) of the present method. As previously described, among the obtained results, 5 out of 8 were between the LOD and the LOQ (1.0 µg kg⁻¹) and the 3 others were less than or equal to the LOD. Therefore, these results are consistent with those reported by Dänicke et al. (2002).

- 3.3. Carry-over of FBs to the liver
- FB1 and FB2 levels were assessed in the 15 control samples and in the 9 samples of the FB treated group. These mycotoxins were not detected in the control samples (LOD = $0.3 \mu g \, kg^{-1}$) (Table 4). FB1 was quantified in all the 9 samples of the FB treated group with a mean level of $30.3 \pm 14.8 \, \mu g \, kg^{-1}$ and a median of 25.8 $\mu g \, kg^{-1}$ (minimum and maximum of 17.9 and 65.5 $\mu g \, kg^{-1}$) (Fig. 3). FB2 was also quantified in all the samples (mean: $2.3 \pm 1.1 \, \mu g \, kg^{-1}$); median 2.1 $\mu g \, kg^{-1}$), with levels ranging from 1.2 to 4.9 $\mu g \, kg^{-1}$ (Fig. 3). The mean level

observed in the liver, 13 times lower for FB2 compared to FB1, may be partially explained by the 5 times lower concentration in feed. Samples with high levels of FB1 also had high levels of FB2. In order to check whether there is a correlation between the FB1 and FB2 levels measured, a Pearson statistical test with a significance level $\alpha = 5\%$ was applied. A significant correlation was found (p value: < 0.001; y = 12.8 x + 0.9; R = 0.97) (Fig. 4). This correlation suggests that even though there was a wide difference in the levels measured between individuals (CV = 46% for FB1 and 43% for FB2), the carry-over of both toxins seems to be very similar between animals. Liver carry-over factors were calculated and compared with previous studies. For FB1, CF was 0.003. This value is very close to the CF of 0.002 reported by Del Bianchi, Oliveira, Albuquerque, Guerra, and Correa (2005) in broiler chickens, after prolonged FB1 oral administration (10 mg kg⁻¹ of feed) from 21 to 41 days of age. Tardieu et al. (2008b) also studied FB1 persistence in poultry tissues after exposure to a diet containing fumonisins at 5, 10 and 20 mg FB1 + FB2 kg⁻¹. After an interval of 8 h between the last ingestion of feed and slaughter of the animals, FB1 concentrations were measured in liver tissues. CFs were slightly higher than the values observed in chickens, with values ranging between 0.004 and 0.007 for the three levels tested. Moreover, with feed less than two times more contaminated, FB levels measured in the liver were about four times higher than the levels measured in the present work, Tardieu et al. (2008b) maintained an interval of 8 hours between the last ingestion of feed and slaughter, whereas in the present study, the interval was 12 hours. Therefore, the difference in terms of fasting period duration could explain the difference in the results obtained. The implementation of a fasting period is a practice also carried out in slaughterhouses to improve the safety of animal products by reducing the incidence of carcass contamination due to rupture or laceration of the intestines, intestinal overflow, or faecal discharge from the rectum.

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To our knowledge, FB2 carry over has so far never been reported in animal tissues. In the 413 present study, we observed a CF three times lower than for FB1 (CF = 0.001 vs. 0.003). 414 Considering the highest FB1 and FB2 concentrations in livers measured in the present study 415 (70.4 µg FB1 + FB2 kg⁻¹), a person of 70 kg would have to eat 1.0 kg per day of poultry liver 416 to exceed the FB tolerable daily intake (TDI) of 1.0 µg kg⁻¹ bw, and 2.3 kg per day of poultry 417 liver for the mean value measured (32.6 µg FB1 + FB2 kg⁻¹) (EFSA, 2018). It is important to 418 note that the levels measured in feed during the experiment reached only 64% of the FB 419 guidance level defined for poultry. If consumption data were available for poultry livers, it 420 could be interesting for this feed contamination level to measure the relative contribution of 421 poultry liver consumption to the total exposure of humans to FBs. 422 In the same way as for ZEA, excretion via the excreta is considered the main route of 423 elimination of FBs (Bailly & Guerre, 2009). In the present study, FB excreta analyses were 424 carried out following the NF EN 16006 Standard. An FB excreta level of 15.2 mg kg⁻¹ (FB1: 425 13.0 ± 3.9 mg kg⁻¹, FB2: 2.2 ± 0.7 mg kg⁻¹) was measured for chickens exposed to the 426 contaminated diet and a level of 0.065 mg kg⁻¹ (FB1: 0.065 ± 0.033 mg kg⁻¹, FB2 ≤ 0.03 mg 427 kg⁻¹) was found for chickens exposed to the control feed. The comparison of these levels with 428 the levels measured in the contaminated feed points to significant elimination of FBs in 429 430 excreta.

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livers

Given that FBs were quantified in all livers in the FB treated group, the chicken muscles belonging to the same animals were also analysed. Results showed that for each sample "liver – muscle" pair, the FB1 and FB2 concentrations were systematically lower in the muscle than

in the liver (Table 4). On average, the FB1 levels measured in meat were 16 times lower than

3.4. Determination of FBs in chicken fillets and comparison with the concentration found in

in the liver, and no significant correlation was highlighted between the levels measured for both tissues (R = 0.08). In a previous study, Tardieu et al. (2008a) studied the FB1 contamination of duck tissues after a single oral dose of 5 and 40 mg kg⁻¹ body weight. Liver levels were between 5 and 18 times higher than muscle levels. Considering the differences in the experimental protocols implemented and in the avian species studied, these results compare well with results presented here. In another study, Tardieu et al. (2008b) did not detect FB1 in muscles of turkeys fed with 20 mg FB1 + FB2 kg⁻¹ feed for 9 weeks, whereas FB1 levels in livers of the same animals reached a mean value of 117 μ g kg⁻¹. The absence of FB1 in meat in this former study and its detection in the present work could be explained by the large difference in terms of method sensitivity. The limit of detection of the method presented here is about 33 times lower. Considering the low levels measured in muscles (all < 5.0 μ g kg⁻¹), high sensitivity was essential to be able to detect and quantify FB1. FB2 was detected but not quantified in one sample and was < LOD for the other eight samples.

4. Conclusions

A sensitive and accurate SIDA-UHPLC-MS/MS method using immunoaffinity clean-up was developed, validated and implemented to study the carry-over of FBs and ZEA and metabolites to meat products after exposure of broiler chickens to contaminated feed. The analysis of livers from chickens exposed to a contaminated diet revealed the presence of α -ZEL in several samples. The presence of this compound, more oestrogenic than ZEA, and the absence of ZEA indicate excreta elimination and metabolisation processes of ZEA by hydroxylation. FB1 and FB2, which have chronic toxicity in humans, were quantified in livers and muscles. From these results, it was possible to calculate carry-over factors from feed to poultry livers. Finally, FB levels in muscle were also investigated and compared with the levels measured in the livers belonging to the same animals. These new carry-over data will

contribute to a better understanding of mycotoxin transfer from feed to meat products. Considering the FB levels measured in poultry tissues, this study could be useful to assess whether the current FB guidance value for poultry is sufficiently protective for consumers. Moreover, given the high excreta elimination of FBs and ZEA, the duration of fasting could be considered an effective parameter for animal product decontamination and deserves further investigation. In future studies, it could also be interesting to adopt this type of approach for the other analogues of FB (FB3 to FB6) and their modified forms, to respond to the recent call for additional data expressed by the European Food Safety Agency (EFSA).

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Declarations of interest

477 None.

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Figure Captions

- 484 Fig. 1: Chromatogram obtained for a chicken liver sample spiked at the quantification limit
- 485 (1.0 μg kg⁻¹ for all mycotoxins): a) quantification transition (Q); b) qualifier transition (q).
- 486 **Fig. 2:** Accuracy profile of FB1 ($\beta = 80\%$; $\lambda = \pm 25\%$).

- Fig. 3: Box Plot of FB1 and FB2 obtained from levels measured in chicken livers from the FB
- 488 treated group.
- 489 Fig. 4: Correlation between FB1 and FB2 levels measured in chicken livers from the FB
- treated group. The grey shaded area represents the confidence interval.

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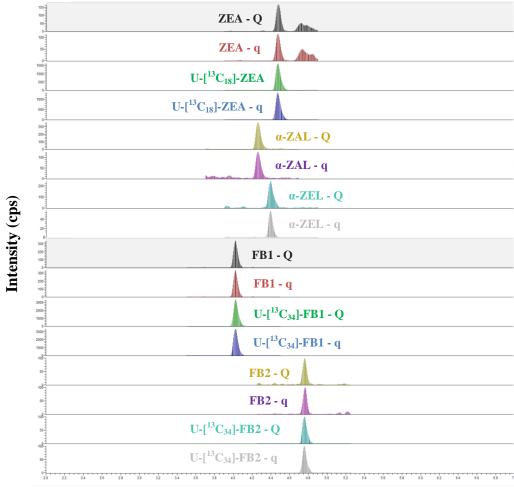
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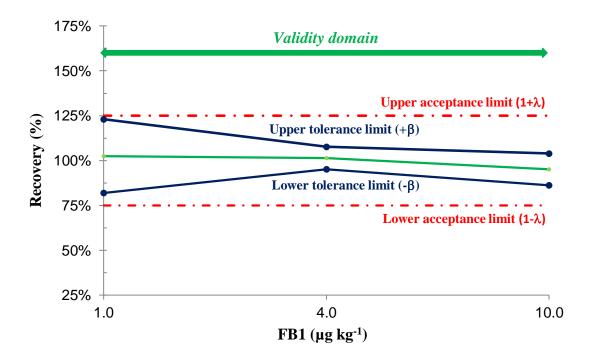
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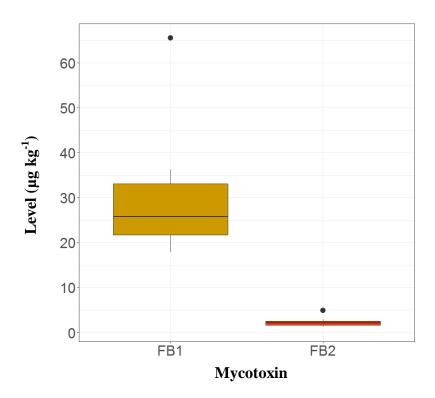
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Retention Time (min)





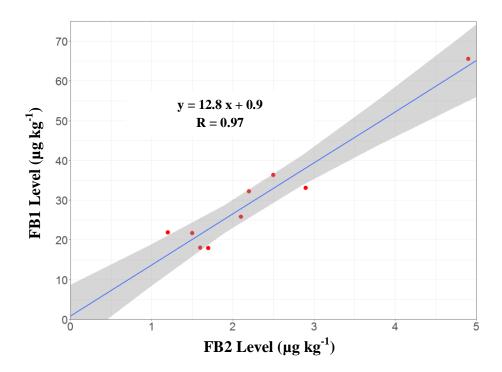


Table 1: Optimised MS/MS parameters

Compounds	Ionisation mode	S-lens voltage (V)	Precursor ion (m/z)	Collision energy (V)	Product ion (m/z)
FB1	ESI+	142	722.4	38	334.3 (Q)
		137		34	352.3 (q)
$U-[^{13}C_{34}]$ -FB1	ESI+	142	756.4	34	374.3 (Q)
		137		36	356.3 (q)
FB2	ESI+	142	706.4	34	336.3 (Q)
		137		36	318.3 (q)
U-[13 C $_{34}$]-FB2	ESI+	142	740.4	34	358.3 (Q)
		137		36	340.3 (q)
ZEA	ESI-	117	317.1	24	175.0 (Q)
		117		30	131.1 (q)
$U-[^{13}C_{18}]$ -ZEA	ESI-	55	335.1	26	185.1 (Q)
		55		32	140.1 (q)
α-ZAL	ESI-	118	321.2	23	277.2 (Q)
		118		22	303.2 (q)
α-ZEL	ESI-	118	319.1	21	277.2 (Q)
		118		26	303.2 (q)

Table 2: Method performance criteria

		LOD (µg kg ⁻¹)	LOQ (µg kg ⁻¹)	Accuracy			
				Trueness	Prec	Precision	
	Levels (µg kg ⁻¹)			Recovery (%)	Repeatability (% RSD)	Intermediate precision (% RSD)	
	1.0	0.3	1.0	102	6.1	13.1	
ZEA	4.0			101	4.3	4.3	
	10.0			95	5.8	6.5	
	1.0	0.3	1.0	102	5.8	10.1	
α - ZAL	4.0			105	6.7	11.7	
	10.0			101	3.8	9.6	
	1.0	0.3	1.0	107	8.9	9.9	
α - ZEL	4.0			96	5.5	7.0	
	10.0			92	5.2	5.6	
	1.0	0.3	1.0	99	6.5	12.2	
FB1	4.0			95	7.8	10.6	
	16.0			102	6.2	10.3	
	1.0	0.3	1.0	103	7.9	9.4	
FB2	4.0			94	10.9	10.9	
	16.0			102	10.1	10.1	

Table 3: Results for ZEA and its associated metabolites in chicken livers in the ZEA treated

group

	Animal _ Number	Levels in liver (µg kg ⁻¹)			
		ZEA	α-ZAL	α-ZEL	
	787	\leq LOD ^a	\leq LOD ^a	\leq LOD ^a	
	800	\leq LOD ^a	\leq LOD ^a	\leq LOQ $(0.4)^{b}$	
77 E A	811	\leq LOD ^a	\leq LOD ^a	\leq LOQ $(0.8)^{b}$	
ZEA	863	\leq LOD ^a	\leq LOD ^a	\leq LOQ $(0.5)^{b}$	
treated	877	\leq LOD ^a	\leq LOD ^a	\leq LOD ^a	
group	895	\leq LOD ^a	\leq LOD ^a	\leq LOD ^a	
	908	\leq LOD ^a	\leq LOD ^a	\leq LOQ $(0.4)^{b}$	
	914	\leq LOD ^a	\leq LOD ^a	\leq LOQ $(0.6)^{b}$	

a: LOD = 0.3 µg kg⁻¹
b: LOQ = 1.0 µg kg⁻¹; results comprised between LOD and LOQ were calculated and are presented in brackets.

Table 4: Results for FBs in chicken livers and muscle in the FB treated group

	Animal Number	Levels in liver (μg kg ⁻¹)		Levels in muscle (μg kg ⁻¹)		
		FB1	FB2	FB1	FB2	
	821	36.3	2.5	1.3	≤ LOD ^a	
	846	32.2	2.2	1.3	\leq LOD ^a	
	855	18.0	1.6	\leq LOQ $(0.5)^{b}$	\leq LOD ^a	
ED 44-1	860	17.9	1.7	1.2	\leq LOD ^a	
FB treated	861	65.5	4.9	1.3	\leq LOD ^a	
group	872	33.1	2.9	2.4	\leq LOD ^a	
	886	21.9	1.2	1.1	\leq LOD ^a	
	897	25.8	2.1	4.7	\leq LOQ $(0.5)^{b}$	
	918	21.7	1.5	2.9	\leq LOD ^a	

a: LOD = 0.3 μ g kg⁻¹; results comprised between LOD and LOQ were calculated and are presented

13 14 15 in brackets