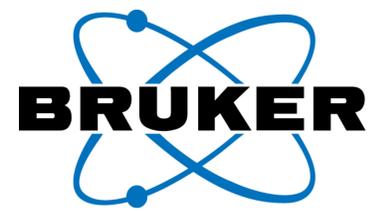


Mycotoxins – analyzing an interesting and challenging group



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D. Brombach¹, L. Klöhn², O. Grundmann², S. Bodendiek¹

¹ Bruker Daltonik GmbH, Fahrenheitstraße 4, 28359 Bremen, Germany

² Impetus GmbH & Co. Bioscience KG, Fischkai 1, 27572 Bremerhaven, Germany

Introduction:

Mycotoxins are a class of compounds that are produced by mold. Due to their high toxicity to vertebrates it is important to monitor the concentrations of mycotoxins in food and feed. Over a hundred mycotoxins are known with Aflatoxins (B1, B2, G1, G2), Zearalenon, DON, T2-Toxin, HT2-Toxin, Ochratoxin A and Fumonisin being the ones of the highest concern.

Here we report a fast, easy and reliable method with the option of screening an unlimited number of mycotoxins in general. We report results for the most important mycotoxins showing that QTOF MS is a very sensitive technique to cover a broad range of compound classes of interest for routine analysis.

Methods:

- Maize samples from a proficiency test were prepared after a QuEChERS method.
- Multistep gradient, runtime 20 min, A (water/ MeOH, 90:10) with 5 mM NH₄HCO₂ and 0.01% formic acid, B (MeOH) with 5 mM NH₄HCO₂ and 0.01% formic acid.
- Analysis were performed using an UltiMate 3000 (Thermo, Dionex) LC system coupled to a compact QTOF mass spectrometer (Bruker Daltonics).
- MS Data acquired from 50-1000 m/z in alternating full-scan and broad-band CID acquisition mode at 2 Hz in positive ionization mode.
- Calibration curves were prepared in solvent A (with ¹³C labeled internal standards). Samples were spiked with ¹³C labeled internal standards to compensate for matrix effects.
- Processing was done in TASQ 1.4 (Bruker Daltonics).

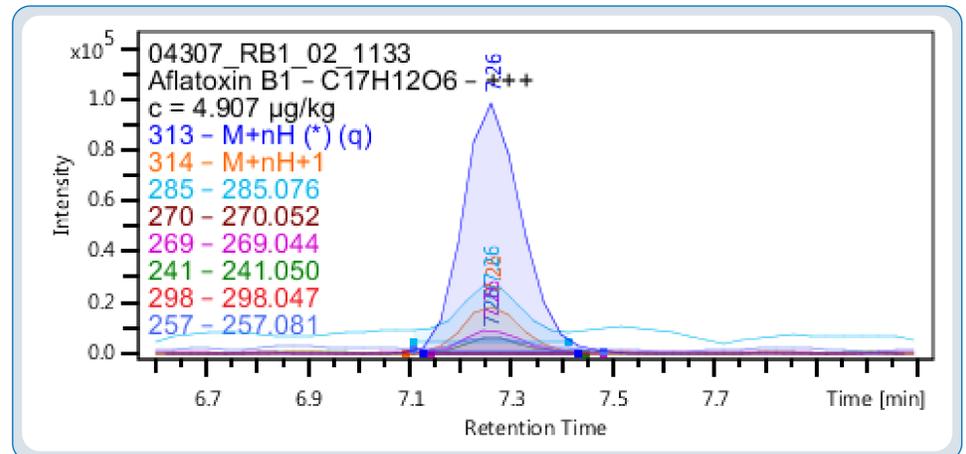


Fig. 2 Chromatogram traces for Aflatoxin B1 of a representative proficiency test sample. It can be clearly seen that besides the principal ion ([M+H]⁺ in dark blue) the different qualifier ions (all other colors) can be clearly seen. A concentration of 4.91 µg/kg suits perfectly to the spiked value of 4.81 µg/kg.

Results:

- The herein described methods are based on the complete solution PesticideScreener 2.1 (Bruker Daltonics). Even though the method is optimized for pesticides it could be perfectly transferred to mycotoxins applying only slight changes (e.g. collision energy). The achieved linear ranges, LOQs and LODs were shown to be suitable for the regulated concentrations.
- Calibration curves were measured in solvent A. To avoid any matrix effects, a fully ¹³C labeled internal standard for each mycotoxin was spiked.
- All mycotoxins could be quantified using a linear calibration curve fit with R² values > 0.9987.
- Fig. 1 shows a calibration curve for Aflatoxin B1. A linear fit for 0.25-50 µg/kg shows a high R² value.
- Fig. 2 shows the chromatogram traces in the proficiency maize test sample for Aflatoxin B1. It can be clearly seen, that the principal ion ([M+H]⁺) was nicely detected and verified by a number of different fragments (qualifier ions).
- Aflatoxin B1 was spiked with 4.81 µg/kg in the sample. Quantitation of this sample resulted in a concentration of 4.91 µg/kg which is in good alignment to the expected value (accuracy: 102%).

- All other mycotoxins were tested and quantified with good accuracies during the proficiency test: Aflatoxin G2 (0.3 µg/kg), DON (1195.2 µg/kg, 104%), Ochratoxin (2.2 µg/kg, 72%) and ZEN (194.8 µg/kg, 105%).



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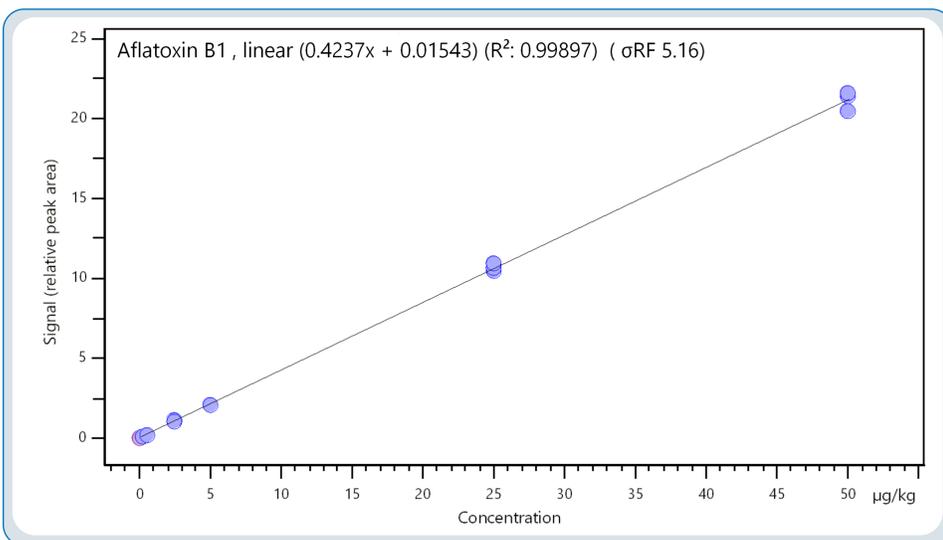


Fig. 1 Calibration curve for Aflatoxin B1 of triplicate measurements from 0.25-50 µg/kg in solvent A.

Mykotoxin	LOD µg/kg	LOQ µg/kg	R2	Linear range µg/kg
Aflatoxin B1	<0.25	0.25	0.99897	0.25-5
Aflatoxin B2	<0.25	0.25	0.99961	0.25-50
Aflatoxin G1	<0.25	0.25	0.99885	0.25-50
Aflatoxin G2	<0.25	0.25	0.99992	0.25-25
DON	0.25	0.50	0.99960	0.5-1000
HT2-Toxin	<0.25	2.50	0.99978	2.5-250
Ochratoxin A	0.5	0.50	0.99876	0.5-2500
T2-Toxin	1	2.00	0.99973	2-2000
Zearalenone	<10	10.00	0.99976	10-2000

Table 1: Result summary of the detected and quantified mycotoxins using TASQ 1.4 (Bruker Daltonics). All calibration curves show a linear fit within a 20% precision.

Conclusions:

- All mycotoxins can be identified and quantified down to concentrations that are regulated by authorities.
- Some mycotoxins were identified and quantified down to 0.25 µg/kg with high accuracies.
- Fully ¹³C labeled internal standards compensate for matrix effects. Therefore, no matrix matched calibration is needed which simplifies the routine work, tremendously



hr-QTOF-MS