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## Mixtures of mycotoxins, phytoestrogens and pesticides co-occurring in wet spent brewery grains (BSG) intended for dairy cattle feeding in Austria

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### ABSTRACT

Spent brewery grains (BSG) are the main by-product of beer production and are incorporated in rations of food-delivering animals, mainly dairy cows. Like other agricultural commodities, BSG can be contaminated by a broad spectrum of natural and synthetic undesirable substances, which can be hazardous to animal and human health as well as to the environment. The co-occurrence of mycotoxins, phytoestrogens, other fungal and plant secondary metabolites, along with pesticides, was investigated in 21 BSG samples collected in dairy farms in Austria. For this purpose, a validated multi-metabolite liquid chromatography/electrospray ionisation tandem mass spectrometry (LC/ESI-MS/MS) was employed. Metabolites derived from *Fusarium*, *Aspergillus*, *Alternaria* and pesticide residues, were ubiquitous in the samples. Zearalenone (ZEN), T-2 and HT-2 toxins were the only regulated mycotoxin detected, albeit at concentrations below the European guidance values for animal feeds. Ergot alkaloids, *Penicillium*-derived metabolites, and phytoestrogens had occurrence rates of 90, 48 and 29%, respectively. *Penicillium* metabolites presented the highest levels among the fungal compounds, indicating contamination during storage. Aflatoxins (AFs), ochratoxins and deoxynivalenol (DON) were not detected. Out of the 16 detected pesticides, two fungicides, ametoctradin (9.5%) and mandipropamid (14.3%) revealed concentrations exceeding their respective maximum residue level (MRL) ( $0.01 \text{ mg kg}^{-1}$ ) for barley in two samples. Although based on European guidance and MRL values the levels of the detected compounds probably do not pose acute risks for cattle, the impact of the long-time exposure to such mixtures of natural and synthetic toxicants on animal health and food safety are unknown and must be elucidated.

**Abbreviations:** BSG: Spent brewery grains; ZEN: Zearalenone; DM: Dry matter; OTA: Ochratoxin A; AFs: Aflatoxins; DON: Deoxynivalenol; EDCs: Endocrine-disrupting chemicals; ZAMG: Agency of Meteorology and Geodynamics, Zentralanstalt für Meteorologie und Geodynamik; LOD: Limit of detection; LOQ: Limit of quantification

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Dietary exposure; feed safety; mixtures toxicology; mycotoxins; pesticide residues; phytoestrogens; spent brewery grains

## Introduction

The use of agro-industrial by-products as valuable feeds in dairy farming is a strategy that can reduce the direct dependence on whole cereals grains and oilseeds, which are essential in human nutrition. Due to the low cost of by-products, their incorporation into livestock rations improves the economics, also contributing to increasing the edible feed

conversion ratio (Bocquier and González-García 2010; Ertl et al. 2015). Beer is the most consumed alcoholic beverage in Europe and worldwide (Violino et al. 2020; Ambra et al. 2021). Spent brewery grains (BSG) are the main by-product of beer production, accounting for around 85% of its agricultural waste and allowing the availability of this by-product throughout the year (Mussatto 2014;

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Petit et al. 2020). Most of the generated BSG are utilised in animal nutrition; however, this product has also been used for biogas production or, in a minor proportion, simply disposed of in landfill (Bianco et al. 2020). The vast potential of this by-product as a valuable feed/foodstuff in animal and human nutrition is because of a high content of digestible fibre (usually fluctuating from 30 to 50% on a dry matter (DM) basis), good quality protein (varying from 19 to 30% on DM basis), lipids and several minerals (José et al. 2013; Mussatto 2014; Lynch et al. 2016). Additionally, the BSG contain arabinoxylans and  $\beta$ -glucans, which can be used as prebiotics, promoting the activity of beneficial bacteria, such as *Bifidobacterium*, *Enterococcus*, and *Lactobacillus* species in monogastric livestock (Lao et al. 2020). In the feeding of ruminants, wet BSG are ideal for mixing with forage rations, offering high-quality protein feed, which is inexpensive and can reduce the dependency on commercial concentrate feeds (Gonzalez Pereyra et al. 2011). Although BSG have been fed to beef cattle, horses, pigs, sheep and poultry, the primary market for wet BSG is as a dairy cattle feedstuff (Westendorf and Wohlt 2002; Mussatto 2014; Kamboh 2017; Pack et al. 2021).

Like other agricultural products, the presence of contaminants and residues in BSG is associated with feed safety issues. Potential hazards of feedstuffs can be of natural (e.g. mycotoxins and plant toxins) as well as synthetic origin (like pesticides) (FAO and WHO 2019). Safety of wet BSG can be jeopardised by hundreds of fungal toxic compounds, which can be produced pre- and postharvest (especially during storage on the farms). However, most studies have investigated only a limited number of them in feedstuffs (including BSG) and other agricultural commodities. Specifically, research on mycotoxin contamination has focused mostly on a limited range of mycotoxins such as aflatoxins (AFs), fumonisins (FBs), trichothecenes, ochratoxin A (OTA) and zearalenone (ZEN) (Lynch et al. 2016; Cinar and Onbaşı 2019; Battilani et al. 2020; Pack et al. 2021), which are regulated by European legislation (EC 2002, 2006, 2012, 2013). Barley is the main cereal utilised for beer production (Palmer 2018). Like other cereal grains, barley is susceptible to mould infection with subsequent mycotoxin contamination and other secondary metabolites during the complete feed-

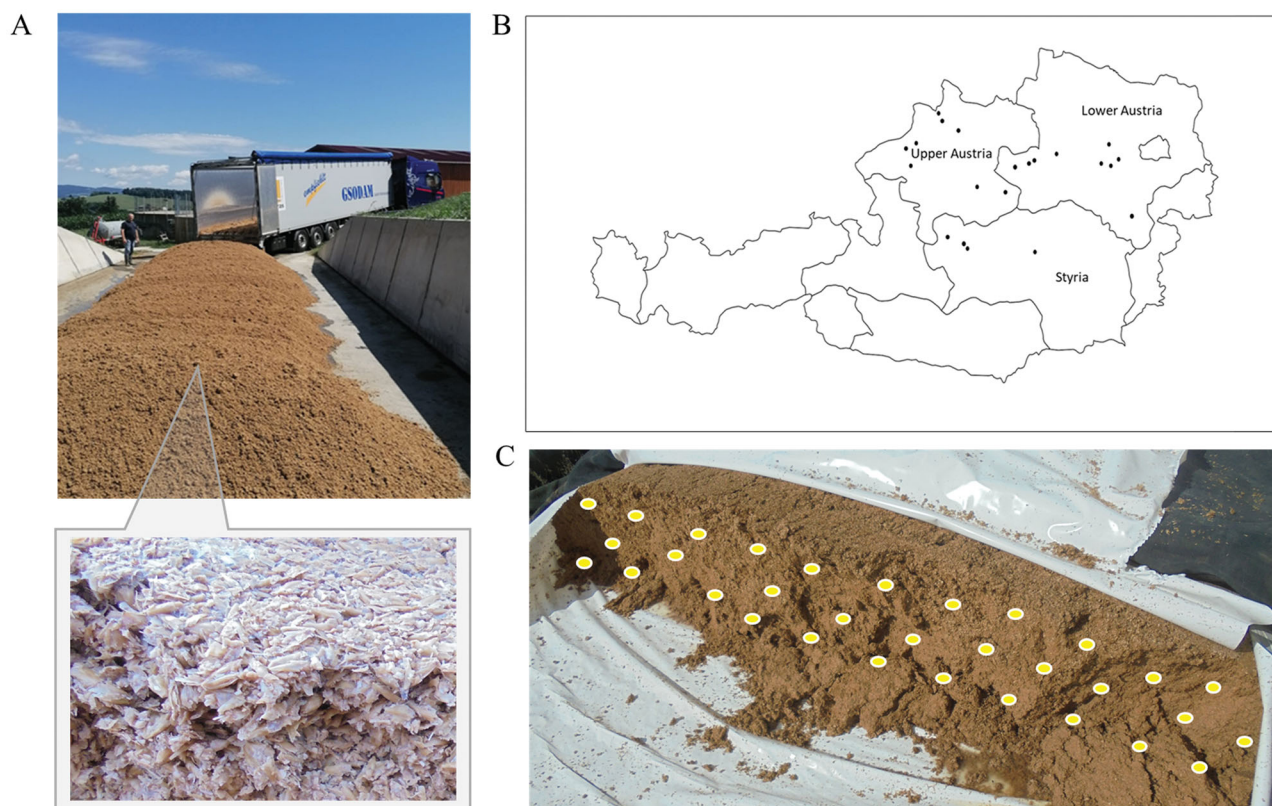
production chain, pre- and postharvest (Pascari et al. 2018). Thus, research concerning other fungal toxins such as emerging and modified mycotoxins from *Fusarium*, *Alternaria*, *Aspergillus* and *Penicillium* is still limited but often advocated (Battilani et al. 2020; Lao et al. 2020). Additionally, other compounds of natural origin like phytoestrogens can occur in feeds, affecting animal farming. For instance, these plant secondary metabolites are known endocrine disruptors that, at certain levels, can impair reproductive functions and reducing the productive efficiency of dairy herds (Wocławek-Potocka et al. 2008, 2013).

The pre- and post-harvest use of pesticides under conventional farming systems protects crops, like barley, from insects, pests, weeds and plant pathogens, improving the production yields. Residues of such pesticides can be accumulated in crops and the environment, with potentially toxic effects on human and animal (including wildlife) health as well as on the soil microorganisms (Igbedioh 1991; Damalas and Eleftherohorinos 2011; Cozma et al. 2017). Additionally, pesticide applications can generate the presence of residues, which are usually at lower levels than the mycotoxins and phytoestrogens, but which should be analysed in order to address the exposure to entire mixtures of toxicants and endocrine-disrupting chemicals (EDCs), which can have public health and environmental implications (Igbedioh 1991; Connolly 2009; Rivera-Becerril et al. 2017; Guo et al. 2020; Geissen et al. 2021; Pires et al. 2021). Like some mycotoxins, pesticides are also regulated by European Union legislation (specifically by Regulation (EC) No 396/2005) (EC 2005). Thus, this study aimed to analyse a broad-spectrum profile of mycotoxins, phytoestrogens, other secondary metabolites, and pesticide residues in wet BSG intended to feed Austrian dairy cattle. This was achieved using a validated multi-analyte method based on liquid chromatography-electrospray ionisation tandem mass spectrometry (LC-ESI-MS/MS).

## Materials and methods

### Sample collection and preparation

Representative samples of wet BSG intended for dairy cattle feeding were collected from batches acquired by farmers from regional breweries



**Figure 1.** Representative sampling of wet barley brewery's spent grains (BSG) intended for feed dairy cattle in Austria. A) The samples were collected from piles or silo bags, which were stored for a maximum of 2 weeks (Picture gently provided by Mr. Alexander Kopper<sup>©</sup>). B) The farms ( $n = 21$ ) were in Lower Austria ( $n = 9$ ), Upper Austria ( $n = 8$ ) and Styria ( $n = 4$ ). C) Subsamples were collected manually (around 20 – 30 handfuls) from the next-to-be-fed section of the BSG charge of each farm, for a final sample of 1 – 1.5 kg.

(Figure 1(A)). The sampling was conducted between June and September 2020 from 21 conventional dairy farms located in Lower Austria ( $n = 9$ ), Upper Austria ( $n = 8$ ) and Styria ( $n = 4$ ) (Figure 1(B)). Upon sample collection, BSG batches had been stored for less than two weeks at the respective farms. Each representative sample was collected manually, consisting of 20–30 incremental subsamples (randomly selected handfuls) using nitrile gloves, collected superficially (not deeper than 20 cm) from the next-to-be-fed section of the wet BSG (Figure 1(C)). The incremental samples were composited, giving a final sample amount of 1.5 kg, vacuum-packed in plastic bags and stored at  $-20^{\circ}\text{C}$  in the dark until sample preparation. Subsequently, the samples were thawed for 12 h, freeze-dried in a SCANVAC CoolSafe<sup>TM</sup> freeze (Labogene, Lillerød, Denmark) for 24 h, and milled through a 0.5 mm sieve using a cutting mill (SM 300, Retsch GmbH, Haan, Germany) at 1500 rpm for approximately 1 min. Five grams ( $\pm 0.01$  g) of the

homogenised samples were added to 50 ml polypropylene conical tubes (Sarstedt, Nümbrecht, Germany) and stored at  $-20^{\circ}\text{C}$  until analysis. The DM content of the sampled wet BSG was, on average, 25.3%, fluctuating from 20.9 to 30.6%. The farms were in altitude ranges between 266 and 814 m. The temperature during the months of sampling of participating farms fluctuated from 14.8 to 21.5 $^{\circ}\text{C}$ , being on average 18.4 $^{\circ}\text{C}$  (According to the Agency of Meteorology and Geodynamics, Zentralanstalt für Meteorologie und Geodynamik (ZAMG), <https://www.zamg.ac.at/cms/de/klima/klimauebersichten/jahrbuch>).

#### **Analysis multiple of metabolites and pesticides**

The previously dried and milled sample ( $5 \pm 0.01$  g) was placed into a 250 ml Erlenmeyer flask with 20 ml of extraction solvent according to the protocol described by Steiner et al. (2020). After agitation with a GFL 3017 rotary shaker (GFL, Burgwedel, Germany) for 90 min, the solvent

solution-sample mixture was centrifuged for 2 min at  $2012 \times g$  on a GS-6 centrifuge (Beckman Coulter Inc., Brea, CA). The extract was diluted 1:1 with dilution solvent. The injection volume of both diluted extracts of the samples and the standard analyte solutions was 5  $\mu$ l. Identification and quantification of each analyte were performed in the mode of multiple reaction monitoring with positive and negative polarity in two separate chromatographic runs using a QTrap 5500 LC-MS/MS system (Applied Biosystems, Foster City, CA) equipped with a TurboV electrospray ionisation (ESI) source coupled to a 1290 series UHPLC system (Agilent Technologies, Waldbronn, Germany). Quantitative analysis of all the analytes was performed using a validated method based on LC-ESI-MS/MS described by Sulyok et al. (2020). Quantification was based on external calibration using a serial dilution of a multi-analyte stock solution. Results were corrected for apparent recoveries determined during method validation according to Steiner et al. (2020). The values of the method performance (apparent recovery, limit of detection [LOD] and limit of quantification [LOQ]) of each analyte are presented in Table 1. The apparent recovery for each pesticide was calculated using the equation proposed by Awapak et al. (2021). The method's accuracy is verified on a routine basis by participating in a proficiency testing scheme organised by BIPEA (Gennevilliers, France) with current z-scores between  $-2$  and  $2$  indicating  $> 95\%$  confidence. All values submitted for a sample of wheat chaff were within this satisfactory range.

### Data analysis

Concentrations of all detected contaminants, residues, and other non-regulated metabolites were presented on a DM basis in  $\mu\text{g kg}^{-1}$ . Descriptive statistics, i.e. frequencies, mean, median and ranges of the concentration of analytes, were calculated considering only the positive results ( $x \geq$  limit of LOD). Results below the LOQ were computed as LOQ/2. All statistical evaluations, tables, and graphs were performed using Microsoft Excel<sup>®</sup> and GraphPad Prism<sup>®</sup> version 9.1 (GraphPad Software, San Diego, CA).

**Table 1.** Performance values of LC-MS/MS analysis targeting fungal and other contaminants as well as pesticide residues detected in wet brewery's spent grains intended for dairy cattle nutrition.

Analyte	Method performance		
	Apparent recovery (%)	LOD ( $\mu\text{g kg}^{-1}$ )	LOQ ( $\mu\text{g kg}^{-1}$ )
15-Hydroxyculmorin	100	0.3	0.9
Alternariol	45	3.4	11.3
Alternariolmethylether	52	3.3	11
Altersetin	75	0.7	2.3
Altetoxin-I	26	22	73
Ametoctradin	57	2.7	8.9
Andrastin A	88	4.3	14
Andrastin B	84	0.4	1.2
Andrastin C	41	3.5	12
Antibiotic Y	80	0.1	0.4
Apicidin	93	0.2	0.8
Apicidin D2	71	0.9	2.8
Asperphenamate	75	0.1	0.4
Aurofusarin	51	0.6	1.9
Azoxystrobin	64	1.7	5.6
Beauvericin	103	0.05	0.2
Benzovindiflupyr	73	0.8	2.8
Bikaverin	72	10	30
Bixafen	56	2.8	9.3
Boscalid	41	2.7	8.9
Brevianamid F	49	2.4	8.1
BTS 44595	67	1.3	4.3
Butenolid	93	7	23.4
Chanoclavin	44	2.6	8.5
Chrysogine	95	5.7	19
Citreorosein	33	2	6.6
Culmorin	65	0.5	1.5
Cyclo (L-Pro-L-Tyr)	31	15	52
Cyclo (L-Pro-L-Val)	31	15	52
Cyclosporin A	56	4.9	16.4
Daidzein	42	50	180
Deoxynortryptoquivalin	52	1.9	6.4
Deoxytryptoquivaline A	52	0.9	3.1
Emodin	71	2.1	7
Enniatin A	37	4.2	13.9
Enniatin A1	52	0.5	1.7
Enniatin B	75	2	6
Enniatin B1	51	17	58
Enniatin B2	92	0.7	2.4
Epiequisetin	138	1	3.2
Equisetin	138	1	3.2
Ergocormine	96	0.3	0.9
Ergocristine	63	1.1	3.8
Ergocristinine	76	0.8	2.7
Ergocryptine	55	0.1	0.4
Ergometrine	61	0.06	0.2
Ergometrinine	77	0.3	1
Ergosin	58	0.2	0.6
Ergosinin	65	0.2	0.6
Ergotamine	65	0.1	0.4
Ergotaminin	65	0.1	0.4
F01 1358-A	100	1	3
Festuclavine	50	0.5	1.5
Flavoglucan	47	0.4	1.3
Fluopyram	6.4	2.3	7.5
Fluxapyroxad	57	2.2	5.3
Fumiquinazolin D	53	1	3.2
Fungerin	72	0.4	1.3
Fusaproliferin	100	10	30
Fusaric acid	80	10	30
Genistein	62	28	92
Gibberellin A12	57	1.2	4.1
Glycitein	59	31	105
HT-2 toxin	74	2.6	8.5
Hydroxyandrastin A	52	2.5	8.5

(continued)

Table 1. Continued.

Analyte	Method performance		
	Apparent recovery (%)	LOD ( $\mu\text{g kg}^{-1}$ )	LOQ ( $\mu\text{g kg}^{-1}$ )
Hydroxyandrastin C	85	2.1	7.1
Infectopyrone	210	0.7	2.5
Isopyrazam	75	1.1	3.7
Kotantin A	55	1.5	5
Macrosporin	79	2	6.7
Mandipropamid	51	2.4	7.8
Marcfortine A	56	1.9	6.2
Marcfortine C	56	1.9	6.2
Metrafenone	64	0.5	1.8
Monocerin	67	1.3	4.2
Mycophenolic acid	52	0.6	1.9
Mycophenolic acid IV	57	0.6	1.9
Neoechinulin A	38	4	14
Patulin	89	0.6	2.1
Phenopyrrozin	75	0.3	0.9
Physcion	45	11	37
Pinselin	57	1.3	4.2
Piperonyl butoxide	100	5	15
Pirimiphos-methyl	97	1	3.3
Pseurotin A	56	7.4	28
Pyraclostrobin	62	2.3	7.7
Pyrenophorol	74	2.4	8.1
Questiomycin	74	2.8	9
Quinadoline A	112	0.8	2.6
Roquefortine C	58	0.9	3.1
Roquefortine D	66	1.3	4.4
Rubellin D	49	1.7	5.6
Rugulosovin	56	1	3.2
Siccanol	72	2.3	7.6
Sporidesmolide II	100	0.02	0.05
T-2 toxin	111	7	23
Tebuconazole	68	1.1	3.7
Tentoxin	58	2.2	7.5
Tenuazonic acid	150	10	30
Trifloxystrobin	67	1.2	4
Tryptophol	30	100	300
Tryptoquialanine derivate	35	1	3.3
Tryptoquialanine A	58	1.8	6
Verrucofortine	57	1.5	5
Viriditoxin	100	2.5	7.5
W493	195	2.1	7
Zearalenone	70	2.8	9.2

## Results and discussion

### General overview of detected groups of analytes

Analytes of natural origin were categorised into groups: *Alternaria*, *Aspergillus*, ergot alkaloids, *Fusarium* and *Penicillium* mycotoxins, other fungal species, phytoestrogens and unspecific metabolites, as in previous reports (Szulc et al. 2019; Hajnal et al. 2020; Penagos-Tabares 2021). Table 2 shows the occurrences and respective average, median, and range concentrations ( $\mu\text{g kg}^{-1}$  DM) of observed secondary metabolites. A total of 107 out of more than 1400 targeted secondary metabolites and pesticides were detected: 78 fungal compounds, three phytoestrogens, 16 pesticides and 10 unspecific metabolites (Figure 2). The

categories of fungal metabolites with the highest number of detected metabolites in this exploratory study were *Fusarium* spp. (26 metabolites), *Penicillium* (18), *Aspergillus* (11), ergot alkaloids (10), *Alternaria* (9), with fewer metabolites from other fungal genera (4). All the samples contained metabolites derived from *Aspergillus*, *Alternaria*, *Fusarium* and other fungal species, while ergot alkaloids and *Penicillium*-derived metabolites occurred in 90 and 48% of BSG, respectively. Phytoestrogens were found in 29% of the analysed BSG, and the detection of pesticide residues was ubiquitous (100%). In this study, *Fusarium* metabolites showed the highest level of diversity, which has been previously observed in other naturally-contaminated samples from Austria and Europe (Reisinger et al. 2019; Penagos-Tabares et al. 2021, 2022). This again suggests the status of *Fusarium* as the most widespread fungal genus in cereal growing areas and as a significant contributor to mycotoxin contamination in animal feeds (Nesic et al. 2014). Additionally, it also corroborates the widespread occurrence of other mycotoxigenic genera (*Aspergillus*, *Alternaria* and *Penicillium*) (Grenier and Oswald 2011). Regarding the detected levels, the group of unspecific metabolites showed the highest average concentration ( $7010 \mu\text{g kg}^{-1}$ ), followed by fungal metabolites ( $5140 \mu\text{g kg}^{-1}$ ), phytoestrogens ( $957 \mu\text{g kg}^{-1}$ ) and pesticides ( $208 \mu\text{g kg}^{-1}$ ). Specifically for the fungal metabolites, the highest average levels were for the category of *Penicillium* ( $5600 \mu\text{g kg}^{-1}$ ), followed by *Fusarium* ( $2200 \mu\text{g kg}^{-1}$ ), *Alternaria* ( $107 \mu\text{g kg}^{-1}$ ), ergot alkaloids ( $69.5 \mu\text{g kg}^{-1}$ ), *Aspergillus* ( $79.6 \mu\text{g kg}^{-1}$ ) and metabolites from other fungi ( $15.5 \mu\text{g kg}^{-1}$ ) (Table 2, Figure 2(A)).

### Regulated mycotoxins and related forms

Among the mycotoxins included in the European legislation were detected ZEN, T-2 toxin, and HT-2 toxin, which have recommended GVVs (EC 2006, 2013). ZEN was the regulated mycotoxins that occurred the most, detected in 57% of the samples with a maximum concentration of  $32.3 \mu\text{g kg}^{-1}$ , whereas T-2 and HT-2 toxins were detected in only one sample (5%), showing concentrations of 3.8 and  $4.25 \mu\text{g kg}^{-1}$ , respectively.

**Table 2.** Occurrences and levels of fungal and other natural contaminants detected in wet brewery's spent grains intended for dairy cattle nutrition.

Group	Metabolite	Occurrence (%) <sup>a</sup>	Concentration ( $\mu\text{g kg}^{-1}$ ) <sup>b</sup>		
			Mean $\pm$ SD	Median	Range (GV) <sup>c</sup>
<i>Alternaria</i> spp.	Alternariol	81	5.99 $\pm$ 1.41	5.65	5.65–11.5
	Alternariolmethylether	86	5.50 $\pm$ 0	5.50	5.50–.50
	Altersetin	14	15.8 $\pm$ 14	8.70	6.85–31.9
	Altetoxin-I	5		–	36.5
	Infectopyrone	100	76.2 $\pm$ 30.1	74.2	25.9–141
	Macrosporin	52	3.64 $\pm$ 0.37	3.75	2.51–3.75
	Pyrenophorol	5			14.9
	Tentoxin	5			1.15
	Tenuazonic acid	48	29.6 $\pm$ 16.8	22.8	15.0–56.7
	Total	100	107 $\pm$ 52.3	95.1	30.9–229
	<i>Aspergillus</i> spp.	Deoxynortryptoquivalin	33	3.20 $\pm$ 0	3.20
Deoxytryptoquivaline A		90	26.5 $\pm$ 15.1	22.9	1.55–53.2
Flavoglucin		71	1.82 $\pm$ 1.42	1.63	0.65–5.19
Fumiquinazolin D		10	4.92 $\pm$ 4.7	4.92	1.60–8.25
Kotanimin A		33	2.50 $\pm$ 0	2.50	2.50–2.50
Pinselin		43	4.40 $\pm$ 3.33	1.55	1.55–10.3
Pseurotin A		5			31.7
Quinadoline A		71	13.1 $\pm$ 5.9	11.0	5.2–24.8
Tryptoquialanine derivate		76	11.6 $\pm$ 5.09	11.7	5.07–20.9
Tryptoquivaline A		14	3.00 $\pm$ 0	3.00	3.00
Viriditoxin		48	65.3 $\pm$ 25.9	60.5	28.9–116
Total		100	79.6 $\pm$ 46.8	72.9	22.2–228
Ergot alkaloids		Ergocornine	52	9.77 $\pm$ 4.95	8.67
	Ergocristine	86	26.8 $\pm$ 22.8	22.6	7.66–104
	Ergocristinine	29	2.23 $\pm$ 1.42	1.35	1.35–4.70
	Ergocryptine	90	13.9 $\pm$ 8.3	12.4	3.37–33.0
	Ergometrine	43	0.11 $\pm$ 0.04	0.1	0.10–3.75
	Ergometrinine	57	1.62 $\pm$ 1.02	1.53	0.37–3.75
	Ergosin	90	7.8 $\pm$ 6.24	7.02	0.30–23.3
	Ergosinin	57	1.92 $\pm$ 1.56	1.24	0.30–5.62
	Ergotamine	76	12.9 $\pm$ 7.06	10.2	4.38–27.9
	Ergotaminin	71	3.7 $\pm$ 3.29	2.41	0.94–12.3
	Total	90	69.5 $\pm$ 52.7	57.4	14.0–210
<i>Fusarium</i> spp.	15-Hydroxyculmorin	14	11.7 $\pm$ 0	11.7	11.7–11.7
	Antibiotic Y	5			9.50
	Apicidin	57	15.1 $\pm$ 13.3	9.64	2.74–43.0
	Apicidin D2	5			6.95
	Aurofusarin	100	137 $\pm$ 107	116	15.4–364
	Beauvericin	100	6.37 $\pm$ 6.25	4.37	1.40–32.4
	Bikaverin	100	13.6 $\pm$ 5.93	11.4	4.61–26.4
	Butenolid	5			237
	Chrysogin	19	3.46 $\pm$ 1.65	3.74	1.20–5.17
	Culmorin	100	390 $\pm$ 210	348	118–802
	Enniatin A	100	5.56 $\pm$ 1.83	5.12	2.06–8.73
	Enniatin A1	100	35.0 $\pm$ 12.8	30.0	12.2–60.9
	Enniatin B	100	201 $\pm$ 77.3	173	84.4–380
	Enniatin B1	100	171 $\pm$ 61.5	151	65.2–321
	Enniatin B2	100	4.40 $\pm$ 1.67	3.91	1.98–7.71
	Epiequisetin	14	1.60 $\pm$ 0	1.60	1.60–1.60
	Equisetin	95	3.91 $\pm$ 6.18	1.60	1.60–29.0
	Fungerin	43	2.38 $\pm$ 2.28	1.41	0.65–7.27
	Fusaproliferin	24	36.2 $\pm$ 24.9	32.5	15.0–75.1
	Fusaric acid	10	481 $\pm$ 617	481	45.1–917
	Gibberellin A12	95	189 $\pm$ 89.6	168	43.1–389
	HT-2 toxin	5			3.80 (250)
	Siccanol	100	966 $\pm$ 297	989	275–1503
	T-2 toxin	5			4.25 (250)
	W493	14	18.4 $\pm$ 16.3	15.8	3.50–35.7
	Zearalenone	57	13.2 $\pm$ 8.92	11.5	4.60–32.3 (500)
	Total	100	2200 $\pm$ 579	2343	1049–3100
<i>Penicillium</i> spp.	Andrastin A	19	2910 $\pm$ 4540	1020	14.7–9570
	Andrastin B	10	5990 $\pm$ 5420	5990	2160–9830
	Andrastin C	10	6400 $\pm$ 3630	6400	3840–8970
	Chanoclavin	5			0.95
	F01 1358-A	5			19.2
	Festuclavine	5			74
	Hydroxyandrastin A	5			46.9
	Hydroxyandrastin C	10	15.4 $\pm$ 12.5	15.4	6.60–24.2

(continued)



Table 2. Continued.

Group	Metabolite	Occurrence (%) <sup>a</sup>	Concentration ( $\mu\text{g kg}^{-1}$ ) <sup>b</sup>		
			Mean $\pm$ SD	Median	Range (GV) <sup>c</sup>
Other fungi	Marcfortine A	14	292 $\pm$ 498	8.48	0.99–867
	Marcfortine C	5		7.25	
	Mycophenolic acid	14	261 $\pm$ 399	33.0	28.6–722
	Mycophenolic acid IV	5			1094
	Patulin	5			9201
	Phenopyrrozin	24	5.33 $\pm$ 5.58	1.30	1.29–12.5
	Questiomycin	19	2.86 $\pm$ 1.67	2.51	1.5–4.92
	Roquefortine C	10	480 $\pm$ 450	480	162–798
	Roquefortine D	10	60.7 $\pm$ 64.8	60.7	14.9–106
	Verrucofortine	29	2.50 $\pm$ 0	2.50	2.50–2.50
	Total	48	5600 $\pm$ 10,400	14.0	1.3–30,300
	Cyclosporin A	5			222
	Monocerin	5			1.86
	Rubellin D	90	5.12 $\pm$ 3.57	2.80	2.8–12.8
Sporidesmolide II	90	0.20 $\pm$ 0.10	0.18	0.09–0.42	
Total	100	15.5 $\pm$ 48.2	3.01	0.10–225	
Sum of total fungal metabolites		100	5140 $\pm$ 7750	2750	1280–34,000
Phytoestrogens	Daidzein	24	501 $\pm$ 682	90.0	90.0–1660
	Genistein	29	508 $\pm$ 707	149	46.0–850
	Glycitein	10	93.6 $\pm$ 58.2	93.6	52.5–135
	Total	29	957 $\pm$ 1400	213	136–3650
	Unspecific metabolites	Asperphenamate	62	0.25 $\pm$ 0.12	0.20
Brevianamid F		100	307 $\pm$ 150	268	138–738
Citreosein		19	9.90 $\pm$ 4.26	8.46	6.62–16.0
Cyclo(L-Pro-L-Tyr)		100	2304 $\pm$ 1387	2080	580–6370
Cyclo(L-Pro-L-Val)		100	1937 $\pm$ 1118	1690	527–4770
Emodin		52	3.99 $\pm$ 1.64	3.50	3.50–8.93
Neoechinulin A		24	10.9 $\pm$ 8.83	7.00	7.00–26.7
Physcion		5			535
Rugulosoavin		100	107 $\pm$ 61.3	99.4	24.2–280
Tryptophol		100	2330 $\pm$ 2480	090	150–8240
Total		100	7010 $\pm$ 3400	6380	2620–16,160

<sup>a</sup> $n = 21$  representative samples of wet barley-derived brewery spent grains from Austria, considered as positive values  $>$  limit of detection (LOD); <sup>b</sup>calculations without data  $<$  LOD. In case values  $>$  LOD and  $<$  limit of quantification (LOQ), LOQ/2 was used for the calculation; <sup>c</sup>GV: Guidance value according to European Commission (EC 2006, 2013)

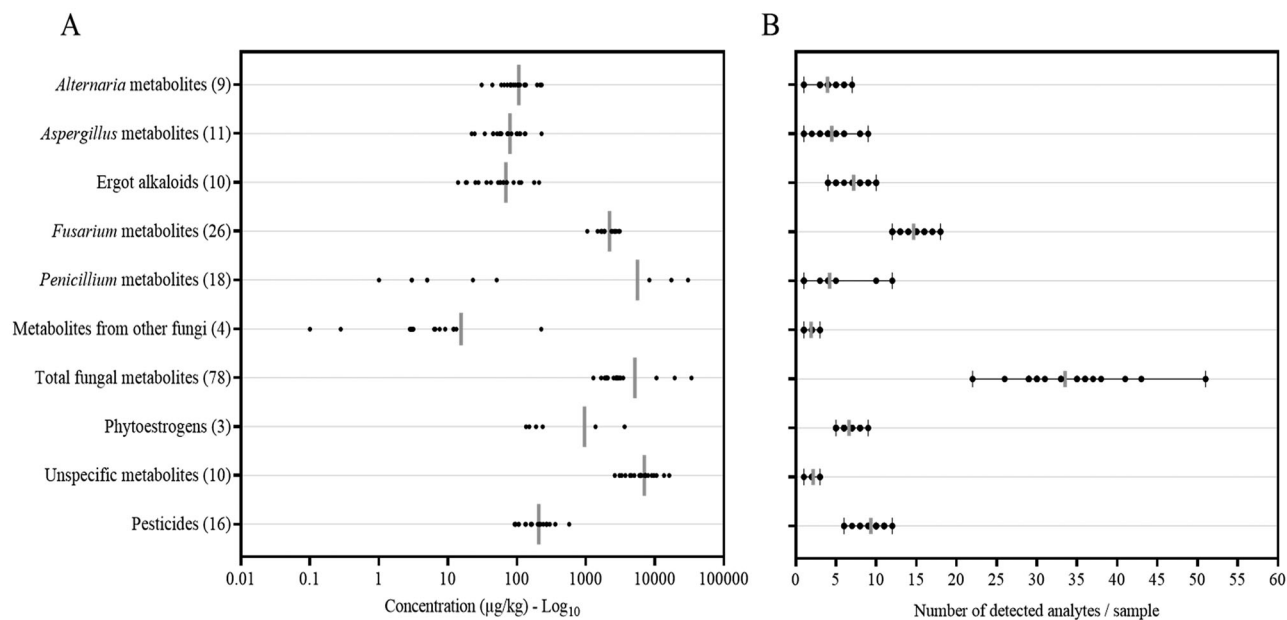


Figure 2. (A) Distribution of concentrations and (B) co-contamination grade (detected analytes per sample) of major categories of analytes detected in wet brewery's spent grains intended for the nutrition of dairy cattle in Austria.

The observed concentrations were still under the GVs of the European Union, which are  $500 \mu\text{g kg}^{-1}$  for ZEN and  $250 \mu\text{g kg}^{-1}$  for the sum of T-2 toxin and HT-2 toxin relative to a feed with moisture of 12% (EC 2002, 2006, 2013). The findings align with a previous report, which detected ZEN contamination in beer (Bauer et al. 2016). Other regulated mycotoxins such as AFB1, deoxynivalenol (DON), OTA and FB1 and FB2, which are regulated in European legislation for dairy cattle, were not detected. Regulated mycotoxins are known to cause adverse health effects. For example, ZEN is linked with hyper-estrogenism, reduced milk production, early abortion and other reproductive abnormalities (Hussein and Brasel 2001; Marczuk et al. 2012). Trichothecenes, like T-2 toxin and HT-2 toxin, are known for inducing inhibition of DNA and RNA synthesis, which can be a secondary effect of the inhibition of protein synthesis or due to apoptosis (Cope 2018). *Fusarium* head blight is a common disease in barley associated with ZEN, type A trichothecenes and other mycotoxins. This disease is a significant threat to the brewing industry, with *Fusarium graminearum* considered the predominant causal species worldwide (Bai and Shaner 2004; Starkey et al. 2007; Schwarz 2017). Other species like *Fusarium culmorum*, *Fusarium poae* and *Fusarium avenaceum* have also been described as some of the most widely occurring in Europe (Becher et al. 2013). *F. graminearum* and *F. culmorum* are important producers of ZEN, DON and nivalenol (Bottalico and Perrone 2002). *F. langsethiae* and *F. sporotrichioides* are producers of HT-2 and T-2 toxins (Thrane et al. 2004). It has been suggested that low levels of fusarial mycotoxins like ZEN and type-B trichothecenes (like DON, 15-acetyldeoxynivalenol and 3-acetyldeoxynivalenol) are retained in BSG (Pack et al. 2021). Previous studies found that the residual concentrations of several fungal toxins including OTA, AFB2, FB2, AFG1, AFB1, ZEN and patulin decreased to less than 20% during the brewing process (Inoue et al. 2011; Piacentini et al. 2019). Although T-2 toxin has received special attention in the malting barley chain, due to its occurrence in this cereal crop and its toxic potency, the contamination levels of this toxin generally decrease during the

brewing process (Edwards et al. 2009; Malachova et al. 2010). A study of mycotoxins in BSG in Argentina evidenced FB1 (100%; range:  $104\text{--}145 \mu\text{g kg}^{-1}$ ) and AFB1 (18%;  $19\text{--}44.5 \mu\text{g kg}^{-1}$ ), whereas AFB2, AFG1, AFG2 and ZEN were not detected (Pereyra et al. 2011).

Regarding ergot alkaloids, which were detected in high occurrence (90%), the European Union recommends the monitoring of these metabolites (Recommendation 2012/154/EC) (EC 2012). There is no specific guidance value for ergot alkaloids in animal feeds. Still, the limit of  $1000 \text{ mg kg}^{-1}$  rye ergot (*Claviceps purpurea*) represents a maximum value relative to a feedstuff with a moisture of 12% described in the directive 2002/32/EC (EC 2012). Stricter regulation is in place for foodstuffs. Since January 2022, the Commission Regulation (EU) 2021/1399 established a maximum level of ergot alkaloids in certain foodstuffs. For instance, for milling barley products intended for human consumption (with an ash content lower than  $9000 \text{ mg kg}^{-1}$ ), a maximum level of  $100 \mu\text{g kg}^{-1}$  ( $50 \mu\text{g kg}^{-1}$ ) will be implemented from 1.7.2024). For milling barley products (with an ash content equal to or higher than  $9000 \text{ mg kg}^{-1}$ ) or barley grains placed on the market for the final consumer, the maximum level set is  $150 \mu\text{g kg}^{-1}$  (EU 2021).

Moreover, our results show that members of the ergopeptine class alone, including ergocryptine, ergosin and ergocristine, presented with occurrences of over 85%. The average concentrations of individual ergot alkaloids were under  $30 \mu\text{g kg}^{-1}$ , and the maximal accumulated concentration of total ergot alkaloids was  $210 \mu\text{g kg}^{-1}$ , which should not be ignored. Feed contaminated with  $250 \mu\text{g kg}^{-1}$  of ergot alkaloids should not be fed to pregnant or lactating animals due to a higher risk of abortion and agalactia syndrome; even low concentrations of alkaloids in the diet ( $<100 \mu\text{g kg}^{-1}$  total) can reduce the growth efficiency of livestock (Coufal-Majewski et al. 2016). Some members of the ergot alkaloids such as ergotamine, ergocristine, ergosine, ergocornine, ergocryptine and ergovaline are responsible for the majority of nervous or gangrenous syndromes in humans and animals, which consume grains, grain products or grasses contaminated with the sclerotia of the

fungus (Gupta et al. 2018). Ingestion of this kind of alkaloids by livestock can trigger a range of impacts from decreased performance and reduced fertility to acute clinical signs of ergotism, including nervous or gangrenous syndromes, hyperthermia, convulsions, necrosis of the extremities and death (Evans 2011). According to the scientific opinion of EFSA, ergotism in ruminants is usually a chronic disease and the result of continued ingestion of minor quantities of the fungus on grass (EFSA 2012). A large proportion of the original peptide alkaloids can be removed during brewing which is believed to result from thermal degradation (Schwarz et al. 2007). However, our data confirm that the reduction is not absolute.

### Other fungal toxins and metabolites

Emerging mycotoxins are the focus of high scientific interest. They are defined as commonly occurring in feed and foods (agricultural commodities) and are legislatively unregulated and non-regularly tested (Vaclavikova et al. 2013). We showed that several emerging and non-regulated mycotoxins and metabolites were detected in samples of BSG intended for cattle feeding on Austrian farms. Several *Fusarium*-derived emerging mycotoxins include culmorin, siccanol, aurofusarin, beauvericin, bikaverin and enniatins (A, A1, B, B1 and B2), were detected in all the samples. The enniatins B and B1 presented the highest concentrations among the fusarial mycotoxins, with averages exceeding  $170 \mu\text{g kg}^{-1}$ . Enniatins and beauvericin have haematotoxic, immunotoxic and antibiotic activities (Sy-Cordero et al. 2012; EFSA 2014; Juan et al. 2019; Křížová et al. 2021). Research on the impact of such fungal antimicrobial compounds on rumen ecology and functionality is essential (Fink-Gremmels 2005, 2008; Reisinger et al. 2019). Siccanol (also called terpestacin) (Chan and Jamison 2003) was the fusarial compound with the highest concentration (average:  $966 \mu\text{g kg}^{-1}$ ) in BSG. Another fusarial metabolite, fusaric acid occurred with a frequency of 19%. This compound can increase the toxicity of other *Fusarium* mycotoxins such as moniliformin, trichothecenes and FBs (Bacon et al. 1996; D'Mello et al. 1999).

Emerging *Alternaria* mycotoxins, such as alternariol (81%), alternariol methyl ether (86%) and tenuazonic acid (48%) were detected in relatively low concentrations. It is known that these compounds have estrogenic activity and genotoxic effects (Escrivá et al. 2017; Aichinger et al. 2019, 2021). The genus *Alternaria* is widely distributed in the environment and is one of the leading causes of disease in cereal crops like wheat, barley and sorghum (Deshpande 2002). However, information is still missing regarding *Alternaria* mycotoxins in the feeds and their toxicological repercussions on animal health (EFSA 2011). Infectopyrone was found in all the BSG samples and the compound with the highest average and maximum concentration among the *Alternaria*-derived metabolites. Is a potential mycotoxin whose biological activities are unknown and should be further explored (Andersen et al. 2002; Larsen et al. 2003). *Alternaria*-derived compounds like altersetin, altertoxin-I, pyrenophorol and ten-toxin were also found. Although at this time, there are no global regulations establishing limits for these toxins in food and feed, the European Food Safety Authority (EFSA) has raised concern about *Alternaria* mycotoxins in relation to public health (EFSA 2011; Escrivá et al. 2017). *Aspergillus*-derived compounds were omnipresent in Austrian BSG evaluated in this study. The most frequently-occurring metabolites from *Aspergillus* were deoxytryptoquivaline A, tryptoquialanine derivate, flavoglaucin and quinadoline A, detected in frequencies above 70%. Viriditoxin and deoxytryptoquivaline A presented the highest average, median and maximum concentration of *Aspergillus*-derived metabolites. Other molecules produced by this genus including deoxynortryptoquivalin (33%), fumiquinazolin D (10%), kotanin A (33%), pinselin (43%), pseurotin (5%) and tryptoquivaline A (14%) were also found in BGS (Table 2). Viriditoxin, fumiquinazolin D, quinadoline A exhibit antibacterial properties (Qian et al. 2019; Urquhart et al. 2019; Almeida et al. 2021). Tryptoquialanines belong to the group of tremorgenic mycotoxins that can be produced by species of *Aspergillus* and *Penicillium* (Clardy et al. 1975; Ariza et al. 2002).

Strongly linked to postharvest contamination, *Penicillium*-derived metabolites presented an occurrence of 48%. This category of fungal compounds showed the highest average and maximal

concentration (average:  $5604 \mu\text{g kg}^{-1}$ ; max:  $30,300 \mu\text{g kg}^{-1}$ ). The metabolites verrucofortine (29%) phenopyrrozin (24%) were the more recurrent compounds from penicillia, but their mean concentrations were very low ( $<10 \mu\text{g kg}^{-1}$ ). The *Penicillium*-derived metabolites detected in highest concentrations were andrastin C (average:  $6,400 \mu\text{g kg}^{-1}$ , occurrence: 10%), andrastin B ( $5990 \mu\text{g kg}^{-1}$ , 10%) and andrastin A ( $2910 \mu\text{g kg}^{-1}$ , 19%). These compounds, among other *Penicillium*-derived substances, are commonly found in silage, with higher concentrations in mouldy hot spots (Reisinger et al. 2019; Gallo et al. 2021; Penagos-Tabares et al. 2022; Manni et al. 2022). The complete spectrum of the biological activities and toxicological effects of the andrastins has not been elucidated. It is known that the andrastins are protein farnesyltransferase inhibitors, which can inhibit the efflux of anti-cancer drugs from multidrug-resistant cancer cells and are devoid of antimicrobial activity. They are commonly found in European blue (mould) cheeses (Uchida et al. 1996). Other compounds detected in this study were mycophenolic acid, and roquefortines, the most investigated *Penicillium* metabolites in silages (Gallo et al. 2015). A common feature of many detected metabolites like mycophenolic acid, roquefortines and patulin, is their immunotoxic properties (Oh et al. 2012; Brennan et al. 2017), which could interfere with the activity of innate and adaptive immune responses, predisposing to secondary infectious diseases (Oh et al. 2015). Phenopyrrozin, along with marcfortines A and C, were also detected in the analysed BSG samples. Several researchers have proposed that *Penicillium* toxins can induce unspecific clinical signs like appetite reduction, affecting nutrient efficiency and increasing the incidence of abomasal ulcers, laminitis, gastroenteritis, abortion and paralysis (Dzidic et al. 2006; Nielsen et al. 2006; Fink-Gremmels 2008; Alonso et al. 2013; Gallo et al. 2015). Additionally, further less-known metabolites produced mainly by other fungal species were detected in this study. Rubellin D and sporidesmolide occurred at a frequency of 90%, whereas cyclosporin A and monocerin were detected in only one sample (5%). The quantified levels of these metabolites

were below  $15 \mu\text{g kg}^{-1}$ , except for cyclosporin A ( $222 \mu\text{g kg}^{-1}$ ) (Table 2). Rubellin D is an anthraquinone derived from *Ramularia collo-cygni* with antibacterial activity (Walters et al. 2008; Miethbauer et al. 2009). Cyclosporin A has a potent immunotoxic activity, which has even been used commercially in human and veterinary medicine as an immunosuppressant (Laupacis et al. 1982; Shevach 1985; Stähelin 1996).

Postharvest infestations with moulds proliferate under aerobic conditions, producing potent toxins, disruptive endocrine substances and antimicrobial compounds. Moreover, fungal growth leads to spoilage, thereby reducing the nutritional value, DM content, intake and palatability (O'Brien et al. 2006). The high proportion of moisture in wet BSG makes this product especially susceptible to microbial growth and spoilage in a short period (7–10 d) (Lilly et al. 1980; Stojceska and Ainsworth 2008; Chanie and FieVez 2017). Strategies suggested for preserving wet BSG include drying with solar radiation and ensiling. Drying by solar radiation is found to be challenging due to costs (Conrad and Rogers 1977). Alternatively, the ensiling of wet brewery grain alone or mixed with dry fodders is the proposed practice for dairy farmers, especially in developing countries (Kindbom 2012; Souza et al. 2012). The preservation of wet BSG by lowering the water activity of the material using beet molasses (30%) and further stabilising the mixture by incorporating an anti-mycotic agent (0.3% of potassium sorbate) has been achieved at both laboratory-scale and pilot-scale. For practical preservation, the stabilised grains should be stored under anaerobic conditions in plastic bags, squeezing the air out and sealing tightly (Lilly et al. 1980). However, more applied research on preservation strategies for BGS is still required.

### **Contamination of phytoestrogens and other secondary metabolites**

In the present research, three isoflavones were detected in medium-low frequencies: Daidzein (24%), genistein (29%) and glycitein (10%) (Table 2). The predominant daidzein and genistein presented average concentrations above  $500 \mu\text{g kg}^{-1}$  and maximum above  $1600 \mu\text{g kg}^{-1}$ . These

metabolites are found primarily in *Leguminosae* plants, such as soy (*Glycine max*) but also occur in clovers (*Trifolium* spp.) and alfalfa/lucerne (*Medicago sativa*) (Reed 2016). Glycitein presented an average concentration of  $93.6 \mu\text{g kg}^{-1}$ , ranging from  $52.5$  to  $135 \mu\text{g kg}^{-1}$ . Liggins et al. (2002) reported daidzein and genistein in cereal-derived products for human consumption. In pearl barley, only genistein was detected with an average concentration of  $86 \mu\text{g kg}^{-1}$ . The concentration of the two isoflavones in the remaining foods ranged from  $33$  to  $11,873 \mu\text{g kg}^{-1}$  (Liggins et al. 2002). Coumetrans like coumestrol, which were not found, seem to have a more potent estrogenic activity than the detected isoflavones here (Romero et al. 1997). The detected concentration of phytoestrogens (isoflavones) found in our study apparently does not represent a potential risk for cattle (Grgic et al. 2021).

In addition, several unspecific secondary metabolites were detected. These analytes can be produced by different and unrelated living systems belonging to diverse kingdoms (Plantae, Fungi, Animalia and/or Eubacteria). Several of the unspecific secondary metabolites detected in our study are biologically active molecules. These compounds could influence the toxicological complexity of the complete cocktails of secondary metabolites evidenced in this investigation. They included, for example, emodin (antibacterial and immunosuppressive) (Kiyoshi et al. 1984; Dong et al. 2016) as well as the diketopiperazines cyclo-(L-Pro-L-Tyr) (synonym: maculosin) and cyclo-(L-Pro-L-Val) (antibacterial) (Park et al. 1993; Rahman et al. 2020; Zin et al. 2020; Paudel et al. 2021). Other detected unspecific metabolites were neoechinulin A, physcion, rugulosovine and tryptophol. The most predominant unspecific metabolites (detected in all the samples) were tryptophol, cyclo-(L-Pro-L-Tyr) and cyclo-(L-Pro-L-Val), which also showed the highest average concentrations of this group ( $>1900 \mu\text{g kg}^{-1}$ ) (Table 2).

### **Pesticide residues**

All the samples presented residues of pesticides, varying from six to twelve different biocides per sample (Figure 2(B)). No illegal compounds (EU-

MRL-Database 2022) were detected. In total, 16 pesticides were found: 14 fungicides, one insecticide (pirimiphos-methyl) and an insecticide synergist (piperonyl butoxide) as classified in previous studies (Table 3) (Huang and Subramanyam 2005; Opalski et al. 2006; Lamberth et al. 2008; Sooväli and Koppel 2010; Harp et al. 2011; Lazzari et al. 2012; Rodrigues et al. 2013; Kanungo and Joshi 2014; Rumbos et al. 2016; EFSA 2017, 2018; McLean and Hollaway 2019; Xu et al. 2020; Basak et al. 2021; Yao et al. 2021; EU-MRL-Database 2022; Rathod et al. 2022). Occurrences along with the corresponding average, median and range concentrations (expressed in  $\mu\text{g kg}^{-1}$  DM) of the detected pesticide residues, their respective uses, the maximum residue levels (MRLs) in barley, and the proportion of samples above the respective MRL are presented in Table 3. The most frequently detected pesticides in the analysed BSG samples were fluopyram, piperonyl butoxide, fluxapyroxad, bixafen, mandipropamid and tebuconazol, which were seen in  $\geq 85\%$  of the samples. The fungicides azoxystrobin, benzovindiflupyr and boscalid as well as the insecticide pirimiphos-methyl showed occurrences between 43 and 62%. Residues of ametoctradin, isopyrazam, pyraclostrobin and trifloxystrobin were found in less than 40% of the evaluated BGS samples. The pesticides with the highest average levels of residues were piperonyl butoxide ( $116 \mu\text{g kg}^{-1}$ ), metrafenone ( $30.4 \mu\text{g kg}^{-1}$ ) and fluopyram ( $24.7 \mu\text{g kg}^{-1}$ ) (Table 3). Notably, 9.5% and 14.3% of the samples exceed the respective current MRLs ( $0.01 \text{ mg kg}^{-1}$ ) of ametoctradin and mandipropamid for barley. The other pesticides were detected in amounts lower than the MRLs (EU-MRL-Database 2022). Piperonyl butoxide occurred frequently and presented the highest levels among the groups of pesticides. Piperonyl butoxide enhances the potency of certain pesticides such as carbamates, pyrethrins and pyrethroids but has no pesticide activity of its own (Basak et al. 2021). It has been demonstrated that this insecticide synergist can induce the formation of liver tumours in mice *via* the constitutive androstane receptor, which is qualitatively not plausible for humans due to the lack of effect on replicative DNA synthesis in human hepatocytes (Lake et al.

**Table 3.** Occurrences and levels of pesticide residues detected in wet brewery's spent grains intended for dairy cattle nutrition.

Analyte	Occurrence (%) <sup>a</sup>		Concentration ( $\mu\text{g kg}^{-1}$ ) <sup>b</sup>					WHO classification by hazard <sup>c</sup> / listed as highly hazardous pesticides by PAN <sup>d</sup> (+)	MRL <sup>e</sup> ( $\mu\text{g kg}^{-1}$ )	Use	References
	Positive samples	$\geq$ MRL <sup>c</sup>	Mean $\pm$ SD	Median	Range						
Ametoctradin	23.8	9.5	12.0 $\pm$ 12.7	4.45	4.45–33.8	III	10	Fungicide	Dreiner et al. (2018)		
Azoxystrobin	61.9	0.0	7.17 $\pm$ 5.15	2.8	6.20–17.3	U	1500	Fungicide	Rodrigues et al. (2013)		
Benzovindiflupyr	47.6	0.0	1.40 $\pm$ 0.00	1.4	1.4–1.40	II	1500	Fungicide	Yao et al. (2021)		
Bixafen	95.2	0.0	7.51 $\pm$ 3.96	4.65	4.65–18.1	–	400	Fungicide	Lazzari et al. (2012)		
Boscalid	42.9	0.0	9.75 $\pm$ 16	4.45	4.45–52.1	U	4000	Fungicide	Xu et al. (2020)		
BTS 44595 (Metabolite of prochloraz)	38.1	0.0	3.65 $\pm$ 4.26	2.15	2.15–14.2	II	30	Fungicide	EFSA (2018)		
Fluopyram	100	0.0	24.7 $\pm$ 15.8	19.8	12.5–75.6	III	200	Fungicide	Rathod et al. (2022)		
Fluxapyroxad	95.2	0.0	8.45 $\pm$ 6.22	6.84	2.65–24.5	III	3,000	Fungicide	McClean and Hollaway (2019)		
Isopyrazam	19.0	0.0	1.85 $\pm$ 0.00	1.85	1.85–1.85	II/+	600	Fungicide	Harp et al. (2011)		
Mandipropamid	90.5	14.3	11.8 $\pm$ 22.1	3.9	3.90–99.1	U	10	Fungicide	Lamberth et al. (2008)		
Metrafenone	71.4	0.0	30.4 $\pm$ 45.6	15.7	2.48–191	U	600	Fungicide	Opalski et al. (2006)		
Piperonyl butoxide	95.2	N/A	116 $\pm$ 58.8	80.1	15.5–254	U	N/A	Insecticide synergist	Basak et al. (2021)		
Pirimiphos-methyl	47.6	0.0	3.57 $\pm$ 2.61	2.54	1.65–9.29	II/+	5000	Insecticide	Huang and Subramanyam (2005)		
Pyraclostrobin	4.8	0.0	3.85 $\pm$ 0	–	3.85	–	1000	Fungicide	Kanungo and Joshi (2014)		
Tebuconazole	85.7	0.0	10.1 $\pm$ 3.35	9.41	1.24–16.7	II/+	2000	Fungicide	Sooväli and Koppel (2010)		
Trifloxystrobin	14.3	0.0	7.04 $\pm$ 8.73	2	2–17.1	U	500	Fungicide	EFSA (2017)		
Total	100	N/A	208 $\pm$ 113	201	92.6–572	N/A	N/A				

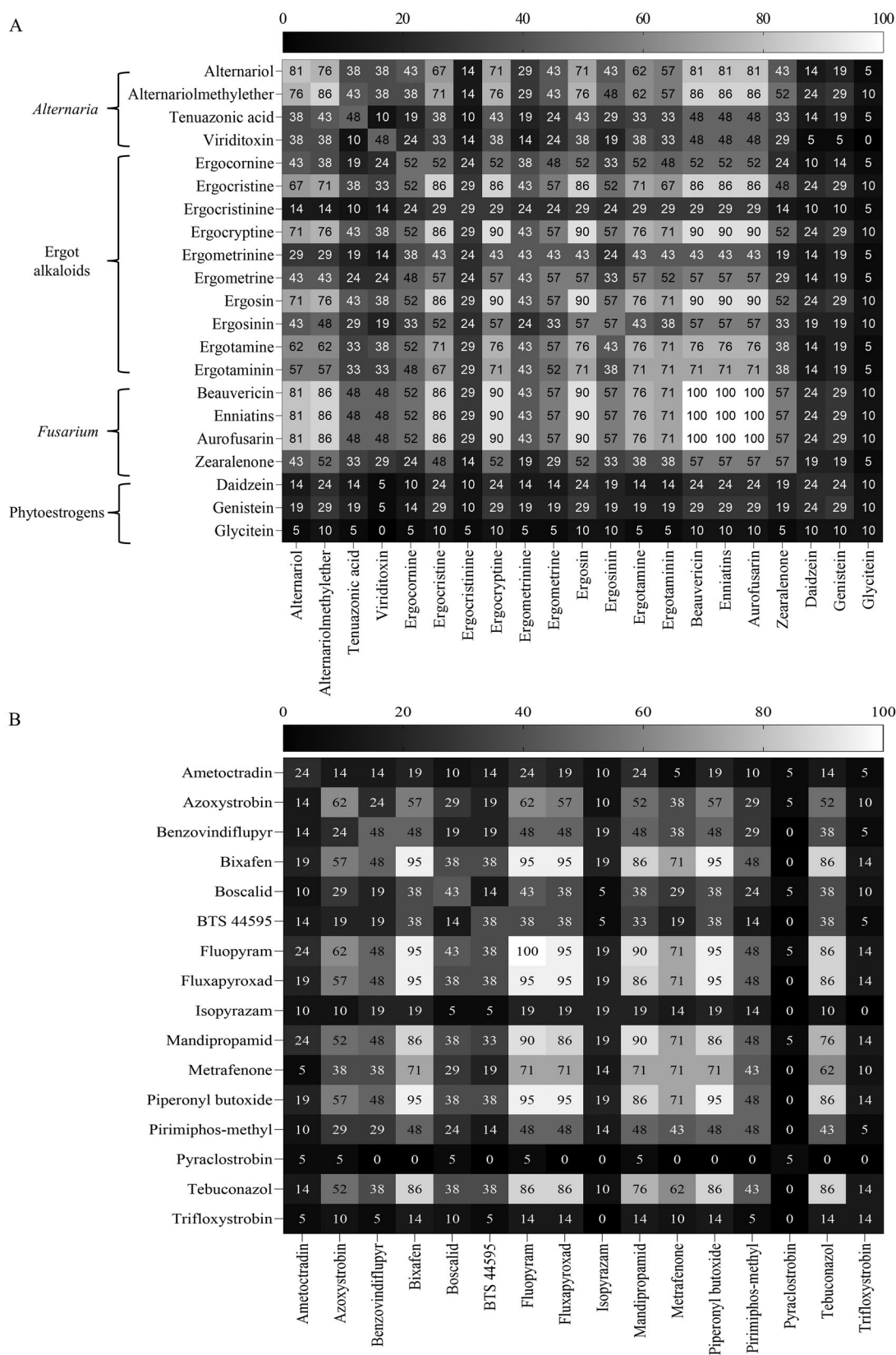
<sup>a</sup>n = 21 representative samples of wet barley-derived brewery spent grains from Austria, considered as positive values > limit of detection (LOD); <sup>b</sup>calculations without data < LOD. In case values > LOD and < limit of quantification (LOQ), LOQ/2 was used for the calculation; <sup>c</sup>WHO classification of pesticides by hazard. Ia (Extremely hazardous), Ib (highly hazardous), II (moderately hazardous), III (slightly hazardous) and U (unlikely to present acute hazard) (WHO 2020); <sup>d</sup>according to pesticide action network international (PAN 2021); <sup>e</sup>maximal residue level for barley according to the European Union guidelines expressed at 88% DM (EU-MRL-Database, 2022), N/A: Not available/not apply.

2020). This insecticide synergist is not a cholinesterase inhibitor and has low toxicity; it is also employed for other purposes than crop protection. It may also be used in conjunction with flea or tick dips, collars and oral medications in farm animals (Keane 1999). A widespread use of fluopyram, a pyridinyl ethylbenzamide, applied as a broad-spectrum fungicide with nematicide activity (Becker et al. 2020; Rathod et al. 2022) was recorded in this study, because it was detected in all the samples. The literature suggests that the high persistence of fluopyram in the environment (soil and water/sediment) can present risks for human, animal and soil health. The fate of this fungicide in diverse soil environments is still to be studied (Rathod et al. 2022). Fluopyram is authorised for use on crops that might be fed to livestock (EFSA 2020). A feeding study (Schoening and Wolters 2008 Ref: MR-07/367 (unpublished) cited by Lunn (2010) investigated the residue depuration of fluopyram in lactating dairy cattle, finding fluopyram and its metabolites in milk and different tissues. Barley has frequently been found contaminated with traces of fungicides (Palladino et al. 2021). In terms of risk, according to the WHO, five of the detected biocides here (benzovindiflupyr, prochloraz [confirmed by its metabolite of BTS 44595], isopyrazam, pirimiphos-methyl and tebuconazole) were classified as moderately hazardous (II) (WHO 2020). The last three are included in the list of highly hazardous pesticides (HHP) of the Pesticide Action Network (PAN 2021). Isopyrazam was added to the HHP list in 2011, it is considered likely carcinogenic for humans, very persistent in water, soil or sediment and very toxic for aquatic organisms (<50 ng/L) (US EPA 2017; Yao et al. 2018). Pirimiphos methyl was added to the HHP list in 2009, which is considered highly toxic for bees (Barnett et al. 2007; PAN 2021). Also, in the HHP list, tebuconazole can induce acute toxicity and long-term effects (PAN 2021) and can have ecotoxicological effects on an aquatic decomposer-detritivore system (Zubrod et al. 2011). Other detected pesticides like ametoctradin, fluopyram and fluxapyroxad are classified as slightly hazardous (III). In contrast, azoxystrobin, boscalid, mandipropamid, metrafenone, piperonyl butoxide and

trifloxystrobin are grouped as unlikely to present acute hazards (WHO 2020). It has been suggested that reduction in pesticide levels during the malting and brewing processes reduced considerably the risks of contaminating beer with pesticides and only a few pesticides remained without being removed or resolved (Navarro et al. 2007; Kong et al. 2016). Several pesticides found during beer production are adsorbed onto the spent grain after mashing. Moreover, some pesticides are degraded or transformed during boiling and fermentation, indicating that such reduction was caused primarily by adsorption, pyrolysis, and hydrolysis (Inoue et al. 2011; Xi et al. 2014). In the European Union, pesticide residue levels in particular plant and animal-derived foods and feeds, have been set by the Commission (EC) No 396/2005 (EC 2005). Information concerning MRLs and toxicity is available in the EU Pesticide database (EU-MRL-Database 2022).

### ***Co-occurrence of fungal toxins, phytoestrogens and pesticide residues***

Bioaccumulation rates and effects of long-term exposure to contaminant mixtures are unpredictable and should be investigated through the feed and food production chain. The co-contamination of several natural and synthetic contaminants was found (Figure 2(B)). All the samples presented co-contamination with several fungal secondary metabolites, fluctuating from 22 to 51 fungal metabolites per sample; 34 on average. Similarly, a broad spectrum of co-contamination with fungal and other metabolites has been observed in different complex matrices of feed-stuffs such as silage, pastures, concentrate feed, and total mix rations (Shimshoni et al. 2013; Nichea et al. 2015; Kemboi et al. 2020; Awapak et al. 2021). Interestingly, we generated data concerning pesticide residues, which occurred with an average of nine compounds per sample, ranging from six to twelve (Figure 2(B)). Additionally, Figure 3 illustrates co-occurrence matrices of mycotoxins (with recurrence over 25%) and phytoestrogens (Figure 3(A)) as well as pesticides (Figure 3(B)). All the samples evidenced that three fusarial emerging mycotoxins, aurofusarin, beauvericin and enniatins, were



**Figure 3.** Heatmaps of the co-occurrence (%) of (A) fungal contaminants (with occurrence rate > 25%) and phytoestrogens as well as (B) of pesticide residues detected in wet brewery's spent grains intended for the nutrition of dairy cattle in Austria.



detected and co-occurred with ZEN in 57% of the samples. More than 50% of the BSG presented co-contamination with several ergot alkaloids. Combinations between mycoestrogens derived from *Alternaria* AOH, AME, and TeA with ZEN were 43%, 52% and 33%, respectively. Co-occurrences between the mentioned mycoestrogens and the detected phytoestrogens (daidzein, genistein and glycitein) were lower than 30% (Figure 3(A)). Concerning pesticide residues, over 60% of the samples presented mixtures of fluopyram, fluxapyroxad, mandipropamid, metrafenone, piperonyl butoxide and tebuconazole. All the samples containing the insecticide for storage pirimiphos-methyl (48%) contained the semisynthetic synergist piperonyl butoxide: both compounds has been recently found in cereal samples from Croatia (Kovač et al. 2021).

These outcomes evidenced the ubiquitous presence of mixtures of multiple natural and synthetic chemicals in this by-product, linked to the feed and food supply chain. Although the occurrence of several contaminants was high, the concentrations found were low and under the legal limits (GVs and MRLs). The individual concentrations indeed do not represent an acute or critical risk for farm animals and human consumers. However, it is known that the combined effect of several co-occurring toxins and endocrine disruptors may be additive, synergistic or antagonistic, varying by type of compound or/and concentration (Guo et al. 2020). Such biological effects of toxin mixtures on animal and human health have been growing notably in recent years, but related knowledge is still overall scarce (Battilani et al. 2020; Gil-Serna et al. 2014; Smith et al. 2016; Weaver et al. 2020). Toxicological interactions have been described among mycotoxins, phytoestrogens and pesticides (Hessenberger et al. 2017; Vejdovszky, Hahn, et al. 2017a, Vejdovszky, Schmidt, et al. 2017a,b; Eze et al. 2019). For example, it is known that the interaction of diverse kinds of natural and synthetic xenobiotics, such mycotoxins, plant metabolites and chemical biocides, can also shape microbiota composition, which influences the health and metabolic status of the host (Lindell et al. 2022). The relevance of the co-occurrence (in real-world situations) of natural and synthetic chemicals has

to be addressed by toxicologists (Warne and Hawker 1995; Groten et al. 2001; Mattsson 2007). Nowadays, advances in analytic methods allow for evaluating hundreds of natural and synthetic pollutants, achieving high performances (LOD, LOQ and recovery) (Steiner et al. 2020; Sulyok et al. 2020; Steiner et al. 2021). Multi-toxin and multi-metabolites analysis has been used during the last decade to bring more insights into the complex field of mixture toxicology (Groten et al. 2001; Battilani et al. 2020; Martin et al. 2021). The evidenced ubiquitous presence of mixtures of pesticides suggests an extended application of this kind of substances in barley intended for beer production. This could also indicate that multiple biocides are being incorporated constantly at low levels in the feed and food chain, which can result in negative ecological and toxicological consequences (Mishra et al. 2014; Rivera-Becerril et al. 2017; Vanbergen 2021; Panico et al. 2022). Pesticide interactions lead mainly to synergic effects. Mixture effects differ depending on the dose and/or physiological target. Thus, more research and data for this important and exciting field are still required (Rizzati et al. 2016).

## Conclusion

This study provides insights into the widespread occurrence of cocktails of mycotoxins, phytoestrogens and pesticides in wet BSG. Mycotoxins/metabolites produced by the genera *Fusarium*, *Aspergillus*, and *Alternaria* were detected in all the samples. Ergot alkaloids were also frequently found (90%). *Penicillium* secondary metabolites, associated primarily with storage contamination, were present in 48% of the samples and showed the highest average concentration among the groups of fungal compounds. The storage-associated contamination leads to the necessity to improve strategies for preserving wet BSG in the farms. Additionally, we demonstrate the ubiquitous co-occurrence of several pesticide residues (at least six per sample, primarily fungicides). Two of them (ametoctradin and mandipropamid) exceeded the EU MRLs. Some pesticides (azoxystrobin, bixafen, fluopyram, fluxapyroxad, mandipropamid, metrafenone, piperonyl butoxide and tebuconazole) showed high occurrences (>60%),

which could suggest a common and extended use on food/feed crops of the mentioned pesticides and incorporation of these biocides into the feed/food chain and into the agroecosystems. Although the vast majority (88%) of the detected pesticides presented low concentrations, the potential combined effects of such biocide mixtures and natural toxins are unpredictable and should be subject to future studies. Further investigations with a larger number of samples and evaluation of BSG together with other feeds/foods is highly advocated.

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## Disclosure statement

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