



# Multitoxin analysis of *Aspergillus clavatus*-infected feed samples implicated in two outbreaks of neuromycotoxicosis in cattle in South Africa

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*Aspergillus clavatus* intoxication is a highly fatal neuromycotoxicosis of ruminants, especially cattle. It is caused by the ingestion of infected sprouting grain and sorghum beer residue. Locomotor disturbances, tremors and paralysis are observed. Histologically, degeneration and necrosis of larger neurons in the medulla oblongata, the midbrain, the thalamus and the ventral horns of the spinal cord are observed. Although a range of mycotoxins such as patulin, cytochalasin E and pseurotin A have been isolated, there is limited information on which specific mycotoxin or group of mycotoxins are involved during outbreaks of intoxication in livestock. In the present study, two outbreaks of *A. clavatus* poisoning in cattle are briefly described. Feed samples were collected for fungal identification, and culture and multitoxin analysis. A range of fungal metabolites were detected, and the estimated concentrations ( $\mu\text{g}/\text{kg}$ ) are provided. Both the sprouting barley and brewer's grain were predominantly infected with *A. clavatus* and, to a lesser extent, *Rhizopus arrhizus*. The only common *Aspergillus* secondary metabolite present in all the samples was pseurotin A. Patulin and cytochalasin E were present in the sprouting barley samples, as well as the *A. clavatus* isolates cultured on malt extract agar for 2 weeks; however, neither of these mycotoxins could be detected in the brewer's grain sample.

## Introduction

Ingestion of *Aspergillus clavatus*-infected feed may induce a highly fatal neuromycotoxicosis in ruminants, especially in cattle; however, sheep have also been poisoned (Gilmour *et al.* 1989; Kellerman *et al.* 2005; Riet-Correa *et al.* 2013; Schlosberg *et al.* 1991). This intoxication occurs sporadically in South Africa (Kellerman *et al.* 2005), but has also been reported in other countries (Riet-Correa *et al.* 2013; Sabater-Vilar *et al.* 2004; Schlosberg *et al.* 1991). Infected sprouting grains and malt are usually the cause, although, in South Africa, sorghum beer residue (*maroek*) has also been implicated. *Aspergillus clavatus* is a saprophytic fungus that often overgrows other fungal contaminants in the feed and is then visible as a greenish-blue mould (El-Hage & Lancaster 2004; Kellerman *et al.* 1976; Kellerman *et al.* 1984).

The clinical signs of *A. clavatus* poisoning in cattle include: hypersensitivity, tremors (which are not consistently present), ataxia, progressive paresis and paralysis. Affected animals walk with a stiff-legged gait, take short steps and sometimes knuckle over at the fetlocks. They may stagger and fall before paralysis and death set in. Neurological signs that are not evident at rest can be precipitated following exercise. In addition, drooling of saliva and constipation have also been reported (El-Hage & Lancaster 2004; Kellerman *et al.* 1976; Kellerman *et al.* 1984). Chronically affected cattle can survive for several weeks. There appears to be an extended latent period or a cumulative effect. During the experimental reproduction of this neurotoxicity, Kellerman and co-workers (1976) noted that cattle only became sick 7–10 days after commencement of feeding the poisonous ration or sorghum beer residue. El-Hage and Lancaster (2004) reported that new cases appeared up to 18 days after removal of infected sprouted barley.

Gross lesions are visible in the skeletal muscles of severely affected animals that exhibit clinical signs for longer periods (Kellerman *et al.* 2005). A greyish-white appearance of the larger muscles of the hindlimbs and forelimbs, especially near their origins and insertions, has been reported (Kellerman *et al.* 1976; Loretto *et al.* 2003; Riet-Correa *et al.* 2013). The most important microscopic lesions, which are also useful for diagnostic confirmation, occur in the brain and spinal cord. Degeneration and necrosis of larger neurons (chromatolysis and cytoplasmic vacuolation) in nuclei of the medulla oblongata, the midbrain, the thalamus and the ventral horns of the spinal cord have been described (Kellerman *et al.* 1976; Loretto *et al.* 2003). Furthermore,

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hyaline degeneration and necrosis of skeletal muscles and myocardial degeneration and necrosis have also been noted (Kellerman *et al.* 1976; Loretta *et al.* 2003).

*Aspergillus clavatus* synthesises mycotoxins such as: patulin (Lopez-Diaz & Flannigan 1977); cytochalasin E (Büchi *et al.* 1973; Lopez-Diaz & Flannigan 1977); tremorgenic metabolites such as tryptoquivalone (Clardy *et al.* 1975) and tryptoquivaline and its related toxic metabolites (Büchi *et al.* 1977; Clardy *et al.* 1975).

## Case histories

### Case 1

In November 2013, a group of young heifers and bulls exhibited neurological signs such as: knuckling over at the fetlocks (Figure 1), muscle fasciculation, hyperaesthesia, weakness of the hind quarters, ataxia, stumbling and recumbency. When handled, some collapsed and displayed mild seizure-like activity with paddling, but when left undisturbed were able to rise again. The farmer produced sprouting barley to feed the animals. The sprouting barley was heavily contaminated with fungi (Figure 2); two samples



Source: Photo provided by authors

**FIGURE 1:** An affected heifer exhibiting flexion (over-knuckling) of the fetlocks.



Source: Photo provided by authors

**FIGURE 2:** *Aspergillus clavatus*-infected sprouting barley.

were collected and submitted for fungal identification and mycotoxicological analysis.

### Case 2

In March 2014, 12 Friesland dairy cows died acutely after exhibiting moderate muscle tremor, aggressive behaviour and paralysis. The cows were fed a total mixed ration containing silage, brewer's grain, maize stover and a commercial concentrate. A set of tissue samples was collected and submitted for histopathology. Distinctive microscopic lesions in the central nervous system confirmed *A. clavatus* poisoning. Later, three animals became recumbent with the hindlimbs extended in an abnormal position behind them. Eventually, a total of 32 highly prized dairy cows succumbed. The brewer's grain and commercial pelleted ration were submitted for mycological identification and mycotoxicological screening.

There is limited information on which specific toxin or group of toxins are involved during outbreaks of intoxication (Lopez-Diaz & Flannigan 1977; Sabater-Vilar *et al.* 2004). Although a range of mycotoxins have been isolated from toxic *A. clavatus* strains, none of the metabolites have been administered to ruminants in order to reproduce the intoxication (McKenzie *et al.* 2004). The objective of the present study was to submit feed samples implicated in the two recent outbreaks and *A. clavatus* isolates cultured from the barley for comprehensive screening for fungal metabolites.

## Materials and methods

### Fungal isolation

Two sprouting barley samples from Case 1 and one brewer's grain sample and one commercial pelleted ration from Case 2, were visually inspected and isolations made directly from surface fungal growth by plating small clumps of fungal spores and/or mycelia onto potato dextrose agar (PDA) (Merck, Germany) amended with penicillin (250 mg/L). Isolations were also made from the plant material. Seeds ( $n = 25$ ) and root segments ( $n = 25$ , 10 mm long) from the sprouting barley and malt residue ( $n = 50$ ) from brewer's grain were surface disinfected with 1% sodium hypochlorite (3 min), rinsed twice with sterile water, blotted dry and plated onto PDA amended with penicillin. Plates were incubated for 7–14 days, at 25 °C under cool-white fluorescent light. The plates were checked regularly and selected colonies were transferred to half-strength PDA. Isolates retrieved were identified based on culture and morphological characteristics. *Aspergillus* isolates were plated to Czapek Yeast Agar (CYA), Czapek Yeast Agar with 20% Sucrose (CY20S) and Malt Extract Agar (MEA) (Klich 2002) and incubated in the dark at 25 °C for 7 days. One plate of CYA was incubated at 37 °C for 7 days. *Rhizopus* isolates were plated to PDA and MEA and incubated in the dark at 25 °C and 37 °C for 7 days (Schipper 1984). Selected single-spore isolates were deposited in the culture collection of the National Collections of Fungi, Biosystematics Division, Plant Protection Research Institute, Agricultural Research Council (ARC-PPRI), Pretoria, South Africa.



## Mycotoxin analysis

To comply with international phytosanitary regulations the feed samples collected from both outbreaks (Barley 1, Barley 2 and Brewers grain; Table 1) were air dried and finely milled before being shipped to Austria for multitoxin screening. In addition, two *A. clavatus* isolates (PPRI 13831 and PPRI 13832; Table 1) were also submitted for mycotoxin screening. A standardised extraction procedure was followed (Sulyok *et al.* 2006). Briefly: 5 g of each sample were extracted for 90 min with 20 mL of acetonitrile/water/acetic acid (79:20:1, v/v/v) on a rotary shaker (GFL 3017, GFL, Burgwedel, Germany). The crude extracts were diluted 1+1 (v+v) with acetonitrile/water/acetic acid (20:79:1, v/v/v) and 5 µL of the diluted extract were injected.

Detection and quantification were performed with a liquid chromatography/tandem mass spectrometry (LC-MS/MS) system (QTrap5500, Applied Biosystems, Foster City, California, United States of America) equipped with an electrospray ionisation (ESI) source (TurboIonSpray, Agilent Technologies, Waldbronn, Germany) and an ultra-high performance liquid chromatography system (1290 Series, Agilent Technologies, Waldbronn, Germany) (Malachova *et al.* 2014; Vishwanath *et al.* 2009). Chromatographic separation was performed at 25 °C on a Gemini C18-column, 150 mm × 4.6 mm inner diameter (i.d.), 5 µm particle size, equipped with a C18 security guard cartridge, 4 mm × 3 mm i.d. (all from Phenomenex, Torrance, California, USA). Elution was carried out in binary gradient mode. Both mobile phases contained 5 mM ammonium acetate and were composed of methanol/water/acetic acid 10:89:1 (v/v/v; eluent A) and 97:2:1 (v/v/v; eluent B), respectively. After an initial time of 2 min at 100% A, the proportion of B was increased linearly to 50% within 3 min. Further linear increase of B to 100% within 9 min was followed by a hold-time of 4 min at 100% B and 2.5 min column re-equilibration at 100% A. The flow rate was 1000 µL/min. The ESI-MS/MS was performed in the scheduled multiple reaction monitoring (sMRM) mode both in positive and negative polarities in two separate chromatographic runs (Malachova *et al.* 2014; Vishwanath *et al.* 2009). The sMRM detection window of each analyte was set to the respective retention time ± 27 s and ± 42 s in positive and negative modes, respectively. The target scan time was set to 1s. Confirmation of positive analyte identification was obtained by the acquisition of two sMRMs per analyte (with the exception of moniliformin and 3-nitropropionic acid that each exhibit only one fragment ion), which yields 4.0 identification points according to commission decision 2002/657/EC (European Union 2002). Analyst® software version 1.5.1 (AB Sciex, Foster City, California, USA) was used to control the LC-MS/MS instrument, as well as for automatic and manual integration of the peak (Malachova *et al.* 2014; Vishwanath *et al.* 2009). Quantification of the > 300 metabolites included in the method was done based on linear, 1/x weighed calibration curves derived from the analysis of serial dilutions of a multi-analyte stock solution. Results were not corrected for apparent recoveries.

## Results

### Fungal identification

Both the sprouting barley and brewer's grain were predominantly infected with *A. clavatus* and, to a lesser extent, *Rhizopus arrhizus*. In contrast, the commercial pelleted ration yielded only *Rhizopus* and no *Aspergillus* colonies. Plated plant material from the sprouting barley resulted in 70% *A. clavatus* and 50% *R. arrhizus* colonies, whereas the brewer's grain resulted in 100% *A. clavatus* and 20% *R. arrhizus* colonies when plated on the respective agars. Several plant material segments were colonised by both fungi. *Aspergillus clavatus* isolates (PPRI 13831, 13832, 14650 and 14651) had colony diameters of 38 mm – 42 mm on CYA, 40 mm – 45 mm on MEA, 43 mm – 48 mm on CY20S and 20 mm on CYA at 37 °C. Stipes were 500 µm – 1500 µm × 20 µm – 30 µm, smooth-walled, colourless, and expanding gradually into clavate vesicles, 50 µm – 75 µm wide. Conidial zone extended from 50 µm – 150 µm down from the top of the vesicles; heads were uniseriate with phialides 7 µm – 9 µm × 3.5 µm. Conidia were smooth-walled, ellipsoidal, 4 µm – 6 µm × 3 µm – 4 µm, and dull turquoise in mass. Colonies of *R. arrhizus* (PPRI 14572, only one isolate selected for preservation) were greyish brown. Rhizoids were brownish. Sporangiophores on stolons were up to 1500 µm in length, up to 18 µm in width, brown, single or aggregated in small groups. Sporangia were greyish black, powdery in appearance, up to 175 µm in diameter. Sporangiospores were subglobose to ellipsoidal, with ridges on the surface, up to 8 µm in length.

### Mycotoxin analysis

The multitoxin screening results are tabulated (Table 1).

## Discussion

Patulin is considered to be a major contributor to the neurotoxicity induced by *A. clavatus* (Sabater-Vilar *et al.* 2004). Riet-Correa *et al.* (2013) suggested that patulin analysis can aid in confirming a diagnosis of *A. clavatus* poisoning. In the current investigations, patulin was detected in the sprouting barley samples collected during the first outbreak and both isolates also synthesised this metabolite. The concentrations were considered to be high, as maximum tolerable levels ranging from 10 µg/kg – 50 µg/kg have been set in the European Union for a range of apple-based and fruit-based commodities, including baby food (Commission Regulation 1881/2006). On the other hand, patulin could not be detected (< limit of detection [LOD]) in the brewer's grain sample collected during the second outbreak, despite histological confirmation of *A. clavatus* poisoning. Kellerman *et al.* (1976) could also not demonstrate patulin in toxic fractions isolated with column chromatography from an extract prepared from sorghum beer residue (*maroek*) collected during an outbreak in South Africa. However, despite mycological verification of *A. clavatus* contamination of the brewer's grain collected during the second outbreak, not all fungal strains are toxic, which might explain the absence of patulin in the sample (Loretti *et al.* 2003). Another common metabolite, cytochalasin

**TABLE 1:** Fungal metabolites and estimated concentrations ( $\mu\text{g}/\text{kg}$ ) in feed samples and *Aspergillus clavatus* isolates.

Analyte	Samples					LOD
	Case 1				Case 2: Brewer's grain	
	Barley 1	Barley 2	PPRI 13831 <sup>‡</sup>	PPRI 13832 <sup>‡</sup>		
Fumonisin B <sub>1</sub> <sup>a</sup>	1263	5107	< LOD	< LOD	< LOD	6
Fumonisin B <sub>2</sub> <sup>a</sup>	283	1384	< LOD	< LOD	< LOD	2
Fumonisin B <sub>3</sub> <sup>a</sup>	250	1362	< LOD	< LOD	< LOD	4
Fumonisin B <sub>4</sub> <sup>a</sup>	389	1244	< LOD	< LOD	< LOD	2
Deoxynivalenol <sup>a</sup>	< LOD	294	< LOD	< LOD	< LOD	5
DON-3-glucoside <sup>a</sup>	< LOD	12.5	< LOD	< LOD	< LOD	0.1
Nivalenol <sup>a</sup>	6.45	< LOD	< LOD	< LOD	< LOD	0.8
Zearalenone-4-sulphate <sup>a</sup>	NT	NT	< LOD	< LOD	< LOD	0.2
Zearalenone <sup>a</sup>	195	53.4	< LOD	< LOD	< LOD	0.2
$\alpha$ -Zearalenol <sup>a</sup>	7.18	< LOD	< LOD	< LOD	< LOD	0.5
$\beta$ -Zearalenol <sup>a</sup>	28.0	< LOD	< LOD	< LOD	< LOD	0.5
Patulin <sup>§</sup>	504	728	30 016	70 784	< LOD	75
Diacetoxyscirpenol <sup>b</sup>	5.47	< LOD	< LOD	< LOD	< LOD	1
Monoacetoxyscirpenol <sup>b</sup>	6.09	< LOD	< LOD	< LOD	< LOD	1.5
Moniliformin <sup>b</sup>	51.0	< LOD	< LOD	< LOD	< LOD	5
Enniatin A <sup>b</sup>	98.3	10.0	< LOD	< LOD	4.31	0.02
Enniatin A <sub>1</sub> <sup>b</sup>	47.7	15.8	< LOD	< LOD	20.13	0.02
Enniatin B <sup>b</sup>	12.5	20.4	< LOD	< LOD	20.03	0.01
Enniatin B <sub>1</sub> <sup>b</sup>	15.8	26.9	< LOD	< LOD	37.98	0.02
Beauvericin <sup>b</sup>	127	145	< LOD	< LOD	1	0.02
Fusarin C <sup>b</sup>	7638	6356	< LOD	< LOD	< LOD	5
Fusaric acid <sup>b</sup>	3754	29 659	< LOD	< LOD	< LOD	15
Fusarinolic acid <sup>b</sup>	8085	13 344	< LOD	< LOD	< LOD	15
Aurofusarin <sup>b</sup>	866	1076	< LOD	< LOD	18.39	1
Bikaverin <sup>b</sup>	19 960	37 463	< LOD	< LOD	15.23	1
Butenolide <sup>b</sup>	12.6	< LOD	< LOD	< LOD	< LOD	10
Culmorin <sup>b</sup>	40.7	83.4	< LOD	< LOD	< LOD	0.8
15-Hydroxyculmorin <sup>b</sup>	< LOD	80.2	< LOD	< LOD	< LOD	3
Epiequisetin <sup>b</sup>	54.3	24.3	< LOD	< LOD	< LOD	0.15
Equisetin <sup>b</sup>	337	202	< LOD	< LOD	8.47	0.15
Antibiotic Y <sup>b</sup>	34.9	< LOD	< LOD	< LOD	< LOD	0.8
Apicidin <sup>b</sup>	1.47	49.4	< LOD	< LOD	< LOD	2
Hydrolysed fumonisin B <sub>1</sub> <sup>b†</sup>	< LOD	220 500	< LOD	< LOD	< LOD	0.3
Siccanol <sup>b†</sup>	18 730	5758	< LOD	< LOD	4870	NA
Alternariol <sup>c</sup>	< LOD	7.72	< LOD	< LOD	< LOD	0.6
Alternariolmethylether <sup>c</sup>	0.45	9.68	< LOD	< LOD	1.73	0.1
Macrosporin <sup>c</sup>	< LOD	< LOD	< LOD	< LOD	21.65	0.2
Infectopyron <sup>c</sup>	1549	752	< LOD	< LOD	< LOD	4
Kojic acid <sup>d</sup>	113 840	1006	< LOD	< LOD	< LOD	10
3-Nitropropionic acid <sup>d</sup>	76.3	24.5	< LOD	< LOD	6.77	1.2
Cytochalasin E <sup>d§</sup>	6791	6734	7560	7860.8	< LOD	0.5
Pseurotin A <sup>d§</sup>	906	7727	1835.2	1148.3	50.30	1.5
Aspterric acid <sup>d</sup>	233	< LOD	< LOD	< LOD	< LOD	5
Gliotoxin <sup>d</sup>	< LOD	< LOD	< LOD	< LOD	< LOD	2
Bis(methylthio)gliotoxin <sup>d</sup>	< LOD	< LOD	< LOD	< LOD	< LOD	5
Tryptoquivaline F <sup>d†</sup>	26 610	47 460	< LOD	< LOD	< LOD	NA
15-dimethyl-2epi-fumiquinazolin A <sup>d†</sup>	50 490	67 300	< LOD	< LOD	< LOD	NA
Fumiquinazoline F <sup>d†</sup>	20 960	36 080	< LOD	< LOD	< LOD	NA
Griseofulvin <sup>e</sup>	122	< LOD	< LOD	< LOD	< LOD	0.2
Dechlorogriseofulvin <sup>e</sup>	118	< LOD	< LOD	< LOD	< LOD	0.2
Dihydrogriseofulvin <sup>e</sup>	42.6	< LOD	< LOD	< LOD	< LOD	0.2
Chaetoglobosin A <sup>e†</sup>	< LOD	153 100	181 000	205 000	< LOD	1
Roquefortine C <sup>e</sup>	28.2	5.53	< LOD	< LOD	< LOD	0.4
Curvularin <sup>e</sup>	5.74	< LOD	< LOD	< LOD	< LOD	0.3
Agroclavin <sup>e</sup>	< LOD	< LOD	< LOD	< LOD	< LOD	0.1

LOD, Limit of detection ( $\mu\text{g}/\text{kg}$ ); < LOD, below limit of detection; NT, not tested; NA, not available; PPRI, culture collection of the National Collections of Fungi, Biosystematics Division, Plant Protection Research Institute, Agricultural Research Council (ARC-PPRI), Pretoria, South Africa.

<sup>a</sup>, Major mycotoxins and derivatives; <sup>b</sup>, *Fusarium* metabolites; <sup>c</sup>, *Alternaria* metabolites; <sup>d</sup>, *Aspergillus* metabolites; <sup>e</sup>, *Penicillium* metabolites; <sup>†</sup>, Metabolites from other fungal species; <sup>§</sup>, Unspecified metabolites.

<sup>‡</sup>, Values denote peak area, as no standard is available; <sup>‡</sup>, isolates were cultured on malt extract agar for 2 weeks; <sup>§</sup>, major *Aspergillus clavatus* metabolites.

Table 1 continues on next page →

**TABLE 1 (Continues...):** Fungal metabolites and estimated concentrations ( $\mu\text{g}/\text{kg}$ ) in feed samples and *Aspergillus clavatus* isolates.

Analyte	Samples					LOD
	Case 1				Case 2: Brewer's grain	
	Barley 1	Barley 2	PPRI 13831‡	PPRI 13832‡		
Fumitremorgin C <sup>e</sup>	NT	NT	< LOD	< LOD	34.43	NA
Atpenin A <sub>5</sub> <sup>e</sup>	0.38	< LOD	< LOD	< LOD	< LOD	0.5
Trichodermin <sup>f</sup>	350	350	< LOD	< LOD	< LOD	10
Brevianamide F <sup>g</sup>	26.0	8.43	299.36	436.32	448.96	0.5
Rugulosovinin <sup>g</sup>	48.1	18.7	198.72	260.16	779.2	0.3
Emodin <sup>g</sup>	33.7	41.9	< LOD	< LOD	19.38	0.2
Tryptophol <sup>g</sup>	4117	646	< LOD	< LOD	3376	15

LOD, Limit of detection ( $\mu\text{g}/\text{kg}$ ); < LOD, below limit of detection; NT, not tested; NA, not available; PPRI, culture collection of the National Collections of Fungi, Biosystematics Division, Plant Protection Research Institute, Agricultural Research Council (ARC-PPRI), Pretoria, South Africa.

<sup>a</sup>, Major mycotoxins and derivatives; <sup>b</sup>, *Fusarium* metabolites; <sup>c</sup>, *Alternaria* metabolites; <sup>d</sup>, *Aspergillus* metabolites; <sup>e</sup>, *Penicillium* metabolites; <sup>f</sup>, Metabolites from other fungal species; <sup>g</sup>, Unspecified metabolites.

†, Values denote peak area, as no standard is available; ‡, isolates were cultured on malt extract agar for 2 weeks; §, major *Aspergillus clavatus* metabolites.

E, was present in the barley samples as well as the isolates, but was also below the limit of detection in the brewer's grain sample.

The only common *Aspergillus* secondary metabolite present in all the samples was pseurotin A, although at a comparatively lower concentration in the brewer's grain. At relatively high concentrations, pseurotin A is reported to be cytotoxic to cell lines (A2780 human ovarian carcinoma cells and rat pheochromocytoma cells [PC12 cells]). On the other hand, at lower concentrations it induces neurite outgrowth in PC12 cells. The authors concluded that pseurotin A may possess neuroprotective properties that may prevent or be useful to treat human diseases related to dysfunction of the nervous system such as senile dementia and Alzheimer's disease (Komagata *et al.* 1996). It is thus impossible to conclude that pseurotin A is involved in the neurotoxicity seen in cattle.

The tremorgenic compound, tryptoquivaline F was also present in the barley samples, but the concentration could not be determined, as no standard is available. No consistent pattern with respect to the presence of mycotoxins could be discerned when the incriminated feed samples collected during the two outbreaks were compared (Table 1).

Although mycological identification alone is usually insufficient to confirm a diagnosis and analytical tests are advisable, the question still remains as to which mycotoxin or combination of mycotoxins are involved. The specific mycotoxin or group of mycotoxins responsible for the neurotoxicity induced during these outbreaks could not be identified. Furthermore, when sampling feed for mycotoxin analysis, it should be remembered that the toxic portion of the feed could have all been consumed and mycotoxins are not homogeneously distributed in a feed sample (Sabater-Vilar *et al.* 2004).

In both cases investigated, the feed was overgrown by *A. clavatus*, possibly because this fungus grows and sporulates more aggressively. Flannigan (1986) reported that the optimum temperature and relative humidity for growth of *A. clavatus* is 20 °C – 35 °C and 93% – 98%, respectively. Higher temperatures in the hydroponically cultivated barley sprouts will favour growth and *A. clavatus* will dominate the fungal population (Lopez-Diaz & Flannigan 1977; McKenzie *et al.*

2004). At 25 °C, the synthesis of specifically cytochalasin E is favoured when compared to patulin production (Lopez-Diaz & Flannigan 1977). In the first outbreak, the stock owner changed the process of producing sprouting barley. Where the hydroponically spouted barley was usually produced in trays in a refrigerated container, the batch implicated as the cause of the neurological signs was not produced in a refrigerated environment where the temperature and humidity could be controlled. The nutritional advantage of feeding sprouting grain produced hydroponically is doubtful (McKenzie *et al.* 2004). This practice is fraught with danger and livestock veterinarians should warn farmers about the risks of fungal contamination.

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## Competing interests

The authors declare that they have no financial or personal relationship(s) that may have inappropriately influenced them in writing this article.

## Authors' contributions

C.J.B. (University of Pretoria) was the principal investigator, M.J.L. (Private practitioner) investigated the first outbreak, M.T. (Agricultural Research Council-Plant Protection Research Institute) performed the mycological culturing and identification and M.S. (University of Natural Resources and Life Sciences) conducted the multitoxin analysis. All authors read and approved the final manuscript.

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