Case study

**CLOSTRIDIUM TERTIUM ISOLATED FROM FEED**

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**Abstract**

Although *Clostridium tertium* is supposed to be a foodborne pathogen, the data on its detection in foodstuffs is scarce, and there are no reports on its isolation from feed. In this communication paper, the isolation of *C. tertium* from a sample of soya semolina is described. *C. tertium* may be important in differential diagnosis, when it is to be distinguished from *Clostridium perfringens*. It is a unique species due to the lack of key characteristics of the genus it belongs to because it grows in the presence of oxygen and does not produce toxins. It has been well-documented as a human pathogen, although its mechanisms of pathogenicity are still unknown. According to sporadic reports in veterinary medicine, it has been identified as a rare causative agent of infections in cattle, pigs, birds and marine mammals.

**Keywords**: *Clostridium tertium*, soya semolina, MALDI-TOF

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INTRODUCTION

Bacterium species of the Clostridium genus are endospore-forming, obligate anaerobes (or relatively oxygen-tolerant) widespread not only in solid and liquid environments (soil, sewage, surface waters, marine sediments, etc), but also in animal and human intestines and, eventually, in animal and plant products. Based on its rRNA structure, the genus comprises extremely heterogeneous species, many of which share phylogenetic similarities with some other bacterial genera (Collins et al., 1994). Owing to its capability to produce enteritis and enterotoxaemia in various domestic animals, Clostridium perfringens is the most important clostridium species in veterinary medicine. Animal feed is one of potential sources of infection. According to regulations on the microbiological criteria for animal feed quality, it is considered safe if
no *Clostridium perfringens* and *Clostridium botulinum* are detected in 50 g of a sample (Regulation on the Quality of Animal Feed, 2010). The isolation and identification of *C. perfringens* should be done in compliance with the EN ISO 7937 standard, which enables the precise identification and enumeration of the target species in food and animal feeding stuff. The identification of other members of *Clostridium* genus is not part of the routine procedure in laboratories for feed analysis in Serbia and is beyond their diagnostic capacity. For the above mentioned reasons, this case report is a result of an aspiration to satisfy the researchers’ curiosity, discover the identity of certain *Clostridium* isolates from feed and to broaden the knowledge about bacterium species (other than *C. perfringens*) present in animal feedstuffs and feed. The isolation and identification of *Clostridium tertium* is presented in this communication paper. To the best of our knowledge, *Clostridium tertium* has not yet been detected in animal feed samples, although it is sometimes present in food of animal origin.

**A REPORT ON A LABORATORY CASE**

**Sample:** Soya semolina.

**Isolation:** Clostridia were isolated following the instructions given in the EN ISO 7937:2010 Standard. For further confirmation, five colonies black in colour due to sulphite reduction - grown on TSC (tryptone-sulfite-cycloserine) agar (Biokar Diagnostics, France) were chosen. They were inoculated into thioglycollate broth and incubated for 24 hours at 37°C (Fig 1.A). After incubation, 5 drops of thioglycollate culture was inoculated into lactose-sulfite (LS) broth (Biokar Diagnostics, France) for *C. perfringens* confirmation. After 24 hours of incubation at 46°C, LS was examined for gas production and the presence of black colour (sediment of iron sulfite). The formation of black colour has been observed, but Durham’s tube was filled with gas to less than a 1/4 of its volume (Figure 1.B). According to ISO standard, the test in LS medium should be repeated in this case by transferring 5 drops of culture grown in LS broth to another test tube with the same medium, repeating the incubation in the same conditions. As the repeated test once again failed to confirm the presence of *Clostridium perfringens* species, the culture which grew in thioglycollate medium was transferred by streaking onto two plates with Columbia blood agar base with the addition of 5% of defibrinated sheep blood. The plates were incubated at 37°C, one in aerobic and the other in anaerobic conditions using GasPak EZ (Becton Dickinson and Company, USA). After 24 hours of incubation, the growth was noted only on the plate which was incubated in anaerobic conditions, which led to the conclusion that the species is a strict
anaerobe. However, after 48 hours, the growth was also observed in the dish incubated in aerobic conditions, which would have led to ambiguity if it had been a *Clostridium* species. The isolate formed little (about 1 mm in diameter), opaque colonies, surrounded by a zone of incomplete (a) haemolysis (Figure 1.C). It was confirmed that it was *Clostridium* genus by a negative test for catalase and the microscopic appearance of the Gram stained smears: Gram-positive rods with rounded ends were found and oval spores located terminally were rarely present in smears made from cultures grown in anaerobic conditions. Based on characteristic black colonies grown on sulfite cycloserine agar, Gram-stain morphology and negative catalase test results, the isolates were presumptively identified as *Clostridium* species.

Figure 1. A. Growth of the isolate in thioglycollate medium; B. Test in lactose-sulfite medium; C. Colonies of *Clostridium* sp. isolate on blood agar.

Due to the lack of tests for biochemical and molecular typing of isolates of *Clostridium* species other than *C. perfringens* in our laboratory, the isolate was further processed at the Institute of Public Health of Vojvodina in Novi Sad. The identification was performed with matrix-assisted laser desorption/ionization time-of-flight mass spectrometry (MALDI-TOF MS). MALDI-TOF MS has been adapted to generate protein mass spectra from whole bacteria and other microorganisms. These spectra can be compared to a reference database for rapid and accurate taxonomic classification of unknown organisms at the genus, species, and, in some cases, at strain levels. The isolate was identified by MALDI TOF as *Clostridium tertium* (Fig 2).
Figure 2. Spectra of *Clostridium tertium* isolate generated by MALDI-TOF Bruker flexControl software.

**COMMENT**

*Clostridium tertium* is abundant in soil, but is also found in animal and human intestines as well as in the commensal microbiome of the mouth cavity (Vanderhofstadt et al., 2010). It is a rare human pathogen. Reportedly, it was first described and its biochemical properties were studied in isolates from war wounds in the First World War (Ray et al., 2003). Therefore, *C. tertium* is considered capable of causing bacteraemia (Ray et al., 2003). In addition, it was found in persons with various ailments: meningitis, septic arthritis, enterocolitis, peritonitis, posttraumatic brain abscess, pneumonia, and necrotizing fasciitis and gangrene. *C. tertium* does not produce exotoxins and the mechanism of its virulence is not known (Ferrell and Tell, 2001; Ray et al., 2003; Vanderhofstadt et al., 2010). Moreover, its clinical importance is questionable since it is not entirely clear if it is a real pathogen or only a contaminant (Vanderhofstadt et al., 2010). It is supposed that *C. tertium* does damage to the gut mucosa when colonizing it (Ferrell and Tell, 2001), meaning it can penetrate into the bloodstream (Ray et al., 2003).

In veterinary medicine, *C. tertium* has been recognized as a causative agent of enteritis in cattle and pigs. AlMashat and Taylor in 1984 isolated similar
bacteria from cattle with enteritis and phenotypically identified it as *Sporolactobacillus* species (Silvera et al., 2003). In artificially infected cattle, this bacterium caused mild diarrhoea. Ferrell and Tell (2001) reported an isolation of *C. tertium* from faeces of *Trichoglossus moluccanus* that vomited and had blood in faeces. It was assumed that contaminated water was the source of infection and that a diet rich in carbohydrates is a favourable medium for bacterial fermentation. Šeol et al. (2006) were the first to accuse *C. tertium* of causing abscesses, osteomyelitis and, finally, death in a dolphin, which was the first detection of this bacterium in marine animals.

Postollec et al. (2012) detected nine *Clostridium* species in various foodstuff, but not *C. tertium* (although they reviewed certain data during its previous detection). A long time ago, in 1965, Goudkov and Sharpe first published a paper on *C. tertium* detected in cheese and milk. They claimed that in spite of unfavourable conditions for *C. tertium* growth in dairy products, it can spoil certain cheese types. Later (Fernández et al., 2015), this bacterium was listed as one of the three clostridial foodborne pathogens in cheese, along with *C. botulinum* and *C. perfringens*. Le Bourhis et al. (2005) even developed and validated PCR primers for *Clostridium* spp. detection in cheese.

*C. tertium* was also detected in meat samples (Ersöz and Coşansu, 2018). Search for *C. difficile* with the API20A (System for the identification of anaerobes and with serological Clostridium Difficile Test Kit) resulted in *C. tertium* detection in one out of 101 samples of meat products (beef and chicken) collected from the market (Ersöz and Coşansu, 2018). The same agent was successfully recovered from foie gras and was proved to be capable of growing during storage at 8°C (Prevost et al., 2013). It was confirmed that *C. tertium* spores can be inactivated in meat by hydrostatic pressure and bacteriocins (Kalchayanand et al., 2003).

*C. tertium* is resistant to high temperatures, it can grow under various atmospheric conditions, can cause diarrhea and is present in both healthy and diseased humans (Silvera et al., 2003). Since it is considered an intestinal commensal in animals, it remains unclear whether animal feed should be regarded as a potential source of infection.

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Authors’ contributions

DM completed the microbiological analyses of Soya semolina samples and isolated *Clostridium tertium*. MD identified *C. perfringens* using MALDI-TOF. DM, NA, MV prepared the manuscript, and NA did the reviewing, editing and supervision.

Competing interest

Authors declared no conflict of interests regarding the present paper.

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