

1 **Black aspergilli and ochratoxin A in foods.**

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10

11 **Abstract**

12 Ochratoxin A (OTA) is a potent nephrotoxin and carcinogen which is found in a wide variety of
13 common foods and beverages. The black aspergilli are distributed worldwide and are regarded as
14 common food spoilage fungi. These fungi are one of the more difficult groups concerning
15 classification and identification. New molecular approaches have shown that there is a high
16 biodiversity, but that species are occasionally difficult to recognise based solely on their phenotypic
17 characters. Only few species have been confirmed to be OTA producers in this group and fewer are
18 known to contaminate foods with this mycotoxin as a natural occurring contaminant. In this paper,
19 the OTA-producing species included in the *Aspergillus* section *Nigri* and the foods that they are
20 able to contaminate are reviewed in depth.

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26 **Introduction**

27 Ochratoxin A (OTA) is a potent nephrotoxin and carcinogen which is found in a wide variety of
28 common foods and beverages [1]. Cereals are considered the major source of human exposure to
29 OTA. For this reason many countries have set a limit for OTA in cereals [2] However, human
30 exposure to OTA is most likely coming from low level contamination of a wide range of different
31 foods. In order to prevent or minimize this risk, the European Union have established maximum
32 OTA levels for many foodstuffs such as cereals, coffee, dried vine fruit, liquorice, spices, wheat
33 gluten and wine [3]. Nonetheless, this regulatory status of OTA lacks consensus in other countries
34 [4].

35
36 More than a half century has elapsed since OTA was discovered as a metabolite of *Aspergillus*
37 *ochraceus* [5]. A few years later, *Penicillium verrucosum* (as *Penicillium viridicatum*) was
38 reported to produce also this mycotoxin [6, 7]. Traditionally, OTA contamination of food and feeds
39 was believed to be restricted to *A. ochraceus* and *P. verrucosum*, which affect mainly dried stored
40 foods and cereals respectively, in different regions of the world [8]. Currently, the list of OTA-
41 producing species has expanded and, consequently, the list of foods which can be contaminated
42 with this mycotoxin has also increased. However, although a high number of additional *Aspergillus*
43 species are able to produce OTA, few of them are known to contaminate foods with this mycotoxin
44 [9].

45
46 Of these latter species, those include in the *Aspergillus* section *Nigri* (black aspergilli) are becoming
47 of interest because some of them are able to colonize foods other than cereals and cereal by-
48 products and to contaminate them with OTA. In this paper, the OTA-producing species included in
49 this section and the foods that they are able to contaminate are reviewed in depth.

50

51 **The black aspergilli (*Aspergillus* section *Nigri*)**

52 The black aspergilli are distributed worldwide and are regarded as common food spoilage fungi.
53 Some species are widely used in the biotechnology industry and a few of them are involved in animal
54 or human mycoses. These fungi are one of the more difficult groups concerning classification and
55 identification, and several taxonomic schemes have been proposed [10]. In the last revision of the
56 genus *Aspergillus*, 27 species were accepted in the *Aspergillus* section *Nigri* [11]. Basic
57 information about these taxa based on the last taxonomic proposals [11,12, 13, 14,15] are
58 summarized in Table1. Some of these species are rare or recently described, and only a few strains
59 of them have been studied. Consequently, the distribution of these species it is not well known.

60

61 The main morphological character of the most common species (e.g. *A. niger*) included in this
62 section is the colour of the conidial head, which is black or some shade of black. For this reason,
63 these fungi form generally characteristic black or dark brown colonies (Figure 1). However, the
64 colour of the colonies varies not only in different species, but in the same strain, depending upon
65 the culture medium, the age of the culture, the abundance of sporulation or the production of
66 sclerotia, among other factors. Consequently, depending on these factors, some species can form
67 purple-brown, reddish-brown, dark greenish, yellow or orange colonies [16]. In the conidial head,
68 some species have a single layer of phialides on the vesicle and are named uniseriate species (Table
69 1 and Figure 1). However, most of them have biseriate conidiophores (Figure 1), showing a layer of
70 metulae on the vesicle and a second layer of phialides over them. These are named biseriate species
71 (Table 1).

72

73 New molecular approaches have shown that there is a high biodiversity in this group. However,
74 these species are difficult to recognise based solely on their phenotypic characters [16]. For
75 example, the taxa related to *A. niger* have always been extremely difficult to distinguish by

76 morphological means. In these fungi, the differences between the described species and varieties in
77 the proposed classifications are very subtle and the number of taxa varies from one author to
78 another [10]. For this group of fungi, it was proposed the *A. niger* aggregate. The term aggregate is
79 used in mycology for groups of closely related morphospecies only distinguishable with difficulty.
80 Nowadays, this group comprise ten species [11], including *A. brasiliensis*, *A. niger* and *A.*
81 *tubingensis* which are the most common species isolated from foods. Although there is a substantial
82 call for searching for new molecular and physiological markers that are usable in the classification
83 of black aspergilli, the concept of several recently described species was only based on genetic
84 differences at one locus [17].
85
86 Nevertheless, new polyphasic taxonomic approaches have been proposed for the classification and
87 identification of these fungi [11, 16]. These polyphasic studies include sequence analysis of some
88 genes such as ITS-5.8S rRNA region (ITS), and β -tubulin (BenA) and calmodulin (CaM) genes,
89 morphological analyses and characterization of extrolite profiles, among other characters. Although
90 ITS is considered the universal DNA barcode of fungi, these sequences do not contain enough
91 variation for distinguishing among all species in the *Aspergillus* section *Nigri*. A secondary
92 barcode or identification marker usually is needed to identify a black aspergilli culture to species
93 level with confidence [11]. For example, the uncommon species *A. lacticoffeatus* is characterized
94 by its hair brown to dark blonde colonies which is an important distinguishing feature in its
95 description [18] (Figure 2). However, *A. lacticoffeatus* is very close molecularly to *A. niger* sensu
96 stricto [18]. In fact, this species has been considered a colour mutant of *A. niger* [12]. They have
97 identical ITS sequences and can not be separated by their BenA sequences but can be distinguished
98 using CaM sequence data and have also a different extrolite profile [16]. This new taxonomic
99 approach has also allowed to determine that some black aspergilli which are used in the production
100 of Asian fermented foods and beverages, such as *A. luchuensis*, *A. coreanus*, *A. kawachii* and *A.*

101 *acidus* were the same species. *A. luchuensis* was selected as the correct name based on priority [11**,
102 12, 14*].

103

104 In the last ten years, following these taxonomical approaches, the number of described uniseriate
105 species has dramatically increased from two to eleven species (see Table 1). Only, *A. aculeatus* and
106 *A. japonicus* were recognized previously [10*, 18]. Consequently, nowadays, in order to accurately
107 identify black aspergilli, in addition to characterize the micromorphology (e.g. conidial head,
108 conidia) of an isolate is necessary to compare ITS and some protein-coding gene sequences such as
109 BenA and CaM with published sequences in curated International Databases [11**, 19**]. The use of
110 CaM as a temporary secondary identification marker in *Aspergillus* has been suggested [11**]. Using
111 these markers, these taxa can be divided into five main molecular clades (e.g. *A. aculeatus*, *A.*
112 *carbonarius*, *A. heteromorphus*, *A. homomorphus* and *A. niger* aggregate clades) (see Table 1).

113

114

115 **OTA producing species in the black aspergilli**

116 Only a few black aspergilli have been confirmed to be OTA producers and fewer are known to
117 contaminate foods with this mycotoxin as a natural occurring contaminant. In the biseriate species,
118 only five species are considered to be able to produce OTA. In the *A. niger* aggregate, the
119 phylogenetically close species *A. niger* and *A. welwitschiae* are considered OTA producers (Figure
120 3). They are black aspergilli morphologically indistinguishable.

121

122 The common species *A. niger* has a worldwide distribution and the reported percentage of OTA-
123 producing strains in this species is usually very low [10*]. It is one the most commonly reported
124 species from foods (e.g. sun dried products, fresh fruits, nuts, cereals) [1]. Nevertheless, since the
125 first description of OTA production by *A. niger* [20*] this species is achieving a greater significance

126 regarding OTA content in some food commodities such as grapes, raisins and wine, and also in
127 coffee. Interestingly, this species is perhaps the most important mold used in biotechnology. It is
128 worth noting that *A. niger* products hold the GRAS (Generally Regarded as Safe) status from the
129 Food and Drug Administration and is a widely applied industrial species for large-scale
130 biotechnological production of organic acids and enzymes in the food industry. However, some of
131 the most frequently used strains in industry were able to produce this toxin on media suggested for
132 citric acid production [21*]. Among other things, due to their potential for mycotoxin production, no
133 filamentous fungi has the QPS (Qualified Presumption of Safety) status proposed by the European
134 Food Safety Authority [22*].

135

136 Regarding *A. welwitschiae*, which was previously called *A. niger* or *A. niger* var. *phoenicis*, most
137 strains produce occasionally OTA. This species has not yet been reported frequently because this
138 old species name has been reintroduced in a recent study about the elucidation of the taxonomic
139 position of some black aspergilli involved in the awamori fermentation [14*]. Consequently, *A.*
140 *welwitschiae* is expected to be more frequently reported in a near future. It has been found on
141 Welwitschia plants, grapes, dried fruits, coffee, cocoa, and other sources and it has also a worldwide
142 distribution from all the continents [14*].

143

144 On the other hand, the rare species *A. lacticoffeatus* which is located in the same molecular subclade
145 as *A. niger* and *A. welwitschiae* (*A. niger/welwitschiae* subclade) [11**] is also able to produce OTA
146 over a wide range of temperatures [18, 23] (Figure 3). However, in this case, *A. lacticoffeatus* which
147 was isolated from coffee beans, forms distinctive coffee-with-milk coloured colonies (Figure 2).

148

149 Outside this group, *A. carbonarius* is very consistent in producing this mycotoxin, and non-OTA-
150 producing strains in this species are very rare [9, 24, 25*] (Figure 3). In some cases, the suspected

151 non-OTA producing isolates previously identified as *A. carbonarius* belonged actually to other
152 different black aspergilli species [26, 27]. The conidia of *A. carbonarius* are much larger ($> 6\mu\text{m}$)
153 than those of most of the species in section *Nigri*. A large number of studies have shown that *A.*
154 *carbonarius* is the main responsible source of OTA in wine or dried vine fruits from main
155 viticultural regions worldwide [28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39]. It is also considered a
156 potential source of OTA in coffee. This species has been reported from coffee beans from various
157 coffee-producing countries such as Brazil [40], Philippines [41], Thailand [42, 43] and Vietnam
158 [44]. However, *A. ochraceus* and *A. westerdijkiae* are considered the most significant source of
159 OTA contamination in some coffee varieties [40, 43]. Although it is not so common as *A. niger*, *A.*
160 *carbonarius* has been also reported from other foods such as figs, maize, paprika, peanuts among
161 others [1].

162

163 On the other hand, *A. sclerotium* which is phylogenetically close to *A. carbonarius*, has been
164 also confirmed as an OTA producer. This species has also large conidia and produce sclerotia.
165 Nevertheless, *A. sclerotium* is known from only one strain (CBS 115572) which was isolated
166 from green Arabica coffee beans in India [18].

167

168 Natural occurrence of OTA in maize and maize-based products is a worldwide problem. The
169 available information on the ochratoxigenic mycobiota and OTA presence in corn, corn based food
170 and feed is limited. Several surveys have been shown that *A. niger* and *A. ochraceus* could be the
171 main source of OTA [45]. It is well known that *P. verrucosum* is the major producer of OTA in
172 cereals such as wheat and barley in temperate and cold climates, and , although *A. ochraceus* has
173 been isolated from a wide range of cereals, records are rather infrequent [46, 47]. However, *A. niger*
174 is frequently isolated from maize [1, 48, 49, 50] and a high incidence of *A. carbonarius* has been
175 also reported in this product [49]. It has been speculated that both species could be a source of OTA

176 in maize and other food products in both tropical and subtropical zones of the world [51]. In fact,
177 both species were able to produce OTA in maize kernels from the fifth day of incubation over a
178 wide range of temperatures and water availabilities [52].

179

180 OTA-producing strains of both species have been also reported from cocoa beans. Several studies
181 have shown that *A. carbonarius* and *A. niger* are important fungal sources of OTA in cocoa [53, 54,
182 55]. However, ochratoxigenic strains of *Aspergillus melleus*, *A. westerdijkiae* and *A. ochraceus*
183 have also been reported [55]. Different molds may contaminate many stages in cocoa processing,
184 and poor practices may have a strong influence on the quality of the beans. OTA is found at all
185 stages of cocoa processing, with the major incidence during drying and storage. In fact, the
186 contamination of cocoa by OTA-producing fungi can already take place in the fermentation, but a
187 considerable increase in the numbers of these species, as well in ochratoxin A contamination is
188 observed during drying and storage [55, 56]. More systematic surveys on different stages of cocoa
189 production in other important cocoa producing countries are needed in order to confirm the OTA
190 fungal sources in these products.

191

192 In addition to cereals, coffee, dried vine fruit and wine, the European Union have established
193 maximum OTA levels also on other foodstuffs such as liquorice and spices, including dried spices,
194 *Piper* spp. (including white and black pepper), *Myristica fragrans* (nutmeg), *Zingiber officinale*
195 (ginger), *Curcuma longa* (turmeric), *Capsicum* spp. (including chillies, chilli powder, cayenne and
196 paprika) and mixtures of spices containing one of the above mentioned spices [3].

197

198 At present, the source of OTA in liquorice is not known. In recent studies in China [57, 58], some
199 *Penicillium* spp., such as *P. polonicum* and *P. chrysogenum*, among other fungal species have been
200 reported to be a possible source of OTA in this product. These authors claimed that *P. chrysogenum*

201 derived from surrounding environments was likely to be a stable contributor to high OTA level in
202 liquorice. However, none of these species are considered OTA producers [47, 59]. This should be
203 confirmed using proper chromatographic detection of OTA and accurate identification of the fungi.

204

205 Currently, the source of OTA in spices is also unknown. However, *A. niger* has been reported as the
206 most frequently isolated species in *Piper* spp [60, 61], *Myristica fragrans* [62], *Zingiber officinale*
207 [61], *Curcuma longa* [61] and *Capsicum* spp. [61, 63, 64, 65]. Some other potential OTA-producing
208 species such as *A. ochraceus* and *A. westerdijkiae* were also reported in some of these studies, but
209 less frequently. Interestingly, solid-state fermentation with *A. niger* is used in the process of pepper
210 peeling. So, some OTA-producing starter strains of this species could be the OTA source in this
211 product. This kind of fermentation is widely used in traditional Chinese food fermentation due to its
212 easy operation, low cost, and wide feasibility on farms and in the countryside [60].

213

214 Some strains of other black aspergilli such as *A. awamori*, *A. usamii* and *A. foetidus* have been also
215 cited as OTA producers [10, 66]. However, their identity has been questioned [14*, 18]. Most *A.*
216 *awamori* strains isolated from oriental food fermentation process could be accommodated into *A.*
217 *niger* group, such as *A. luchuensis*, *A. niger*, *A. tubingensis* or *A. welwitschiae*. In fact, the neotype
218 of *A. awamori* (CBS 557.65) did not originate from awamori fermentation and it was shown to be
219 identical with *A. welwitschiae* [14*].

220

221 The ability of *A. tubingensis* to produce OTA remains a controversial issue. Some strains of this
222 species have been cited as OTA producers [67, 68, 69]. However, *A. tubingensis* is not considered
223 an OTA producer [10*, 18, 21] (Figure 3). In fact, a high number of strains of this species tested by
224 other authors were not able to produce OTA [28*, 29, 37, 70, 71, 72, 73, 74]. Very recently, none of
225 the 261 *A. tubingensis* strains isolated from wine grapes was found to be ochratoxigenic when they

226 were analysed with UPLC-MS/MS [74]. Nevertheless, some of these isolates were initially
227 considered to be able to produce the toxin when they were screened by HPLC-FLD. Consequently,
228 OTA production by strains of *A. tubingensis* and their taxonomical identity, should be confirmed
229 using proper techniques. Similarly, none of the uniseriate species are considered to be able to
230 produce OTA [66, 71, 73, 75, 76, 77]. Although the ability of some of these species to produce
231 OTA has been mentioned [30, 78, 79, 80], this fact needs to be confirmed.

232

233 As an unusual OTA-producing species is proposed, an accurate identification of the isolate and a
234 proper detection of the OTA production must be carried out in different culture media and
235 conditions in order to confirm that is a new OTA fungal source (Figure 1). OTA confirmation by
236 mass spectrometry is recommended (Figure 4). In order to avoid the chronic problem of
237 misidentification of mycotoxigenic fungi some useful recommendations have been proposed [9, 72,
238 81].

239

240 **Conclusions**

241

242 *Aspergillus* section *Nigri* is one of the more difficult fungal groups concerning classification and
243 identification. Only a few of these fungi are able to produce OTA. Some of these OTA-producing
244 species can contaminate a wide variety of foods and beverages. Nowadays, we know that *A.*
245 *carbonarius* is the main responsible source of OTA in wine or dried vine fruits from main
246 viticultural regions worldwide. Although there is clear evidence of the participation of *A.*
247 *carbonarius* and *A. niger* on the OTA contamination of cocoa and coffee, their exactly role have not
248 been stated. On the other hand, it is not always clear which black aspergilli species are responsible
249 for OTA contamination in other foods. While *A. niger* and *A. welwitschiae* are usually reported
250 from a wide variety of foods, their role as a source of OTA is not well known. In fact, *A. niger* is

251 frequently isolated from some EU regulated foods such as liquorice and spices. Nevertheless, the
252 source of OTA in these foods has not been identified yet. More systematic research is needed to
253 confirm which black aspergilli species are responsible for the OTA contamination in these and other
254 commodities.

255

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584

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586

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589

590 **Conflict of interests**

591 None.

592

Table 1. Current accepted species in the *Aspergillus* section *Nigri*, their main ecological characteristics and OTA production.

Species, authorities and year of the description	Molecular clade	Conidial head	Ecology, main habitats and distribution	OTA production ^a / OTA source ^b
<i>A. aculeatinus</i> , Noonim <i>et al.</i> , 2008	<i>A. aculeatus</i>	Uniseriate	Coffee beans, Thailand	-
<i>A. aculeatus</i> , Iizuka, 1953	<i>A. aculeatus</i>	Uniseriate	Tropical soil, unknown, air, USA, <i>Lactuca sativa</i> , Indonesia, dead branches, Papua	-
<i>A. brasiliensis</i> , Varga <i>et al.</i> , 2007	<i>A. niger</i> aggregate	Biseriate	Soil, wine grapes, worldwide	-
<i>A. brunneoviolaceus</i> , Batista & Maia, 1957	<i>A. aculeatus</i>	Uniseriate	Air, Trinidad and Tobago, and USA	-
<i>A. carbonarius</i> , (Bainier) Thom, 1916	<i>A. carbonarius</i>	Biseriate	Soil, wine grapes, coffee, figs, maize, paprika, peanuts, worldwide	+ / Wine, dried vine fruits and other grape products (Cocoa, coffee) ^b
<i>A. costaricaensis</i> , Samson & Frisvad, 2004	<i>A. niger</i> aggregate	Biseriate	Soil, Costa Rica	-
<i>A. ellipticus</i> , Raper & Fennell, 1965	<i>A. heteromorphus</i>	Biseriate	Soil, Costa Rica	-
<i>A. eucalypticola</i> , Varga <i>et al.</i> , 2011	<i>A. niger</i> aggregate	Biseriate	Leaves of <i>Eucalyptus</i> sp. , Australia	-
<i>A. floridensis</i> , Jurjević <i>et al.</i> , 2012	<i>A. aculeatus</i>	Uniseriate	Air, USA and Martinique, soil, Japan, almonds USA	-
<i>A. heteromorphus</i> , Batista & Maia, 1957	<i>A. heteromorphus</i>	Biseriate	Fungal culture contaminant, Brazil	-
<i>A. homomorphus</i> , (Steiman <i>et al.</i>) Samson & Frisvad, 2004	<i>A. homomorphus</i>	Biseriate	Soil of death sea area, Israel	-
<i>A. ibericus</i> , Serra <i>et al.</i> , 2006	<i>A. carbonarius</i>	Biseriate	Wine grapes and raisins, Portugal and Spain	-
<i>A. indologenus</i> , Frisvad <i>et al.</i> , 2011	<i>A. aculeatus</i>	Uniseriate	Soil, India	-
<i>A. japonicus</i> , Saito, 1906	<i>A. aculeatus</i>	Uniseriate	Soil, Brazil	-
<i>A. labruscus</i> , Fungaro <i>et al.</i> , 2017	<i>A. homomorphus</i>	Uniseriate	Grapes for juice production, Brazil	-
<i>A. lacticoffeatus</i> , Frisvad & Samson, 2004	<i>A. niger</i> aggregate	Biseriate	Coffee beans, Indonesia and Venezuela	+ / (coffee) ^b
<i>A. luchuensis</i> , Inui, 1901	<i>A. niger</i> aggregate	Biseriate	East Asia, food fermentation environment	-
<i>A. neoniger</i> , Varga <i>et al.</i> , 2011	<i>A. niger</i> aggregate	Biseriate	Mangrove water, Venezuela, and desert sand, Namibia	-
<i>A. niger</i> , van Tieghem, 1867	<i>A. niger</i> aggregate	Biseriate	Worldwide, cosmopolitan fungus	+ / (many foods) ^b
<i>A. piperis</i> , Samson & Frisvad, 2004	<i>A. niger</i> aggregate	Biseriate	Grounded black pepper of tropical origin	-
<i>A. saccharolyticus</i> , Sørensen <i>et al.</i> , 2011	<i>A. homomorphus</i>	Uniseriate	Oak wood, Denmark	-
<i>A. sclerotii carbonarius</i> , Noonim <i>et al.</i> , 2008	<i>A. carbonarius</i>	Biseriate	Coffee beans, Thailand	-
<i>A. sclerotioniger</i> , Samson & Frisvad, 2004	<i>A. carbonarius</i>	Biseriate	Green Arabica coffee, India	+ / (coffee) ^b
<i>A. trinidadensis</i> , Jurjević <i>et al.</i> , 2012	<i>A. aculeatus</i>	Uniseriate	Air, Trinidad and Tobago, and USA	-
<i>A. tubingensis</i> , Mosseray, 1934	<i>A. niger</i> aggregate	Biseriate	Worldwide, cosmopolitan fungus	-
<i>A. uvarum</i> , Perrone <i>et al.</i> , 2008	<i>A. aculeatus</i>	Uniseriate	Wine grapes, Europe and Israel , air, USA	-
<i>A. vadensis</i> , Samson <i>et al.</i> , 2005	<i>A. niger</i> aggregate	Biseriate	Air, Egypt	-
<i>A. welwitschiae</i> , (Bresadola) Hennings apud Wehmer, 1907	<i>A. niger</i> aggregate	Biseriate	Grapes, dried fruits, coffee, cocoa, worldwide	+ / (many foods) ^b

594

^a OTA production: +, OTA producing species; -, non-OTA producing species

595

^b (potential source)

596

Data are from Cabañes & Bragulat [9], Samson *et al.* [11], Varga *et al.* [12], Jurjević *et al.* [13], Hong *et al.* [14], Fungaro *et al.* [15], Samson *et al.* [18], Taniwaki *et al.* [40], Copetti *et al.* [56].

Legends

Figure 1. Importance, detection and identification of black aspergilli in foods.

(This figure summarizes the main concepts discussed)

Figure 2. Colonial morphology of *A. lacticoffeatus* (CBS 101883), grown on Czapek Yeast extract Agar at 30°C for 10 days. Note the distinctive coffee-with-milk colour of the colony.

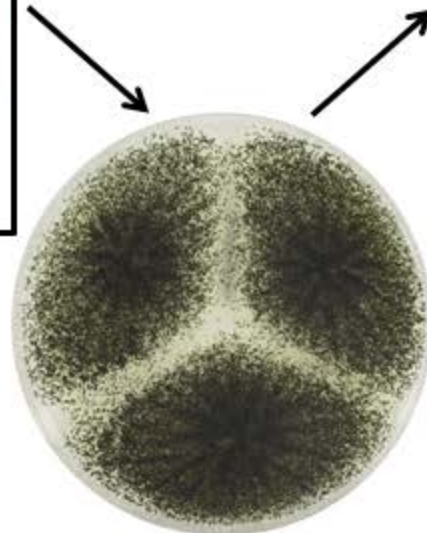
Figure 3. Selected chromatograms of fungal extracts analysed using HPLC coupled to a fluorescence detector of (a) an OTA-producing strain of *A. carbonarius*, (b) an OTA-producing strain of *A. niger*, (c) an OTA-producing strain of *A. lacticoffeatus* and (d) a non-OTA-producing strain of *A. tubingensis*.

Figure 4. Electrospray ionisation–mass spectrometry spectrum of OTA (a) (major ions: m/z 358.08 [MH-HCOOH]⁺, m/z 404.09 [MH]⁺ and m/z 426.07 [MNa]⁺) and selected extracted ion chromatograms of fungal extracts of an OTA-producing strain of *A. carbonarius* (b) and a non-OTA-producing strain of *A. carbonarius* (c) analysed using HPLC-MS. OTA standard (d).

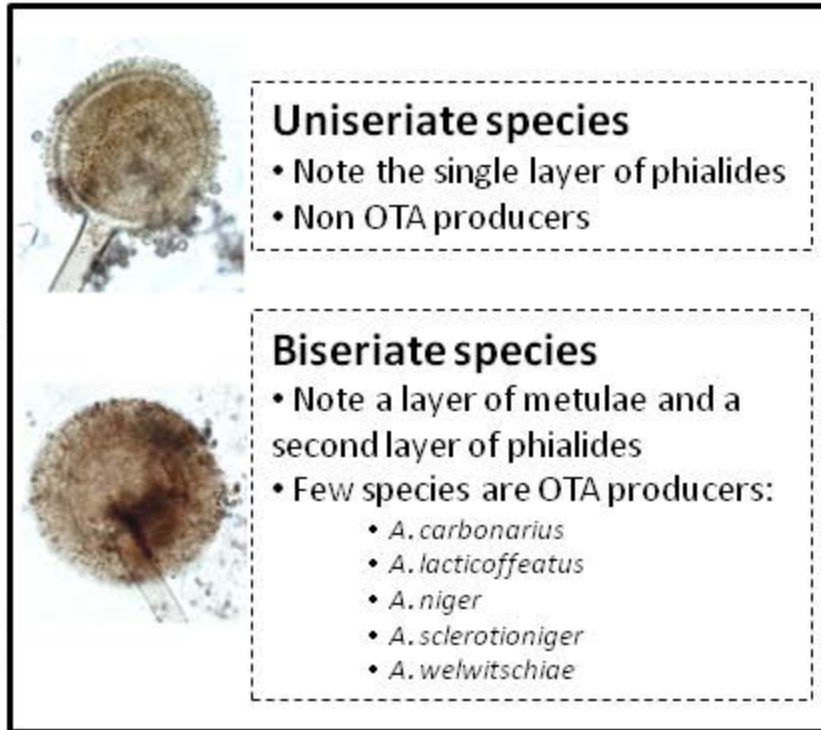
OTA contamination of food



A. carbonarius is the main OTA source in grape and grape products



Form black colonies



Uniseriate species

- Note the single layer of phialides
- Non OTA producers

Biseriate species

- Note a layer of metulae and a second layer of phialides
- Few species are OTA producers:
 - *A. carbonarius*
 - *A. lacticoffeatus*
 - *A. niger*
 - *A. sclerotiumniger*
 - *A. welwitschiae*

Assessment of a new OTA producing species

- Detection of OTA by HPLC-FLD
- Confirmation by MS
- Deposit the strain in a international culture collection

Identification

- DNA sequencing is needed
- β -tubulin and calmodulin genes
- BLAST search
- Curated International Databases
- Phylogenetic analysis



