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Suppression of Aflatoxin Production in *Aspergillus* Species by Selected Peanut (*Arachis hypogaea*) Stilbenoids

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Suppression of Aflatoxin Production in *Aspergillus* Species by Selected Peanut (*Arachis hypogaea*) Stilbenoids

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ABSTRACT: *Aspergillus flavus* is a soil fungus that commonly invades peanut seeds and often produces carcinogenic aflatoxins. Under favorable conditions, the fungus-challenged peanut plant produces and accumulates resveratrol and its prenylated derivatives in response to such an invasion. These prenylated stilbenoids are considered peanut antifungal phytoalexins. However, the mechanism of peanut–fungus interaction has not been sufficiently studied. We used pure peanut stilbenoids arachidin-1, arachidin-3, and chircanine A to study their effects on the viability of and metabolite production by several important toxigenic *Aspergillus* species. Significant reduction or virtually complete suppression of aflatoxin production was revealed in feeding experiments in *A. flavus*, *Aspergillus parasiticus*, and *Aspergillus nomius*. Changes in morphology, spore germination, and growth rate were observed in *A. flavus* exposed to the selected peanut stilbenoids. Elucidation of the mechanism of aflatoxin suppression by peanut stilbenoids could provide strategies for preventing plant invasion by the fungi that produce aflatoxins.

KEYWORDS: peanut, *Arachis hypogaea*, groundnut, phytoalexin, stilbenoid, arachidin-1, arachidin-3, chircanine A, aflatoxin, aflatoxin production, aflatoxin suppression, aflatoxin inhibition, *Aspergillus*, *Aspergillus flavus*, *Aspergillus parasiticus*, *Aspergillus nomius*

INTRODUCTION

Aflatoxins are among the most potent human and animal carcinogens known in nature.¹ It is estimated that over half of the world's population is chronically exposed to aflatoxins 7–10 (Figure 1).² Preharvest aflatoxin contamination of peanuts caused by the toxigenic soil fungi *Aspergillus flavus* and *Aspergillus parasiticus* has been a serious health and economic problem since the early 1960s, the period of aflatoxin discovery.^{3,4} Slow progress in resolving the issue is explained mainly by the complexity of peanut–fungus interactions.^{5,6} The mechanism of peanut defense is poorly understood, although there is sufficient evidence that the peanut plant protects itself from fungal invasion by promptly producing stilbene-derived phytoalexins.^{6–8} Fungi, in turn, produce phytoalexin-detoxifying enzymes to successfully invade the plant host.⁵ The involvement of fungal secondary metabolites in these intricate interactions has not been explored. However, the actions of peanut-derived resveratrol and various antioxidative compounds on *A. flavus* morphology and toxin formation has been reported by different scientific groups.^{9–11} Caffeic acid, **1** (Figure 1), at a 12 mM concentration added to a fat-based growth medium reduced >95% of aflatoxin production by *A. flavus* NRRL 3357, without affecting fungal growth.¹⁰ The action of caffeic acid¹⁰ and other antioxidants tested (gallic and tannic acids and methyl gallate) was attributed to the alleviation of oxidative stress in fungi.¹¹ Treatment of *A. flavus* with resveratrol, **2** (Figure 1), decreased aflatoxin production and the formation of conidia, the asexual spores of a fungus. In addition, this stilbenoid caused abnormal mycelial development and directly inhibited the expression of aflatoxin-biosynthetic-pathway cluster genes.⁹ Resveratrol content in resistant peanut lines was significantly higher than that in susceptible lines. At the same time, aflatoxin content was lower in the resistant lines

compared with that in the susceptible lines.¹² Based on that negative correlation, the authors suggested that resveratrol is strongly related to the resistance to aflatoxin production in peanut seeds. However, the production of resveratrol in peanut seeds is often accompanied by a prompt accumulation of prenylated stilbenoids¹³ with significantly higher antifungal activities compared with resveratrol.¹⁴ The authors did not report any prenylated stilbenoids in their samples, although the presence of these kinds of compounds is very likely on the basis of reports from research groups.^{5,13,15} A contribution to aflatoxin inhibition by these stilbenoids is also expected. However, the information on this issue is lacking.

The objective of the present research was to evaluate the abilities of some peanut prenylated stilbenoids (arachidin-1, **3**; arachidin-3, **5**; and chircanine A, **6**) to influence aflatoxin production in selected strains of important aflatoxin producers, namely, *A. flavus*, *A. parasiticus*, and *A. nomius*.

MATERIALS AND METHODS

Reagents, Materials, and Basic Apparatus. The HPLC-grade solvents used in the preparation of mobile phases and separations on silica gel were obtained from Fisher (Suwanee, GA). The HPLC-grade H₂O was prepared with a ZD20 four-bowl Milli-Q water system (Millipore, Burlington, MA). The HPLC-grade methanol used for media extraction was purchased from VWR (Suwanee, GA), and 48-well cell-culture plates (Nunc, Roskilde, Denmark) were used in all feeding experiments.

Reference Compounds. Pure, individual stilbenoids *trans*-arachidin-1, **3**; *trans*-arachidin-2, **4**; and *trans*-arachidin-3, **6**, were

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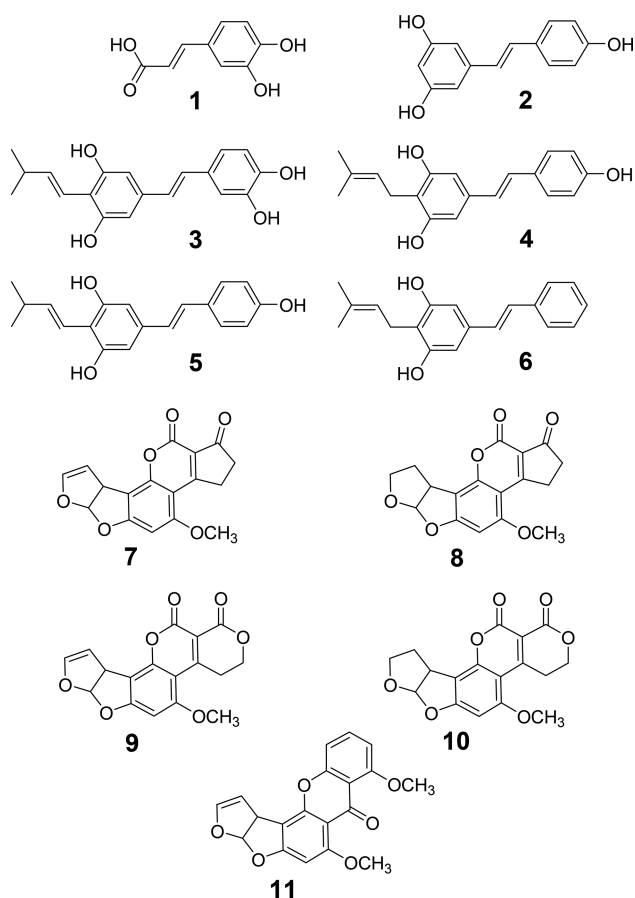


Figure 1. Structures of compounds discussed in the text. 1, *trans*-caffeic acid; 2, *trans*-resveratrol; 3, *trans*-arachidin-1; 4, *trans*-arachidin-2; 5, *trans*-arachidin-3; 6, chiricanine A; 7, aflatoxin B₁; 8, aflatoxin B₂; 9, aflatoxin G₁; 10, aflatoxin G₂; 11, O-methyl sterigmatocystin.

obtained as previously described¹⁶ except that preparative HPLC was used as a final purification step rather than preparative TLC. HPLC separation was achieved by using a 100 × 19 mm i.d., 5 μm, XTerra Prep RP18 OBD column (Waters, Milford, MA) and an isocratic mobile phase composed of CH₃CN, 2% HCOOH in H₂O, and H₂O (55, 3, and 42%, respectively). The flow rate was 8.0 mL/min.¹⁷ Chiricanine A, 6, was prepared as described¹⁷ (Figure 1). O-Methyl sterigmatocystin was purchased from Cayman Chemical (Ann Arbor, MI), and a certified solution of combined aflatoxins B₁, 7; B₂, 8; G₁, 9; and G₂, 10, was purchased from Supelco (Bellefonte, PA).

Fungi. *A. flavus* NRRL 3357, *A. flavus* NRRL 29487, *A. nomius* NRRL 13137, *A. parasiticus* NRRL 29580, and *A. parasiticus* 29602 were made available by the fungal collection of the National Peanut Research Laboratory, Agricultural Research Service (ARS), USDA (Dawson, GA).

Feeding Technique. The experiments comprised a total of 546 cell-culture wells (2 mL each) containing the essential stilbenoids arachidin-1, 3; arachidin-3, 5; and chiricanine A, 6, at 0.3 mM concentrations in 0.5 mL of potato dextrose broth or agar (each well received 46.8 μg of arachidin-1, 44.4 μg of arachidin-3, or 42.0 μg of chiricanine A). To prepare the spiked potato dextrose agar (PDA), the stilbenoids were dissolved in 96% EtOH, added to an appropriate amount of molten PDA agar, and then cooled to 45–50 °C, followed by intensive mixing with a magnetic stirrer. The concentration of EtOH in the media did not exceed 1%. Twenty microliters of fungal spores (10⁶/mL) were applied to each experimental well and thoroughly distributed on the surface of the agar with a glass rod. For comparison, a set of controls containing broth alone, broth with the individual phytoalexins but without the fungi, and broth with the individual fungi but without the phytoalexins was added to the

experimental setup. All the experiments were performed in duplicate or triplicate. Samples were collected every 24 h and kept frozen at –28 °C. The extraction of each sample was performed with 5 mL of MeOH at 22 ± 2 °C for 18 h without agitation in the dark. The filtered extracts were evaporated to dryness in a stream of N₂, redissolved in 300 μL of MeOH, and filtered again through a glass-fiber filter, and then aliquots of the filtrates were analyzed by HPLC-MS.

HPLC-DAD-MS Analyses. Separations of the well extracts were performed using a tandem HPLC-MS Surveyor system equipped with an MS Pump Plus, an Autosampler Plus, and a PDA Plus Detector (Thermo Electron Corporation, San Jose, CA) covering the 200–600 nm range, and a 100 × 4.6 mm i.d., 3.5 μm, XSelect HSS C18 analytical column (Waters, Milford, MA) was used. H₂O (A), MeOH (B), and 2% HCOOH in H₂O (C) were used in the following gradient: the initial conditions were 59% A/40% B/1% C, which was changed linearly to 10% A/89% B/1% C in 11 min, changed to 0% A/99% B/1% C in 0.01 min, held isocratic for 3 min, then changed to the initial conditions in 0.01 min, and held for 4 min before the next injection. The flow rate was 1.2 mL/min. The column was maintained at 40 °C.

The MS analyses were performed using a Finnigan LCQ Advantage MAX ion-trap mass spectrometer equipped with an ESI interface and operated with the Xcalibur version 1.4 software (Thermo Electron Corporation, San Jose, CA). The data were acquired in the full-scan mode (MS) from *m/z* 100–2000. The heated-capillary temperature was 250 °C, the APCI-vaporizer temperature was 380 °C, the sheath gas flow was 60 units, the auxiliary gas flow was 5 units, the capillary voltage was 53 V, and the source voltage was 4.5 kV. In the MS² analyses, the [M + H]⁺ ions observed for each chromatographic peak in the full-scan analyses were isolated and subjected to source collision-induced dissociation (CID) using a He buffer gas. In all CID analyses, the isolation width, relative fragmentation energy, relative activation *Q*, and activation time were: 1.2, 30 or 35%, 0.25, and 30 ms, respectively. The concentrations of *trans*-arachidin-1, *trans*-arachidin-3, chiricanine A, and O-methyl sterigmatocystin in the extracts were calculated by reference to the peak areas of the corresponding pure standards at 340, 335, 312, and 314 nm, respectively. High concentrations of all aflatoxins were determined at 362 nm. To determine the low concentrations of the toxins, the extracts were purified as previously described¹⁸ and subjected to aflatoxin analysis using an Acquity UPLC instrument equipped with a matching UPLC H-class Quaternary Solvent Manager; UPLC Sample Manager; UPLC Fluorescent Detector (FLR); and 50 × 2.1 mm i.d., 1.7 μm, Acquity UPLC BEH C18 column (Waters, Milford, MA). The mobile phase was composed of a water/MeOH/CH₃CN (64:23:13, v/v/v) mixture, and the flow rate was 0.25 mL/min. The column was maintained at 35 °C in the system column heater. The concentrations of the aflatoxins were determined by reference to the peak areas of the corresponding commercial standards (calibration curve). The detection limits were 0.15 ng/g for aflatoxins G₁ and B₁ and 0.02 ng/g for aflatoxins G₂ and B₂.

Preparation of Medium and Estimation of Morphological Changes in *A. flavus* NRRL 3357. Spore germination and hyphal growth of *Aspergillus flavus* NRRL 3357 were monitored over 20 h in the presence of the individual peanut phytoalexins arachidin-3, arachidin-2, and chiricanine-A. Stock solutions of these phytoalexins were prepared separately at 150 mM in ethanol and added to 5 mL test tubes containing 1 mL of potato dextrose broth (PDB), using only one phytoalexin per well at a final concentration of 0.3 mM; 2 μL of ethanol was added to 1 mL of PDB to be used as a control. *A. flavus* NRRL3357 was grown on potato-dextrose-agar (PDA) medium for 6 days at 30 °C, and its spores were harvested in sterile distilled water and passed through frits that were placed into a matching 1.5 mL SPE reservoir (both from Grace Davison Discovery Science, Deerfield, IL) in order to remove fragments of hyphae, the threadlike filaments forming the mycelium of a fungus. A suspension of 10⁶ spores/mL was prepared in sterile distilled water, and 40 μL of the spore suspension was added to each 5 mL test tube containing 1 mL of PDB supplemented with a phytoalexin and to the PDB control. For each phytoalexin and for the control, duplicate test tubes were used during

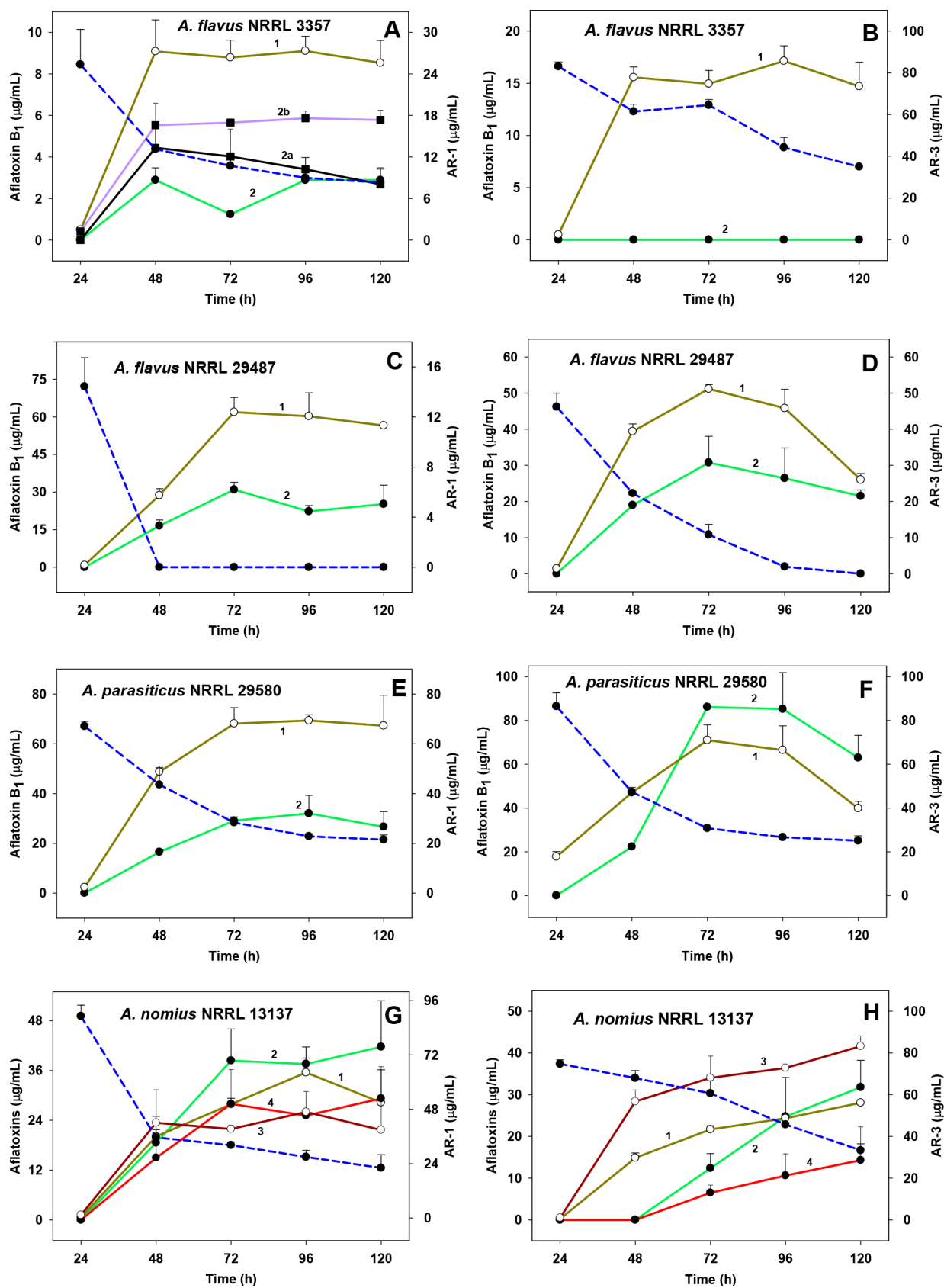


Figure 2. continued

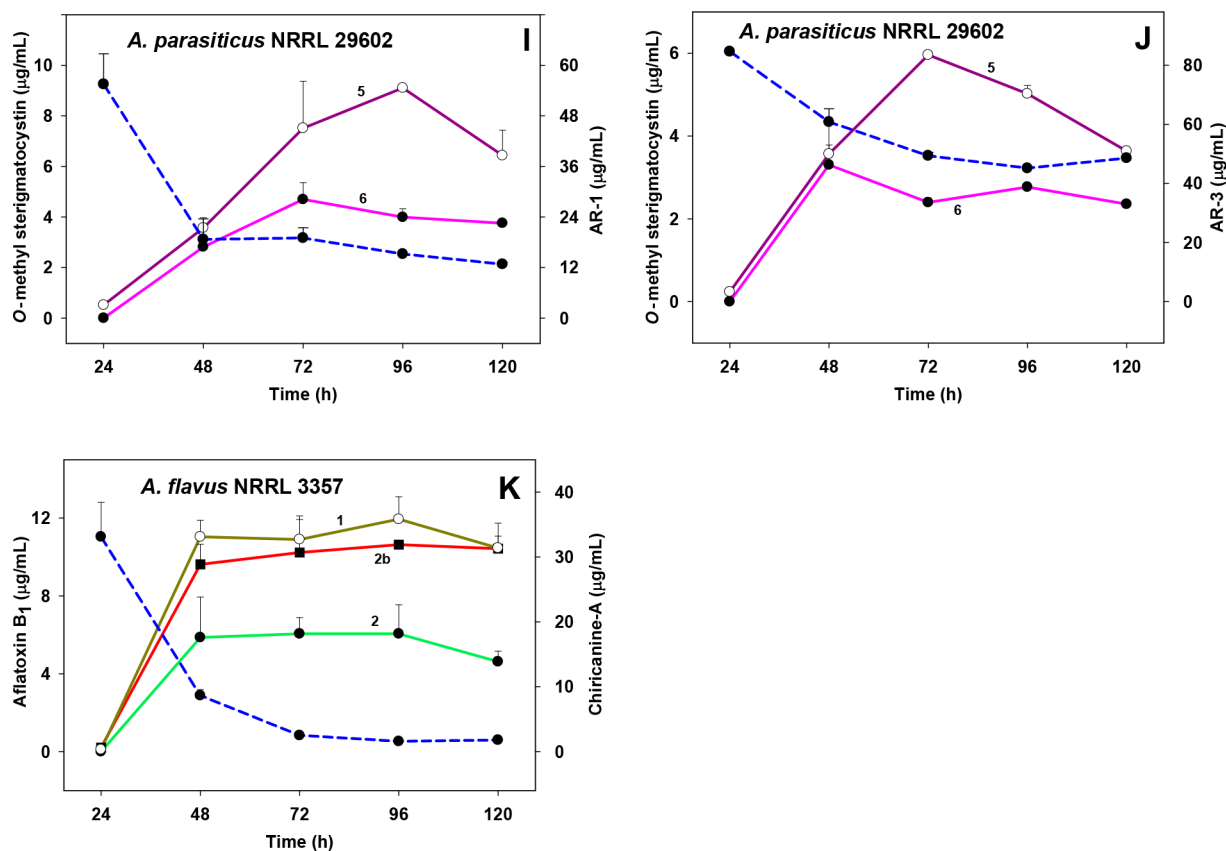


Figure 2. Dynamics of aflatoxin inhibition by peanut stilbenoids. Line 1 in all graphs represents aflatoxin B₁ formation in the control (without stilbenoids) samples. Line 2 in all graphs shows aflatoxin B₁ formation in the experimental samples. (A) Lines 2a and 2b show aflatoxin B₁, 7, formation after the treatment with 0.15 mM and 0.075 mM arachidin-1, respectively. (G,H) Line 3 represents aflatoxin G₁, 9, formation in the control samples (without the arachidins), and line 4 shows aflatoxin G₁ formation in the experimental samples. (I,J) Line 5 shows the dynamics of accumulation of *O*-methyl sterigmatocystin, 11, without the arachidins, and line 6 shows the concentrations of 11 in the experimental samples. (K) Line 2b shows aflatoxin B₁ formation after treatment with 0.075 mM chiricanine A, 6. In all graphs, dashed blue lines show the dynamics of the stilbenoid-concentration decline. AR-1 means arachidin-1, and AR-3 means arachidin-3.

the experiment. The inoculated test tubes were incubated at 30 °C in the dark. Using wide bore tips, 12 μ L of each sample was collected at 2 h intervals starting after 6 h of incubation. The samples were observed under the microscope using a hemocytometer for spore counting, and hyphal-length quantitation was performed using the Live Measurement module of LAS software, ver. 4.3.0, in a Leica DM 2500 microscope (Vashaw Scientific Inc., Roswell, GA). Each sample was evaluated on 5–10 \times 4 nL fields, and the number of fields was increased over time as the samples became more heterogeneous.

Data Analysis. Data were analyzed by ANOVA procedures using SAS 2000, ver. 7 (SAS Institute, Inc., Cary, NC). Multiple comparisons of the various means were carried out by the least significant difference (LSD) test at $p = 0.05$. Comparisons of the means of two groups of data were performed using the t test; the Mann–Whitney Rank Sum Test was applied when the normality test failed ($p < 0.05$).

RESULTS AND DISCUSSION

It is reasonable to suggest that there is appreciable inhibition of toxin formation in *Aspergillus* species by peanut prenylated stilbenoids on the basis of the published findings.^{9,10,14} To study the influence of peanut phytoalexins on aflatoxin accumulation, we used three prenylated stilbenoids, arachidin-1, 3; arachidin-3, 5; and chiricanine A, 6 (Figure 1), in our feeding experiments. Because an insufficient quantity was available, another important stilbenoid, arachidin-2, 4, was used only in the morphological study. The rationale for the choice was based on the fact that arachidin-1, 3, and arachidin-3, 5, are

the major prenylated stilbenoids that are formed in fungus-challenged peanut seeds. In addition, these stilbenoids, as well as chiricanine A, 6, and arachidin-2, 4, demonstrated appreciable biological activity compared with the other stilbenoids tested.¹⁴ The soil fungi used in this research included five strains of *Aspergillus* that differed from each other by their metabolite profiles and toxigenic potential: *A. flavus* NRRL 3357, a moderate producer of aflatoxins B₁ (7) and B₂ (8); *A. flavus* NRRL 29487, a high producer of aflatoxins B₁ and B₂; *A. nomius* NRRL 13137, a high producer of aflatoxins B₁, B₂, G₁ (9), and G₂ (10); *A. parasiticus* NRRL 29580, a very high producer of aflatoxins B₁, B₂, G₁, and G₂; and *A. parasiticus* NRRL 29602, a producer of *O*-methyl sterigmatocystin (11). The data on the fungal toxigenic potentials (not listed here) were provided by Dr. B. Horn of the National Peanut Research Laboratory, ARS, USDA (Dawson, GA). Preliminary feeding experiments demonstrated that arachidins 3 and 5 as well as chiricanine A, 6, were substantially more potent inhibitors of aflatoxin formation than caffeic acid, 1,¹⁰ and resveratrol, 2.⁹ Therefore, about a 30–40-fold lower initial concentration of each stilbenoid was suggested on the basis of their activity. The final concentration of 0.3 mM was chosen on the basis of the highest full solubility of 3, 5, and 6 in the PDA medium. This concentration is equivalent to 88.8 μ g/mL of arachidin-3 and is about 50-fold lower than the concentrations of the fungus-induced stilbenoids detected in alive, wounded peanut seeds.¹³

The current research demonstrated that the selected stilbenoids were capable of inhibiting the formation of aflatoxins 7–10 and the precursor *O*-methyl sterigmatocystin, 11, in all of the *Aspergillus* species tested. The most dramatic results were obtained when *A. flavus* NRRL 3357 was grown in the presence of arachidin-3, 5 (Figure 2B). At all sampling times, aflatoxin accumulation was almost completely (>98%) suppressed (line 2, Figure 2B), whereas the control without arachidin-3 (line 1) demonstrated substantial accumulation of aflatoxin B₁, 7, from 48 to 120 h of incubation. Aflatoxin B₂, 8, was also detected, but the insignificant productions of aflatoxin B₂ by the *A. flavus* strains and of aflatoxins B₂ and G₂, 10, by the *A. parasiticus* strains used in this research were not accounted for in the simplicity of the presentation. A gradual decrease of arachidin-3, 5, from its initial concentration of 88.8 to 35.0 µg/mL occurred over the course of the experiment within 120 h (Figure 2B). *A. flavus* NRRL 3357 growth inhibition in the presence of arachidin-3 was obvious from 24 to 120 h of incubation compared with the growth of the control (Figure 3B). Both mycelial and conidial growth was compromised in the experimental wells. However, there is no sufficient evidence to conclude that aflatoxin inhibition occurred because of the compromised fungal growth and development. The morphological observations and measurements were performed after 6 h of incubation of *A. flavus* NRRL 3357 conidia with arachidin-3 at 30 °C. At 6 h and throughout the course of the experiment, in the presence of arachidin-3, conidia formed clustered masses (Figures 4C), unlike the control samples without arachidin-3 (Figure 4A,B). After allowing the spores to germinate for 13 and 15 h in the presence of arachidin-3 (Figure 4C,D), a higher degree of the mycelial branching was demonstrated compared with that in the control, whose branching was not so obvious and frequent (Figure 4A,B). The white arrows show clusters of germinating and branching spores (Figure 4C). The black arrows show hyphae, the branching filaments that form the fungal mycelium. Figure 5A shows that there was a significant difference in the degree of spore germination after 8 and 10 h of incubation. A significant difference in hyphal length was also observed from 8 h to the end of the experiment (Figure 5B).

A similar experiment with the same fungal strain, *A. flavus* NRRL 3357, but arachidin-1, 3, as an inhibitor (Figure 2A) at 0.3 mM concentration, demonstrated that aflatoxin accumulation was detected from 24 h to the end of the experiment (line 2), although it was 4–5-fold lower compared with that in the control (line 1). Lower arachidin-1, 3, concentrations of 0.15 mM (line 2a) (Figure 2A) and 0.075 mM (line 2b) also inhibited aflatoxin accumulation in a concentration-dependent pattern. The concentration of arachidin-1 rapidly dropped from the initial 93.6 to 25.4 µg/mL at 24 h and then gradually declined to 8.3 µg/mL at 120 h. The difference in growth of the treated and control set of samples was not obvious (Figure 3A).

A stronger aflatoxin producer, *A. flavus* NRRL 29487, grown on the medium spiked with arachidin-3, was also affected by the stilbenoid, but its aflatoxin-inhibition pattern (Figure 2D) differed from that of *A. flavus* NRRL 3357. The peak of aflatoxin production in the control as well as in the treated sample was observed at 72 h, and then production gradually declined. Arachidin-3 concentration gradually dropped from the original concentration to zero within 120 h. When treated with arachidin-1, the same strain showed a different toxin-formation pattern, as seen from Figure 2C. Suppression of aflatoxin (2- to 3-fold) was observed from 72 to 120 h. At the same time, the arachidin-1 concentration rapidly dropped from

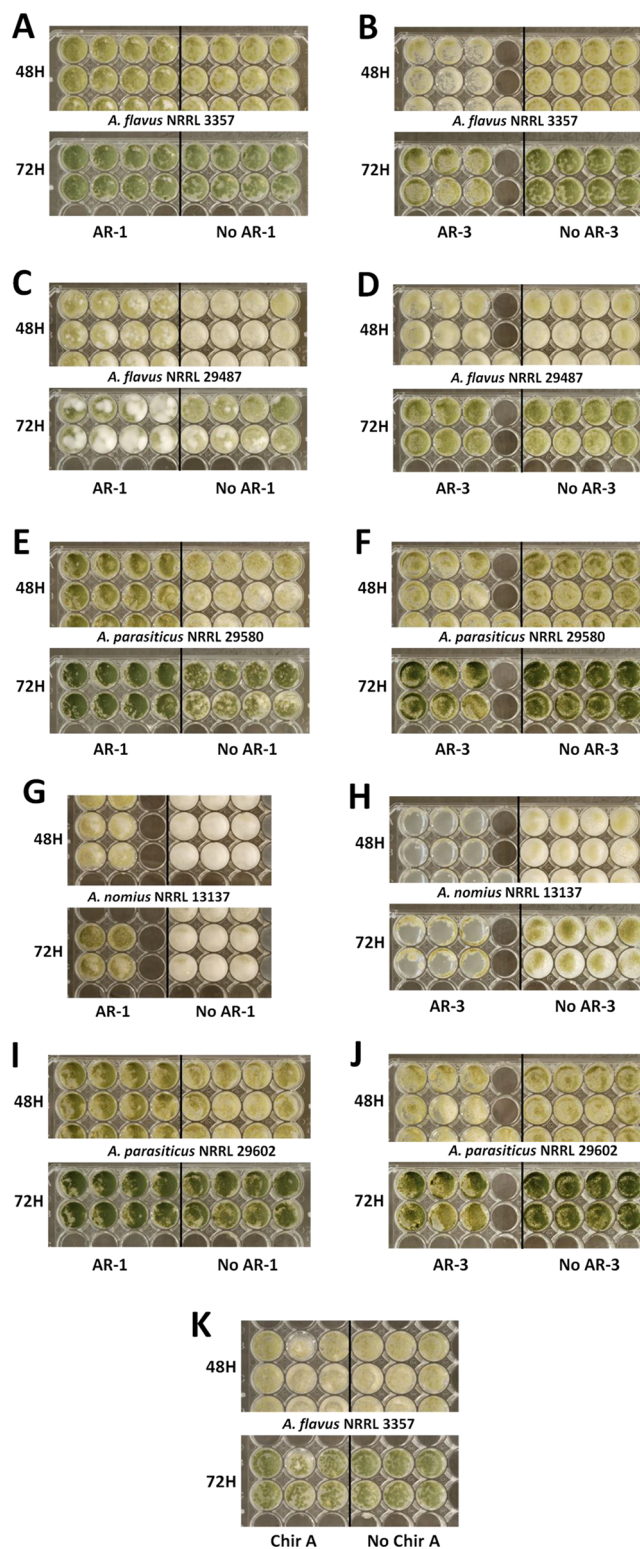


Figure 3. Experimental and control wells with the fungal species *A. flavus*, *A. parasiticus*, and *A. nomius* grown for 48 and 72 h on PDA medium with arachidin-1, 3 (B,D,F,H,J); arachidin-3, 5 (A,C,E,G,I); and with chircanine A, 6 (K). AR-1 means arachidin-1, AR-3 means arachidin-3, and Chir A means chircanine A.

the original concentration to almost zero at 48 h (Figure 2C). There was no visual difference in growth and development patterns between the control and the experiment in the case of arachidin-3 (Figure 3D), and only a slightly favorable formation

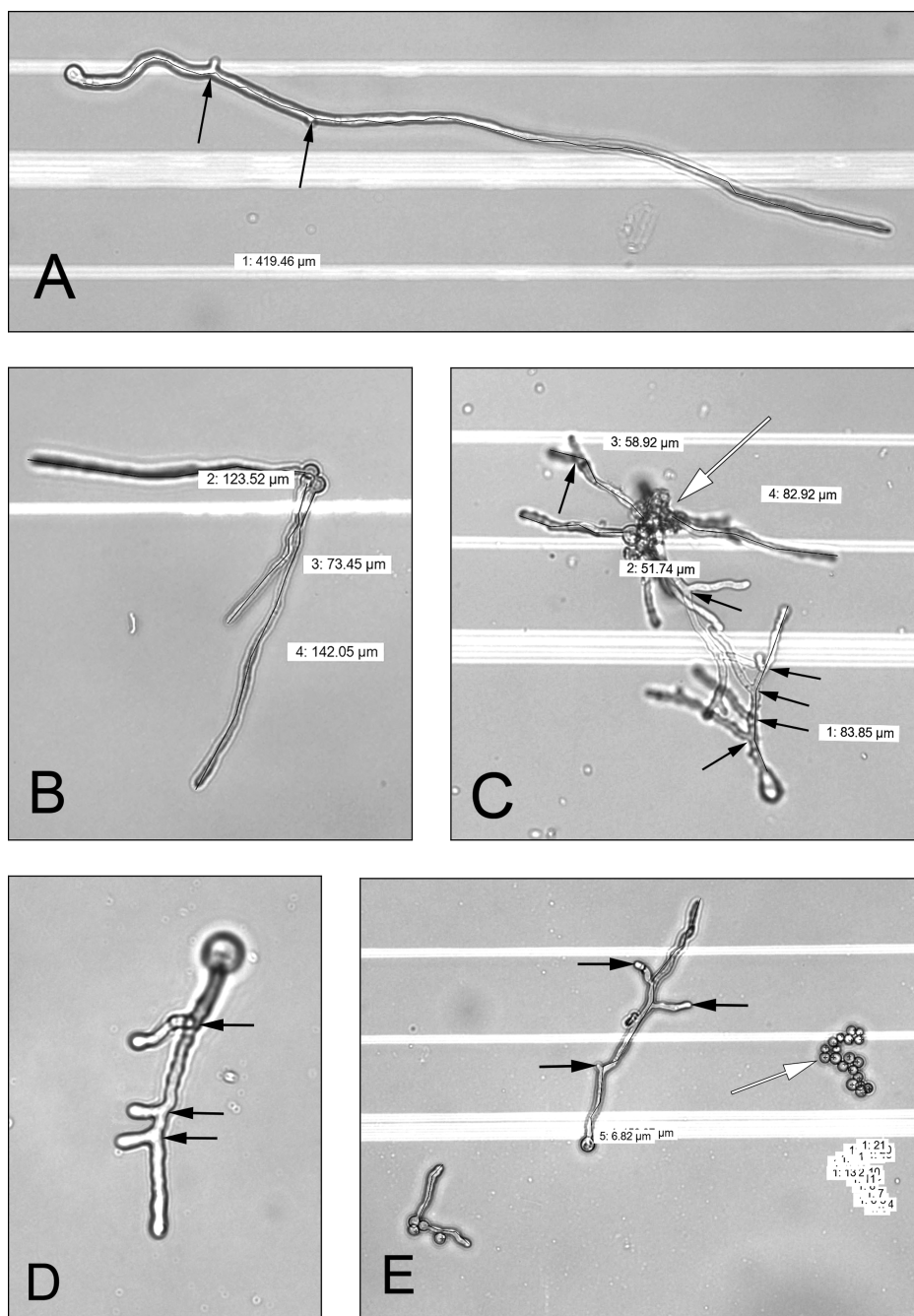


Figure 4. Microscopic view of the *A. flavus* NRRL 3357 fungal structures developed (A) without the stilbenoids (control) after 14 h of incubation; (B) without the stilbenoids (control) after 18.5 h of incubation; (C) with arachidin-3, **5**, after 13 h of incubation; (D) with arachidin-3 after 15 h of incubation; and (E) with chiricanine A, **6**, after 16 h of incubation. The black arrows show hyphae, the branching filaments that form the fungal mycelium; the white arrow shows a cluster of nonviable spores.

of conidia and mycelia was observed in the wells with arachidin-1 (Figure 3C).

The growth and development of the highest aflatoxin producer tested, *A. parasiticus* NRRL 29580, was not affected by arachidin-3, **5** (Figure 3F). Also, there was no significant difference in aflatoxin inhibition between the control samples and samples with arachidin-3 (Figure 2F). However, aflatoxin suppression by arachidin-1, **3**, was significant (about 3-fold) throughout the duration of the experiment with a concomitant decrease in the arachidin-1 concentration (Figure 2E). Although there was no distinct difference in appearance between the control and experimental wells with the fungus

grown from 24 to 120 h in the absence or presence of arachidin-3, **5**, respectively, (Figure 3F), there was a distinct difference when the fungus was treated with arachidin-1, **3**. Surprisingly, more active fungal growth was observed in the wells treated with **3** compared with that in the control wells (Figure 3E). In addition, arachidin-1 seemed to promote conidial (green colored) formation rather than mycelial (white colored) formation from 48 to 120 h.

Arachidin-1, **3**, and arachidin-3, **5**, displayed even more dramatic influences on fungal formation in *A. nomius* NRRL 13137 from 24 to 120 h (Figure 3G,H). Although the effect of arachidin-3 on fungal development was predictable, the effect of

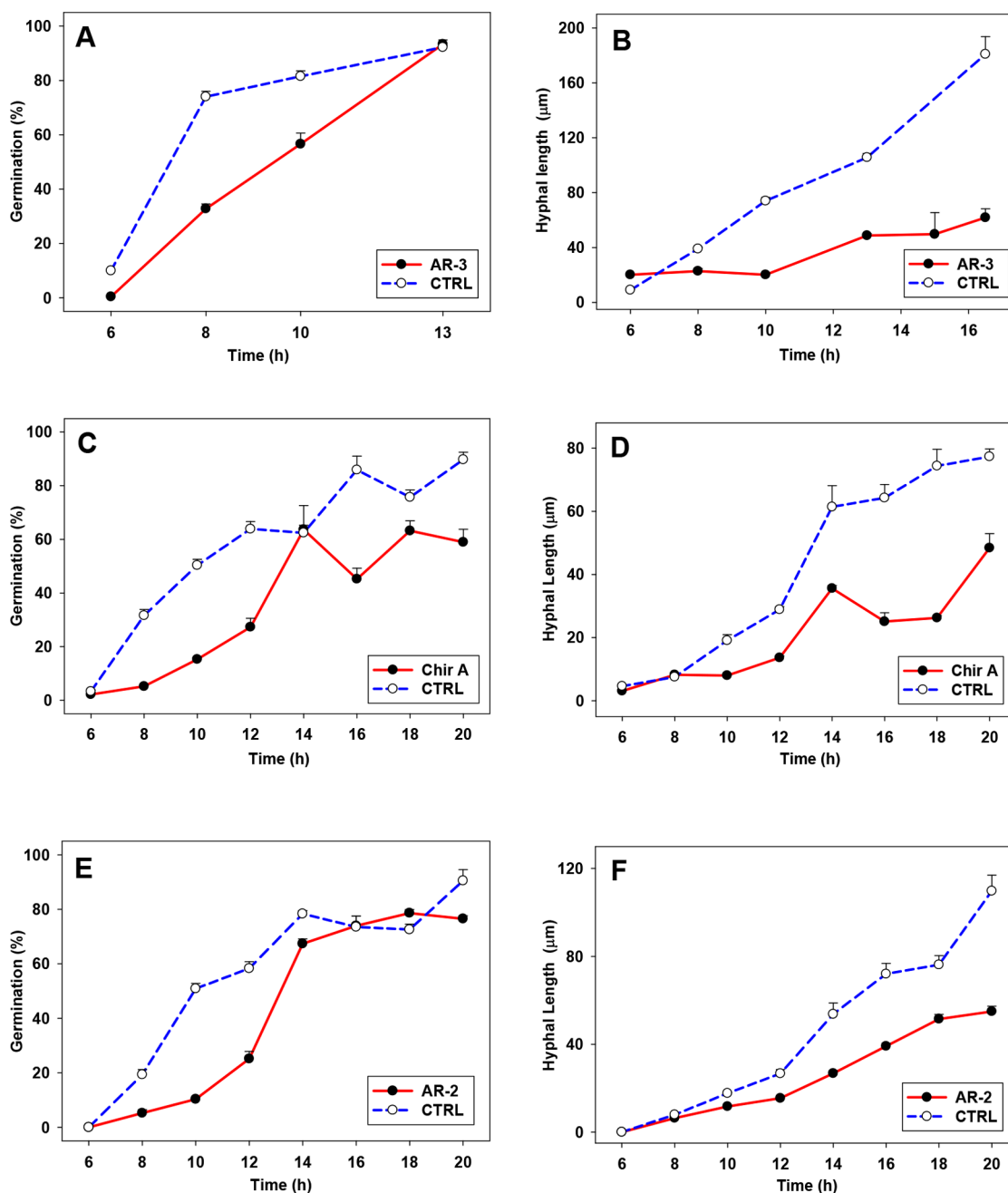


Figure 5. Morphological changes in *A. flavus* NRRL 3357 exposed to (A,B) arachidin-3, 5; (C,D) chiricanine A, 6; and (E,F) arachidin-2, 4. AR-3 means arachidin-3, Chir A means chiricanine A, and AR-2 means arachidin-2. CTRL means control.

arachidin-1 was unexpected. Arachidin-3 almost completely suppressed fungal formation at 48 h and substantially did so at later times (Figure 3H). In contrast, arachidin-1 favored conidial formation, whereas the control showed basically mycelial growth from 24 to 72 h of incubation (Figure 3G). At the same time, the formation of all aflatoxins, B₁, B₂, G₁, and G₂, was not significantly inhibited by arachidin-1 (Figure 2G). On the other hand, arachidin-3 demonstrated appreciable activity and substantially inhibited the formation of aflatoxin B₁ at 48 and 72 h (Figure 2H, lines 1 and 2) and significantly suppressed aflatoxin G₁ formation (lines 3 and 4). Degradation of arachidin-3 occurred at a slower rate compared with that of arachidin-1. The initial concentrations of 3, 5, and 6 in all the control samples remained unchanged within the statistical error throughout the course of the experiments. Therefore, the

significant decrease of the stilbenoid concentrations in the presence of the fungi tested allows us to suggest that the most likely fate of 3, 5, and 6 is degradation by fungal enzymes.¹⁹

A natural *O*-methyl sterigmatocystin, 11, producer, *A. parasiticus* NRRL 29602, was also tested against the arachidins. It was reasonable to test the potential inhibition of 11 by the same stilbenoids as this mycotoxin is a precursor in the aflatoxin biosynthetic pathway. Both arachidin-1 and arachidin-3 suppressed the formation of 11 most actively between 72 and 120 h of incubation (Figure 2I,J). At the same time, there were no noticeable differences between fungal growth and appearance at all times (Figure 3I,J) with the exception of slightly increased spore formation in the presence of 3 at 48 h.

The limited quantity of chiricanine A, 6, allowed us to test its action only on one strain, *A. flavus* NRRL 3357. In the presence

of this stilbenoid, significant aflatoxin reduction was observed at all times starting at 48 h at 0.3 mM (Figure 2K, line 2). Inhibition at 0.075 mM was statistically insignificant but noticeable (Figure 2K, line 2b). The concentration of chircanine A gradually reached a level of almost zero at 72 h from its original value of 84.0 $\mu\text{g}/\text{mL}$. Visually, the control and experimental wells were indistinguishable at all times of fungal growth (Figure 3K). The morphological differences between the control and experimental samples of the fungus were obvious and significantly different in terms of germination rates and hyphal lengths (Figure 5C,D) starting at 8 h of incubation for the germination rates and 10 h for the hyphal lengths. At 6 h, large numbers of spores clustered together (Figure 4E, white arrow) were observed in the presence of chircanine A, whereas the control samples had single, loose spores. The fungal spores in the samples with chircanine A stayed as clusters even after they started to form hyphae. In the samples with chircanine A at 14–16 h of incubation, some branching was observed in the growing mycelia (Figure 4E). However, the branching was not as evident as it was in the experiments with arachidin-3. The control spores began to form clusters at 12 h of incubation, but the spore growth and elongation occurred without branching throughout the remainder of the observation (Figure 4B).

A pure sample of arachidin-2 was also tested using the same experimental setup. *A. flavus* NRRL 3357 treated with arachidin-2 demonstrated morphological changes (Figure 5E,F) similar to those that were observed with arachidin-3 and chircanine A (Figure 5A–D). Conclusions on morphological changes were made on a statistically significant number of microscopic observations. The total number of fungal structures measured was 5496 in 329 fields of observation.

The present research demonstrated that the most abundant and highly biologically active peanut phytoalexins, arachidin-1, 3; arachidin-3, 5; and chircanine A, 6, had appreciable capacity to inhibit aflatoxin formation in the important toxigenic *Aspergillus* species tested, *A. flavus*, *A. parasiticus*, and *A. nomius*. Aflatoxin inhibition was not necessarily accompanied by visible changes in fungal growth and development. At present, a mechanism of aflatoxin inhibition by peanut stilbenoids in *Aspergillus* cannot be suggested. Additional experiments with a larger number of peanut stilbenoids and fungal species may help to elucidate the stilbenoid structure–aflatoxin inhibition relationship. New in vivo research is also needed to ensure that aflatoxin inhibition occurs in the course of the peanut–fungus interaction. Such research is planned. Knowledge on the mechanism of the plant–fungus interaction could lead to new strategies for preventing plant invasion by the fungi that produce aflatoxins.

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Notes

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