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Research Article

Endophytic fungus

(Peak 536 nm) AuNPs

Biogenesis of nanoparticles with inhibitory effects on aflatoxin B_1 production by *Aspergillus flavus* $\stackrel{\star}{\approx}$

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Silver and gold nanoparticles

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flavus growth inhibit at 10000-125 µg/ml

0-1.42 g

G R A P H I C A L A B S T R A C T

UV-Vis



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ABSTRACT

Background: Fungal nanofactories have been utilized to synthesize silver and gold nanoparticles. This study was designed to mycosynthesize and characterize silver and gold nanoparticles (AgNPs and AuNPs) and to study their effect on aflatoxin B_1 production by *Aspergillus flavus*.

Results: Silver and gold nanoparticles were synthesized by endophytic *Aspergillus versicolor* and then analyzed by UV–vis spectroscopy. The results revealed surface plasmon resonance peaks at 432 and 536 nm for Ag and Au nanoparticles, respectively. The obtained transmission electron microscopy results revealed the fashioning of spherical AgNPs and spherical and hexagonal AuNPs with a mean particle magnitude of 5–37 and 37–62 nm, respectively. X-ray diffraction showed the typical face-centered cubic structure of the mycosynthesized Ag and Au nanoparticles. An *in vitro* investigation showed that AgNPs, AuNPs, and their mixture at different concentrations (10000, 5000, 3000, 1000, 750, 500, 250, and 125 μ g/mL) could inhibit or reduce the outgrowth and production of aflatoxin B₁ (AFB₁) by *A. flavus*. The concentration that showed no AFB₁ production was less than those for the inhibition of fungal growth. AgNPs, AuNPs, and their mixture also exhibited promising antiradical scavenging activity. *Conclusions:* The use of fungi in the metallic nanoparticle's fabrication and the utilization of mycosynthesized is promising as a substitute of chemicals to control antiaflatoxigenic fungi.

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Abbreviations: AFB₁, aflatoxin B₁; AgNPs, silver nanoparticles; AuNPs, gold nanoparticles; BHT, butylated hydroxytoluene; DPPH, 1,1-Diphenyl-2-picryl-hydrazyl; FTIR, Fourier Transform Infrared Spectroscopy; HPLC, high performance liquid chromatography; ITS, Internal Transcribed Spacer; PDA, potato dextrose agar; TEM, Transmission Electron Microscopy; TLC, thin-layer chromatography; XRD, X-ray diffraction; YES, yeast extract sucrose.

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1. Introduction

Nanoparticles, which are known to possess a dimension of 100 nm or less in size, [1] have gained growing awareness as inorganic antimicrobial agents in nonfood applications [2] because of their distinctive physical and chemical attributes, which vary considerably from their traditional coordinates [3]. Physical and chemical methods, which converge on utilizing a vast amount of chemicals and elevated temperature through radiation, are used to achieve the synthesis of metal nanoparticles. Modern investigations have shown the use of easy, neat, eco-friendly, inexpensive, and safe techniques for bio-fabrication of nanoparticles [4]. This involves the use of various biological resources that include bacterial, fungal, algal, plant, and animal metabolites to synthesize different metal nanoparticles [5,6,7,8,9]. Fungal nanofactories have been utilized to synthesize silver, platinum, silica, zirconium, iron titanium, and gold nanoparticles [1,10,11,12]. Green chemistry is a simple eco-friendly technique for the fabrication of metal nanoparticles in which fungi are employed for nanoparticle biosynthesis [13].

Aflatoxins are a small group of closely related heterocyclic secondary metabolites [14]. It was found as difuranocoumarin derivatives biosynthesized through a polyketide pathway [15]. Of the 20 well-characterized diverse sorts of aflatoxins, aflatoxins B_1 , B_2 , G_1 , G_2 , M_1 , and M_2 are considered to be the most important ones [16], and along with aflatoxin B_1 (AFB₁), they are the most common and toxic toxins that cause a serious risk to human and animal health [17]. Aflatoxin B_1 is produced mainly by aflatoxigenic species of filamentous fungi, common saprophytes, and opportunistic pathogens *Aspergillus* (*A. flavus, A. parasiticus,* and *A. nomius*) [18]. Aflatoxin B_1 is the most common globally and reported in about 75 % of food and feeds contamination with aflatoxins [19,20].

Aflatoxin B_1 , whose mutagenic, hepatic-carcinogenic, teratogenic, and immunosuppressive activity to a broad spectrum of living organisms has been widely studied [21], while the International Agency of Research on Cancer was classified AFB₁ as Group 1A carcinogen [22]. The European Commission has established 2 ng/g as the maximum level for AFB₁, 4 ng/g for combined aflatoxins in foodstuffs, and 0.05 ng/mL for AFM₁ in liquid milk [23]. Inhibiting the outgrowth of aflatoxigenic fungi is first necessary to prevent aflatoxins production in agricultural merchandises [24].

Physical, chemical, and biological methods to reduce aflatoxins can be employed for detoxification or removing from contaminated food and nutrients [25]. Larger surface area of nanoparticles, because of their size, interact with microorganisms rather than the larger shape of particles [26]. Thus, it may be functional to utilize nanoparticle-based modes as a substitutional remedy versus fungi [27]. The influence of silver nanoparticles on the development and AFB₁ biosynthesis of *Aspergillus parasiticus* was studied and determined in an earlier report [28]. This study aimed to evaluate antiaflatoxigenic activities using different concentrations of mycosynthesized AgNPs and AuNPs by *A. versicolor* individually/mixture on outgrowth and AFB₁ accumulation by *A. flavus*.

2. Materials and methods

2.1. Isolation of fungal isolates

Aspergillus versicolor was utilized for the biosynthesis of nanoparticles as it exhibited a great potency to biosynthesize both investigated nanoparticles among the 10 tested fungal isolates. *A. versicolor* was isolated as an endophyte from sand lily (*Pancratium maritimum* L.) leaves [29], as previously described by Kumar et al. [30], while aflatoxigenic *A. flavus* was isolated from peanut (*Arachis hypogaea* L.) seeds according to earlier report [31].

2.2. Molecular identification of isolated fungi

Czapek Dox broth medium was used to culture the pure fungal isolates for 5 d at 28°C. Total DNA was extracted from each isolate using a specific kit "Norgen Plant/Fungi DNA Isolation Kit (Sigma, Thorold, Canada)" [32]. To amplify the ITS region of the selected strains, universal primers ITS-1 (5'-TCC GTA GGT GAA CCT GCG G-3') and ITS-4 (5'-TCC TCC GCT TAT TGA TAT GC-3') were employed, as previously described by Mohamed et al. [33]. PCR amplification was performed following the same protocol as earlier described by Hassan et al. [32].

2.3. Preparation of mycelial-free culture filtrate

The fungal biomass was cultivated aerobically in PD broth and incubated for 5 d at 28°C in an orbital shaker with constant shaking at 150 rpm/min. Subsequently, the fungal biomass obtained was then filtered by using a Whatman no. 1 filter paper, and the resultant biomass was washed numerous times with double distilled water. Then, 10 g of fungal mycelia were soaked into 200 mL sterile double distilled water in a 500 mL Erlenmeyer flask and was further incubated at 28°C for 48 h with gentle shaking at 120 rpm. The filtrate was then used for nanoparticle synthesis.

2.4. Mycosynthesis of silver and gold nanoparticles

Stock solutions of both silver nitrate and gold chloride (hydrogen tetrachloroaurate (III) hydrate (HAuCl₄·3H₂O) were prepared in deionized water, and the solutions were then individually added to the mycelial-free filtrate, resulting in a final concentration of 1 mM. The flask was then kept in dark condition at 28°C for 72 h. Gold chloride solution and fungal filtrate served as negative controls. The resulted nanoparticles were collected by centrifugation and freeze-dried by lyophilization, and a certain weight was disseminated by using ultra-sonication in sterile double distilled water for use as stock solutions.

2.5. Characterization of AuNPs and AgNPs

2.5.1. UV-visible (UV-vis) spectroscopy

The formation of AgNPs and AuNPs was determined by visual observation of color change that indicates the reduction reaction, and AgNPs and AuNPs were established by UV–vis spectroscopy (Jasco V-530) for the occurrence of distinguishable surface plasmon resonance bands of AgNPs and AuNPs.

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2.5.2. Transmission electron microscopy (TEM)

The morphology of newly synthesized AgNPs and AuNPs were analyzed by a transmission electron microscope (JEOL/JEM-2100, HRTEM, Tokyo, Japan). The examined TEM specimen was concocted by drop-casting and dissemination of AgNPs and AuNPs on copper grids, and the samples were then dried at ambient temperature.

2.5.3. Fourier transform infrared spectroscopy (FTIR)

FTIR (6100, Perkin-Elmer, Germany) was utilized to investigate the functional groups on the surface of AgNPs and AuNPs at 4 cm^{-1} resolution and 400–4000 cm⁻¹ wavelength range.

2.5.4. X-ray diffraction (XRD) analysis

The freeze-dried powders were subjected for XRD analysis for the presence of elemental AgNPs and AuNPs, and the experiments were performed on a X-ray diffractometer (Panalytical X'PERT PRO) with Cu k α - radiation ($\lambda k \alpha = 1$. 1.540562 Å) run at 30 mA and 40 kV and scanning diffraction style in the 2 \emptyset of 5-90° range.

2.6. Impact of AgNPs and AuNPs and their mixture on the growth and aflatoxin B_1 accumulation of aflatoxigenic A. flavus strain

Dilutions of the stock solutions of AgNPs and AuNPs were prepared using sterilized deionized H_2O to attain the desired concentrations (10000, 5000, 3000, 1000, 750, 500, 250, and 125 µg/mL) used for antimycotic and antiaflatoxigenic investigation. The dry weight of the fungal biomass in each NPs treatment was then recorded and compared with the control. All experiments were repeated three times [27]. Aflatoxin B₁ was purified and detected using the thin-layer chromatographic (TLC) technique [34,35].

2.6.1. Determination of AFB₁ by high performance liquid chromatography (HPLC)

The concentration of AFB1 was determined by HPLC [36].

2.7. Radical scavenging activity by 1,1-diphenyl-2-picryl-hydrazyl (DPPH) assay

The antioxidant properties of AgNPs and AuNPs were measured as scavenging activity or as a hydrogen donating form based on the procedure reported by Brand-Williams et al. [37] and Adebayo et al. [38] with minor modifications (Fig. S1).

2.8. Data analysis

Statistical analysis was performed using SPSS (version 16), and the differences were analyzed by one-way ANOVA. The values were expressed as mean \pm SE, and the significance level was considered as 0.05.

3. Results

3.1. Fungal isolates and phylogenetic analysis

According to the phenotypic and phylogenetic characteristics, *A. versicolor* was identified as an endophyte from the leaves of *Pancratium maritimum*, while *A. flavus* was identified as an isolate from the seeds of *Arachis hypogaea*. Both the isolates were identified by performing ITS region sequencing, and the sequences obtained for the ITS regions were further subjected to a BLAST search at the NCBI database. The two isolates were identified as *Aspergillus versicolor* TU-63 (GenBank accession no. OL411612) and *A. flavus* TU-61(GenBank accession no. OL411610) (Fig. 1).

Nucleotide comparisons of the ITS regions among the *Aspergillus* and other similar strains revealed that the *A. versicolor* TU-63 strain exhibited 100% and 98% similarity with the strains *A. versicolor* MK722085 and *A. versicolor* MT609910 from the GenBank, respectively. On the other hand, 99 and 98% identity was observed between *Aspergillus flavus* TU-61 and *A. flavus* MZ052072 and *A. flavus* MZ018647, respectively. *Aspergillus flavus* TU-61 also showed approximately 97% and 96% similarity with *A. flavus* MW193211 and *A. flavus* MW193310, respectively.

3.2. Characterization of mycosynthesized AuNPs and AgNPs

The bioreduction of $AgNO_3$ was noticed by visual examination of the culture filtrate after 2 d of incubation with a color change to deep brown, while the color change in the solution containing gold chloride and mycelia filtrate to purple indicated biogenesis of AuNPs (Fig. 2A).

UV–Vis wave analysis of AgNPs over the range of 200–600 nm revealed that the surface plasmon resonance band originated at 432 nm. Absorption measurements over the range of 300–700 nm for AuNPs produced a peak at 536 nm, coinciding to the plasmon absorbance of AuNPs (Fig. 2B).

The size and morphology examined by TEM showed that the formed AgNPs were monodispersed with a relatively regular spherical structures and an average particle size of 5–37 nm. TEM analysis of the biosynthesized AuNPs revealed that they were polydispersive showing spherical and hexagonal structures with a particle size range of 37–62 nm (Fig. 2C).

The crystalline quality of AgNPs was determined by the XRD analysis that exposed the typical face-centered cubic (fcc) structure of the metal silver. The XRD spectra revealed four major Bragg diffraction peaks at 2θ values of 38.121° , 44.308° , 64.458° , and



Fig. 1. Neighbor-joining phylogenetic tree of the ITS sequences of the isolated fungal strains with the sequences of closely related strains: (A) the phylogenetic tree of *A. versicolor* and (B) the phylogenetic tree of *A. flavus*.



Fig. 2. (A) Color change of mycosynthesized AgNPs and AuNPs (a) silver nitrate, (b) AgNPs, (c) *A. versicolor* mycelial filtrate, (d) AuNPs, and (e) gold chloride; (B) UV–Vis absorption spectrum of biosynthesized (a) AgNPs and (b) AuNPs; (C) TEM images of (a) AgNPs and (b) AuNPs.

77.415° that matched to the (111), (200), (220), and (311) planes of fcc Ag, respectively. The diffraction peaks were harmonious with the (JCPDS card No. 04–0783) standard database files, which confirm that the mycosynthesized AgNPs had a crystalline nature (Fig. 3A). XRD analysis of AuNPs exhibited Bragg diffraction peaks and planes of fcc as for AgNPs by employing (space group Fm3 m, JCPDS File NO. 89–3697) standard database files (Fig. 3B).

The FTIR spectrum of AgNPs showed a number of different bands including strong bands that appeared in the spectrum such as the band centered at 2965.86 and 2903.53 cm⁻¹ for aliphatic C—H stretching and the strong band at 1547.36 cm⁻¹ for carbonyl group C=O stretching. The band centered at 1292.03 is attributed for the C–O stretching group, while the strong band centered at 1053.73 is attributed to the S=O stretching functional group in the sample (Fig. 4A).

The FTIR spectrum of AuNPs showed a number of different bands including strong bands that appeared in the spectrum such as the band centered at 2972.08 and 2900.92 cm⁻¹ for aliphatic C–H stretching. The strong band centered at 1053.73 was attribu-

ted to the S=O stretching functional group in the sample. A small peak was observed at 3658.98 cm⁻¹, which could be because of the presence of a -O-H hydroxyl group. The band centered at 1596.09 cm⁻¹ was attributed to carbonyl group C=O stretching (Fig. 4B). These bands indicated that those functional groups were able to bind ions of the silver and gold metals and can biosynthesize and capping AgNPs and AuNPs to block clustering and settle the medium.

3.3. Impacts of AgNPs and AuNPs and Ag-AuNPs treatment on the growth of Aspergillus flavus and its capability to produce aflatoxin B_1

The impacts of AgNPs, AuNPs, and their mixture to inhibit the outgrowth and aflatoxins B₁ biosynthesis by *A. flavus* were observed (Table 1 and Fig. S1-S20). The present study revealed that the impacts of AgNPs, AuNPs, and their mixture in inhibiting the growth and aflatoxin B₁ production capability of *A. flavus*. The growth inhibition was determined by dry weight, where it ranged between 0–1.55 g at 10000–125 μ g/mL AgNPs, 0–1.42 g at 10000–



Fig. 3. XRD micrograph of mycosynthesized (A) AgNPs and (B) AuNPs.

125 µg/mL AuNPs, and 0–1.62 g at 10000–125 µg/mL Ag-AuNPs in comparison to the control (1.73 g). Our study also revealed that the inhibition of growth and aflatoxin B₁ production capability of A. flavus was directly linked to the concentrations of the nanoparticles tested. The quantities of accumulated aflatoxin B₁ decreased when the used nanoparticles concentration increased. The growth of A. flavus was completely inhibited at 10000 µg/mL concentration, while no aflatoxin B₁ production occurred at 10,000 and 5000 µg/mL of AgNPs, AuNPs, and their mixture, respectively. The concentration that showed no aflatoxin B1 production was less than those for the reduction of fungal growth. The highest concentration for the inhibition of aflatoxin B_1 production (99.55, 99.999, and 99.59%) for AgNPs, AuNPs, and Ag-AuNPs, respectively, was 3000 μ g/ml, whereas fungal growth was decreased significantly at these concentrations. The lowest concentration for the inhibition of aflatoxin B₁ formation (60.50, 67.33, and 23.24%) for AgNPs, AuNPs, and Ag-AuNPs, respectively, was 125 μ g/mL, whereas fungal growth was reduced significantly at these concentrations (Fig. 5A, B).

3.4. Antioxidant activity of AgNPs, AuNPs, and Ag-AuNPs

The mycosynthesized AgNPs exhibited satisfactory antiradical activities as compared to BHT with inhibitions of 27.02–79.58%, AuNPs displayed inhibitions of 40.19–80.22%, while Ag-AuNPs displayed inhibitions of 33.59–76.85% at the tested concentrations (12.5–100 μ g/mL) (Table 2).

4. Discussion

In this study, we attempted to synthesize AuNPs and AgNPs by *Aspergillus versicolor* isolated as an endophyte from the leaves of *Pancratium maritimum*, while aflatoxigenic *A. flavus* was isolated from the seeds of *Arachis hypogaea*. Both fungi were identified on the basis of the phenotypic and phylogenetic characteristics. The ITS loci are the most credible regions for identifying strains at the species level. Not surprisingly, ITS region sequencing is reported as a powerful alternative method for accurate molecular identification of *Aspergillus* strains to the species level [32].

AgNPs biosynthesis was indicated by the change in color of the culture filtrate from faint xanthous to deep brown color, while pur-



Fig. 4. FTIR analysis of biosynthesized (A) AgNPs and (B) AuNPs.

Table 1	
Effects of different concentrations of biosynthesized AgNPs, AuNPs, and their mixture on the growth of A. flavus and accumulation of AFB ₁ .	

NPs conc.	AgNPs			AuNPs			Ag-AuNPs		
(µg/mL)	Growth AFB ₁			Growth	AFB1		Growth	AFB ₁	
	Dry weight (g)	Quantity (ng/100 ml)	Reduction (%)	Dry weight (g)	Quantity (ng/100 ml)	Reduction (%)	Dry weight (g)	Quantity (ng/100 ml)	Reduction (%)
Control	1.73 ± 0.02^{a}	8787	0	1.73 ± 0.02^{a}	8787	0	1.73 ± 0.02^{a}	8787	0
10,000	0.00 ± 0.00^{p}	0	100	0.00 ± 0.00^{p}	0	100	0.00 ± 0.00^{p}	0	100
5000	0.49 ± 0.01^{n}	0	100	0.38 ± 0.01°	0	100	0.49 ± 0.01^{n}	0	100
3000	0.67 ± 0.02 k	39.6	99.55	0.62 ± 0.01 ¹	0.11	99.999	0.57 ± 0.01 ^m	36	99.59
1000	0.85 ± 0.01^{j}	91.5	98.96	0.82 ± 0.01^{j}	42.3	99.52	0.67 ± 0.01 k	247	97.19
750	0.97 ± 0.01 ^h	274	96.88	0.92 ± 0.01^{i}	230	97.38	0.83 ± 0.02^{j}	797	90.93
500	1.09 ± 0.01^{f}	308	96.49	1.08 ± 0.01^{f}	271	96.92	1.02 ± 0.01 ^g	1280	85.43
250	1.45 ± 0.01^{d}	1414	83.91	1.20 ± 0.01^{e}	859	90.22	1.19 ± 0.03^{e}	6138	30.15
125	1.55 ± 0.03 ^c	3471	60.50	1.42 ± 0.01^{d}	2871	67.33	1.62 ± 0.02^{b}	6745	23.24

The data are expressed as mean value of three replicates \pm SE. Values followed by different letters are significantly different at p < 0.05.



Fig. 5. Effects of different concentrations of biosynthesized AgNPs, AuNPs, and their mixture on (A) the outgrowth of A. flavus and (B) AFB1 production.

Table 2 DPPH radical scavenging activities (%) of mycosynthesized AgNPs, AuNPs, and Ag-AuNPs.

Test NPs	Ps Concentration (μg/mL)								
	1000	500	300	100	75	50	25	12.5	
AgNPs AuNPs Ag-AuNPs BHT	79.58 ± 0.36^{b} 80.22 ± 0.22^{b} 76.85 ± 0.97^{c} 94.17 ± 0.23^{a}	76.18 ± 0.30^{c} 76.82 ± 0.08^{b} 69.17 ± 0.07^{d} 90.57 ± 0.17^{a}	65.87 ± 0.17^{c} 69.64 ± 0.18^{b} 65.13 ± 0.52^{c} 83.96 ± 0.54^{a}	60.14 ± 0.18^{c} 67.22 ± 0.21^{b} 58.49 ± 0.16^{d} 78.87 ± 0.37^{a}	56.98 ± 0.29^{c} 64.05 ± 0.17^{b} 55.59 ± 0.50^{d} 73.21 ± 0.51^{a}	$48.08 \pm 0.26^{\circ}$ $60.82 \pm 0.01^{\circ}$ $44.68 \pm 0.19^{\circ}$ $69.17 \pm 0.07^{\circ}$	$41.68 \pm 0.60c$ $56.28 \pm 0.24b$ $32.58 \pm 0.53d$ 65.13 ± 0.52^{a}	27.02 ± 0.64^{c} 40.19 ± 1.13^{b} 33.59 ± 0.33^{d} 60.14 ± 0.18^{a}	

The data are expressed as mean values of three replicates ± SE. Values followed by different letters are significantly different at p < 0.05.

ple color mycelial filtrate was indicative of AuNPs biosynthesis; these observations are in line with those reported by earlier studies [39,40]. The bioreduction of silver ions into silver-NPs is monitored by extracellular enzymes secreted by fungi [41] and in the case of bioreduction of gold ions [42].

UV-Vis spectroscopic analyses of AgNPs revealed the surface plasmon resonance band at 432 nm, while for AuNPs, surface plasmon absorbance peak was observed at 536 nm. These observations are in accordance with previous studies, where UV-visible absorbance peak at 420 nm was observed for the fungal filtrate containing AuNPs [43,44], which reported that mycosynthesized gold nanoparticles exhibited a peak corresponding to 550 nm in the absorption spectrum. The size and morphology analysis of AgNPs and AuNPs by TEM showed that AgNPs formed were monodispersed, showing relatively stable spherical structures with an average size of 5–37 nm. while AuNPs were polydispersive, showing spherical and hexagonal structures with a particle size range of 37–62 nm. These observations concur with Elgorban et al. [45] who synthesized AgNPs by using a soil isolate of Aspergillus versicolor, which were characterized as spherical shape nanoparticles with the size ranging from 5 to 30 nm, and with Abd El-Kareem and Zohri [40] who analyzed TEM images of mycosynthesized AuNPs and showed that they were well dispersed with spherical, pentagonal, and hexagonal shapes covering a size range of 5-30 nm.

Next, XRD analyses of the newly formed AgNPs and AuNPs were conducted. The crystalline quality of AgNPs was confirmed by XRD analyses, which revealed the typical fcc structure of the metal silver, while AuNPs exhibited Bragg diffraction peaks and planes of fcc in XRD analyses.

We also performed FTIR analyses of AgNPs and AuNPs to ascertain the major functional groups bound to the AgNPs and AuNPs. Several different bands were observed for both AgNPs and AuNPs at different wavelengths. For AgNPs, the band centered at 2965.86 and 2903.53 cm⁻¹ indicated aliphatic C–H stretching; a strong band at 1547.36 cm⁻¹ represented carbonyl group C=O stretching, while bands centered at 1292.03 and 1053.73 cm⁻¹ were attributed to the C–O stretching and S=O stretching functional groups in the sample. Similarly, for AuNPs, the band centered at 2972.08 and 2900.92 cm⁻¹ showed aliphatic C-H stretching, while the band at 1053.73 cm⁻¹ is linked to a S=O stretching. The other functional groups identified in the sample were -O-H hydroxyl group and carbonyl group C=O based on the presence of a weak band centered at 3658.98 cm-¹ (for hydroxyl group) and a band centered at 1596.09 cm⁻¹ (for carbonyl group). These bands indicated that these functional groups are better able to bind ions of the silver and gold metals and can biosynthesize and cap AgNPs and AuNPs to block clustering and settle the medium. According to an earlier report [46], AgNPs are bound to proteins through free carboxylic or amine groups, whilst the incidence of a peak at C=O stretching confirms the existence of carboxylic groups in the substances that are attached to AuNPs. The role of extracellular proteins in the biosynthesis of AuNPs is supported by the presence of standard FTIR spectra of proteins and peptides [47].

The effectiveness of the AgNPs, AuNPs, and their mixture to inhibit the growth and aflatoxin B_1 synthesis ability of *A. flavus* was analyzed next. Our data clearly show that the nanoparticles inhibited the growth of *A. flavus* and aflatoxin B_1 in a concentration-dependent manner. These results were expected because when AgNPs come in contact with the fungal plasma membrane, they perforate the membrane, resulting in the seepage of ions and low molecular weight compounds. Subsequently, the silver-NPs then disrupt the respiratory chain, which is then followed by halting of cell division, ultimately resulting in cell death [48]. After AgNPs treatment, the fungal genome loses its capacity

to replicate, and transcription and adenine triphosphate synthesis could be promoted [49]. When *A. flavus* is treated with AgNPs, the expression levels of *omt-A* gene, which encodes enzymes encompassed in the aflatoxins biosynthesis pathway, are reduced [50]. The antimicrobial specificity of the nanoparticles is also correlated to their total surface area; thus, the smaller the particles in size, the quicker will be their penetration in microbial cells [51].

Antioxidants are vital in decreasing the intracellular levels of reactive oxygen species, which lowers the intracellular oxidative level that in turn affects the signaling of the aflatoxin biosynthesis pathway, followed by significant suppression of aflatoxins production by *A. flavus*. In our study, at the concentrations between 12.5–100 μ g/mL, AgNPs showed antiradical activity of 27.02–79.58%, while AuNPs and Ag-AuNPs showed antiradical activity of 40.19–80.22% and 33.59–76.85%, respectively. The functional groups of the biomolecules attached to the particle's surface could be accountable for the antioxidant potency of the biofabricated AgNPs through their bioreductant capacity [52]. Antioxidants exerted a significant influence on AFB₁ accumulation by *A. flavus* [53]. Phytosynthesized silver, gold, and silver-gold nanoparticles displayed satisfactory DPPH-scavenging activities that were dosedependent at the tested concentrations [38].

5. Conclusions

In the present study, *Aspergillus versicolor* was used to biosynthesize AgNPs and AuNPs. Mycosynthesized NPs were characterized by UV–vis spectroscopy, XRD, TEM, and FTIR. Data obtained from the present investigation suggest that mycosynthesized AgNPs, AuNPs, and Ag-AuNPs exerted antifungal and antiaflatoxigenic ability against *A. flavus* and antioxidant activity using DPPH assay. More studies are required to estimate the influence and application of mycosynthesized metallic nanoparticles against different mycotoxigenic fungi, particularly for food products.

Author Contributions

- Study conception and design: H Sheikh; MF Awad
- Data collection: H Sheikh; MF Awad
- Analysis and interpretation of results: H Sheikh; MF Awad
- Draft manuscript preparation: H Sheikh; MF Awad
- Revision of the results and approved the final version of the manuscript: H Sheikh; MF Awad

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Conflict of interest

The authors declare no conflict of interest.

Supplementary material

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