Statically improved fungal laccase-mediated biogenesis of silver nanoparticles with antimicrobial applications

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ABSTRACT
Green synthesis of silver nanoparticles (AgNPs) using microbial enzymes has gained great attention owing to its advantages, such as cost-effectiveness, biocompatibility, eco-friendliness, and simplicity, over conventional physical and chemical methods. In our study, the laccase enzyme of an unconventional source like the endophytic strain Alternaria arborescens MK629314 was optimized and applied for AgNP biogenesis. Alternaria arborescens laccase enzyme production was first improved using a three-factor, five-level central composite design of 20 trials; the low-cost medium of rice bran increased the laccase production to 6.5-fold. Second, the semipurified laccase enzyme was applied as a reducing and capping agent for the synthesis of AgNPs, and all the characteristics of the formed nanoparticles were studied. The applied techniques were transmission electron microscopy, UV-vis spectroscopy, Fourier transform infrared (FTIR), and dynamic light scattering (DLS). The results of all the analyses emphasized the formation of AgNPs. AgNPs biosynthesized from 25% and 50% semipurified enzyme fractions had size ranges of 6.51 ± 5.12 and 2.51 ± 3.45 nm and showed UV-characteristic peaks at wavelengths of 416 and 404 nm, respectively. DLS analysis presented good peaks with particle size means of 96.20 and 82.30 nm and zeta potentials of −58.3 and −98.5 mv. FTIR analysis showed the appearance of distinctive bands at 3,849.22, 3,440.39, 1,637.27, 1,432.85, 1,159.01, 1,031.73, 597.825, and 520.686 cm⁻¹. The well diffusion method showed a good inhibition zone diameter of antibacterial efficiency against Escherichia coli and Staphylococcus aureus (15–17 mm) and antifungal activity against filamentous fungi Aspergillus niger and Fusarium solani (11–13 mm). The achieved results add to the growing relevance of A. arborescens as a potential applied endophytic fungus used for diverse biomedical applications.

INTRODUCTION
Metal nanoparticles showed immense popularity due to their characteristic physicochemical properties, as well as containing catalyzing, anticancer, antimicrobial, and magnetic properties (Ovais et al., 2018). Silver nanoparticles (AgNPs) are considered one of the most potential researched metal nanoparticles with high biological activities due to their unique physical properties, especially their large surface area, stability, and small tunable size. In addition, AgNPs have many advantages and are greatly demonstrated in the fields of biotechnology, pharmaceuticals, and medicine (Ammar et al., 2021; Elyamny et al., 2021).

AgNPs are generally synthesized using photochemical, electrochemical, and γ-irradiation hazardous methods, which depend on the application of toxic chemicals, resulting in environmental problems and biological hazards. In recent times, much effort has been focused, by researchers, on preparing AgNPs using green chemistry approaches (Alshammari and Abd El Aty, 2022; Eltarahony et al., 2021; Raju et al., 2014). Enzyme-mediated formation of nanoparticles is one of the recent techniques used for the safe synthesis of nanoparticles. Enzymes can catalyze the biosynthesis of nanoparticles in their active form and by amino acids released from denatured enzymes, where they act as reducing and stabilizing agents in nanoparticle biosynthesis. Also, the enzyme itself may act as a reducing and capping agent in the formation of nanoparticles (Adelere and Lateef, 2016; Sanket...
Laccase, ligninase, α-amylase, nitrate reductase, cellulase, and sulfite reductase are examples of relevant microbial enzymes in nanotechnology (Behera et al., 2022; Kumar et al., 2007; Rai and Panda, 2015; Rangnekar et al., 2007; Sanghi et al., 2011).

Laccase (EC 1.10.3.2) is the enzyme superfamily of multicopper oxidases, which is a widely distributed protein family among eukaryotes and prokaryotes (Devasia and Nair, 2016). Laccases are known for their several biological important roles in medicinal applications, organic synthesis, carbohydrate chemistry, polymer chemistry, food industries, bioremediation, and degradation of different recalcitrant compounds (Abd El Aty and Ammar, 2016; Bassanini et al., 2021; Mayolo-Deloisa et al., 2020; Sousa et al., 2021). In addition, fungal laccase may also be a talented biocatalyst for the formation of a metal nanoparticle, and very few researches have been carried out on the laccase-catalyzed synthesis of metal nanoparticles (Chaurasia et al., 2022).

In this direction, white-rot fungi were known as a potential source for biosynthesizing metal nanoparticles due to their high excretion of enzymes, such as peroxidases and laccases (Tortella et al., 2008), and some of these species, *Lentinus edodes*, are applied in the formation of AgNPs depending on laccase enzyme secretion (Lateef and Adedayo, 2015). Moreover, Sanghi et al. (2011) demonstrated that the extracellular laccase production by *Phanerochaete chrysosporium* was responsible for the biosynthesis of gold nanoparticles. Also, the laccase from *Paraconiothyrium variabile* was able to form gold nanoparticles (Faramarzi and Forootanfar, 2011).

Currently, using laccase on a commercial scale is limited due to its low productivity in fungal fermentation. For this reason, optimization of the fermentation medium is preferred for high laccase enzyme productivity. The statistical experimental central composite design (CCD) with the analysis of variance (ANOVA) is one of the useful applications which helps to study the maximum number of factors affecting enzyme production at different levels with a minimum number of experiments (Abd El Aty et al., 2016). The results analyzed by a statistically planned experiment of response surface methodology (RSM) have many advantages over those carried out by the traditional one-variable-at-a-time, with less development time and low overall costs of production (Abd El Aty et al., 2018; Shehata and Abd El Aty, 2014).

The number of reports on the eco-safe synthesis of AgNPs from fungal ascomycete enzymes was very limited. Therefore, the main goal of the present study was to evaluate the biosynthesis of AgNPs using the laccase enzyme of the endophytic strain *Alternaria arborescens* MK629314. Economic optimization of the agriculture residue medium, for high laccase production with low cost, depends on the statistical design (CCD). Finally, AgNPs, were characterized and evaluated for their antibacterial and antifungal activities.

**MATERIALS AND METHODS**

**Microorganism and qualitative assay for laccase production**

The endophytic fungus *A. arborescens* used in the present study was isolated from medicinal plants of Wadi Abu Matir, Saint Katherine Protectorate, South Sinai, Egypt, and identified based on both morphological and genetic 18S-rDNA analysis and deposited in GenBank with specific accession number MK629314 (Shaheen and Abd El Aty, 2018). The endophytic fungus was screened for laccase production using the guaiacol oxidation qualitative assay (Abd El Aty et al., 2015; Abd El Aty and Mostafa, 2013).

**Analysis of the agricultural waste rice bran (RB)**

The agricultural waste RB obtained from the local market was evaluated for the macronutrients (nitrogen, phosphorus, potassium, calcium, magnesium, and sodium) and micronutrients (iron, zinc, copper, and manganese) according to that described by Cottennie et al. (1982). The total organic carbon and ash were also evaluated in the Unit of Land Resources Evaluation and Mapping, National Research Centre, Cairo, Egypt.

**Low-cost production of laccase enzyme**

RB is used as a natural culture medium [medium of RB (RBM)] for laccase production by the endophytic fungus *A. arborescens* MK629314. The fermentation medium contained about 2 g of RB and 50 ml of distilled water in a 250 ml flask, with the addition of copper sulfate at a concentration of 1 mM on day 5 to each culture flask as a laccase inducer. The flasks were incubated for 12, 14, and 17 days at a temperature of 28°C under shaking (150 rpm) conditions.

**Three-factor five-level CCD**

The 3-factor, 5-level CCD of 20 trails was carried out to optimize the low-cost RBM for maximum laccase production. The correlation between the three variables, RB weight (A), CuSO₄ concentration (inducer) (B), and incubation period (C), and enzyme activity (Y) was studied. All three variables were investigated at the low level (−1), zero level (0), and high level (+1), respectively, with $\alpha = 1.682$, as shown in Table 1. Statistical analysis of the model was performed to evaluate the ANOVA, and the quadratic models were represented as a contour. The statistical Design-Expert® 8 software from Stat Ease, Inc., was used for the purpose of matrix construction and analysis.

<table>
<thead>
<tr>
<th>Variable code</th>
<th>Variable</th>
<th>Levels</th>
<th></th>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>A</td>
<td>RB (g/flask)</td>
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<td>3</td>
<td>4</td>
<td>5</td>
<td>5.68</td>
</tr>
<tr>
<td>B</td>
<td>CuSO₄ (mM)</td>
<td>1.32</td>
<td>2</td>
<td>3</td>
<td>4</td>
<td>4.68</td>
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<tr>
<td>C</td>
<td>Incubation period (days)</td>
<td>8.64</td>
<td>10</td>
<td>12</td>
<td>14</td>
<td>15.36</td>
</tr>
</tbody>
</table>

$\alpha = 1.682$.  

**Table 1.** CCD design for optimization of *A. arborescens* MK629314 laccase enzyme production.
Laccase assay

According to Mostafa and Abd El Aty (2018), extracellular laccase activity was measured based on the oxidation of the substrate 2,2’-azino–bis(3-ethylbenzothiazoline)6-sulphonic acid (ABTS). One unit of laccase activity was defined as the activity of an enzyme that catalyzes the conversion of 1 μmol of ABTS per minute.

Protein assay

In accordance with Lowry et al.’s (1951) description, protein concentration was determined.

Partial purification of laccase enzyme

Fractional precipitation of the laccase enzyme with acetone was applied at three different concentrations of 25%, 50%, and 75%. The specific activity (SA) and purification fold of each fraction were calculated according to the following equations (Abd El Aty et al., 2017):

\[ SA = \frac{\text{Enzyme activity} \times \text{U}}{\text{protein content} \times \text{mg}} \]  

\[ \text{Purification fold} = \frac{\text{SA of partial purified fraction}}{\text{SA of crude culture filtrate}} \]

Applications of laccase enzyme in biosynthesis of nanosilver

The partially purified fractions of laccase (25%, 50%, and 75%) in comparison with the crude enzyme were used for the synthesis of AgNPs. The reaction mixture contains 3 ml of the equal volume of the enzyme fraction and 1 mM AgNO₃, incubated in a 150 rpm rotating shaker at 30°C–33°C for 24 hours duration in dark conditions.

Characterization studies of biosynthesized AgNPs

The formation of AgNPs was preliminarily indicated by changing the color of the mixture to reddish-brown (Abd El Aty and Ammar, 2016).

UV-visible spectroscopy

The bioreduction of silver ions was monitored by a UV-visible spectrophotometer. The absorption spectrum of the reaction mixture was scanned in the range of 200–700 nm, and the sharp peak given by the UV-visible spectrum at the absorption range of 400–450 nm indicated the formation of AgNPs.

Transmission electron microscopy (TEM)

AgNPs were characterized by TEM (JEOL-2100) to detect the size and shape of nanoparticles. 2 μl of the AgNPs sample was loaded on a formvar-coated 200-mesh copper grid (Ted Pella, Redding, CA), air-dried before examination, and loaded onto a specimen holder. The size and shape of AgNPs were clarified from the TEM micrographs.

Dynamic light scattering (DLS)

A DLS was used for the analysis of the hydrodynamic diameter of particles and distribution of particles in solution. The distribution of solid particles of nanosilver in the liquid solution was determined by Nicomp Particle Sizing Systems CW388 (Nicomp 380, Inc., Santa Barbara, CA). Zetasizer Nano ZS ZEN3600 (Malvern Instruments Ltd., Worcestershire, UK) was used to analyze the AgNPs’ hydrodynamic diameters.

Fourier transform infrared (FTIR) spectroscopy

FTIR (JASCO, FT/IR-6100) was used to detect the structural characteristic and functional groups of the nanosilver solution. The peaks were recorded by drop-coating AgNPs solution into a KBr disk under a hydraulic press at 10,000 psi. Each KBr disk was scanned at the range of 400–4,000 cm⁻¹ and a resolution of 4 cm⁻¹.

Antimicrobial characters of biosynthesized AgNPs

Nanosilver biosynthesized from 25%, 50%, and 75% semipurified enzyme fractions and the crude enzyme was screened in vitro against different pathogenic strains of Gram-positive bacteria (Staphylococcus aureus ATCC29213), Gram-negative bacteria (Escherichia coli ATCC25922), and fungi (Aspergillus niger NRC53, Fusarium solani NRC15) by the agar well diffusion technique (Amnar et al., 2021; Abd El Aty et al., 2020).

RESULTS AND DISCUSSION

Laccase enzyme production

Alternaria arborescens MK629314 has a good ability to oxidize guaiacol; as shown in Figure 1A, a reddish-brown zone formed under and around a 10 mm fungal disk indicating the production of the laccase enzyme. The endophytic fungus A. arborescens MK629314 showed the ability to produce an appreciable amount of the laccase enzyme (41.05 U/ml) after incubation for 12 days in an RB medium under shaking conditions of 150 rpm (Fig. 1B). The analysis study of RB agricultural waste indicated that it contains many valuable nutrients of (9.44%) protein, (86%) organic carbon, (2.3%) phosphorus, (1.51%) nitrogen, (0.8%) potassium, (0.65%) magnesium, (240 ppm) calcium, (11,000 ppm) sodium, (380 ppm) iron, (0.384 ppm) manganese, (52 ppm) zinc, (9 ppm) copper, and (14%) ash. The results obtained are in agreement with Chawachart et al.’s (2004) study, who found that cultivation of the thermotolerant Coriolus versicolor strain RC3 on RB exhibited the highest laccase production compared to glucose, wheat bran, and rice straw. Using RB as an efficient substrate for laccase production may be related to its content of phenolic compounds (ferulic acid and vanillic acid), which were reported as inducers for laccase production (Munoz et al., 1997). On the other hand, the lignocellulosic substrate wheat straw was found
to be the most suitable for laccase production using *Tricholoma giganteum* AGHP (Patel and Gupte, 2016).

**Optimization of *A. arborescens* MK629314 laccase production**

The economic production medium (RBM) of the laccase enzyme was optimized using the RSM of CCD. The combined effect of factors tested RB weight (A), CuSO$_4$ concentration (B), and incubation period (C), and was optimized using a three-factor, five-level CCD with 20 trails for maximum laccase production. In Table 2, all the observed and predicted values of responses along with the matrix of the design were shown; the lowest response of 59.14 U/ml enzyme activity was obtained at long period incubation for 15 days with 4 g/flask RB and 3 mM CuSO$_4$ concentration. On the other hand, the highest response of 266.65 U/ml enzyme activity was noticed in the experimental trial of 4 g/flask RB, 3 mM CuSO$_4$, and an incubation period of 9 days. Various response values of laccase activity obtained with different factors’ concentrations and different interactions validate the applied CCD in accordance with a previous study (Eltarahony et al., 2020).

The results obtained indicated the great effect of incubation time, where the proteins may lose a significant amount of their catalytic activity over the long period of incubation, as indicated by Muthukumarasamy et al. (2015), who obtained the maximum laccase enzyme from *Bacillus subtilis* MTCC 2414 by using RB (267 ± 2.64 U/ml) as a substrate with a 96 hours incubation period, but the enzyme activity declined at a higher incubation time of 144 hours.

In comparison with the basal medium, the statistical design indicated that increasing the RB weight and CuSO$_4$ concentration to 4 g/flask and 3 mM, respectively, with a 9-day incubation period is more favorable for high laccase production. After the application of RSM, it was obvious that a 6.5-fold improvement in laccase enzyme production was being achieved. This result emphasized the value and necessity of applying the statistical optimization design.

The following quadratic model Equation (3) in terms of coded factors was given by the multiple regression analysis of the experimental data:

$$Y \text{ activity (U/ml)} = +188.07 + 12.38 \times A − 20.94 \times B − 43.32 \times C + 4.97 \times A \times B + 1.59 \times A \times C + 15.42 \times B \times C − 37.35 \times A^2 − 2.11 \times B^2 − 6.98 \times C^2 \quad (3)$$

$Y$ activity was the response (laccase activity). A–C were the code values of the tested variables (RB, CuSO$_4$, and incubation period, respectively). The regression equation was graphically represented by the three-dimensional response surface and the two-dimensional contour plots for understanding the interaction effects of the independent factors on the response value (Figs. 2 and 3).

Figures 2 and 3A–C show the response surface of RB and CuSO$_4$ at $C = 0$ (A), RB and incubation period at $B = 0$ (B), and CuSO$_4$ and incubation period at $A = 0$ (C) on laccase activity, respectively, keeping the other components at the fixed zero level. The 3D plot showed that when the concentrations of RB and CuSO$_4$ increased, the enzyme activity gradually increased until it reached the center point, and the elliptical 2D contour plot confirmed the significant synergistic interaction between the two factors (Fig. 3A).

Ibrahim et al. (2022) demonstrated that the contour plot shape usually points out the nature and extent of the interactions between the tested variables. In this context, a circular contour plot (Fig. 3B and C) reveals an insignificant interaction between the incubation period and concentrations of RB and CuSO$_4$. Maximum laccase activity could be achieved by increasing the concentration of RB and CuSO$_4$ while decreasing the incubation period.

The satisfactory correlation between the actual (experimental) and predicted values of laccase enzyme activity ($R$) showed as clusters near the diagonal line. The parity plot indicated a suitable fit of the model (Fig. 4).
ANOVA indicated that the applied model is highly significant, as represented in Table 3. The analysis indicated that the model’s F-value of 10.42 implies the model is significant. There is only a 0.05% chance that a “model F-value” this large could occur due to noise. In addition, goodness of fit and suitability of the model were also shown from the high values of $R^2 = 0.9037$ and Adj $R^2 = 0.8170$. Values of “Prob > F” less than 0.0500 indicate that model terms are significant. In this case, B, C, and A2 are significant model terms. In Figure 3, the relationship between the expected and actual values of laccase activity was shown. The cluster of calculations formed near the diagonal line in the plot showed the goodness of fit of the model and indicated an excellent correlation between the expected and actual values.

The validation was carried out with the optimum fermentation conditions, RB 4 g/flask with the addition of 3 mM CuSO$_4$, and incubated for 9 days. The actual value of laccase production (266.65 U/ml) was observed, which is to some extent closer to the predicted value (241.21 U/ml). The obtained results emphasized the effectiveness and validity of the applied experimental model.

**Precipitation of the laccase enzyme by acetone**

*Alternaria arborescens* laccase was partially purified from the culture filtrate by acetone at 25%, 50%, and 75% concentrations. Figure 5 shows a good purification fold (3.14) with 50% acetone, followed by a purification fold (1.36) at 25% acetone and 75% acetone, which showed the highest recovered protein and the lowest purification fold (0.49).

### Table 2. RSM of CCD of predicted and experimental values of enzyme production.

<table>
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<th>Trials</th>
<th>RB (A, g/flask)</th>
<th>CuSO$_4$ (B, mM)</th>
<th>Incubation period (C, days)</th>
<th>Laccase activity U/ml</th>
</tr>
</thead>
<tbody>
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<td>Levels</td>
<td>Levels</td>
<td>Levels</td>
<td>Experimental</td>
</tr>
<tr>
<td>1</td>
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<td>0</td>
<td>0</td>
<td>81.57 ± 1.62</td>
</tr>
<tr>
<td>2</td>
<td>$+\alpha$</td>
<td>0</td>
<td>0</td>
<td>72.40 ± 1.87</td>
</tr>
<tr>
<td>3</td>
<td>$-1$</td>
<td>$-1$</td>
<td>$+1$</td>
<td>95.09 ± 12.18</td>
</tr>
<tr>
<td>4</td>
<td>$+1$</td>
<td>$-1$</td>
<td>$+1$</td>
<td>144.79 ± 3.61</td>
</tr>
<tr>
<td>5</td>
<td>$-1$</td>
<td>$+1$</td>
<td>$-1$</td>
<td>108.60 ± 9.07</td>
</tr>
<tr>
<td>6</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>188.39 ± 8.34</td>
</tr>
<tr>
<td>7</td>
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</tr>
<tr>
<td>8</td>
<td>0</td>
<td>$-\alpha$</td>
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</tr>
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</tr>
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<td>20</td>
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<td>0</td>
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</table>

Biosynthesis of nanosilver using laccase enzyme

The semipurified enzyme fractions precipitated at 25%, 50%, and 75% acetone and crude enzyme were applied for AgNP biosynthesis. The formation of nanosilver was firstly detected by optical characterization, where the color of the solution mixture was changed from a pale yellow to reddish color with different degrees (Fig. 6). The change in color formed as the result of the silver ions reduction by the laccase enzyme. Also, the variation between the four mixtures is due to the difference between the biomolecules involved in the biosynthesis of AgNPs, in addition to the surface plasmon excitation in nanoparticles (Chaurasia *et al*., 2022; Mulvaney, 1996; Lateef and Adeeyo, 2015).

**Characterization of AgNPs**

The biosynthesized AgNPs were further characterized by a UV-visible spectrophotometer. AgNPs were scanned in ranges from 200 to 700 nm. The result obtained indicated the surface plasmon resonance of AgNPs biosynthesized by 25%, 50%, and 75% appearing at wavelengths of 416, 404, and 400 nm, respectively, which is considered a characteristic plasmon peak for AgNPs (Shaheen and Abd El Aty, 2018). No characterized peak formed by using the crude enzyme at the absorption range of 400–450 nm, indicating the absence of differentiated nanoparticles, which may be formed according to the color change of the reaction mixture but are unstable and highly aggregated (Fig. 7). This observation has further been confirmed by the TEM analysis.
Figure 2. 3D response surface plots of laccase activity by *A. arborescens* showing the interactive effects between independent significant variables.
Figure 3. 2D contour plots of laccase activity by *A. arborescens* showing the interactive effects between independent significant variables.
The size and structure of AgNPs were analyzed by high rate-TEM. The results showed well-dispersed cubic to rounded particles with different sizes biosynthesized from 25% and 50% semipurified enzyme fractions in the ranges of 6.51 ± 5.12 and 2.51 ± 3.45 nm (Fig. 8A and B). Various sizes of spherical-shaped AgNPs have been reported in previous studies (Ammar et al., 2021; Abd El Aty and Zohair, 2020). The TEM image of the AgNPs 75% enzyme fraction at 100–200 nm showed some aggregations of AgNPs with a variable size range of 28.65 ± 7.48 nm (Fig. 8C). On the other hand, TEM analysis indicated the presence of different fungal metabolites in the crude extract interferes with the good reduction and stabilization of the AgNO₃, where the image showed no dispersed nanoparticles (Fig. 8D). The obtained results indicated the importance of the purification process for perfect AgNO₃ reduction and biosynthesis of well-dispersed stable AgNPs.

The best nanoparticles formed with the 25% and 50% semipurified enzyme fractions were more characterized, and a perfect reduction process was confirmed by DLS analysis of the Zetasizer analysis and FTIR spectrum.
In this study, the DLS analysis of AgNPs of 25% and 50% presented characteristic peaks with the particle size means of 96.20–82.30 nm, with zeta potentials of −58.3 and −98.5 mV, respectively (Fig. 9). High negative values of dispersed biosynthesized AgNPs of 25% and 50% enzyme fractions appeared clearly to be advantageous for long-term colloidal stability, as mentioned by Eltarahony et al. (2018), who indicated that values greater than +25 mV or less than −25 mV display higher electrical charge on nanoparticles surface, which prevents agglomeration and flocculation by the action of potent repulsive forces between the ultrafine particles. On the other hand, Elyamny et al. (2021) showed slightly low surface charges (−5.34 mV) of Ag @ Ag₂O NCs, which tend to form aggregations.

In addition, the nanoparticle biosynthesis by 25% and 50% enzyme fractions was extended to analyze their FTIR spectrum. FTIR analysis showed the appearance of distinctive bands at 3,849.22, 3,440.39, 1,637.27, 1,432.85, 1,159.01, 1,031.73, 597.825, and 520.686 cm⁻¹ (Fig. 9). The primary amine (NH₂) was indicated by peaks that appeared at 3849.22 and 3,440.39 cm⁻¹ (Shaligram et al., 2009), where the 1,159.01 and 1,031.73 cm⁻¹ bands are due to OH and aliphatic amine deformation (Gopinath and Velusamy, 2013). In addition, the bands that appeared at 1,637.27–1,432.85 cm⁻¹ were an indication of hydroxyl group deformation, which in turn causes the silver-
metal ion bioreduction to form nanoparticles (Gopinath and Velusamy, 2013; Thirumurugan et al., 2011). Eltarahony et al. (2021) reported that the peaks in the low range of 400–700 cm$^{-1}$ denote the presence of metals in the examined sample. In the present study, characteristic peaks were observed at 597.825 and 520.686 cm$^{-1}$ that showed the P–O–C groups in phospholipids and/or S–S stretch band of protein (Eid et al., 2020).

Generally, the following study reflected that the biomolecules of the *A. arborescens* semipurified enzyme were responsible for perfect capping and stabilization of AgNPs, depending on free amine groups in the laccase enzyme proteins, which are able to bind to nanoparticles to protect them from aggregation.

**Antibacterial and antifungal activity of AgNPs**

AgNPs biosynthesized from different semipurified enzyme fractions were tested against pathogenic bacteria and fungi using the agar well diffusion technique (Fig. 10). The results showed that both AgNPs of 25% and 50% enzyme fractions had good antimicrobial efficiency against all tested pathogens (Table 4). AgNPs 25% and 50% induced a maximum inhibitory zone of 15–17 mm inhibition zone diameter (IZD) against Gram-positive and Gram-negative pathogenic bacteria strains and exhibited the inhibitory activity of 11–13 mm IZD as an antifungal effect against filamentous fungi. It has been reported that AgNPs biosynthesized from extracellular laccase of *L. edodes* showed selective antimicrobial activities against 10 clinical bacterial isolates (Lateef and Adeeyo, 2015). In this...
context, AgNPs synthesized from xylanases of *A. niger* L3 (NEA) and *Trichoderma longibrachiatum* L2 (TEA) inhibited the growth of bacteria (*E. coli*, *Klebsiella granulomatis*, *S. aureus*, and *Pseudomonas aeruginosa*) and fungi (*A. niger*, *Aspergillus flavus*, and *Aspergillus fumigatus*) (Elegbede et al., 2018).

The obtained results showed that all tested pathogens, *E. coli*, *A. niger*, and *F. solani*, are able to resist AgNPs of 75% and the crude enzyme, except *S. aureus*, which showed a weak IZD of 12 mm. Bhat et al. (2011) emphasized the bactericidal effect of AgNPs depending on the nanoparticle size. The obtained results are in agreement with this study’s findings, where small-sized AgNPs of 25% and 50% in the ranges of 6.51 ± 5.12 and 2.51 ± 3.45 nm, respectively, with a larger surface area can easily interact with containing constituents (sulfur and phosphorus) of the bacterial cell causing cell killing by attacking the respiratory chain and cell division in accordance with Lateef et al. (2016).

AgNPs of 25% and 50% showed zeta potential of high negative charges (−58.3 and −98.5 mV) on their surface, directly resulting in an interaction of these charges with the electric charge present on the surfaces of the microbes. Generally, this effect seems to be the most important factor affecting AgNPs antimicrobial activities (Ammar et al., 2021).

## CONCLUSION

This work showed the first report on *A. arborescens* laccase enzyme-mediated synthesis of AgNPs. The laccase enzyme of *A. arborescens* endophytic fungus was economically improved and optimized through a three-factor, five-level CCD of an economic agricultural waste medium. Application of RSM improved enzyme production to 266.65 U/ml in the experimental trial of 4 g/flask RB, 3 mM CuSO₄, and an incubation period of 9 days. Eco-safe synthesis of AgNPs by 25% and 50% semipurified enzyme was demonstrated, leading to the formation of well-dispersed cubic to rounded particles with 6.51 ± 5.12 and 2.51 ± 3.45 nm size ranges, and UV analysis exhibited characteristic peaks at wavelengths of 416–404 nm, respectively. In addition, good peaks with particle size means of 96.20–82.30 nm and zeta potentials of −58.3 and −98.5 mV were observed by DLS analysis. The FTIR characterization showed the proteins’ responsibility for the biosynthesized AgNPs, as shown from the obtained bands at 3,849.22, 3,440.39, 1,637.27, 1,432.85, 1,159.01, 1,031.73, 597.825, and 520.686 cm⁻¹. The antimicrobial results showed potential activity against Gram-positive *S. aureus*, Gram-negative *E. coli* (15–17 mm), and fungi *A. niger* and *F. solani* (11–13 mm). The antimicrobial properties demonstrated by the biosynthesized AgNPs can be applied as antimicrobial agents for some biomedical applications.

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## CONFLICTS OF INTEREST

The authors declare that they have no conflicts of interest.

## AUTHOR CONTRIBUTIONS

All authors made substantial contributions to conception and design, acquisition of data, or analysis and interpretation of data; took part in drafting the article or revising it critically for important intellectual content; agreed to submit to the current journal; gave final approval of the version to be published; and agree to be accountable for all aspects of the work. All the authors are eligible to be an author as per the international committee of medical journal editors (ICMJE) requirements/guidelines.

## ETICAL APPROVALS

This study does not involve experiments on animals or human subjects.

## DATA AVAILABILITY

All data generated and analyzed are included within this research article.

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## REFERENCES


## Table 4. Antimicrobial assay of biosynthesized AgNPs against pathogens.

<table>
<thead>
<tr>
<th>Treatment</th>
<th>Gram +ve bacteria</th>
<th>Gram +ve bacteria</th>
<th>Fungi</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td><em>E. coli</em> ATCC25922</td>
<td><em>S. aureus</em> ATCC29213</td>
<td><em>A. niger</em> NRC53</td>
</tr>
<tr>
<td>AgNPs 25%</td>
<td>17 ± 0.92</td>
<td>15 ± 0.71</td>
<td>11 ± 0.14</td>
</tr>
<tr>
<td>AgNPs 50%</td>
<td>16 ± 0.06</td>
<td>15 ± 0.14</td>
<td>11 ± 0.78</td>
</tr>
<tr>
<td>AgNPs 75%</td>
<td>ND</td>
<td>12 ± 0.83</td>
<td>ND</td>
</tr>
<tr>
<td>Crude enzyme</td>
<td>ND</td>
<td>ND</td>
<td>ND</td>
</tr>
</tbody>
</table>

*IZD (mm)*


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