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Immobilization and Characterization of Purified *Aspergillus flavus* Peroxidase Mediated Silver Nanoparticle Synthesis: Peroxidase Surface Reactive Residues are Implemented for Reduction of Silver Ions, More than Its Active Sites

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Abstract

Silver nanoparticles (AgNPs) have been used extensively in various therapeutic applications, molecular diagnostics. Chemical methods for synthesis of AgNPs are frequently used, however, the expenses and releases of toxic byproducts are the main limitations. Biological methods provide an alternative source for AgNPs synthesis for its cost and environmental safety. Fungi possess unique system for Ag⁺ resistance via secretion of multiple proteins for ions reduction to their nanoparticles. The objective of this study was to deciphered the mechanisms of Ag⁺ reduction to Ag⁰ by *A. flavus*. The synthesized AgNPs by *A. flavus* have molecular size ranged from 9-15 nm with visual molecular stability for abundance of stabilizing agents. *A. flavus* peroxidase was purified, immobilized on chitosan, polyacrylamide and sodium alginate, and biochemically characterized with strong potency for Ag⁺ reduction to AgNPs that are significantly stable (-30.6 mv) and UV-Vis analysis (A₃₉₀ nm). The immobilized peroxidase has a significant operative stability for continuous production of AgNPs with homogeneus molecular stability till the 5th catalytic cycle, emphasizing the efficiency of this technology for industrial scale synthesis of AgNPs. The reducing properties of peroxidase to Ag⁺ to AgNPs is mainly due to the protein functional surface reactive groups than the enzyme functional active sites. Scalingup potency and molecular stability of AgNPs produced by immobilized peroxidase provides a new feasible technical avenue for AgNPs exploitation.

Keywords: *Aspergillus flavus*; Peroxidase immobilization; Silver nanoparticles synthesis

Introduction

Silver nanoparticles (AgNPs) containing compounds have been used in various pharmaceutical and biological applications as bactericidal agents, in formulation of dental resin, sanitizer sprays, respirators, socks, toothpastes, in addition to electronics and optics industries [1]. Physical and chemical methods depending on top-down and bottom-up approaches have been mainly used for production of AgNPs. The top-down method based on mechanical grinding of bulk metals to nano-size particles followed by addition of stabilizing agents [2], while the bottom-up technique includes reduction of metals, electrochemical and sonodecomposition [3]. Other methods were recently validated for AgNPs synthesis, including chemical reduction, electrostatic complexing with anionic surfactant aerosol with NaHB,, microwave-based methods, thermal decomposition and photoreduction [4-7]. However, the major problem for the physical and chemical methods are the extremely expenses, release of toxic byproducts, constrains significant environmental risks.

Microbial biosynthesis (green chemistry) of AgNPs being the most feasible, clean, eco-friendly technology to fulfil the huge requirements of AgNPs [8], in addition to their homogeneity on shapes, optical, electrical and chemical properties [9]. Microbial synthesis of nanoparticles has been studied extensively [10,11]. Metal ion reduction has been recognized as one of the heavy metal resistance mechanisms in microorganisms. Fungal biosynthesis of nanoparticles is usually initiated by trapping of the Ag^+ ions to the surface of fungal cells, subsequent reduction by surface enzymes, peptides and reducing compounds [12], in addition to secretion of capping agents to stabilize the synthesized nanoparticles. The reduction has been

J Nanomed Nanotechnol, an open access journal ISSN: 2157-7439

implemented as the basic mechanism for nanoparticles production in all microorganisms [13]. Biosynthesis of AgNPs in fungi has been attributed to various reducing enzymes, peptides, amino acids, sugars and organic acids [14]. The possibility of large-scale production of extracellular enzymes, economic viability and simplicity in handling are the distinctive advantages of using fungi in NPs synthesis. Amino acids of enzymes and proteins especially with free amino groups, thiols are mainly implanted with particles interaction, acting as reducing and stabilizing agents of synthesized AgNPs [11,14]. Various proteins as bovine serum albumin [15], glucosidase [16], trypsin [17], pepsin and laccase [18] are mainly involved in reduction of metals to their nanoparticles. It was claimed that enzymes and proteins that have free amino groups like phenylalanine, tyrosine, tryptophan and histidine and free thiol groups of cysteine are able to catalyse the reduction of metals [19,20]. Nitrate reductases as well as peroxidase had been

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Received January 05, 2017; Accepted January 19, 2017; Published January 27, 2017

Citation: EI-Sayed ASA, Rabie GH, EI-Gazzar NS, Ali GS (2017) Immobilization and Characterization of Purified *Aspergillus flavus* Peroxidase Mediated Silver Nanoparticle Synthesis: Peroxidase Surface Reactive Residues are Implemented for Reduction of Silver Ions, More than Its Active Sites. J Nanomed Nanotechnol 8: 420. doi: 10.4172/2157-7439.1000420

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remarked as potential enzymes/proteins, having abundant reactive amino and thiols groups that participate in metals reduction [21].

The objective of this study was to emphasis the potency of *A. flavus* for AgNPs synthesis. To purify and immobilize *A. flavus* peroxidase, comparatively kinetic characterization of the free and immobilized enzymes. To assess the operative stability of the immobilized peroxidases for bulk synthesis of AgNPs.

Materials and Methods

Fungal isolates, identification, growth conditions and silver nanoparticles synthesis

Fourteen fungal species were isolated from soils contaminated with wastes from Ceramics and Photographic Industries (10th of Ramadan, El-Sharkia, Egypt). These fungal isolates were screened for their ability to reduce silver ions to silver nanoparticles according to Ahmad, Mukherjee [14] with minor modifications. Briefly, the fungal isolates were grown on liquid medium containing 0.7% KH₂PO₄, 0.2% K₂HPO₄, 0.01% MgSO₄.7H₂O, 0.06% yeast extract and 2% glucose. The cultures were incubated for 28°C for 5 days then filtered and the collected mycelia were washed thoroughly with sterile distilled water to remove any particulate. The fungal mycelia (10 gm) was suspended in 100 ml sterile potassium phosphate buffer (1 mM, pH 7.5) containing 1 mM PMSF and 1 mM EDTA, grinded thoroughly, the mixture was centrifuged at 6000 rpm for 10 min, the supernatant was used as the source of crude intracellular enzymes for synthesis of AgNPs. The crude extracts were amended with 1 mM AgNO₃, incubated for 24 h at 25°C, then the developed AgNPs was visually monitored from the intensity of yellow-brown color [14]. Distilled water supplemented with silver nitrate at the same conditions were used as control. The concentration and physical properties of the developed AgNPs were determined.

The recovered fungal isolates were identified based on morphological features [22,23]. The most potent fungal isolate producing AgNPs was molecularly identified based on the ITS sequences [24-26].

Properties of silver nanoparticles

The developed AgNPs was assessed by UV-Vis analysis using JASCO-V630 Spectrophotometer at λ 200-600 nm, normalizing to controls. The size of the developed AgNPs was determined using Zetasizer Nano S (ZEN 1600, Malvern, UK). The morphology and size of AgNPs were assessed by Transmission Electron Microscopy (TEM) (FP 5018/40, Tecnol G2 Spirit Bio TWIN) according to Ahmed et al. (2003). The presence of functional groups of amino acids as protecting agents to AgNPs was detected by Fourier Transform Infrared Spectroscopy (FTIR) (Thermo Nicolet model 6700 spectrum). The charge of silver nanoparticles was determined by Zeta potential analysis (ZEN 1600, Malvern, UK). Sterile distilled water with silver nitrate was used as negative control.

Implementation of various coenzymes for induction of AgNPs synthesis

The impact of NADH and pyridoxal 5'-phosphate (PLP) on the potency of *A. flavus* to reduce silver ions to AgNPs was assessed. Different concentrations of NADH and PLP (10-500 mM) was added to the fungal extract with 1 mM AgNO₃. The mixture was incubated at 30° C for 24 h, then the concentration and properties of the developed AgNPs was determined as described above.

Peroxidase and alcoholic dehydrogenase activities

The crude peroxidase activity was measured using O-dianisidine as substrate [27]. Briefly, the reaction contains O-dianisidine (50 mM), H_2O_2 (0.5%) and 500 µl of the crude enzyme extracts, in 1 ml total volume. The reaction was incubated for 20 min at 37°C, then the intensity of developed color was measured at 460 nm. Blanks of enzyme and substrate were used separately. One activity unit of peroxidase was defined as the amount of enzyme oxidizing 1 µmol of O-dianisidine per min under standard assay [28]. The activity of crude alcoholic dehydrogenase was determined (Vallee and Hoch, 1955). The protein concentration of the crude fungal extract was determined by Bradford assay [29] using bovine serum albumin as standard. The specific activity of peroxidase (µmol/mg) was expressed by its activity (µmol/ml) per protein concentration (mg).

Purification, molecular subunit structure and Zymogram analysis of *A. flavus* peroxidase

The total crude protein was extracted from the mycelia of *A*. *flavus* by grinding in buffer containing EDTA and PMSF as described above. The crude peroxidase was fractionated by ammonium sulfate saturation (20-70%), incubation at 4°C for 2 h, then the mixture was centrifuged for 10 min at 4000 g. The precipitated protein was collected in potassium phosphate buffer containing 1 mM EDTA and dialyzed against the same buffer till complete removal of salt traces. The enzyme was purified by gel filtration chromatography using Sephadex G₁₀₀ and G₂₀₀ columns [26,30-32]. After column equilibration with the same buffer, the sample was loaded to the column and the enzyme fractions were eluted at flow rate 0.2 ml/ min. The activity of peroxidase and its protein were assayed by the standard assay.

The molecular homogeneity and subunit structure of the purified peroxidase were assessed using SDS-PAGE [33]. The protein sample (20 μ g) was boiled in dissociation buffer Tris-HCl (50 mM, pH 7.0), 1.0% glycerol, 2.0% SDS, 5.0% 2-mercaptoethanol and 0.1% bromophenol blue. After gel running, staining with Coomassie brilliant blue R-250, the gel was decolorized with methanol-acetic acid and photographed. The molecular subunit structure was determined from the inference of authentic protein marker (PageRulerTM Plus, ThermoScientific).

Zymogram of peroxidase was analysed on native-PAGE (without SDS on gel and running buffer) [34]. After running, the gel was soaked on substrate solution (10 mM *O*-diansidine, 1.6 mM H_2O_2 in 50 mM succinate buffer of pH 5.5). The gel was incubated for 1 h at 30°C, the position of enzyme was allocated from the developed yellow color of oxidized *O*-diansidine, the gel was photographed by Digital Camera.

Synthesis of AgNPs by the partially purified A. *flavus* peroxidase

The potency of partially purified peroxidase to reduce silver nitrate to silver nanoparticles was assessed [21]. Peroxidase dissolved in potassium phosphate buffer was incubated with silver nitrate solution (1 mM) for 1 h at 30°C, distilled water with the same AgNO₃ concentration was used as blank. The evolved yellow- brown colouring of the solution reveals the development of AgNPs. The synthesized AgNPs was collected by centrifugation at 5000 rpm for 5 min and suspended in 10 ml of sterile distilled water. The synthesized AgNPs were characterized by scanning UV-Vis analysis at 200-600 nm range. The morphology, aggregation, size and topography of the developed AgNPs were determined by TEM analysis. The size of the AgNPs was measured by the DLS analysis.

Covalent and entrapment immobilization of the purified *A*. *flavus* peroxidase

To evaluate the potency of peroxidase for bulk synthesis of AgNPs, the enzyme was immobilized on different solid supports. The enzyme was covalently immobilized on chitosan [35] with slight modifications. Briefly, chitosan (1.0 g) was activated by soaking in 100 ml of 0.1 N HCl of 2.5% glutaraldehyde for 3 h at 30°C, then neutralizing the mixture pH to 7.0 by 1 N NaOH. The mixture was centrifuged for 10 min at 5000 rpm, the activated chitosan was washed with distilled water, collected and suspended in 10 ml potassium phosphate buffer (pH 7.0). The glutaraldehyde-activated chitosan has free reactive aldehyde groups that form Schiff base with the reactive amino groups of peroxidase[32]. One ml of peroxidase (35.0 µmol/mg) was added to the 5 ml activated chitosan, incubated overnight at 4°Cwith gently shaking. The mixture was centrifuged and protein concentration of the supernatant was determined by the standard assay. The amount of conjugated protein with chitosan was estimated from the difference between the residual and initial protein concentration. Activity of chitosan-peroxidase conjugates was determined by the standard assay.

The purified peroxidase was entrapped on polyacrylamide and sodium alginate [35] and its potency to synthesize AgNPs was determined. Briefly, 5 ml of acrylamide solution (30%), dissolved in potassium phosphate buffer, 100 μ l of ammonium persulfate (10%), mixed thoroughly with 1 ml of peroxidase (35.0 μ mol/mg) for 5 min, then 50 μ l of TEMED was add. The gel was polymerized in casting cassette, uniformly cut into discs, washed by buffer. The activity polyacrylamide entrapped peroxidase was determined using 0.1 g of resin, incubated with the substrate solution, similar to the native enzyme assay.

For sodium alginate immobilization, 1 ml of peroxidase (35.0 μ mol/mg) was pipetted into 5 ml of 1% sodium alginate in potassium phosphate buffer, mixed thoroughly, then dropped into 5 ml of 1 M CalCl₂ with gentle stirring. The developed pellets were collected, washed and the activity of alginate immobilized peroxidase was determined, using 0.1 g of resin amended with the substrate solution, at standard assay conditions.

Biochemical properties of the purified free and immobilized peroxidase

The biochemical properties of the native and immobilized peroxidase were assessed. The optimum pH was studied using

reaction mixture of different pHs with buffers potassium phosphate (pH 5.2-8.0) and Tris-HCl buffer (pH 8.0-10.2) at 50 mM. The activity of free peroxidase was assessed by the standard assay. The activity of immobilized enzyme was estimated using 0.1 g of the resin containing enzyme as chitosan, polyacrylamide and sodium alginate was supplemented with the 5 ml of substrate solution at the same concentration, and the activity was measured by the standard assay. The pH stability of native and immobilized peroxidase was determined by pre-incubation of the enzymes at different pH range (5.8-10.2) for 2 h at 4°C then measuring their residual activity. The effect of reaction temperature on the peroxidases activity was determined by incubating the reaction mixture at different temperature (30-50°C) and measuring the activity by standard assay.

Results and Discussion

Potency of the isolated fungal isolates for AgNPs biosynthesis

Fourteen fungal isolates were isolated from soils enriched with Ceramics Industrial Wastes. The fungal isolates were identified based on morphological features as Aspergillus niger, A. candidus, A. carneus, A. flavus, A. fumigatus, A. flavipes, A. terreus, A. oryzae, Penicillium citrinum, Cladosporiumsp, Microphomoniasp, Fusarium oxysporum, Cephalosporiummaydis and Sclerotium rolfsii. Among the fourteen isolates, only seven isolates (A. niger, A. candidus, A. flavus, A. fumigatus, A. terreus, P. citrinum, F. oxysporum) have the potentiality to reduce Ag⁺ to Ag⁰ as revealed from the UV-Vis, TEM and DLS analyses as summarized in Table 1. While, A. carneus, A. flavipes, A. oryzae, Cladosporium sp., Microphomonia sp. and S. rolfsii lack the potentiality to synthesize Ag⁰ as revealed from the measured physical parameters. Five species of the genus Aspergillus namely; A. flavus, A. fumigatus, A. candidus, A. niger and A. terreus had the highest reducing potency to silver ions synthesizing AgNPs based on the visual color, UV-Vis and DLS analyses (Figure 1). The appearance of yellow-brown color suggesting the formation of AgNPs, the lack of precipitations or agglomeration assure the homogeneity and stability of developed AgNPs. Fungal extract without AgNO₃ and sterile distilled water with AgNO₃ as controls had no change in color of solution under the same conditions. From the UV-Vis analysis, the maximum absorption of the developed AgNPs by A. flavus, A. fumigatus, A. candidus, A. niger and A. terreus was centred at 436, 420, 430, 400 and 400 nm, respectively. The absorption peak at 260-280 nm on the fungal extracts reveals the presence of DNA, proteins or aromatic amino acids that

No.	Fungal isolate	Absorption peak (λ nm)	Zeta potential (mv)	Size of AgNPs (nm)	
				DLS	TEM
1	Aspergillus niger	430	-	78	-
2	Aspergillus candidus	400	-18.9, -39.9	23	14-18
3	Aspergillus carneus	-	-	-	-
4	Aspergillus flavus	436	-30.4	9	6-13
5	Aspergillus fumigatus	420	-14.9, -38.9	33	13-21
6	Aspergillus flavipes	-	-	-	-
7	Aspergillus terreus	400	-8.7, 18.9, 30.4	36	14-27
8	Aspergillus oryzae	-	-	-	-
9	Penicillium citrinum	390	-	80	24-27
10	Cladosporium sp.	-	-	-	-
11	Microphomonia sp.	-	-	-	-
12	Fusarium oxysporum	430	-29.2	11	13-19
13	Cephalosporium maydis	-	-	-	-
14	Sclerotium rolfsii	-	-	-	-

Table 1: Physical properties of AgNPs synthesized by various fungal isolates isolated from heavy metals contaminated samples.



grown for 5 days, the mycelial pellets were collected, washed and grinded in buffer containing PMSF and EDTA (protease inhibitors). The mycelium was amended with 1 mM AgNO₃ for 10 hrs then the visual appearance was photographed (A), UV-Vis analysis (B), Size analysis (C), and zeta potential (D).

involved in reduction of Ag^+ , stabilizing these particles [36]. The average hydrodynamic diameter and polydispersity indexes of the developed AgNPs for the experimented fungi was evaluated by DLS analysis (Figure 1C). The average diameter of AgNPs from *A. flavus*, *A. fumigatus*, *A. candidus*, *A. niger* and *A. terreus* was 9, 33, 23, 78 and 36 nm, respectively. To assess the degree of stability of synthesized AgNPs, the charge was measured by Zeta potential. Zeta potential of AgNPs from *A. flavus* was -30.4 mV with one sharp peak, suggesting the higher stability of this NPs. Two zeta potential peaks of AgNPs from *A. fumigatus* (-14.9 mV, -38.9 mV) and *A. candidus* (-18.9 mV, 39.9 mV) were detected, while, AgNPs suspension from *A. terreus* had three major peaks at -8.7, 18.9 and 30.4 mV. The higher negative charge of AgNPs as measured by zeta potential confirm the repulsion of the synthesized particles, subsequently led to stability and mono-dispersity of the synthesized AgNPs solution [37,38].

The size and shape of the synthesized AgNPs by the experimented fungi were determined from the TEM analysis (Figure 2). AgNPs from different fungal species had variable shapes that being depends on the identity of reducing agents and stabilizing compounds from each species, however, almost of the fungal synthesized AgNPs had spherical and triangular shapes. AgNPs diameter from A. flavus, A. fumigatus, A. candidus, A. terreus, and A. niger ranged from 6-13 nm, 13-21 nm, 14-18 nm, 14-27 nm and 25-39 nm, respectively. Visually, AgNPs synthesized by A. flavus extracts was relatively stable till more than 25 days with no aggregations that might be due to the presence of stabilizing protein, in contrary to other Aspergilli which the color of AgNPs solution turned to deep brown, black due to the molecular aggregations. The colloidal stability of AgNPs synthesized by A. flavus extracts are consistent with those reported for other fungi [14] and Azospirillum [21]. To further validate the presence of stabilizing proteins, A. flavus extracts was analysed by FTIR, confirming the presence of various functional groups at 1643.2 cm⁻¹, 1604.7 cm⁻¹, 1801.4 cm⁻¹, 3994.5 cm⁻¹, and 3201.6 cm⁻¹ corresponding to carbonyl residues and peptide bonds of proteins (Figure 2F). The band at 1801.4 cm⁻¹ referred to C=C alkenyl group



Figure 2: Transmission Electron Microscope (TEM) image of silver nanoparticle synthesized by the crude intracellular extracts of *A. flavus* (A), *A. fumigatus* (B), *A. candidius* (C), *A. terreus* (D), and *A. Niger* (E).

stretch of protein in solution, whereas the bands at 1643.2 cm⁻¹ and 1604.7 cm⁻¹ are belonging to the amide I and II bonds. Similar results showing the presence of aldehyde and keto groups as stabilizing and capping agents to AgNPs synthesized from fungi [37,39]. The bands at 1643.2 cm⁻¹ are belongs to carbonyl stretch of unsaturated aldehyde, ketone and alkene group, while the bands at 1604.7 cm⁻¹ are belongs to NH₂ stretch vibrations in the amide linkages of proteins. Consistently, AgNPs biosynthesis by filamentous fungi was extensively documented [40-42]. The strong fungal AgNPs productivity has been attributed to secretion of abundant extracellular redox proteins, diverse metal reductases, thus reducing soluble metal ions into their insoluble nanocrystals [43].

Thus, *A. flavus* was selected for further biochemical studies to elucidate its reducing potency which implemented for reduction of silver nitrate to AgNPs.

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Biochemical identity of *A. flavus* metabolites implicated with synthesis of AgNPs

To determine the coenzyme dependence of A. flavus reducing agents that implemented for AgNPs synthesis, various coenzymes as NADH and PLP were supplemented to fungal extracts and their potency to reduce Ag+ was determined. Apparently, the reducing properties of A. flavus extracts to synthesize AgNPs was proportionally enhanced with NADH concentration (0, 1, 5, 10, 50 mM) as revealed from visual appearance and UV-Vis analysis (Figure S1). Distilled water containing the same NADH concentrations was used as negative controls. At 50 mM NADH, the synthesized AgNPs was increased by three folds comparing to zero NADPH. The significant reduction of AgNO₂ to AgNPs with NADPH endorsing the implementation of NADPH-dependent reductases for catalysing this reaction. However, incorporation of PLP to A. flavus extracts had no significant effect on enhancing AgNPs synthesis revealing the lack of participation of PLP-dependent enzymes on this process (Data not shown). To validate the hypothesis of NADH dependency of reductases on enhancement of AgNPs synthesis, the activity of NADH-dependent reductases particularly peroxidase and dehydrogenase were determined. In absence of NADPH, the activity of crude peroxidase and dehydrogenase were 8 µmol/ml and 2.7 µmol/ml, respectively (Figure S1). However, the activity of peroxidase and dehydrogenase was increased by about 5 times at 50 mM NADPH comparing to positive control. Consistently with these results, peroxidase was reported frequently as efficient reducing agents to Ag⁺ to AgNPs [21,36,44]. For the higher activity of peroxidase and statistically significant correlation with incorporation of NADH, we have been motivated to further purify and characterize this enzyme to synthesize AgNPs for more technological exploitations.

Purification, molecular mass and AgNPs biosynthesis by *A*. *flavus* peroxidase

NADPH-dependent peroxidase was purified from A. flavus extract by salting out and gel filtration chromatography, by the salting out (20-70% saturation), the activity of peroxidase was increased to 20 µmol/ mg comparing to 8 µmol/mg of the crude enzyme, i.e. purified by about 2.5 fold. By the gel filtration chromatography using Sephadex G100, followed by Sephadex G200, the specific activity of peroxidase was increased to 35 µmol/mg, with 4.4 purification fold. The active fractions of peroxidase were gathered, concentrated by dialysis and its molecular homogeneity and subunit structure was analysed by SDS-PAGE. The molecular mass of purified A. flavus peroxidase was 70 kDa under SDS-PAGE, while its entire molecular mass was 140 kDa under native-PAGE (Figure S2). Peroxidase from A. flavus has two identical subunits on their molecular mass, i.e. homodimer identity. From the Zymogram analysis, the activity of peroxidase was authenticated from the development of brown color due to the oxidation of substrate O-diansidine, allocating the enzyme on gel.

The potency of purified peroxidase to reduce AgNO₃ for synthesis AgNPs was verified, two ml of the purified peroxidase (35 μ mol/mg/ ml) was amended with 1 mM AgNO₃, incubated at 30°C for 1 h. From the visual appearance, a greyish plasmon color was clearly developed revealing the synthesis of AgNPs in contrary to the colourless solution of control (Figure S2).

Immobilization and biochemical properties of the free and immobilized peroxidase

To evaluate the potency of peroxidase for bulk synthesis of AgNPs, the enzyme was immobilized on different solid supports such as

chitosan, polyacrylamide and sodium alginate. Overall, the activity of purified A. flavus peroxidase was slightly reduced upon chemical conjugation. The activity of free peroxidase was 35.0 $\mu mol/mg,$ while the activity of enzyme conjugated with chitosan, sodium alginate and polyacrylamide was 24.3, 23.2 and 21.5 $\mu mol/mg,$ respectively. The yield of enzyme immobilization on chitosan, polyacrylamide and sodium alginate was 69.2, 68.4 and 66.2%, respectively. Consistently, reduction of enzymes activity upon chemical conjugation was frequently reported [32]. Subsequently, to increase the yield of immobilization, different amino acids were used as active site protectants to block/mask the catalytic sites during the conjugation process [35]. Various amino acids such as lysine, arginine, cysteine and tyrosine as active sites containing residues of peroxidase [45] were incorporated into the enzyme solution at 100 mM, prior the immobilization process. Interestingly, lysine and arginine have an obvious effect on increasing the yield of peroxidase immobilization. Upon addition of lysine and arginine (separately), the yield of peroxidase immobilization was increased by 20% for chitosan, polyacrylamide and sodium alginate than the corresponding controls (Figure S2). Moreover, by using lysine and arginine together as active site protectants, the immobilization yield of peroxidase was dramatically increased by about 33% than corresponding controls. Thus, the specific activity of peroxidase immobilized on chitosan, polyacrylamide and sodium alginate was 31.2, 30.5 and 29.1 µmol/mg, respectively, comparing to the free enzyme (35 µmol/mg).

To further evaluate the effect of chemical conjugation on the catalytic and structural properties of peroxidase, the biochemical properties of the free and immobilized enzyme was concurrently determined. The effect of medium pH on the activity of free and immobilized peroxidase was assessed. From the obtained results (Data not shown), the free and immobilized peroxidase have the same catalytic pattern in response to the reaction pH ranged from pH 5.8-10.2. The maximum activity of free and immobilized peroxidase was reported at reaction pH 7.5, with significant reduction on their activities at higher acidic and alkaline pHs. At reaction pH 4.2 and pH 10.2, the activity of free and immobilized peroxidases was reduced by about 40% comparing to control at 7.5 pH, suggesting the negligible effect of chemical modification by the current compounds on the ionic states of peroxidase. The pH stability of free and immobilized peroxidase was evaluated at pH range from 6.0-9.0. The highest stability of free and immobilized peroxidase was detected at pH range 7.4-7.8, with an obvious decreasing to the enzymes activities by about 55% at pH 6.0 and 9.0, comparing to control. The significant reduction of the enzymes activities at pH 6.0 and 9.0, might be due to the enzyme denaturing or changing the ionic state of the enzymes thus masking the proper interaction with their substrates. The current approaches of peroxidase modification have no significant effect on the kinetics of catalysis and structure of peroxidase.

The effect of reaction temperature $(30-50^{\circ}\text{C})$ on activity of the free and immobilized peroxidase was evaluated. The maximum activity of the free and immobilized peroxidase was recorded at reaction temperature 35°C and 40°C, with obvious decreasing to the enzymes activities at higher temperature (50°C). At 40°C, the activity of peroxidase immobilized on chitosan, polyacrylamide and sodium alginate was increased to 34.0, 32.3 and 30.9 µmol/mg, respectively, i.e. by about 6.0%-9.0% more than control (at 35°C). However, the activity of peroxidases was dramatically decreased by 55% by incubation at 50°C, suggesting the enzyme denaturation. The slight increasing on optimum reaction temperature for immobilized peroxidase over the free enzyme seems to be reasonable due to increasing the diffusion of substrate influx and product efflux during catalysis process [32]. The thermal stability of free and immobilized peroxidase was evaluated by

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pre-incubation of the enzymes at different temperature then measuring their residual activity. From the kinetics of thermal stability (Table 2), the immobilized peroxidase exhibited a higher thermal structural stability comparing to free peroxidase. At 40°C, the half-life time $(T_{\mu\nu})$ of peroxidase immobilized on chitosan, polyacrylamide and sodium alginate was 3.6, 7.8 and 7.8 h, respectively, while it was 2.7 h for the free enzyme. At 40°C, the enzyme structural stability was increased by 1.4, 2.9 and 2.9 fold upon conjugation with chitosan, polyacrylamide and sodium alginate, respectively. Immobilization has a dramatically positive effect on catalytic and structural stabilities of peroxidase against thermal denaturing as revealed from thermal treatment at 60°C. The structural stability of peroxidase conjugated with chitosan, polyacrylamide and sodium alginate was increased by 33.6%, 40.2% and 35.0%, respectively, normalizing to the free enzyme at 70°C. The half-life temperature (T_m) of the chemically conjugated peroxidase was 2.8-3.2 h comparing to 2.4 h to free peroxidase. In addition, the thermal inactivation rate (K) of peroxidase was dramatically reduced by chemical conjugation than the free enzyme. From the thermal kinetic properties of the free and immobilized enzyme, the current chemical modification approaches strongly stabilize the structural and catalytic properties of enzymes as consistent with those reported for other enzymes [32].

AgNPs synthesis and operative stability of the free and immobilized peroxidase

The actual potency of free and immobilized peroxidase to reduce silver ions to AgNPs was investigated. Five ml of free and immobilized peroxidase (at the same concentration 6.0 μ mol/mg) in 15 ml tubes were amended with 1 mM AgNO₃ incubated for 2 h at 40°C with gentle stirring. The synthesis of AgNPs was visually monitored from the development of yellow color comparing to control. The reaction mixture was withdrawn, the developed AgNPs was centrifuged, washed and re-suspended in distilled water. The concentration and size of AgNPs were measured by UV-Vis, DLS and TEM analyses. The size of developed AgNPs from the activity of immobilized peroxidase on chitosan, polyacrylamide and sodium alginate was ranged from 13-16 nm, 16-22 nm and 25-39 nm, respectively, while it was 15-30 nm for the free enzyme. From the DLS analysis, the size of AgNPs synthesized by native peroxidase and immobilized on chitosan, polyacrylamide and sodium alginate was around 80 nm (Figure 3).

Enzyme	Temp. (°C)	T _{1/2} (hr)	SF (%)	Tm (°C)	
	40	5.31	100		
Free Derevidere	50	2.7	100	- A	
Free Peroxidase	60	2.38	100	54	
	70	1.94	100		
	30	9.01	169.7	62	
	40	3.57	132.3		
Chitosan-peroxidase	50	2	84.03		
	60	1.86	95.9		
	30	6.77	127.5	68.7	
Sodium- alginate	40	7.78	288.1		
Peroxidase	50	2.37	100		
	60	3	154.6		
	30	6.6	127.5		
Debreendenside	40	7.7	285.1		
peroxidase	50	2.4	100.84	66.6	
	60	3	154.6	00.0	

 Table 2: Thermal stability kinetic parameters of the purified peroxidase immobilized on chitosan, sodium alginate and polyacrylamide.

The operative stability of immobilized peroxidase for continuous reduction of Ag+ was assessed. The pellets of peroxidase immobilized on various resins was washed, supplemented with 1 mM AgNO₃ incubated under the standard conditions, their catalytic stability for mass production of AgNPs was evaluated via five successive reaction cycles. The physical properties of synthesized AgNPs were assessed for each cycle. For the free peroxidase, the developed AgNPs have maximum absorption at A $_{_{390}}$ nm, with the actual size 70-81.0 nm by DLS analysis, while from the TEM micrographs the AgNPs appears spherical, oval shaped with size ranged from 15-30 nm (Figure 3). From the operative stability, the immobilized peroxidase has the potency for synthesis of AgNPs till the 5th catalytic cycle. Due to the presence of free reactive aldehyde groups that might participates in reduction of Ag⁺ to Ag⁰ [46], blanks of chitosan, polyacrylamide and sodium alginate were used. By the 3rd catalytic cycle, the reductive potency of immobilized peroxidase to Ag⁺ was reduced by about 50%, and reduced by about 70% by the 5th catalytic cycle, normalizing to the first cycle. The AgNPs concentration for each cycle was calculated from the intensity of absorption at 390-420 nm. For the immobilized peroxidases, the yield of synthesized AgNPs was decreased sequentially with the reaction cycling, that could be due to the release of enzyme during washing or even enzyme denaturing. Practically, the immobilized peroxidase being economically more feasible than free enzyme that it can be used for bulk production of AgNPs for at least three catalytic cycles, thus, producing AgNPs by three times more than the free enzyme. However, we had assumed that part of the synthesized AgNPs was adsorbed to the surface of chitosan, or entrapped to the polyacrylamide or sodium



Figure 3: DLS and TEM micrographs of the synthesized AgNPs by the free and immobilized peroxidase. The size of AgNPs synthesized by free and immobilized peroxidase by DLS analysis (Right panel), and morphology and size of AgNPs by TEM analysis (Left panel).

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alginate resins. To validate this hypothesis, after the 2nd reaction cycle, the pellets of chitosan immobilized peroxidase were washed thoroughly by sterile distilled water and analysed by TEM. The TEM micrographs (Figure S3) ensure the presence of AgNPs adsorbed to the surface of chitosan pellets. Thus, in spite of applicable feasibility of immobilized peroxidase as a new technology for Ag⁰ biosynthesis, adsorption of Ag⁰ to the surface of chitosan pellets is one of raised technical arguments.

Implementation of surface thiols or surface reactive amino groups of enzymes on reduction of silver ions to AgNPs, is one of the technical arguments. To understand the identity of reducing agents to Ag⁺, the native enzyme was denatured by autoclaving for 30 min, then amended with silver nitrate solution at 1 mM, incubated under authentic conditions and the development of AgNPs was visually monitored (Figure S3). Strikingly, the autoclaved and non-autoclaved (control) peroxidase have the same reducing effect of silver ions to AgNPs, that implies the surface thiols or amino groups or aldehydes of amino acids act as reducing agents of silver ions to AgNPs in addition to its stabilization. Thus, reduction of Ag⁺ to Ag⁰ is mainly due to surface reactive groups of peroxidase as protein but does not to its functionality as enzyme. Similar results reporting the responsibility of surface free amino groups, cysteine residues and carboxyl groups on electrostatic interaction with Ag⁺ ions to form Ag⁰ [12]. Consistently, several enzymes such as glucosidase [16], trypsin [17], pepsin [47], laccase [18], lysozyme, and fibrinolytic enzymes [48] were used as reducing peptides for nanoparticles synthesis due to their reactive amino acids, but not for their functionality as enzymes. Amino acids as phenylalanine, tyrosine, tryptophan, cysteine, histidine, and lysine are the major amino acids depicted for reduction of ions to the corresponding nanoparticles and their stabilization [20].

Conclusion

The potency of *Aspergillus flavus* for reduction of silver ions to its nanoparticles has been emphasized in terms of size, quantity and morphology. The identity of silver ions reducing agents from *A. flavus* was disclosed as NADPH-dependent reductases, peroxidase. Peroxidase was purified from *A. flavus* and immobilized into solid support and biochemically characterized. To the best of knowledge, this is first report emphasize the operative stability of immobilized peroxidase for synthesis of AgNPs on industrial scale for various technological exploitations.

Acknowledgements

This work was supported by Botany and Microbiology Department, Faculty of Science, Zagazig University, Egypt.

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