The synergistic effect of biosynthesized silver nanoparticles from a combined extract of parsley, corn silk, and gum arabic: in vivo antioxidant, anti-inflammatory and antimicrobial activities

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Abstract

Microbial resistance, oxidative stress, and inflammatory conditions are among the leading causes of death worldwide. In the current work, silver nanoparticles (AgNPs) were biosynthesized using the aqueous extracts of parsley, corn silk (CS), gum Arabic (GA) or combination of the three extracts. The formed nanoparticles were characterized using three techniques including transmission electron microscopy (TEM), UV-visible spectrophotometer and Fourier-transform infrared spectroscopy (FTIR). The antioxidant, anti-inflammatory, and antimicrobial activities were tested for the formed nanoparticles, the aqueous extracts of each of the three plants and their combination. Oxidative stress was induced by alloxan which promoted the development of diabetes mellitus in rats. Inflammation was induced by injecting carrageenan in rats’ paws. Pathogenic microorganisms causing serious urinary tract infection (UTI) were selected for the antimicrobial assay. All aqueous extracts and the biosynthesized AgNPs showed variable degrees of antioxidant, anti-inflammatory and antimicrobial activities, however, the AgNPs biosynthesized by the combination of the three aqueous extracts was the most effective one. LC/MS was done to identify the compounds present in the crude extracts that may be responsible for the observed biological activities. LC/MS resulted in the identification of 13 compounds. Docking experiments on COX-1 (cyclooxygenase-1) and COX-2 (cyclooxygenase-2) were performed to determine the compounds responsible for the anti-inflammatory activity of the extracts. The results showed that silver nanoparticles synthesized by the combination of the three aqueous extracts are considered promising candidates for the development of antioxidant, anti-inflammatory and antimicrobial agents.

1. Introduction

Nanotechnology is a rapidly expanding field which involves the synthesis and characterization of noble metals such as silver, gold, and platinum as nanoparticles. The importance of nanotechnology relies on its wide applications in drug delivery, bioengineering, textile engineering, biological labeling, biotechnology, catalysis, water treatment and the detection of genetic disorders [1–4]. Nanoparticles are of maximum size of 100 nm showing different chemical, optical, mechanical, electronic and magnetic properties than larger particles due to the variation in specific characteristics such as shape, size and atomic distribution [1, 5]. Many methods are employed for the synthesis of nanomaterials such as heat evaporation [6], chemical reduction [7–9],
photochemical [10], electrochemical [7, 11], reverse micelle [12], thermal decomposition [13], radiation [7, 14] and microwave-assisted [7] methods. Most of these methods require the use of hazardous chemicals and high energy for the preparation of nanoparticles. Biological synthesis of nanoparticles involves the use of natural materials such as plants, bacteria, fungi [1, 5, 15]. The use of plants for the synthesis of nanoparticles have an advantage over other biological methods as it does not involve the use of cell culture, and does not need longer incubation time required for the reduction of metal ions [16, 17]. Plants are known to contain various secondary metabolites such as alkaloids, terpenoids, flavonoids and tannins which provide suitable reducing and surface agents for the nanoparticle synthesis and stabilization. Biopolymers such as cellulose, chitosan, alginate, dextran and tree gums are another family of natural sources which were used for the reduction and stabilization of nanoparticles [17, 18].

Recently, pathogenic bacteria and fungi such as Staphylococcus spp., Enterococci spp., Klebsiella pneumoniae, and Pseudomonas spp. demonstrated resistance to commercially available antimicrobial agents at an increasing rate and has become a global threat especially in developing countries. To obviate this, nanoparticles were recently used which provide a new strategy for the development of new antimicrobial agents [2, 19–22]. Among noble metals, silver shows disinfecting, antimicrobial properties and tremendous medicinal value [7, 23]. AgNPs show antimicrobial activity by the direct attachment to the cell wall, disturbing cell-wall permeability, and cellular respiration. These nanoparticles penetrate microbial cells causing damage by interacting with DNA and proteins. AgNPs perform this activity without affecting normal cells and does not easily provoke microbial resistance [24]. AgNPs possess strong anti-inflammatory and antioxidant activities. AgNPs showed effective antibacterial efficacy against Klebsiella pneumoniae, Escherichia coli, and S. aureus in previous reports [25–27]. Thus, the green synthesis of spherical AgNPs with potent antimicrobial, anti-inflammatory, and antioxidant properties is an interesting target. These nanoparticles can be utilized in food, cosmetics, biomedical and pharmaceutical industries [28].

Plants were used for the green synthesis of nanoparticles and the developed green protocols demonstrated many advantages over chemical synthesis. Certain plants rich in antioxidants were superior in the green synthesis of biologically active nanoparticles. The use of local plants, plants exudates, and plants waste products for the synthesis of nanoparticles emerged as an interesting methodology for the preparation of biologically active nanoparticles with minimal cost and adverse effects on human beings and environment. Parsley (Petroselinum crispum) leaves are a rich source of ascorbic acid which plays a vital role as a strong reducing agent in the synthesis of nanoparticles. AgNPs synthesized from silky hairs of corn (Stigma maydis) displayed potent antibacterial activity against foodborne pathogenic bacteria such as Listeria monocytogenes, Bacillus cereus, E. coli, Staphylococcus aureus and Salmonella Typhimurium, exhibited potent synergistic antibacterial activity with standard antibiotics and antifungal activity with amphotericin B. In previous reports, the corn silk (CS) AgNPs showed potent antioxidant activity as demonstrated by 2,2-diphenyl-1-picrylhydrazyl (DPPH) radical scavenging, 2,2’-azino-bis(3-ethylbenzothiazoline-6-sulfonic acid (ABTS) radical scavenging, nitric oxide scavenging and reducing power assays [28].

Tree gums were used as a template for the synthesis of various metals and metal oxide nanoparticles due to the presence of polysaccharides which are effective reducing agents. Also, gums are non-toxic and biodegradable materials used for the stabilization of nanoparticles [18, 29]. The use of gum Arabic (GA) for coating of AgNPs stabilizes the nanoparticles against aggregation, which enhance the transportation and toxicity of the synthesized nanoparticles. GA AgNPs are effective antimicrobial agents against E. coli, Pseudomonas aeruginosa and Micrococcus luteus bacteria [30, 31].

The current study focuses on the green synthesis of AgNPs using three different aqueous plant extracts namely; parsley, CS, or GA. We selected these plants because they are locally available, inexpensive and generally used in folk medicine as a combination therapy for the treatment of urinary tract disorders resulting from inflammation, oxidative stress or infection. We compared the effectiveness of each plant extract before and after the synthesis of AgNPs as antimicrobial, anti-inflammatory and antioxidant agents. A combined extract was formulated of the three previously mentioned aqueous extracts to investigate the synergistic effect of the three different extracts. LC/MS was done to identify the possible compounds that may be responsible for the synthesis and stabilization of the AgNPs. Docking experiment was employed on the identified compounds to determine the possible compounds responsible for the anti-inflammatory activity.

2. Materials and methods

2.1. Plant material
Fresh aerials parts of P. crispum (parsley), S. maydis (CS) and gummy exudate of Acacia senegal (GA) were purchased from the local markets, identified and authenticated by Dr Nada Mostafa, Pharmacognosy Department, Faculty of Pharmacy, Ain-Shams University. Voucher samples were deposited at the Herbarium of
Pharmacognosy Department, Faculty of Pharmacy, Ain-Shams University with the numbers: Parsley (PHG-P-PC 198), corn silk (PHG-P-ZM 197) and gum Arabic (PHG-P-AS 199). Parsley herb and CS were carefully washed with tap water followed by distilled water to remove impurities. They were left to dry for several days at room temperature to remove any moisture. GA was sieved to remove any foreign matters. The dried plant parts and the sieved GA were then ground with an electric blender to obtain fine powder that was stored in amber-colored and airtight bottles in the fridge until further use.

2.2. Animals
All animal procedures and care were conducted according to the general guidelines of the Research Ethics Committee of the National Research Center, Cairo, which conformed with the guiding principles of the International Council on Harmonization and the Islamic Organization for Medical Sciences, the United States Office for Human Research Protections and the United States Code of Federal Regulations and operated under Federal Wide Assurance No.FWA00006444.

2.3. Chemicals
All reagents were purchased of analytical grade and used without any further purification. Silver nitrate (AgNO₃) ≥ 99.0% purity, carrageenan, alloxa were purchased from Sigma-Aldrich (Sigma Aldrich, St. Lewis, USA), Indomethacin was purchased from Epic (Epic, Egyptian Int. Pharmaceutical industries Co., Cairo, Egypt), Metformin (Cidophage)® was obtained from Chemical Industries Development (Chemical Industries Development (CID), Giza, Egypt), Vitamin E was obtained from Pharco (Pharco Pharmaceutical Co., Alexandria, Egypt). Distilled water was used for the preparation of the aqueous extracts in all the experiments.

2.4. Biochemical kits
Biodiagnostic kit for the assessment of blood glucose level and Biodiagnostic Glutathione kit for the assessment of the antioxidant activity were supplied by Diamond Diagnostics (Diamond Diagnostics Chemical Company, Cairo, Egypt).

2.5. Microorganisms
Bacterial and yeast cultures used in this study were obtained from the Microbial Type Culture Collection (MTCC) IMTECH, Chandigarh and from the American type culture collection (ATCC). The bacterial strains were Staphylococcus saprophyticus (ATCC 15305), Enterococcus faecalis (ATCC 29212), Escherichia coli (MTCC 443) and Pseudomonas aeruginosa (MTCC 1034), and the yeast strains of Candida were Candida albicans (MTCC 183, 227) and for Cryptococcus, the used strain was Cryptococcus neoformans (RCMB 0049001). These strains were used for the in vitro evaluation of the antimicrobial activity. All the tested microorganisms were supplied by the Microbiology Department, Faculty of Sciences, Al-Åzhar University.

2.6. Synthesis of silver nanoparticles by the use of aqueous plants’ extract
Parsley herb, CS and GA (5 grams) were added to 100 ml distilled water and kept in a water bath at 60 °C for 30 min, except for GA as the water bath time was extended to 60 min to deactivate the oxidase enzyme. The decoction was then filtered through Whatman No.1 filter paper. The filtrate (aqueous extract) was used as a control and for comparison in the characterization and biological studies. The synthesis of silver nanoparticles was done by the addition of different aqueous extracts to 1 mM (0.001 M) silver nitrate in a ratio (2:10) followed by the addition of two drops of 1 N NaOH and kept in a water bath for 10 min at 60 °C.

2.7. Characterization of the biosynthesized AgNPs
Characterization of the biosynthesized AgNPs was done using three techniques: TEM, UV-visible spectrophotometer and FTIR. TEM was performed by the addition of a drop of AgNPs solution on a copper grid, which was coated with carbon-supported film then left to dry under ambient conditions. After 10 min, the shape and size of the synthesized AgNPs were analyzed using TEM (JEOL model JEM-1010EX). The formation of AgNPs was monitored by measuring the UV-visible spectrum of the reaction medium in the wavelength range from 300 to 600 nm using a UV-visible spectrometer (Shimadzu A16351) using distilled water as the reference. FTIR was used to identify the functional groups in the aqueous extracts responsible for the capping and stabilization of AgNPs. We used Shimadzu FTIR-8400S in the wavelength range 4000–500 cm⁻¹.

2.8. Experimental design
2.8.1. Antioxidant activity
The antioxidant activity of parsley, CS or GA aqueous extracts and their biosynthesized AgNPs, as well as the combination formula, were evaluated in terms of the reduced glutathione (GSH) level and compared with that of
vitamin E as the standard [32]. This method is based on the reduction of 5,5'-dithiobis(2-nitrobenzoic acid) (DTNB) with GSH to produce a yellow-colored compound. The reduced chromogen is directly proportional to the GSH concentration and its absorbance can be measured at 405 nm.

Calculation of Glutathione conc.: GSH in blood = \( A_{\text{sample}} \times 66.66 \) (mg/dl)

\( A_{\text{sample}} \) = absorbance of the sample at 405 nm.

Male albino rats of Sprague Dawley Strain (130–140 g) were kept under the same hygienic conditions and were given an intraperitoneal injection of alloxan (150 mg kg\(^{-1}\) body weight) to induce diabetes mellitus [33]. Hyperglycemia was assessed after 72 h by measuring the blood glucose level [34]. Animals were randomly classified into 12 groups and each group contained five animals. The groups were treated according to the following scheme: Group I: Normal group that received 1 ml of saline and served as the normal healthy control; Group II: Diabetic group that received 1 ml saline and served as the control; Group III: Diabetic group that received 1 ml AgNO\(_3\) and served as the positive control group; Group IV: Diabetic group that received 7.5 mg kg\(^{-1}\) body weight of vitamin E; Groups V–VIII: Diabetic group that received 200 mg kg\(^{-1}\) body weight of the aqueous plant extracts of parsley, CS, GA and combination from the three aqueous extracts, respectively; Groups IX–XII: Diabetic group that received 200 mg kg\(^{-1}\) body weight of the biosynthesized AgNPs by parsley, CS, GA and combination from the three aqueous extracts, respectively. Data are presented as mean ± S.E from five animals in each group. Statistical significance was evaluated using the ANOVA test followed by post hoc Duncan’s multiple range test. A probability value of less than 0.05 was considered statistically significant (\( P < 0.05 \)).

2.8.2. Anti-inflammatory activity
The anti-inflammatory effect was determined through the induction of inflammation by sub plantar injection of carrageenan [35]. Fifty-five male albino rats, weighing (130–150 g), were divided into eleven groups, each group of five animals: Group I: Rats that received 1 ml of saline and served as the control; Group II: Received 1 ml AgNO\(_3\) and served as the positive control group; Groups III–IV: Rats that received 200 mg kg\(^{-1}\) body weight of the aqueous plant extracts of parsley, CS, GA and combination from the three aqueous extracts, respectively; Group VII–X: Rats that received 200 mg kg\(^{-1}\) body weight of the biosynthesized AgNPs by parsley, CS, GA and combination from the three aqueous extracts, respectively; Group XI: Rats that received 20 mg kg\(^{-1}\) of the reference drug indomethacin.

One hour later, all the animals received a sub plantar injection of 0.1 ml of 1% carrageenan suspension in saline in the right hind paw. The thickness of the right hind paw (mm) was measured immediately before and 1, 2, 3 and 4 h post carrageenan injection with a micrometer caliper. The results were expressed as mean S.E. (n = 5). The statistical comparison between the control group and the treated groups was carried out using two-way ANOVA followed by Duncan’s multiple range test. The results were statistically significant at \( P < 0.05 \).

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\% \text{Edema} = \frac{\text{Thickness of the right paw} - \text{Thickness of the left paw}}{\text{Thickness of the left paw}} \times 100
\]

2.9. Antimicrobial activity
2.9.1. Sensitivity of bacteria and yeast to the prepared extracts
The antimicrobial screening was done by the agar well diffusion method [36, 37]. Twenty milliliters (20 ml) of molten sterile Mueller–Hinton agar was poured into the sterile Petri dishes and allowed to solidify. Bacterial or fungal inoculum containing \( 5 \times 10^6 \text{CFU/mL} \) suspensions were uniformly streaked on the surface of the agar using sterile cotton swabs. The agar was then punched aseptically using a sterile tip to make a hole 5 mm in diameter. Twenty microliters of the different extracts were added to each well. Three replicates were performed for each treatment and incubated for 18–22 h at 37 °C for bacteria and for 24 h at 25 °C for yeasts. At the end of the incubation period, the inhibition zones diameters of all three replicates were measured in millimeters using a measuring slide and the mean of the three measurements was determined. The antimicrobial activity was indicated by a clear zone in the agar, which was measured after treatment. AgNO\(_3\) and was used as the control.

2.9.2. Determination of the minimum inhibitory concentration (MIC) of the prepared extracts by micro-dilution method
To characterize the antibacterial activity of the synthesized AgNPs and the aqueous plants’ extracts, broth microdilution method was performed according to the Clinical and Laboratory Standards Institute (CLSI) standards using four quality control strains (CLSI, 2015). The set contained two Gram-negative strains: E. coli and P. aeruginosa, as well as two Gram-positive strains: Staphylococcus saprophyticus and Enterococcus faecalis. An inoculate of the microbial strains was prepared from the 24 h broth cultures and the suspensions were adjusted.
TEM analysis showed that the mean size ranged between 8.16 ± 3.1 nm. Characterization of the synthesized AgNPs to 0.5 McFarland standard turbidity. Each sample (100 μg mL⁻¹) stock solution. A number of wells were reserved in each plate for the positive and negative controls. Sterile broth (100 μl) was added to the well from row B to H. The stock solutions of the samples (100 μl) were added to the wells in rows A and B. The mixture of the samples and sterile broth (100 μl) in row B was transferred to each well in order to obtain a twofold serial dilution of the stock samples. The inoculums (100 μl) were added to each well and a final volume of 200 μl was obtained in each well. Plates were incubated at 37 °C for 24 h. Experiments were repeated in duplicates and the results were determined as an average value. The MIC endpoint was considered as the lowest concentration of the extract or fraction inhibiting the total growth of the microorganisms. MIC was detected by the lack of visual turbidity (matching the negative growth control).

2.10. Metabolomic profiling of P. crispum, S. maydis and a combination of the three crude extracts
Metabolomic profiling was performed on the crude extracts of P. crispum, S. maydis and a combination of the three crude extracts to deliver general qualitative and quantitative profiles of metabolites that may be involved in the activity of the extracts [38]. Dereplication refers to the rapid identification of the known secondary metabolites and their quantification in crude unfractionated extracts [39, 40]. Identification of metabolome was done on an Acquity Liquid Chromatography (LC) system coupled to a Synapt G2 HDMS quadrupole time-of-flight hybrid mass spectrometer (Waters, Milford, USA). Chromatographic separation was carried out on a BEH C18 column (2.1 × 100 mm, 1.7 μm particle size; Waters, Milford, USA) with a guard column (2.1 × 5 mm, 1.7 μm particle size) and a linear binary solvent gradient of 0%–100% eluent B over 6 min at a flow rate of 0.3 ml min⁻¹, using 0.1% formic acid in water (v/v) as solvent A and acetonitrile as solvent B. The injection volume was 2 μl and the column temperature was 40 °C. To convert the raw data into separate positive and negative ionization files, Ms converter software was used. The files were then imported to the data mining software MZmine 2.10 for peak picking, deconvolution, deisotoping, alignment, and formula prediction. The database used for the identification of compounds was the Dictionary of Natural Products (DNP) 2015.

2.11. Molecular docking
Docking was done on the ovine COX-1 enzyme complexed with ibuprofen (PDB ID: 1EQG) and human COX-2 enzyme complexed with mefenamic acid (PDB ID: 5IKR) to suggest which compounds identified in LC/MS that might be responsible for the anti-inflammatory activity of the crude extracts. In all docking experiments, a grid box of dimensions 50 grid points and spacing 0.375 was centered on the given co-crystallized ligand. A total of 13 molecules were docked with four starting conformations. These conformers were generated for each ligand using OpenBabel [41] and the docking experiments were performed via Autodock4 [42] implementing 100 steps of genetic algorithm while keeping all the default setting provided by Autodock Tools. Visualization was done using Molecular Operating Environment MOE.

3. Results and discussion

3.1. Characterization of the synthesized AgNPs
3.1.1. Characterization using TEM
TEM analysis showed that the mean size ranged between 8.16–13.35 nm for AgNPs prepared by parsley aqueous extract, 5.28–9.39 nm for AgNPs prepared by CS extract and 6.56–21.47 nm for AgNPs prepared by GA aqueous extract (figure 1). TEM suggested that the morphology of the synthesized nanoparticles is spherical with a mean size in the nano range which indicated the synthesis of AgNPs.

3.1.2. Characterization of AgNPs using UV-Visible spectrophotometer
On the addition of different volumes of the aqueous extracts (1, 2, 3, 4 and 5 ml) to a constant concentration of AgNO₃ solution (10 ml of 1 mM solution), a preliminary visual observation showed that the initial color of the reaction mixture was faint, then changed from pale-yellow to light brown then to dark brown exponentially with reaction-time (figure 2). Changing the color of the reaction from light to dark brown with the reaction-time as aggregation proceeds confirmed the formation of silver nanoparticles [43]. The solutions were analyzed using UV spectra. AgNPs solutions prepared by parsley, CS and GA aqueous extracts showed absorbance peaks at 418, 419 and 422 nm, respectively (figure 3). The absorbance bands were observed at 410–450 nm, which were similar to those reported in the literature. The intensity of the absorption increased with increasing the concentration of different extracts. These changes in the absorbance confirmed the changes in the particles’ sizes during nanoparticles synthesis [44–46].
3.1.3. Characterization of AgNPs using FTIR

FTIR spectra (figure 4) showed major peaks at 3645.46 till 3275.13, 2368.59, 2306.86, 2083.12, 2075.41, 2067.69, 2052.26, 1917.24, 1913.39, 1643.35, 1392.61, 1284.59, 1230.58, 1199.72, 1153.43, 1099.43, 1060.85 and
The peaks in the range of 3200–3700 cm$^{-1}$ were assigned as stretching of N–H group and OH stretching in alcohols and phenolic compounds with strong hydrogen bonds. The peaks in the range 2368.59–2052.26 cm$^{-1}$ is relevant to the triple bond stretching. The peaks in the range of 1986.68–1913.39 cm$^{-1}$

Table 1. Antioxidant activity of the aqueous extracts of parsley, CS, GA, the combination of the three aqueous extracts, AgNPs synthesized by the three extracts and their combination in male albino rats ($n = 5$).

| Group                              | Blood glutathione (mg/dl) | % of change from the control |
|------------------------------------|----------------------------|-----------------------------|
| Control                            | 36.7 ± 1.3                 | —                           |
| Diabetic                           | 21.3 ± 0.4$^*$             | 41.96%                      |
| Diabetic + AgNO$_3$                | 30.4 ± 0.5                 | 17.16%                      |
| Diabetic + vitamin E               | 36.2 ± 1.1                 | 1.36%                       |
| Diabetic + Parsley aqueous extract | 34.1 ± 0.6                 | 7.08%                       |
| Diabetic + CS aqueous extract      | 32.9 ± 0.3                 | 10.35%                      |
| Diabetic + GA aqueous extract      | 33.6 ± 0.4                 | 8.44%                       |
| Diabetic + Combination of the three aqueous extracts | 35.1 ± 1.1 | 4.35% |
| Diabetic + Parsley AgNPs           | 34.5 ± 0.9                 | 5.99%                       |
| Diabetic + CS AgNPs                | 33.4 ± 0.7                 | 8.99%                       |
| Diabetic + GA AgNPs                | 34.1 ± 0.8                 | 7.08%                       |
| Diabetic + AgNPs prepared a combination of the three aqueous extracts | 35.9 ± 0.2 | 2.17% |

*Statistically significant different from control group at $P < 0.05$. 

Figure 4. FTIR spectra of $P$. crispum aqueous extract before (A) and after (B) synthesis of AgNPs, $S$. maydis aqueous extract before (C) and after (D) synthesis of AgNPs and GA aqueous extract before (E) and after (F) synthesis of AgNPs.
| Group                                | Zero   | 1 h         | 2 h         | 3 h         | 4 h         |
|-------------------------------------|--------|-------------|-------------|-------------|-------------|
|                                     | Paw diameter (mm) | Paw diameter (mm) | Percentage of edema | Paw diameter (mm) | Percentage of edema | Paw diameter (mm) | Percentage of edema | Paw diameter (mm) | Percentage of edema |
| Control                            | 3.49 ± 0.07  | 4.67 ± 0.1*  | 33.8        | 4.76 ± 0.2*  | 36.38       | 4.81 ± 0.1*  | 37.8        | 4.94 ± 0.1*  | 41.5        |
| AgNO₃                              | 3.41 ± 0.04  | 4.48 ± 0.1*  | 31.3        | 4.39 ± 0.1*  | 28.7        | 4.33 ± 0.6*  | 26.9        | 4.28 ± 0.01* | 25.5        |
| Parsley aqueous extract            | 3.45 ± 0.04  | 4.28 ± 0.07* | 24.05       | 4.21 ± 0.07* | 22.02       | 4.18 ± 0.06* | 21.15       | 4.14 ± 0.08* | 20         |
| Corn silk aqueous extract          | 3.42 ± 0.04  | 4.16 ± 0.04* | 21.6        | 4.14 ± 0.1*  | 21.05       | 4.11 ± 0.09* | 20.17       | 4.05 ± 0.08* | 18.4        |
| GA aqueous extract                 | 3.53 ± 0.06  | 4.44 ± 0.1*  | 25.8        | 4.41 ± 0.13* | 24.9        | 4.35 ± 0.01* | 22.9        | 4.31 ± 0.09* | 22.1        |
| Combination of the three aqueous extracts | 3.51 ± 0.05  | 4.23 ± 0.57* | 20.51       | 4.12 ± 0.1*  | 17.3        | 4.11 ± 0.08* | 17.1        | 4.05 ± 0.09* | 15.38       |
| Parsley AgNPs                      | 3.58 ± 0.07  | 4.22 ± 0.1*  | 17.8        | 4.14 ± 0.1*  | 15.6        | 4.06 ± 0.08* | 13.4        | 4.02 ± 0.07* | 12.3        |
| Corn silk AgNPs                    | 3.49 ± 0.06  | 4.08 ± 0.1*  | 16.9        | 4.02 ± 0.1*  | 15.1        | 3.95 ± 0.6*  | 13.2        | 3.9 ± 0.4*   | 11.7        |
| GA AgNPs                           | 3.47 ± 0.08  | 4.03 ± 0.09* | 16.1        | 3.98 ± 0.06* | 14.7        | 3.91 ± 0.01* | 12.7        | 3.86 ± 0.01* | 11.2        |
| AgNPs prepared by the combination of the three aqueous extracts | 3.54 ± 0.08  | 4.06 ± 0.1*  | 14.68       | 4.03 ± 0.1*  | 13.8        | 3.93 ± 0.08* | 11.01       | 3.87 ± 0.07* | 9.3         |
| Indomethacin                       | 3.48 ± 0.08  | 3.96 ± 0.09* | 13.79       | 3.90 ± 0.06* | 12.06       | 3.85 ± 0.01* | 10.6        | 3.76 ± 0.01* | 8.04        |

*Significantly different from zero time at $P < 0.05$. 

Table 2. Anti-inflammatory activity of the aqueous extracts of parsley, CS, GA, the combination of the three aqueous extracts, AgNPs synthesized by the three extracts and their combination in male albino rats (n = 5).
Table 3. Antimicrobial activity as indicated by the growth-inhibition zone (mm) of different extracts against different strains of fungi and bacteria against standard antimicrobial agents.

| Sample               | Microorganisms                  | AgNO₃ (C) | Parsley AgNPs | Parsley aqueous extract | GA AgNPs | GA aqueous extract | CS AgNPs | CS aqueous extract | Comb. aqueous extract | Comb. AgNPs | Standard antimicrobial agents |
|----------------------|---------------------------------|-----------|----------------|-------------------------|----------|---------------------|----------|---------------------|------------------------|-------------|-------------------------------|
| Gram Positive bacteria: |                                |           |                |                         |          |                     |          |                     |                        |             |                               |
| Staphylococcus saprophyticus |                                | NA        | 13             | NA                      | 18       | NA                  | 19       | 13                  | NA                     | 19          | 26 ± 0.11                      |
| Enterococcus faecalis |                                | NA        | 17             | 14                      | 19       | NA                  | 14       | 16                  | 13                     | 19          | 29 ± 0.22                      |
| Gram negative bacteria: |                                |           |                |                         |          |                     |          |                     |                        |             |                               |
| Escherichia coli     |                                | NA        | 18             | 13                      | 17       | 12                  | 18       | NA                  | 14                     | 22          | 27 ± 0.12                      |
| Pseudomonas aeruginosa |                                | NA        | 19             | 14                      | 14       | 14                  | 18       | NA                  | 13                     | 19          | 31 ± 0.89                      |
| Fungi:               |                                |           |                |                         |          |                     |          |                     |                        |             |                               |
| Candida albicans     |                                | NA        | 16             | NA                      | 19       | NA                  | 15       | NA                  | NA                     | 18          | 20.7 ± 0.22                     |
| Cryptococcus neoformans |                            | NA        | NA             | NA                      | 14       | NA                  | NA       | NA                  | NA                     | 17          | 21 ± 0.14                      |

Abbreviations: (C), Control; Comb., Combination formula; NA, No activity.
indicated C-H bending of aromatic compounds. The peaks in the range of 1643.35 cm$^{-1}$ indicated C=O bond of the carbonyl group, C=C bond stretching and the stretching vibrations of amides. The peak in the range 1392.61 cm$^{-1}$ indicated the presence of tertiary amide, C-N stretching and N-H bending. Peaks appeared in the range 1284.59-1060.85 cm$^{-1}$ is relevant to C-O stretching, while the peaks in the range 509.21-408.91 cm$^{-1}$ suggested alkyl halides bond stretching.

There are many functional groups present in the aqueous extract of the tested plants which may have been responsible for the reduction of Ag$^+$ ions. FTIR spectrum illustrated different peaks positions which were attributed to flavonoids, glycosides, terpenoids, tannins, and phenolics. The presence of such functional groups increased the stability of the synthesized nanoparticles by preventing the clotting and aggregation of the nanoparticles. The similarities between the spectra, before and after synthesis of nanoparticles, with some marginal shifts in peak positions indicated the presence of residuals from the different plant extracts in the sample which acted as a capping agent to the synthesized AgNPs$^{44, 47}$. Therefore, it can be concluded that these secondary metabolites are responsible for the effective stabilization and capping of the synthesized nanoparticles.

3.2. Antioxidant activity

The assessment of GSH level showed that AgNPs prepared by the combination of the three extracts showed the highest antioxidant activity with the change percentage (2.17%), followed by solution of the combined three aqueous with the change percentage (4.35%) and AgNPs prepared by aqueous extract of parsley with the change percentage (5.99%), while the CS aqueous extract showed the least antioxidant activity with change percentage of 10.35% in comparison with Vitamin E as the standard drug with change percentage (1.36%) shown in (table 1). From this table, we can conclude that the antioxidant activity of AgNPs synthesized by the three different aqueous extracts has the highest blood GSH, suggesting the possible synergistic activity of the AgNPs synthesized by the three aqueous plant extracts.

3.3. Anti-inflammatory activity

The anti-inflammatory activity of AgNPs prepared by the combination of the three aqueous extracts was the highest as demonstrated by a time-dependent reduction in edema. The edema was reduced by 14.7%, 13.8%, 11.01%, 9.3% after 1, 2, 3 and 4 h, respectively. GA AgNPs was also active in reducing edema showing 16.1%, 14.7%, 12.7%, and 11.2%, respectively. The GA aqueous extract showed the least anti-inflammatory activity after 1, 2, 3 and 4 h with the following reduction percentages in edema 25.8%, 24.9%, 22.9%, and 22.1%, respectively (table 2). AgNPs synthesized by the three different aqueous extracts showed potent anti-inflammatory activity after four hours resulting in edema reduction suggesting the optimum use of this combination formula as an anti-inflammatory agent. This effect was further confirmed by the aid of the docking technique on two of the well-known enzymes involved in the inflammatory cascade, COX-1, and COX-2.
Table 4. MIC (mg/ml) values of the samples against the tested microorganisms were performed for the most active samples.

| Sample Microorganisms | Parsley AgNPs | Parsley aqueous extract | GA AgNPs | GA aqueous extract | CS AgNPs | CS aqueous extract | Comb. aqueous extract | Comb. AgNPs |
|-----------------------|---------------|------------------------|---------|-------------------|---------|-------------------|-----------------------|------------|
| Gram Positive bacteria: |               |                        |         |                   |         |                   |                       |            |
| Staphylococcus saprophyticus | ND           | —                      | 25      | —                 | ND      | —                 | 100                   |            |
| Enterococcus faecalis | 50            | ND                     | 25      | —                 | 25      | ND                 | ND                    | 25         |
| Gram negative bacteria: |               |                        |         |                   |         |                   |                       |            |
| Escherichia coli      | 50            | 100                    | 50      | 100               | 50      | —                 | ND                    | ND         |
| Pseudomonas aeruginosa| 50            | ND                     | 50      | ND                | 100     | —                 | ND                    | 50         |
| Fungi:                |               |                        |         |                   |         |                   |                       |            |
| Candida albicans      | ND            | —                      | ND      | —                 | 50      | —                 | 50                    |            |
| Cryptococcus neoformans| —            | —                      | 50      | —                 | —       | —                 | 100                   |            |

**Abbreviations:** Comb., combination; ND, not detected.
3.4. Antimicrobial activity

3.4.1. Sensitivity of bacteria to the prepared extracts and AgNPs

The antibacterial activity was studied against *S. saprophyticus*, *E. faecalis*, *E. coli* and *P. aeruginosa*. The results indicated that some extracts were active against Gram-positive and Gram-negative bacteria involved in the study (table 3 and figure 5). In case of *S. saprophyticus*, both CS AgNPs and AgNPs prepared by the combination of the three aqueous extracts showed the most potent activity with inhibition zones of 19 mm for both, followed by GA AgNPs with inhibition zone of 18 mm, while all aqueous extracts, except of CS, did not show any activity against the tested microorganisms. For *E. faecalis*, the highest antibacterial activity was seen with GA AgNPs and AgNPs prepared by the combination of the three aqueous extracts with inhibition zones of 19 mm for both, while the

### Table 5. Dereplication of the metabolomics of the crude extracts of *P. crispum*, *S. maydis* and the combined extract formulated from *P. crispum* and *S. maydis* and GA aqueous extracts assembled according to their molecular weight.

| m/z       | Rt. (min.) | M.wt.   | Name                                         | Source                | Molecular formula | References |
|-----------|------------|---------|----------------------------------------------|-----------------------|-------------------|------------|
| 195.15    | 3.1        | 194.1423587 | Ferulic acid                                  | Corn silk             | C_{10}H_{10}O_{4} | [48]       |
| 271.061   | 3.7        | 270.0333351 | Imperatorin                                    | Parsley               | C_{14}H_{14}O_{4} | [49]       |
| 277.148   | 1.8        | 276.1405506 | Muurolene dihydrochloride                     | Corn silk             | C_{15}H_{26}Cl_{2} | [50]       |
| 295.227   | 4.8        | 294.2200031 | 13-Hydroxy-10-oxo-11-octadecenoic acid; (±)-(E)-form, Lactone | Corn silk             | C_{19}H_{30}O_{3} | [51]       |
| 326.187   | 4.8        | 327.1942075 | Sophazrine                                    | Corn silk             | C_{19}H_{25}N_{3}O_{2} | [52]       |
| 329.233   | 3.9        | 330.2402922 | 9,18-Dihydroxy-10-octadecenoic acid; (9E,10E)-form | Combination           | C_{16}H_{34}O_{5} | [53]       |
| 329.233   | 3.9        | 330.2402922 | 5,8,12-Trihydroxy-9-octadecenoic acid         | Combination           | C_{16}H_{34}O_{5} | [54]       |
| 338.212   | 6.4        | 337.2049716 | Holadienine; 18-Oxo, 14,15-didehydro           | Parsley               | C_{22}H_{27}N_{3}O_{2} | [55]       |
| 340.202   | 5.1        | 341.2094539 | Iboluteine; Oxime                             | Combination           | C_{22}H_{27}N_{3}O_{2} | [56, 57]  |
| 356.259   | 4.3        | 355.2514057 | 20-Amino-18-hydroxypregna-1,4-dien-3-one; (20 S,14R)-form, 18-Aldehyde, N,N-di-Me | Combination           | C_{22}H_{27}N_{3}O_{2} | [58]       |
| 431.134   | 3.8        | 430.1267871 | Alternanthin                                   | Corn silk             | C_{22}H_{27}O_{4} | [59]       |
| 441.322   | 6.4        | 440.3139993 | 13-Hydroxy-10-oxo-11-octadecenoic acid; (±)-(E)-form, Lactone | Corn silk             | C_{22}H_{27}O_{4} | [51]       |
| 469.134   | 2.8        | 470.141348  | 4-Hydroxy-5-methyl-2H-1-benzopyran-2-one; O-[β-D-Apiofuranosyl-(1 → 6)-β-D-glucopyranoside] | Corn silk             | C_{21}H_{26}O_{12} | [61]       |

*Abbreviations: Rt, Retention time (min.); M.wt., Molecular weight.

### Table 6. The docking scores of the top 10 ligands in Kcal/mol, as generated by Autodock4.

| Ligand                             | COX-1  | COX-2  |
|------------------------------------|--------|--------|
| Cocrystallized ibuprofen           | −7.1   | −7.4   |
| Cocrystallized mefenamic acid      | −11.1  | −7.1   |
| 20-Amino-18-hydroxypregna-1,4-dien-3-one; (20 S)-form, 18-Aldehyde, N,N-di-Me | −10.7  | −8.7   |
| Holadienine; 18-Oxo, 14,15-didehydro | −10.6  | −8.3   |
| 8,13-Epoxy-14,15,16,19-labdanetetrol; (ent-8α,13 R,14 S)-form, 19-(3-Methylbutanoyl) | −10.1  | −7.2   |
| Iboluteine; Oxime                  | −10.1  | −7.2   |
| 4-Hydroxy-5-methyl-2H-1-benzopyran-2-one; O-[β-D-Apiofuranosyl-(1 → 6)-β-D-glucopyranoside] | −9.9   | −10.3  |
| Imperatorin                         | −9.7   | −9.3   |
| 13-Hydroxy-10-oxo-11-octadecenoic acid; (±)-(E)-form, Lactone | −9.6   | −9.0   |
| Sophazrine                          | −9.4   | −6.3   |
| Muurolene dihydrochloride          | −9.4   | −8.3   |
| Alternanthin                        | −9.4   | −7.8   |

*Results are assembled according to their docking scores.

3.4. Antimicrobial activity

3.4.1. Sensitivity of bacteria to the prepared extracts and AgNPs

The antibacterial activity was studied against *S. saprophyticus*, *E. faecalis*, *E. coli* and *P. aeruginosa*. The results indicated that some extracts were active against Gram-positive and Gram-negative bacteria involved in the study (table 3 and figure 5). In case of *S. saprophyticus*, both CS AgNPs and AgNPs prepared by the combination of the three aqueous extracts showed the most potent activity with inhibition zones of 19 mm for both, followed by GA AgNPs with inhibition zone of 18 mm, while all aqueous extracts, except of CS, did not show any activity against the tested microorganisms. For *E. faecalis*, the highest antibacterial activity was seen with GA AgNPs and AgNPs prepared by the combination of the three aqueous extracts with inhibition zones of 19 mm for both, while the
least antibacterial activity was seen with the combination of the three aqueous extracts with inhibition zone of 13 mm and no activity was seen with GA aqueous extract. AgNPs prepared by the combination of the three aqueous extracts showed the highest inhibitory activity against *E. coli* with inhibition zone of 22 mm followed by parsley AgNPs and CS AgNPs with inhibition zones of 18 mm for both, while the least inhibitory activity was seen with GA aqueous extract with inhibition zone of 12 mm and no activity was detected with CS aqueous extract. For *P. aeruginosa*, AgNPs prepared by the combination of the three aqueous extracts, parsley AgNPs and CS AgNPs showed the highest inhibitory activity with inhibition zones of 19, 19 and 18 mm respectively, while the combination of the three aqueous extracts showed the least inhibitory activity with inhibition zone of 13 mm with no activity was detected for CS aqueous extract.

3.4.2. Sensitivity of yeast to the prepared extracts and AgNPs

The antifungal activity was studied against *C. albicans* and *C. neoformans* (table 3 and figure 5). In case of *C. albicans*, all the synthesized AgNPs showed inhibitory activity with GA AgNPs and AgNPs prepared by the combination of the three aqueous extracts showing the highest activity with inhibition zones of 19 and 18 mm, respectively. All aqueous extracts did not show any activity. For *C. neoformans*, only the AgNPs prepared by the combination of the three aqueous extracts and GA AgNPs showed inhibitory activity without any activity from the aqueous extracts with inhibition zones of 19 and 14 mm, respectively.

From table 3 and figure 5, we can conclude that AgNPs biosynthesized by the three different aqueous extracts showed a broad-spectrum activity against both bacteria and fungi involved in the study, followed by GA AgNPs which also showed inhibitory activity against both bacteria and fungi. These results suggested the possible use of the combination formula as a broad-spectrum antimicrobial agent.

3.4.3. Minimum inhibitory concentration (MIC) of the prepared extracts by microdilution method

The microdilution method, standardized by the Clinical and Laboratory Standards Institute (formerly NCCLS) allowed the determination of the minimal inhibitory concentrations (MICs) of the prepared extracts alone. MIC was determined for some extracts that showed obvious inhibitory activity against the tested microorganisms (table 4).

3.5. Metabolomic profiling of the aqueous extracts of parsley, CS and combined extract of parsley, CS and GA

Dereplication of the secondary metabolites of the aqueous extracts of parsley, CS and the combination formulated from parsley, CS and GA aqueous extracts resulted in the identification of different classes of
compounds. The positive mode showed the majority of the identified compounds and revealed the presence of a variety of alkaloids, terpenes, flavonoids, furanocoumarins, polyphenolic and steroidal compounds (table 5). These identified compounds are responsible for the capping and stabilization of the synthesized AgNPs and might be responsible for the synergistic activity seen in the combination formula.

3.6. Molecular docking
Docking was performed in an attempt to rationalize the observed anti-inflammatory activity of the crude extracts. The top-scoring 10 ligands (according to docking in COX-1) are listed with their docking scores in (table 6). Most of the compounds identified in the LC/MS spectra showed better or similar scores to the co-crystallized ligands. The scores were significantly better in case of COX-1. According to the scores, the ligands were expected to have more affinity towards COX-1 than COX-2, with the exception of 4-hydroxy-5-methyl-2H-1-benzopyran-2-one; O-\([\beta-D-apiofuranosyl-(1 \rightarrow 6)]-\beta-D-glucopyranoside\]. The interactions between the two high-scoring compounds and the surrounding amino acids in the COX-1 active sites are shown in figure 6. Both ligands are expected to form H-bonds with Arg120 in addition to one extra H-bond. The differences between COX-1 and COX-2 active sites are well-studied in literature [62]. It is now common knowledge that COX-2 has an extra binding site that is not present in COX-1. Interactions with Arg513 in COX-2 are also thought to be essential for the selective inhibition of COX-2.41 Interestingly, some of our studied compounds showed excellent binding which was very similar to the known selective COX-2 inhibitors. Figure 7 illustrates how 4-hydroxy-5-methyl-2H-1-benzopyran-2-one; O-\([\beta-D-apiofuranosyl-(1 \rightarrow 6)]-\beta-D-glucopyranoside\] assumes a binding conformer that is very similar to that of celecoxib (from the crystal structure 3LN1). As clearly illustrated in the 2D diagram, 4-hydroxy-5-methyl-2H-1-benzopyran-2-one; O-\([\beta-D-apiofuranosyl-(1 \rightarrow 6)]-\beta-D-glucopyranoside\] makes many H-bonds including the crucial one with Arg513.

4. Conclusion
The synthesis of AgNPs from aqueous extracts of parsley, CS and GA and combination thereof provided nanoparticles with potent antioxidant, anti-inflammatory, and antimicrobial activity. AgNPs formed by the combined three plants aqueous extracts showed the most potent activity. LC/MS served to identify various compounds that were responsible for the capping and stabilization of the synthesized AgNPs. The docking study pointed out the justification of the anti-inflammatory activity of various extracts in terms of binding to COX-1 and COX-2 enzymes. These findings presented the AgNPs synthesized by the combined extracts of parsley, CS and GA as effective formula for reversing oxidative stress, treatment of various inflammatory conditions and as antimicrobial agents against pathogenic bacteria and fungi causing serious urinary tract infections.

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Disclosure

The author reports no conflicts of interest in this work.

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