

Biosynthesis and characterization of silver nanoparticles produced by plant extracts and its antimicrobial activity

Abstract

Solanum tuberosum is the fourth most challenging plant in Egypt, affected by several fungi, viral and bacterial diseases. Bacterial and fungal isolates (Brown rot disease (*Ralstonia solaniserum*), soft root disease (*Pectobacterium carotovora*) and dry rot disease (*Fusarium oxisporum*)) were collected. The green extracts of silver nanoparticles were prepared by means of aqueous extracts of three wild plants, *Physalis peruviana* (leaves, red and green fruits) (N1, N2 and N3), *Solanum nigrum* (fruit) (N4) and *Moringa oliefera* (leaves) (N5). SEM, TEM, FT-IR and X-RD obtained the characterization of the biosynthesis of silver nanoparticles. The results indicated that nanoparticles were spherical, smooth and the sizes varied between 12 and 33 nm. The activity of the nanoparticle formulations was tested against the bacterial isolates using agar diffusion method and one fungus using mycelial growth method. The results also elucidated that N5 formulation showed a significantly potent antibacterial activity against *R. solanacearum*. However, N1 formulation was the highest active one against *P. carotovra*. In addition, the antifungal activity indicated that N1 had the highest effect ($EC_{50} = 687.03$ mg/L) followed by N3 ($EC_{50} = 981.61$ mg/L) against *F. oxysporum*. Nanoparticles synthesized by wild plants could be used as safe alternatives to harmful microbicides.

Keywords: Biosynthesis, Silver nanoparticles, *Physalis peruviana*, *Solanum nigrum*, *Moringa oliefera*, Plant extract, Antifungal, Antibacterial, SEM, TEM, FT-IR, XRD.

1. Introduction

Solanum tuberosum (family Solanaceae) is a worldwide-cultivated tuber-bearing plant, which is the fourth main food crop in the world after rice (*Oryza sativa*), maize (*Zea mays*) and wheat (*Triticum aestivum*), in terms of both area cultivated and total production (Douches et al., 1996; Czajkowski et al., 2011). Potato does not need exceptional growth circumstances; it has been for a long time a most important field crop in temperate regions, and progressively in warmer areas (Haverkort, 1990). It is presently the second greatest significant vegetable crop after tomatoes in Egypt, and Egypt is one of Africa's prime potato producers and exporters. Potato is vulnerable to a numeral of diseases, enclosing late blight triggered by *Phytophthora infestans*, numerous viruses and bacterial wilt caused by *Ralstonia solanacearum*. Bacteria and fungi are played a primary character in the harvest losses, particularly *Erwinia* the causal agent

36 of soft rot in potato (Rashid et al. 2012) and *Alternaria spp* the causal agent of early blight of
37 potato (Belosokhov et al., 2017). *Ralstonia solanacearum*, the causative agent of bacterial wilt
38 in potatoes, is soilborne and can persist in soil for a long time in infected host plant debris or by
39 colonizing potato volunteer plants, alternative hosts or even non-host plants (Aliye et al., 2008).
40 To infect a plant effectively, the pathogen first has to be capable to penetrate and colonize host
41 tissues and overcome active plant defense responses to encourage the set of actions finally that
42 leads to disease symptoms. Furthermore, *Pectobacterium carotovra* is a gram-negative
43 phytopathogenic bacterium, which attack carrots, cucumber, onions, potatoes and tomatoes. It
44 produced black leg (soft rot) to these plants through farming, transportation and storage (Leite et
45 al., 2014). *Pectobacterium* produced damage of the cell wall of the plants then cause death of the
46 plants. *Fusarium* wilt diseases are accountable for imperative harvest damages on several crops.
47 *Fusarium oxysporum* causes dry rot, stem-end rot and wilt of potatoes. *Fusarium* dry rot is
48 mainly a post-harvest disease and can turn into a foremost problem when infected stored
49 potatoes. Chemical control of potato brown rot with currently existing crop protectants is not
50 effective (Lopez and Biosca, 2004). Improvement of additional effective chemical control
51 techniques is not fortified owing to the universal awareness about adverse impacts of synthetic
52 crop protectants on human health and the environment; this has led to the phasing out of an
53 increasing quantity of crop protectants. Consequently, there is a perfect necessity to improve
54 alternative practical, harmless and effective managing approaches that can condense the time that
55 no host plants can be grown. Plant extracts of many higher plants have been described to display
56 antibacterial and antifungal properties under laboratory trails (Okigbo and Ogbonnaya, 2006;
57 Shariff et al., 2006). Plant metabolites and plant-based pesticides seem to be one of the
58 improved alternatives, as they are known to have minimal environmental impact and hazard to
59 consumers in contrast to the synthetic pesticides (Varma and Dubey, 1999). Nanotechnology
60 has been used widespread in plant pathogens and the application of nanoparticles become
61 essential in the managing of plant diseases (Sastry et al., 2010). Silver nanoparticles exhausting
62 plant extracts are a significant distinction chemical and biosynthetic using gold, platinum and
63 silver in the synthesized of nanoparticles (Patil and Hooli, 2013). Therefore, the current study
64 goals to synthesize silver nanoparticles by a green biological route, using an extract derived from
65 *Physalis peruviana* (leaves, red and green fruits), *Solanum nigrum* (fruit) and *Moringa oliefera*
66 (leaves). Characterization of the synthesized nanoparticles achieved using scanning electron
67 microscope (SEM), transmission electron microscope (TEM), X-ray diffraction (XRD) and
68 Fourier transform infrared spectroscopy (FT-IR) analysis. Besides, their antimicrobial activity
69 against representatives of plant pathogenic bacteria (*Pectobacterium carotovra* and *Ralstonia*
70 *solaniserum*) and fungus (*Fusarium oxysporium*) was investigated.

71

72 2. Materials and methods

73 2.1. Cultures and growth conditions

74 The potato plants were grown at two localities in Abo-Homous and Borg-Elarb, El-Behera
75 and Alexandria Governorates, **respectively, Egypt** during the growing **season 2016**. **The bacteria**
76 were isolated from infected potato tubers and purified on Luria Bertani medium (LB) (**Maniatis**
77 **et al. 1982**), and incubated for 24 hours at 30°C. In addition, the fungi was grown on Potato
78 dextrose agar (PDA) and Kelman's TZC media (**Kelman, 1954**), then incubated at 28°C for 7
79 days. The microbes (bacteria and fungi) were identified using different methods including
80 microscopically extension and molecular identification.

81 **2.2. Pathogenicity test**

82 According to (**Zhang et al., 2014**) with some modification, healthy potato tubers selected
83 and washed carefully in water. **Tubers dipped in ethanol 70% for 5 min and washed** in distilled
84 water. Sterilized tuber was inoculated by syringe in plates containing a piece of sterile cotton
85 saturated with water. The suspension concentration of bacteria and fungi were 10⁸ and 10⁶
86 CFU/mL, respectively. Control tubers were inoculated by distilled water and incubated at the
87 same conditions.

88 **2.3. Preparation of the plant extracts**

89 Three medicinal plants, *P. peruviana* (leaves, red and green fruits), *S. nigrum* (fruit) and
90 *M. oliefera* (leaves) were selected from Abo-Homous and Borg-Elarb, El-Behera and Alexandria
91 Governorates, **respectively, Egypt**. Fresh and healthy leaves and fruits were collected locally and
92 rinsed thoroughly **first with tap water followed by** distilled water to remove all the dust and
93 unwanted visible particles, cut into small pieces and dried at room temperature. About 10 g of
94 **these** finely incised leaves of each **plant type** were weighed **separately**, 100 mL distilled water
95 was added and boiled for about 20 min. The extracts were then filtered thrice to get clear
96 solutions, which were **then, refrigerated** (4°C) for further experiments (**Banerjee et al., 2014**).

97 **2.4. Green synthesis of silver nanoparticles formulations**

98 Plant extract was added to aqueous solution (10 mM) of silver nitrate (AgNO₃) in dark flask
99 with shaking at 250 rpm and the **changes in the color was** observed. The reduction of Ag solution
100 was subjected to UV- Visible spectrophotometer at 540 nm (Beckman, model Du 540), and the
101 reaction stopped when the value of optical density was decreased. The solution was centrifuged
102 at 12000 rpm for 30 min, the supernatant **was discard** and the **pellet washed 3 times** by sterile
103 water. The pellet was dried at 50°C and then dissolved in sterile water (**Banerjee et al., 2014**).

104 **2.5. Characterization of silver nanoparticles formulations**

105 **2.5.1. Scanning electron microscopy (SEM)**

106 Scanning electron microscopy (SEM) is a method for high-resolution imaging of surfaces.
107 SEM analysis was done by using a JEOL JSM-5410 (Japan) electron microscope with a W-

108 source and operating at 80 kV. Sample was prepared on a glass slide (1 × 1 cm) after washing it
109 with ethanol. A tiny drop of nanoparticles was spreaded evenly over glass slide and allowed to
110 air dry. In order to make it conductive, gold coating with Jeol Quick Auto Coater was performed
111 (JFC-1500). The NPs were then subjected to SEM analysis under ambient conditions.

112 **2.5.2. Transmission electron microscopy (TEM)**

113 Morphology of the nanoparticles usually determined by transmission electron microscopy
114 (TEM). A combination of bright-field imaging at increasing magnification and of diffraction
115 modes use to reveal the form and size of the nanoparticles. To perform the TEM observations,
116 the nanoparticles formulation dilute with water (1/100). A drop of the diluted nanoparticles
117 directly deposited on the film grid and observed after dry.

118 **2.5.3. Fourier transform infrared spectroscopy (FT-IR)**

119 FTIR spectra of nanoparticles were taken with potassium bromide pellets on a Thermo
120 Nicolet AVATAR 300 FTIR spectrometer in the range 400-4000 Cm^{-1} .

121 **2.5.4. X-ray diffraction analysis (XRD)**

122 X-ray powder diffraction patterns of nanoparticles were obtained by a D/max-ra
123 diffractometer. The X-ray source was CuK radiation (40 kV, 80 mA). Samples were scanned
124 at a scanning rate of 4°/min.

125 **2.6. Assessment of antimicrobial assay**

126 **2.6.1. Antibacterial activity of nanoparticles formulations**

127 The antibacterial activity of the nanoparticles was evaluated against *P. carotovra* and *R.*
128 *solaniserum* by the agar diffusion method with LB agar media. A 20 mL of LB agar media was
129 poured into sterilized petri dishes and the plates were leaved for solidification then bacterial
130 suspension of the two tested bacteria was streaked. The paper discs of 6 mm size were saturated
131 with 20 μL of silver nanoparticles solutions (100, 200, 400 and 600 mg/L) or Doxycycline (30
132 μg) as standard antibacterial agent and plated on the surface of each plates at equivalent distance
133 with control. Bacteria was stand by 30 min, then incubated at 30° C for 24 h and the formed
134 inhibition zone was measured and three replicates were used (Abbassy et al., 2016).

135 **2.6.2. Antifungal activity of nanoparticles formulations**

136 The antifungal activity was tested using mycelia radial growth technique (Badawy et al.,
137 2014). The compounds were dissolved and serial concentrations ranged from 1000 to 3000 mg/L
138 were tested. Standard fungicide, gold plus was used at 0.25, 0.5 and 1.0 fold of field application
139 (200 g/100 L). The aliquots (quantity???) of the stock solutions were added to the PDA medium,
140 and then transferred to Petri dishes. After solidification, the mixtures were inoculated with a 5
141 mm in diameter mycelium fungi at the center of Petri dishes and these were incubated in the dark
142 at 27 ± 2°C. Fungal growth was measured when the control had grown to the edge of the plate.
143 The inhibition of fungal growth was calculated as the percentage of inhibition of radial growth

144 compared to the control. The effective concentration that inhibits 50% of mycelial growth (EC₅₀)
145 for each compound was estimated by probit analysis (**Finney 1971**) using SPSS 21.0 software.

146 **2.7. Molecular identification of obtained isolates using specific PCR, sequencing and** 147 **phylogenetic analysis**

148 DNA was isolated from the two bacterial isolates and the fungus isolate using QIAgene
149 DNA extraction kit according to the manufacture procedures (QIAgene, Germany). PCR
150 amplification for the bacteria was performed using the 16S rRNA primers (forward;
151 AGAGTTTGATCCTGGCTCAG and reverse; AAGGAGGTGATGCAGCC) according to
152 (**weisburg et al., 1991**). On the other hand, the fungus DNA was subjected to PCR amplification
153 using ITS specific primers (ITS1; TCCGTAGGTGAACCTGCGG and ITS4;
154 TCCTCCGCTTATTGATATG) according to (**White et al., 1990**). The 25 µL PCR **reaction**
155 **components were**; 12.5 µL master mix (Applied Biotechnology, Egypt), 1 µL DNA (30 ng),
156 1 µL for each primer (10 p mol/µL) and the volume **completed up to 25 µL** with sterile H₂O.
157 The PCR program was applied as follow; initial denaturation at 95°C for 2 min; 34 cycles of
158 94°C for 1 min; annealing at 55°C for 1 min; extension at 72°C for 1 min and a final extension
159 step at 72°C for 5 min; A 5 µL of PCR products were separated on 2% (w/v) agarose gel
160 electrophoresis in 0.5x TBE buffer. The molecular weight of band was estimated using DNA
161 marker (**marker size???**). Finally, the gel was photographed using gel documentation system.
162 PCR products were purified using PCR clean up column kit (Maxim biotech INC, USA). The
163 purified PCR products were subjected to DNA sequencing using the forward primer of 16S
164 rRNA and ITS (**Sigma company, Korea**). The DNA nucleotide sequences **were alignment using**
165 **BLASTn** (<http://www.ncbi.nlm.gov/BLAST>) and then the **clean sequences was submitted to**
166 **Gene Bank**. **Phylogenetic tree** was constructed using Mega 4 program, to examine the origin of
167 the obtained microbial strains (**Tamura et al., 2007**).

168 **2.8. Statistical analysis**

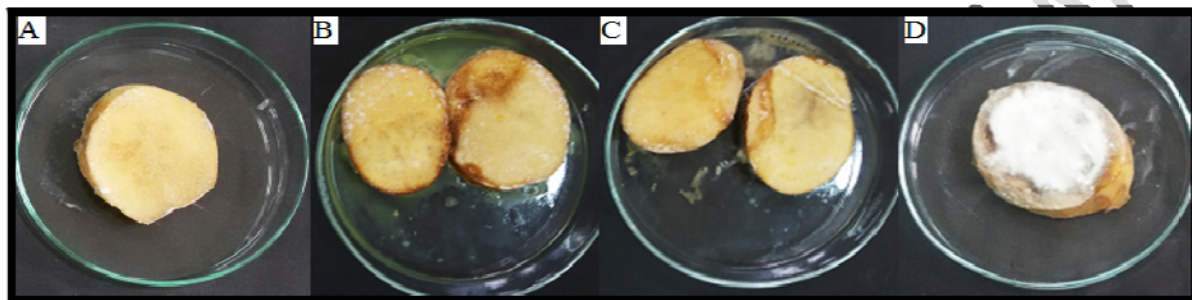
169 Statistical analysis was performed using SPSS 21.0 software (Statistical Package for Social
170 Sciences, USA). **All experiments** were repeated at least 3 times. The data were expressed as the
171 mean ± standard error (SE). **The log dose-response curves allowed determination of the EC₅₀**
172 **values for the fungal bioassay according to the probit analysis (Finney 1971)**. The 95%
173 confidence limits for the range of EC₅₀ values were determined by the least-square regression
174 analysis of the relative growth rate (% control) against the logarithm of the compound
175 concentration.

176 **3. Results and Discussion**

177 **3.1. Pathogenicity test**

178 Many different bacteria and fungi were successfully isolated from collected potato tuber,
179 they include; *P. carotovra*, *R. solaniserum* and *F. oxysporum*, all of which were implicated as
180 pathogens when tested on healthy tubers. The bacterial isolate showed high capability for
181 infection the healthy potato tubers. *P. carotovra* causes soft rot disease symptoms in the

182 inoculated healthy potato tubers after 4-5 days post inoculation. The appeared symptoms were;
183 chlorosis, wilting, tuber rot, blackleg and haulm desiccation. These results are in agreement with
184 those obtained by **Motyka et al., (2017) and Onkendi and Moleleki, (2014)**. While, healthy
185 potato tuber inoculated with *R. solanacearum* was showed the wilt disease symptoms; vascular
186 browning, dark brown streaks and grey-white bacterial ooze was observed on tuber surfaces.
187 Moreover, the *F. oxysporium* was isolated and used in inoculation of the healthy tubers and it
188 was observed that the isolate succeeded to cause dry rot disease for the tubers after 7 days. The
189 observed symptoms were; dry rot, sunken, wrinkled and a white mold was visible on tuber
190 surfaces (**Fig. 1**).

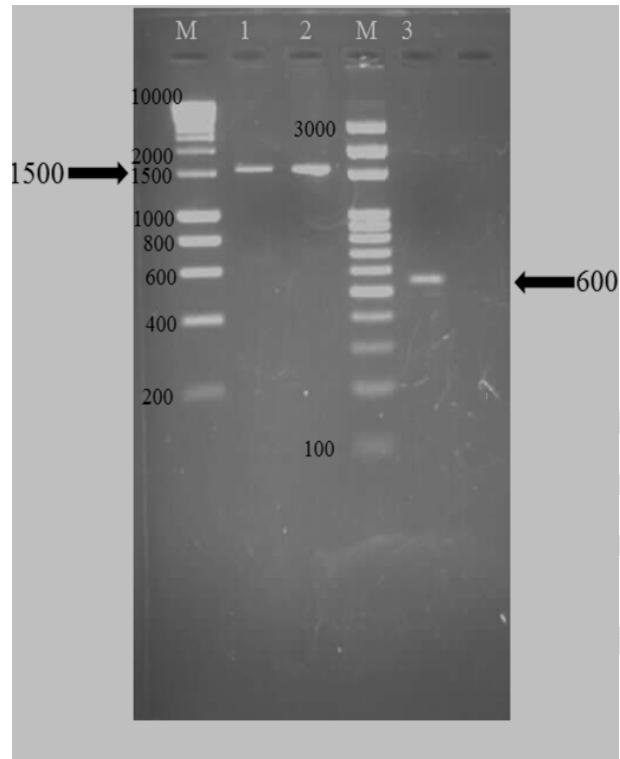


191
192 **Fig. 1:** Pathogenicity test of the potato tubers. Tuber control (A); tuber infected with *P. carotovra*
193 (B); tuber infected with *R. solanacearum* (C) and tuber infected with *F. oxysporium* (D).

194 **3.2. Molecular identification of the obtained isolates**

195 Approximately 1500 bp region of the 16SrRNA gene was amplified for *P. carotovra* and *R.*
196 *solanacearum*, while, PCR product of ITS gene amplified 550 bp for *F. oxysporium* (**Fig. 2**)
197 using universal primers. The DNA sequence results revealed that the examined two bacteria and
198 one isolate of fungi. The phylogenetic tree constructed based on the obtained DNA sequence
199 revealed that *P. carotovra* contained of two cluster; cluster one was divided into two sub cluster,
200 sub cluster one was divided into two group, group one divided to two sub group which contain *P.*
201 *carotovra* that similar with investigated isolate with different percentage. *R. solanacearum* had
202 phylogenetic tree contained two cluster; cluster one contain *R. solanacearum* isolate whereas
203 cluster two consist two sub cluster that divided into two group which divided into two sub group
204 that contain different strains of *R. solanacearum*. The phylogenetic tree of *F. oxysporium*
205 contains two cluster; cluster one divided to two sub cluster, sub cluster one contain to two group
206 that divided into two sub group which contain strains of *F. oxysporium*. While cluster two
207 contain two sub cluster, cluster two divided into two group, group two contain two sub group
208 while sub group two contain detected isolate of Fusarium as shown in **Fig. (3)**.

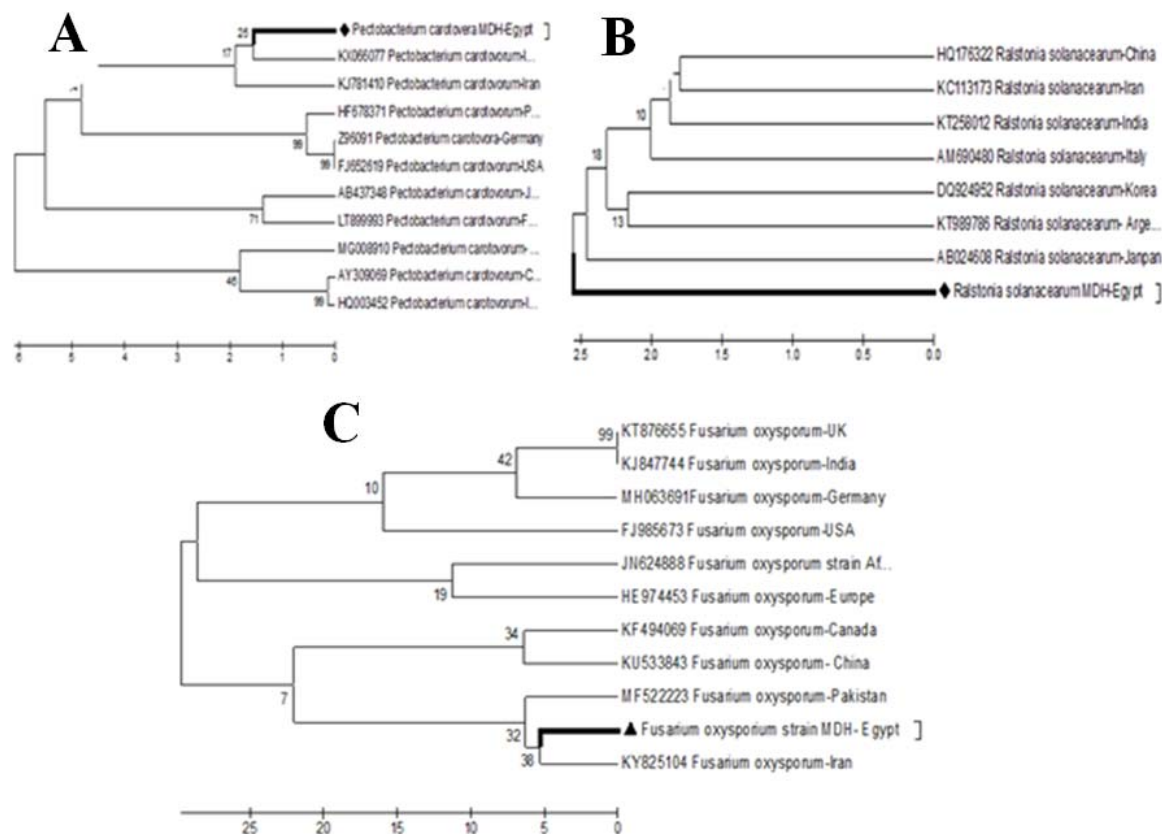
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210

211 **Fig. 2:** PCR products of 16s RNA gene for both two bacterial isolates and iTs gene of fungi,
 212 respectively. M, 10000 Kbp DNA marker; Lane 1, *P. carotovra*; Lane 2, *R. solanacearum*; M,
 213 3000 Kbp DNA marker; *F. oxysporium*

214



215
 216 **Fig. 3:** Phylogenetic tree of 16s RNA and ITS genes: *P. carotovora* (A); *R. solanacearum* (B) and
 217 *F. oxysporium* (C). Based on the DNA nucleotide sequencing and comparing with the other
 218 species listed in the Gene Bank.

219 3.3. Green synthesis of silver nanoparticles using plant extracts

220 The synthesized nanoparticles using the five different aqueous plant extracts; *M. oleifera*
 221 (leaves) *S. nigrum* (fruits) and *P. peruviana* (leaves, red and green fruits) were obtained after
 222 incubation period lasts for 24h. It was observed that the solution color changed from yellow to
 223 dark brown within the first 10 hrs. Silver nanoparticles exhibit yellowish brown color in aqueous
 224 solution due to excitation of surface plasmon vibrations in silver nanoparticles. Thus, plant
 225 extracts act as reducing agents as well as capping agents.

226 The papaya fruit extract was mixed in the aqueous solution of the silver ion complex; it
 227 started to change the color from watery to yellowish brown due to reduction of silver ion, which
 228 indicated formation of silver nanoparticles (Jain et al., 2009). UV-Vis spectroscopy could be
 229 used to examine size- and shape-controlled nanoparticles in aqueous suspensions. Five plant leaf
 230 extracts (Pine, Persimmon, Ginkgo, Magnolia and Platanus) were used and compared for their
 231 extracellular synthesis of silver nanoparticles (Song and Kim, 2009). Stable silver nanoparticles
 232 were formed by treating aqueous solution of AgNO_3 with the plant leaf extracts as reducing
 233 agent of Ag^+ to Ag^0 . Magnolia leaf broth was the best reducing agent in terms of synthesis rate

234 and conversion to silver nanoparticles. The average particle size ranged from 15 to 500 nm.
235 Silver nanoparticles were rapidly synthesized using leaf extract of *Acalypha indica* and the
236 formation of nanoparticles was observed within 30 min with the size of 20–30 nm (Krishnaraj
237 et al., 2010). Ali et al., (2011) showed that the leaf extract of menthol is very good bioreductant
238 for the synthesis of silver nanoparticles and synthesized nanoparticles were found to be spherical
239 in shape with 90 nm.

240 3.4. Characterization of silver nanoparticles formulations

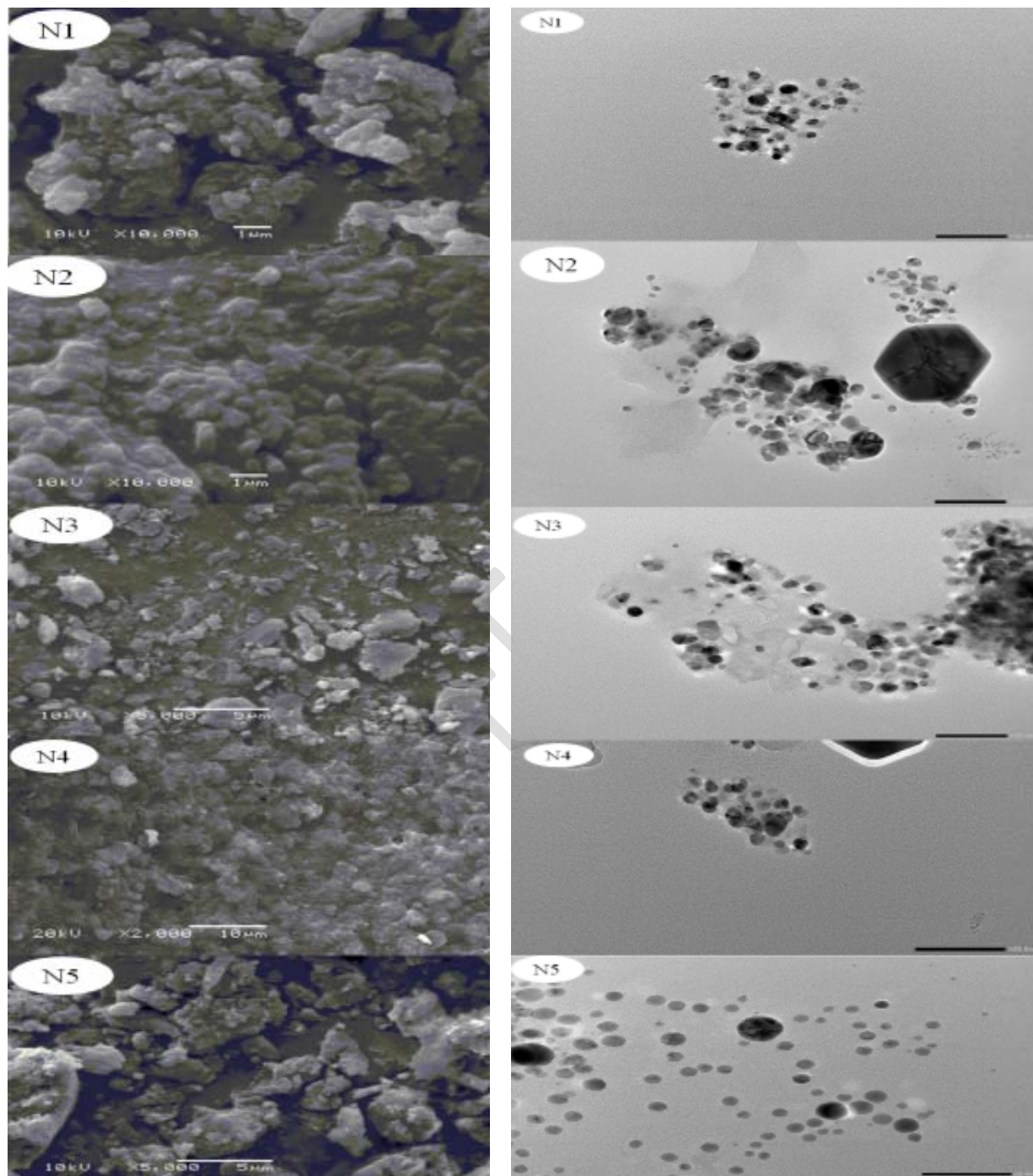
241 The obtained silver nanoparticles was subjected to different characterization methods; SEM,
242 TEM, XRD and FT-IR. From the SEM and TEM micrograph of AgNPs, different extracts
243 produce different size and different crystals, which occurs different effective of the activity of
244 nanoparticles on organisms (Fig. 4). FT-IR results revealed that the obtained particles are silver
245 nanoparticles when compared with the standard nanosilver profile (Fig. 5). It was noticed that
246 extract which produced silver nanoparticles in the range of 12-33 nm, and detected the function
247 group which coated on the surface of particles by X-RD and FT-IR.

248 The biosynthesised silver nanoparticles by using papaya fruit extract was confirmed by XRD
249 and SEM (Jain et al., 2009). The characteristic peaks observed in the XRD image showed in
250 (Fig. 6) three intense peaks in the whole spectrum of 2θ value ranging from 10 to 80. The XRD
251 pattern average size of the particles synthesized was 15 nm with size range 10 to 50 nm with
252 cubic and hexagonal shape. The SEM image showing the high-density silver nanoparticles
253 synthesized by the papaya extract further confirmed the development of silver nanostructures.

254 FT-IR analysis was used for the characterization of the extract and the resulting
255 nanoparticles (Bar et al., 2009). The peaks near 3450 cm^{-1} and near 2933 cm^{-1} were assigned to
256 O–H stretching and aldehydic C–H stretching, respectively. The weaker band at 1643 cm^{-1}
257 corresponds to amide I, arising due to carbonyl stretch in proteins. The peak at 1031 cm^{-1}
258 corresponds to C–N stretching vibrations of the amine. IR spectroscopic study confirmed that the
259 carbonyl group from amino acid residues and proteins has the stronger ability to bind metal
260 indicating that the proteins could possibly form a layer covering the metal nanoparticles (i.e.,
261 capping of silver nanoparticles) to prevent agglomeration and thereby stabilize the medium. This
262 suggests that the biological molecules could possibly perform dual functions of formation and
263 stabilization of silver nanoparticles in the aqueous medium.

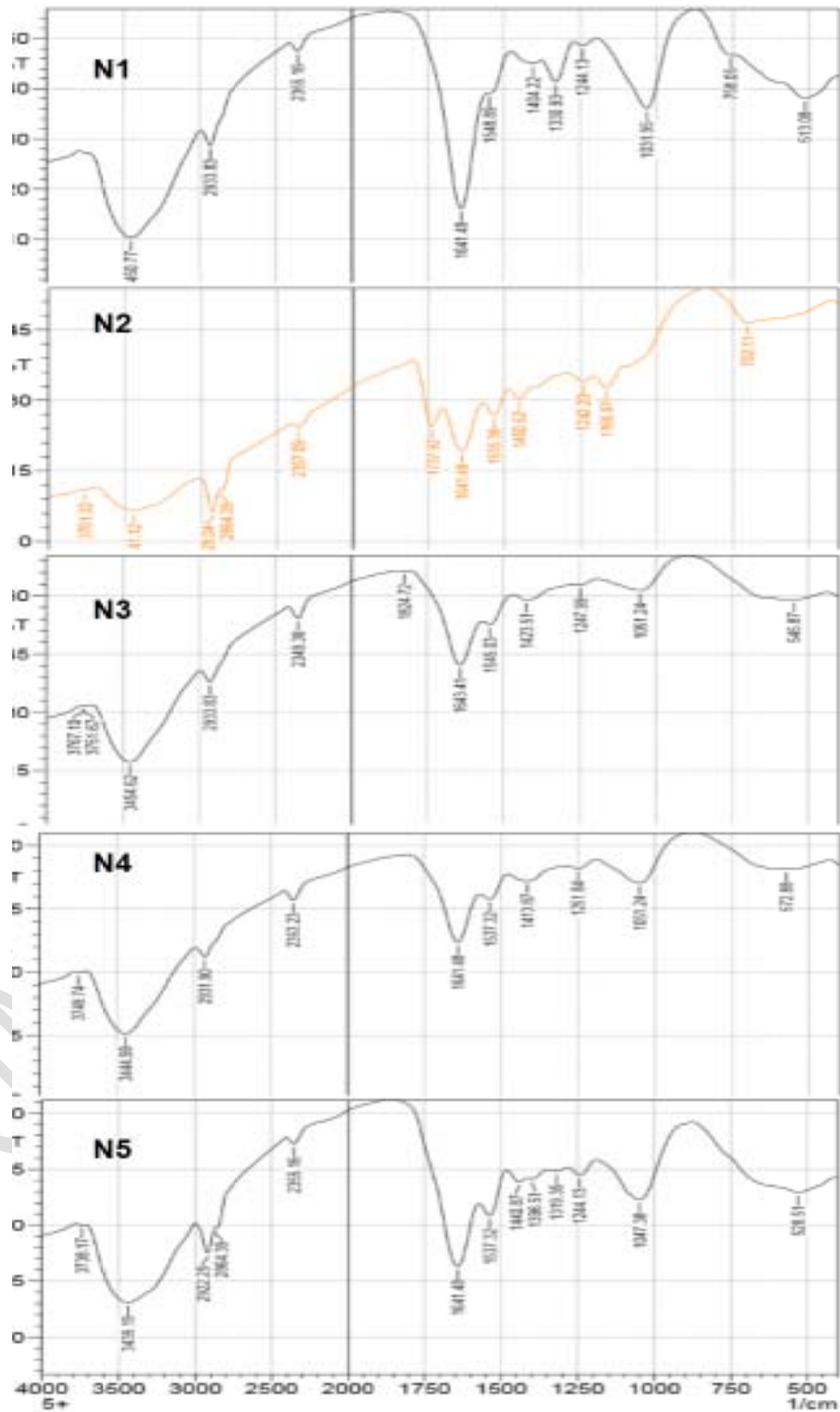
264 FTIR analysis was used for the characterization of the silver nanoparticles using *Garcinia*
265 *mangostana* leaf extract (Veerasingam et al., 2011). Absorbance bands were observed at 1619,
266 1522, 1340, 1160 cm^{-1} . These absorbance bands are known to be associated with the stretching
267 vibrations for –C–C– [(in-ring) aromatic], C–O–C (ethers) and C–O (–C–OH). In particular, the
268 1160 cm^{-1} band arises most probably from the C–O of aromatic-OH group (such as
269 hydroxyflavones and hydroxyxanthenes). The total disappearance of this band after the
270 bioreduction may be due to the fact that the polyols are mainly responsible for the reduction of

271 Ag ions, whereby they themselves get oxidized to unsaturated carbonyl groups leading to a broad
272 peak at 1660 cm^{-1} (for reduction of Ag) (Jain et al., 2009).
273



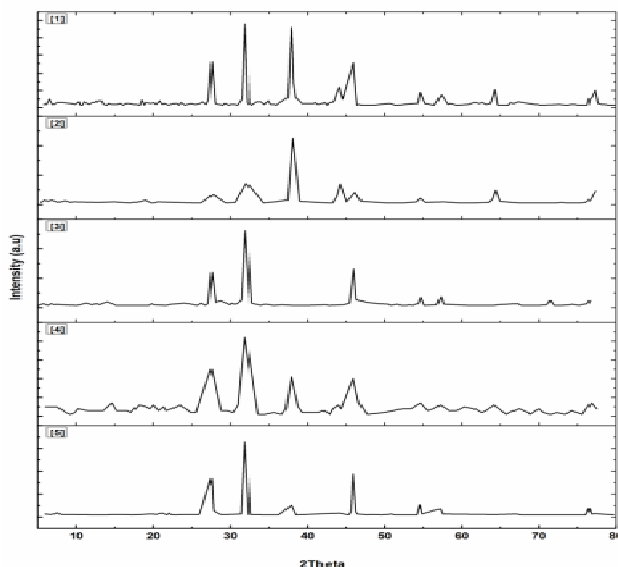
274 **Fig. 4:** SEM (right) and TEM (left) of silver nanoparticles formulations, N1 to N5. The SEM
275 was performed on a JEOL JSM-1200EX II scanning electron microscope operating at an
276 acceleration voltage of 80.0 kV with 20 µm aperture.

277
278



279

280 **Fig. 5:** The FT-IR spectra of the five biosynthesized silver nanoparticles: *S. nigrum* (N1); *P.*
281 *peruviana* Red (N2); Leave *P. peruviana* (N3); Green *P. peruviana* (N4); *M. oliefera* (N5).



282
283 **Fig. 6:** The XRD analysis of the five biosynthesized silver nanoparticles: *S. nigrum* (N1); *P.*
284 *peruviana* Red (N2); Leave *P. peruviana* (N3); Green *P. peruviana* (N4); *M. oliefera* (N5).

285 3.5. Antibacterial activity of silver nanoparticles formulations

286 The *in vitro* antibacterial activity of silver nanoparticles formulations against *R.*
287 *solanacearum* and *P. carotovra* is presented in Table 1 using the agar diffusion method. The
288 measured zone of inhibition of the silver nanoparticles formulations showed significantly
289 different inhibitory effects. The results demonstrated that all formulations showed good
290 inhibition (Inhibition (%) ranged from 20.56 to 29.26 %) against the tested bacteria and the
291 inhibitory effects were concentrations dependent. For the five silver nanoparticles formulations,
292 N5 formulation exerted significantly potent antibacterial activity against *R. solanacearum*.
293 Followed by N4 in the descending order. However, N1 formulation was the lowest active as
294 showed in Fig. 7. Against *P. carotovra*, N1 formulation exerted significantly potent antibacterial
295 activity. Followed by N5 in the descending order. However, N4 formulation was the lowest
296 active as showed in Fig. 7. When we consider the susceptibility of the microorganisms, another
297 point deserves attention; it can be noticed that bacterium of *P. carotovra* was more susceptible
298 than *R. solanacearum* to all formulations (Table 1 and Fig. 7). It appears that the antibacterial
299 activity of the silver nanoparticles formulations increased with increase in surface-to-volume
300 ratio, due to the decrease in size of nanoparticles.

301 Antibacterial effects of Ag nanoparticles obeyed a dual action mechanism of antibacterial
302 activity, i.e., the bactericidal effect of Ag⁺ and membrane-disrupting effect of the polymer
303 subunits. The antibacterial activities of Ag nanoparticles, Ag⁺ ions were blocked by
304 thiolcontaining agents. Silver was also known to cause pits in bacterial cell walls, leading to an

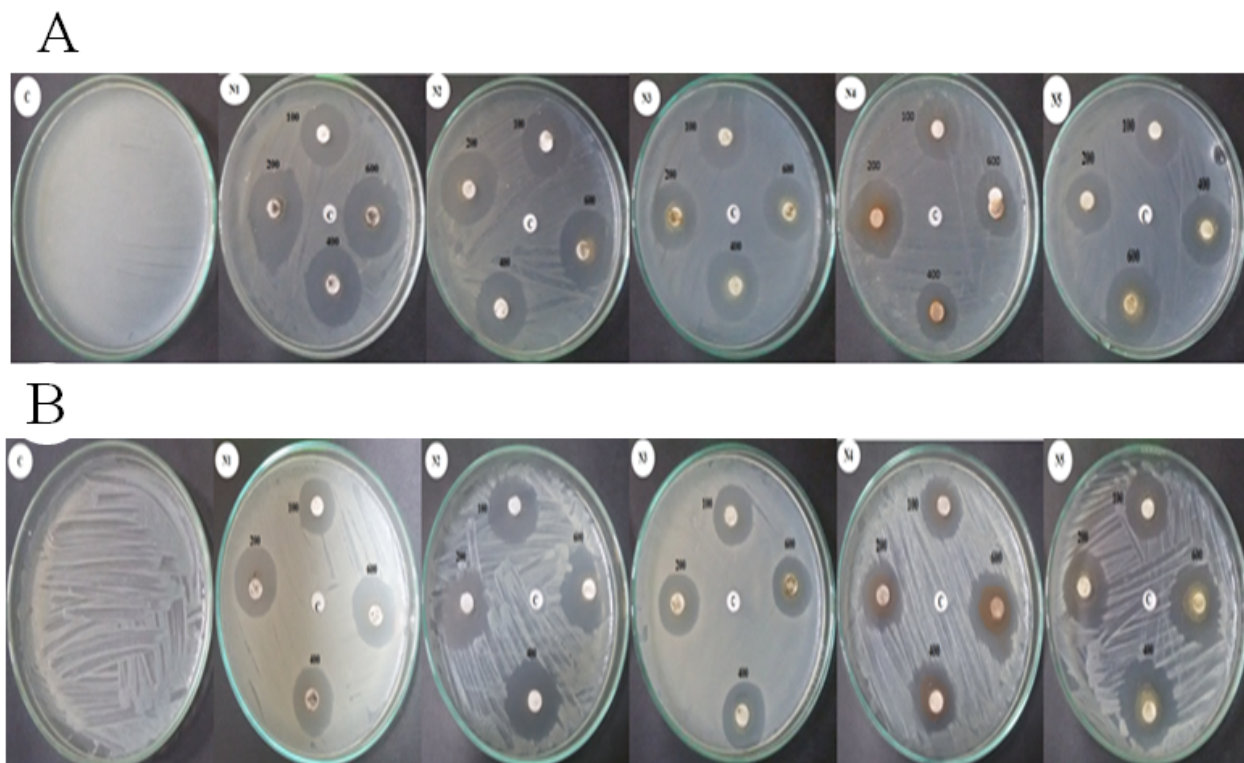
305 increased cell-membrane permeability and cell death (Sambhy et al., 2006). The antibacterial
 306 activity of synthesized silver nanoparticles using leaf extract of *Acalypha indica* showed
 307 effective inhibitory activity against water borne pathogens, *Escherichia coli* and *Vibrio cholera*
 308 (Krishnaraj et al., 2010). Silver nanoparticles 10 g/ml were recorded as the minimal inhibitory
 309 concentration (MIC) against *E. coli* and *V. cholerae*. Alteration in membrane permeability and
 310 respiration of the silver nanoparticle treated bacterial cells were evident from the activity of
 311 silver nanoparticles.

312 **Table 1:** The *in vitro* antibacterial activity of biosynthesized silver nanoparticles against *R.*
 313 *solanacearum* and *P. carotovra* by the agar diffusion method

Formulations	Conc. (mg/L)	Inhibition (%)	
		<i>R. solanacearum</i>	<i>P. carotovra</i>
N1	100	20.95 ± 1.39	23.70 ± 0.64
	200	21.48 ± 0.64	24.81 ± 1.70
	400	21.48 ± 2.24	27.04 ± 1.70
	600	22.22 ± 1.46	29.26 ± 0.64
N2	100	20.56 ± 1.11	21.07 ± 1.11
	200	23.04 ± 1.79	23.37 ± 1.89
	400	23.14 ± 2.62	25.9 ± 2.79
	600	24.63 ± 2.50	26.26 ± 1.45
N3	100	22.04 ± 0.32	23.70 ± 1.69
	200	22.96 ± 1.69	24.07 ± 1.89
	400	23.15 ± 1.15	25.00 ± 0.56
	600	23.26 ± 1.67	25.22 ± 0.91
N4	100	20.74 ± 2.56	16.85 ± 6.62
	200	21.85 ± 2.56	22.22 ± 3.33
	400	22.04 ± 4.01	22.59 ± 0.64
	600	25.00 ± 1.11	23.89 ± 2.00
N5	100	22.41 ± 0.32	21.85 ± 0.84
	200	26.30 ± 2.74	23.15 ± 1.60
	400	26.67 ± 1.11	26.67 ± 1.11
	600	27.04 ± 0.64	28.52 ± 2.31
Doxycycline	30	12.77 ± 0.40	22.89 ± 0.99

314 Green *P. peruviana* (N1); Red *P. peruviana* (N2); leaves of *P. peruviana* (N3); *S. nigrum* (N4)
 315 and *M. olieferia* (N5).

316



317
 318 **Fig. 7:** The *in vitro* antibacterial activity of silver nanoparticles formulations against *P.*
 319 *carotovra* (A) and *R. solanacearum* (B) by the agar diffusion method with different
 320 concentrations (0, 100, 200, 400 and 600 mg/L, respectively). Green *P. peruviana* (N1); Red *P.*
 321 *peruviana* (N2); leaves of *P. peruviana* (N3); *S. nigrum* (N4) and *M. oliefera* (N5).

322 3.6. Antifungal activity of silver nanoparticles synthesized with plant extracts

323 The *in vitro* antifungal activity of silver nanoparticles formulations against the plant
 324 pathogenic fungus *F. oxysporum* is presented in Table 2 and the results are expressed as EC₅₀.
 325 Most of the tested compounds showed inhibitory effect against tested fungus. For the five silver
 326 nanoparticles formulations, N1 formulation exerted significantly potent antifungal activity with
 327 EC₅₀ of 687.03 mg/L against *F. oxysporum*. Followed by N3 in the descending order with EC₅₀
 328 of 981.61 mg/L. However, N2 formulation was the lowest active (EC₅₀ = 1474.86 mg/L against
 329 tested fungus as showed in Fig. 8. Standard fungicide, Ridomil gold showed the highest
 330 fungicidal activity (EC₅₀ = 204.02 mg/L). From statistical analysis, there is no significant
 331 difference between standard fungicide and N1 formulation (see Table 2).

332 Different concentrations of biosynthesized silver nanoparticles were tested to know the
 333 inhibitory effect of fungal plant pathogens namely *Alternaria alternata*, *Sclerotinia sclerotiorum*,
 334 *Macrophomina phaseolina*, *Rhizoctonia solani*, *Botrytis cinerea* and *Curvularia lunata*.
 335 Remarkably, 15 mg concentration of silver nanoparticles showed excellent inhibitory activity
 336 against all the tested pathogens (Krishnaraj et al., 2012). Narayanan and Park, (2014)
 337 demonstrated the synthesis of silver nanoparticles using turnip leaf extract and its interaction

338 with wood-degrading fungal pathogens, *Gloeophyllum abietinum*, *G. trabeum*, *Chaetomium*
 339 *globosum*, and *Phanerochaete sordida*. The synthesized silver nanoparticles showed broad-
 340 spectrum antifungal activity against wood-degrading fungi by inhibiting growth.

341 Reports on the mechanism of inhibitory action of silver ions on microorganisms have shown
 342 that upon treatment with Ag⁺, DNA loses its ability to replicate resulting in inactivated
 343 expression of ribosomal subunit proteins, as well as certain other cellular proteins and enzymes
 344 essential to ATP production (Feng et al., 2000; Yamanaka et al., 2005). It has also been
 345 hypothesized that Ag⁺ primarily affects the function of membrane-bound enzymes, such as those
 346 in the respiratory chain (McDonnell and Russell, 1999).

347
 348 **Table 2.** The *in vitro* antifungal activity of biosynthesized silver nanoparticles against *F.*
 349 *oxysporium* by mycelia radial growth technique.

350

Formulations	EC ₅₀ ^a (mg/L)	95% confidence limits		Slope ^b ± SE	Intercept ^c ± SE	(χ ²) ^d
		Lower	Upper			
N1	687.03	39.36	687.03	1.588±0.60	-4.42±1.95	0.41
N2	1474.86	1087.44	1709.49	1.942±0.57	-6.152±1.84	1.83
N3	981.61	99.52	1321.13	1.404±0.57	-4.20±1.85	0.27
N4	1319.49	685.56	1588.39	1.629±0.57	-5.08±1.83	0.03
N5	999.61	257.19	1306.04	1.596±0.58	-4.79±1.87	0.78
Ridomil gold	204.02	138.44	680.25	0.976±0.31	-2.26±0.62	0.58

351 ^aThe concentration causing 50% mycelial growth inhibition.

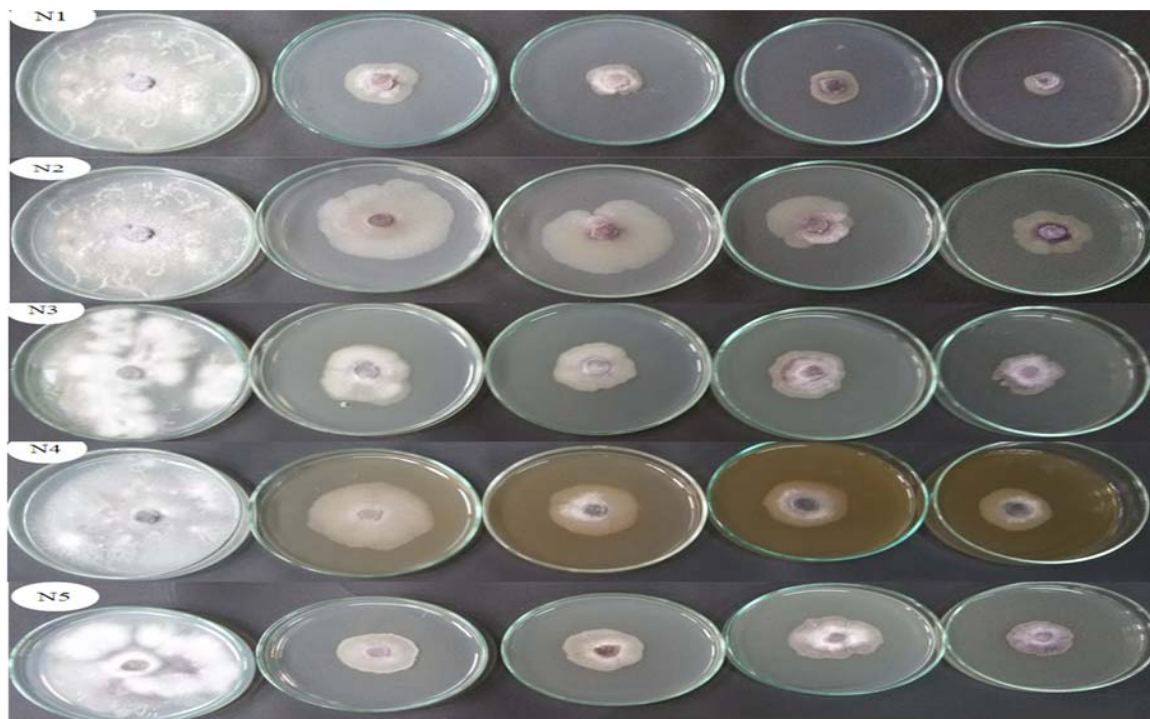
352 ^bSlope of the concentration-inhibition regression line ± standard error.

353 ^cIntercept of the regression line ± standard error.

354 ^dChi square value.

355 Green *P. peruviana* (N1); Red *P. peruviana* (N2); leaves of *P. peruviana* (N3); *S. nigrum* (N4)
 356 and *M. oliefera* (N5).

357



358
359

360 **Fig. 8:** The antifungal activity of the silver nanoparticles formulations (from left to right, 0,
361 1200, 1600, 2000 and 2400 mg/L, respectively) against *F. oxysporium*. Green *P. peruviana* (N1);
362 Red *P. peruviana* (N2); leaves of *P. peruviana* (N3); *S. nigrum* (N4) and *M. oliefera* (N5).

363 **4. Conclusion**

364 The silver nanoparticles have been formed by *P. peruviana*; *S. nigrum* and *M. oliefera*
365 extracts, which is an efficient, eco-friendly and economical process. FT-IR spectrophotometer,
366 XRD, SEM and TEM techniques have confirmed the reduction of silver nitrate to silver
367 nanoparticles. The zones of inhibition were formed in the antimicrobial screening test showed
368 that the Ag NPs synthesized in this process has the efficient antimicrobial activity against the
369 tested pathogenic bacteria and fungi. The biologically synthesized silver nanoparticles could be
370 of immense use in agriculture field for their efficient antimicrobial function. Nanoparticles may
371 be particularly effective delivery systems for plant extracts due to their ability to facilitate
372 antimicrobial application and increase antimicrobial efficacy.

373

374 **Ethical approval:** NA

375 **Consent:** NA

376

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