

## Original Research Article

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## Antifungal Activity of Copper Oxide Nanoparticles against the Fungal Pathogens Isolated from Arid Environment of Jodhpur

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### ABSTRACT

Five plant pathogenic fungi were isolated from infected plants of groundnut, castor and cumin growing under arid environment. The isolates belonged to two fungal genera and were identified as *Alternaria tenuissima* (CZC-2), *Alternaria alternata* (CZCU-1), *Fusarium equiseti* (CZCU-2), *Fusarium oxysporum* (CZCU-4) and *Fusarium solani* (CZGN-9) based on ribosomal DNA sequencing. The effect of copper oxide nanoparticles on growth inhibition of these fungal pathogens was carried out in solid and liquid media. In the plate method *A. tenuissima* (CZC-2) showed the maximum diameter of colony growth in control plates in both the media, whereas the minimum diameter of colony growth was shown by *Alternaria tenuissima* CZC-2 and *Fusarium solani* CZGN-9 in the potato dextrose agar (PDA) and rose bengal agar (RBA) supplemented with 500 ppm of copper oxide nanoparticles (CNP), respectively. In the plate method the maximum inhibition of 68.8 % was reported against *A. tenuissima* CZC-2 in plates supplemented with 500 ppm of copper nanoparticles. Whereas in liquid medium there was complete growth inhibition of *A. tenuissima* CZC-2 and *F. oxysporum* CZCU-4 at 250 ppm concentration of copper oxide nanoparticles.

#### Keywords

Pathogenic fungi,  
Inhibition,  
*Alternaria* and  
*Fusarium*

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### Introduction

To feed the increasing world population by improving the agricultural production in sustainable manner is a great challenge being faced agri-researchers. Despite of great strides in agriculture in the last few decades many developing countries are still vulnerable in achieving the food security (Husen and Siddiqi, 2014; Kasana *et al.*, 2017). Agricultural crops face various abiotic and biotic stresses adversely affecting the crop yields which are more severe in arid environments resulting in overall reduction in

crop yield. Along with others factors growth of fungal pathogens in plants is one of the main causes for considerable economic loss during the production and postharvest handling of food grains. Prevalence of many plant pathogenic diseases and insect injuries results in the loss of about one-third of plant harvest worldwide (Bramhanwade *et al.*, 2016). India contributes more than 90 % of cumin (*Cuminum cyminum* L.), 65 % of Castor (*Ricinus communis* L.) and is the major producer of peanut (*Arachis hypogaea* L.) at global level (FAOSTAT 2016). The castor oil is commercially valued for

ricinoleic acid and non-edible oil (Ladda and Kamthane 2014; Jeong and Park 2009), Cumin for cuisines having therapeutic significance (Bettaieb *et al.*, 2011) and as an antioxidant (Nadeem and Riaz, 2012). Peanut is a known major source of edible oil. Nevertheless, incessant attack by various pathogenic fungi causing wilt, root rot and blight diseases is major limiting factors in production of these crops (Chattopadhyay *et al.*, 2016; Sharma and Rana, 2017; Lodha and Mawar, 2007; Singh *et al.*, 2020a, 2020b). The correct molecular identification is a prerequisite for the management of the target pathogens. Nucleotide variations in ITS regions encompassing ITS-1, 5.8S rRNA gene and ITS-2 have been used extensively for molecular diagnostics of pathogens (Jain *et al.*, 2013; Singh *et al.*, 2013; Gautam *et al.*, 2016). Various agrochemical and fungicides are used to control the plant pathogenic fungi resulting in decrease in outbreak of fungal diseases. At the same time continuous application of conventional fungicides has resulted in development of resistance to many fungicides making it difficult to control the plant pathogens. With the advent of biotechnology various metal nanoparticles has been used as antibacterial and antifungal agents for inhibiting the growth of plant pathogens. The scope of copper oxide nanoparticles has been a keen area of interest for the researchers due to their distinguished antibacterial and antifungal activity which has opened new frontiers to biological sciences (Kasana *et al.*, 2017). Inhibition efficiency of copper nanoparticles against fungi is significant than other metal nanoparticles (Servin *et al.*, 2015, Viet *et al.*, 2016). Various pathogenic bacteria belonging to *Bacillus subtilis*, *Enterococcus faecalis*, *Escherichia coli*, *Klebsiella pneumonia*, *Legionella pneumophila*, *Pseudomonas aeruginosa*, *Pseudomonas fluorescens*, *Shigella sonnei*, *Staphylococcus aureus*, *Staphylococcus epidermidis* and

*Xanthomonas campestris* has been found to be inhibited by copper nanoparticles (Mahmoodi *et al.*, 2018). Previous research on antifungal effects of copper nanoparticles against plant pathogenic fungi has mostly been carried out without stating the concentrations used (Kanhed *et al.*, 2014, Shende *et al.*, 2015, Bramhanwade *et al.*, 2016). However, there is one report where copper nanoparticles at 450 ppm showed the maximum inhibition of *Fusarium sp.* growth in the solid medium after 9 days of incubation (Viet *et al.*, 2016). The present study was conducted to observe the effect of copper oxide nanoparticles in both solid and broth culture media against five fungal pathogens isolated from three crops growing in arid environments of Jodhpur.

## **Materials and Methods**

### **Isolation of pathogens**

The fungal pathogens were isolated from infected plants of groundnut, castor and cumin. Diseased plant samples were brought to the laboratory and washed thoroughly with distilled water to remove the surface contaminants. The infected plant parts were cut into about 1 inch pieces and surface sterilized using 2 % sodium hypochlorite for 2 minutes and washed three times in sterilized distilled water and aseptically transferred on to petri plates containing PDA medium. The inoculated Petri plates were incubated at  $28 \pm 2^\circ\text{C}$  in a BOD incubator for five days. Pure cultures were raised from peripheral mycelium with a piece of culture media from petri plates into test tubes and again incubated at  $28 \pm 2^\circ\text{C}$  for 10 days. Pure cultures were then stored at in a refrigerator until used.

### **DNA isolation**

Fungal mycelium from each pure culture was aseptically transferred to the broth culture

medium and incubated in a BOD incubator at  $28 \pm 2^\circ\text{C}$  for one week. The genomic DNA was extracted from approximately 100 mg of fresh mycelium by crushing it in micro centrifuge tubes using micro-pestles in liquid nitrogen. The HiPura kit of HiMedia Company and protocols suggested by Birren & Lai 1993; Sambrook *et al.*, 1989 were followed for genomic DNA isolation. The DNA was eluted in 200  $\mu\text{l}$  of Tris EDTA buffer (TE buffer). The yield and purity of extracted DNA was determined electrophoretically on 0.8 % agarose gel and spectrophotometrically at 260 and 280 nm wavelength using BioPhotometer (D30, Eppendorf Germany)

### Identification of pathogens

Molecular identification of the fungal strains was done by amplifying the internal transcribed spacer (ITS) region of ribosomal DNA, which encompasses the 5.8S gene from genomic DNA in a thermocycler using ITS-1 (5' TCC GTA GGT GAA CCT GCG G 3') and ITS-4 (5' TCC TCC GCT TAT TGA TAT GC 3') universal primers from each fungal pathogen. Each PCR reaction of 50  $\mu\text{l}$  contained: 1x PCR buffer (Sigma Aldrich), 2.5 mM  $\text{MgCl}_2$ , 150 mM dNTP mix (Thermo Scientific), 100 pmol of each ITS-1 and ITS-4 primers, 1U *Taq* DNA polymerase (Sigma Aldrich) and 50 ng genomic DNA. The PCR was initiated by incubating the reaction mixture at  $94^\circ\text{C}$  for 3 min, followed by 36 cycles of 1 min at  $94^\circ\text{C}$ , annealing at  $55^\circ\text{C}$  for 1 min and extension at 1 min at  $72^\circ\text{C}$  and finally extension for 10 min at  $72^\circ\text{C}$ . The PCR products were separated on 1.6 % agarose gel prepared in  $1\times$  TAE buffer in an electrophoresis run for 100 min at 60 V. The agarose gel was pre-stained with 12  $\mu\text{l}$  of ethidium bromide and photographs captured under UV light using gel documentation system.

The ITS amplified region of each PCR product was sequenced by Sanger di-deoxy sequencing technology employing ABI prism DNA sequencer using universal primers. The BLASTN program <http://www.ncbi.nlm.nih.gov/BLAST/>; National Center for Biotechnology Information, Bethesda, MD) was used for homology searches with the standard program default for identification of fungi.

### Phylogenetic analysis

For phylogenetic analysis the ITS1–5.8S–ITS2 gene sequences of the species of genus *Fusarium* and *Alternaria* were downloaded from NCBI data bases. Multiple sequence alignments of the sequence was performed using CLUSTAL W (Thompson *et al.*, 1994). The evolutionary history was inferred using the Neighbor-Joining method (Saitou and Nei 1987). The evolutionary distances were computed using the Maximum Composite Likelihood method (Tamura *et al.*, 2004). *Mucor irregularis* was used as out group. All ambiguous positions were removed for each sequence pair and evolutionary analyses were conducted in MEGA X (Kumar *et al.*, 2018).

### Antifungal activity

Copper oxide nanoparticles with <50 nm particle size (Sigma Aldrich) were used for antifungal activity. Growth of fungi and antifungal activity/inhibition assays were performed in different media; solid media and liquid medium.

### In solid media

To study the antifungal activity of copper oxide nanoparticles in solid media the growth of five pathogenic fungi in following media was checked and compared i) PDA ii) PDA having 250 ppm CNP iii) PDA having 500 ppm CNP iv) rose bengal agar (RBA) v) RBA

having 250 ppm CNP vi) RBA having 500 ppm CNP. Solidified plates of media described above prepared and with then with the help of cork borer fungal discs of 7 mm diameter were transfer on to these plates. Plates incubated at  $28 \pm 2^{\circ}\text{C}$  and the diameter of the fungal colonies was determined after the incubation periods of 9 days. The inhibition efficiency of fungal plant pathogens was calculated by using formula as suggested by Vincent (1947).

$$I = \frac{C - T}{C} \times 100$$

Where, I is inhibition percentage; C is growth of fungal plant pathogens in control plates (cm), and T is growth of fungal pathogens in CNP supplemented plates (cm).

### **In liquid medium**

To study the effect of nanoparticles on growth of fungi in the liquid media the different pathogenic fungi were grown i) potato dextrose broth medium (PDB), ii) PDB with 250 ppm CNP iii) PDB with 500 ppm CNP.

Different fungal cultures were inoculated into 100 ml of the above medium and incubated at  $28 \pm 2^{\circ}\text{C}$ . The flasks were observed for growth up to 7 days of incubation then the fungal samples were filtered through filter paper. Then the paper with fungal filtrate was observed to determine the fungal biomass.

## **Results and Discussion**

### **Isolation and identification**

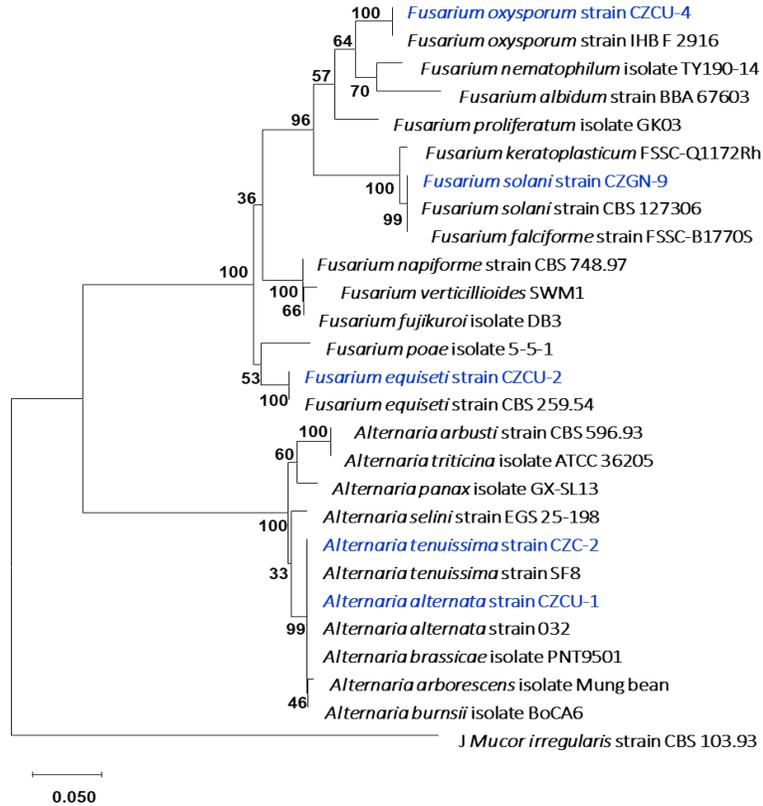
Five pathogenic fungal isolates designated as CZC-2 from castor, CZCU-1, CZCU-2 and CZCU-4 from cumin and CZGN-9 from groundnut were selected for the studies. After preliminary microscopic observations the identification of fungal pathogens; was conducted based on the sequencing ITS1–5.8S–ITS2 region. The different fungal

isolates CZC-2, CZCU-1, CZCU-2, CZCU-4 and CZGN-9 showed a homology of 100, 100, 99.81, 99.81 and 100 % with *Alternaria tenuissima* strain SF8, *Alternaria alternata* strain 032, *Fusarium equiseti* strain CBS 259.54, *Fusarium oxysporum* strain IHB F 2916 and *Fusarium solani* strain CBS 127306 respectively with published species in the literature (Zhao *et al.*, 2018, Oliveira *et al.*, 2018, Vu *et al.*, 2019). Based on their homology the isolates were identified as *Alternaria tenuissima* CZC-2, *Alternaria alternata* CZCU-1, *Fusarium equiseti* CZCU-2, *Fusarium oxysporum* CZCU-4 and *Fusarium solani* CZGN-9. Fungal pathogens belonging to various species of genus *Alternaria* and *Fusarium* have been isolated from castor, cumin and groundnut, and identified by morphological and molecular methods (Zaman and Ahmed, 2012; Ozer and Bayraktar, 2015; Sharma and Rana, 2017). Sequences of all the fungi were submitted to NCBI with accession numbers MF166760, MF166764, MF166765, MF166767 and MF166779 respectively. Ssequences of the isolates were then aligned and compared with previously published sequences of the species from genus *Fusarium* and *Alternaria* and a neighbor-joining phylogenetic trees was constructed using MEGA X (Kumar *et al.*, 2018). In phylogenetic analysis the three species of *Fusarium* formed three different clusters while the two species of *Alternaria* were grouped in a single cluster (Fig. 1).

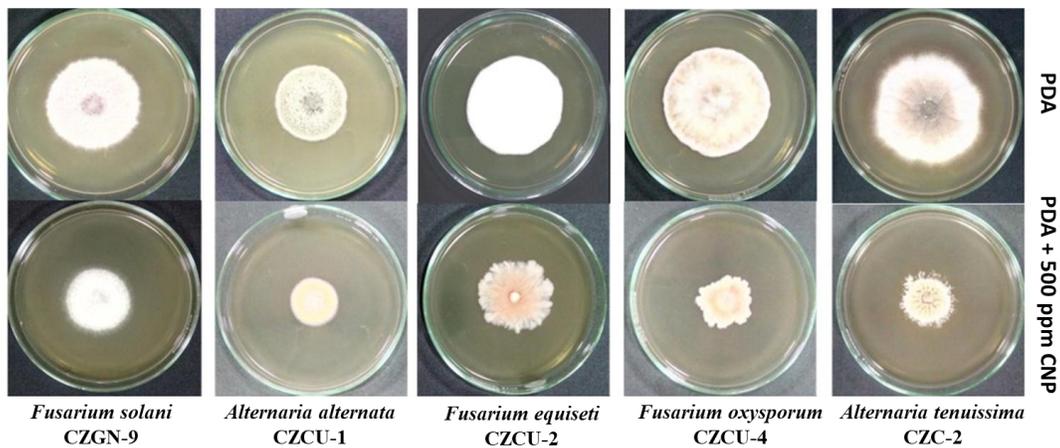
### **Antifungal properties of copper nanoparticles against various pathogens**

The effect of copper oxide nanoparticles of > 50 nm size on development of five pathogens belonging to two genera and five species was observed by determining the diameter of the fungal colonies on two fungal culture media. The growth of each pathogenic fungi was observed both in control medium and treatment medium supplemented with copper oxide nanoparticles.

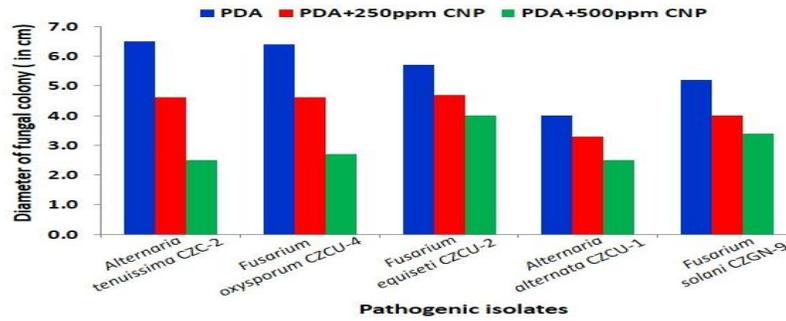
**Fig.1** Phylogenetic tree based on ITS1–5.8S–ITS2 region sequences, drawn using the neighbor joining method and showing the relationship between *Alternaria tenuissima* CZC-2, *Alternaria alternata* CZCU-1, *Fusarium equiseti* CZCU-2, *Fusarium oxysporum* CZCU-4 and *Fusarium solani* CZGN-9 and species from genus *Alternaria* and *Fusarium*. The sequences were downloaded from NCBI database. *Mucor irregularis* was used to root the tree. Bar, 0.05 substitutions per site. Evolutionary analyses were conducted in MEGA X



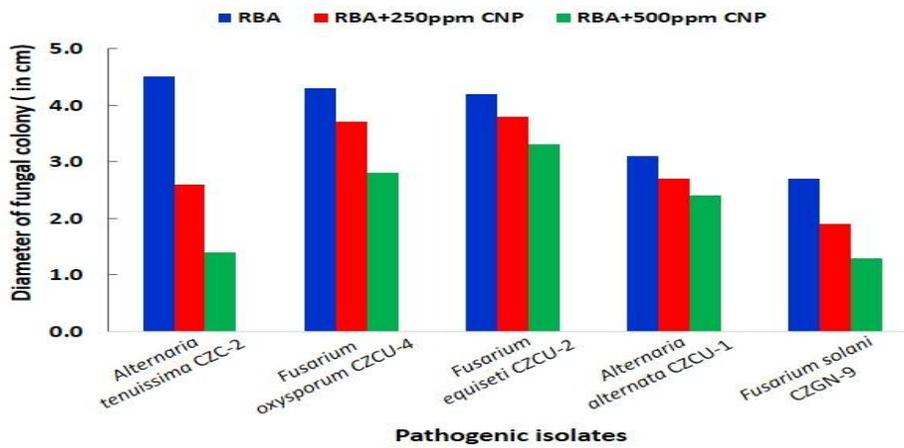
**Fig.2** The diameter of colonies of various fungi in the control (PDA) and treatment (PDA+500 ppm of CNP) after 9 days of incubation



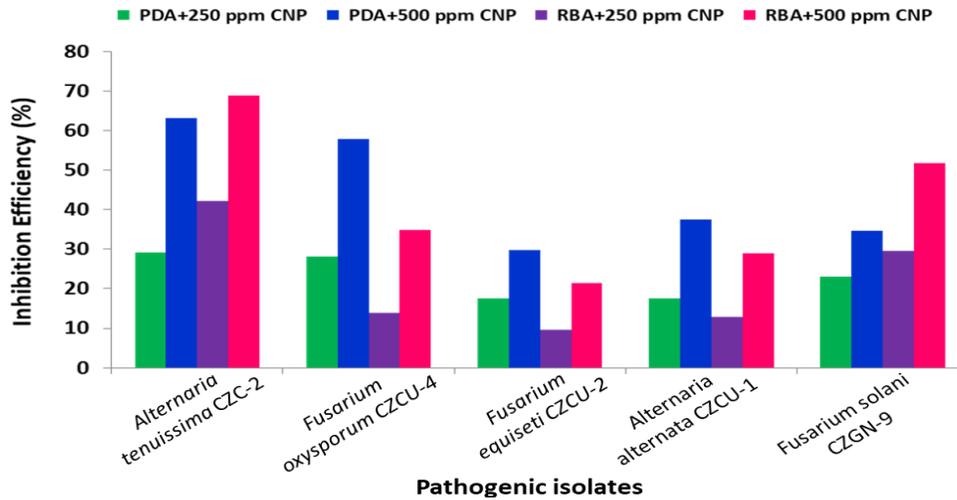
**Fig.3** Colony diameter of various pathogenic fungi in potato dextrose agar (control and treatment plates) after incubation for 9 days



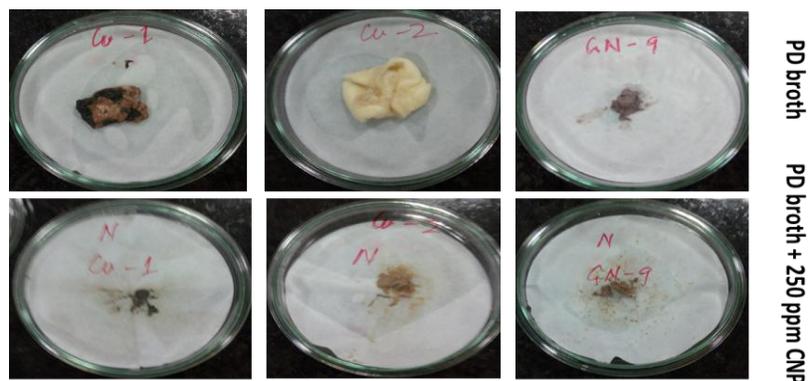
**Fig.4** Colony diameter of various pathogenic fungi in rose bengal agar (control and treatment plates) after incubation for 9 days



**Fig.5** The inhibition efficiency of copper oxide nanoparticles at 250 ppm and 500 ppm against various pathogenic fungi



**Fig.6** The fungal biomass of pathogenic fungi in potato dextrose broth and potato dextrose broth +250 ppm of CNP



The representative images of growth of colonies of various test pathogenic fungi in PDA medium (control) and medium supplemented with 500 ppm of copper oxide nanoparticles are shown (Fig. 2). Similar pattern was observed in case of rose bengal agar culture medium supplemented with copper oxide nanoparticles. From the results it is evident that addition of copper oxide nanoparticles inhibited the growth of all the five pathogens tested; *A. tenuissima*, *A. alternata*, *F. equiseti*, *F. oxysporum* and *F. solani*. The diameter of fungal colonies of various test pathogens on PDA and PDA supplemented with 250 ppm and 500 ppm CNP and RBA and RBA supplemented with 250 ppm and 500 ppm CPM is shown in figures (Fig. 3 and Fig. 4). The maximum diameter of colony growth was observed in case of *A. tenuissima* CZC-2 in control plates in both the media. The minimum diameter of colony growth was shown by *A. tenuissima* CZC-2 and *F. solani* CZGN-9 in the PDA and RBA supplemented with 500 ppm of CNP respectively. It was also observed that the fungal mat was thin in case of medium mixed with copper oxide nanoparticles as compared to the control.

From the data on growth the inhibition efficiency of copper oxide nanoparticles calculated at two different concentrations in

two different media against five pathogenic fungi is shown in figure (Fig. 5). Results showed good inhibition efficiency of copper nanoparticles against all the pathogens in Petri plate method. The inhibition efficiency of copper oxide nanoparticles was concentration dependent and found to increase with increase in the concentration used. The maximum inhibition of 68.8 % and 63.3 % respectively were reported against *Alternaria tenuissima* CZC-2 in rose bengal agar and potato dextrose agar supplemented with 500 ppm of copper oxide nanoparticles. Further the growth of different pathogenic fungi was observed in potato dextrose broth supplemented with copper oxide nanoparticles. It was interesting to report that in none of the fungi was able to grow in the broth medium supplemented with 500 ppm of copper oxide nanoparticles. Even at 250 ppm there was no growth of two fungal isolates (*A. tenuissima* and *F. oxysporum*) while other three isolates also showed very less growth at this concentration (Fig. 6). Antifungal activity of copper oxide nanoparticles against many pathogenic fungi belong to various genera *Phoma*, *Curvularia*, *Alternaria*, and *Fusarium* using the inhibition zone method have been carried without providing the concentrations of copper nanoparticles (Kanhed *et al.*, 2014, Shende *et al.*, 2015, Bramhanwade *et al.*, 2016). In another study on effect of copper

oxide nanoparticles against a single isolate of *Fusarium* sp. showed maximum inhibition of growth at 450 ppm in the solid medium after 9 days of incubation (Viet *et al.*, 2016).

In conclusion the five pathogenic fungi belonging to two genera and five species were isolated from infected plants of groundnut, castor and cumin growing under arid environment and molecularly identified. In the plate assay inhibition of 68.8 % and 63.3 %, respectively was observed against *Alternaria tenuissima* in rose bengal agar and potato dextrose agar supplemented with 500 ppm of copper oxide nanoparticles. While in case of liquid medium at 250 ppm concentration there was complete inhibition of *Alternaria tenuissima* and *Fusarium oxysporum* and other three isolates also showed very less growth.

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