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## RESEARCH ARTICLE

ANTIBACTERIAL ACTIVITY AND PHYTOCHEMICAL SCREENING OF WILD EDIBLE  
MUSHROOM *PLEUROTUS OSTREATUS* COLLECTED FROM HIMACHAL PRADESH.

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### Abstract

**Objective:-**The objective of present study was to identify the unknown wild edible mushroom based on morphological characteristics and DNA based sequencing by using universal primers ITS 1 and ITS 4, antibacterial activity, phytochemical screening and FTIR analysis of mushroom extract spectra.

**Methods:-** The wild edible mushroom specimen was tentatively identified morphologically and was cultured on 2% MEA. The isolate thus obtained was screened for its antimicrobial activity. Methanol extract was prepared in order to carry out antibacterial activity. The extract was subjected to qualitative phytochemical and FTIR analysis.

**Results:-**The isolates on the basis of morphological and molecular characterization was identified as *P. ostreatus*. Agar-well diffusion method was used to carry out antimicrobial assay. The methanol extract of *P. ostreatus* showed antimicrobial activity against *E.coli* having zone inhibition diameter  $24.3 \pm 0.5$ , against *S. aureus* having zone inhibition diameter  $26.6 \pm 0.5$ , against *P. aeruginosa* having zone inhibition diameter  $20.3 \pm 0.5$  and against *K.pneumoniae* having zone inhibition diameter  $28.6 \pm 0.5$ . Fourier transform infrared (FT-IR) analysis indicated that all mushroom's extract spectra showed typical carbohydrate pattern with strong and broad absorption near  $3300 \text{ cm}^{-1}$  which indicates the presence of OH group.

**Conclusion:-**The isolate analysis have been showed to be good source of antimicrobial agent and phytoconstituents, therefore could be potential source of bioactive compounds with beneficial biological activities.

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### Introduction:-

The growing emergence of drug resistance microorganism now a days has become a serious threat to effective treatment of infections (Klein *et al.*, 2007). Resistance to drug happens usually due to long term indiscriminate use of antimicrobial agents which results in acquisition of mutation in the genome and genes of disease causing microorganisms that assist their survival (Mazodier and Davies, 1991, Gao *et al.*, 2003). Whilst it is not possible to prevent the bacterial evolution, but it is important to choose the most appropriate and reliable antibiotics and use them properly to minimize the development of drug resistant strains (Alves *et al.*, 2012).

Mushrooms are source of nutritionally functional food as well as physiologically beneficial medicines. (Sagakami *et al.*, 1991, Wasser and Weis, 1999). The fruiting body and mycelium constitutes the compounds with a

wide range of antimicrobial activity. In mushrooms the cell wall glucans are well known for their immunomodulatory properties along with many of the externalized secondary metabolites that combat bacteria and viruses, hence are known to be rich source of antibiotics (Benedict and Brady, 1972, Kupraet *al.*, 1979, Suzukiet *al.*, 1990, Collins and Ng, 1997, Eoet *al.*, 1999, Brandt and Piraino, 2000). It is also believed that many mushrooms species consist of antimicrobial compounds that help them to protect themselves and survive in their natural environment and some of them proved to be beneficial for humans (Lindequist, 2005). The effect of mushroom's extract against pathogenic bacteria are studied in different parts of the world by a number of researchers (Gbolagadeet *al.*, 2007, Turkoglu *et al.*, 2007). The active domain of present study was to investigate the antimicrobial properties of wild edible mushroom having variety of beneficial biological properties which may play important role in the human health maintenance and to search for new products having these properties.

## **Material and methods:-**

### **Collection and isolation of sample:-**

Mushroom fruiting body was collected from the forest of Badu Sahib, Distt, Sirmour, Himachal Pradesh and was labelled as sample no. 105. The fruiting body was brought to laboratory in a sterile plastic bag washed thoroughly with sterile distilled water and 0.001% mercuric chloride and with the aid of sterile blade it was aseptically break lengthwise exposing the inner tissue. A small piece of tissue was aseptically transferred on to the plates of Malt extract agar.

### **Identification of sample:-**

#### **Morphological identification**

The identification of fruiting body of wild mushroom was based upon sporocarp morphology and macroscopic characteristics.

### **Molecular identification:-**

#### **Genomic DNA isolation and PCR amplification:-**

Genomic DNA isolation was done by CTAB (Cetyltrimethyl ammonium lysis buffer) method (Zolan and Pukkila 1986). The entire region of isolate 105 was amplified by using universal primers ITS 1 and ITS 4. PCR reaction were performed in 50 µl reaction mixture containing 1 µl of DNA, 1 X PCR buffer, 1.5 mM MgCl<sub>2</sub>, 0.2 mM of each dNTPs, 2.5 units of Taq DNA polymerase, 0.5 µM of each primers i.e., ITS1 (5' TCCGGTAGGTGAACCTGCGG 3') ITS 4 (5' TCCTCCGCTTATTGATATGC 3'). The tubes were placed in a thermal cycler (Agilent sure cycler 8800) for amplification under following conditions: One cycle Initial denaturation at 94 °C for five minutes, 35 cycles of denaturation at 94 °C for 30 seconds, annealing at 62 °C for 30 seconds, elongation at 72 °C for one minute and final extension at 72 °C for five minutes. Forward and reverse DNA sequencing reaction of PCR amplicon was carried out with ITS1 and ITS4 primers using BDT v3.1 Cycle sequencing kit on ABI 3730xl Genetic Analyser at Eurofins genomics, Bangalore. The 18S rDNA gene sequence was used to carry out BLAST with the database of NCBI gene bank database. Based on maximum identity score first ten sequences were selected and aligned using multiple alignment software program Clustal W. Distance matrix was generated using RDP database and the phylogenetic tree was constructed using MEGA 4.

### **Antimicrobial activity:-**

#### **Source of test microorganism used:-**

Four bacterial strains *Klebsiella pneumoniae* MTCC109, *Pseudomonasaeruginosa* MTCC741, *Staphylococcus aureus* MTCC 737 and *E. coli* MTCC739 were obtained from Microbial Type Culture Collection (MTCC) IMTECH, Chandigarh (INDIA) and stored at 4°C in refrigerator and sub cultured at regular intervals of 48 h until use.

### **Extract Preparation:-**

The mycelia grown on MEA plates were inoculated in a conical flasks having liquid medium (pH 6.6) culture containing glucose 2%, peptone 1%, and yeast extract 2%. The flasks were shaken at 150 rpm at 25°C. The culture fluid was separated from pellets by filtration after 7 days of fermentation and then extract three times with methanol solution. The methanol fractions was evaporated and the extracted material was dissolved in DMSO (Merck). All extracts and fractions from mycelial cultures were stored at 4°C until use for assaying their activities (Rojas *et al.*, 2006).

### **Antibacterial activity:-**

Muller Hinton agar plates with 4% NaCl supplementation were prepared. Sterilized swabs were dipped in standardized bacterial suspension with inoculum size of  $1.5 \times 10^8$  cfu/ml prepared above and excess culture was removed by turning the swab against the side of the tube. Inoculum was spread evenly over the entire surface of Muller Hinton agar plates. These plates were allowed to dry for at least 15 min and then wells (7mm diameter) were made on petri dish using sterile cork borer. About 25 $\mu$ l extract was introduced into bore agar wells using sterile dropping pipette. These plates were kept inside the refrigerator at 4°C for 6 hours to allow proper diffusion of extracts into medium. The plates were then examined for antibacterial activities of extracts after 24 hours of incubation at 37°C. Antimicrobial activity was determined by measuring the diameter zone of inhibition in mm (Moshi *et al.*, 2006).

#### **Preliminary phytochemical screening:-**

The freshly prepared extract was subjected to phytochemical analysis in order to ensure the presence of phytochemical constituents.

#### **Test for alkaloids:-**

**Mayer's test:-** 5ml of extract was taken and few drops of Mayer's reagent was added by the side of test tube. White creamy precipitate were observed indicating positive results.

**Wagner's test:-** 5ml of extract was taken and few drops of Wagner's reagent was added by the side of test tube. Reddish brown precipitate appeared confirming the test as positive.

#### **Test for Carbohydrates:-**

**Molisch's test:-** 5ml of extract was mixed with two drop of alcoholic solution of  $\alpha$  naphthol and shaken well. 1 ml of concentrated H<sub>2</sub>SO<sub>4</sub> was added along the side of the test tube and allowed to stand. The violet ring appeared indicating the presence of carbohydrates.

**Benedict's test:-** To the 3ml of extract 1 ml of Benedict's reagent was added. The mixture was heated on boiling water bath for two minutes. The characteristic colour precipitate was observed confirming the presence of sugar.

#### **Test for detection of phenolic compounds and tannin:-**

**Ferric chloride test:-** Few drops of neutral ferric chloride was added to 5ml of extract. The dark green colour was appeared.

**Lead acetate test:-** To the 5ml of extract 4 ml of 10 % lead acetate was added. Bulky white precipitate was appeared confirming the presence of phenolic compounds.

#### **Test for flavonoids:-**

**Alkaline reagent test:-** Few drops of NaOH was added to test solution intense colour was formed which turn colourless on the addition of dilute acid confirming the presence of flavonoids.

#### **Fourier transform infrared spectroscopy (FTIR):-**

FTIR spectroscopy of the samples was tested by using FTIR spectrometer (Aligent Cary- 630) (Grosveet *et al.*, 2001). 10 mg of sample was put on mirror stage of spectrometer and spectrum was obtained by tightening the lens over sample. After analysis the lens was cleaned to remove the sample completely (Jouraiphyet *et al.*, 2008).

#### **Results:-**

##### **Identification of sample:-**

##### **Morphological characterization:-**

**Cap:** 4-10 cm; convex, becoming flat or somewhat depressed; kidney-shaped, pale brown to dark brown in colour having in-rolled margins.

**Gills:** Running down the stem; close; with a gray tinge.

**Stem:** Usually rudimentary and lateral.

**Flesh:** Thick; white.

Spores 6-9 x 3.5-5  $\mu$ ; smooth; cylindrical. Mycelia are septate and consist of clamp connections as shown in Fig 1.

##### **Molecular identification:-**

In the present study genomic DNA was isolated from mycelial culture of isolate and was amplified with ITS 1 and ITS 4 primers. PCR products obtained from couple of primers ITS1 and ITS 4 provided a single band at 500 bp. The sequences for sample was recorded and shown in table 1. (<http://blast.ncbi.nlm.nih.gov/Blast.cgi>), sequence of sample was homologous with partial sequence of 5.8 S ribosomal RNA gene, internal transcribed spacer 2 complete sequence and 28S ribosomal gene partial sequence of species. *Pleurotus ostreatus* accession number HM 770898.1, GU722280.1, KR908742.1, KF724525.1, KJ010873.1. The aligned nucleotide sequence of sample was submitted to NCBI and nucleotide sequence is provided with gene bank accession no. KU 892064. The phylogenetic tree constructed (Fig 2) also revealed that the sample 105 was classified in Kingdom Fungi, division Basidiomycota, Class Agaricomycetes, Order Agaricales, Family Pleurotaceae, Genus *Pleurotus* and Species *Pleurotus ostreatus*.

#### **Antibacterial activity:-**

Screening for antibacterial activity was done by agar well diffusion method. Results of antibacterial activity was shown in Table 2. Methanol was used as solvent for the preparation of extract and finally dissolved in DMSO. The extract showed varying degree of inhibition on test microorganisms taken for screening. It was observed from the results that extract was highly effective against *K. pneumoniae* having zone inhibition diameter  $28.6 \pm 0.5$  followed by *S. aureus*  $26.6 \pm 0.5$ , *E. coli*  $24.3 \pm 0.5$  and *P. aeruginosa*  $20.3 \pm 0.5$  respectively.

#### **Preliminary phytoconstituents analysis:-**

The preliminary phytoconstituents analysis of *P. ostreatus* revealed that the extract contain alkaloid, carbohydrates, flavonoids, Phenolic compounds and Tannins as shown in Table 3.

#### **Fourier transform infrared spectroscopy (FTIR):-**

An infrared spectrum of samples was recorded with an FT-IR spectrometer (Agilent technologies) in the range of  $4000-800 \text{ cm}^{-1}$ . The infrared spectrum helps to evaluate the vibrations of molecule and polar bonds present between the atoms of sample molecules. As shown in Figure 3 and table 4. The mushroom extract spectra showed typical carbohydrate pattern with strong and broad absorption near  $3264 \text{ cm}^{-1}$ , which indicates the presence of OH groups. Other regular features was also identified in FT-IR spectra of extract. The peaks at  $2921 \text{ cm}^{-1}$  in sample 105, are interpreted due to  $\text{CH}_2$  asymmetric stretch/ symmetric stretch. The signal at  $1547 \text{ cm}^{-1}$  is indicative of Secondary amine N-H bending. The region around  $1208$  and  $1156 \text{ cm}^{-1}$  in sample is dominated by sugar ring vibrations overlapping with stretching vibration of (C-OH) side group and (C-O-C) glycosidic bond vibration. The peak near  $1374 \text{ cm}^{-1}$  corresponds to C-H in plane bending vibrations.

#### **Discussion:-**

Resistance to antibiotic among bacteria has been increasing drastically, therefore nutraceuticals and drugs having antibacterial activities are needed. Microorganisms such as *S. aureus*, *K. pneumoniae*, *E. coli*, *P. aeruginosa* are examples of resistance to antibiotics now a days. (Alves *et al.*, 2005) 14 mushroom isolates with higher antimicrobial activity against target microorganism was detected by Rosa *et al.*, in 2003. In another study by Zjawiony, 2004 it was observed that about 75% of polypore fungi showed strong antimicrobial activity. In the present study the extract showed good antimicrobial activity against all the pathogenic bacteria taken which are similar to the results reported from extracts prepared from *P. eryngii*, *P. ostreatus*, *P. sajor-caju*, *L. squarrosulus* and *Agaricus bisporus* against *K. pneumoniae* and *S. aureus*. (Alves *et al.*, 2012). Yamac and Bilgili (2006) in his study found that mycelial extract of many fungi possess high inhibitory activity against gram negative bacteria our results are in collaboration with this finding. The mycelial extract was found to have good inhibitory effect against *E. coli*, *P. aeruginosa* in the present study.

Phytochemical analysis of *P. ostreatus* has revealed that it consist of alkaloids, carbohydrates, flavonoids, phenolic compounds and tannins. The results are in accordance to the previous literature reported (Okwulehie and Ogoke, 2013). In order to investigate the molecular properties of extract of mushroom and to evaluate the co- relation between their structural characteristics and bioactivity in the present study FTIR spectra technique was used. The peak at  $2925$  is due to H-C-H stretching of alkane (Kozarskiet al., 2012). The  $1200-1000 \text{ cm}^{-1}$  region is dominated by sugar ring vibration overlapping with stretching vibration of (C-OH) side groups. The peaks near  $1374 \text{ cm}^{-1}$  corresponds to C-H in plane bending vibration and at  $1080 \text{ cm}^{-1}$  the stretching band is curved by the signal of pyranose form of the O substitute residue (C-O) (Mathlouthi and Koenig, 1986, Heet *et al.*, 2012). The band near  $1030 \text{ cm}^{-1}$  are indicative of  $\alpha$ -linkage which correlates the study of Kozarskiet al., 2012. The extract therefore considering the criteria that it consist of both  $\alpha$  and  $\beta$  glycosidic conformations.



**Table 1:** Nucleotide sequence of wild edible mushroom.

<i>P. ostreatus</i>	GCTTTATCGTTCGTAGTGACCTGCGGAGGATCATTAATGAATTCACATGGAGTT GTTGCTGGCCTCTAGGGGCATGTGCACGCTTCACTAGTCTTTCAACCACCTGTGA ACTTTTGATAGATCTGTGAAGTCGTCTCTCAAGTCGTCAGACTTGGTTGCTGGGA TTAAACGCTCTCGGTGTGACTACGCAGTCTATTTACTTACACACCCCAAATGTAT GTCTACGAATGTCATTTAATGGGCCTTGTGCCTATAAACCATAATAACAACCTTCA ACAACGGATCTCTTGGCTCTCGCATCGATGAAGAACGCAGCGAAATGCGATAAG TAATGTGAATTGCAGAATTCAGTGAATCATCCGAATCTTTGAACGCACCTTGCG CCCCTTGGTATTCCGAGGGGCATGCCTGTTTGAGTGTGCATTAATTTCTCAAACCTC ACTTTGGTTTCTTTCCAATGTGATGTTTGGATTGTTGGGGGCTGCTGGCCTTGA CAGGTCGGCTCCTCTTAAATGCATTAGCAGGACTTCTCATTGCCTCTGCGCATGA TGTGATAATTATCACTCATCAATAGCACGCATGAATAGAGTCCAGCTCTCTAAT CGTCCGCAAGGACAATTTGACAATTGACCTCAA
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**Table 2:** Zone inhibition of methanolic extract of *P.ostreatus* against pathogenic bacteria.

Antibacterial activity of Methanol extract			
Zone inhibition diameter in mm			
Microorganisms	Extract	DMSO (-ve control)	Ciprofloxacin (+ve control)
<i>E. coli</i>	24.3±0.5	-	21.5±0.5
<i>S.aureus</i>	26.6±0.5	-	28.6±0.5
<i>P. aeruginosa</i>	20.3±0.5	-	22.3±0.5
<i>K. pneumoniae</i>	28.6±0.5	-	30.6±0.5

Each value is expressed as mean±SD of three replicates, - indicates no activity

**Table 3.** Phytochemical analysis of edible mushroom.

Phytochemicals	Test	Observation
Alkaloids	Mayer's test	+
	Wagner's test	+
Carbohydrates	Molisch's test	+
	Benedict's	+
Phenolic compounds and Tannin	Ferric chloride test	+
	Lead acetate test	+
Flavonoids	Alkaline reagent test	+

**Table 4:**Infrared spectroscopy correlation table.

<b>Pleurotus ostreatus</b>		
1.	3275	Hydroxyl group (O-H stretching)
2.	2921	(H-C-H asymmetric stretch/ symmetric stretch) Alkane group
3.	1547	Secondary amines (N-H bending )
4.	1030	α-linkage
5.	1156	(C-O-C) glycosidic bond vibration
6.	1208	sugar ring vibrations overlapping with stretching vibration of (C-OH) side group
7.	1374	C-H in plane binding vibrations.
8.	1320	O-H group of phenolic compound
9.	1644	Amide band indicating presence of some residual protein

### Conclusion:-

The methanol extract of *P. ostreatus* found to have good antimicrobial activity against gram positive and gram negative bacteria and being a polar solvent it was found to extract maximum phytoconstituents. The molecular properties and compounds detected by FTIR analysis warrant investigation for the compounds having potential to improve human health, and application as dietary supplements.



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