INTRODUCTION

Natural environment is still the most important contributor of novel drugs in the face of development of combinatorial chemistry, which quickly generated thousands of new chemicals [1]. Microbial sources from marine environments are prolific producers of natural antibiotics, with massive prevalence of drug resistant pathogens and microorganisms. It is crucial to search for novel antibiotics with wide applications in chemotherapy, plant pathology, food preservation, veterinary medicine, biotechnology and molecular biology. The applicable antibiotics may affect the screening of bioactive compounds, with selective and sensitive bioactivity detection methods of their importance [2]. The biosynthesis of antibiotics like other microbial metabolites is determined by a number of regulatory factors [3]. Including growth conditions, carbon, nitrogen, mineral salt levels and physical parameters like temperature, pH and agitation during production. These conditions may vary from species to species from many of the microbes that live in extreme environments like high temperatures, high salt concentrations, low pH, and high radiation etc., influence the specifically fungal growth and metabolite productions. Usually the bioactive metabolite production from microorganisms depends on their special adaptations to their environment [4]. Some of primary metabolic components may serves as a branching point of metabolic pathways which leads to the end products of secondary metabolic products. Growth media and incubating conditions have a very strong influence on secondary metabolite production. Secondary metabolism is regulated by the factors like carbon sources, nitrogen sources, phosphate, NaCl, trace elements and different parameters like temperature, p[H] and incubating time intervals [5]. These regulatory factors involves in balancing of biosynthesis and yield of antibiotic activity. Conventional practice of single factor optimization maintains other factors to get involved in improvement of unspecified level to optimum constant level. The study of optimization for secondary metabolite production from marine fungi is to determine the minimum inhibitory concentration of the fungal metabolite against the test pathogens. So our present study is focussed on optimization of antibiotic production by marine fungi.

MATERIALS AND METHODS

Isolation of marine fungi

The marine soil sediments are collected from the Nizampatnam mangroves soil, Guntur district (Andhra Pradesh, India) are serially diluted with sterile distilled water. Aliquots of 0.2 ml from each sample are spread onto potato dextrose agar plates (containing 20% potato, 2% dextrose, 2% agar). The antibiotics about 50 mg were added to molten media to inhibit the growth of bacteria. Two replicate agar plates are used for each sample. After 5 d of incubation at 30 °C, the inoculated agar plates were examined daily for the presence of fungal hyphae, using a dissecting microscope at 20× magnification. Distinct fungal colonies on the agar plates were then transferred to new agar plates for further isolation and purification [6].

Basal medium

According to literature Potato dextrose broth medium is used as a basal medium. The fungal species about 5 mm mycelial disk obtained from 7 d old spore culture are inoculated in 100 ml of the medium dispersed in 250 ml sterilized conical flasks and incubated at 27 °C for 5 d. After incubation the mycelia are harvested by filtration using whatman no 1 filter paper, washed thoroughly with distilled water to remove medium components. Excess of water is removed by blotting with filter papers. The mycelia are allowed to get dry at 50 °C to express as dry weight (in mg/250 ml). The production of bioactive metabolites are expressed by measuring the diameter of the inhibition zone against test organisms including Salmonella enterica MTCC 10248, Escherichia coli ATCC 8739,
Identifications of fungal isolates

Morphological studies of Mycelia is done by using lacto phenol cotton blue staining to identify genus level [7]. The effects of salt concentration on the growth of isolated fungi were determined by [8]. The selected strains are cultured with potato dextrose Broth medium at 27 °C for 5 d.

Production of antimicrobial substances from marine fungi

All the potent fungal isolates were inoculated in to each potato dextrose broth and incubated at 27 °C for 5 d. Then biomass is removed aseptically using sterile spatula, and crude was extracted by using double amount (1:1) of chemical solvent (ethyl acetate) for 24 h. The organic solvent layer was collected and concentrated by evaporation. The concentrated crude extract was collected for antibacterial assay.

Antimicrobial activity

Antimicrobial activity of marine fungal crude extracts was tested by using agar well diffusion method [9]. A 50 μl of crude extract from stock solution is added in to each well (6 mm diameter). The crude extract diffused in to nutrient agar plates inoculated with test organisms (Salmonella enterica MTCC 10248, Escherichia coli ATCC 8739, Pseudomonas aeruginosa ATCC 27853, Staphylococcus aureus ATCC 25923, Vibrio cholera ATCC 15748). All the plates are incubated at 37 °C for 24 h. The zone of inhibition was measured and expressed diameter was in millimetre.

Minimum inhibitory concentrations (MIC)

MIC of secondary metabolite was performed by agar-cup diffusion method. The crude extract was dissolved in different concentrations ranging from highest concentration of 10,000 μg/ml and then dilution were performed at concentration of 5000 μg/ml, 4000 μg/ml, 3500 μg/ml, 3000 μg/ml, 2500 μg/ml, 2000 μg/ml, 1500 μg/ml, 1000 μg/ml, 750 μg/ml, 500 μg/ml and 250 μg/ml. MIC value of marine fungal crude extract was determined against five different human pathogens belonging to both gram positive and gram negative groups such as Salmonella enterica MTCC 10248, Escherichia coli ATCC 8739, Pseudomonas aeruginosa ATCC 27853, Staphylococcus aureus ATCC 25923, Vibrio cholera ATCC 15748. All the plates are incubated at 37 °C for 24 h. The zone of inhibition was measured and the expressed diameter was in millimetre.

Optimization of growth and active metabolite production

Optimization for the production of bioactive secondary metabolites was made with potato dextrose broth and these studies are used to estimate the potency of the compound by antimicrobial assay to check the activity of the compound from marine fungi. These studies are useful distinguish the effect of physical and chemical parameters on the growth and bioactive metabolite. So, these optimization studies are carried out through each parameter and its dependence on the other parameter on the biomass and metabolite. The optimization studies were done as following.

Effect of optimization on mycelia and bioactive metabolite production

Effect of culture media on mycelia growth and bioactive metabolite production

To select a suitable growth medium for marine fungal isolates (gaps-1 and gaps-2) are grown in different culture media such as Potato dextrose broth, Czapek's Dox broth, Sabouraud dextrose broth, Malt extract broth and Peptone Yeast Glucose broth. The medium which exhibits maximum biomass accumulation and bioactive metabolite (antibiotic) production in terms of zone of inhibition was used as the optimized medium for further study. All the selected media compositions are Hi-media procured from Hi-Media Laboratories, Mumbai, India.

Effect of agitation on mycelia growth and bioactive metabolite production

Effect of agitation on mycelia growth and bioactive metabolite production was studied by incubating culture in conical flasks on an orbital shaker at different rpm (50, 100, 150 and 200 rpm). Control experiment was done by incubating the inoculated media at static condition. The metabolite production of the inoculated media was determined after seven days of incubation at 280C at 150 rpm in an orbital shaker. After incubation the mycelia growth as well as the bioactive metabolite production for each rpm concentration was estimated and recorded.

Effect of salinity on mycelia growth and bioactive metabolite production

The effect of salinity on mycelia growth and bioactive metabolite production by the isolates were carried out by incubating in various NaCl concentrations (3%-7%) at room temperature with optimum incubation time period. The biomass as well as the bioactive metabolite production for each sodium chloride concentration were estimated and recorded.

Effect of incubation time on mycelia growth and bioactive metabolite production

To study the effect of incubation time (h 24–240) 100 ml of potato dextrose broth flasks were inoculated with 5 mm mycelia disks of seven days old fungal culture under aseptic condition and incubated. The mycelia weight and antimicrobial production were recorded at the end of the incubation period [10].

Effect of pH on mycelia growth and bioactive metabolite production

The effect of pH on the mycelia growth and bioactive metabolite production of the isolates were tested by using liquid cultures containing different pH levels (pH 4-9). 100 ml of liquid medium was poured into a 250 ml conical flask under aseptic conditions along with optimized parameters. The medium was adjusted to the desired pH by adding 1N NaOH [11]. Flasks were sterilized at 121 °C at 15 psi for 20 min. Each flask was inoculated with 5 mm diameter mycelia disc in sterile conditions. Inoculated flasks were incubated at 27±1 °C for seven days and the dry mycelia weight and bioactive metabolite productions were recorded.

Effect of temperature on mycelia growth and bioactive metabolite production

Theungal isolates were subjected to different temperature (from 15 °C to 45 °C) with parameters like salinity, incubation and pH to study the optimum temperature required for mycelia growth and bioactive metabolite yield. 100 ml of basal medium is prepared and sterilized at 121 °C at 15 psi for 20 min. Under aseptic conditions 5 mm diameter of the culture discs were inoculated and incubated for seven days. After incubation the dry mycelial weight and the antimicrobial productions were recorded [12].

Effect of carbon sources on mycelia growth and bioactive metabolite production

The fungal isolates were subjected to different carbon temperature (from 15 °C to 45 °C) with parameters like salinity, incubation and pH to study the optimum temperature required for mycelia growth and bioactive metabolite yield. 100 ml of basal medium is prepared and sterilized at 121 °C at 15 psi for 20 min. Under aseptic conditions 5 mm diameter of the culture discs were inoculated and incubated for seven days. After incubation the dry mycelial weight and the antimicrobial productions were recorded [12].

Effect of carbon sources on mycelia growth and bioactive metabolite production

To study the effect of different carbon sources (glucose, lactose, sucrose, fructose, galactose, xyllose, maltose, arabinose, raffinose) about 1% of each is added to the selected medium individually. Each flask with different carbon sources was inoculated with a 5 mm mycelia disc of three days old fungal cultures and incubated for seven days. After incubation period biomass (mycelia weight) and the production of bioactive metabolites are recorded [13].

Effect of nitrogen source on mycelia growth and bioactive metabolite production

To study the effect of different nitrogen sources (beef extract, yeast extract, peptone, ammonium chloride and sodium nitrate) are used about 1% of each to the selected medium individually and the dextrose was used as the source of carbon in all the treatments. Flasks were inoculated with 5 mm mycelia disks of seven days old

fungal culture under aseptic condition and incubated for seven days. The mycelia weight and antimicrobial production were recorded at the end of the incubation period [14-17].

Effect of micro nutrients on mycelia growth and bioactive metabolite production

To study the effect of mineral salts (trace elements) like CaCl₂, FeSO₄, MnCl₂, MgSO₄, ZnCl₂, K₂HPO₄, the experiments were done by adding about 0.01% of each to the selected medium individually and the dextrose was used as the source of carbon and nitrogen sources in all the treatments. Flasks were inoculated with 5 mm mycelia disks of seven days old fungal culture and incubated. The mycelia weight and antimicrobial production were recorded.

RESULTS AND DISCUSSION

Isolation and morphological characterization of marine fungi

Totally 43 fungal isolates were recovered, among these 10 morphologically different strains were sub cultured and maintained for further analysis. According to Microscopic and Cultural characteristics the isolates were belonging to Aspergillus species (date not disclosed) were found predominantly in the marine soil from Nizamapatnam, Guntur district (Andhra Pradesh).

Extraction of secondary metabolite

Extraction of the secondary metabolite was effectively done with ethyle acetate and broth culture in ratio 1:1, and allowed to vacuum drying. The extracted compound was stored in the same 1 ml of solvent and stored for further analysis.

Media optimization

Production of growth and secondary metabolite from marine fungi strongly influenced by parameters like salinity, temperature, pH, incubation time and media components, carbon, nitrogen sources and trace elements. It is significant in order to study the influence of various physico-chemical parameters to find out optimized media for the production of potent secondary metabolite.

In present study, initially one parameter was assessed and it was then incorporated at its optimized level in the subsequent experiments using potato dextrose broth medium.

Effect of Culture media on Secondary metabolite production and biomass by gaps-1, gaps-2 showing maximum growth and antibiotic production in Potato dextrose broth comparatively to other media. Fig. 1 Reveals that the effect of different growth media on biomass and bioactive metabolite production of inoculated gaps-1 and gaps-2 within 6 d of incubation (on orbital shaker incubator). According to fig. 1 among the tested media, maximum mycelial dry weight and bioactive metabolite production was recorded in potato dextrose broth medium.
Effect of Culture media on Secondary metabolite production and biomass by gaps-1, gaps-2 showing maximum growth and antibiotic production in Potato dextrose broth comparatively to other media. Fig. 2 Reveals that the effect of Nacl on Secondary metabolite production and biomass by gaps-1 and gaps-2 showing maximum growth and antibiotic production in pdb at Nacl3% with agitation 150 rpm.

Fig. 2: Effect of Nacl on secondary metabolite production and biomass by GAPS 1 and 2 showing maximum growth and antibiotic production in pdb at Nacl3% with agitation 150 rpm.

Fig. 3: Effect of incubation on microbial growth of marine fungi

Fig. 3 reveals that the effect of incubation period on biomass and bioactive metabolite production of inoculated gaps 1 and gaps 2 within 6d.

Fig. 4: Effect of pH on microbial growth of marine fungi

Fig. 4 explains about effect of pH on Secondary metabolite production and biomass by GAPS 1 and 2 showing maximum growth and antibiotic production in pdb at Nacl3% with incubation time 144h and pH interval from 3.0 to 10.0.

Selected potential marin fungal metabolite were subjected to various concentrations of pH ranging from 3.0 to 10.0 were taken for the study of an interval of pH 1.0. Maximum growth at ph 7.0 (gaps 1) ph 8.0 (gaps-2) and decreased further increases the pH beyond 7.0. No growth was registered at ph range below 4.
Fig. 5: Effect of temperature on microbial growth on marine fungi

From fig. 5 effect of Temperature on Secondary metabolite production and biomass by gaps-1, gaps-2 showing maximum growth and antibiotic production in PDB at Nacl 3%; incubation time 144 h; pH-7(GAPS-1), 8(GAPS-2); temperature-25 °C(GAPS-1), 30 °C(GAPS-2). The requirements for specific physical temperature (25°C and 30°C).

Fig. 6: Effect of carbon sources on microbial growth on marine fungi

Effect of carbon sources on Secondary metabolite production and biomass by gaps-1gaps-2 showing maximum growth and antibiotic production in peb at Nacl 3%; incubation time 144 h; pH-7(GAPS-1), 8(GAPS-2); temperature-25 °C (GAPS-1), 30 °C (GAPS-2). 1% of carbon sources: Glucose (GAPS-1), sucrose (GAPS-2). From fig 6 of all the tested carbon sources (1% w/v) in the selected medium inoculated with gaps-1 and gaps-2 incubated shaken for 6 d, glucose and sucrose supported the highest level of mycelia growth and bioactive metabolite production.
According to fig. 7, the effect of nitrogen sources on secondary metabolite production and biomass by GAPS-1 and GAPS-2 showed maximum growth and antibiotic production in PDB at NaCl 3%; incubation time 144 h; pH 7 (GAPS-1), 8 (GAPS-2); temperature 25°C (GAPS-1), 30°C (GAPS-2). 1% of carbon sources: Glucose (GAPS-1), sucrose (GAPS-2); 1% of nitrogen sources: Beef extract (GAPS-1), Ammonium chloride (GAPS-2). Addition of equimolar amounts of various nitrogen sources in the selected medium supplemented with 1.0% (w/v) showed that beef extract and ammonium sulphate were good nitrogen sources on the mycelia growth and bioactive metabolite production.

Fig. 8 shows the effect of trace elements on secondary metabolite production and biomass by GAPS-1, GAPS-2 showing maximum growth and antibiotic production in PDB at NaCl 3%; incubation time 144 h; pH 7 (GAPS-1), 8 (GAPS-2); temperature 25°C (GAPS-1), 30°C (GAPS-2). 1% of carbon sources: Glucose (GAPS-1), sucrose (GAPS-2); 1% of nitrogen sources: Beef extract (GAPS-1), Ammonium chloride (GAPS-2).
chlordane (GAPS-2) but trace elements shown the effect on growth and antibiotic production by reducing the activity of gaps-1, gaps-2.

Antimicrobial activity and minimum inhibitory concentration

Antimicrobial resistance had become a serious threat in treating of infectious diseases, to overcome this problem many academicians and scientists are search of antimicrobials against the clinical pathogens from marine environments. The present study also can be useful for the survey of antimicrobials from marine fungi. In this present study about 43 isolates were isolated from the marine soil samples which act as antibacterial against the pathogens like Salmonella enterica MTCC 10248, Escherichia coli ATCC 8739, Pseudomonas aeruginosa ATCC 27853, Staphylococcus aureus ATCC 25923, Vibrio cholerae ATCC 15746. Among these isolates 10 strains were selected as they have morphological variations. About 5 samples were collected with highest antibacterial activity and the minimum concentration ranges from 250 μg/ml to 10000 μg/ml but the sample gaps-1 had more activity 250 μg/ml than gaps-2 300 μg/ml table 1.

Table 1: Minimum inhibitory concentration

<table>
<thead>
<tr>
<th>Pathogens</th>
<th>Mic value(μg/ml) Gaps-1</th>
<th>Mic value(μg/ml) Gaps-2</th>
</tr>
</thead>
<tbody>
<tr>
<td>Salmonella enterica</td>
<td>1500</td>
<td>1500</td>
</tr>
<tr>
<td>E. coli</td>
<td>250</td>
<td>300</td>
</tr>
<tr>
<td>Pseudomonas aeruginosa</td>
<td>1000</td>
<td>1000</td>
</tr>
<tr>
<td>Staphylococcus aureus</td>
<td>1500</td>
<td>1500</td>
</tr>
<tr>
<td>Vibrio cholerae</td>
<td>1500</td>
<td>1500</td>
</tr>
</tbody>
</table>

According to the above figures, strains are having high microbial activity when compared with streptomycin as standard. The factor dependent optimization shows that there is a great effect of different types of parameters on the marine fungal growth and production of metabolites against bacterial pathogens. The parameters like pH, temperature, incubation period, macro and micronutrients. In the present study role of trace elements like magnesium, manganese, ferrous and zinc had less metabolite production when correlate with other parameters. The stability of metabolite seems to be about five days and maximum activity is within 24 h of time.

CONCLUSION

The optimization Process of present study concludes that optimum conditions are required for the production of bioactive metabolite from marine fungi gaps-1 and gaps-2. The optimum conditions of the marine fungi were incubation period 144 h pH 8.0 and 7.0 and temperature 25 °C and 30 °C. Aeration also played a crucial role in terms of product formation. Agitation speed of 150rpm showed maximum yield. Secondary metabolites had showed broad spectrum activity against bacteria. Hence these bioactive metabolites can be used as source of antimicrobial. Further studies of these metabolites will be carried on purification, characterization and identification of bioactive metabolites of marine fungi.

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CONFLICT OF INTERESTS

Declare none

REFERENCES


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