INTRODUCTION

Human health is developing complicated diseases day by day due to current lifestyle and changes in their food habits, which leads to the search of new antibiotics for multiple drug-resistant microbes. 1-3. Urgent search for newer and more effective antibiotics that should be cheap and have a minimal environmental impact to deal with these disease problems is now underway. Hence, plant-associated endophytes that are relatively unexplored can be studied for the extraction of a substance containing secondary metabolites. Endophytes have a mutual relationship inside healthy plant tissue and are found abundantly in some part or in the whole plant which spends their whole life or a period of life cycle, without causing any disease symptoms. Endophytic fungi have been widely studied and discovered to produce a significant variety of natural products. They often synthesize several novel metabolites such as alkaloids, terpenoids, steroids, quinones, isocoumarin derivatives, and flavonoids with various pharmaceutical applications such as herbicidal, anticancer, anti-diabetic, anti-microbial, and immunosuppressant. Secondary metabolites produced by the endophytes are similar to those produced by the host, as a consequence of genetic exchange. Thus, it can be said that endophytic fungi have emerged as an alternative source for the production of novel drugs.

From ages, plants were used as traditional medicine due to their curative properties to treat various diseases. Studies have shown that the curative property of medicinal plant is not only because of chemicals present in the plant but also because of the endophytes that reside within the plants hence search for new bioactive strains of endophytic fungi played a very important role in the present scenario. Plectranthus amboinicus, is a perennial herb know as Indian borage, it is widely used as medicinal plant commonly found in tropical countries. Traditionally used to treat variety of medicinal applications such as fever, cough, skin infections, throat infection, constipation and digestive problems. Early research data discloses that the plant possesses antimicrobial, antioxidant and phytochemical properties having 70 volatile and 30 non-volatile compounds with significant pharmacological properties. The leaves of this herb are also used for flavouring and in preparation of traditional food hence, the present study was carried out to isolate endophytic fungi from Plectranthus amboinicus and to investigate their biological properties such as antimicrobial, antioxidant and anti-diabetic activities.

Globally, complicated diseases have been of concern such as Diabetes mellitus (DM), a metabolic disorder considered as a major health concern due to its prevalence that results in the deficiency in insulin secretion, insulin action, or both promoting disturbance of carbohydrate, fat, and protein metabolism by alpha-amylase. These antioxidants also gaining importance in protecting the body against reactive
oxygen species (ROS) toxicity either by removing or by converting those to less reactive molecules, and it is also produced in numerous biochemical reactions in the body and has been implicated as mediators of many diseases, including cancer, atherosclerosis and heart diseases. In recent years, herbal medicine is used all over the world for health care management due to fewer side effects compared with synthetic medicine. In an effort the search for possible sources of novel therapeutic compounds, the present study was carried out to isolate endophytic fungi from *Plectranthus amboinicus* and to investigate their biological properties such as antimicrobial, antioxidant, and anti-diabetic activities.

**MATERIALS AND METHODS**

**Sample Collection**

Fresh and symptomatic leaves of *Plectranthus amboinicus* (Indian borage) were collected from Chikka Aluvara, Kodagu District, Karnataka, India. Leaves were collected in sterile polythene bags and immediately washed with tap water and processed for isolation of endophytic fungi.

**Isolation of endophytic fungi**

Leaves were washed thoroughly in running tap water for 10 min to remove dust, debris, soil and minimize the microbial load from the sample surface. The surface sterilization process was carried out as per the procedure described by with minor modification. The leave samples were surface sterilized with 70 % ethanol for 1 min, 4% sodium hypochlorite for 3 min followed by 70% ethanol for 30 seconds, and finally washed thoroughly in sterile distilled water to remove the traces of alcohol and excess water dried in sterile blotting sheet. Surface sterilized leaves were chopped into small segments and transferred to Saboraud Dextrose Agar (SDA) plates supplemented with streptomycin (100µg/ml) to suppress bacterial growth. These plates were sealed with parafilm and incubated at 28ºC for 4-5 days. The plates were monitored every day to check the growth of endophytic fungal colonies from the segments. After the incubation period, the hyphal tips of the developing fungal colonies were transferred to fresh SDA slants and also SDA plates to maintain the pure cultures.

**Identification of endophytic fungi**

**Morphological and microscopic identification**

In the sporulation stage, fungal mycelium was stained in cotton blue and observed under a microscope. According to their microscopic and morphological appearance, initial identification was done using standard manuals described by .

**18S rRNA gene sequencing**

Total genomic DNA of selected G3 isolated endophytic fungi was isolated directly from actively growing mycelium scraped from SDA plates, using EXpure Microbial DNA isolation kit according to manufacturer’s protocol (Bogar Bio Bee stores Pvt Ltd.,). DNA amplification was performed by PCR using Taq master mix polymerase and the primer pair ITS 1: 5’ AGAGTTGTGATCCTGGTCAG 3’; and ITS 4: 5’ TACGGTACCTTGTTACGACTT 3’ (Millipore) (White et al. 1990) in a thermocycler. PCR was carried out according to the following protocol: initial denaturation 95ºC for 2 min; denaturation 95ºC for 30 sec; annealing 50C for 30s; extension 72ºC for 2 min; final extension 72ºC for 10 min and steps 2-4 were repeated 25 cycles. Each sample prepared for PCR consisted of 5 µL of isolated DNA in 25 µL of PCR reaction solution (1.5 µL of Forward Primer and Reverse Primer), 5 µL of deionized water, and 12 µL of Taq Master Mix.

Purification of PCR Product was done by using Montage PCR clean up kit (Millipore) protocol. The PCR product was sequenced using the primers. Sequencing reaction were performed using a ABI PRISM® BigDyeTM terminator Cycle Sequencing Kits with AmpliTaq® DNA polymerase (FS enzyme) (Applied Biosystems). The fluorescent-labelled fragments were purified from the unincorporated terminators with an ethanol precipitation protocol. The samples were resuspended in distilled water and subjected to electrophoresis in an ABI 3730xl sequencer (Applied Biosystems). To identify the isolates, sequences were subjected to the BLAST search with the NCBI database. Multiple sequence alignments of sequences was used by the program MUSCLE 3.7 . The resulting aligned sequences were cured using th program Gblocks 0.91b. Finally, the program PhyML 3.0 Airt was used for phylogeny analysis and HKY85 as substitution model. The obtained sequences were then analysed using the BLAST algorithm and closely related phylogenetic sequences obtained from the National Centre of Biological Information (NCBI) database. The phylogenetic tree was constructed using Neighbour Joining method in MEGA 5 .

**Preparation of fungal extract**

The cultivation of fungus was done on 400 ml Saboraud Dextrose Broth (SDB) by placing agar blocks of actively growing pure fungal strains in a 500 ml Erlenmeyer flask. Each flask was incubated at room temperature for 21 days with periodical shaking at 150 rpm. The cultures were taken out and filtered through sterile muslin cloth to remove the mycelia mats. The culture filtrate was then filtered according to with some modification by centrifugation at 3,600 rpm for 10 mins and supernatant was extracted thrice with equal volumes of solvent ethyl acetate. The organic phase was collected and the solvent was removed by evaporation under reduced pressure at 45ºC using a rotary vacuum evaporator. The dry solid residue was re-dissolved in ethyl acetate or dimethyl sulphoxide (DMSO) and subjected to bioassays.

**Bioassays**

**Anti-bacterial activity**

**Test microorganisms**

The common foodborne human pathogenic microorganisms were obtained from medical college,
Yenepoya University, Mangalore, India. One Gram-positive (*Enterococcus faecalis*) and four Gram-negative (*Escherichia coli*, *Salmonella typhi*, and *Shigella sonnei*) strains were used throughout the study. Tested bacteria were inoculated in nutrient broth and incubated for 24hrs at 37°C in aerobic condition. The suspension was adjusted to 0.5 McFarland turbidity standard correspondingly (~1.5×10⁸ CFU/ml).

**Antibacterial activity by agar well diffusion method**

The Agar-well diffusion method was used to screen the antibacterial activity of fungal extracts using nutrient agar (NA) medium according to 27. A fresh bacterial suspension adjusted to 0.5 McFarland turbidity standard ~1.5×10⁸ CFU/ml was spread on NA plates with a sterile cotton swab. Using sterile cork borer 6 mm wells were made in which 100 µl of crude DMSO extract was loaded on seeded plates. Positive control was maintained using standard antibiotics, viz Streptomycin at a concentration of 1mg/ml according to Kirby-Bauer disk diffusion susceptibility test protocol with minor modification, and Negative control was also maintained using DMSO. The culture plates were then incubated at 37°C for 24h. The experiment was carried out in three replicates and an inhibition zone was recorded.

**Determination of minimum inhibitory concentrations (MICs) and minimal bactericidal concentrations (MBCs).**

The crude extract G3 isolate that showed a potent antimicrobial activity was further assessed for MIC and MBC assays. The test samples of the crude extracts were dissolved in DMSO. Dilutions were prepared to get final concentrations ranging from 0 to 50 mg/mL. Each well received 10 µl suspension of microorganisms and 100 µl liquid culture media. Plates were incubated at 37°C for 24h, 15 µl of 0.01% resazurin was added to characterize cell viability then re-incubated for 4h. The absence of bacterial growth with the lowest concentration was taken as the MIC. Using the results of MIC were determined by sub-culturing which showed no bacterial growth on agar plates after incubated at 37°C for 24h in an incubator. The complete absence of growth on the agar surface at the lowest sample concentration was defined as the MBC. All experiments were conducted in triplicate. The results were represented as mean ± sd 28.

**Anti-diabetic activity**

**α-Amylase Inhibition Assay**

Alpha-amylase activity was carried out using a modified method described previously by 29. Briefly, 50µl, 100µl, and 200µl of different sample concentrations dissolved in DMSO were added to 150µl starch solution (containing 1% starch and 17mM NaCl). The reaction was initiated by adding 10µl α-amylase (1mg/ml in 0.1M sodium phosphate buffer, pH 7.0) to the mixture and was incubated at room temperature. After 30min, the reaction was stopped by adding 20µl of NaOH solution (2M). Subsequently, 20µl of dinitrosaliclyic acid was added to the reaction mixture and placed in a water bath for 20 min. The enzymatic hydrolysis of substrate was monitored at 540 nm using a spectrophotometer. For each concentration, blank tubes were prepared by replacing the enzyme solution with buffer. Acarbose as a positive control was taken and control representing enzyme activity was replacing the plant extracts with DMSO. The experiments were repeated thrice using the same protocol.

The α-amylase inhibitory activity was calculated using the formula:

\[
\% \text{ inhibitory activity} = \frac{\text{Absorbance(Control) - Absorbance(Extract)}}{\text{Absorbance (Control)}} \times 100
\]

The mode of inhibition of endophytic extract on alpha-amylase action was determined by increasing substrate (starch) concentration. Kinetic parameters, namely Michaelis-Menten constant affinity (Km) and maximum velocity (Vmax) were derived from appropriate Lineweaver-Burk plots.

**Phytochemical Screening**

Preliminary phytochemical screening was performed on the endophytic fungal extracts to detect different phytochemicals such as phenols, flavonoids, alkaloids, terpenoids, saponins, tannins, glycosides, and steroids using standard methods 30.

**Antioxidant assay**

**Total phenolic content (TPC)**

Determination of Total Phenolic Content (TPC) of ethyl acetate extract was determined by Folin-Ciocalteu method 31. One ml of the extract was added to 2 ml of distilled water and 1 ml of Folin Ciocalteu phenol reagent. The mixture was allowed to stand at room temperature for 5 min and then 2 ml of sodium carbonate was added to the mixture. The resulting mixture was read at 765 nm using a UV visible spectrometer against blank. Gallic acid was used to calculate the standard curve (0.01- 0.08 mM; Y = 0.026X + 0.121; R² = 0.979) and the results were expressed as mg of gallic acid equivalents (GAEs).

**DPPH free radical scavenging assay**

DPPH (2,2-Diphenyl-1-picrylhydrazyl) was performed on selected fungal extracts to measure their ability to decolorize the purple-colored methanol solution of DPPH, as described by 32. In brief 3 ml of 0.3 mM methanolic solution (4mg/100ml) of DPPH solution was added to 3ml of extract and shaken vigorously, the tubes were kept in dark for incubation at room temperature for 30 min. Ascorbic acid (0.5 to 1.5µM) was used as a positive control. The absorbance was checked against the blank at 516 nm. The scavenging activity was measured based on the extent of bleaching of purple-colored methanol solution of DPPH treated with fungal extract and the percentage (%) of radical scavenging activity was calculated according to the equation.
%DPPH radical scavenging activity  
\[
\text{Absorbance of blank} - \text{Absorbance of sample} \times 100 \\
\text{Absorbance of blank}
\]

**Ferric Reducing Antioxidant Power (FRAP) Assay**

In this assay, 10 µl of the sample was mixed with 15 µl of 0.1 M sodium phosphate buffer (pH 6.6) and 15 µl of potassium ferricyanide (1% w/v) followed by incubation at 50°C for 20 min. After 20 min incubation, 15 µl of trichloroacetic acid (10 %) was added in the reaction mixture and mixed well. Distilled water (55 µl) and ferric chloride (110 µl, 10 % w/v) were added, and the absorbance was measured at 700 nm. Increased absorbance of the reaction mixture indicates increased reducing capability. L-ascorbic acid was taken as a reference standard. All experiments were carried out in triplicates.

**Statistical Analyses**

All the assays were performed in triplicate, and the results were expressed as mean values with standard deviations (SD). The significant differences between means represented by letters were obtained by one-way analysis of variance (ANOVA) followed by Tukey’s honestly significant difference (HSD) post hoc test (P < 0.05). Correlations were calculated on a honey mean basis, according to Pearson’s test.

**RESULTS AND DISCUSSION**

In the present investigation, after surface sterilization of leaves from *Plectranthus amboinicus* different endophytic fungi were isolated on SDA medium respectively. A total of five endophytic fungal isolates were classified into five different taxa namely *Aspergillus sp.*, *Fusarium sp.*, *Acremonium sp.*, *Trichoderma sp.* and *Penicillium sp.* based on their microscopic and morphological appearance. Similar species were also reported from the same plant by34,35,36, where host relevance specific to host was observed such as *Aspergillus terreus* and *Penicillium* sp. Further, all the isolates were subjected to antibacterial activity against four foodborne pathogens. The preliminary screening results of antibacterial activity showed that the tested endophytic fungi *Aspergillus sp.*, *Fusarium sp.*, *Acremonium sp.*, *Trichoderma sp.*, and *Penicillium sp.* exhibit inhibition to all the test pathogens (Table 1) and revealed significant differences among the fungal isolates (P < 0.05).

**Table 1: Preliminary antibacterial activity screening of fungal endophytes**

<table>
<thead>
<tr>
<th>Fungal Isolates</th>
<th>Zone of inhibition*</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td><em>Escherichia coli</em></td>
</tr>
<tr>
<td><em>Aspergillus sp.</em></td>
<td>8.0 ± 0.0d</td>
</tr>
<tr>
<td><em>Fusarium sp.</em></td>
<td>9.0± 0.3c</td>
</tr>
<tr>
<td><em>Acremonium sp.</em></td>
<td>8.0 ± 0.0</td>
</tr>
<tr>
<td><em>Trichoderma sp.</em></td>
<td>10.0 ± 0.1b</td>
</tr>
<tr>
<td><em>Penicillium sp.</em></td>
<td>8.0 ± 0.0d</td>
</tr>
<tr>
<td>Streptomycin (1mg/ml)</td>
<td>20.0 ± 0.3a</td>
</tr>
</tbody>
</table>

*Values are means of three independent replicates ± standard deviation; Means in the same column with different letters are significantly (P < 0.05) different.

The isolates which showed antibacterial activity in preliminary screening were further subjected for fermentation assay and the crude extracts were obtained through ethyl acetate extraction. The obtained crude extract was subjected to secondary screening and showed a better zone of inhibition against tested pathogens (Table 2) compared to the preliminary assay. A similar study was found by37 on *Basella rubra* L. This could be due to the antimicrobial potency of the extract to the high concentration of unidentified active principle in the extract. The crude extracts of endophytic fungi yielded more potent compounds once they had undergone some purification38.

**Table 2: Secondary antibacterial screening of fungal endophytes**

<table>
<thead>
<tr>
<th>Fungal Isolates</th>
<th>Zone of inhibition*</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td><em>Escherichia coli</em></td>
</tr>
<tr>
<td><em>Aspergillus sp.</em></td>
<td>10.0 ± 0.0c</td>
</tr>
<tr>
<td><em>Fusarium sp.</em></td>
<td>10.0± 0.6c</td>
</tr>
<tr>
<td><em>Acremonium sp.</em></td>
<td>9.0 ± 0.0d</td>
</tr>
<tr>
<td><em>Trichoderma sp.</em></td>
<td>11.0 ± 0.1b</td>
</tr>
<tr>
<td><em>Penicillium sp.</em></td>
<td>11.0 ± 0.0b</td>
</tr>
<tr>
<td>Streptomycin (1mg/ml)</td>
<td>25.0 ± 0.3a</td>
</tr>
</tbody>
</table>

*Values are means of three independent replicates. ± Standard deviation; Means in the same column with different letters are significantly (P < 0.05) different.
The crude extract Aspergillus sp. was selected to determine the MIC and MBC by microdilution technique against E. coli, S. typhi, Sh. sonnei and E. faecalis (Table 3) based on the good zone of inhibition against all tested pathogens in secondary antibacterial screening. The result revealed 10mg/ml low concentration of MIC and 40mg/ml high concentration of MBC. The observed variations are due to the synergistic effect of the different congener components within one sample\(^5\). Upon serially diluting the content of the well where no resazurin color change was observed.

**Table 3: MIC and MBC (mg/ml) values of crude Aspergillus extract against foodborne pathogens.**

<table>
<thead>
<tr>
<th>Fungal Isolates</th>
<th>E. coli</th>
<th>S. typhi</th>
<th>Sh. sonnei</th>
<th>E. faecalis</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>MIC</td>
<td>MBC</td>
<td>MIC</td>
<td>MBC</td>
</tr>
<tr>
<td>Aspergillus sp.</td>
<td>10</td>
<td>20</td>
<td>15</td>
<td>10</td>
</tr>
</tbody>
</table>

The presence of phytochemicals in endophytes is an indicator that they can be a potential source of precursors in the development of synthetic drugs\(^44\). Phytochemical analysis of ethyl acetate solvent extract revealed the presence of phenol, flavonoids, alkaloids, terpenoids, steroids, saponin, tannins, and glycosides. The crude fungal extracts gave a wide variety of presence and absence of phytochemicals given in Table. 4.

**Table 4: Phytochemical screening of Aspergillus terreus ethyl acetate extract.**

<table>
<thead>
<tr>
<th>Fungal Isolates</th>
<th>Phenol</th>
<th>Flavonoids</th>
<th>Alkaloids</th>
<th>Terpenoids</th>
<th>Steroids</th>
<th>Saponin</th>
<th>Tannins</th>
<th>Glycosides</th>
</tr>
</thead>
<tbody>
<tr>
<td>Aspergillus sp.</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>-</td>
</tr>
<tr>
<td>Fusarium sp.</td>
<td>-</td>
<td>+</td>
<td>-</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>Acremonium sp.</td>
<td>+</td>
<td>-</td>
<td>+</td>
<td>-</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>+</td>
</tr>
<tr>
<td>Trichoderma sp.</td>
<td>+</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>Penicillium sp.</td>
<td>+</td>
<td>-</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>+</td>
<td>+</td>
<td>-</td>
</tr>
</tbody>
</table>

+: presence, -: absent.

The endophytes which revealed the presence of different phytochemicals are known to possess strong antimicrobial and antioxidant activity. The antioxidant capacities of the endophytic fungal cultures were correlated with their total phenolic contents, suggested that phenolic were also the major antioxidant constituents of the endophytes\(^46\). The ethyl acetate extract of Aspergillus sp. revealed good phenolic content as well as radical scavenging activity and suggested phenolic content represent the major group of compounds that act as primary free radical scavengers. Total phenolic content was found to be 5.08 ± 0.6 expressed in gallic acid equivalents (GAE) mg/ml and the antioxidant potential was evaluated by DPPH and FRAP assay. The effect of antioxidants on DPPH is thought to be to their hydrogen donating ability\(^42,43\) where freshly prepared DPPH solution exhibits a deep purple color which disappears when an antioxidant is present in the medium indicating that the metabolites of Aspergillus sp. could be potential agents in scavenging free radicals and treating diseases related to free radical reactions. The FRAP assay measures antioxidants in a sample at 593 nm compared to other assays measuring inhibition of free radicals exhibit 68.13 ± 0.26 (AAE mg/ml) indicating their electron-donating potential. Where, the value of FRAP is directly proportional to its antioxidant capacity, as in this assay, antioxidants act as reducing agents by donating electrons to ferric ion. The results obtained in the DPPH and FRAP assay indicated high activity (Figure. 1).

**Figure 1: Antioxidant activity of Aspergillus terreus ethyl acetate extract**

A significant positive correlation between total phenolic content and antioxidant activity was observed in our study (Table. 5). A positive correlation was found between FRAP and TPC, DPPH and FRAP was R= 0.942 and R= 0.975 respectively. The correlation between TPC and DPPH free radical scavenging activity was also studied and it was observed to be highly positively correlated with R = 0.987. The antioxidant activity of phenolics can be interpreted due to the increase with which they can act as reducing agents, hydrogen donors or singlet oxygen quenchers\(^44\). These results indicate that extract having more phenolic content tends to serve a strong antioxidant activity. Earlier studies on the screening of endophytic extracts for antioxidants from medicinal plants have revealed the chemical diversity of phenolic compounds from endophytes\(^45\).
Table 5: Correlation matrix (Pearson’s correlation coefficients)

<table>
<thead>
<tr>
<th>Variable</th>
<th>TPC</th>
<th>FRAP</th>
<th>DPPH</th>
</tr>
</thead>
<tbody>
<tr>
<td>DPPH</td>
<td>0.987**</td>
<td></td>
<td></td>
</tr>
<tr>
<td>FRAP</td>
<td>0.942**</td>
<td></td>
<td></td>
</tr>
<tr>
<td>TPC</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

** Correlation is significant at the 0.01 level

The study on synergistic antidiabetic activity was performed to detect the inhibition by alpha-amylase assay. Alpha-amylase is an enzyme that hydrolyses carbohydrates into glucose, and easily absorbable. The crude extract Aspergillus sp. isolates in different concentrations of 50, 100, and 200 µg/ml gave a potent antidiabetic activity. The antidiabetic activity in the concentrations 50, 100, 200 µg/ml was 68.12%, 34.18% and 75.31% respectively. The highest antidiabetic activity was observed in the concentration of 200 µg/ml (Figure 2). This study reveals that endophytic fungi inhibit the activity of alpha-amylase could be a potential source of developing remarkable antidiabetic drugs. These drugs could control diabetes by inhibiting a major enzyme alpha-amylase that hydrolyzes carbohydrates into sugar.

Figure 2: Showing amylase activity in different concentration of Aspergillus terreus ethyl acetate extract

Finally, Aspergillus sp. (Figure. 3) were considered for molecular characterization. The rDNA-ITS region was amplified, sequenced, and submitted to the NCBI GenBank with accession number MW391576. A BLAST search of ITS gene sequences reveals the endophytic fungal isolates to be the closest homolog of Aspergillus terreus (Figure. 4). To the best of our knowledge, this study considers a first report conducted on anti-diabetic and antioxidant activity of ethyl acetate extract of endophytic fungi isolated from Plectranthus amboinicus revealed both hosts, as well as its endophyte, are good sources of phytochemicals, antioxidants, antibacterial and antidiabetic activity. Further, elucidating the profile would reveal the presence of similar compounds both in host and endophyte.

Figure 3: Growth pattern A-Front view; B-Back view and microscopic view of endophytic fungi Aspergillus terreus.

Figure 4: Evolutionary positions of the endophytic fungal isolate E1 with other related fungal species based on internal transcribed spacer sequence similarity.
CONCLUSION

The study concludes that *Plectranthus amboinicus* harbor several endophytic fungi that showed their promising biological activity. The *Aspergillus terreus* isolate exhibited a promising antibacterial activity with a quite low MIC and high MBC value by exposing the bactericidal nature of its extract. The extract of *Aspergillus terreus* also revealed good anti-diabetic activity. Therefore, the anti-diabetic compounds can be isolated for a potential treatment for DM. The study suggests *Aspergillus terreus* has a multi biological activity and could be a promising therapeutic agent. These findings will facilitate further studies to gain a better understanding and production of bioactive metabolites.

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