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Antimicrobial, anti-cancer, anti-filarial and anti-inflammatory activities of Cowabenzophenone A extracted from the endophytic fungus *Aspergillus terreus* isolated from a mangrove plant *Bruguiera gymnorrhiza*

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ABSTRACT

Cowabenzophenone A was isolated from an endophytic fungus *Aspergillus terreus* isolated from a mangrove plant *Bruguiera gymnorrhiza*. The structure was determined by analysis of 1D and 2D NMR spectra and mass spectrometric data as a tetracyclo[7.3.3.3.3,11.03,7]tetradecane-2,12,14-trione skeleton. When the compound was tested in vitro for its ability to inhibit inflammations, α -glucosidase inhibitory activities and its antimicrobial properties, it showed an anti-inflammatory activity amounting to an IC₅₀ of 12.1 μ g/mL, α -glucosidase inhibitory activity, with IC₅₀ values of 7.8 ± 0.5 μ M, and antibacterial activity against *Bacillus subtilis* TISTR 088, and *Bacillus cereus* TISTR 688 with MIC values of 1 μ g/mL and 2 μ g/mL respectively. The compound showed cytotoxicity against HCT 116 colon cancer cell line with an IC₅₀ value of 10.1 μ M, and also showed a considerably high potential towards anti-filarial activities by resulting MIC, IC₅₀ and LC₅₀ values of 0.358 ± 0.02 mg/mL, 0.708 ± 0.021 mg/mL and 3.89 ± 0.18 mg/mL, respectively, in comparison to the standard drug Ivermectin (IVM).

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Introduction

There is a need to search for novel drug-leads to fight diverse complications in health, because infectious diseases are still a global problem due to development and spread of drug-resistant pathogens (Anaissie et al. 1996). Novel anti-cancer drugs leads are in massive demand due to high worldwide mortality (Parkin et al. 1999). Over the last few decades, endophytic fungi have been recognised as a rich source for pharmacologically active metabolites. Sri Lanka has a high rate of endemic speciation in both plants and microbes (Gunatilleke and Gunatilleke 1990), which offers great potential for the discovery of plant-associated microbes that can produce new natural products with medicinal or agricultural importance (De Silva et al., 2011; Chandra 2012).

On this scheme to identify and isolate novel drug leads, our research team had successfully obtained many compounds from endophytic fungi resident in terrestrial and aquatic plants. Biological activities associated with mycoleptodiscin B, isolated from the endophytic fungus *Mycoleptodiscus* sp., (Dissanayake et al. 2016b) chaetoglobosin A and C, produced by the endophytic fungus *Chaetomium globosum*, (Dissanayake et al.

2016a) equisetin, from endophytic *Fusarium* sp. and solanionic acid, an antibacterial-degraded steroid produced in culture by the fungus *Rhizoctonia solani* are some of them. (Ratnaweera et al. 2015b)

As a part of this ongoing programme aimed at discovering bioactive secondary metabolites of endophytic fungi obtained from Sri Lankan mangrove plants (Centko et al. 2014), the extracts of laboratory cultures of the endophytic fungi isolated from *Bruguiera gymnorrhiza* were investigated.

Mangroves are the most extensively distributed habitat in the Jaffna lagoon area and are found along lagoon borders, estuaries and along the coastline of islands (Myers et al., 2000; Mittermeier et al. 2005). Two types of mangrove communities are recorded in this area, namely riverine mangroves and fringing mangroves. A total of 13 true mangrove species and 18 mangrove-associated species have been reported from this lagoon and neighbouring islands. (Karunathilake 2003)

Bruguiera gymnorrhiza belongs to the *Rhizophoraceae* family. Trees would grow up to about 10 m in height and possess knee roots but lack prop roots. New growth of green leaves becoming grey, glossy and hairless. (Perera and Amarasinghe 2018)

Fungi of the genus *Aspergillus* were reported to produce a high variety of secondary metabolites, exhibiting diverse and remarkable biological properties such as antifungal, anti-tubercular, anticancer, antibacterial and farnesyl protein transferase inhibitory activities (Coles et al. 2002).

Therefore, as a promising source of naturally bioactive metabolites, *Aspergillus terreus* has attracted much attention. *Aspergillus terreus* has usually been found as a saprophytic fungus, producing mycotoxins such as patulin and cytochalasin E (Dong et al. 2008). Different species of *Aspergillus* grown in different media had given rise to distinct types of compounds. Extracts of *Aspergillus terreus* grown in liquid culture had given rise to several pyrones such as ampelopyrone, sulphated derivatives of macrosporin and 3-O-methylalaternin (Yates et al. 2012). In addition to these compounds, lactones such as methyltriacetic lactone had also been isolated (Yeasmin and Shamsi 2013). Previous results, as well as the findings in the present investigation, indicate that fungus *Aspergillus terreus* as an interesting source for new bioactive metabolites.

Materials and methods

Collection of plant material

Fresh healthy leaves, stem, flowers, bark and fruits of *Bruguiera gymnorhiza* were collected from Jaffna lagoon, Northern Province, Sri Lanka (90 50' 0 N and 790 50' 0 E to 90 20' 0 N and 800 30' 0 E; 1,087,208 N and 372,066 E to 1,031,744 N and 445,094 E). The collection was made in October 2014, transported to the laboratory in a tightly sealed Zip bag and stored at room temperature (30°C) under humid conditions. The identity of the plant was confirmed and authenticated by comparison with the voucher specimen in the National Herbarium at Royal Botanical Gardens, Peradeniya, Sri Lanka. The voucher specimen (No. UOC/NPL/KUBG12) of the plant used for the current study was deposited in the Pathology Laboratory, Department of Plant Sciences, University of Colombo, Sri Lanka. Collected plant parts were used for isolation of endophytic fungi within 16 h after collection.

Isolation of endophytic fungi from the collected plant specimens

Samples were cleaned under running tap water and then air-dried. Plant parts were cut into approximately

5 cm long pieces, and the parts of the leaves, flowers, roots and stem bark were surface sterilised by immersing in 70% ethanol for 1 min, 5% sodium hypochlorite solution for 1 min and sterile distilled water for 1 min. The surface sterilised leaves, flowers, roots and stem bark were cut into 0.5 cm² pieces using a sterile blade and placed separately on potato dextrose agar (PDA) medium under sterile conditions and incubated at 30°C for 21–27 days until the emergence of hyphae. Isolation of pure cultures was achieved from the original growths following serial inoculation method.

Identification of fungal isolates

Isolation of genomic DNA from the mycelia of each fungal isolate was carried out according to the published protocol of (Kariyawasam et al. 2012). The ITS region of the isolated genomic DNA was amplified by polymerase chain reaction (PCR) using the forward primer ITS 1 (5'TCCGTAGGTGAACCTGCGG3') and the reverse primer ITS 4 (5'TCCTCCGCTTATTGATATGC3') under the conditions, initial denaturation of 5 min at 94°C, followed by 35 cycles of 30 s at 94°C, 1 min at 55°C and 2 min at 72°C, with a final extension of 7 min at 72°C (Diaz et al. 2012). The amplified DNA was sequenced and BLAST analysed [National Centre for Biotechnology Information (NCBI)]. The acquired gene sequences were submitted to the NCBI GenBank database and accession numbers were obtained.

Extraction of crude extracts

Each single endophytic fungal isolate was grown in small scale on PDA (15 plates – 100 × 15 mm). The solid medium with the mature fungi was cut into small pieces and extracted into ethyl acetate. The ethyl acetate extract was concentrated by evaporating the solvent under reduced pressure at room temperature to obtain crude organic extracts.

Antimicrobial activities of the crude extracts

The antibacterial and antifungal activities of the crude extracts were determined using disc-diffusion assays. Selected fungus (KUBGACJ-14) which expressed the highest antimicrobial activity was grown in large scale, on sterile PDA in 600 petri dishes (size, 100 mm × 120 mm) for 16 days at 30°C, and this culture was

extracted thrice with EtOAc (3 x 2 L). The combined extracts were filtered and were concentrated *in vacuo*.

Crude extracts were assayed for antimicrobial activity against Gram-positive bacteria, *B. subtilis* (UBC 344), *S. aureus* (ATCC 43,300) and methicillin-resistant *S. aureus* (MRSA, ATCC 33,591) at 10 µg/disc and two Gram-negative bacteria, *Escherichia coli* (*E. coli*) (UBC 8161), *P. aeruginosa* (ATCC 27,853) at 100 and 25 µg/disc. The crude extracts were qualitatively assayed against pathogenic fungi *C. albicans* (ATCC 90,028), *F. oxysporum* (UBC 3512), *R. microsporus* (UBC 6314), *C. gloeosporoides* (UBC 9312) and *A. niger* (UBC 6514) at 50.0 µg/disc.

The major compound in the extract was assayed for antimicrobial activity against the same Gram-positive, Gram-negative bacteria and pathogenic fungi except for *F. oxysporum* (UBC 3512), *R. microsporus* (UBC 6314) and *A. niger* (UBC 6514) to obtain the minimum inhibitory concentrations (MICs). The MIC values were determined using broth microdilution method according to National Committee for Clinical Laboratory Standards 2003. The MIC was taken as the lowest concentration with more than 90% growth inhibition by measuring optical density. The commercial antimicrobial agents, polymyxin B for *B. subtilis*, *E. coli* and *P. aeruginosa*, rifamycin for *S. aureus* and MRSA, amphotericin for *C. albicans* and *C. gloeosporioides* were used as positive controls.

Isolation and structure elucidation of bioactive compounds

The crude extracts were next subjected to bioassay-guided fractionations in order to isolate the active compounds. The fractionation schemes were preceded from size exclusion chromatography (LH20) and polarity-based separation techniques (Silica columns and C18 Sep pak) by confirming the consistency of bioactivity using the bioautography technique. After every separation scheme, each individual extract was further guided by the examination of their thin layer chromatography (TLC) behaviour. Structure elucidation of the isolated active compound was done by using spectroscopic data obtained from UV, IR, and NMR (1D & 2D) spectroscopy.

To isolate the principal bioactive component from the complex mixture of the fungal extract, an advanced series of bioassay-guided purification steps were performed. The crude extract (720 mg) was first subjected

to solvent/solvent partitioning between hexane and MeOH/H₂O, 7:1 (1:1 300 mL of each) and left for 12 h; after the separation of the hexane layer, the polarity of the aqueous layer was increased to MeOH/H₂O, 2:1 by addition of H₂O and extracted with CHCl₃ (300 mLx2). The CHCl₃ layer was separated and the aqueous layer was concentrated at reduced pressure and was partitioned between H₂O and EtOAc (200 mLx3). All four fractions (MeOH, H₂O, EtOAc and CHCl₃) were tested for bioactivity and continued with the combined CHCl₃ and EtOAc soluble fractions which had retained bioactivity. Next, the combined fraction (91 mg) was eluted through a Sephadex LH-20 size exclusion chromatography (2.5 x 175-cm column) with MeOH/CHCl₃; 5:4 as the eluting solvent. Resulting fractions were combined according to the thin layer chromatography (TLC) profiles and the combined fractions were tested for antibacterial/antifungal activity. The most active fraction (27 mg) was run on a normal phase silica (2.5 x 90-cm) column using gradient elution (10% to 60% MeOH: CH₂Cl₂). The resulting active fraction (21.2 mg) was then run on Waters Sep pak C18 (2 g) reversed-phase cartridge (70%: 30% H₂O: MeOH) to further purify the active component. Finally, the active fraction (11 mg) was purified by C18 reversed-phase high-performance liquid chromatography (HPLC) using a CSC-Inertisil 150A/ODS2, 5µm 25 x 0.94 cm column with 1:3 MeCN/H₂O as an eluting solvent with a flow rate of 2 mL min⁻¹ to yield the major compound.

The melting point was measured on a Buchi melting point B-540 visual thermometer. The optical rotations were measured with a Bellingham-Stanley APD440 polarimeter. The UV spectra were recorded with Perkin Elmer UV-vis or Varian Cary 5000 UV-vis-NIR spectrophotometers. Infrared (IR) spectra were recorded on a PerkinElmer FTS FT-IR or Perkin Elmer Frontier Optica FT-IR spectrophotometers. ECD spectra were recorded on a JASCO J-815 spectrometer. The 1D and 2D NMR spectra were recorded on a 400 MHz Bruker FT-NMR Ultra Shield and 600 MHz Bruker AV-600 spectrometers in CDCl₃ and/or acetone-d₆ (CO) 206.2 and (CH₃) 30.6, with TMS as the internal standard. Chemical shifts are reported in parts per million (δ), and coupling constants (J) are expressed in hertz. ESI-QIT-MS and HR-El-MS spectra were measured on a Bruker-Hewlett-Packard 1100 Esquire-LC and a MAT 95 XL mass spectrometer, respectively. HPLC was performed on a Waters 515 HPLC pump system liquid chromatography using the

following columns: RP C18 CSC-Inertsil 150A/ODS-2 column (25 × 0.94 cm) and a CHIRALCEL OD-H column (4.6 × 250 mm). Quick column chromatography (QCC) and column chromatography (CC) were carried out on silica gel 60 H (5 – 40 µm, SiliCycle Inc.) and silica gel 100 (63 – 200 µm, SiliCycle Inc.), respectively. Sephadex LH-20 was also used for CC in determining elutes of different size fractions. Precoated plates of silica gel 60 F254 were used for TLC analysis.

Screening for anti-inflammatory activity

For cell-based THP-1 cytokine-release assay; the human monocytic cell line THP-1 (American Type Culture Collection, Manassas, Va.) was maintained in Roswell Park Memorial Institute, USA (RPMI) supplemented with 2 mM of L-glutamine, 100 U/mL penicillin, 100 mg/mL streptomycin, 25 µL of 12% (v/v%) 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid, as the zwitterionic buffering agent (HEPES), and 10% (v/v%) foetal bovine serum (FBS). Induction of cell differentiation was done with 100 µL of 18% (v/v%) para-methoxyamphetamine (PMA) for 24 h. After incubation, non-adherent cells were removed by aspiration, and the adherent cells were washed with RPMI three times. For cell stimulation, the cells were further incubated both with and without lipopolysaccharide (LPS) for 24 h in fresh complete medium with 10% FBS. After cell plating, the test compounds dissolved in 0.5% DMSO were added to each well, and the plate was incubated for 30 min at 37.8°C. Finally, 20 mL (10mg/mL) of LPS per well was added, to obtain a final concentration of 1 mg/mL.

Blood was withdrawn from human volunteers and aliquoted to 96-well plate (50 µL each). Immunomodulator was then added (10 µL), incubated at 37°C, 5% O₂ for 15 min before the addition of 10 µL *E. coli* (final concentration 10⁵ cells/mL). They were then incubated for 7 h, followed by the addition of 100 µL of PBS. After mixing, the plate was centrifuged at 1200 rpm for 5 min, and the supernatant was obtained and used for IL6 analysis. (Ukwatta et al. 2019)

Detection of secreted cytokines – A glucosidase activities

Human TNF-α and IL-6 were quantified by an enzyme-linked immunosorbent assay (ELISA) (Ukwatta et al.,

2019). Briefly, 96-well ELISA plates (Maxisorp; Nunc, Naperville, Ill.) were coated with an antihuman TNF-α/IL-6 monoclonal antibody in a coating buffer (carbonate/bicarbonate buffer, pH 9.6), followed by overnight incubation at 48°C. The wells were blocked for 2 h, at room temperature with 10% FBS prepared in assay buffer. Biotinylated anti-human TNF-α/IL-6 polyclonal antibody was added, followed by avidin-horseradish peroxidase conjugate, which used tetramethylbenzidine as the substrate. The reaction was stopped by the addition of 2M H₂SO₄, and optical density was recorded in a Sapphire microplate reader (Tecan) at 490 to 600 nm.

Cytotoxic assay

HCT116 colon cancer cells (1 × 10⁴ cells/well) were cultured in a 96-well plate and allowed to adhere for 24 h at 37°C. The cells were treated with testing compounds (10 µM or µg/mL) in Dulbecco's modified eagle's medium (DMEM) for 24 h. The medium was removed and fresh DMEM containing 0.5 mg/mL of 3-(4,5-Dimethylthiazol-2-Yl)-2,5-Diphenyltetrazolium bromide (MTT) solution was added to each well for 2 h. After that, the medium was discarded by an aspirator. The violet formazan crystals in the viable cells were dissolved in 100 µL of DMSO. The absorbance of each well was read at a wavelength of 570 nm using a microplate reader. Doxorubicin was used as a positive control with an IC₅₀ value of 9.74 µM. Percentage cell viability and percentage cytotoxicity were calculated by using the following equations.

$$\% \text{cell viability} = \frac{\text{absorbance of treated well}}{\text{absorbance of control well}} \times 100$$

$$\% \text{cytotoxicity} = 100 - \% \text{cell viability}$$

Screening for anti-filarial assay materials and mechanisms

Primary evaluation: *In vitro* assays: Based on viability of the parasites, *in vitro* motility and 3-(4, 5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) reduction assays were carried out using ursolic acid (UA), ivermectin (IVM) and diethylcarbamazine-citrate (DECC) as reference drugs. For incubation of microfilaria (mf) and adult worms cell culture plates (Nunc, Denmark) were used with Hank's balanced salt solution (HBSS) of pH 7.2.

The UA and IVM were dissolved in DMSO, whereas 2 mg of DECC was dissolved in 10 mL of sterile triple-distilled water (STDW). The anti-filarial agents were used at twofold serial dilutions ranged from 15.63–1000 mM (DECC), 1.56–100 mM (UA) and 0.31–20 mM (IVM). The final concentration of DMSO in the incubation medium was kept below 0.1% to ensure maintaining buffer capacity of test agent's solution as control assay.

Motility assay: Efficacy of UA and reference drugs was assessed *in vitro* on microfilaria and adult worms of *B. malayi* (as target parasites) using motility (microfilaria and adult parasite) and MTT (adult parasite only) reduction assay. Duplicate wells containing 40–50 mf/100 µl/well (of 96-well plates) and 1 female worm/mL/well (of 48-well plate) were used. UA (100 mM) or reference drugs IVM (20 mM), or DECC (1000 mM) were added accordingly based on the mf motility indices to duplicate wells and incubated. The wells with a test compound and DEC were incubated for 24 h and those with IVM were incubated for 24 and 48 h as it has a slow action on the parasites. All incubations were at 37°C in 5% CO₂ atmosphere. The effect on the motility of the parasite stages was examined under a microscope and scored. The experiment was repeated twice. In case of microfilaria, only motility assay was performed.

Motility assessment: Parasite motility was assessed under a microscope after 24/48 h exposure to test substance and scored as: 0 = dead; 1–4 = loss of motility (1 = 75%; 2 = 50%; 3 = 25% and 4 = no loss of motility). Loss of motility is defined as the inability of the worms to regain pretreatment level of motility even after incubating in fresh medium without the test agent at 37°C for 1 h, and was expressed as percentage (%) inhibition of control.

MTT (formazan colourimetric assay for the viability of worms): The same female worms used in motility were then gently blotted and transferred to 0.1 mL of 0.5% MTT in 0.01 M phosphate-buffered saline (pH 7.2) and incubated for 1 h at 37°C. The formazan formed was extracted in 1 mL of DMSO for 1 h at 37°C and its absorbance was measured at 510 nm. The mean absorbance value obtained for treated worms was compared with the controls. The viability of the treated worms was assessed by calculating the inhibition percentage of motility and MTT reduction over DMSO control worms.

Criteria for assessment of *in vitro* hits: 100% inhibition in the motility of female adults, mf and 50% inhibition in MTT reduction ability of female parasites were considered acceptable anti-filarial (microfilaricidal/adulticidal)

activity and picked up shifts and subjected to further testing *in vivo*.

Secondary evaluation – Determination of IC₅₀: (the concentration at which the parasite motility was inhibited by 50%). The parasites were incubated with twofold serial dilutions from 1.56 to 100 (UA), 0.31 to 40 (IVM) and 15.63 to 1000 mM (DECC) using triplicate wells of the cell culture plate. Experiments were run in duplicate and incubations were carried out in replicates for 24/48 h as above. After incubation, inhibition in motility (mf and female worm) and MTT reduction potential of the parasites were assessed as above. The experiment was repeated twice.

Determination of cytotoxic concentration 50 (CC₅₀): The cytotoxicity assay of the test substances was carried out as published in literature with some modifications (Ukwatta et al. 2019). Briefly, VERO Cell line C1008 (African green monkey kidney cells) was plated in 96-well plates (Nunc, Denmark) at 0.16106 cells/mL (100 µl per well) in DMEM supplemented with 10% heat-inactivated FBS. A three-fold serial dilution of the test substances (starting from .20 x LC100 conc. of the test agent) in buffer was added. The plates with a final volume of 100 µL/well were incubated in 5% CO₂ atmosphere at 37°C. After 72 h of incubation 10 µL of 0.025% Resazurin in phosphate-buffered saline (PBS; pH 7.2) was dispensed as indicator for viability followed by an additional incubation for 4 h and the plate was then read in a fluorescence reader (Synergy HT plate reader, Biotek, USA) at excitation wavelength of 530 nm and an emission wavelength of 590 nm. The assay was run in replicates in each of two independent experiments.

Data of IC₅₀ and CC₅₀ fed to statistical software (SPSS) and were analysed as described in the literature by linear interpolation between the two concentrations above and below 50% inhibition. (Paige et al. 2011)

Selectivity Index (SI) of the samples was computed by the formula as:

$$SI = \frac{CC_{50}}{IC_{50}}$$

Results and discussion

Isolation of endophytic fungi

A total of 29 morphologically distinct endophytic fungi, 18 from leaves, 04 from stems and 07 from

bark were isolated from the collected plant specimens.

Identification of the fungus

The endophytic fungus was identified as *Aspergillus terreus* based on the DNA sequence of the ITS ribosomal RNA gene. The closest accession number obtained when the sequences were deposited in the GenBank was KP131622.

Microbial activities of crude endophytic fungal extracts

The mean diameter ($\pm$ Standard error (SE)) of the inhibition zones of the active crude fungal extracts in the antibacterial and antifungal bioassays are shown in Tables 1 and 2 respectively. Four of the extracts were active against Gram-negative bacteria tested (KUBGACJ-07, KUBGACJ-11, KUBGACJ-14 and KUBGACJ-21) at 25 μ g/disc, while 09 fungal extracts inhibited the growth of at least one Gram-positive bacterial strains at 10 μ g/disc. As mentioned in the literature, (Hugo 1998) the lesser potential activity towards Gram-negative bacteria would be due to the rigid, stable and complex cell wall structures in them. Among them KUBGACJ-15, KUBGACJ-16 and KUBGACJ-27 were only active against *S. aureus* and other six extracts were active against *S. aureus*, MRSA and *B. subtilis*. KUBGACJ-14 showed antibacterial activity against Gram-positive and negative bacteria at 25 μ g/disc concentrations and also on 10 μ g/disc, respectively, further anti-fungal activity at 50 μ g/disc; thus, this fungus was selected for the isolation of bioactive metabolites after large scale culturing. Two crude

extracts, KUBGACJ-02 and KUBGACJ-05, exhibited antifungal activity against *F. oxysporum* (UBC 12,145), *R. microporus* (UBC 6314) and *C. albicans* (ATCC 90,028) only but KUBGACJ 14 showed antifungal activity against all tested five fungal strains by showing activities against *C. gloeosporioides* (UBC 3110) and *A. niger* (UBC 9214) as well. It was selected for sequential isolation of the bioactive compound. The three extracts that showed antifungal activity also showed activity against Gram-positive bacteria. Although antifungal activities of plant endophytic fungi have been reported occasionally, the occurrence of antifungal activities of plant endophytic fungi is not as common as antibacterial activities.

Isolation and structure elucidation of the major compound (KUBGACJ-14)

The bioassay-guided fractionation of the EtOAc extract led to the isolation of the active compound (4.3 mg) as a colourless crystalline solid eluting at a retention time of 37 min under the HPLC conditions (60%:40% H₂O:MeOH) used. This compound showed a molecular ion peak at $[M]^+m/z$ 568.3547 in HR-ESI-MS corresponding to the molecular formula of C₃₈H₄₈O₄ (Figure 1). Analysis of ¹H and ¹³C NMR data as well as 2D NMR (COSY, HSQC, HMBC, tROESY and HSQC ¹⁵N) spectral data in CDCl₃ revealed that the structure of the active compound to be cowabenzophenone. A matching with the data reported in the literature.²³

The UV spectrum of the compound had maximum absorbance readings at 207, 242, 282 and 308 nm, while the IR spectrum had absorption bands corresponding to three carbonyl groups (1736, 1703 and 1686 cm⁻¹). The basic skeletal

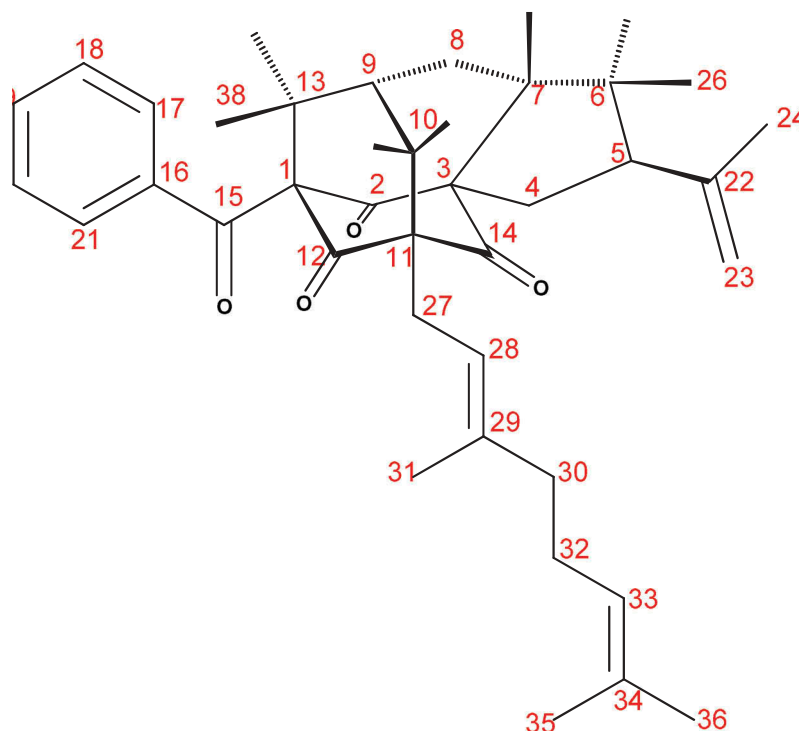
Table 1. Antibacterial activities of the crude extracts of endophytic fungi of the plant *Bruguiera gymnorrhiza*.

Fungal strain	Antibacterial activity – Gram-negative (mean diameter of the inhibition zone in mm $\pm$ SE)				Antibacterial activity – Gram-positive		
	<i>E. coli</i>		<i>P. aeruginosa</i>		MRSA	<i>B. subtilis</i>	<i>S. aureus</i>
	100 μ g/disc	25 μ g/disc	100 μ g/disc	25 μ g/disc	10 μ g/disc		
KUBGACJ-02	04.1 $\pm$ 0.1	-	05.1 $\pm$ 0.3	-	04.3 $\pm$ 0.2	02.1 $\pm$ 0.2	03.4 $\pm$ 0.1
KUBGACJ-05	05.2 $\pm$ 0.3	-	-	-	02.3 $\pm$ 0.3	07.3 $\pm$ 0.3	-
KUBGACJ-07	07.3 $\pm$ 0.4	05.3 $\pm$ 0.3	03.3 $\pm$ 0.2	02.3 $\pm$ 0.1	03.2 $\pm$ 0.1	04.3 $\pm$ 0.4	02.6 $\pm$ 0.1
KUBGACJ-11	07.9 $\pm$ 0.1	05.3 $\pm$ 0.4	04.3 $\pm$ 0.1	02.3 $\pm$ 0.2	05.3 $\pm$ 0.2	03.1 $\pm$ 0.2	02.1 $\pm$ 0.2
KUBGACJ-14	20.3 $\pm$ 0.2	16.3 $\pm$ 0.4	20.7 $\pm$ 0.1	18.4 $\pm$ 0.2	08.3 $\pm$ 0.4	06.3 $\pm$ 0.2	07.1 $\pm$ 0.4
KUBGACJ-15	09.3 $\pm$ 0.3	-	-	-	-	-	06.3 $\pm$ 0.3
KUBGACJ-16	05.4 $\pm$ 0.6	-	-	-	-	-	04.2 $\pm$ 0.1
KUBGACJ-21	16.3 $\pm$ 0.2	12.3 $\pm$ 0.4	14.1 $\pm$ 0.1	10.5 $\pm$ 0.2	04.2 $\pm$ 0.1	03.3 $\pm$ 0.4	03.4 $\pm$ 0.3
KUBGACJ-27	-	-	-	-	-	-	06.3 $\pm$ 0.4
+ Control (1 μ g/disc)	23.3 $\pm$ 0.3	23.6 $\pm$ 0.3	21.2 $\pm$ 0.4	23.4 $\pm$ 0.3	20.6 $\pm$ 0.3	21.6 $\pm$ 0.3	22.3 $\pm$ 0.4
- Control	-	-	-	-	-	-	-

Table 2. Antifungal activities (Qualitative)^a of the crude extracts (50 µg/disc) of endophytic fungi of the plant *Bruguiera gymnorrhiza*.

Fungal strain	<i>C. albicans</i>	<i>F. oxysporum</i>	<i>R. microporus</i>	<i>C. gloeosporioides</i>	<i>A. niger</i>
KUBGACJ-02	+	++	-	-	+
KUBGACJ-05	+	+	+	-	-
KUBGACJ-07	-	-	-	-	-
KUBGACJ-11	-	-	-	-	-
KUBGACJ-14	++	+++	++	+++	+++
KUBGACJ-15	-	-	-	-	-
KUBGACJ-16	-	-	-	-	-
KUBGACJ-21	-	-	-	-	-
+ Control (1 µg/disc)	++++	++++	++++	+++	+++
- Control	-	-	-	-	-

aKey – + – inhibition zone <3 mm, ++ – inhibition zone <5 mm, +++ – inhibition zone <7 mm, ++++ – inhibition zone <9 mm, (-) – No inhibition

**Figure 1.** Structure of cowabenzophenone A.

structure determination was performed using the COSY, HMQC and HMBC spectra (Table 3). The sequence of C-7 to C-10 was confirmed in the COSY spectrum by the cross-peaks. Thus, the structure of Figure 1 was identified as cowabenzophenone A.

Antimicrobial activity of cowabenzophenone a

Cowabenzophenone A showed strong selective antibacterial activities against Gram-positive *B. subtilis* (UBC 344), and *S. aureus* (ATCC 43,300) with MIC values of 1 and 2 µg/mL, respectively. It also showed promising activity against MRSA (ATCC 33,591), as 4 µg/mL.

From the MIC assay it showed activity against Gram-negative *E. coli* (UBC 8161) and *P. aeruginosa* (ATCC 27,853), as 4 and 2 µg/mL, respectively (Table 4). Cowabenzophenone A also showed potential activities against the pathogenic fungi *C. albicans* (ATCC 90,028) *C. gloeosporioides* (UBC 3110) and *A. niger* (UBC 9214), 4, 2 and 2 µg/mL, respectively. Polymixin B, amphotericin and rifamycin were used as positive controls and their respective activities are shown in Table 4.

Anti-inflammatory activities of cowabenzophenone a

This compound showed a value of 37% (results were expressed as percentages of IL6 relative to *E. coli*-

Table 3. ^1H and ^{13}C NMR data of cowabenzophenone A.

Position	Cowabenzophenone A		
	δ_{H}	δ_{C}	HMBC ($^1\text{H}^{13}\text{C}$)
1	-	82.5	-
2	-	203.8	-
3	-	72.4	-
4a	2.89 (t)	34.0	2, 3, 5, 14, 22
4b	2.07 (m)	-	2, 3, 5, 14
5	3.03 (dd)	56.0	6, 22, 23
6	-	44.6	-
7	2.20 (m)	55.0	2, 14
8a	2.07 (m)	22.6	10, 13
8b	1.76 (m)	-	3, 6, 7, 10, 13
9	2.07 (m)	42.7	1, 7, 11
10a	2.51 (dd)	35.1	8, 9, 11, 12, 27
10b	2.20 (m)	-	11, 12
11	-	67.9	-
12	-	204.5	-
13	-	47.5	-
14	-	205.0	-
15	-	193.2	-
16	-	135.2	-
17	7.07 (d)	128.6	15, 19, 20
18	7.27 (t)	128.5	16, 17
19	7.39 (t)	132.4	17, 18
20	7.27 (t)	128.5	16, 17
21	7.07 (d)	128.6	15, 19, 20
22	-	144.5	-
23a	4.90 (s)	113.2	5, 24
23b	4.77 (s)	-	-
24	1.77 (s)	23.7	22, 23
25	0.87 (s)	26.7	5, 6, 7, 26
26	1.01 (s)	26.7	5, 6, 7, 25
27	2.62 (m)	29.2	11, 12, 28, 29
28	5.29 (t)	119.1	27, 30, 31
29	-	139.1	-
30	2.07 (m)	40.2	28, 33
31	1.66 (s)	16.5	28, 29, 30
32	2.07 (m)	26.0	29, 30, 34
33	5.06 (m)-	124.3	32, 36
34	1.66 (s)	132.3	-
35	1.59 (s)	25.9	33, 34
36	1.39 (s)	17.8	33, 34
37	1.41 (s)	22.6	1, 9, 13
38	-	25.3	1, 9, 13

stimulated blood, therefore, lower the number showed greater anti-inflammatory activity.) This compound showed an IC_{50} of $12.1 \mu\text{g/mL}$.

Alpha-glucosidase activities

This compound showed an α -glucosidase inhibitory activity exhibiting an IC_{50} value of $7.8 \pm 0.5 \mu\text{M}$.

Cytotoxicity of cowabenzophenone a

This compound showed a cell viability value of 32%, resulting the cytotoxicity to be 68%. It showed cytotoxicity against HCT 116 colon cancer cell line with IC_{50} values of $10.1 \mu\text{M}$.

Anti-filarial activities of cowabenzophenone a

This compound showed considerably high potential anti-filarial activity due to its significantly low MIC, IC_{50} and LC_{50} values $0.358 \pm 0.02 \text{ mg/mL}$, $0.708 \pm 0.021 \text{ mg/mL}$ and $3.89 \pm 0.18 \text{ mg/mL}$, respectively, in comparison to the standard drug Ivermectin (IVM) (MIC, IC_{50} and LC_{50} values $3.12 \pm 0.78 \text{ mg/mL}$, $6.25 \pm 0.14 \text{ mg/mL}$ and $16.57 \pm 0.21 \text{ mg/mL}$, respectively) against microfilariae and adults. This compound showed a 50% inhibitive cytotoxic concentration value (CC_{50}) of 0.521 mg/mL . Therefore, the calculated selectivity index (SI) of this compound was 69.

Conclusions

Cowabenzophenone A, first described in December 2013, was isolated in a sequence of bioassay-guided fractionations, as a crimson crystalline solid. The respective structure elucidation was completed by spectroscopic techniques, which was shown to have a molecular formula of $\text{C}_{38}\text{H}_{48}\text{O}_4$. This is the first report of extraction of cowabenzophenone A, from a mangrove endophytic fungus *Aspergillus terreus* isolated from a *Bruguiera gymnorrhiza* mangrove plant.

This is also the first publication of antibacterial, anti-fungal, anti-inflammatory, anti-filarial, α -glucosidase inhibitory and cytotoxic activities of cowabenzophenone A. Due to the promising biological activities exhibited of inflammation and α -glucosidase inhibitions, the compound can be further investigated to develop potential drug leads.

Table 4. MIC values obtained for cowabenzophenone A and for positive controls.

Compound name	Minimum inhibitory concentration values (MIC) – $\mu\text{g/mL}$						
	<i>B. subtilis</i>	<i>S. aureus</i>	<i>E. coli</i>	MRSA	<i>C. albicans</i>	<i>P. aeruginosa</i>	<i>C. gloeosporioides</i>
Cowabenzophenone A	1	2	4	4	4	2	2
Polymixin B	4	-	1	-	-	1	-
Amphotericin	-	-	-	-	1	-	1
Rifamycin	-	1	-	1	-	-	-

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References

- Anaissie EJ, Paetznick VL, Ensign LG, Espinel-Ingroff A, Galgiani JN, Hitchcock CA, LaRocco M, Patterson T, Pfaller MA, Rex JH, et al. **1996**. Microbroth antifungal susceptibility testing of *Candida albicans* and *Cryptococcus neoformans* with and without agitation: an eight-center collaborative study. *Antimicrob Agents Chemother.* 40:2387–2391.
- Centko RM, Williams DE, Patrick BO, Akhtar Y, Chavez MAG, Wang YA, Isman MB, De Silva ED, Andersen RJ. **2014**. Dhilirilolides E–N, meroterpenoids produced in culture by the fungus *penicillium purpurogenum* collected in Sri Lanka: structure elucidation, stable isotope feeding studies, and insecticidal activity. *J Org Chem.* 79:3327–3335.
- Chandra S. **2012**. Endophytic fungi: novel sources of anticancer lead molecules. *Appl Microbiol Biotechnol.* 95:47–59.
- Coles BF, Chen G, Kadlubar FF, Radominska-Pandya A. **2002**. Interindividual variation and organ-specific patterns of glutathione S-transferase alpha, mu, and pi expression in gastrointestinal tract mucosa of normal individuals. *Arch Biochem Biophys.* 403:270–276.
- De Silva ED, Williams DE, Jayanetti DR, Centko RM, Patrick BO, Wijesundera RLC, Andersen RJ. **2011**. Ionization of carbon acids in liquid ammonia. *Org Lett.* 13:1174–1177.
- Diaz PL, Hennell JR, Sucher NJ. **2012**. Genomic DNA extraction and barcoding of endophytic fungi. In: Sucher NJ, Hennell JR, Carles MC, editors. *Plant DNA fingerprinting and barcoding: methods and protocols*. Methods in Molecular Biology. 862:171–179. doi:10.1007/978-1-61779-609-8_14.
- Dissanayake R, Ratnaweera PB, Williams DE, de Silva ED, Wijayarathne CD, Wijesundera RLC, Andersen RJ. **2016a**. Antimicrobial activities of endophytic fungi of the Sri Lankan aquatic plant *Nymphaea nouchali* and chaetoglobosin A and C, produced by the endophytic fungus *Chaetomium globosum*. *Mycology.* 7(1):1–8.
- Dissanayake RK, Ratnaweera PB, Williams DE, de Silva ED, Wijayarathne CD, Wijesundera RLC, Andersen RJ. **2016b**. Antimicrobial activities of endophytic fungi of Sri Lankan *Calamusthwaitesii* Becc. and mycoleptodiscin B, produced by the endophyte *Mycoleptodiscus* species. *J Appl Pharm Sci.* 6(1):22–28.
- Dong ER, Schulz B, Boyle C, Draeger S, Römmert A-K, Krohn K. **2008**. Endophytic fungi: A source of novel biologically active secondary metabolites. *Micol Res.* 106:996–1004.
- Gunatilleke IAUN, Gunatilleke CVC. **1990**. Distribution of Floristic Richness and Its Conservation in Sri Lanka. *Conserv Biol.* 4:21–31.
- Hugo WB. **1998**. Bacteria. In: Hugo WB, Russell AD, editors. *Pharmaceutical microbiology*. sixth ed. United Kingdom: blackwell Science Ltd; p. 4–8.
- Kariyawasam GK, Mithrasen YJPK, Fernando THPS, Wijesundera RLC, Wijesundera WSS. **2012**. A new cost effective method for extracting genomic DNA from fungi. Abstracts of papers, 5th Annual sessions, Institute of Biochemistry Molecular Biology and Biotechnology, Colombo. Colombo: Institute of Biochemistry, Molecular Biology and Biotechnology, 5:49.
- Karunathilake KMBC. **2003**. Status of Mangroves in Sri Lanka. *J Coastal Dev.* VII(1):5–9. ISSN: 1410-5217.
- Mittermeier RA, Gil RP, Hoffman M, Pilgrim J, Brooks T, CG M, Lamoreux J, Fonseca GAB. **2005**. Hotspots revisited: earth's biologically richest and most endangered terrestrial ecoregions. Boston (USA): University of Chicago Press.
- Myers N, Mittermeier RA, Mittermeier CG, Gustavo A, da Fonseca B, Kent J. **2000**. Biodiversity hotspots for conservation priorities. *Nature.* 403:853–858.
- Paige AL, Aly AH, Debbab A, Proksch P. **2011**. Fungal endophytes: unique plant inhabitants with great promises. *Appl Microbiol Biotechnol.* 90:1829–1845.
- Parkin DM, Pisani P, Munˆ Oz N, Ferlay J. **1999**. Estimates of the worldwide incidence of 25 major cancers in 1990. *Int J Cancer.* 80:827–841.
- Perera KARS, Amarasinghe MD. **2018**. Ecosystem carbon stock of Mangroves at the Batticaloa Lagoon, Sri Lanka. *Ousl J.* 13 (2):81–100.
- Ratnaweera PB, Williams DE, Patrick BO, de Silva ED, Andersen RJ. **2015b**. Solanolic acid, an antibacterial degraded steroid produced in culture by the fungus *Rhizoctonia solani* from tubers of the medicinal plant *Cyperus rotundus*. *Org Lett.* 17:2074–2077.
- Sriyatep T, Maneerat W, Sripisut T, Cheenpracha S, Machan T, Phakhodee W, Laphookhieo S. **2014**. Cowabenzophenones A and B, two new tetracyclo[7.3.3.3.3,11.03,7]tetrade-cane-2,12,14-trione derivatives, from ripe fruits of *Garcinia cowa*. *Fitoterapia.* 92:285–289.
- Ukwatta KM, Lawrence JL, Wijayarathna CD. **2019**. The study of antimicrobial, anti-cancer, anti-inflammatory and α -glucosidase inhibitory activities of Nigronephthalphenyl, isolated from an extract of *Nigrospora sphaerica*. *Mycology.* 10(2):1–10.
- Yates LA, Fleurdeˆ Pine S, Rissland OS, De Colibus L, Harlos K, Norbury CJ, Gilbert RJ. **2012**. Structural basis for the activity of a cytoplasmic RNA terminal uridylyl transferase. *Nat Struct Mol Biol.* 19:782–787.
- Yeasmin F, Shamsi S. **2013**. Phylloplanemycoflora of *Gerbera spp.* and their pathogenic potentiality. *J Bangladesh Aca Sci.* 37:211–217.