Secondary Metabolites from Endophytic Fungus
_Penicillium pinophilum_ Induce ROS-Mediated Apoptosis through Mitochondrial Pathway in Pancreatic Cancer Cells

**Abstract**

The endophytic fungus strain MRCJ-326, isolated from _Allium schoenoprasum_, which is also known as Snow Mountain Garlic or Kashmiri garlic, was identified as _Penicillium pinophilum_ on the basis of morphological characteristics and internal transcribed spacer region nucleotide sequence analysis. The endophytic fungus extract was subjected to 2D-SEBOX bioactivity-guided fractionation and purification. The anthraquinone class of the bioactive secondary metabolites were isolated and characterized as oxyksyrin (1), skyrin (2), dicatenarin (3), and 1,6,8-trihydroxy-3-hydroxy methylanthraquinone (4) by spectral analysis. Dicatenarin and skyrin showed marked growth inhibition against the NCI60/ATCC panel of human cancer cell lines with least IC_{50} values of 12 µg/mL and 27 µg/mL, respectively, against the human pancreatic cancer (MIA PaCa-2) cell line. The phenolic hydroxyl group in anthraquinones plays a crucial role in the oxidative process and bioactivity. Mechanistically, these compounds, i.e., dicatenarin and skyrin, significantly induce apoptosis and transmit the apoptotic signal via intracellular reactive oxygen species generation, thereby inducting a change in the mitochondrial transmembrane potential and induction of the mitochondrial-mediated apoptotic pathway. Our data indicated that dicatenarin and skyrin induce reactive oxygen species-mediated mitochondrial permeability transition and resulted in an increased induction of caspase-3 apoptotic proteins in human pancreatic cancer (MIA PaCa-2) cells. Dicatenarin showed a more pronounced cytotoxic/proapoptotic effect than skyrin due to the presence of an additional phenolic hydroxyl group at C-4, which increases oxidative reactive oxygen species generation. This is the first report from _P. pinophilum_ secreting these cytotoxic/proapoptotic secondary metabolites.

**Abbreviations**

<table>
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<tr>
<th>Abbreviation</th>
<th>Definition</th>
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<tr>
<td>ACN</td>
<td>acetonitrile</td>
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<tr>
<td>COSY</td>
<td>correlation spectroscopy</td>
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<td>DAPI</td>
<td>4',6-diamidino-2-phenylindole</td>
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<td>DCFH-DiACET</td>
<td>2',7'-dichlorofluorescein diacetate</td>
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<td>DMSO</td>
<td>dimethyl sulfoxide</td>
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<td>DNA-PK</td>
<td>DNA dependent protein kinase</td>
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<td>EDTA</td>
<td>ethylene diaminetetraacetic acid</td>
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<td>ELISA</td>
<td>enzyme linked immunosorbent assay</td>
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<td>ESIMS</td>
<td>electrospray ionization mass spectrometry</td>
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<td>HPLC</td>
<td>high-performance liquid chromatography</td>
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<td>HMBC</td>
<td>heteronuclear multiple bond correlation</td>
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<td>HRMS</td>
<td>high-resolution mass spectrometry</td>
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<td>ITS</td>
<td>internal transcribed spacer region</td>
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<tr>
<td>LC</td>
<td>liquid chromatography</td>
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<td>LC-MS</td>
<td>liquid chromatography mass spectrometry</td>
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<td>LC-MS/MS</td>
<td>liquid chromatography mass spectrometry/mass spectrometry</td>
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<td>LC-QTOF</td>
<td>liquid chromatography quadrupole time-of-flight</td>
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<tr>
<td>MTP</td>
<td>mitochondrial transmembrane potential</td>
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<tr>
<td>mTOR</td>
<td>mammalian target of rapamycin</td>
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<tr>
<td>MW</td>
<td>molecular weight</td>
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<tr>
<td>PBS</td>
<td>phosphate buffered saline</td>
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<tr>
<td>PCR</td>
<td>polymerase chain reaction</td>
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<tr>
<td>PDA</td>
<td>potato dextrose agar</td>
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<tr>
<td>PI</td>
<td>propidium iodide</td>
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<tr>
<td>rDNA</td>
<td>ribosomal DNA</td>
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<tr>
<td>ROS</td>
<td>reactive oxygen species</td>
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<td>UV</td>
<td>ultraviolet</td>
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<td>one dimensional nuclear magnetic resonance</td>
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<td>2D NMR</td>
<td>two dimensional nuclear magnetic resonance</td>
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Introduction

Natural products have been exploited world over in hope of finding new pharmaceuticals. Development of the pharmaceutical industry has stimulated the search for novel natural products from diverse environments and organisms. In this context, secondary metabolites of microbial origin deserve special consideration. Microorganisms are able to produce a variety of secondary metabolites including cytotoxic and antiproliferative compounds, which are valuable in cancer drug discovery. Particularly, natural metabolites such as secondary metabolites of microbial origin deserve special consideration. In this context, the pharmaceutical industry has stimulated the search for novel natural products from the metabolites of microbial origin. Substantial work has been carried out worldwide with regard to endophytism in the secondary metabolites of microbial origin.

Results

Isolated endophytic fungal strain MRCJ-326 from the bulb of Allium sp. (snow mountain garlic) has shown morphological distinct characteristic features like hypha and spores of the genus Penicillium. The corresponding ITS-rDNA sequence of the endophytic fungus was blasted in NCBI and revealed 100% sequence similarity with Penicillium pinophilum. In our initial screening, the ethyl acetate extract of P. pinophilum exhibited cytotoxicity against human pancreatic cancer (MIA PaCa-2) cells (Table 1). Further LC-MS/MS analysis of the ethyl acetate extract showed the presence of an anthraquinone class of compounds and identified a series of four compounds, oxyskyrin (1), skyrin (2), dicatenarin (3), and 1,6,8-trihydroxy-3-hydroxymethylanthraquinone (4). These compounds were further purified by detailed spectroscopic analysis. Compound 1 was obtained as an orange red powder. It showed UV absorption maxima at 251, 256, 300, and 462 nm, which indicated the anthraquinone-type moiety. The 13C NMR data of 1 (Table 1, Supporting Information) showed signals corresponding to 25 carbons. 1H NMR showed proton signals of one aromatic singlet at δ 7.66 and 7.51 ppm, and one oxymethylene at δ 4.87 ppm. The 13C NMR data showed signals corresponding to 25 carbons. 1H NMR showed proton signals of one aromatic methyl group resonating at δ 2.48 ppm, one oxygenaldehyde at δ 4.87 ppm, one aromatic singlet at δ 6.81 ppm (2H), two aromatic broad singlets at δ 7.66 and 7.51 ppm, and two aromatic doublets at δ 7.66 and 7.51 ppm. The 13C NMR data of 1 exhibited a total of 25 carbon peaks attributable to one methyl group at δC 22.1 (C-11), one oxygenaldehyde at δC 63.5 (CH2-OH), five methine carbons, and eighteen quaternary carbon atoms. Detailed analysis of the 13C spectrum disclosed signals for four carbon atoms and identified a series of four compounds, oxyskyrin (1), skyrin (2), dicatenarin (3), and 1,6,8-trihydroxy-3-hydroxymethylanthraquinone (4). These compounds were further purified by detailed spectroscopic analysis.
eighteen aromatic carbons at δC 120.8 (C-2), 126.6 (C-2′), 152.2 (C-3), 148.6 (C-3′), 120.1 (C-4), 122.0 (C-4′), 133.7 (C-4a), 132.3 (C-4a′), 124.2 (C-5/5′), 108.1 (C-7/7′), 107.2 (C-8a/8a′), 118.6 (C-9a), 115.1 (C-9a′), 133.3 (C-10a), and 132.2 (C-10a′). Further analysis of 1D together with 2D data revealed that we were dealing with a dimeric anthraquinone-like emodin. Comparison of NMR data and UV spectra of 1 with those of emodin, as well as consideration of the MW, suggested that 1 is a non-symmetrical anthraquinone including one emodin and one 1,6,8-trihydroxy-3-hydroxyanthraquinone unit. The 1H-1H COSY spectrum showed that the aromatic methyl signal δH 2.48 correlated to the meta-coupled protons H-2 and H-4. A methyl signal at δH 2.21 and a methoxyl group at δH 2.35 p.p.m., one aromatic singlet at δH 7.21 and 7.50 p.p.m., and two aromatic broad singlets at δH 6.63 p.p.m., and two aromatic broad singlets at δH 7.21 and 7.50 p.p.m. The 13C NMR data exhibited 15 carbon peaks attributable to one methyl group, three methine carbons, and eleven quaternary carbon atoms. Further analysis of the 13C spectrum disclosed signals for two carbonyl groups resonating at δC 190.1 (C-9/9′) and 186.1 (C-10/10′), three oxygenated aromatic carbon atoms at δC 162.0 (C-1/1′), 164.7 (C-6/6′), and 164.1 (C-8/8′), and nine aromatic carbons at δC 121.4 (C-2/2′), 149.1 (C-3/3′), 123.1 (C-4/4′), 134.2 (C-4a/4a′), 125.1 (C-5/5′), 107.3 (C-7/7′), 109.1 (C-8a/8a′), 113.1 (C-9a/9a′) and δC 131.5 (C-10a/10a′), and one methyl at 21.7 (C-11/11′). Comparison of the NMR data and UV spectra of 2, as well as consideration of the MW, suggested that 2 is a symmetrical dimer of two emodin units. The linkage between the emodin building blocks was confirmed by 2D NMR analysis, including 1H-1H COSY and HMBC correlations. The 1H-1H COSY spectrum showed that the aromatic methyl signal correlated to the meta-coupled protons H-2 and H-4. Furthermore, in the HMBC spectrum, the aromatic proton at δH 6.63 (H-7/7′) showed a strong correlation to a carbon resonating at δC 125.1, which was attributed to C-5/5′ (Fig. 3S, Supporting Information). The above data was fully comparable to the data of skyrin reported in the literature, and confirms the structure of 2 as skyrin [11, 12].

Compound 3 was obtained as an orange powder and had an identical pattern in UV absorption maxima to those of 1 and 2, isolated along with this compound. LC-HRMS exhibited a prominent peak at m/z 569.3 [M – H]− with a 34 mass unit compared with the MW of 2, which established the molecular formula C30H17O12 for 3. Comparison of the 1H NMR spectra of 3 and 2 showed a close relationship, except for the absence of one aromatic signal. Further analysis of the 1D and 2D NMR data of 3 and 2 confirmed that both share a similar skeleton and substituents, except for the additional presence of a free hydroxyl group at C-4 in 3. In analogy to 2, two isolated aromatic protons were detected in the 1H-1H COSY spectrum of 3. In addition, the HMBC spectrum confirmed the 13C assignments of all of the systems in the molecules (Table 1S, Supporting Information). In contrast to 2, the signal of a methyl group (δH 2.21, CH3-11) in 3 exhibited a strong J HMBC correlation to a carbon resonating at δC 159.9 p.p.m., which was attributed to C-4. Hence, an additional free hydroxyl group was connected to C-4. Thus, a planar structure of 3 was accomplished and confirmed as dicatenarin. The data was fully analogous to the data of dicatenarin reported in the literature [11].
Compound 4 was obtained as a yellow-orange needle with an m.p. between 285–290 °C and UV absorption maxima quite similar to those of 1–3. LC-HRMS exhibited a prominent peak at m/z 285.3 [M – H]− corresponding to the molecular formula C15H9O6. The 13C NMR (Table 1S, Supporting Information) spectrum of 4 showed a close relationship with 1. Further analysis of the 1D and 2D NMR data of 4 confirmed that it has the same skeleton and substituents as 1; however, it is a monomer, as revealed from the molecular formula. A careful literature search revealed that 4 is a well-known 1,6,8-trihydroxy-3-hydroxymethylanthraquinone previously isolated from Penicillium janthinellum [13].

Of the four isolated compounds from the ethyl acetate extract, two, dicatenarin and skyrin, showed significant cytotoxicity with least IC50 values of 12 and 27 µg/mL, respectively, against human pancreatic cancer (MIA PaCa-2) cells. However, oxyskyrin and 1,6,8-trihydroxy-3-hydroxymethylanthraquinone, a significant cytotoxic effect was not observed (Table 1).

Induction of apoptosis in MIA PaCa-2 cells was investigated by microscopic analysis of the DAPI-stained cells (Fig. 4). After 48 h of treatment, dicatenarin- and skyrin-treated MIA PaCa-2 cells demonstrated significant chromatin condensation and frag-
mentation within the nucleus in a concentration-dependent manner. However, chromatin condensation and fragmentation was more prominent in dicatenarin- compared to skyrin-treated cells. In paclitaxel-treated MIA PaCa-2 cells, significant chromatin condensation was observed, and no chromatin condensation was observed in untreated controls.

To determine the extent of apoptosis of dicatenarin- and skyrin-induced cell death in MIA PaCa-2 cells, the annexin V-FITC and PI dual staining was determined by laser scanning confocal microscopy. The data showed that untreated cells were negative for annexin V-FITC and PI staining, whereas positive cells were seen following both dicatenarin and skyrin treatment. The data also revealed that a significant amount of annexin V positive cells were seen in the case of dicatenarin, more than in skyrin, in the treated cells (Fig. 5).

We determined ROS generation is induced by dicatenarin and skyrin in MIA PaCa-2 cells. Both dicatenarin and skyrin increased ROS generation in a concentration-dependent manner after 48 h treatment, but the phenomenon of ROS generation was much higher in dicatenarin-treated cells compared to skyrin. In H$_2$O$_2$-treated (0.05%) MIA PaCa-2 cells, prominent ROS generation was observed, whereas in the untreated control, no ROS generation was observed (Fig. 6).

Loss of MTP ($\Delta \psi_m$) in MIA PaCa-2 cells was measured with a laser scanning confocal microscope. MTP loss has been seen considerably in MIA PaCa-2 cells when exposed to different concentrations of dicatenarin and skyrin after 48 h treatment as compared to untreated cells. Moreover, dicatenarin treatment induced significant MTP loss compared to skyrin. In paclitaxel-treated (1 µg/mL) MIA PaCa-2 cells, prominent MTP loss was observed (Fig. 7).

We further carried out immunofluorescence microscopic studies to determine the cytochrome c release from the mitochondria into the cytosol in MIA PaCa-2 cells during apoptosis. The distribution of cytochrome c was visualized by immunostaining the cells with anti-cytochrome c antibody. The localization of the mitochondria was revealed by staining the cells with MitoTracker Red. As we compared the distribution pattern of the endogenous cytochrome c and the morphological pattern of mitochondria, we found a prominent release of cytochrome c from the mitochondria in dicatenarin-treated MIA PaCa-2 cells, whereas skyrin-treated MIA PaCa-2 cells showed a mild cytochrome c release at higher concentrations after 48 h treatment. In doxorubicin (0.5 µg/mL), the release of cytochrome c was very prominent (Fig. 8).

We confirmed the participation of activated caspase-3 in the events of apoptosis in MIA PaCa-2 cells induced by dicatenarin and skyrin after 48 h treatment. Caspase-3 activity was shown to be significantly elevated in dicatenarin-treated MIA PaCa-2 cells in a concentration-dependent manner (Fig. 9). However, in comparison to dicatenarin, there was the least increment in caspase-3 activity in skyrin-treated MIA PaCa-2 cells, even at a higher concentration. In staurosporine-treated (0.5 µg/mL) MIA PaCa-2 cells, significant caspase-3 activity was observed.

The clonogenic assay revealed that both dicatenarin and skyrin treatment significantly induced a reduction in the colony formation ability of MIA PaCa-2 cells. However, the reduction was less observed in skyrin-treated cells compared to dicatenarin treatment. In untreated MIA PaCa-2 cells, overlapping colonies were observed, whereas no overlapping cells were observed following treatment of both of the compounds (Fig. 10).

Discussion

Endophytic fungi have been a promising source for bioactive compounds, especially their anticancer potential. As great demand arises for new drug leads for cancer, there arises a need to exploit the endophytic fungi associated with medicinal plants. In
In the present study, we collected A. schoenoprasum bulbs from snow mountain regions of the Western Himalayas for the isolation of endophytic fungi. In 1965, Chopra et al. reported an enhancement of seven times in the medicinal properties of the garlic species growing at higher elevations due to a high content of organosulphur compounds [14]. It is also known to have abundant biological activities including immunomodulatory and anticancer properties [15]. The endosymbiotic nature of many endophytic fungi is to inherit their bioactive secondary metabolite-producing capability from their host. Considering this a major reason, we selected it as a host plant [4,16] and isolated the endophytic fungus P. pinophilum (MRCJ-326) growing from its bulbs.

Endophytic fungal research is focused on Penicillium sp. for its secondary metabolites displaying various pharmacological effects including anticancer activity [17]. Previously, it has been reported that a multitude of anticancer agents have been produced from Penicillium sp., e.g., brefeldin A, wortmanin, and chloctans-
nases and related enzymes such as mTOR, DNA mannin, which is a specific inhibitor of phosphoinositide 3-kinase, which has also produced other steroid metabolites, i.e., wortmannin were merely killing the cells through cytotoxic action or...
also the release of cytochrome c, which recruits effector caspases leading to programmed cell death. For further analysis of the apoptotic pathway, the release of cytochrome c and the induction of caspase(s) was studied. Our immunofluorescence data revealed colocalization between mitochondria and cytochrome c. Cytochrome c was found to be highly colocalized with mitochondria in the untreated control cells. In contrast, MIA PaCa-2 cells exposed to dicatenarin and doxorubicin (positive control) treatment exhibited diffused cytoplasmic distribution of cytochrome c from the transition pores of the mitochondria into the cytosol, whereas skyrin-treated human pancreatic cancer (MIA PaCa-2) cells showed mild cytochrome c release at a higher concentration. Since it is a key event in initiating the caspase cascade involving caspase-3, we confirmed the participation of activated caspase-3 in the events of apoptosis induced by dicatenarin and skyrin. Caspase-3 activity was significantly elevated in dicatenarin-treated human pancreatic cancer cells (MIA PaCa-2). Comparatively, there is the least increase in caspase-3 activity in skyrin-treated MIA PaCa-2 cells after 48 h treatment. To assess the effect of the test material(s) on proliferation, growth, survival, and colonogenic ability, human pancreatic cancer (MIA PaCa-2) cells were treated with different concentrations of test material(s) for colony formation. Both of the compounds produced a concentration-dependent inhibitory effect on the ability of the cells to reproduce and form large colonies. The untreated control showed overlapping colonies, whereas no overlapping was observed in both of the compounds. Considering the true pharmacological potential of isolated and well-characterized secondary metabolites, i.e., dicatenarin (3) and skyrin (2) of P. pinophilum isolated from the bulb of snow mountain garlic (Allium sp.), further studies are required to mechanistically evaluate and optimize the identified leads for cancer.

In conclusion, we have isolated and characterized four compounds (1–4) from the endophytic fungus P. pinophilum (MRCJ-326) associated with the bulbs of A. schoenoprasum. All of the isolated compounds were tested against a panel of human cancer cell lines. As shown in Table 1, two compounds, dicatenarin (3) and skyrin (2), exhibited marked cytotoxic and proapoptotic potential, while compounds 1 and 4 were found inactive against the tested cell lines. Our results indicate that dicatenarin induced apoptosis through an apoptotic signal via intracellular ROS generation, and thereby a loss of MTP, and the induction of the mitochondrial transition pore further resulted in an increased expression of apoptotic effector protein caspase-3 for the execution of programmed cell death. In compounds 1 and 4, the methyl group at the C-3 position was oxidized to primary alcohol and converted into CH₂OH, which leads to the complete loss of its activity. These results also suggested that in 1,6,8-trihydroxy anthraquinone, the C-3 methyl group played a significant role in the cytotoxic potential as in skyrin and dicatenarin. Based on the bioactivity potential of these compounds, they can be studied further mechanistically and optimized for development as anticancer leads.

Materials and Methods

Chemicals and Reagents
Cetyltrimethylammonium bromide, deuterated solvents, and CD₃OD used in this study were obtained from Sigma-Aldrich.

Materials and Methods

Cells, cell culture, growth conditions, and treatment
The human lung cancer cell line (A549), human colon cancer cell line (HCT-116), human breast cancer cell line (T47D), human prostate cancer cell line (PC3), and human ovarian cancer cell line (OVCAR-3) were obtained from the National Cancer Institute (NCI). The human pancreatic cancer cell line (MIA PaCa-2) was obtained from American Type Culture Collection (ATCC). The human pancreatic cancer cell line (MIA PaCa-2) and human colon cancer cell line (HCT-116) were maintained in Dulbecco’s modified Eagle’s medium. The human lung cancer cell line (A549), human breast cancer cell line (T47D), human prostate cancer cell line (PC3), and human ovarian cancer cell line (OVCAR-3) were maintained in RPMI. The media was supplemented with fetal bovine serum (10%), penicillin (100 units/mL), streptomycin (100 µg/mL), and glutamine (2 mM). Cells were grown in CO₂ (Thermo Scientific) and all other chemicals were purchased from Sigma-Aldrich.

Isolation and identification of endophytic fungi
Plant samples, i.e., bulbs of A. schoenoprasum, were collected from the Western Himalayas from the snow mountain regions of Natha top, J&K, India. Samples were tagged, transported to the laboratory in sterile plastic bags, and stored at 4°C until processed for isolation of the endophytic fungi. Samples were identified taxonomically by Dr. Sumeet Gairola, the Herbarium and Plant Systematics Section, Plant Biotechnology Division, CSIR-IIIM, Plant Systematics Section, Plant Biotechnology Division, CSIR-IIIM, and given Crude Drug Repository Accession Number (CDR) 4029. Isolation of the endophytic fungi was carried out based on the procedures described earlier by Erza et al. [21]. The bulb samples were washed thoroughly with tap water to remove soil particles, sectioned into small pieces of 2–3 cm, and surface sterilized with 1% sodium hypochlorite, followed by a rinse with sterile distilled water to remove traces of sodium hypochlorite and 90% ethanol. The outer tissues of the bulb samples were removed under sterile laminar air flow. Small sections of 0.5 to 1 cm were plated on the culture media, such as PDA and water agar, after passing through a flame. The plates were incubated at 25°C for 3 weeks. Hyphal tips of fungi, emerging out of the plant bulb samples, were picked and grown on a PDA plate until pure cultures were obtained. The chemical data for the protons was reported in parts per million (p.p.m.) downfield from tetramethylsilane and are referenced to the residual proton in the NMR solvent (CD₃OD: 3.31 ppm). ESIMS were recorded on an Agilent 1100 LC-QTOF mass spectrometer. HPLC grade solvents like ACN and MeOH used for 2D-HPLC were purchased from SD Fine Chemical Limited. Water used for extractions was obtained from a Milli-Q Advantage A10 high-purity water system (Millipore), PDA, 1% sodium hypochlorite, lactophenol-cotton-blue, 15% glycerol, and DMSO, were purchased from Difco. The DNA purification kit was purchased from Qiagen. The caspase activity kit was purchased from Cell Signaling Technology. Dulbecco’s modified Eagle’s medium, Rosewell Park Memorial Institute medium (RPMI-1640), fetal bovine serum (10%), rhodamine 123 (Rh 123), DAPI (Dilactate), DMSO, gentamycin sulphate, trypsin, EDTA, PBS penicillin, streptomycin, paclitaxel (purity > 95% by HPLC), mitomycin (purity > 95% by HPLC), doxorubicin (purity > 98% by HPLC), gemicatube (purity > 98% by HPLC), staurosporine (purity > 95% by HPLC), and all other chemicals were purchased from Sigma-Alfrich. Tris buffer was procured from Himedia. Glacial acetic acid was purchased from Fisher Scientific and trichloroacetic acid was from Merck Specialties Private Ltd.
cultures were stored in 15% glycerol and stored at −70°C. The isolated cultures were also submitted to the IIIM repository (IIIM Microbial repository No. MRJ-326) for lyophilization and preservation.

Identification and characterization of the MRJ-326 isolate was carried out by using morphological and molecular procedures. The morphological examination was performed by observing the microscopic characteristics of the fungal culture on PDA. Microscopic slides were prepared by staining with lactophenol cotton blue [22] and were examined under a light microscope (Olympus). Total genomic DNA of the fungal isolate was extracted according to Ausubel et al. [23] using CTAB (cetyltrimethylammonium bromide) as per the manufacturer’s protocol. Fungal identification methods were based on their ITS of the fungal ribosomal DNA (ITS1–5.8S–ITS2), and were amplified using universal primers ITS4 (5’TTCCTCCGCTTATTGATATGC3’) and ITS5 (5’GGAAGTAAAAGTCGTAACAA3’). The thermal cycling program used was as follows: 3 min initial denaturation at 94°C, followed by 30 cycles of 15 sec denaturation at 94°C, 30 sec primer annealing at 55°C, 45 sec extension at 72°C, and a final 7-min extension at 72°C. The PCR amplified products were visualized on 1% (w/v) agarose gel. The amplicons were purified by using a DNA purification kit (Qiagen) and the purified product was used for sequencing the DNA. The corresponding ITS-rDNA sequence of the endophytic fungus was then used for similarity analysis using a BlastN algorithm against the public database at the National Center for Biotechnology Information (NCBI; http://www.ncbi.nlm.nih.gov).

Extraction and isolation of fungal metabolites

The endophytic fungal isolate P. pinophilum (MRJ-326) was cultured in 25 L of potato dextrose broth for 10 days at 25°C and 180 rev/min. The fermentation culture was then filtered using muslin cloth. The filtrate was extracted with ethyl acetate (10L, 16 h) successively five times. The extract was evaporated separately to dryness using rotavapor at 30°C. LC-MS is the preferred technique for the separation and detection of the large and often unique group of semi-polar secondary metabolites in natural product extracts. Specifically, LC-MS enables the detection of large numbers of parent ions present in a single extract and provides valuable information on the chemical composition and thus the putative identity of large numbers of metabolites. In our study, the ethyl acetate extract was analyzed using LC-MS (Agilent), a reversed-phase C18 (RP-C18) HPLC equipped with a photodiode array detector and triple quadrupole MS. The detection of large and often unique group of semi-polar secondary metabolites by using two detection systems [ultraviolet (UV) detection and an optional evaporative light scattering detection, ELSD]. This purification was performed for the ethyl acetate extract. The extract (150 mg) was dissolved in methanol (15 mL), membrane filtered, and then mixed with sorbent C4 silica (10 g, sepiatec, 360–200 mesh). The mixture was dried under vacuum to remove the residual solvent, thus yielding a free-flowing powder. The sample was then loaded onto a stainless steel column (8 mm D × 60 mm L), leaving one side open and 1 g of blank C4 silica material (1.0 g) was added. After levelling the surface of the stationary phase, the adsorbed extract was funnelled onto the top of the stationary phase, the surface was levelled, the remaining volume of the injection column was filled with blank C4 silica material, the column end fitting was attached, and then the injection column was connected to the SEPB-2D-250-HPLC system (Sepia tec). Using HPLC grade water as solvent A and methanol as solvent B, the effluent of the injection column was continuously pumped onto a 150 × 16 mm, i.d., 5 μm, VP150/16 Nucleoprep 100–12 C4 end-capped column at a flow rate of 4.5 mL/min. Chromatography was performed with the following gradient: 0 min, 20% B; 10 min, 50% B; 20 min, 100% B; 30 min, 100% B; 31 min, 20% B; 37 min, 20% B. Concomitant with the separation, a flow gradient was run with water, which was mixed with the C4 column effluent using the following flow rates: 0 min, 0 mL/min; 1 min, 50 mL/min; 12 min, 25 mL/min; 16 min, 20 mL/min; 21 min, 20 mL/min; 24 min, 1 mL/min. The effluent was collected in 16 subfractions, namely EA/1-EA/16, which were trapped onto VP30/32 Nucleoprep 100–12 C4 end-capped solid-phase extraction columns preconditioned with water (10 min, 8 mL/min) prior to the separation. Each subfraction was further separated on a 250 × 10 mm, 5 μm, Luna Phenyl-Hexyl column (Phenomenex) using the following gradient of water (solvent A), methanol (solvent B), and ACN (solvent C) at a flow rate of 4.5 mL/min: fraction EA/1, 0 min, A/B/C (100:0:0), 60 min, A/B/C (100:0:0), 65 min, A/B/C (9:0:10), 72 min, A/B/C (50:50:0); fraction EA/2, 0 min, 38% B, 29 min, 58% B, 30 min, 100% B, 34 min, 100% B; fraction EA/3, 0 min, A/B/C (56:44:0), 29 min, A/B/C (37:63:0), 30 min, A/B/C (100:0:0), 34 min, A/B/C (100:0:0); fraction EA/4, 0 min, A/B/C (48:52:0), 29 min, A/B/C (23:67:0), 30 min, A/B/C (100:0:0), 34 min, A/B/C (100:0:0); fraction EA/5, 0 min, A/B/C (32:68:0), 29 min, A/B/C (28:72:0), 30 min, A/B/C (100:0:0), 34 min, A/B/C (100:0:0); fraction EA/6, 0 min, A/B/C (31:69:0), 29 min, A/B/C (20:80:0), 30 min, A/B/C (100:0:0), 34 min, A/B/C (100:0:0); fraction EA/7, 0 min, A/B/C (22:78:0), 29 min, A/B/C (18:82:0), 30 min, A/B/C (100:0:0), 34 min, A/B/C (100:0:0); fraction EA/8, 0 min, A/B/C (22:78:0), 29 min, A/B/C (9:91:0), 30 min, A/B/C (100:0:0), 34 min, A/B/C (100:0:0); fraction EA/9, 0 min, A/B/C (21:79:0), 29 min, A/B/C (9:91:0), 30 min, A/B/C (100:0:0), 34 min, A/B/C (100:0:0); fraction EA/10, 0 min, A/B/C (22:88:0), 29 min, A/B/C (7:93:0), 30 min, A/B/C (100:0:0), 34 min, A/B/C (100:0:0); fraction EA/11, 0 min, A/B/C (7:93:0), 29 min, A/B/C (5:95:0), 30 min, A/B/C (100:0:0), 34 min, A/B/C (100:0:0); fraction EA/12, 0 min, A/B/C (3:97:0), 16 min, A/B/C (3:97:0), 17 min, A/B/C (3:97:0), 29 min, A/B/C (9:93:7), 30 min, A/B/C (50:50:0), 34 min, A/B/C (50:50:0); fraction EA/13, 0 min, A/B/C (3:97:0), 16 min, A/B/C (3:97:0), 17 min, A/B/C (100:0:0), 29 min, A/B/C (9:93:7), 30 min, A/B/C (50:50:0), 34 min, A/B/C (50:50:0); fraction EA/14, 0 min, A/B/C (9:98:2), 29 min, A/B/C (89:11:0), 30 min, A/B/C (50:50:0), 34 min, A/B/C (50:50:0); fraction EA/15, 0 min, A/B/C (98:2:9), 29 min, A/B/C (80:20:0), 30 min, A/B/C (50:50:0), 34 min, A/B/C (50:50:0); fraction EA/16, 0 min, A/B/C (100:0:0), 29 min, A/B/C (34:66:0), 30 min, A/B/C (10:90:0), 34 min, A/B/C (10:90:0). Column equilibration for each separation was 10 min at the initial gradient conditions. Each subfraction, EA/1–16, was separated/collected into subfractions in 48-well plates (4 mL/well) based on its purity. Well numbers 7, 8, 11, and 44 contained the compounds dictaranin (3), oxyskyrin (1), skyrin (2), and 1,6,8-trihydroxy-3-hydroxymethylantraquinone (4) sequentially. Com-
In vitro cytotoxicity sulphorhodamine B assay

Potential cell cytotoxicity of the crude ethyl acetate extract and isolated compounds were evaluated against a panel of human cancer cell lines. Test materials (extract and isolated compounds) were dissolved in DMSO and serially diluted at different concentrations (1, 10, 30, 50, 100 µg/mL). Human cancer cell lines were seeded in 96-well plates at a cell density of 10^4 cells/well for 24 h before treatment with the test material to allow the attachment of the cells to the plate. Serially diluted test materials were added to the cells in triplicate after being prepared for each individual concentration and were further incubated with the compound for 48 h at 37°C in atmosphere of 5% CO2. After 48 h, the cells were fixed with chilled 50% trichloroacetic acid, dried, and stained with sulphorhodamine B dye. The excess dye was washed with 1% glacial acetic acid and the remaining cell bound dye was dissolved in DMSO and the fluorescence was measured in a Biotek ELISA reader [24, 25].

Nuclear morphology studies

The presence of apoptotic cells were determined by staining MIA PaCa-2 cells with DAPI. Human pancreatic cancer (MIA PaCa-2) cells (2 × 10^5 cells/well) were fixed with chilled 50% trichloroacetic acid, dried, and stained with sulphorhodamine B dye. The excess dye was washed with 1% glacial acetic acid and the remaining cell bound dye was dissolved in Tris EDTA buffer and colorimetrically measured at 540 nm in a Biotek ELISA reader [24, 25].

Detection of apoptosis by annexin V-fluorescein isothiocyanate and propidium iodide

Annexin V-FITC and PI dual staining is usually used to detect the early and late apoptotic cells. For measuring apoptosis, human pancreatic cancer (MIA PaCa-2) cells were seeded in 6-well plates (2 × 10^5 cells/well) and were incubated with dicatenarin (20 µg/mL) and skyrin (50 µg/mL) for 48 h. Paclitaxel (1 µg/mL) was used as a positive control. After 48 h treatment, the cells were collected and resuspended in binding buffer. Then, the cells were stained with annexin V/PI and PI for 15 min in the dark and analyzed by a laser scanning confocal microscope (Olympus Fluoview FV1000) [27].

Detection of intracellular reactive oxygen species accumulation

Intracellular ROS levels were monitored by fluorescence microscopy after staining with DCFH-DA. Human pancreatic cancer (MIA PaCa-2) cells (2 × 10^5 cells/well) were seeded in 60 mm culture dishes. After 24 h, the cells were incubated with different concentrations of dicatenarin (5, 10, 20 µg/mL) and skyrin (25, 50 µg/mL) for 48 h. H2O2 (0.05%) was used as a positive control. Following treatment, the medium was removed and the cells were loaded with 5 µM DCFH-DA diluted in incomplete medium for 30 min at 37°C. The cells were washed three times with incomplete medium and were observed by a laser scanning confocal microscope at 10× [20].

Loss of mitochondrial transmembrane potential

Loss in MTP (ΔΨm) as a result of mitochondrial perturbation was studied using confocal microscopy after staining with rhodamine 123 (Rh123). Human pancreatic cancer (MIA PaCa-2) cells (2 × 10^5 cells/well) were seeded in 6-well plates and were treated with different concentrations of dicatenarin (5, 10, 20 µg/mL), skyrin (10, 25, 50 µg/mL) and paclitaxel (1 µg/mL) for 48 h. The cells were trypsinized and washed twice with PBS. The cell pellets were then suspended in 2 mL fresh medium containing Rh123 (1.0 µM) and incubated at 37°C for 20 min with gentle shaking. The cells were collected by centrifugation and washed twice with PBS, then analyzed by a laser scanning confocal microscope (Olympus Fluoview FV1000) [28].

Immunofluorescence confocal microscopy for detection of cytochrome c

The status of cytochrome c (intact or released) was examined by immunofluorescence microscopy. After 48 h treatment with dicatenarin (5, 10, 20 µg/mL), skyrin (25, 50 µg/mL), paclitaxel (1 µg/mL) and doxorubicin (0.5 µg/mL), human pancreatic cancer (MIA PaCa-2) cells (2 × 10^5 cells/well) were processed for immunofluorescence microscopic studies for the detection of cytochrome c release. The cells were incubated with MitoTracker Red (100 nm) for 10 min in the dark. Then cells were fixed in 4% paraformaldehyde for 10 min at room temperature and permeabilized using 0.5% Triton-X in PBS for 5 min. The cells were blocked with 10% goat serum for 20 min at room temperature. The expression of cytochrome c proteins was detected by incubating the cells with specific primary antibody anti-cytochrome c (mouse) for 1 h at room temperature (dilution 1:100). The Alexa Fluor 488 conjugated secondary antibody was diluted 1:500 in PBS and incubated for 1 h at room temperature. The cells were then washed three times in PBS and stained with DAPI (1 µg/mL in PBS). The coverslips were mounted over glass slides and the cells were im-

pounds from these were first collected and dried under vacuum for further NMR and MS spectroscopic analysis. Oxyskyrin (1): Orange-red amorphous solid; m.p. > 361°C; UV λmax (MeOH): 215, 256, 300, and 462; 1H NMR (400 MHz, methanol-d4): δH 2.48 (3 H, brs, CH3-11), δH 4.87 (2 H, s, CH2-11), δH 7.26 (1 H, brs, H-2), δH 7.09 (1 H, brs, H-2), δH 8.81 (2 H, s, H-2,7,7), δH 7.66 (1 H, d, J=2.6 Hz, H-4), 7.51 (1 H, d, J=2.6 Hz, H-4); ESIMS: m/z 554.9, LC-HRMS (full scan) calcd. for C30H17O12 [M + H]+ m/z 569.2 (100%) [12].

Oxyskyrin (2): Orange red powder; m.p. > 300°C; UV λmax (MeOH): 212, 256, 302, and 459; 1H NMR (400 MHz, methanol-d4): δH 2.35 (6 H, brs, CH2-11,11), δH 7.21 (2 H, brs, H-2,2′), δH 6.63 (2 H, s, H-7,7′), δH 7.50 (2 H, brs, H-4,4′); LC-HRMS (full scan) calcd. for C30H17O10 [M + H]+ m/z 537.2 (100%) [11, 12].

Dicatenarin (3): Orange amorphous powder, m.p. > 320°C; UV λmax (nm) (MeOH): 212, 265, 307, and 465; 1H NMR (400 MHz, methanol-d4): δH 2.21 (6 H, brs, CH3-11,11′), δH 7.25. (brs, 2 H, 2-H,2′), δH 6.60 (brs, 2 H, H-7,7′); LC-HRMS (full scan) calcd. for C30H17O12 [M + H]+ m/z 569.2 (100%) [12].

1,6,8-trihydroxy-3-hydroxymethylanthraquinone (4): Yellow-orange needles; m.p. 285–290°C; UV λmax (nm) (CH2Cl2): 232, 260, 308, and 372; 1H NMR (400 MHz, methanol-d4): δH 4.76 (2 H, s, CH2-11), δH 6.75 (1 H, d, J=2.1 Hz, H-7), δH 7.20 (1 H, d, J=2.1 Hz, H-5), δH 7.25 (1 H, brs, H-2), δH 7.69 (1 H, brs, H-4); LC-HRMS (full scan) calcd. for C13H9O6 [M + H]+ m/z 285.3 (100%) [13].
aged by a laser scanning confocal microscope (Olympus, Fluoview FV 1000) at 40× with an oil immersion lens [29,30].

Caspase-3 activity
Caspase-3 activity was determined using a commercial kit (Cell Signaling Technology) according to the manufacturer’s instructions. Human pancreatic cancer (MIA PaCa-2) cells (2 × 10^5/mL/well) were seeded in 6-well plates and treated with dicatenarin (5, 10, 20 µg/mL), skyrin (25, 50 µg/mL), and staurosporine (0.5 µg/mL) for 48 h following the same treatment as above. After treatment, the cells were harvested by centrifugation and the pellets were washed with PBS before lysis in chilled lysis buffer. The mixture was left on ice for 10 min and then centrifuged at 14000 g (Eppendorf Centrifuge, Sigma) at 4°C for 10 min. The resulting supernatant was used for the determination of caspase activity.

In vitro clonogenic assay
A clonogenic assay was performed to determine the capability of a cell to grow into a colony. Human pancreatic cancer (MIA PaCa-2) cells were plated at a seeding density of 1 × 10^4 cells/mL/well in 6-well plates. After 24 h, the culture medium was changed and new medium was added, and the cells were exposed to different concentrations of dicatenarin (5, 10, 20 µg/mL) and skyrin (10, 25, 50 µg/mL) for 7 days in a 37°C incubator in 5% CO_2. Later on, the obtained colonies were fixed with 4% paraformaldehyde and were stained with 0.5% crystal violet solution [32, 33]. The colonies from the plates were counted and photographed.

Statistical analysis
Results are expressed as mean ± SD of three individual experiments. Standard deviations were calculated using Graphpad prism and Microsoft excel. P-values < 0.05 were considered significant.

Supporting information
1^C-NMR data of compounds 1-4 in CD_3OD. LC-MS dereplication data of the ethyl acetate extract, and key 1H-^13C HMBC spectra of 1 and 2 are available as Supporting Information.

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Conflict of Interest
The authors have no conflict of interest to declare.

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