

Full Original Paper

## **Cytoskyrins and Cytosporones Produced by *Cytospora* sp. CR200: Taxonomy, Fermentation and Biological Activities**

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**Abstract:** In screening endophytic fungi from Costa Rica for bioactivity, fungal culture CR200, isolated from a buttonwood tree, was found to contain compounds that initiate DNA damage in a test strain of *E. coli* (Biochemical Induction Assay, BIA) and inhibit growth of Gram-positive bacteria, including antibiotic-resistant strains. Two new bisanthraquinones (cytoskyrins A and B) and five new related octaketides (cytosporones A-E) were isolated from fermentation broths of this fungus. Cytoskyrin A exhibited potent *in-vitro* antibacterial (MICs against Gram-positive bacteria, 0.03 - 0.25 µg/mL) and DNA-damaging activities (10 ng/spot), whereas cytoskyrin B was inactive in these assays. Among the cytosporones, only D and E exhibited Gram-positive activity, but they were inactive in the BIA. Mechanistically, cytoskyrin A specifically inhibited DNA synthesis in *E. coli imp* at its MIC; however, it also moderately inhibited protein synthesis at 2x its MIC. Cytoskyrin A exhibited poor cytotoxicity against tumor cell lines (IC<sub>50</sub> > 5 µg/mL) compared to known antitumor agents. The nuclear ribosomal internal transcribed spacer region of CR200 was found to share highest similarity (94-96%) with *Cytospora* spp. Micro- and macroscopic morphological observations of the conidia and conidiomata, respectively, also suggested this fungus to be a *Cytospora* sp.

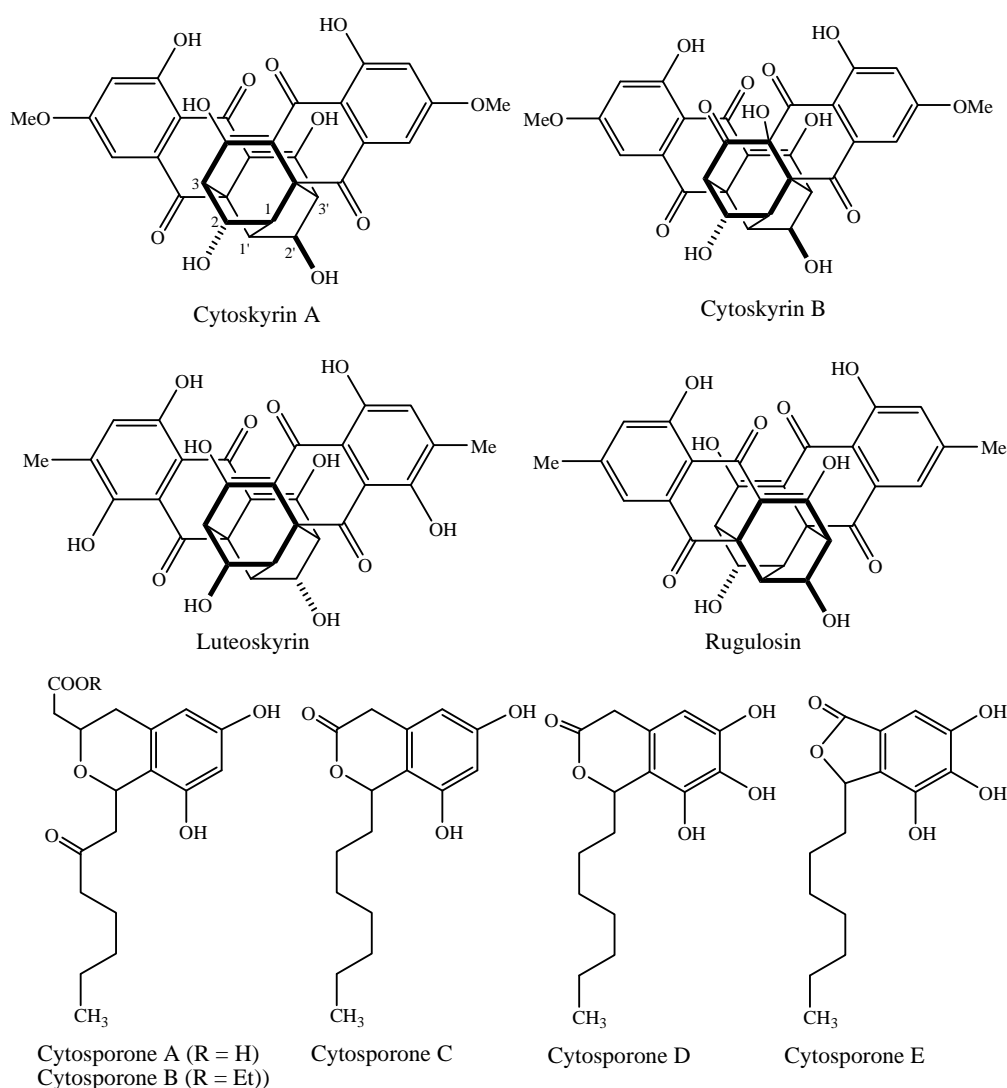
**Keywords:** *Cytospora* sp. CR200, cytoskyrins, cytosporones, antimicrobial, cytotoxic

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## 1. Introduction

Fungi, one of the most diverse groups of organisms, are known to produce a wide variety of medically important metabolites. Of the 1.5 million species estimated to be in existence only a small fraction (ca. 1%) have been studied [1]. In our continued efforts to discover novel compounds from filamentous fungi, culture CR200, obtained from a Costa Rican buttonwood tree, was found to be active in the Biochemical Induction Assay (BIA), which detects compounds that initiate DNA damage [2] and against a panel of antibiotic-resistant bacteria. As previously reported, two new bisanthraquinones (cytoskyrins A and B, Figure 1) and five new related octaketides (cytosporones A-E) were isolated from fermentation broths of this culture [3, 4]. Since then similar bisanthraquinones have been isolated from a number of different fungi [5, 6] and even synthesized [7-10]. In this paper, we report the biological activities of the isolated compounds as well as the taxonomy and fermentation of CR200. Some of these findings were previously presented at the 40<sup>th</sup> Interscience Conference on Antimicrobial Agents and Chemotherapeutic Agents and the Society for Industrial Microbiology meetings [11, 12].

**Figure 1.** Chemical structures of cytoskyrins and related bisanthraquinones, and cytosporones.

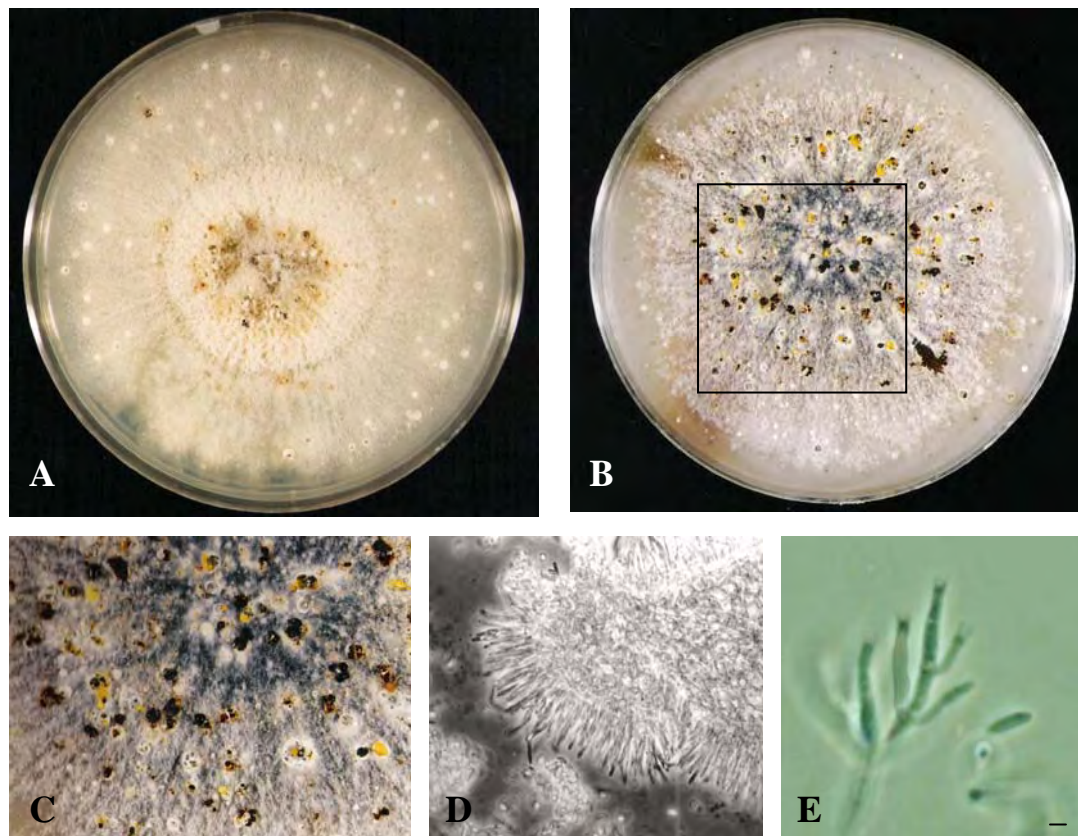


## 2. Results and Discussion

### *Culture Morphology*

Growth of CR200 on potato dextrose agar (PDA), cornmeal agar (CMA) and Leonian's agar (LA) were examined. The organism grew rather well on all three media, but the most differentiation and texture occurred on PDA (Table 1). CR200 grew robustly on PDA at 22°C reaching a diameter of 74 mm after only 7 days of growth and a maximum of 81 mm by 14 days (Figure 2A). Surface morphology was generally floccose to felty throughout the incubation period. Colony color progressed from white to taupe and buff with some yellow-brown at the colony center. Conidiomata (pycnidia) were visible by day 14 as white to light brown dots partially submerged into the agar (Figure 2A). By the 21<sup>st</sup> to 28<sup>th</sup> days, some conidiomata appeared dark brown (Figures 2B). About five to ten percent of them oozed a spore mass as yellow cirrhi in globules (Figures 2C) while others leaked a clear to brownish exudate. The conidiomata displayed elongated necks/beaks, some of them with extensive branching after an extended incubation of six weeks to several months.

**Figure 2.** Morphological characteristics of culture CR200 grown on PDA for 14 days (A), PDA for 28 days (B), enlarged view of the boxed area on B (C), CR200 phase contrast image of conidioma excised from a 28 day culture (D), and phase contrast image of conidiogenous cells with collerettes and a conidium (bar = 2.0 µm) (E).



**Table 1.** Morphological characteristics in different fungal media.

Medium	Temp (°C)	Day	Size (mm)	Color top and reverse, same if not indicated	Morphology
PDA	22	7	74	top: yellow-brown to white; reverse: gray-yellow to white	floccose to felty with submerged mycelia at margins
		14	81	top: taupe to buff; reverse: taupe to yellow-gray	floccose to felty with frequent conidiomata forming near margins
		21	81	top: taupe to beige to buff; reverse: yellow-gray	felty, floccose to fuzzy with submerged mycelia and frequent conidiomata
CMA	22	7	58	translucent	moist, flat, wispy
		14	76	translucent	thin, moist, smooth
		21	78	translucent to pale yellow	thin, moist, smooth
LA	22	7	82	translucent	flat, moist, radiate
		14	83	light olive-gray to translucent	thin, moist, smooth with some large submerged conidiomata
		21	83	light olive-gray to translucent	thin, moist, smooth with some large submerged conidiomata
PDA	33	7	26	yellow-brown to white	low raised, fuzzy, leathery to felty
		14	33	top: light brown to tan-gray to white; reverse: red-brown	floccose, powdery with sulcated margin and soluble red-brown pigment
		21	42	top: brown to taupe to light tan; reverse: medium brown to orange-brown	dusty, felty to floccose with sulcation at margins and brown soluble pigment
CMA	33	7	30	white	flat, radiate, moist
		14	44	translucent	thin, smooth, moist
		21	56	light buff to translucent	thin, smooth, moist with submerged mycelia
LA	33	7	28	brown to tan to white	moist, leathery
		14	46	red-brown to buff	leathery, gritty to smooth, submerged mycelia
		21	57	medium brown to dark tan to buff	smooth leathery to slightly gritty with submerged mycelia

CR200 grew most rapidly on LA, but remained flat or submerged in the agar. Mycelia were colorless or translucent becoming light olive-gray by day 14. Conidiomata also formed on LA by day 14, but were completely submerged, olive-gray and larger than those that formed on PDA.

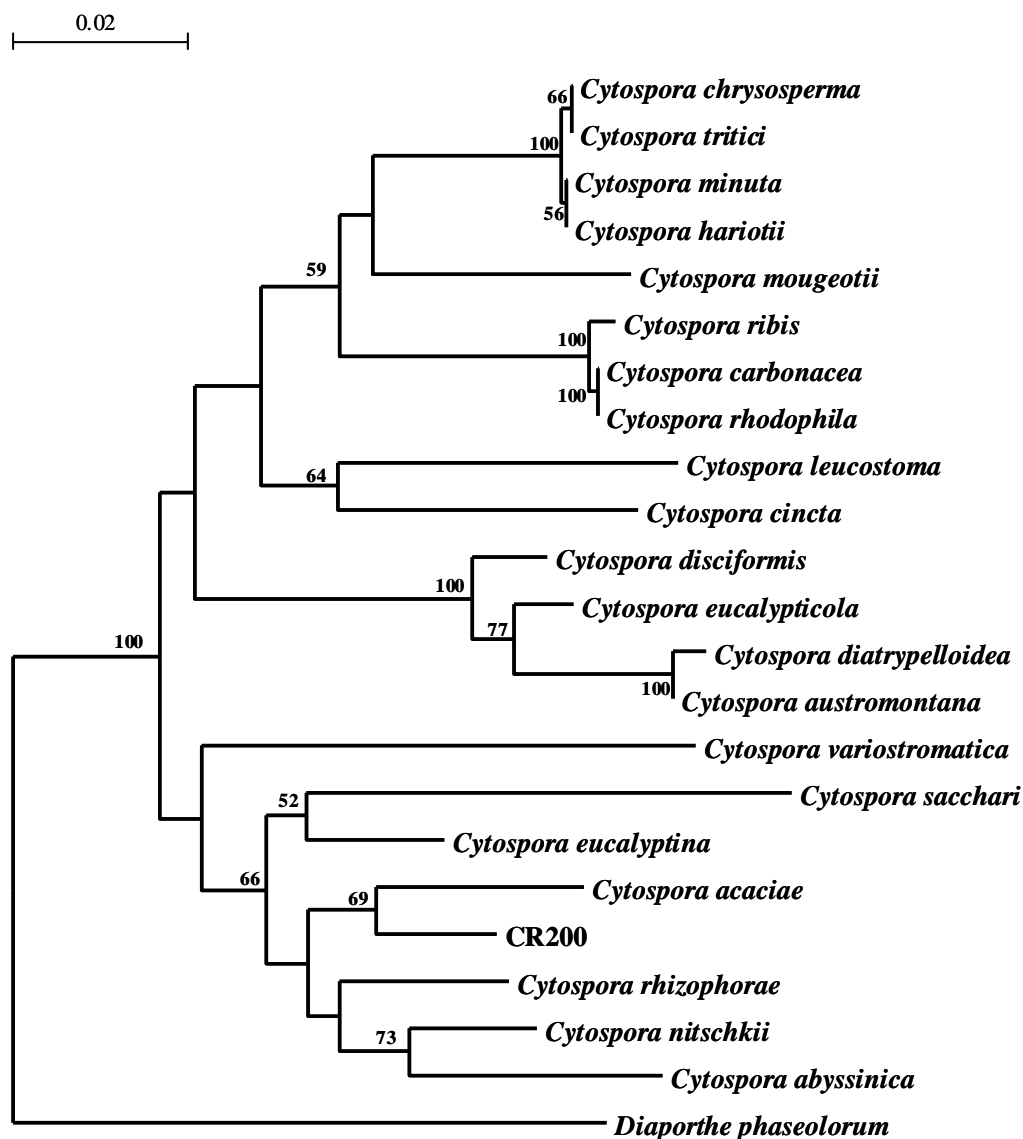
CR200 grew more slowly on all three media at 33°C. The growth stress at the higher temperature was indicated by altered morphologies. On PDA, a red-brown to brown soluble pigment was produced,

while on LA both the surface and reverse displayed red-brown to brown pigmentation. Conidiomata did not form on any media at 33°C. Conidiophores were hyaline, unbranched to highly branched at the base and/or mid-height and formed a continuous layer (Figure 2D, E). Conidiogenous cells were enteroblastic, phialidic, tapering to the apices with collerettes up to 2 µm long (Figure 2E). Conidia were hyaline, allantoid and measured about 5 to 6 µm long and 1.5 µm wide (Figure 2E).

### Phylogenetics

The ITS of CR200 was found to be 517 bp in length. A BLAST search of the ITS of CR200 revealed that it is most similar to *C. acaciae* (96%). A phylogenetic tree (Figure 2) of the ITS of 22 *Cytospora* spp. placed CR200 in a clade with *C. acaciae*, *C. rhizophorae*, *C. nitschkii*, and *C. abyssinica*. However, CR200 branched deeply from *C. acaciae* (supported by 69 bootstrap replicates), indicating that it is a distinct species.

**Figure 2.** ITS region phylogenetic tree of CR200, bootstrap values above 50 appear at nodes.



### Fermentation

Production of cytoskyrin A as measured by BIA activity was quite heterogeneous in the 100-flask fermentation. Approximately 87% of the flask fermentations were active in the BIA, with 42% displaying very good activity and 45% with fair to poor activity. The remaining 13% of the flask fermentations did not exhibit activity in the BIA, suggesting no production of cytoskyrin A. Separation of the cells from the broth and extraction of both with ethyl acetate revealed that the majority of cytoskyrin A was associated with the cells, but is also present in the fermentation broths. The higher shaking speed of 200 rpm produced activity detected by the BIA sooner than the lower shaking speed of 100 rpm, whether incubated at 22°C or 28°C. Interestingly, BIA activity was not detected from the whole broths of CR200 fermented at 22°C and 100 rpm, even after 21 days of incubation.

### Biological activities

DNA-damaging activities of the CR200 fermentation broth and pure compounds were tested by the biochemical induction assay (BIA). Cytoskyrin A demonstrated a minimum inducing concentration of 12 ng/spot in this assay, suggesting this compound as a potent DNA-damaging agent in bacteria (Table 2). Related bisanthraquinones such as rugulosin and luteoskyrin, known DNA intercalating agents, were toxic but did not induce the positive BIA response under the used assay conditions. These agents exhibit induction in the BIA only after a prolonged preincubation period. Cytoskyrin B, also produced by CR200, was inactive in the BIA. Cytoskyrin A appears to have the structural requirements to exhibit potent DNA-damaging activity in bacteria. Cytoskyrin A also exhibited excellent activity against the Gram-positive bacteria and the *E. coli imp*. However, it did not have activity against other Gram-negative bacteria, indicating that it may have difficulty crossing the outer membrane barrier (Table 3). Cytoskyrin B was inactive against the bacterial isolates tested (Tables 2 and 3). Minor differences in the chemical structures appear to affect the bioactivity of this class of compound (Figure 1). Among the five novel cytosporones A-E isolated from the CR200 fermentations, only D and E were active against Gram-positive bacteria and yeast (Table 3).

**Table 2.** BIA activity of Cytoskyrins and known anticancer drugs.

Concentration (µg/spot)	Cytoskyrin		Rugulosin	Luteoskyrin	Bleomycin	Mitomycin C	Doxorubicin
	A	B					
100	1+:T	0	T	T	1+:T	nt	nt
25	1+:T	0	T	T	1+:T	nt	nt
2.5	3+	0	T	T	3+	1+:11T	2+
1.25	3+	0	T	T	3+	2+:10T	1+
0.6	3+	0	T	T	3+	2+:7T	+/-
0.3	3+	0	T	T	3+	2+	0
0.15	2+	0	T	T	3+	2+	0
0.08	+/-	0	T	T	2+	1+	0

Test organism is *E. coli* with a lambda lysogen whose P<sub>L</sub> promoter is fused to *lacZ*

T = toxic zone, nt = not tested, 0 = inactive

3+ to +/- = level of β-galactosidase induction from strongest to weakest

**Table 3.** Antimicrobial Activity of Cytoskyrins and Cytosporones.

Organism	MIC (MBC) in µg/mL		
	Cytoskyrin A	Cytosporone D	Penicillin G
<i>S. aureus</i> ATCC 29923	0.031	8	<.062
<i>S. aureus</i> ATCC6538P	0.06 (0.12)	8	<.062
<i>S. aureus</i> #310 (MRSA)	0.03 (0.06)	8	64
<i>E. faecium</i> #379 (VREF)	0.25	16	32
<i>E. faecium</i> #436 (VSEF)	0.12	16	2
<i>B. subtilis</i> BGGS1A1	< 0.06 (0.06)	16	0.125
<i>E. coli imp</i> BAS849	0.12 (1)	64	2
<i>E. coli</i> BAS849	>128	128	32
<i>E. coli</i> ATCC25922	>64	64	16
<i>K. pneumoniae</i> ATCC 10031	>64	64	16
<i>P. aeruginosa</i> ATCC 27079	>64	>128	>64
<i>C. albicans</i> #54	16	4	>128

Test organisms #310 (methicillin-resistant), #379 (vancomycin-resistant), #436 (vancomycin-sensitive), and #54 are clinical isolates maintained in our lab. MIC of cytosporone E was same as cytosporone D. Cytoskyrin B and cytosporones A, B and C were inactive at the highest concentration tested.

Inhibition of DNA, RNA, and protein synthesis was determined by measuring the incorporation of  $^3\text{H}$ -Tdr,  $^3\text{H}$ -Udr, and  $^3\text{H}$ -AA, respectively, into TCA-precipitable material of a logarithmic-phase culture of *E. coli imp*. Control drugs affected the anticipated macromolecular processes (Table 4). Cytoskyrin A specifically inhibited the incorporation of  $^3\text{H}$ -thymidine into DNA (Table 4). Incorporation of amino acids into protein was also affected at higher concentrations. Based on the BIA activity and the specific inhibition of DNA synthesis in *E. coli imp*, cytoskyrin A appears to be a potent DNA-damaging agent. Since DNA-damaging agents also constitute an important class of anticancer drugs, cytoskyrin A was tested for antiproliferative activity against four human tumor cell lines and it was found to have moderate cytotoxicity with  $\text{IC}_{50}$  values in the range of 4 to 24 µg/mL (Table 5).

### 3. Conclusions

The cytoskyrins and cytosporones isolated from *Cytospora* sp. CR200 were first disclosed as novel compounds in 2000 by Brady et al [3,4]. Since then other researchers have reported the production of similar compounds by several different fungi [5, 6]. Unlike the related compound luteoskyrin, cytoskyrin A exhibited potent activity in the bacterial BIA and is therefore capable of directly or indirectly initiating DNA damage. Selective inhibition of DNA synthesis (measured by the incorporation of radiolabeled thymidine) in *E. coli* further confirmed its mode of action. Unfortunately, cytoskyrin A was less toxic to human cell lines than luteoskyrin and could not be used as an anticancer agent. Although luteoskyrin and cytoskyrins are chemically related, the mechanisms-of-action of these compounds appear to be different in prokaryotic and eukaryotic cells.

**Table 4.** Effects on the incorporation of radiolabeled precursors into macromolecules in *E. coli imp.*

Compound	Conc. ( $\mu\text{g/mL}$ )	$^3\text{H-Tdr}$	$^3\text{H-Udr}$	$^3\text{H-AA}$
Cytoskyrin A	2	3	102	54
	1	6	98	66
	0.50	19	100	89
	0.25	97	95	94
Ciprofloxacin	0.25	3	90	95
Rifampin	0.25	93	2	12
Chloramphenicol	8	98	95	16
Polymyxin B	8	1	2	3

Data presented represent % of untreated control after 10 min drug treatment and 5 min pulse labeling; Incorporation = precursor incorporated into TCA-precipitable material;  $^3\text{H-Tdr}$ ,  $^3\text{H-Udr}$  and  $^3\text{H-AA}$  are tritiated thymidine, uridine and amino acids, respectively.

**Table 5.** Cytotoxicity ( $\text{IC}_{50}$  in  $\mu\text{g/mL}$ ) of Cytoskyrin A and known antitumor agents.

Cell line (human)	Cytoskyrin A	Luteoskyrin	Taxol	Adriamycin
Epidermoid Carcinoma A431	4.1-9.6	0.08	0.0064	0.09
Breast Carcinoma SKBR3	5.2-24.3	0.12	0.0039	0.06
Carcinoma MDA435	3.6-13.0	0.13	0.0034	0.33
Colon Carcinoma SW620	5.2-16.6	0.18	0.0096	0.16

CR200 was isolated as an endophyte from a tissue sample of a branch of *Conocarpus erecta*. Although the fruiting structures of this fungus were never observed on its host, morphology of CR200 in culture and phylogenetic analysis of the ITS identified CR200 as a *Cytospora* sp. Further classification of CR200 to the species level would be difficult because no comprehensive keys are available for distinguishing *Cytospora* spp. in culture. Adams *et al.* [21] have shown that sequencing and comparative analysis of the ITS can be important to the identification of *Cytospora* spp., yet ITS data exists for less than 30 of the more than 300 species of *Cytospora* that have been described. Considering the importance of these fungi in the production of useful natural products such as antibiotics [22], angiotensin inhibitors [23] and HIV integrase inhibitors [24] efforts to sequence the ITS of validly described species would greatly aid in the future classification of *Cytospora* spp.

#### 4. Experimental

##### *Morphological Taxonomy*

Fungal culture CR200 was isolated from a branch of *Conocarpus erecta* (Buttonwood tree) in the Guanacaste National Park, Costa Rica (collection permit number 246.94). CR200 was streaked on Difco potato dextrose agar (PDA) and grown for 7-14 days at 22 °C. Mycelia and conidia were washed with potato dextrose broth (PDB) containing 25 % glycerol, and 1 mL aliquots were transferred into

cryovials for long-term storage at -140°C at Wyeth Research, Pearl River, NY. An additional copy is kept at the repository of the National Institute of Biodiversity, INBio Code 12652.

For macroscopic morphology, an agar plug of CR200 was placed in duplicate onto PDA, cornmeal agar (CMA, Oxoid) and Leonian's agar (LA, 1.2 g KH<sub>2</sub>PO<sub>4</sub>, 0.6 g MgSO<sub>4</sub>, 6.25 g maltose, 6.25 g Difco malt extract, 15.0 g Bacto agar/L of distilled water) and the culture plates were incubated at 22 °C and 33 °C. Colony morphology and size (in mm) were noted on the 7<sup>th</sup>, 14<sup>th</sup>, and 21<sup>st</sup> days of incubation. For microscopic analysis, CR200 was grown on PDA for 21 days. The conidiomata were observed at 1x to 6.3x magnification under an Olympus brand stereoscope. A smash prep was prepared by aseptic excision of several conidiomata, which were placed on a microscope slide with water (10 µL) and gently flattened with a coverslip. Microscopic observations were performed with an Olympus B071 phase-contrast microscope at 50x and 1,250x magnification.

### *Molecular Taxonomy*

Isolation of the genomic DNA, PCR amplification and direct sequencing of the nuclear ribosomal ITS1-5.8S-ITS2 region have been described previously [13]. Briefly, mycelia (50-100 mg) were lysed in lysis buffer (0.5 mL, 2% Triton X-100, 1% SDS, 100 mM NaCl, 10 mM Tris pH 8.0, 1 mM EDTA). Proteins were precipitated by the addition of 25:24:1 phenol/chloroform/isoamyl alcohol (0.5 mL, Sigma). The aqueous and organic layers were separated by centrifugation at 14,000 rpm for 5 minutes. The aqueous layer was pipetted into a new microfuge tube and the DNA was precipitated by addition of 3 M sodium acetate (0.1 mL) and isopropanol (0.8 mL). The nuclear ribosomal ITS1-5.8S-ITS2 region was amplified with primers ITS1 and ITS4 [14]. Amplification of the ITS was confirmed by gel electrophoresis. The PCR product was cleaned and sequenced directly with an ABI 3700 sequencer with the ABI Prism DNA sequencing kit and Big Dye terminators version 3.0 (Applied Biosystems).

For this study, the ITS were sequenced for *Cytospora* sp. CR200, *Cytospora cincta* ATCC 32673, *Cytospora chrysosperma* ATCC 56609, *Cytospora leucostoma* ATCC 42549 (deposited as *Leucostoma personii*), *Cytospora rhizophorae* ATCC 38475, and *Cytospora sacchari* ATCC 32322 and assigned the following GenBank accessions: DQ996039-DQ996044.

To determine the most closely related fungi, the ITS of CR200 was compared to other sequences in the GenBank database by BLASTN 2.2.2 analysis. The ITS of CR200 and several *Cytospora* spp. (Table 6) were aligned in Clustal X 1.81 with the multiple alignment parameters set at default values. A phylogenetic tree (Figure 2) was generated in TREECON 1.3b [15] with the distances calculated according to the Jukes and Cantor [16] method and insertions and deletions not taken into account. Bootstrap values were calculated using 1,000 bootstrap samples, and neighbor-joining was used to infer tree topologies. The phylogenetic tree was rooted with *Diaporthe phaseolorum*.

**Table 6.** *Cytospora* spp. used in the phylogenetic analysis.

<b>Culture name</b>	<b>Isolate</b>	<b>GenBank accession number</b>
<i>Cytospora abyssinica</i>	CMW 10181	AY347353
<i>Cytospora acaciae</i>	CBS 468.6	DQ243804
<i>Cytospora austromontana</i>	CMW 6735	AY347361
<i>Cytospora carbonacea</i>	CBS 219.54	DQ243805
<i>Cytospora chrysosperma</i>	ATCC 56609	DQ996043
<i>Cytospora cincta</i>	ATCC 32673	DQ996041
<i>Cytospora diatrypelloidea</i>	CMW 8549	AY347368
<i>Cytospora disciformis</i>	CMW 6509	AY347374
<i>Cytospora eucalypticola</i>	ATCC 56123	AF192314
<i>Cytospora eucalyptina</i>	CMW 5882	AY347375
<i>Cytospora hariotti</i>	CBS 195.49	DQ243807
<i>Cytospora leucostoma</i>	ATCC 42549	DQ996042
<i>Cytospora minuta</i>	CBS 134.25	DQ243808
<i>Cytospora mougeotii</i>	ATCC 44994	AY347329
<i>Cytospora nitschkii</i>	CMW 10184	AY347355
<i>Cytospora rhizophorae</i>	ATCC 38475	DQ996040
<i>Cytospora rhodophila</i>	ATCC 38695	DQ243809
<i>Cytospora ribis</i>	CBS 187.36	DQ243810
<i>Cytospora sacchari</i>	ATCC 32322	DQ996044
<i>Cytospora</i> sp.	CR200	DQ996039
<i>Cytospora tritici</i>	CBS 827.84	DQ243812
<i>Cytospora variostromatica</i>	CMW 6766	AY347366
<i>Diaporthe phaseolorum</i>	CBS 116018	AY745017

### Fermentation

Fermentation of CR200 for the production of the cytoskyrins was performed in a 6-liter batch. For seed fermentation, culture mycelia grown for seven days on Bennetts' agar (pH 7.0, containing per liter: dextrose 10 g, beef extract 0.77 g, yeast extract 1 g, NZ amine 2 g, agar 15 g) were scraped into thirty 25x150 mm Pyrex culture tubes each containing PDB (11 mL). Seed tubes were incubated for four days at 22 °C and 160 rpm. On the fourth day the contents of the seed tubes were combined into a sterile vessel and gently macerated to obtain a homogeneous inoculum. One hundred Erlenmeyer

flasks (250 mL), each containing sterile PDB (60 mL) were inoculated with the seed culture (5% v/v), and fermented for 21 days at 24 °C and 200 rpm. Additional flasks (five flasks for each condition) were inoculated to study the effects of temperature (22 °C versus 28 °C) and aeration (shaking at 100 rpm versus 200 rpm) on the production of cytoskyrin A, which was measured by testing whole broths in the BIA. Fermentation of CR200 for the production of the cytosporones was performed in a 10-liter batch. First-stage seed conditions were the same as described above except that only three seed tubes were inoculated. For the second stage, the contents of the seed tubes were added to PDB (500 mL in a 2.8 L Fernbach flask) and incubated at 22°C, 200 rpm for three days. For production fermentation, 10 L of PDB in a 10 L bioreactor (Bioflo 3000, New Brunswick Scientific) were inoculated at 5% v/v with second-stage and incubated at 24 °C, 350 rpm for 15 days.

### *Bioactivities*

#### *In vitro Antimicrobial Susceptibility Testing*

The minimum inhibitory concentrations (MICs) were determined by the microbroth dilution method [17, 18]. Briefly, inocula were adjusted to a density of  $1 \times 10^7$  to  $5 \times 10^7$  CFU/mL and 5  $\mu$ L were added to minimal medium or Mueller-Hinton broth (MHB) (100  $\mu$ L, containing the drug at concentrations ranging from 0.03 to 128  $\mu$ g/mL) in the wells of a microtiter plate. The MIC was defined as the lowest concentration of the antibiotic that prevented visual turbidity after 18 h of incubation at 37 °C.

#### *Biochemical Induction Assay (BIA)*

A slightly modified version of the previously described agar plate BIA method was used [2]. Briefly, an overnight culture of *E. coli* BR513 grown in LBE broth (100 mL/500 mL Erlenmeyer flask) at 37 °C and 200 rpm was inoculated into molten soft agar (40 mL, 1 %) at 50 °C to give a final  $A_{600}$  of 0.15 ( $1-3 \times 10^7$  cfu/mL). The inoculated soft agar was immediately poured onto a LBE agar base layer previously poured into a 9"x9" Nunc plate. After 15 min, five  $\mu$ L of each of the two-fold serially diluted solutions of the test compounds were spotted on the agar surface. The plate was incubated at 37 °C for 3.5 h and then overlaid with soft-agar (40 mL), containing fast blue RR (87 mg) and 6-bromo-2-naphthyl- $\beta$ -D-galactopyranoside (BNG, 13 mg). A red-violet color produced around the sample spot was read as a positive response.

#### *Incorporation of Radiolabeled Precursors*

Macromolecular synthesis in *E. coli imp*, an outer membrane permeability mutant [19], was studied by measuring the incorporation of appropriate radiolabeled precursors into trichloroacetic acid (TCA)-precipitable material [18]. *E. coli imp* was grown at 37 °C, 200 rpm in modified minimal medium (50 mL medium/250 mL Erlenmeyer flask) to an  $A_{450}$  of 0.20. Aliquots (100  $\mu$ L) were dispensed into microtiter wells containing antibacterial agents and the plates were incubated for 10 min at 37 °C with vigorous agitation. Cells were pulse-labeled for 5 min by adding the following radiolabeled precursors

at the indicated final concentrations:  $^3\text{H}$ -thymidine (Tdr), 2  $\mu\text{Ci/ml}$  with 0.05  $\mu\text{g}$  unlabeled thymidine/mL;  $^3\text{H}$ -Uridine (Udr), 2  $\mu\text{Ci}$  with 2.5  $\mu\text{g}$  unlabeled uridine/mL; or  $^3\text{H}$ -amino acid mixture (AA), 2.5  $\mu\text{Ci/mL}$ . To determine specific incorporation into DNA, RNA, and protein, chilled (4 °C) TCA (100  $\mu\text{L}$ , 10%) supplemented with unlabeled precursors (0.5 mg/mL) was added to each well, and the plate was immediately refrigerated for 1 h. The precipitate was collected on a glass fiber filter (Wallac filtermat B, Wallac 1205-404) using a Skatron 96-well cell harvester (Model 11050) programmed for a 3 sec prewet with chilled DI water, a 12 sec wash with 5% chilled TCA and a 5 sec drying cycle. Filter mats were dried for 7 min at high power in a microwave oven (Quasar, 700 Watts), solid scintillant (MeltilexB, Pharmacia 1205-402) was applied, and the isotope that was retained on the filter was quantitated in an LKB Betaplate scintillation counter (Wallac 1205). The levels of incorporation of  $^3\text{H}$ -Tdr,  $^3\text{H}$ -Udr, and  $^3\text{H}$ -AA were expressed as the percent of the untreated control.

### Antiproliferative Activity

Four human carcinoma cell lines, A431 (epidermoid carcinoma), SKBR3 (breast carcinoma), MDA-MB-435 (breast carcinoma), and SW620 (colon carcinoma) were used for cell proliferation assay. All cell lines were obtained from the American Type Culture Collection (ATCC). Cells were maintained in RPMI-1640 medium supplemented with 5% fetal bovine serum. Cells were plated in 96-well plates at the densities of  $5.0 \times 10^4/\text{mL}$ . On the next day, compounds were dosed at 0.5, 5, 50, 500, and 5000 ng/mL range and cultured for 2 days. At the end of incubation, cell survival was determined by the sulforhodamine B assay as previously described [20]. The  $\text{IC}_{50}$  values were obtained from the growth curves.

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### References and Notes

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