Viriditoxin, from a Jellyfish-derived Fungus, is Antibiotic to Fish Pathogens

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Abstract – A bioassay-guided fractionation of the extract of the fungus *Paecilomyces variotii*, which was derived from the giant jellyfish *Nemopilema nomurai*, led to the isolation of antibacterial compounds viriditoxin and its monomeric subunit semi-viriditoxin. Viriditoxin showed significant antibacterial activity against several marine fish and human pathogens including MDR strains. Significant potencies against resistant pathogens such as VRE *Enterococcus faecium*, VRE *Enterococcus faecalis*, and MRSA were highly interesting. Viriditoxin also showed notable antibacterial activity against the fish pathogen *Streptococcus iniae*. Its potency was over 100-fold higher than oxytetracycline which is employed as a general antibiotic for aquaculture.

Keywords - Paecilomyces variotii, Viriditoxin, Antibacterial activity, Jellyfish-derived fungus.

Introduction

In nature, the 6,6'-binaphthopyran-1-ones are rare metabolites and so far only five analogues have been reported (Suzuki et al., 1990; Suzuki et al., 1992; Arnone et al., 1995; Elix et al., 2004). Most of 6,6'binaphthopyran-1-ones were isolated as fungal metabolites except pigmentosin A, which was obtained from the lichen Hypotrachyna immaculata (Elix et al., 2004). Talaroderxines A and B are atropisomers isolated from a heterothallic ascomycetous fungus, Talaromyces derxii, and have strong antibacterial activity against Bacillus subtilis (Suzuki et al., 1990). Asteromine, a metabolite from a strain of Mycosphaerella asteroma, exhibits phytotoxic effects on Cucumis sativus, mild antibacterial and antifungal activity, and lethality to the brine shrimp Artemia salina (Arnone et al., 1995). Of these analogues, viriditoxin was the most interesting due to its broadspectrum antibiotic activity against Gram-positive pathogens, including methicillin-resistant Staphylococcus aureus (MRSA) and vancomycin-resistant Enterococci (VRE). The mechanism of action and broad-spectrum antibacterial properties of viriditoxin was studied and identified as an inhibitor of bacterial cell division by blocking FtsZ, which

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presents an excellent novel target for antibacterial drug discovery (Wang *et al.*, 2003).

Viriditoxin was first isolated from a mycelia extract of *Aspergillus viridi-nutans* (Lillehoj and Ciegler, 1972), and the structure was revised in 1990 (Suzuki *et al.*, 1990). This compound was also isolated from other fungi, such as *A. brevipes* (Lillehoj and Milburn, 1973), *A. fumigatus* (Mukhopadhyay *et al.*, 1987), and *Paecilomyces variotii* (Jiu and Mizuba, 1974).

In our study on the antibacterial components of the fungus *P. variotii* derived from the giant jellyfish *Nemopilema nomurai*, viriditoxin (1) and its monomeric subunit semi-viriditoxin (2) were isolated as causative components.

Experimental

General – ¹H and ¹³C NMR spectra were recorded on UNITY 400 and Varian INOVA 500 instruments. Chemical shifts were reported with reference to the respective residual solvent or deuterated solvent peaks. FABMS data were obtained on a JEOL JMS SX-102A. HRFABMS data were obtained on a JEOL JMS SX-101A. CD spectra were recorded on a JASCO J-715 instrument in CHCl₃; HPLC was performed with an YMC ODS-H80 column (250 × 10 mm i.d., 4 µm, 80 Å) using a Shodex RI-71 detector.

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Fungal materials – The fungus Paecilomyces variotii was isolated from the jellyfish Nemopilema nomurai collected off the southern coast of Korea in June 2007. The specimen was deposited at the Marine Natural Product Laboratory, PNU. Following a rinse with sterile sea water, the jellyfish tissue was homogenized and then inoculated on malt extract agar (MEA), which was prepared with 75% sea water, containing glucose (20 g/L), malt extract (20 g/L), agar (20 g/L), peptone (1 g/L), and antibiotics (10,000 units/mL penicillin and 10,000 µg/mL streptomycin, 5 mg/L). Fungi growing out of the jellyfish tissue were separated on the same MEA medium until a pure culture was obtained. Twelve pure fungal strains (J08NF-1~J08NF-12) were isolated from the jellyfish. The fungal strain J08NF-1 was selected on the basis of significant antibacterial activity against the gram-positive strain Staphylococcus aureus SG 511 (zone of inhibition 14 mm at 400 µg/disc), and it was identified as Paecilomyces variotii by Dr. K. S. Bae using morphological and biochemical analyses. P. variotii was then cultured in MEA medium (prepared with 75% sea water) containing glucose (20 g/L), malt extract (20 g/L), and peptone (1 g/ L) at 30 °C on a shaker platform at 155 rpm for 21 days, in total of 22 L.

Extraction and isolation - The culture medium and mycelia were extracted with EtOAc at room temperature. The antibacterial activity of the crude EtOAc extract of Paecilomyces variotii was tested against a panel of human pathogens (Staphylococcus aureus SG 511, Salmonella typhimurium, Klebsiella aerogenes 1522 E, Escherichia coli 078, and Enterobacter cloacae 1321 E) and marine pathogens (Edwardsiella tarda FP 5060, Listonella anguillarum FP 5208, Streptococcus iniae FP 5228, and Vibiro ichthyoenteri FP 4004) by disc diffusion method. The results showed that S. aureus SG 511 and 2 marine strains, S. iniae FP 5228 and V. ichthyoenteri FP 4004 were sensitive at an exact concentration of 400 µg/disc. Guided by the lethality to brine shrimp larvae (LD₅₀ $2 \mu g/$ mL) and antibacterial activity against these strains, the EtOAc extract (10.2 g) was partitioned between aqueous MeOH and *n*-hexane, whose zones of inhibition against S. aureus SG 511 were 13 and 7 mm at 30 µg/disc, respectively. Some yellow precipitate cannot be dissolved in both aqueous MeOH and n-hexane, and exhibited significant antibacterial activity against S. aureus SG 511 (Inhibition zone: 22 mm at 30 µg/disc). By filtration of the yellow precipitate, compound 1 (1.8 g) was obtained. The aqueous MeOH layer was subjected to a stepgradient MPLC (ODS-A, 120 Å, S-30/50 mesh) eluting with 50% to 100% MeOH to afford 21 fractions. Each

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fraction was tested for lethality to brine shrimp larvae and for antibacterial activity against *S. aureus* SG 511, *S. iniae* FP 5228, and *V. ichthyoenteri* FP 4004. Fraction 5, one of the bioactive fractions (Inhibition zone: 20 mm at 30 µg/ disc), was subjected to reversed-phase HPLC (YMC ODS-H80, 250 × 10 mm i.d., 4 µm, 80 Å) eluting with 75% MeOH + 0.4% HCOOH (v/v) to afford 9 subfractions. Compound **2** (2.6 mg) was obtained by purification of subfraction 3 on reversed-phase HPLC (YMC ODS-H80, 250 × 10 mm i.d., 4 µm, 80 Å) eluting with 60% ACN.

Viriditoxin (1) – Light yellow, amorphous powder; $[\alpha]_D^{25}$ –163.6 (c = 0.51, CHCl₃); CD ($c = 5.5 \times 10^{-5}$ M, CHCl₃): $\Delta \varepsilon$ (nm) + 7.7 (325.0), 0 (308.0), – 109.2 (275.5), 0 (264.5), 100.6 (255.5), 0 (227.0). ¹H NMR (DMSO-*d*₆, 500 MHz) δ 6.84 (H, s, H-8), 6.16 (H, s, H-5), 4.90 (H, m, H-3), 3.69 (3H, s, 7-OC<u>H</u>₃), 3.61 (3H, s, COC<u>H</u>₃), 2.84 (2H, m, H-4), 2.77 (2H, m, H-11); ¹³C NMR (DMSO-*d*₆, 100 MHz) δ 170.6 (C-1), 170.5 (<u>C</u>OCH₃), 163.5 (C-10), 160.8 (C-7), 159.1 (C-9), 139.4 (C-5a), 133.9 (C-4a), 113.5 (C-5), 110.0 (C-6), 108.5 (C-9a), 99.6 (C-10a), 98.8 (C-8), 76.4 (C-3), 56.6 (7-O<u>C</u>H₃), 52.3 (CO<u>C</u>H₃), 39.4 (C-11), 32.4 (C-4). HRFABMS *m*/z 661.1574 [M – H]⁻ (calcd for C₃₄H₂₉O₁₄, 661.1557).

Semi-viriditoxin (2) – Light yellow, amorphous powder; $[\alpha]_D^{25}$ –67.4 (c = 0.25, CHCl₃); ¹H NMR (pyridine- d_6 , 500 MHz) δ 6.89 (H, s, H-5), 6.87 (H, d, J = 1.5 Hz, H-8), 6.78 (H, d, J = 1.5 Hz, H-6), 5.19 (H, m, H-3), 3.80 (3H, s, 7-OC<u>H₃</u>), 3.64 (3H, s, COC<u>H₃</u>), 3.08 (H, m, H-4_a), 3.03 (H, m, H-4_b), 3.01 (H, m, H-11_a), 2.91 (H, m, H-11_b); ¹³C NMR (Pyridine- d_6 , 100MHz) δ 169.2 (<u>C</u>OCH₃), 169.1 (C-1), 163.4 (C-9), 162.0 (C-7), 158.8 (C-10), 140.0 (C-5a), 132.8 (C-4a), 114.5 (C-5), 108.5 (C-10a), 100.5 (C-8), 99.0 (C-9a), 98.3 (C-6), 74.8 (C-3), 54.2 (7-O<u>C</u>H₃), 50.6 (CO<u>C</u>H₃), 38.4 (C-11), 31.6 (C-4).

Bacterial strains and antibiotics – The human pathogens, S. aureus SG 511, S. aureus SG 503, Salmonella typhimurium, Klebsiella aerogenes 1522 E, Escherichia coli DC 0, E. coli TEM, Pseudomonas aeruginosa 1771, Enterobacter cloacae P 99, and E. cloacae 1321 E, were donated by the Korea Institute of Science and Technology (KIST). The marine strains, Edwardsiella tarda FP 5060, Listonella anguillarum FP 5208, S. iniae FP 5228, and V. ichthyoenteri FP 4004, were provided by National Fisheries Research & Development Institute, Korea. The methicillinresistant S. aureus 3089 (MRSA), multidrug-resistant (MDR) P. aeruginosa 2200, MDR S. typhimurium 8173, MDR E. cloacae 0252, MDR E. coli 1137, VRE Enterococcus faecium 5207, VRE Enterococcus faecalis 5210 and MDR Vibrio parahaemolyticus 7001 were purchased from Korea National Reasearch Resource Bank (KNRRB). All standard antibiotics were purchased from the Sigma Aldrich Co.

Antibacterial assay – MIC values of the compounds were determined by the modified 0.5 Mcfarland standard method. Two-fold dilutions of the compounds in the range of (40 - 0.16 µg/mL) were prepared in 0.5% DMSO. Antibiotics were similarly diluted in 0.5% MeOH to generate a series of concentrations ranging from 40 to 0.16 µg/mL per well. The turbidity of the bacterial suspensions was measured at 600 nm, and adjusted with medium to match the 0.5 McFarland standards $(10^5 - 10^6)$ colony forming units/mL). Subsequently, 180 µL of bacterial culture was inoculated into each well, and the test solutions (20 µL) were added to 96-well plates. Finally, the plates were incubated at 36 °C for 24 h, and the MIC values were determined in triplicates and reexamined at appropriate times. To ensure that these vehicles had significant effect on the bacterial growth, each bacterial species was additionally cultured in a blank solution containing LB or BHI broth media at concentrations equivalent to those of the test solutions.

Result and Discussion

The culture medium and mycelia were extracted with EtOAc. The EtOAc extract (10.2 g) was further partitioned between aqueous MeOH and *n*-hexane. Yellow precipitate which cannot be dissolved in both solvents was obtained and exhibited significant antibacterial activity against *Staphylococcus aureus* SG 511 (Inhibition zone: 22 mm at 30 μ g/disc). After filtration, the compound (1) showed high purity. The high yield (81.8 mg/L) and significant antibacterial activity of compound 1 were of interest.

Compound 1 was a yellow, amorphous powder. The exact mass of the $[M - H]^-$ ion at m/z 661.1574 matched well with the expected formula $C_{34}H_{29}O_{14}$ (Δ +1.7 mmu). The NMR data of compound 1 was almost identical to that of viriditoxin (Suzuki et al., 1990). The chirality of the axis between the naphtha- α -pyrone moieties in compound 1 was confirmed by exciton chirality method. The CD spectrum exhibited a strong negative ($\Delta \varepsilon - 109.2$ at 275.5 nm) and a positive second ($\Delta \varepsilon$ +100.6 at 255.5 nm) Cotton effects, which was similar to the reported (Fig. 2). This was due to the coupling between the ${}^{1}B_{b}$ transitions of two naphthalene chromophores, the long axis of naphtho- α -pyrone moieties in the compound was twisted in a counter-clockwise manner. Thus, the chirality of compound 1 was defined as R_a -configuration. Based on biogenetic consideration and specific rotation $([\alpha]_D^{25}$ -163.6) of 1 (Park et al., 2011; Weisleder and Lillehoj,



Fig. 1. Viriditoxin (1) and semi-viriditoxin (2).



Fig. 2. CD spectrum of viriditoxin.

1971), the configuration at C-3 was defined to be *S*, which is the same as that of the co-isolated compound **2** (*vide infra*).

Various bioactivities, especially antibacterial activity, of viriditoxin (1) were reported. Viriditoxin has been shown to block FtsZ polymerization leading to an arrest of cell division, and finally to cell death. Viriditoxin showed broad-spectrum antibiotic activity against Gram-positive pathogens, including methicillin-resistant *S. aureus* and vancomycin-resistant *Enterococci* (Wang *et al.*, 2003). Viriditoxin was lethal to mice with LD₅₀ of 2.8 mg/kg (Weisleder and Lillehoj, 1971). Viriditoxin can activate ATP hydrolysis (ATPase) and swelling in rat liver mitochondria (Wong and Hamill, 1976).

Considering possible biological role of compound **1** in marine ecology, it was evaluated for antibacterial activity against marine fish pathogens in addition to human

Strains	1	А	В	С	D
Edwardsiella tarda FP 5060 ^a	>40		0.63		
Vibrio ichthyoenteri FP 4004	0.63		0.16		
Streptococcus iniae FP 5228 ^a	0.31		40		
Staphylococcus aureus SG 511	0.16	0.16			
Staphylococcus aureus SG 503	40	0.16			
Escherichia coli DC 0	40	0.63			
Escherichia coli DC TEM	> 40	1.25			
Pseudomonas aeruginosa 1771	0.16	0.16			
Salmonella typhimurium	> 40	1.25			
Klebsiella aerogenes 1522 E	>40	1.25			
Enterobacter cloacae P 99	>40	40			
Enterobacter cloacae 1321 E	>40	1.25			
MDR Pseudomonas aeruginosa 2200	> 40			20	
MDR Salmonella typhimurium 8173	> 40				0.31
MRSA 3089	1.25			0.63	
MDR Vibrio parahaemolyticus 7001	> 40				0.63
MDR Enterobacter cloacae 0252	> 40				0.16
MDR Escherichia coli 1137	> 40				0.63
VRE Enterococcus faecium 5207	1.25				1.25
VRE Enterococcus faecalis 5210	1.25				1.25

Table 1. MICs (µg/mL) of viriditoxin (1) and antibiotics

^a The strains *Edwardsiella tarda* FP 5060, *Vibrio ichthyoenteri* FP 4004, and *Streptococcus iniae* FP 5228 were derived from marine fishes. A: tetracycline, B: oxytetracycline, C: vancomycin, D: levofloxacin

pathogens (Table 1). Compound **1** showed significant antibacterial potency against fish pathogens *Streptococcus iniae* FP 5228 and *Vibrio ichthyoenteri* FP 4004. It was especially notable that compound **1** inhibited the fish pathogen *S. iniae* FP 5228 with potency over 100-fold higher than that of oxytetracycline. Oxytetracycline is one of most frequently employed antibiotic in aquaculture. Compound **1** inhibited growth of human pathogens *S. aureus* SG 511 and *Pseudomonas aeruginosa* 1771, with MIC values of 0.16 µg/mL, which is comparable to that of tetracycline. Viriditoxin also inhibited the growth of three MDR strains, MRSA 3089, VRE *Enterococcus faecium* 5207, and VRE *Enterococcus faecalis* with similar potency (MIC 1.25 µg/mL) to that of levofloxacin.

Compound **2** was isolated as a yellow, amorphous powder, and it was identified as defined semi-viriditoxin by comparison of NMR data those reported. The absolute configuration at C-3 was defined as *S* by comparison of the optical rotation with the reported one (Tan and Donner, 2009). Semi-viriditoxin was previously isolated from the fungus *Paecilomyces variotii* and weak antibacterial activity against a number of bacteria was reported (Ayer *et al.*, 1991). Compound **2** showed weak

inhibitory activity against the human pathogen S. aureus SG 511 with a MIC value of 40 μ g/mL.

Viriditoxin (1), from a jellyfish-derived fungus, showed potent antibacterial activity, especially against the fish pathogen *S. iniae* FP 5228. The potency was over 100fold higher than that of oxytetracycline, and this suggest a potential ecological role of 1 in the jellyfish *N. nomurai*. Compound 1 also showed antibacterial activity against VRE with potency comparable to that of levofloxacin. Further study on antibacterial activity of 1 against fish pathogens and VRE would yield valuable informations.

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