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Possible Involvement of Pseudomonas Sphingolipid Ceramide N-deacylase in Red Spot Disease of Eels

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Strain TK4 capable of producing sphingolipid ceramide N-deacylase (SCDase) was isolated from pond in which eel culture was conducted. TK4 was assigned to the genus Pseudomonas on the basis of morphological, physiological and biochemical characteristics and found to be very similar to P. anguilliseptica, a known pathogen for red spot disease in eels. 16S rDNA sequence analysis revealed that TK4 and P. anguilliseptica should be classified into Group I of the genus Pseudomonas. Interestingly, not only TK4 but also P. anguilliseptica had the ability to produce SCDase which hydrolyzes glycosphingolipids and sphingomyelin to generate lyso-forms of sphingolipids. Since lyso-sphingolipids are toxic to many cell lines, SCDase-producing Pseudomonas spp. are suspected to be a potential virulence factor for fish disease including red spot disease (Sekiten-byo) in eels.

INTRODUCTION

Red spot disease (Sekiten-byo) has been reported as an epizootic disease among pond-cultured eels caused by an infection of *Pseudomonas anguilliseptica* (Wakabayashi and Egusa, 1972). This disease is distinguished from red fin disease, which is caused by an infection of *Aeromonas hydrophila* (Hoshina, 1962). Eels with red spot disease show a remarkable petechial hemorrhage in the skin of the mouth region, operculla and ventral side of the body. However, how the bacteria cause the disease has not vet been clarified at the molecular level.

We isolated a bacterium Pseudomonas sp. TK4 from eel cultured pond, which produced a novel sphingolipid-degrading enzyme. The enzyme was found to cleave the N-acyl linkage of ceramides of various glycosphingolipids as well as sphingomyelin to generate lyso-sphingolipids and fatty acids. The enzyme was designated as sphingolipid ceramide N-deacylase (SCDase) based on its unique specificity (Ito $et\ al$, 1995). Lyso-sphingolipids have been known to be toxic; causing not only hemolysis but also various cellular dysfunctions possibly via inhibition of protein kinase C (Taketomi and Kawamura, 1970; Hannun and Bell, 1987). These results may lead to the hypothesis that

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SCDase–producing bacteria are a potential virulence factor for disease in fish. We report here not only TK4 but also *Pseudomonas anguilliseptica* produce SCDase.

MATERIALS AND METHODS

Materials.

[¹⁴C]–Stearic acid (54 mCi/mmol) was purchased from American Radiolabeled Chemicals Inc. (U.S.A.) and silica Gel 60 thin–layer chromatography (TLC) plate was from Merck (Germany). [¹⁴C]–Galactosylceramide (GalCer) was prepared by the reverse hydrolysis reaction of SCDase as described previously (Kita *et al.*, 2001; Mitsutake *et al.*, 1998).

Bacterial strains.

Pseudomonas sp. TK4 was isolated from an eel pond in Miyazaki (Japan) by enrichment culture using a synthetic medium (NH₄Cl 0.05%, K₂HPO₄ 0.05%, and 0.2% NaCl, pH7.2) containing 0.05% bovine brain crude gangliosides as the sole source of carbon. TK4 was cultivated at 25 °C in Tryptone–yeast medium containing bovine–brain acetone powder extracts (medium A: 0.5% polypeptone, 0.5% NaCl, 0.1% yeast extract and 0.1% acetone powder, pH 7.2) for a specific period.

Physiological and biochemical tests.

The identification of TK4 was conducted according to the 9th edition of Bergey's Manual of Systematic Bacteriology (Krieg, 1984). Motility, morphology, and gram–staining characteristics were determined by light microscopy. A transmitting electron microscope was employed for observation of flagella with 2% p–tungstic acid. Utilization of each carbohydrate was determined with Hugh–Leifson medium (Hugh and Leifson, 1953). King's medium was used to test for diffusible pigment production (King $et\ al$, 1954). Kovacs' oxidase test was employed (Kovacs, 1956). The sensitivity of isolates to antibiotics or other antimicrobial agents was examined by using the disc method with Tri Disc (Eiken, Japan) or Showa Disc (Nissui Seiyaku, Co., Ltd.) on Muller–Hinton agar medium (Difco, USA). The sensitivity to vibriostatic agent 0/129 was tested according to Shewan $et\ al$. (Shewan $et\ al$., 1954).

Determination of G+C content.

Bacterial DNA was purified by the method of Marmur (Marmur, 1961). Guanine–plus–cytosine (G+C) content was determined using HPLC as reported by Tamaoka and Komagata (Tamaoka and Komagata, 1984) with some modification. In brief, purified bacterial DNA was dissolved in 10mM sodium acetate buffer, pH 5.3, containing $2\,\text{mM}$ ZnCl₂ (II) at a concentration of $1\,\text{mg/ml}$. The solution was heated at $100\,^{\circ}\text{C}$ for $15\,\text{min}$ and then cooled rapidly in an ice bath. An equal volume of nuclease P1 solution ($1\,\text{mg/ml}$ in $40\,\text{mM}$ sodium acetate buffer, pH 5.3, containing $2\,\text{mM}$ zinc chloride) was added to the DNA solution, and the preparation was incubated at $50\,^{\circ}\text{C}$ for $1\,\text{h}$. The reaction mixture was centrifuged at $15,000\,\text{rpm}$ for $15\,\text{min}$ followed by HPLC analysis with a reverse–phase column (YMC–Pack ODS–AQ, $150\times6.0\,\text{mm}$ I. D., YMC Co., Ltd., Japan). The nucleotides were eluted with $10\,\text{mM}$ phosphate buffer, pH 3.5, at a flow rate of

1.5 ml/min at 25 °C. Nucleotides were detected with a UV detector (Hitachi Co., Ltd., Japan) at 270 nm and peak areas were measured with a HITACHI model L-7200 data processor. A standard mixture of the four deoxyribonucleotides was purchased from Yamasa Shoyu Co., Ltd. (Japan).

Phylogenetic analysis.

A partial DNA sequence for 16S ribosomal RNA (16S rDNA, ca 1.5–kbp fragment) was amplified by using universal primers, p27f (5'–AGA GTT TGA TCM TGG CTC AG–3') and p1492f (5'–GGC TAC CTT GTT ACG ACT–3'). The amplified PCR product was gel purified, and directly sequenced as reported by Hiraishi (Hiraishi, 1992). Phylogenetic analysis was performed by using both distance and parsimony methods. The distance analysis and trees were generated by neighbor–joining in the program CLUSTAL W (Thompson *et al.*, 1994).

Assay of SCDase activity.

For the measurement of SCDase productivity, each strain was cultured in 5 ml of medium A at 25 °C for predetermined periods, the supernatant was obtained as described above, and the enzyme activity was assayed by the hydrolysis of [14 C]–GalCer. Twenty μ l of enzyme was incubated with the same volume of substrate solution (400 pmol of [14 C]–GalCer in 50 mM phosphate buffer, pH 6.0, containing 1.6% Triton X–100) at 37 °C for 6 h. One enzyme unit was defined as the amount capable of catalyzing the hydrolysis of 1 μ mol substrate for 1 min under the conditions described above. A value of 10 ° and 10 ° units of enzyme was expressed as 1 milliunit and 1 microunit, respectively.

TLC analysis.

Each sample was dried up, dissolved in $15\,\mu$ l of chloroform/methanol=2/1 (v/v) and applied onto the TLC plate. After the development (solvent system; chloroform/methanol/12 mM MgCl₂=65/25/4, v/v), the TLC plate was exposed to the IP plate (Fuji Film, Japan) for 3 h following analysis with a BAS-1500 imaging analyzer (Fuji Film, Japan).

RESULTS AND DISCUSSIONS

A SCDase-producing bacterium, strain TK4, was isolated by enrichment culture using synthetic medium containing crude bovine brain gangliosides as the sole source of carbon from pond water in which eel culture was conducted. Fig. 1 shows a photograph of *Pseudomonas* sp. TK4 under an electron microscope. TK4 was a rod-shaped bacterium with a single polar flagellum. The biochemical and physiological properties of TK4 are summarized in Table 1. Judging from these results, TK4 was assigned to the genus *Pseudomonas* and was found to be very similar to *P. anguilliseptica*, *P. alcaligenes*, and *P. pseudoalcaligenes*. 16S rDNA analysis revealed that these 4 strains belong to *Pseudomonas* Group I (Fig. 2). It is worth noting that *P. anguilliseptica* is known to be a pathologen of eel red spot disease (Wakabayashi and Egusa, 1972) and TK4 was isolated from the eel-cultured pond, although the G+C content (Table 1) and antibiotic sensitivity (Table 2) of the two species were clearly different.

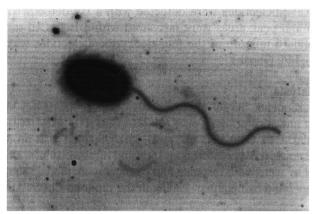


Fig. 1. Electron micrograph of *Pseudomonas* sp. strain TK4. The cells were stained with 2% *p*-tungstic acid solution (pH 7.2) and observed under an electron microscope (JEM 100 CXII, JEOL, Japan).

Table 1. Morphological, physiological, and biochemical properties of strain TK4

Characteristics	TK4	P. anguilliseptica	P. alcaligenes	P. pseudoalcaligenes
Shape	Short rod ·	Short rod	Short rod	Short rod
Motility	+	+	+	+
Flagellum	Polar monotrichous	Polar monotrichous	Polar monotrichous	Polar monotrichous
Gram staining	-	-	-	-
Growth in air	+	+	+	+
Catalase	+	+	+	+
Oxidase	+	+	+	+
O–F test	-	-	_	_
Color of colony	Yellowish white	Yellowish white	Yellowish white	Yellowish white
Growth at 41 °C	+	+	+	+
Arginine dehydrolase	+	+	+	d
Production of fluorescence pigm	ents –	_	_	_
Denitrification	+	+	+	d
Degradation of gelatin	+	+	d	d
Utility of:				
Glucose	-	_	_	-
Trehalose	_	_	_	-
D-xylose	_	_	_	_
Maltose	_	-	_	-
Saccarose	-	-	_	-
Mannitole	_	-	-	-
Ethyleneglycole	- '		_	_
Arginine	+	+	+	+
Citric acid	+	+	d	d
Urea	_	- ,	-	-
Mol% G+C in DNA	66.4	62.4	64-68	62-64

^{+:} positive, -: negative, d: weakly positive

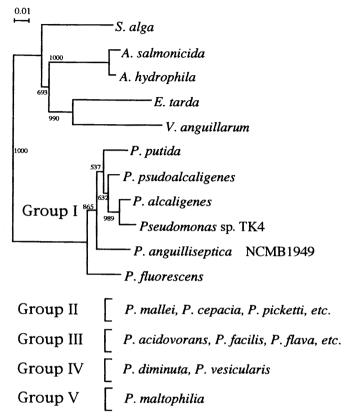


Fig. 2. Phylogenetic tree of *Pseudomonas* spp. by 16S rDNA analysis. The analysis was based on a comparison of approximately 1500 nucleotide bases of 16S rDNA partial sequences in which three different primers were used. The tree was constructed by using the CLUSTAL W algorithm (Thompson *et al.*).

Table 2. Sensitivity to antimicrobial agents

Antimianshial adapts	Strains		
Antimicrobial agents	TK4	P. anguilliseptica	
Penicillin	_	+	
Erythromycin	_	++	
Oleandomycin	-	+	
Leucomycin	_	++	
Novoviocin		+++	
Chloramphenicol	++	+++	
Tetracycline	+++	+++	
Kanamycin	+++	+++	
Colistin	+++	+++	

^{+++:} Strongly sensitive, ++: Moderately sensitive, +: Weakly sensitive,

^{-:} Insensitive

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Pseudomonas sp. TK4 has been isolated as an SCDase-producing bacterium (Ito et al., 1995) and found to be very similar to P. anguilliseptica based on the physiological and biolochemical characteristics in this study. Thus, we examined whether P. anguilliseptica produces SCDase or not. Interestingly, not only TK4 but also P. anguilliseptica NCMB1949 released SCDase into the culture supernatant when cultivated in medium A. However, the other 3 species of Pseudomonas and known fish pathogens tested did not produce the enzyme under the conditions (Table 3). Fig. 3 shows the time course for the production of SCDase by TK4 and P. anguilliseptica. The enzyme activity released in the culture supernatant by TK4 was detected at 1 day after inoculation of the bacterium and reached a plateau at 2–4 days, while that by P. anguilliseptica reached a maximum in 2 days. It was found that the production of SCDase by the latter was relatively low (Fig. 3), possibly due to the lower growth rate of P. anguilliseptica in the medium used.

Lyso-sphingolipids have been known to exert hemolytic (Taketomi and Kawamura, 1970) or cytotoxic activity (Sugiyama *et al.*, 1990). Lyso-sphingolipids were also found to be a potent inhibitor for protein kinase C (Hannun and Bell, 1987) which is known to play an important role in an intracellular signaling cascade. Recently, we reported the apoptosis of Neuro2a cells induced by lyso-sphingolipids with a naturally occurring stereochemical configuration (Sueyoshi *et al.*, 1997; Sueyoshi *et al.*, 2001). In the present study, it was observed that sphingosylphosphorylcholine (lyso-sphingomyelin) and galactosylsphingosine (psychosine) have strong hemolytic activity for fish erythrocytes, although their parental sphingolipids (galactosylceramide and sphingomyelin) nor fatty acids had no effects (Fig. 3).

As previously reported, SCDase can hydrolyze not only glycosphingolipids but also sphingomyelin to generate their lyso– forms (Ito *et al.*, 1995). Lyso–sphingolipids are hemolytic for various erythrocytes including those of fish and possibly toxic to vertebrates at the cellular level. Thus, SCDase–producing bacteria could be a virulence factor for fish disease, especially red spot disease of eel, because *P. anguilliseptica*, a pathogen causing the disease, was also found to produce SCDase in this study.

Species	Strain	Source	SCDase activity (μ U/ml)
E. tarda	NE8003 (FPC498)	flatfish ascites	_
E. tarda	NE8030 (FPC500)	red sea bream gill	
E. tarda	SU226	eel-cultured pond	- -
E. tarda	SU100	eel-cultured pond	
E. tarda	SU138	eel	
E. tarda	E381	tilapia kidney	_
V. anguillarum	PT87050	ayu	_
A. hydrophila	A9	eel	_
A. hydrophila	A10	eel	_
A. salmonicida	2-637-1	greenling	_
P. fluorescens	FPC185	carp	_
P. anguilliseptica	NCMB1949	eel blood	5.2
Pseudomonas sp.	TK4	eel-cultured pond	19.0

Table 3. Activity of SCDase from fish pathogenic bacteria.

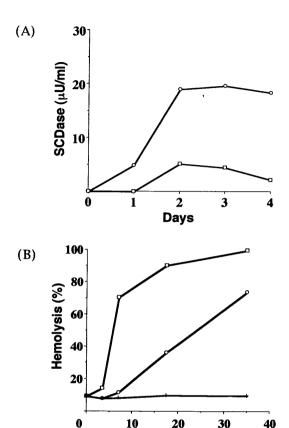


Fig. 3. Production of SCDase by TK4 and *P. anguilliseptica* (A) and hemolysis of carp erythrocytes by lyso–sphingolipids (B).

(A), Each strain was cultured at 25°C for the periods indicated and the enzyme activity was assayed as described under "MATERIALS AND METHODS". \bigcirc , TK4; \square , *P. anguilliseptica*. (B), Carp erythrocytes (7.5×10° cells) were incubated with appropriate concentrations of lyso–sphingolipids or their parental sphingolipids in 200 μ l of 20 mM PBS (pH 7.4) at 37°C for 30 min. After incubation, 400 μ l of PBS was added and centrifugation performed at 4°C (800 g×5 min). The supernatants were transferred into wells of a 96–well microtiter plate and then the absorbance of each well was measured at 541 nm. \square , galactosylsphingosine; \bigcirc , sphingosylphosphorylcholine; +, stearic acid, galactosylceramide or sphingomyelin.

Sphingolipids (µM)

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