

CHARACTERIZATION AND ANTIMICROBIAL ACTIVITY OF STREPTOMYCES STRAINS FROM THAI MANGROVE SOILS

Sirikan Hunadanamra¹, Ancharida Akaracharanya¹ and Somboon Tanasupawat^{2*}

¹Department of Microbiology, Faculty of Science, Chulalongkorn University, Bangkok 10330, Thailand ²Department of Biochemistry and Microbiology, Faculty of Phamaceutical Sciences, Chulalongkorn University, Bangkok 10330, Thailand

Received for publication: February 21, 2013; Accepted: April 05, 2013

Abstract Four actinomycete isolates, SAM2-1, SMP3-1, and J8-1 and J17-2 were isolated from mangrove soils collected in Samut Prakarn and Samut Songkram provinces, the inner gulf of Thailand. These isolates were identified as *Streptomyces* based on their phenotypic and chemotaxonomic characteristics. They contained L-diaminopimelic acid in cell wall. Their major menaquinone components were MK-9 (H6) and MK-9 (H8). On the screening of antimicrobial activity, they could inhibit *Staphylococcus aureus* ATCC 6538P and *Bacillus subtilis* ATCC 6633 but showed weak inhibitory activity against *Kocuria rhizophila* ATCC 9341, *Escherichia coli* ATCC 25922, *Pseudomonas aeruginosa* ATCC 27853 and no activity against *Candida albicans* ATCC 10231. Only isolate J8-1 showed strong inhibitory activity against *Bacillus subtilis* ATCC 6633. The isolates SAM2-1, SMP3-1, J8-1 and J17-2 were closely related to *S. sundarbansensis* DSM 42019T (98.4%), *S. diastaticus* subsp. *ardesiacus* JCM 5815T (97.1%), *S. iranensis* JCM 17327T (96.4%) and *S. seoulensis* NBRC 16668T (95.3%), respectively based on 16S rRNA gene analyses.

Keywords: Actinomycetes, Streptomyces, Antimicrobial Activity, Mangrove Soils

INTRODUCTION

Mangroves are the wet land forests mainly found in tropical and subtropical latitudes of the world coastal area¹⁻². The environment of the mangrove ecosystem is saline, and highly rich in organic matters consistent with high sulfur and nitrogen which can be used by the living microorganisms³. Mangrove rhizosphere is also full of decayed organic matters originated from alluvium, with a pH value range of acid to alkaline⁴. The genus Streptomyces and the rare genera actinomycetes in Micromonospora, Microbispora, Actinoplanes, Actinomadura and Pseudonocardia isolated from mangrove sediments and from the mangrove swamps have been studied for diversity as well as antimicrobial activity ⁵⁻⁸. Screening for the actinomycete species is an important aspect as there is a remarkable source for the production of diverse bioactive metabolites that possess pharmaceutically relevant biological activities⁹. In the course of our investigation of actinomycetes distributed in mangrove forests along the inner gulf of Thailand, the actinomycete isolates from soils in Samut Prakan and Samut Songkarm provinces were isolated, screened for antimicrobial activity and identified based phenotypic and chemotaxonomic on the characteristics including 16 S rRNA gene sequencing.

MATERIALS AND METHODS

Isolation of actinomycetes: The mangrove soil samples collected from Samut Prakan and Samut Songkarm provinces, Thailand (Table 1) were dried

*Corresponding Author:

Dr. Somboon Tanasupawat, Department of Biochemistry and Microbiology, Faculty of Pharmaceutical Sciences, Chulalongkorn University, Bangkok 10330, Thailand. both at room temperature for 1 week and at 110°C for 1 h.

Table.1: L	ocation,	pH of soi	il, isolate	number,	sequence
similarity	(%) and cl	losest spe	cies		

, , ,				
Location (Province)	pH of soil	Isolate no.	(%) Similarity	Closest species
Samut Songkram	7.5	SAM2-1	98.4	S. sunderbansensis DSM 42019 ^T
Samut Prakarn	7.8	SMP3-1	97.1	S. diastaticus JCM 5815 ^T
Samut Prakarn	6.3	J8-1	96.4	S. iranensis JCM 17327 ^T
Samut Prakarn	5.4	J17-2	95-3	S. seoulensis NBRC 16668 ^T

One gram of dried soil samples was suspended in 2.5 ml of sterile distilled water in a test tube, mixed for 30 seconds, left for 30 min, gently mixed 2-3 seconds and measureed the pH. The dried soil samples (0.5 g)were suspended in 4.5 ml of sterile distilled water and were heated at 60-65°C (15 min) to reduce nonthermotolerant microorganisms. The diluted soil suspension of 1:100 and 1:1000 (0.1 ml.) were spreaded on starch-casein nitrate agar (SCA) medium containing $15 \mu g/ml$ novobiocin and $25 \mu g/ml$ nistatin and incubated at 30°C for 14 days¹⁰. Actinomycete colonies were pick up and further purified by streak plate technique on yeast extract-malt extract agar (ISP 2) medium¹¹, and incubated at 30°C for 7-14 days. The purified cultures were maintained on ISP 2 agar slant at 4°C for further studies.



Identification methods: The phenotypic characteristics were determined as described by Shirling and Gottlieb¹¹ and Arai¹². Cell wall diaminopimelic acid (DAP) isomers were determined as described by Kutzner¹³. Menaquinone system was analysed as described by Komagata and Suzuki¹⁴. Scanning electron microscope was used for determining the morphology of strains grew on ISP 2 agar. DNA of the strains was isolated from cells grown in Yeast extract-Malt extract broth (ISP 2) with 0.2% of glycine reported by Yamada and Komagata¹⁵ and purified as described by Saito and Miura¹⁶. The 16S rDNA was amplified by PCR using primers, 8-27f and 1492r. The amplified 16S rDNA was used as templates for sequencing with Big Dve Terminator sequencing Kit (Perkin Elmer) and analyzed by AB1377 automated DNA sequencer (Perkin Elmer). The sequencing reaction for each sample was performed in DNA Thermal Cycler (Gene Amp PCR System 2400; Perkin Elmer) by using primers, 27F (5'-GTTTGATCCTGGCTCAG-3') and 1541R (5'-AAGGAGGTGACCAGCC-3'). The obtained sequence was compared with all sequences from GenBank using the BLAST program. The ClustalW2 program was used for multiple alignments with selected sequences for calculating evolution distances¹⁷ by Sea View version 4.2¹⁸. The phylogenetic tree was constructed using the neighbour-joining method¹⁹. Data were resampled with 1000 bootstrap replications. The values for sequence similarities among the most closely related strains were determined using the EzTaxon server²⁰.

Screening of antimicrobial activity: Primary screening of antimicrobial activities was performed on ISP 2 agar plates²¹ and S. aureus ATCC 6538P, B. subtilis ATCC 6633, K. rhizophila ATCC 9341, E. coli ATCC 25922, P. aeruginosa ATCC 27853, and C. albicans ATCC 10231 were used as indicator strains. All tested microorganisms were cultivated on Mueller-Hinton agar slants at 37°C for 24 h, except for the yeast strain that was cultivated on Sabouraud's dextrose agar slant at 30°C for 24h. The antimicrobial producing actinomycetes exhibited inhibitory distance against microorganisms tested. The inhibitory distances were measured and recorded. In addition, they were cultivated in ISP 2 broth on rotary shaker (200 rpm) for 4 days. Then, the selected cultures was transferred into the same medium containing calcium carbonate, and incubated for 10 days. The culture filtrate was extracted by ethyl acetate, rotary evaporator dried, and dissolved in methanol. Vacuum dried and redissolved in methanol were applied on a paper disc (0.6 cm diameter) with the amount of 1 mg/disc. The applied discs were placed on agar media spreaded over by the indicator strains using the agar disc diffusion method²². Streptomycin (20µg/disc) was used as positive control.

RESULTS AND DISCUSSION

Isolation and identification: The mangrove soil samples collected from Samut Prakarn and Samut Songkram provinces, Thailand. The pH of the soil samples ranged from 5.4 to 7.8. The actinomycete isolates could grow in wide range of pH. The isolate SAM2-1 was isolated from the soil collected in Samut Songkram while isolates SMP3-1, J8-1 and J17-2 were from the samples in Samut Prakarn. On the basis of their phenotypic characteristics (Tables 2 and 3), they were belonged to Streptomyces²³.

Table.2: Cultural charac	teristics of S	Streptomyces	isolates
--------------------------	----------------	--------------	----------

Isolate	Medium	Growth	Spore color	Colony color		
no.				Upper colony	Lower colony	
SAM2-1	YMA	+++	White	Pale reddish	Light reddish	
				yellow	yellow	
	TSA	++	White	White	White	
	OMA	++	White	White	White	
	AGA	++	White	White	White	
	ISA	+++	White	White	White	
SMP3-1	YMA	+++	Grayish white	Rose	Strong yellow	
	TSA	+++	Gray	Rose	Rose	
	ОМ	+++	Gray	Rose	Rose	
	AGA	+++	Purplish pink	White	Pink	
	ISA	+++	Vivid red	Vivid red	Vivid red	
			purple	purple	purple	
J8-1	YMA	+++	Yellowish	Yellowish	Brownish	
	TSA	+	grey White	grey White	gold White	
	ОМА	+++	White	White	Yellowish	
	AGA	+	White	White	Brownish gold	
	ISA	-	-	-	-	
J17-2	YMA	+++	Grayish	Yellowish	Yellowish	
	TSA	+++	Grayish brown	Grayish brown	Pale beige	
	OMA	+++	Gray	Dark brown	Black	
	AGA	+	Gravish	Gravish	Brown	
	-		brown	brown		
	ISA	+++	Grayish	Yellowish	Pale beige	
			white	white		

Note: YMA, Yeast extract-Malt extract agar; TSA, Tyrosine agar; OMA, Oatmeal agar; AGA, Glycerol-asparagine agar; ISA, Inorganic salt-starch agar. +++, good; ++, moderate; +, poor; -, no growth.

Isolate SAM2-1 produced powdery colony with spiral spore chain (Fig.1), and with rectiflexibile spore chain. This isolate grew well on ISP 2 agar plate, produced white spore while the upper colony color was pale reddish yellow and the lower side was light reddish yellow. The cultural characteristics on tyrosine, oatmeal, asparagine and inorganic salt agar media are shown in Table 2.

Isolate SMP3-1 produced powdery colony with spiral spore chain (Fig.1), and with rectiflexibile spore chain. This isolate grew well on ISP 2 agar plate, produced grayish white spore while the upper colony color was rose and the lower side was strong yellow. The cultural characteristics on tyrosine, oatmeal, asparagine and inorganic salt agar media are shown in Table 2.

Table.3: Characteristics and antibacterial activity of

 Streptomyces isolates

Characteristics	SAM2-1	SMP3-1	J8-1	J17-2
Max. NaCl (%,w/v)	5	4	2	6
Growth at pH 5-9	+	+	+	+
Growth at 45 °C	W	+	+	+
Nitrate reduction	+	+	+	+
Milk peptonization	-	-	+	+
Milk coagulation	+	-	+	-
Gelatin liquefaction	+	+	+	+
Starch hydrolysis	W	+	+	+
H₂S formation	-	-	-	-
Melanin formation	-	-	-	-
Utilization of :				
L-Arabinose	w	-	+	w
Fructose	-	-	+	-
D-Glucose	+	+	+	+
Glycerol	+	w	w	+
Mannitol	+	w	+	-
Raffinose	+	-	+	-
Rhamnose	+	-	+	+
Sucrose	-	+	W	+
D-Xylose	W	-	+	w
L-Diaminopimelic acid	+	+	+	+
Indicator strain Inhibitory against (mm)				
S. aureus	24.0	15.0	25.0	25.1
B. subtilis	-	15.0	25.1	14.0
K. rhizophila	-	13.0	-	-
P. aeruginosa	10.0	-	-	-
E. coli	-	8.0	-	-

Note: +, positive; -, negative; w, weakly positive.

Isolate J8-1 produced powdery colony with spiral spore chain (Fig.1), and with rectiflexibile spore chain. This isolate grew well on ISP 2 agar plate, produced yellowish grey spore while the upper colony color was yellowish grey and the lower side was brownish gold. The cultural characteristics on tyrosine, oatmeal, asparagine and inorganic salt agar media are shown in Table 2.

Isolate J17-2 produced powdery colony with spiral spore chain (Fig.1), and with rectiflexibile spore chain. This isolate grew well on ISP 2 agar plate, produced grayish white spore while the upper and the lower colony color were yellowish white. The cultural characteristics on tyrosine, oatmeal, asparagine and inorganic salt agar media are shown in Table 2.









Figure.1: Scanning electron micrograph of Streptomyces sp. SAM2-1 (a), SMP3-1 (b), J8-1(c) and J17-2 (d) grown on YMA medium (7-14 days)

The four isolates grew on ISP 2 medium containing 2% to 6% NaCl, at pH 5.0 to 9.0 and at 45°C. They

reduced nitrate, hydrolysed starch and liquefied gelatin, whereas some strains coagulated skim milk and showed milk peptonization but all did not produce H₂S, form melanin and hydrolyse chitin. All strains utilized glucose and glycerol but some did utilize L-arabinose, fructose, mannitol, rhamnose, raffinose and sucrose, as single carbon sources (Table 3). Isolates SAM2-1, SMP3-1, J8-1 and J17-2 contained L-diaminopimelic acid (L-DAP) indicating that these strains had cell wall chemotype I as described by Lechevalier & Lechevalier²⁴, which is the cell wall type of Streptomyces as described by Schleifer & Kandler²⁵. The major menaquinones of strains were $MK-9(H_6)$ (33.2-67.6%), and MK-9(H₈) (32.4%-66.8%). On the basis of 16 S rRNA gene analyses, the isolates SAM2-1 (1130 bp), SMP3-1 (1070 bp), J8-1 (1088 bp) and J17-2 (1050 bp) were closely related to Streptomyces sundarbansensis DSM 42019^{T} (98.4%), S. diastaticus subsp. ardesiacus JCM 5815^T (97.1%), S. iranensis JCM 17327^T (96.4%) and S. seoulensis NBRC 16668^T (95.3%), respectively ²⁶⁻²⁸.



2010

Figure.2: Neighbour-joining tree based on 16S rRNA gene sequences showing the phylogenetic relationships between the isolates and *Streptomyces* species. Based on 1000 resamplings, bootstrap percentages above 50% are shown. Bar, 0.01 substitutions per nucleotide position

Screening of antimicrobial activity: Four isolates showed inhibitory activity against *S. aureus* ATCC 6538P, isolates SMP3-1, J8-1 and J17-2 showed inhibitory activity against *B. subtilis* ATCC 6633, and only isolate SAM2-1 could inhibit against *P. aeruginosa* ATCC 27853 and and SMP3-1 inhibited *K. rhizophila* ATCC 9341 and *E. coli* ATCC 25922. As mentioned, the isolates could exhibit antimicrobial activities against Gram-positive bacteria and some of them could do against Gram-negative bacteria. The antimicrobial activity test by agar disc diffusion method of the isolates SAM2-1and SMP3-1 revealed that all the culture extracts inhibited Gram positive bacteria; *S. aureus* ATCC 6538P, *B. subtilis* ATCC 16633 and *K. rhizophila*

ATCC 9341 while only the culture extracts of isolate SMP3-1 inhibited *E. coli* ATCC 25922, and those of isolate SAM2-1 and SMP3-1 inhibited *P. aeruginosa* ATCC 27853. None of them inhibited *C. albicans* ATCC 10231.

Recently, the actinobacteria which identified as *S. exfoliatus*, *S. vinaceusdrappus*, *S. tendae*, *S. aureus*, *S. atriruber*, *S. olivochromogenes*, *S. malaysiensis*, *S. purpeofuscus*, *S. sparsogenes*, *S. aldersoniae*, *S. rapamycinicus* and *S. youssoufiensis* were isolated from terrestrial soils collected in Nakhon Si Thammarat, the southern part of Thailand. Most of them showed inhibitory activity against *B. subtilis* ATCC 6633 and *K. rhizophila*²⁷ that they were different from our *S. sundarbansensis*, *S. diastaticus*, *S. iranensis* and *S. seoulensis* strains isolated from mangrove soils. The strains in this study, however they are required to further study on DNA-DNA hybridization for clarifying their taxonomic position.

CONCLUSION

The mangrove actinomycete strains, SAM2-1 isolated from Samut Songkram and SMP3-1, J8-1 and J17-2 isolated from Samut Prakarn were identified as *Streptomyces* based on their phenotypic and chemotaxonomic characteristics including 16S rRNA gene analyses. They could inhibit Gram-positive bacteria, *S. aureus* ATCC 6538P but showed weak inhibitory activity against *K. rhizophila* ATCC 9341, Gram-negative bacteria, *E. coli* ATCC 25922, *P. aeruginosa* ATCC 27853. Only isolate J8-1 showed strong inhibitory activity against *B. subtilis* ATCC 6633.

REFERENCES

- 1. Alongi, DM, Present state and future of the world's mangrove forests, Environ. Conserv, 2002, 29, 331-349.
- 2. Sahoo K, Dhal NK, Potential microbial diversity in mangrove ecosystem: A review, Indian J Mar Sci, 2009, 38, 249-256.
- 3. Kizhekkedathu NN, Parukuttyamma P, Mangrove actinomycetes as the source of Lignolytic enzymes. *Actinomycetologica*, 2005, 19, 40-47.
- 4. Aksornkoae S, Mangroves-ecology and management, 3rd ed., Kasetsart University Press, Bangkok, 1999, 278 pp. (in Thai).
- Ravikumar S, Suganthi P, Moses F, Crude bioactive compounds of actinomycetes from manakkudy mangrove sediment, J Pharm Res, 2011, 4, 877-879.
- 6. Patil RCP, Mule AD, Mali GV, Tamboli RR, Khobragade RM, Gaikwad SK, Katchi VI, Patil D, Isolation of marine actinomycetes from the mangrove swamps for biotechnological exploration, J. Life Sci., 2011, 5, 1030-1036.
- 7. Naikpatil SV, Rathod JL, Selective isolation and antimicrobial activity of rare actinomycetes from mangrove sediment of Karwar, J Ecobiotechnol, 2011, 3, 48-53.
- 8. Mangamuri UK, Muvva V, Poda S, Kamma, S, Isolation, identification and molecular characterization of rare

actinomy cetes from mangrove ecosystem of Nizam patnam, Malay sian J Microbiol, 2012, 8, 83-91.

- Berdy J, Bioactive microbial metabolites: Review article, J Antibiot, 2005, 58, 1-26.
- Thawai C, Tanasupawat S, Itoh T, Suwanborirux K, Kudo, T, Micromonospora auratinigra sp.nov., isolated from a peat swamp forest in Thailand, Actinomycetologica, 2004,18, 8-14.
- 11. Shirling, EB and Gottlieb D, Methods for characterization of *Streptomyces* species, Int J Syst Bacteriol, 1966, 16, 313-340.
- 12. Arai T, Tamotsu F, Masa H, AkihiroM, Yuzuru M, Akio S, Akira S, Culture media for Actinomycetes, The Society for Actinomycetes, Japan, 1976, p.1-31.
- Kutzner H J, The Prokaryotes: A handbook on habitats, isolation and identification of bacteria, Vol. 2, Springer-Verlag, Berlin, 1981, p.2028- 2029.
- 14. Komagata K, Suzuki K, Lipid and cell-wall analysis in bacterial systematics. Methods Microbiol, 1987, 19, 161-207.
- 15. Yamada, K. and Komagata, K. Taxonomic studies on coryneform bacteria. III. DNA base composition of coryneform bacteria. J Gen Appl Microbiol, 1970, 16, 215-224.
- Saito H, Miura K, Preparation of transforming deoxyribonucleic acid by phenoltreatment. Biochim Biophys Acta, 1963, 72, 619-629.
- 17. Kimura M, A simple method for estimating evolutionary rates of base substitutions through comparative studies for nucleotide sequences, J Mol Evol, 1980, 16, 111-120.
- Gouy M, Gascuel S, Gascuel O, SeaView version 4.2: a multiplateform graphical user interface for sequence alignment and phylogenetic tree building, Mol Biol Evol, 2010, 27, 221-224.
- 19. Saitou N, Nei M, The neighbor-joining method: a new method for reconstructing phylogenetic trees, Mol Biol Evol, 1987, 4, 406-425.

- 20. Chun J, EzTaxon: a web based tool for the identification of prokaryotes based on 16ribosomal RNA gene sequences, Int J Syst Evol Microbiol, 2007, 57, 2259-2261.
- 21. Anansiriwattana W, Tanasupawat S, Amnuoypol S, Suwanborirux K, Identification and antimicrobial activities of actinomycetes from soils in Samed Island, and gedanamycin from strain PC4-3, Thai J Pharm Sci, 2006, 30 49-56.
- 22. Lorian V, Antibiotics in Laboratory Medicine, The Williams and Wilkins Company, Baltimore, 1991, p.1-51.
- Holt JG, Filamentous actinomycetes and related bacteria, Vol. IV, In Bergey's Manual of systemic Bacteriology, The Williams and Wilkins Co, Baltimore, 1989.
- 24. Lechevalier MP, Lechevalier HA, Chemical composition as a criterion in the classification of aerobic actinomycetes, Int J Syst Bacteriol, 1970, 20, 435-443.
- 25. Schleifer KH, Kandler O, Peptidoglycan types of bacterial cell walls and their taxonomic implications, Bacteriol Rev, 1972, 36, 407-477.
- 26. Arumugam M, Mitra A, Pramanik A, Saha M, Gachhui R, Mukherjee J, *Streptomyces sundarbansensis* sp. nov., an actinomycete that produces 2-allyloxyphenol, Int J Syst Evol Microbiol, 2011, 61, 2664-2669.
- Hamedi J, Mohammadipanah F, Klenk HP, Pötter G, Schumann P, Spröer C, Vonjan M, Kroppenstedt RM, Streptomyces iranensis sp. nov., isolated from soil, Int J Syst Evol Microbiol, 2010, 60, 1504-1509.
- Chun J, Youn HD, Yim YI, Lee H, Kim MY, Hah YC, Kang SO, Streptomyces seoulensis sp. nov., Int J Syst Bacteriol, 1997, 47, 492-498.
- 29. Sripreechasak P, Tanasupawat S, Matsumoto A, Inahashi, Y, Suwanborirux K, Takahashi Y, Identification and antimicrobial activity of actinobacteria from soils in southern Thailand, Trop Biomed, 2013, 30, 46-55.

Source of support: Nil Conflict of interest: None Declared