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# Preliminary screening for antibacterial and antimycobacterial activity of actinomycetes from less explored ecosystems

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**Abstract** Actinomycetes from less explored ecosystems were screened for antibacterial and antimycobacterial activity. Crude bioactive compounds were produced by growing these strains by shake flask fermentation using soybean meal medium. Culture supernatant and mycelia were extracted with ethyl acetate and methanol, respectively. Antibacterial activity of crude extracts was tested by disc diffusion method against gram positive and gram negative bacteria. Actinomycete strains D10, D5, NEK5, ANS2, M104 and R2 showed prominent activity. Culture filtrates and crude extracts were tested against standard strain Mycobacterium tuberculosis H<sub>37</sub>Rv and drug sensitive and drug resistant clinical isolates of M. tuberculosis by luciferase reporter phage (LRP) assay. Considerable variation was observed in antimycobacterial activity between actinomycete culture filtrates and solvent extracts. Actinomycete strains viz., D10, D5 (desert), CSA14 (forest), CA33 (alkaline soil), NEK5 (Neem plant), MSU, ANS2, R2 and M104 (marine) screened in the present study were found to be highly potent showing good antibacterial and antimycobacterial activity. Five of them such as A3, CSA1, EE9, ANS5 and R9 were exclusively active against M. tuberculosis. Secretary products of actinomycetes of rare ecosystems are meant to antagonize organisms in their respective environments. These are likely to be novel antimycobacterial compounds as they unknown to human pathogens.

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## Introduction

The worldwide problem caused by tuberculosis (TB) and the lack of new drugs in the market makes it imperative to search for novel drugs to fight efficiently against the rapid spread of multidrug-resistant TB. In this context, there is an urgent need for new antiTB drugs with less toxic side effects, improved pharmacokinetic properties, with extensive and potent activity against resistant strains and also drugs able to reduce the total duration of treatment (De Souza 2006).

Programmes aimed at the discovery of antibiotics and other bioactive metabolites from microbial sources have yielded an impressive number of compounds over the past 50 years, many of which have found applications in human medicine and agriculture (Busti et al. 2006). Actinomycetes are common soil inhabitants with an unprecedented ability to produce clinically useful antibiotics. Though more than 50% of the microbial antibiotics discovered so far originate from actinomycete bacteria, only a few soil-derived genera (Streptomyces and Micromonospora) account for most of these compounds. Of the 22,500 antibiotics reported from microbial sources, about 10,100 are reported from actinomycetes in which 800-1000 antibiotics are reported as antimycobacterials (Berdy 2005). From the discovery of streptomycin (the first antibiotic used for antiTB therapy) from Streptomyces griseus, numerous antiTB antibiotics such as kanamycin and rifampicin have been reported from actinomycetes of terrestrial origin (Kobarfard 2004).

Recently, the number of new compounds from terrestrial actinomycetes has decreased, whereas the rate of reisolation

of known compounds has increased (Lam 2006). Thus it is crucial that actinomycetes from unexplored or underexploited habitats be pursued as sources of novel bioactive secondary metabolites (Berdy 2005). With this view, the present investigation was initiated to screen actinomycetes isolated from less explored ecosystems like desert, forest soil and marine sediments for antibacterial and antimycobacterial activity.

# Materials and methods

### Actinomycete strains

Fifteen actinomycete strains used in this study were randomly selected from the Actinomycetes research laboratory of Sri Sankara Arts & Science College, Kanchipuram, Tamilnadu and were isolated from less explored ecosystems (Table 1) and maintained on yeast extract malt extract agar (ISP2 medium). They were characterized by the method described by Shirling and Gottileb (1966) and tentatively identified with the help of Nonomura's keys (1974). For marine actinomycete strains, all the media were prepared in filtered 50% sea water. Media used for the strain CA33 was prepared with the pH 10, since it is an alkalophilic actinomycete.

Production of bioactive compounds by submerged fermentation

All the actinomycete strains were inoculated on ISP2 agar plates and incubated at 28°C for 7 days for mass preparation of mycelium. After the growth, actinomycete mycelium was scraped by adding 10% Tween 80 and aseptically transferred into 50 ml of soybean meal inoculation medium (soybean meal 1%; glucose 1%; glycerol 1%; NaCl 0.5%; pH 7.0  $\pm$  0.2) prepared in 250-ml Erlenmeyer flasks and incubated in a rotary shaker at 120 rev/min for 48 h at 28°C. Then 10% of actinomycete inoculum was transferred to soybean meal production medium (soybean meal 1.5%; glucose 1.5%; glycerol 0.5%; NaCl 0.5%; CaCo<sub>3</sub> 0.10%; pH 7.0  $\pm$  0.2) and incubated in rotary shaker at 120 rev/ min for 72–96 h at 28°C (Radhakrishnan et al. 2007).

#### Preparation of crude extracts

After fermentation, the production medium was collected and centrifuged at 10,000 rev/min for 30 min at 4°C to separate the supernatant and mycelium. Dried mycelium was soaked with 50 ml of methanol for 12–18 h for the liberation of intracellular metabolites. Extracellular metabolites present in fermentation broth were extracted by liquid–liquid extraction using equal volume of ethyl acetate in a separating funnel for 24 h. Then the ethyl acetate and methanol portions were kept under reduced pressure for evaporation to obtain crude extracellular and intracellular metabolites, respectively (Radhakrishnan et al. 2007). Before extraction, about five ml of culture supernatant was collected in sterile vials to study its antimycobacterial activity.

Antibacterial activity of crude extracts

Crude ethyl acetate and methanol extracts were tested for antimicrobial activity by disc diffusion method against bacterial strains viz., *S. aureus*, *B. subtilis*, *E. coli*, *S. typhi*, and *P. aeruginosa*. All these strains were obtained from the Christian Medical College, Vellore, Tamil Nadu, India. About 250  $\mu$ g crude extract was impregnated on sterile 5 mm diameter filter paper disc and placed on the nutrient agar plates seeded with test organisms. The diameter of zone of inhibition was measured after 24 h of incubation at 37°C (Radhakrishnan et al. 2007).

Screening of actinomycetes for antimycobacterial activity

Actinomycete culture filtrates and solvent extracts were screened for antimycobacterial activity by luciferase reporter phage assay. Standard laboratory strain *M. tuberculosis*  $H_{37}$ Rv, SHRE sensitive and SHRE resistant clinical strains were used as test organisms. Both SHRE sensitive and SHRE resistant clinical *M. tuberculosis* isolates used in this study were isolated from TB patients at the Tuberculosis Research Centre, Chennai and maintained on Lowenstein-Jensen medium.

About five ml of actinomycete culture broth and crude solvent extracts (10 mg/ml dimethyl sulfoxide) were filtered using 0.45  $\mu$ m membrane filter and used for LRP assay. About 175  $\mu$ l of culture filtrate was added to 175  $\mu$ l of 2× Middlebrook 7H9 (supplemented with 10% albumin dextrose complex and 0.5% glycerol) medium. Working solutions of crude extracts were prepared in such a way that 350  $\mu$ l of 7H9 broth contained 500  $\mu$ g/ml of crude extracts. Sterile distilled water added to 2× medium served as control. Solvent control was also included. Suspension equivalent to 2 MacFarland units was prepared from log phase culture and 100  $\mu$ l of the same was added to all vials before incubating for 72 h at 37°C.

After incubation, 50  $\mu$ l of luciferase reporter phage phAE129 and 40  $\mu$ l of 0.1 M CaCl<sub>2</sub> were added to test and control vials. All the vials were incubated at 37°C for 4 h. After incubation 100  $\mu$ l from each vial was transferred to luminometer cuvette. About 100  $\mu$ l of D-Luciferin was added and relative light unit (RLU) was measured in luminometer (Sivakumar et al. 2007).

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Strains	Habitat	Cultural chan	acteris	tics		Micr	o morph	ology	Car	bon 1	tiliz	ation					Genera	
		AMC	RSP	SP	MP	SM	AM	SCM	ŋ	Α	Х	I	Μ	н	R	S R	I	
D10	Soil, Thar desert, Rajasthan	Dirty white	Ι	+	Ι	+	+	RA	+	Ι	+	+	+	+	+	 	Streptomyces	iygroscopicus
D5	Soil, Thar desert, Rajasthan	Dirty white	+	+	+	+	+	RF	+	+	+	+	+	+	+	+	S. griseorubig	snsou
CSA1	Soil, Coffee plantation, Kerala	Dirty white	+	+	Ι	+	+	RF	+	Ι	I	Ι	+	I		+	S. carnosus	
CSA14	Soil, Coffee plantation, Kerala	Gray	+	+	+	+	+	S	+	+	+	+	+	I		 +	S. griseofuscu.	7.4
CA33	Alkaline soil, cement factory area, Arakkonam, Tamil Nadu (TN)	Dirty white	I	I	I	+	+	RF	+	+	+	I	+	I	+	1	S. setonii	
NEK5	Leaves, Neem plant, Kanchipuram, TN	Gray	+	Ι	I	+	+	RF	+	Т	Т	Т	+	I	+	 +	S. finlayi	
EE9	Leaves, Eucalyptus plant, Kanchipuram, TN	White	+	Ι	Ι	+	+	RF	+	+	+	Ι	+	+	, T	 +	S. californicus	
MSU	Marine sediment, Palk strait, Thondi, TN	Ash	Ι	Ι	Ι	+	+	RF	+	+	+	+	+	+	+	+	S. libani	
ANS2	Marine sediment, Andaman	Ash	+	+	Ι	+	+	S	+	+	+	+	+	+	+	+	S. purpureus	
ANS5	Marine sediment, Andaman	Gray	Ι	Ι	Ι	+	+	RF	+	+	+	I	Т	+	· ·	1	S. halstendii	
R2	Marine sediment, Rameswaram, TN	Dirty white	Ι	+	Ι	+	+	RF	+	I	Ι	I	+1	I	+I	1	S. alboniger	
R9	Marine sediment, Rameswaram, TN	Gray	+	+	+	+	+	RF	+	+	+	+	+	+	+	+	S. phaeoviridi.	7.4
A3	Marine sediment, Pichavaram, TN	White	+	Ι	Ι	+	+	RF	+	Ι	$+\!\!\!+\!\!\!$	+	+	+		1	S. gedanensis	
RH5	Marine sediment, Pichavaram, TN	White	Ι	Ι	Ι	+	+	RF	+	Ι	Ι	Ι	Ι	+		1	S. goitieri	
M104	Marine sediment, Parangipettai, TN	Brown	+	Ι	T	+	I	М	+	Ι	I	Ι	+	+	, T	+	Micromonospo	<i>ra</i> sp.
<i>AMC</i> ає apertum	rrial mass color, $RSP$ reverse side pigment, $SP$ solul, $RF$ rectifications, $S$ spiral, $M$ monosporic, $+$ press	ole pigment, <i>M</i> ent, – absent.	P mela G gluc	anoid ose, ∕	pigme A arab	ent, <i>SM</i> inose,	' substrat X xylose	te myceliu , <i>I</i> inosito	$\operatorname{Im}, A$	M ae man	rial 1 nitol,	nycel F fri	ium, actos	SCM S, R 1	spor	e cha iose, j	n morphology, RA sucrose, Ra raffi	Retinaculum 10se

# **Results and discussion**

All the 15 actinomycete strains showed good growth on ISP2 agar as well as on soybean meal medium. Cultural, micromorphological and carbon utilization characteristics of all the 15 actinomycete strains were given in Table 1. Based on phenotypic characteristics 14 actinomycetes were identified as *Streptomyces* species and the other as *Micromonospora* species.

Traditionally, soil-derived actinomycetes have been most frequently screened for bioactive compounds. Unfortunately, the frequency of finding structurally new compounds from normal soil-derived actinomycetes is declining because of the redundancy in the isolation of known actinomycetes and antibiotics. Alternatively, actinomycetes from previously unexplored or under explored environments such as marine, desert and forest ecosystems are screened (Berdy 2005). In recent years, pathogenic microorganisms are gaining resistance against antimicrobial agents; hence the search for new, safe and more effective antimicrobial agents is an urgent need. Though there are many reports on actinomycete diversity in unique environments from deep sea to tall mountains, there are only few reports on bioactive compounds from actinomycetes of such origin. As a part of our continuous search, actinomycetes were isolated from various less explored ecosystems such as desert, forest, marine, plants and alkaline soils and screened for antibacterial and antimycobacterial activity.

In total 30 extracts (15 ethyl acetate extracts and 15 methanol extracts) prepared from 15 actinomycete strains were tested against bacterial strains, in which 10 ethyl acetate extracts and 7 methanol extracts showed antibacterial activity against at least one of the organisms tested

Strain No Extracts Test organisms (inhibition zone measured as diameter in mm) S. aureus B. subtilis E. coli S. typhi Pseu. aeruginosa D10 EAE 20 20 20 21 ME 14 16 15 11 D5 12 9 10 9 EAE 7 8 ME \_ CSA1 EAE \_ ME CSA14 EAE 14 ME \_ 9 10 CA33 EAE \_ ME 11 13 \_ \_ \_ NEK5 EAE 12 16 20 13 17 ME EE9 EAE ME \_ MSU EAE 8 \_ \_ ME 8 8 \_ \_ ANS2 EAE 14 15 10 13 15 ME 14 14 8 ANS5 EAE ME \_ \_ \_ 18 10 12 12 R2 EAE 9 12 ME 10 \_ R9 EAE ME A3 EAE ME RH5 EAE 12 ME \_ \_ \_ \_ M104 EAE 15 13 14 \_ \_ ME 16 9 12

**Table 2** Antibacterial activityof actinomycete extracts

*EAE* ethyl acetate extract, *ME* methanol extract

(Table 2). Both ethyl acetate and methanol extracts produced from 7 actinomycete strains showed antibacterial activity. Actinomycete strains viz., D10, D5, NEK5, ANS2, R2 and M104 showed prominent activity against the bacterial strains tested. Only NEK5 and CA33 were found to be active against *P. aeruginosa*. A great part of antibiotic compounds exhibit exclusive activity against gram-positive bacteria (app. 30% of all) and only 1.5% of the compounds (some 250 metabolites) showed activity only against gram-negative bacteria (Berdy 2005). In general most of the antibiotics are extracellular in nature (Augustine et al. 2005). But ethyl acetate and methanol extracts from actinomycete strains *viz.*, D10, D5, CA33, ANS2, MSU, M104 and R2 indicates the production of both extracellular and intracellular bioactive compounds.

In LRP assay, reduction in RLU by 50% or more compared to control was considered as antimycobacterial activity. Culture filtrates from seven actinomycete strains (CSA1, CSA14, CA33, NEK5, ANS2, MSU and R2) and ethyl acetate extracts of culture supernatants of 10 actinomycete strains (D10, CSA1, CSA14, NEK5, EE9, MSU, ANS2, R2, R9 and A3) showed good activity against all the three *M. tuberculosis* strains tested (Table 3). Mycelial methanol extracts of 12 actinomycete strains (D10, D5, CSA14, CA33, NEK5, MSU, ANS2, ANS5, R2, A3, Rh5 and M104) also inhibited all the three *M. tuberculosis* strains. Extraction using ethyl acetate additionally brought out the antimycobacterial activity of three more cultures, thus validating its capability as an ideal solvent for extraction of compounds from culture supernatants. This

finding corroborates an earlier report by Franco and Countinho (1991).

At present, the chemotherapeutic options for TB treatment have been restricted to a handful of compounds introduced 40-50 years ago, which must be administered in combination for extended periods. Finding of new lead compounds with novel antimycobacterial activity is urgent (De Souza 2006). M. tuberculosis is a slow growing organism and conventional methods require 3 to 4 weeks of incubation. Luciferase reporter phage (LRP) assay is a rapid, less laborious and less time-consuming method for high throughput screening of a large number of compounds for antimycobacterial activity. It shows great promise for the detection and drug susceptibility testing of mycobacteria with high sensitivity, specificity, in a timely and relatively 'low-tech' manner (Riska et al. 1999). Plant products and their derivatives were screened for antimycobacterial activity by adopting LRP assay (Sivakumar et al. 2007).

There are only few reports so far on in vitro antituberculous activity of organisms of marine origin. Messetolide A and viscosin are cyclic depsipeptides from cultures of two *Pseudomonas* species isolated from a marine alga and tube worm, respectively (El Sayed et al. 2000). Marinederived antibiotics may be more efficient at fighting infections because the terrestrial bacteria have not developed resistance against them (Saha et al. 2006). Actinomycetes strains MSU and ANS2 reported in this study show promise as novel antimycobacterial compounds from marine origin.

Table 3 Percentage RLU reduction by actinomycetes extracts in LRP assay

Actinomycete	Culture fi	ltrate		Ethyl ace	tate extract		Methanol extract		
strains	H <sub>37</sub> Rv	SHRE resistant MTB	SHRE sensitive MTB	H <sub>37</sub> Rv	SHRE resistant MTB	SHRE sensitive MTB	H <sub>37</sub> Rv	SHRE resistant MTB	SHRE sensitive MTB
D10	21.6	0	57.6	97.4	97.6	95.2	83.7	74.3	52.5
D5	35.3	78.4	67	75.7	36.2	69.5	90.6	97.1	97.9
CSA1	64.7	78.9	78.4	88.1	84.2	75.4	27.4	0	13
CSA14	90	62.6	87.3	91.8	92.1	63.8	79.2	97.3	98.5
CA33	97.9	98.1	91.9	59.7	38.8	55.2	88.7	52.5	66.6
NEK5	88.8	91.7	91.7	88.4	79.7	54.2	87.1	77.7	70.0
EE9	46.1	46.8	67.6	91.6	83.4	55.2	33.1	0	0
MSU	92.4	98.3	97.2	90.9	97.0	84.7	80.1	96.3	84.7
ANS2	93.1	98.8	96.9	82.6	96.2	98.7	91.8	97.4	98.2
ANS5	0	0	0	81.3	36.2	64.3	60.6	76.3	76.2
R2	83.8	87.5	79.3	62.1	73.9	74.4	82.1	84.2	83.1
R9	0	20.2	0	84.9	84.1	76.5	56.3	35.4	55.3
A3	37.2	50.7	0	88.9	70.5	84.3	75.5	51.1	65.7
RH5	37.2	50.7	0	2.2	3.1	0	81.1	77.7	75.6
M104	76.2	32	26.5	17.7	51.1	57.6	78.3	84.1	83.2

The restricted role of INH in therapy of infections caused by other diseases restricts the emergence of cross resistance due to rampant use. While crude extracts of some strains demonstrate antimycobacterial and antibacterial activity, extracts from actinomycete strains A3, CSA1, EE9, ANS5 and R9 showed antimycobacterial activity alone. These strains are promising source of antimycobacterial compounds that can act specifically on mycobacteria.

Early stage drug discovery is a key bottleneck in the pipeline to find novel drugs. Of the 1,556 new chemical entities marketed worldwide between 1975 and 2004, only three were for TB (Casenghi et al. 2007). Actinomycetes strains viz., D10, D5 (desert), CSA14 (forest), CA33 (alkaline soil), NEK5 (Neem plant), MSU, ANS2, R2 and M104 (marine) screened in the present study were found to be highly potent. Purification of active compounds that selectively act on conserved targets may pave the way for highly effective antibiotics.

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