



# Unexplored hypersaline habitats are sources of novel actinomycetes

Polpass Arul Jose and Solomon Robinson David Jebakumar\*

Department of Molecular Microbiology, School of Biotechnology, Madurai Kamaraj University, Madurai, India  
\*Correspondence: jsolomon\_mrna@yahoo.com

**Edited by:**

Luis Cláudio Nascimento Da Silva, University of Copenhagen, Denmark

**Reviewed by:**

Atte Von Wright, University of Eastern Finland, Finland  
Dmitri Debabov, NovaBay Pharmaceuticals, USA

**Keywords:** actinomycetes, hypersaline environments, antibiotic, *Streptomyces*, drug discovery

Pathogenic microorganisms have evolved sophisticated mechanisms to inactivate antibiotics and rendered an urgent need for new antibiotics that would target the emerging multidrug-resistance (Butler et al., 2013). Consequently, search for novel sources of potent antibiotics is desperately needed to develop potent drugs. Microbial resources have made an incredible contribution to the antibiotic drug discovery and development process over the last seven decades (Demain and Sanchez, 2009). In particular, actinomycetes are the most important source of bioactive natural compounds with a long track record of producing novel molecules, including several commercially important drugs like Streptomycin, Gentamicin, Vancomycin, Clindamycin, Erythromycin, Amphotericin, Rifampicin, and Tetracycline. More novel molecules with potential therapeutic applications are still on the row to be discovered from these actinomycetes (Baltz, 2007).

Microbial assemblages and their species distributions in specific environments are mostly determined by their specific environmental conditions, which may translate into novel chemistry (Genilloud et al., 2011). The actinomycetes that produce antibiotics are abundant in soil. In recent decades, isolation and exploitation of actinomycetes for novel compounds from conventional environments have led to rediscovery of known compounds (Walsh, 2003; Tulp and Bohlin, 2005). However, diverse actinomycetes from poorly studied unusual environments promises a raise in the prospect of discovering novel compounds with potential activities that can be developed as a resource for drug

discovery (Peraud et al., 2009; Jose et al., 2011; Poulsen et al., 2011; Carr et al., 2012; Subramani and Aalbersberg, 2012; Hamed et al., 2013; Jose et al., 2013; Yuan et al., 2014). In this respect, current actinomycetes isolation programs are reoriented toward largely unexplored, unusual and extreme environments like hypersaline marine environments, extreme inland saline zones, volcanic zones, hyperarid and glaciers (Hamed et al., 2013; Jose and Jebakumar, 2013b).

The extreme habitats are characterized by chemical or physical conditions that differ significantly from those found in environments that support more abundant and varied life forms (Zhao et al., 2011). Hypersaline environments are typical extreme habitats, in which the high salt concentration, alkalinity, and low oxygen concentrations are environmental factors that may limit their biodiversity (Ventosa, 2006). There are several reports on inhabitation of microorganisms including actinomycetes in diverse hypersaline environments such as salt lakes (Swan et al., 2010; Guan et al., 2011; Phillips et al., 2012), solar salterns (Baati et al., 2008; Sabet et al., 2009), salt mines (Chen et al., 2007) and brine wells (Xiang et al., 2008). However, these hypersaline environments remain largely unexplored as a source of novel actinomycetes in many tropical and subtropical regions throughout the world. In a recent review, Hamed et al. (2013) described that actinomycetes form a stable, metabolically active and persistent population in various marine hypersaline ecosystem and it is accepted that halophilic actinomycetes will provide a valuable resource for novel

products of industrial interest, including antimicrobial, cytotoxic, neurotoxic, antimitotic, antiviral and antineoplastic activities. Series of bioactive metabolites like Pyrostatins, Salinosporamides, Abyssomicins, Trioxacarcin A, Gutingimycin, Sporolides, Marinomycins, Himalomycins, Diazepinomicin, Helquinoline, Lajollamycin, Tetrodotoxin, Mechercharmycins, Cyanosporasides, Erythronolides, and Ammosamide D have been reported from this physiological group of actinomycetes (Hamed et al., 2013). In continued efforts to discover novel natural products from new extremophilic actinomycetes, Zhao et al. (2011) discovered new linear polyketides, actinopolysporins A, B, and C, as well as the known antineoplastic antibiotic tubercidin from the halophilic actinomycete *Actinopolyspora erythraea* YIM 90600. Tian et al. (2013) reported the isolation and characterization of *p*-Terphenyls with antifungal, antibacterial, and antioxidant activities from halophilic actinomycete *Nocardiopsis gilva* YIM 90087. Jang et al. (2013) declared a significant discovery of new antibiotic, anthracimycin, which is produced by a marine-derived actinomycete in saline culture. More recently, an antimicrobial quinoline alkaloid has been isolated from a novel halophilic actinomycete, *Nocardiopsis terrae* YIM 90022 (Tian et al., 2014).

In accordance with the above reports, we initiated a research program which was designed to discover new natural products from novel actinomycetes originating from previously unexplored hypersaline environments. Saltpan soil samples were collected aseptically from

**Table 1 | Range of antimicrobial activities of the actinomycete isolates obtained from Indian inland (Jose and Jebakumar, 2013a) and coastal (Jose and Jebakumar, 2013b) solar salterns.**

Isolates	Antimicrobial activities							
	Antibacterial activity					Antifungal activity		
	Gram—positive		Gram—negative			<i>Aspergillus niger</i>	<i>Fusarium oxysporum</i>	<i>Alternaria alternata</i>
	<i>Staphylococcus aureus</i>	<i>Bacillus subtilis</i>	<i>Klebsiella pneumoniae</i>	<i>Proteus vulgaris</i>	<i>Salmonella typhi</i>			
<i>Streptomyces</i> sp. JAJ06	+	+	+	+	+	—	—	—
<i>Streptomyces</i> sp. JAJ07	+	+	+	+	+	—	—	—
<i>Streptomyces</i> sp. JAJ13	+	+	+	+	+	—	—	—
<i>Nonomuraea</i> sp. JAJ18	+	+	+	+	+	—	—	—
<i>Streptomyces</i> sp. JAJ19	+	+	+	+	+	—	—	—
<i>Micromonospora</i> sp. JAJ20	+	+	+	+	+	+	+	+
<i>Streptomyces</i> sp. JAJ28	—	—	—	—	—	+	+	+
<i>Streptomyces</i> sp. JAJ41	+	+	—	—	—	—	—	—
<i>Streptomyces</i> sp. JAJ59	+	+	+	+	+	—	—	—
<i>Actinoalloteichus</i> sp. JAJ70	+	+	+	+	+	+	+	+
<i>Streptomyces</i> sp. JAJ73	—	—	—	—	—	+	+	+
<i>Pseudonocardia</i> sp. JAJ77	+	+	+	+	+	—	—	—
<i>Pseudonocardia</i> sp. JAJ82	—	—	—	—	—	+	+	+

+, positive for antimicrobial activity; —, negative for antimicrobial activity.

saltpans of coastal solar saltern ponds at Arumuganeri, Tuticorin (about Latitude 8.43 N and Longitude 78.60 E) and inland solar salterns at Sambhar Lake, Jaipur (about Latitude 26.94561 N and Longitude 75.20968 E) in India. Physicochemical characteristics from this study confirmed that this environment is a hypersaline zone. In the course of our program, diverse actinomycetes were isolated by using selective isolation methods which employed stamping, heat and dilution, and succeeded with isolation of both *Streptomyces* (common actinomycete) and non-*Streptomyces* including rare actinomycetes *Nonomuraea*, *Actinoalloteichus* and *Pseudonocardia* (Jose and Jebakumar, 2012, 2013a). A total of 83 actinomycetes were isolated and assigned to eight genera (*Streptomyces*, *Micromonospora*, *Nocardia*, *Nocardioiopsis*, *Nonomuraea*, *Saccharopolyspora*, *Pseudonocardia* and *Actinoalloteichus*) on the basis of their 16S rDNA sequences and it was the first step toward better understanding of actinomycete community from solar saltern ponds in India. Our preliminary biological activity screening and subsequent structural characterisation experiments suggested the occurrence of antimicrobial

compounds producing *Streptomyces* and rare actinomycetes and confirmed that hypersaline solar salterns are reservoirs of antibiotic producing actinomycetes (Jose and Jebakumar, 2013a,b). The range of antimicrobial activities exerted by different actinomycete isolates were summarized in **Table 1**. In concise, our continuing study contributes in acquaintance of solar saltern associated actinobacteria and further augments the array of actinomycetes available for antibiotic discovery programs.

-In conclusion, escalating reports on discovery of diverse natural compounds from halophilic and halotolerant actinomycetes which inhabit in hypersaline environments suggested that this physiological group has enormous capacity to produce array of secondary metabolites with disparate activities. Indeed, hypersaline environments warrant significant attention as habitats of novel actinomycetes and unlock new avenues for natural products discovery.

#### ACKNOWLEDGMENTS

Our research program on actinomycetes is being supported by University Grants Commission (UGC) and Council of

Scientific and Industrial Research (CSIR), India.

#### REFERENCES

- Baati, H., Guermazi, S., Amdouni, R., Gharsallah, N., Sghir, A., and Ammar, E. (2008). Prokaryotic diversity of a Tunisian multipond solar saltern. *Extremophiles* 4, 505–518. doi: 10.1007/s00792-008-0154-x
- Baltz, R. H. (2007). Antimicrobials from actinomycetes: back to the future. *Microbe* 2, 125–131.
- Butler, M. S., Blaskovich, M. A., and Cooper, M. A. (2013). Antibiotics in the clinical pipeline in 2013. *J. Antibiot.* 66, 571–591. doi: 10.1038/ja.2013.86
- Carr, G., Poulsen, M., Klassen, J. L., Hou, Y., Wyche, T. P., Bugni, T. S., et al. (2012). Microtermolides A and B from Termite-associated *Streptomyces* sp. and structural revision of vinylamycin. *Org. Lett.* 14, 2822–2825. doi: 10.1021/ol301043p
- Chen, Y. G., Cui, X. L., Pukall, R., Li, H. M., Yang, Y. L., Xu, L. H., et al. (2007). *Salinicoccus kunmingensis* sp. nov., a moderately halophilic bacterium isolated from a salt mine in Yunnan, south-west China. *Int. J. Syst. Evol. Microbiol.* 57, 2327–2332. doi: 10.1099/ijs.0.64783-0
- Demain, A. L., and Sanchez, S. (2009). Microbial drug discovery: 80 years of progress. *J. Antibiot.* 62, 5–16. doi: 10.1038/ja.2008.16
- Genilloud, O., González, I., Salazar, O., Martín, J., Tormo, J. R., and Vicente, F. (2011). Current approaches to exploit actinomycetes as a source of novel natural products. *J. Ind. Microbiol. Biotechnol.* 38, 375–389. doi: 10.1007/s10295-010-0882-7
- Guan, T. W., Wu, N., Xia, Z. F., Ruan, J. S., and Zhang, X. P. (2011). *Saccharopolyspora lacisalsi* sp.

- nov., a novel halophilic actinomycete isolated from a salt lake in Xinjiang, China. *Extremophiles* 15, 373–378. doi: 10.1007/s00792-011-0369-0
- Hamed, J., Mohammadipanah, F., and Ventosa, A. (2013). Systematic and biotechnological aspects of halophilic and halotolerant actinomycetes. *Extremophiles* 17, 1–13. doi: 10.1007/s00792-012-0493-5
- Jang, K. H., Nam, S.-J., Locke, J. B., Kauffman, C. A., Beatty, D. S., Paul, L. A., et al. (2013). Anthracimycin, a potent Anthrax antibiotic from a marine-derived actinomycete. *Angew. Chem. Int. Ed. Engl.* 52, 7822–7824. doi: 10.1002/anie.201302749
- Jose, P. A., and Jebakumar, S. R. D. (2012). Phylogenetic diversity of actinomycetes cultured from coastal multipond solar saltern in Tuticorin, India. *Aquat. Biosyst.* 8:23. doi: 10.1186/2046-9063-8-23
- Jose, P. A., and Jebakumar, S. R. D. (2013a). Phylogenetic appraisal of antagonistic, slow growing actinomycetes isolated from hypersaline inland solar salterns at Sambhar salt Lake, India. *Front. Microbiol.* 4:190. doi: 10.3389/fmicb.2013.00190
- Jose, P. A., and Jebakumar, S. R. D. (2013b). Diverse actinomycetes from Indian coastal solar salterns - a resource for antimicrobial screening. *J. Pure Appl. Microbiol.* 7, 2569–2575.
- Jose, P. A., Santhi, V. S., and Jebakumar, S. R. D. (2011). Phylogenetic-affiliation, antimicrobial potential and PKS gene sequence analysis of moderately halophilic *Streptomyces* sp. inhabiting an Indian saltpan. *J. Basic Microbiol.* 51, 348–356. doi: 10.1002/jobm.201000253
- Jose, P. A., Sivakala, K. K., and Jebakumar, S. R. D. (2013). Formulation and statistical optimization of culture medium for improved production of antimicrobial compound by *Streptomyces* sp. JAJ06. *Int. J. Microbiol.* 2013:526260. doi: 10.1155/2013/526260
- Peraud, O., Biggs, J. S., Huguen, R. W., Light, A. R., Concepcion, G. P., Concepcion, G. P., et al. (2009). Microhabitats within venomous cone snails contain diverse Actinobacteria. *Appl. Env. Microbiol.* 75, 6820–6826. doi: 10.1128/AEM.01238-09
- Phillips, K., Zaidan, F., Elizondo, O. R., and Lowe, K. L. (2012). Phenotypic characterization and 16S rDNA identification of culturable non-obligate halophilic bacterial communities from a hypersaline lake, La Sal del Rey, in extreme South Texas (USA). *Aquat. Biosyst.* 8:5. doi: 10.1186/2046-9063-8-5
- Poulsen, M., Oh, D.-C., Clardy, J., and Currie, C. R. (2011). Chemical Analyses of Wasp-Associated *Streptomyces* bacteria reveal a prolific potential for natural products discovery. *PLoS ONE* 6:e16763. doi: 10.1371/journal.pone.0016763
- Sabet, S., Diallo, L., Hays, L., Jung, W., and Dillon, J. G. (2009). Characterization of halophiles isolated from solar salterns in Baja California, Mexico. *Extremophiles* 13, 643–656. doi: 10.1007/s00792-009-0247-1
- Subramani, R., and Aalbersberg, W. (2012). Marine actinomycetes: an ongoing source of novel bioactive metabolites. *Microbiol. Res.* 167, 571–580. doi: 10.1016/j.micres.2012.06.005
- Swan, B. K., Ehrhardt, C. J., Reifel, K. M., Moreno, L. I., and Valentine, D. L. (2010). Archaeal and bacterial communities respond differently to environmental gradients in anoxic sediments of a California hypersaline lake, the Salton Sea. *Appl. Environ. Microbiol.* 76, 757–768. doi: 10.1128/AEM.02409-09
- Tian, S., Yang, Y., Liu, K., Xiong, Z., Xu, L., and Zhao, L. (2014). Antimicrobial metabolites from a novel halophilic actinomycete *Nocardiopsis terrae* YIM 90022. *Nat. Prod. Res.* 28, 344–346. doi: 10.1080/14786419.2013.858341
- Tian, S.-Z., Pu, X., Luo, G., Zhao, L.-X., Xu, L.-H., Li, W.-J., et al. (2013). Isolation and characterization of new *p*-Terphenyls with antifungal, antibacterial, and antioxidant activities from halophilic actinomycete *Nocardiopsis gilva* YIM 90087. *J. Agr. Food Chem.* 61, 3006–3012. doi: 10.1021/jf400718w
- Tulp, M., and Bohlin, L. (2005). Rediscovery of known natural compounds: nuisance or goldmine? *Bioorg. Med. Chem.* 13, 5274–5282. doi: 10.1016/j.bmc.2005.05.067
- Ventosa, A. (2006). “Unusual microorganisms from unusual habitats: hypersaline environments,” in *Prokaryotic Diversity – Mechanisms and Significance* (Society for General Microbiology Symposium no. 66), eds N. A. Logan, H. M. Lappin-Scott, and P. C. F. Oyston (Cambridge: Cambridge University Press), 223–253.
- Walsh, C. (2003). Where will new antibiotics come from? *Nature Rev. Microbiol.* 1, 65–70. doi: 10.1038/nrmicro727
- Xiang, W., Guo, J., Feng, W., Huang, M., Chen, H., Zhao, J., et al. (2008). Community of extremely halophilic bacteria in historic Dagong Brine Well in southwestern China. *World J. Microbiol. Biotechnol.* 24, 2297–2305. doi: 10.1007/s11274-008-9744-0
- Yuan, M., Yu, Y., Li, H.-R., Dong, N., and Zhang, X.-H. (2014). Phylogenetic diversity and biological activity of actinobacteria isolated from the chukchi shelf marine sediments in the Arctic Ocean. *Mar. Drugs* 12, 1281–1297. doi:10.3390/md12031281
- Zhao, L. X., Huang, S. X., Tang, S. K., Jiang, C. L., Duan, Y., Beutler, J. A., et al. (2011). Actinopolysporins A–C and Tubercidin as a Pcd4 stabilizer from the halophilic actinomycete *Actinopolyspora erythraea* YIM 90600. *Nat. Prod.* 74, 1990–1995. doi: 10.1021/np200603g

**Conflict of Interest Statement:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Received: 07 April 2014; accepted: 04 May 2014; published online: 27 May 2014.

Citation: Jose PA and Jebakumar SRD (2014) Unexplored hypersaline habitats are sources of novel actinomycetes. *Front. Microbiol.* 5:242. doi: 10.3389/fmicb.2014.00242

This article was submitted to *Antimicrobials, Resistance and Chemotherapy*, a section of the journal *Frontiers in Microbiology*.

Copyright © 2014 Jose and Jebakumar. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) or licensor are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.