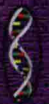




N
o
v
a

B
i
o
m
e
d
i
c
a
l



MOLECULAR BIOLOGY OF BACTERIA

RAJARSHI KUMAR GAUR
HEMANT K. GAUTAM
EDITORS

Microbiology Research Advances

NOVA

MICROBIOLOGY RESEARCH ADVANCES

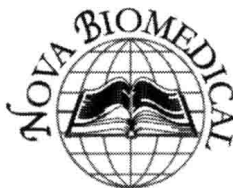
MOLECULAR BIOLOGY OF BACTERIA

RAJARSHI KUMAR GAUR

AND

HEMANT K. GAUTAM

EDITORS



New York

Copyright © 2013 by Nova Science Publishers, Inc.

All rights reserved. No part of this book may be reproduced, stored in a retrieval system or transmitted in any form or by any means: electronic, electrostatic, magnetic, tape, mechanical photocopying, recording or otherwise without the written permission of the Publisher.

For permission to use material from this book please contact us:

Telephone 631-231-7269; Fax 631-231-8175

Web Site: <http://www.novapublishers.com>

NOTICE TO THE READER

The Publisher has taken reasonable care in the preparation of this book, but makes no expressed or implied warranty of any kind and assumes no responsibility for any errors or omissions. No liability is assumed for incidental or consequential damages in connection with or arising out of information contained in this book. The Publisher shall not be liable for any special, consequential, or exemplary damages resulting, in whole or in part, from the readers' use of, or reliance upon, this material. Any parts of this book based on government reports are so indicated and copyright is claimed for those parts to the extent applicable to compilations of such works.

Independent verification should be sought for any data, advice or recommendations contained in this book. In addition, no responsibility is assumed by the publisher for any injury and/or damage to persons or property arising from any methods, products, instructions, ideas or otherwise contained in this publication.

This publication is designed to provide accurate and authoritative information with regard to the subject matter covered herein. It is sold with the clear understanding that the Publisher is not engaged in rendering legal or any other professional services. If legal or any other expert assistance is required, the services of a competent person should be sought. FROM A DECLARATION OF PARTICIPANTS JOINTLY ADOPTED BY A COMMITTEE OF THE AMERICAN BAR ASSOCIATION AND A COMMITTEE OF PUBLISHERS.

Additional color graphics may be available in the e-book version of this book.

Library of Congress Cataloging-in-Publication Data

Molecular biology of bacteria / [edited by] Rajarshi Kumar Gaur (Department of Science, Faculty of Arts, Science and Commerce, Mody Institute of Technology and Science, Rajasthan, India) and Hemant K. Gautam (Institute of Genomics and Integrative Biology, Delhi, India).

pages cm

Includes bibliographical references and index.

ISBN 978-1-62618-189-2 (hardcover)

1. Bacterial genetics. 2. Bacterial genomes. 3. Bacteria. I. Gaur, Rajarshi Kumar, editor of compilation. II. Gautam, Hemant K., editor of compilation.

QH434.M638 2013

572.8'6293--dc23

2013005769

Published by Nova Science Publishers, Inc. † New York

MICROBIOLOGY RESEARCH ADVANCES

MOLECULAR BIOLOGY OF BACTERIA

MICROBIOLOGY RESEARCH ADVANCES

Additional books in this series can be found on Nova's website
under the Series tab.

Additional e-books in this series can be found on Nova's website
under the e-book tab.

BACTERIOLOGY RESEARCH DEVELOPMENTS

Additional books in this series can be found on Nova's website
under the Series tab.

Additional e-books in this series can be found on Nova's website
under the e-book tab.

Preface

Bacteria are not just important as laboratory tools to understand higher organisms; they are important and interesting in their own right. For instances, they play an essential role in the ecology of earth. They are the only organisms that can fix atmospheric nitrogen, that is convert N_2 to ammonia, which can be used to make nitrogen-containing cellular constituents such as proteins and nucleic acids. Moreover bacteria produce most of the naturally occurring so-called greenhouse gases, such as methane and carbon dioxide, which are in turn used by other types of bacteria.

Bacteria are also worth studying because of their role in disease. They cause many human, plant and animal diseases and new disease are continuously appearing. Knowledge gained from the molecular genetics of bacteria will help in the development of new ways to treat or otherwise control old diseases, as well as new ones.

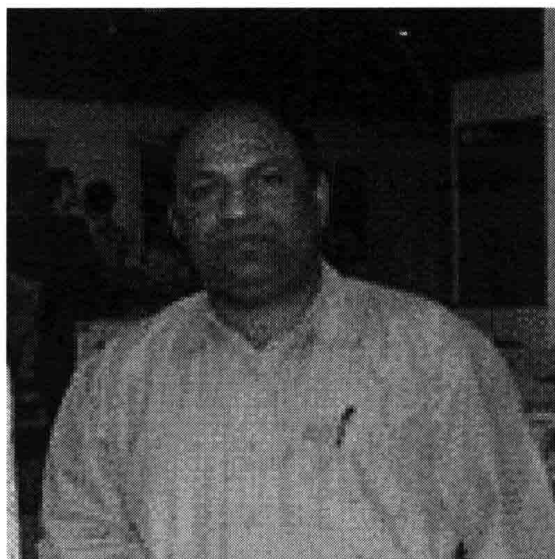
Molecular Biology has proved to be one of the more fruitful technological approaches sciences, being both very powerful and able to generate valuable intellectual property. This book aims to present examples in the application molecular biology and genetic engineering in bacteriology. This book shows diverse roles of bacteria in the ecosystems and it gives significant contributions from biotechnology approaches. The challenges of these aspects and methodologies inspired us to edit the book entitled *Molecular Biology of Bacteria*.

Rajarshi Kumar Gaur
and
Hemant K. Gautam
Editors

About the Editors



Dr. R.K. Gaur is presently working as Head and Assistant Professor, Department of Science, Mody Institute of Technology and Science (Deemed University), Lakshmangarh, Sikar, Rajasthan. He did his Ph.D on molecular characterization of sugarcane viruses of India. He partially characterized three sugarcane virus viz., sugarcane mosaic virus, sugarcane streak mosaic virus and sugarcane yellow luteovirus. He received MASHAV fellowship in 2004 of Israel government for his post-doctoral studies and joined The Volcani Centre, Israel and then shifted to Ben Gurion University, Negev, Israel. In 2007 he received the Visiting Scientist Fellowship from Swedish Institute Fellowship, Sweden for one year to work in the The Umeå University, Umeå, Sweden. He is also a recipient of ICGEB, Italy Post-Doctoral fellowship in 2008. He worked on development of marker-free transgenic plant against cucumber viruses. He has made significant contributions on plant microbes and published 56 national/international papers and presented near about 45 papers in the national and international conferences. He has also visited Thailand, New Zealand, London and Italy for the sake of attending the conference/workshop. He is also a member of national and international societies.



Dr. Hemant K. Gautam obtained his Masters and Ph.D. degree in microbiology from the Indian Agricultural Research Institute, New Delhi. He did post doctorate from France and Israel. He is currently working as a Principal Scientist at Institute of Genomics and Integrative Biology (CSIR), Delhi. He is associated with number of Universities and members of various scientific and academic organizations. He is a recipient of Bharat Excellence award, Biotechnology award, IFP France & Israel Govt. fellowship and UNESCO fellow. He published more than 60 research papers and over 75 new sequences have been submitted to NCBI database. He has a distinguish background in the field of microbial biotechnology and natural products.

List of Contributors

Arinthip Thamchaipenet	Actinobacterial Research Unit, Department of Genetics, Faculty of Science, Kasetsart University, Chatuchak, Bangkok 10900, Thailand Center for Advanced Studies in Tropical Natural Resources, National Research University-Kasetsart University (CASTNAR, NRU-KU), Chatuchak, Bangkok 10900, Thailand
Chakrit Bunyoo	Actinobacterial Research Unit, Department of Genetics, Faculty of Science, Kasetsart University, Chatuchak, Bangkok 10900, Thailand
Ratchaniwan Jaemsaeng	Actinobacterial Research Unit, Department of Genetics, Faculty of Science, Kasetsart University, Chatuchak, Bangkok 10900, Thailand Center for Advanced Studies in Tropical Natural Resources, National Research University-Kasetsart University (CASTNAR, NRU-KU), Chatuchak, Bangkok 10900, Thailand
Karan Lomaneeratana	Actinobacterial Research Unit, Department of Genetics, Faculty of Science, Kasetsart University, Chatuchak, Bangkok 10900, Thailand
Siriwan Rungin	Actinobacterial Research Unit, Department of Genetics, Faculty of Science, Kasetsart University, Chatuchak, Bangkok 10900, Thailand
Somkiat Phornphisutthimas	Department of Biology, and Research Unit on Science, Technology and Environment for Learning, Faculty of Science, Srinakharinwirot University, Sukhumvit 23, Bangkok 10110, Thailand

Pankaj Kumar Jain	Biotechnology Laboratory, Department of Science, MITS University, Lakshmangarh, Sikar-332311, Rajasthan, India
Vivek Bajpai	Department of Microbiology, ITM University, Gwalior, MP, India
Madan Lowry	Center for advance studies, Department of Zoology, University of Rajasthan, Jaipur- 302004, Rajasthan, India
Rajal Debnath	Biotechnology Division, CSIR- North East Institute of Science & Technology, Jorhat-785006, Assam, India
Rupak K. Sarma	Biotechnology Division, CSIR- North East Institute of Science & Technology, Jorhat-785006, Assam, India
Ratul Saikia	Biotechnology Division, CSIR- North East Institute of Science & Technology, Jorhat-785006, Assam, India
Archana Yadav	Biotechnology Division, CSIR- North East Institute of Science & Technology, Jorhat-785006, Assam, India
Tarun C Bora	Biotechnology Division, CSIR- North East Institute of Science & Technology, Jorhat-785006, Assam, India
Kashyap Kumar Dubey	Microbial Biotechnology Laboratory, University Institute of Engineering and Technology, M.D. University Rohtak, Rohtak-124001 Haryana, India
Dhirendra Kumar	Microbial Biotechnology Laboratory, University Institute of Engineering and Technology, M.D. University Rohtak, Rohtak-124001 Haryana, India
Anamika Jha	Ashok and Rita Patel Institute of integrated Study and Research in Biotechnology and Allied Sciences, New Vallabh Vidyanagar, Anand- 388121 (Gujarat), India
Hemant K. Gautam	Institute of Genomics and Integrative Biology Mall Road, University Campus, Delhi-110 007 India
Nisha Daxini	Ashok and Rita Patel Institute of integrated Study and Research in Biotechnology and Allied Sciences, New Vallabh Vidyanagar, Anand- 388121 (Gujarat), India
Sakshi Issar	Department of Science, faculty of Arts, Science and Commerce, Mody Institute of Technology and Science- Lakshmangarh, Sikar-332311, Rajasthan, India

D.K.Choudhary	Department of Science, faculty of Arts, Science and Commerce, Mody Institute of Technology and Science-Lakshmangarh, Sikar-332311, Rajasthan, India
R.K.Gaur	Department of Science, faculty of Arts, Science and Commerce, Mody Institute of Technology and Science-Lakshmangarh, Sikar-332311, Rajasthan, India
Komal Saxena	Institute of Biomedical Education & Research, Mangalayatan University, Aligarh – Uttar Pradesh, India
Girijesh Kumar Patel	Department of Biotechnology, Graphic Era University, 566/6, Bell Road, Clement, Town, Dehradun, Uttarakhand, India
G.K. Aseri	Amity Institute of Biotechnology, Amity University Rajasthan, Jaipur, India
Vikash Babu	Department of Biotechnology, Graphic Era University, 566/6, Bell Road, Clement, Town, Dehradun, Uttarakhand, India
Chand Ram	Bioremediation Laboratory, Dairy Microbiology division, National Dairy Research Institute, Karnal-132001 (Haryana), India
Vijay Kumar	Bioremediation Laboratory, Dairy Microbiology division, National Dairy Research Institute, Karnal-132001 (Haryana), India
M.K.Chattopadhyay	Centre for Cellular and Molecular Biology (CSIR) Hyderabad 500007 India

Contents

Preface		vii
About the Editors		ix
List of Contributors		xi
Chapter 1	Intergeneric Conjugation: A Practical Method for Genetic Manipulation in Actinomycetes <i>Arinthip Thamchaipenet, Chakrit Bunyoo, Ratchaniwan Jaemsaeng, Karan Lohmaneeratana, Siriwan Rungin and Somkiat Phornphisutthimas</i>	1
Chapter 2	Metagenomics: A Hunting Expedition in Microbial Diversity <i>Rajal Debnath, Rupak K. Sarma, Ratul Saikia, Archana Yadav and Tarun C. Bora</i>	19
Chapter 3	Molecular Biology of Nitrile Metabolizing Enzymes <i>Komal Saxena, Ashish Deep Gupta, Girijesh Kumar Patel and Vikash Babu</i>	31
Chapter 4	Multistress-Protective Molecules in Bacteria <i>M. K. Chattopadhyay</i>	55
Chapter 5	Food Microbiome Research: New Insights into Function <i>Anamika Jha, Nisha Daxini and Hemant K. Gautam</i>	63
Chapter 6	Microorganism and Key Degradative Enzymes: Tools of Remediation of Heavy Metals and PAHs <i>Pankaj Kumar Jain, Vivek Bajpai and Madan Lowry</i>	77
Chapter 7	Nanocatalysis: A Prospective and Sustainable Approach for the Chemical World <i>Kashyap Kumar Dubey and Dharendra Kumar</i>	117
Chapter 8	Probiotics and Herbs: Emerging Treatment Options for Non Alcoholic Fatty Liver Diseases <i>Chand Ram and Vijay Kumar</i>	125

Chapter 9	Understanding the Mysterious Radioresistance and Genomic Integrity of <i>Deinococcus radiodurans</i> against Gamma Radiation: Implications in Radiation Response Modulation in Higher Organisms	143
	<i>Raj Kumar and Hemant K. Gautam</i>	
Chapter 10	Nitrogen Fixation Networking in <i>Pseudomonas Putida</i>	157
	<i>Sakshi Issar, Hemant Gautam, D.K. Choudhary and R.K. Gaur</i>	
Chapter 11	Quorum Sensing in <i>Pseudomonas Aeruginosa</i> : Cell to Cell Signaling at Molecular Level	173
	<i>Sakshi Issar and R.K. Gaur</i>	
Index		191

Chapter 1

Intergeneric Conjugation: A Practical Method for Genetic Manipulation in Actinomycetes

***Arinthip Thamchaipenet^{1,2*}, Chakrit Bunyoo¹,
Ratchaniwan Jaemsaeng^{1,2}, Karan Lohmaneeratana¹,
Siriwan Rungin¹ and Somkiat Phornphisutthimas³***

¹Actinobacteria Research Unit, Department of Genetics,
Faculty of Science, Kasetsart University, Chatuchak, Bangkok, Thailand

²Center for Advanced Studies in Tropical Natural Resources,
National Research University-Kasetsart University
(CASTNAR, NRU-KU), Chatuchak, Bangkok, Thailand

³Department of Biology, and Research Unit on Science,
Technology and Environment for Learning, Faculty of Science,
Srinakharinwirot University, Bangkok, Thailand

Abstract

The system for intergeneric conjugation has been developed for many actinomycete species including *Streptomyces* and rare actinomycetes. *E. coli* donor strains generally used are the methylation proficient strain, S17-1 harbouring *tra* genes on its chromosome and the methylation-deficient strain, ET12567 containing *tra* genes in a self-transmissible plasmid pUB307 or a non-transmissible plasmid pUZ8002. Integrative plasmids used are normally non-replicative plasmids carrying the *oriT* for the propose of gene disruption (e.g. pSET151, pIJ8671) or with additional ϕ C31 *attP-int* system in the case of gene expression (e.g. pSET152, pIJ8600). Actinomycete recipients used for mating with *E. coli* could be in spore suspension with or without pre-germination process, or mycelia in liquid culture. Several agar media including MS and ISP 4 were generally used for

* Correspondence: Arinthip Thamchaipenet, Actinobacteria Research Unit, Department of Genetics, Faculty of Science, Kasetsart University, Chatuchak, Bangkok 10900, Thailand. Tel.: +66 2 562 5444 ext. 4208; fax.: +66 2 579 5528; e-mail: arinthip.t@ku.ac.th.

streptomycete strains while other more specific media were proposed for rare actinomycetes. Concentration of $MgCl_2$, ratio of donor and recipient and other parameters are significantly important to increase conjugation efficiency. The intergeneric conjugation approach has become a powerful method to genetically manipulate actinomycete species with high efficiency. It turns out to be a common tool to characterise functions of genes/gene clusters in these pharmaceutically important bacteria.

Introduction

Actinomycetes are high G+C content (74 mol%) filamentous Gram-positive bacteria found abundantly in soil, sand and sediments. They have a complex life cycle with an extended substrate mycelia on which aerial hyphae is mostly formed with exospores. Genus *Streptomyces* in particular is a key producer of valuable secondary metabolites including anti-bacterials, anti-fungals, anti-cancers, immunosuppressants, herbicides and insecticides used nowadays in medicine, pharmaceuticals and agriculture [1, 2]. Actinomycetes, therefore, are the most attractive microorganisms to study genes involved in biosynthesis of such important secondary metabolites. Recently, genome analysis of *Streptomyces* chromosomes revealed that they carry vast numbers of uncharacterised genes and biosynthetic gene clusters for undiscovered secondary metabolites [3]. Therefore, the establishment of efficient genetic manipulation systems to introduce homologous/heterologous DNAs into actinomycete species has become a critical requirement for achieving functional analysis of the genes/gene clusters of interest.

Genetic manipulation in actinomycetes was developed intensively for the genus *Streptomyces*. Recombinant plasmids construction for gene expression and disruption/replacement can be generally transferred into streptomycete hosts using polyethylene glycol (PEG) mediated protoplast [4]. The procedure is based on the protocol defined for *S. coelicolor* and *S. lividans* [5]. However, this method is time-consuming procedure as it requires high productivity of protoplast formation and regeneration to achieve good transformation efficiency. With a number of different streptomycete species, the method has to be optimised e.g. mycelial age, concentration of the lysozyme to give good protoplast formation and conditions for protoplast regeneration. PEG mediated protoplast transformation does not easily applicable to every species and frequently results in low efficiency particularly with rare actinomycete strains. In addition, due to the strong host restriction modification system of *Streptomyces* [6], it is essential to find a proficient method to transfer DNA with high efficiency. Although high efficient transformation by electroporation was proposed to avoid the problematic protoplast formation and regeneration [7, 8], it requires luxurious equipment and has limited application not to every species of actinomycetes.

Nowadays, the method generally employed for genetic manipulation in *Streptomyces* and other genera is plasmid-mediated intergeneric conjugation with *E. coli* as a donor [9]. This method is proved to be simple and less time consuming and likely to be applied to various actinomycete species. However, the efficiency of this system varies significantly from one species to another, and it is necessary to improve experimental procedures for a new species.

Bacterial Conjugation

Conjugation is a common phenomenon in both Gram-negative and Gram-positive bacteria. It requires close physical contact between donor and recipient cells and is usually mediated by plasmid-encoded proteins which provide transfer (*tra*) functions [10, 11, 12].

The conjugation by self-transmissible plasmid is related in a mechanism sense to that of F plasmid in *E. coli*. Donor creates F pilus to make cell-to-cell contact to recipient. The mechanism is initiated by creating a single-stranded nick at the origin of transfer (*oriT*). The 5'-end of the nicked DNA is transported linearly through the F pilus into the recipient cell where it is replicated; while, the unnicked DNA is replicated in the donor cells [13].

Naturally, some self-transmissible plasmids could transfer among the same bacterial species but not to other different species. However, some plasmids were able both to transfer and to replicate in effectively different species. Some conjugation phenomena with non-pilus mechanisms were also found in Gram-positive bacteria including *Streptomyces* [10].

Conjugation within *Streptomyces* Species

The conjugation process with non-pilus mechanism naturally appears in *Streptomyces*. It is a very distinct process from other bacteria in its molecular mechanism and its visible phenotypic appearance. It occurs through intermycelial transfer from the hyphal tips of a plasmid carrying donor to recipient [14, 15]. It seems that such single-stranded DNA transfer that appears in all conjugation systems in unicellular bacteria does not involve in *Streptomyces* conjugation.

Conjugative plasmid was transferred as a double-stranded DNA molecule [15, 16] by *tra* gene products, e.g. *traB* for pSN22 [17] and *traSA* for pSAM2 [16] during mating [18]. Then the plasmid is intramycelial spread through a septum to the next cell of the recipient mycelium. At this point, the growth is retarded in order to obtain sufficient copy number of the plasmids for the next spreading to another compartment of the mycelium [14, 19]. This delay causes a pock formation, a visible retardation zone of plasmid-acquiring recipient cells grown in a confluent lawn of plasmid-free recipients [20, 21].

Intergeneric Conjugation between *E. coli* and *Streptomyces*

Conjugal transfer from *E. coli* to Gram-positive bacteria with F plasmid-like system was demonstrated by Trieu-Cuot et al. [22]. In 1989, intergeneric conjugation from *E. coli* to *Streptomyces* was firstly described [9] and later has been developed and applied to several genera of *Actinomycetales*. This method has proved to be easy, convenient and less time-consuming compared to the original PEG-mediated protoplast transformation method.

The system of intergeneric conjugation between these two taxonomically different microorganisms is absolutely required an *oriT* and *tra* genes for transfer functions in the donor *E. coli* host. It is hypothetically explained that a plasmid was transferred from *E. coli* to *Streptomyces* by single-stranded DNA forms generated by *oriT*-endonuclease function of

donor [10]. This transfer method then could overcome problems of the restriction barrier to foreign DNA in *Streptomyces* hosts by the transfer of single-stranded DNA [23].

E. coli Donor and Mobilisable Plasmids

E. coli donor strains generally used for intergeneric conjugation are the methylation proficient strain, S17-1 harbouring RP4 *tra* genes on its chromosome [9], the methylation-deficient strain, ET12567 (*dam-13::Tn9 dcm-6 hsdM*) [24] containing *tra* genes in a self-transmissible plasmid pUB307 [23] and ET12567 carrying a non-transmissible plasmid pUZ8002 [25]. pUZ8002, a helper plasmid RK2 derivative, is not efficiently transferred itself because of a mutation in its own *oriT* but can supply transfer functions to mobilise *oriT*-carrying non-conjugative replicon plasmids [26]. By using methylation deficient donor such *E. coli* ET12567, the restriction barriers of foreign DNA could be reduced and efficiency of DNA introduction could be highly increased in *Streptomyces* host [6, 23].

The plasmids used for intergeneric conjugal transfer are mobilisable plasmids that required *tra* genes function of F plasmid located on either host chromosome (e.g. *E. coli* S17-1) or helper plasmid (e.g. pUZ8002). There is no evidence yet whether a pilus mechanism is required for mating pair formation between *E. coli* and *Streptomyces*.

Non-replicative plasmids are used on propose of integration of genetic materials into *Streptomyces* chromosomes. These plasmids carry the origin of replication for *E. coli* and the *oriT* region of the broad host range conjugative plasmid RK2 [9, 10]. If the plasmids required for gene expression propose, integrase gene (*int*) and attachment site (*attP*) of *Streptomyces* phage ϕ C31 were inserted in order to drive site-specific recombination at *attB* site of *Streptomyces* chromosome [27, 28]. These plasmids, for example, are pIJ8600 (*oripUC*, *oriT*, ϕ C31 *attP-int*, *Apr^r*, *Thio^r*, *tipAp*) [29], pSET152 (*oripUC*, *oriT*, ϕ C31 *attP-int*, *Apr^r*) [28], pIJ6902 (*oripUC*, *oriT*, ϕ C31 *attP-int*, *Apr^r*, *Thio^r*, *tipAp*) [30] and pIJ10257 (*oripUC*, *oriT*, ϕ BT1 *attB-int*, *Hyg^r*, *ermEp**) [31]. If the plasmids are to use for integration by homologous recombination, no *int* and *attP* of ϕ C31 was required e.g. pIJ8671 (*oripUC*, *oriT*, *Apr^r*, *Thio^r*, *tipAp*) [29] and pSET151 (*oripUC*, *oriT*, *Thio^r*) [32].

The non-replicative recombinant plasmid is forced to integrate at the target homologous gene to achieve disruption. Some shuttle plasmids were designed for replication purpose in *Streptomyces* then the origin of replication for both *E. coli* and *Streptomyces* was inserted such as pJN100 (*oripUC*, *oripIJ101*, *oriT*, *Apr^r*, *snpAp*) [33].

Pre-Treatments for Streptomycete-Actinomycete Recipients

Pre-germination of spores of *Streptomyces* recipients by heat treatment is critical for intergeneric conjugation procedure [5, 23]. The temperature and time used for spore pre-germination treatment effect on the efficiency of conjugation in individual species. In general, pre-germination of *Streptomyces* spores activated by heat treatment at 50°C for 10 min before mating with an *E. coli* donor was recommended [5, 9, 23]. The spore pre-germination of *S. lividans* at this condition increased the efficiency of conjugation by 5-10 times [9]. It was practically used for spores of *S. albus*, *S. anibioticus*, *S. aureofaciens*, *S. bambergiensis*, *S.*