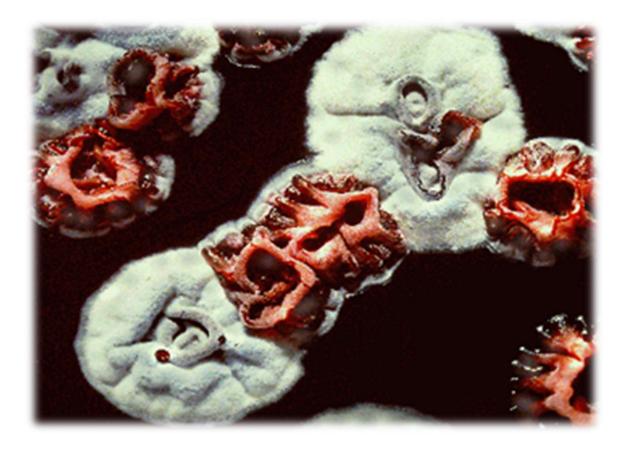
The 4th A3 Foresight Symposium on

Chemical & Synthetic Biology of Natural Products

Dates: July 5-8, 2019

Shanghai Jiao Tong University, Shanghai, China





Aim of this program

To construct an Asian academic research hub for the chemical & synthetic biology of natural products through 3-way international collaboration on the 'chemical & synthetic biology of natural products through *Streptomyces* genome mining (Japan), artificial chromosome engineering (Korea), and synthetic cell factory design (China)'.



http://a3-chemical-biology.jp/en/

Symposium

The 1st A3 Foresight Symposium on "Chemical & Synthetic Biology of Natural Products" at Shanghai Jiao Tong University, Shanghai, China. (August 22-23, 2016)



The 2nd A3 Foresight Symposium on "Chemical and Synthetic Biology of Natural Products" at Grabel Hotel and Jeju ICC, Korea. (May 22-24, 2017)



The 3rd A3 Foresight Symposium on "Chemical and Synthetic Biology of Natural Products" at Hotel North City, Sapporo, Japan. (July 9-12, 2018)



Schedule

July 5 th Fri.	Registration
	Location: Leadingmen Collection Hotel
July 6 th Sat.	
	(07:45 Bus Leadingmen Collection Hotel)
	Loaction: Shuhua Auditorium, Shanghai Jiao Tong University
08:30-08:45	Opening Remarks
	Zixin Deng (Shanghai Jiao Tong University, China)
	Eung-Soo Kim (Inha University, Korea)
	Yasuo Ohnishi (The University of Tokyo, Japan)
08:45-09:00	Group Photo
09:00-10:00	Session 1: Presentation by invited speakers (1)
10:00-10:20	Coffee Break
10:20-11:20	Session 2: Presentation by invited speakers (2)
11:25	Lunch (Bus to Canteen 2, SJTU)
13:00-15:01	Session 3: Group Presentation from Japan (1)
15:01-15:20	Coffee Break
15:20-16:50	Session 4: Group Presentation from Japan (2)
16:50-17:20	Session 5: Group Presentation from China (1)
17:25	Dinner (Bus to Huhua International Hotel)

July 7 th Sun.	(08:00 Bus Leadingmen Collection Hotel)
	Location: Shuhua Auditorium, Shanghai Jiao Tong University

- 08:30-10:15 Session 6: Group Presentation from Korea (1)
- 10:15-10:30 Coffee Break
- 10:30-10:55 Session 7: Group Presentation from Korea (2)
- 10:55-11:30 Session 8: Group Presentation from China (2)
- 11:30 **Lunch** (Bus to Canteen 2, SJTU)
- 13:00-15:00 Session 9: Group Presentation from China (3)
- 15:00-15:20 Coffee Break
- 15:20-17:38 Session 10: Group Presentation from China (4)
- 17:38-18:10 Panel Discussion, Award and Closing Ceremony
- 18:15 **Dinner** (Bus to Academic Exchange Center Hotel, SJTU)

July 8th Mon. Departure

- 07:45 Two Bus (Leadingmen Collection Hotel)
- 08:00 Bus 1 Hongqiao International Airport, Terminal 1, Terminl 2 (40 min) Bus 2 Pudong International Airport, Terminal 1, Terminl 2 (60 min)

Program

July 6th Sat.

Opening Remarks (Chair: Prof. Linquan Bai)

08:30-08:45

Zixin Deng (Shanghai Jiao Tong University, China) Eung-Soo Kim (Inha University, Korea) Yasuo Ohnishi (The University of Tokyo, Japan)

08:45-09:00 Group Photo

Session 1: Presentation by Invited Speakers (1) (Chair: Prof. Shuangjun Lin)

09:00-09:30

Streptomyces metabolites in microbial interaction

Prof. Kenji Ueda

Life Science Research Center, College of Bioresource Sciences, Nihon University

09:30-10:00

Characterization and application of agar hydrolyzing system of *Streptomyces* coelicolor A3 (2) <u>Prof. Soon-Kwang Hong</u> Myongji University

10:00-10:20 Coffee Break

Session 2: Presentation by Invited Speakers (2) (Chair: Prof. Shuangjun Lin) 10:20-10:50

Structural insights into pericyclase catalysis

<u>Prof. Jiahai Zhou</u> Shanghai Institute of Organic Chemistry, Chinese Academy of Sciences

10:50-11:20

Compartmentalized biosynthesis of mycophenolic acid

Prof. Shengying Li Shandong University

11:25 Lunch (Bus to Canteen 2, SJTU)

Session 3: Group Presentation from Japan (1) (Chair: Prof. Yasuo Ohnishi)

13:00-13:05

Introduction

Yasuo Ohnishi^{1,2}, Yohei Katsuyama^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo

²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

13:05-13:25

Studies on the unique life cycle of the rare actinomycete *Actinoplanes missouriensis* Takeaki Tezuka^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo

²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

13:25-13:35

Genome analysis of the industrial fungus Aspergillus sojae Atsushi Sato^{1,2}, Yasuji Koyama² ¹Graduate School of Agricultural and Life Sciences, The University of Tokyo ²Noda Institute for Science Research

13:35-13:45

Genome mining for novel nitrating cytochrome P450s provides insights into lajollamycin biosynthesis

Hiroya Tomita¹, Yohei Katsuyama^{1,2}, Yasuo Ohnishi^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo ²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

13:45-13:55

X-ray structural analysis of the KS-CLF heterodimer and the ACP-KS-CLF ternary complex of a highly reducing type II PKS

Danyao Du^{1,†}, Yohei Kastuyama^{1,2}, Shinya Fushinobu^{1,2}, Aochiu Chen³, Tony D. Davis³, Michael D. Burkart³, Yasuo Ohnishi^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo

²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

³Department of Chemistry and Biochemistry, University of California, San Diego, USA [†]Novozymes Japan Ltd.

13:55-14:00

Biochemical and structural analysis of geranyl pyrophosphate C6methyltransferase BezA

Hayama Tsutsumi¹, Yohei Katsuyama^{1,2}, Yasuo Ohnishi^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo

²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

14:00-14:05

Altering the regioselectivity of the toluic acid-recognizing biosensor XylS through directed evolution

<u>Yuki Ogawa</u>¹, Yohei Katsuyama^{1,2}, Kento Ueno¹, Yasuo Ohnishi^{1,2} ¹Graduate School of Agricultural and Life Sciences, The University of Tokyo ²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

14:05-14:08

Biosynthesis of the aromatic polyketide yoropyrazone in *Streptomyces* sp. IFM11307

<u>Kasumi Fujita</u>¹, Yohei Katsuyama^{1,2}, Kazufumi Toume³, Masami Ishibashi⁴, Yasuo Ohnishi^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo

²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

³Institute of Natural Medicine, University of Toyama

⁴Graduate School of Pharmaceutical Sciences, Chiba University

14:08-14:11

Identification of *Kutzneria albida* secondary metabolites synthesized using the secondary metabolism-specific nitrous acid biosynthetic pathway <u>Akito Yamada</u>¹, Yohei Katsuyama^{1,2}, Yasuo Ohnishi^{1,2} ¹Graduate School of Agricultural and Life Sciences, The University of Tokyo ²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

14:11-14:14

Analysis of cross-talk of ACPs among type II FAS and PKSs

Masanobu Horiuchi¹, Danyao Du¹, Yohei Katsuyama^{1,2}, Yasuo Ohnishi^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo ²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

14:14-14:17

Production of methylgeranylated polyphenols using BezA and NphB

<u>Naoki Urano</u>¹, Hayama Tsutsumi¹, Yohei Katsuyama^{1,2}, Yasuo Ohnishi^{1,2} ¹Graduate School of Agricultural and Life Sciences, The University of Tokyo ²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

14:17-14:20

Biosynthesis of the nonribosomal peptide, cirratiomycin from *Streptomyces* cirattus

Jiafeng Li¹, Yohei Katsuyama^{1,2}, Kazuo Shin-ya³, Yasuo Ohnishi^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo

²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

³National Institute of Advanced Industrial Science and Technology (AIST)

14:20-14:25

Introduction

Hiroyasu Onaka^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo ²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

14:25-14:45

Intergeneric cell-cell contact dependent conversion of bioactive natural products in combined-culture between *Streptomyces* sp. HEK616 and *Tsukamurella*

pulmonis

Shumpei Asamizu^{1,2}, Hiroyasu Onaka^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo ²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

14:45-14:50

Interrogating the relationship between filamentous actinomycetes and mycolic acid-containing bacteria in ecosystem

Manami Kato¹, Shumpei Asamizu^{1,2}, Hiroyasu Onaka^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo

²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

14:50-14:53

Toward understanding the mechanism of intergeneric cell-cell contact induced activation of secondary metabolism using *Streptomyces coelicolor* and *Tsukamurella pulmonis* as model system

Masashi Kida¹, Shumpei Asamizu^{1,2}, Hiroyasu Onaka^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo

²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

14:53-14:58

Tokyo

Enzymatic condensation of RiPP and very long chain fatty acid to synthesize unprecedented lipopeptide

Ryosuke Kozakai¹, Shumpei Asamizu^{1,2}, Hiroyasu Onaka^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo ²Collaborative Research Institute for Innovative Microbiology, The University of

12

14:58-15:01

Toward understanding the mechanism of goadsporin induced activation of secondary metabolism

Sachiko Kawano¹, Shumpei Asamizu^{1,2}, Hiroyasu Onaka^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo

²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

15:01-15:20 Coffee Break

Session 4: Group Presentation from Japan (2) (Chair: Prof. Yasuo Ohnishi)

15:20-15:25

Introduction

Tomohisa Kuzuyama^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo

²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

15:25-15:35

Biogasoline production by engineered polyketide synthases

Satoshi Yuzawa¹ and Tomohisa Kuzuyama^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo

²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

15:35-15:38

Biosynthetic studies on unusual methylbenzene-containing polyenes Jie Zhang¹ and Tomohisa Kuzuyama^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo ²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

15:38-15:41

Biosynthetic studies on Streptomyces meroterpenoid

<u>Tomohiro Noguchi¹</u> and Tomohisa Kuzuyama^{1,2}

¹Graduate School of Agricultural and Life Sciences, The University of Tokyo ²Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

15:41-15:46

Introduction:

Biosynthetic studies of macrocyclic polyketide antibiotics in actinomycetes

Fumitaka Kudo

Tokyo Institute of Technology

15:46-15:51

Biosynthetic studies on 3-aminononanoic acid starter unit of cremimycin

Daisuke Kawasaki, Taichi Chisuga, Akimasa Miyanaga, Fumitaka Kudo, Tadashi Eguchi

Tokyo Institute of Technology

15:51-15:56

Engineering of macrocyclic polyketides biosynthesis in *Streptomyces* graminofaciens

Kosuke Kishikawa, Akimasa Miyanaga, Fumitaka Kudo, Tadashi Eguchi

Tokyo Institute of Technology

15:56-16:01

Engineering of hitachimycin biosynthesis

<u>Sotaro Takahashi</u>, Yuichiro Nakazawa, Koichi Kawamura, Akimasa Miyanaga, Fumitaka Kudo, Tadashi Eguchi Tokyo Institute of Technology

16:01-16:06

Introduction:

Polycationic peptides - discovery, biosynthesis, and application

Yoshimitsu Hamano

Graduate School of Bioscience and Biotechnology, Fukui Prefectural University

16:06-16:11

Biosynthesis of peptide natural products

Chitose Maruyama

Graduate School of Bioscience and Biotechnology, Fukui Prefectural University

16:11-16:14

Identification and characterization of bacterial 1-aminocyclopropane-1carboxylic acid synthase

<u>Yukiko Chinone</u>¹, Yoshimitsu Hamano¹, Junko Hashimoto², Ikuko Kozone², Kazuo Shin-ya³, Chitose Maruyama¹

¹Graduate School of Bioscience and Biotechnology, Fukui Prefectural University

²Japan Biological Informatics Consortium (JBIC)

³National Institute of Advanced Industrial Science and Technology (AIST)

16:14-16:17

Biosynthesis of a streptothricin analogue possessing O-acylpeptide side chain

Seren Nagashima¹, Chitose Maruyama¹, Yukiko Chinone¹, Yasushi Ogasawara², Junko

Hashimoto³, Kazuo Shin-ya⁴, Tohru Dairi², Yoshimitsu Hamano¹

¹Department of Bioscience, Fukui Prefectural University

²Department of Engineering, Hokkaido University
³Japan Biological Informatics Consortium (JBIC)
⁴National Institute of Advanced Industrial Science and Technology (AIST)

16:17-16:20

Intracellular delivery of macromolecules modified with ε-poly-L-lysine <u>Yamato Takeuchi</u>¹, Chitose Maruyama¹, Yasuo Kato², Yoshimitsu Hamano¹ ¹Graduate School of Bioscience and Biotechnology, Fukui Prefectural University ²Graduate School of Engineering, Toyama Prefectural University

16:20-16:23

Chemical modification of the bioactive small molecules with ε-poly-L-lysine for improving cell membrane permeability and water solubility

<u>Fumika Matsumura</u>¹, Chitose Maruyama¹, Yamato Takeuchi¹, Yasuo Kato², Yoshimitsu Hamano¹

¹Graduate School of Bioscience and Biotechnology, Fukui Prefectural University ²Graduate School of Engineering, Toyama Prefectural University

16:23-16:26

Chemical modification of the bioactive small molecules with oligo(L-Lys) for improving cell membrane permeability and water solubility

<u>Kohei Kaneda</u>¹, Chitose Maruyama¹, Yamato Takeuchi¹, Yasuo Kato², Yoshimitsu Hamano¹

¹Graduate School of Bioscience and Biotechnology, Fukui Prefectural University ²Graduate School of Engineering, Toyama Prefectural University

16:26-16:31

Introduction:

Overview of biosynthetic machinery, structural redesign, regulatory network, and genome mining in actinomycetes

Kenji Arakawa

¹Department of Molecular Biotechnology, Graduate School of Advanced Sciences of Matter, Hiroshima University

²Unit of Biotechnology, Graduate School of Integrated Sciences for Life, Hiroshima University

16:31-16:34

Metabolite analysis of biosynthetic gene mutants of the signaling molecule SRB in *Streptomyces rochei*

Aiko Teshima¹, and Kenji Arakawa^{1,2}

¹Department of Molecular Biotechnology, Graduate School of Advanced Sciences of Matter, Hiroshima University

²Unit of Biotechnology, Graduate School of Integrated Sciences for Life, Hiroshima University

16:34-16:37

A practical genome mining using butenolide-type signaling molecules in *Streptomyces* species

Yuto Eguchi¹, Miho Sumiyoshi², Aiko Teshima², and Kenji Arakawa^{1,2}

¹Unit of Biotechnology, Graduate School of Integrated Sciences for Life, Hiroshima University

²Department of Molecular Biotechnology, Graduate School of Advanced Sciences of Matter, Hiroshima University

16:37-16:40

Analysis of metabolic production through rational modification of signalingmolecule regulatory genes coded on the *Streptomyces rochei* chromosome

Yuya Misaki¹, Miyuki Iwakuni², Toshihiro Suzuki², and Kenji Arakawa^{1,2}

¹Unit of Biotechnology, Graduate School of Integrated Sciences for Life, Hiroshima University ²Department of Molecular Biotechnology, Graduate School of Advanced Sciences of Matter, Hiroshima University

16:40-16:43

Comprehensive metabolome analysis to explore the biologically active metabolites based on genome sequences

<u>Rikito Nishiura</u>¹, Amirudin Akhmad Fauzi¹, Yosi Nindita¹, Junko Hashimoto², Kazuo Shin-ya³, Haruo Ikeda⁴, and Kenji Arakawa^{1,5}

¹Department of Molecular Biotechnology, Graduate School of Advanced Sciences of Matter, Hiroshima University

²Japan Biological Informatics Consortium (JBIC)

³National Institute of Advanced Industrial Science and Technology (AIST)

⁴Kitasato Institute for Life Sciences, Kitasato University

⁵Unit of Biotechnology, Graduate School of Integrated Sciences for Life, Hiroshima University

16:43-16:47

Biosynthetic investigation of unique carbocyclic polyketides

<u>Hazuki Ogata</u>¹, Yuji Yukiyoshi¹, Yuki Inahashi^{2,3}, Yoko Takahashi², Satoshi Ōmura², Tomohisa Kuzuyama^{4,5}, Chin Piow Wong⁶, Hiroyuki Morita⁶, Takuji Nakashima^{2,3}, and Kenji Arakawa^{1,7}

¹Department of Molecular Biotechnology, Graduate School of Advanced Sciences of Matter, Hiroshima University

²Kitasato Institute for Life Sciences, Kitasato University

³Graduate School of Infection Control Sciences, Kitasato University

⁴Graduate School of Agricultural and Life Sciences, The University of Tokyo

⁵Collaborative Research Institute for Innovative Microbiology, The University of Tokyo

⁶Institute of Natural Medicine, University of Toyama

⁷Unit of Biotechnology, Graduate School of Integrated Sciences for Life, Hiroshima

University

16:47-16:50

Analysis of the biosynthetic machinery of azoxyalkene compounds

<u>Kota Fujita</u>¹, Ayaka Tatsukawa², Takuya Kishimoto², Atsushi Fukumoto³, Yojiro Anzai³, and Kenji Arakawa^{1,2}

¹Unit of Biotechnology, Graduate School of Integrated Sciences for Life, Hiroshima University

²Department of Molecular Biotechnology, Graduate School of Advanced Sciences of Matter, Hiroshima University

³Faculty of Pharmaceutical Science, Toho University

Session 5: Group Presentation from China (1) (Chair: Prof. Linquan Bai)

16:50 -16:55

Introduction

Shuangjun Lin, Tingting Huang

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

16:55-17:00

Production of hydroxypyruvate in Escherichia coli

Xiaolai Lei, Qi Liu, Jingjie Jiang, Tingting Huang, Shuangjun Lin

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

17:00-17:05

Biosynthesis of coprisidin

<u>Feiyan Lin</u>, Yingyi Duan, Tingting Huang, Manchen Tang, Shuangjun Lin State Key Laboratory of Microbial Metabolism, School of Life Sciences and

Biotechnology, Shanghai Jiao Tong University

17:05-17:10

Biosynthesis of β -carboline alkaloid intermediate in streptonigrin pathway

<u>Xiaozheng Wang</u>, Wenli Guo, Dekun Kong, Tingting Huang, Shuangjun Lin State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

17:10-17:15

Genome mining for the flavinprotein as coenzyme for StnD catalyzed quinone hydroxylation

<u>Xinyue Xie</u>, Xiaozheng Wang, Tingting Huang, Shuangjun Lin State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

17:15-17:20

C-H hydroxylation of aromatics catalyzed by dioxygenase

<u>Chong Yin</u>, Wangli Peng, Shuangjun Lin, Rubin Liang State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

17:25 Dinner (Bus to Huhua International Hotel)

July 7th Sun

Session 6: Group Presentation from Korea (1) (Chair: Prof. Eung-Soo Kim) 08:30-08:35 Introduction: Overview of isolation and expression of large-sized natural product biosynthetic gene cluster Prof. Eung-Soo Kim Inha University

08:35-08:47

Development of novel NPP antifungal polyene macrolides in a rare *Actinomycetes* <u>Dr. Sisun Choi</u>, Heung-Soon Park, Chi-Young Han, Hye-Jin Kim and Eung-Soo Kim* Inha University

08:47-08:59

Cell factory design and culture process optimization for dehydroshikimate biosynthesis in *Escherichia coli*

<u>Dr. Han-Na Lee</u>, Eunhwi Park, Seung-Yeul Seo, Sisun Choi, Sang joung Lee, and Eung-Soo Kim*

Inha University

08:59-09:11

Thioesterase domain modification for production of cyclized tautomycetin

Dr. Hee-Ju Nah, Seunghee Choi, Sisun Choi, Eung-Soo Kim*

Inha University

09:11-09:16

Isolation of novel polyene compounds via antifungal bioassay-based screening in

21

Actinomycetes species

<u>Heung-soon Park</u>, Si-Sun Choi, and Eung-Soo Kim* Inha University

09:16-09:21

High production of tautomycetin enhancing expression of regulatory genes in heterologous host <u>Seunghee Choi</u>, Hee-Ju Nah, Sisun Choi, and Eung-Soo Kim* Inha University

09:21-09:26

Construction of *Corynebacterium* cell factory for shikimic acid overproduction by metabolic pathway engineering

Eunhwi Park, Han-Na Lee, Seung-Yeul Seo, Sang joung Lee, and Eung-Soo Kim* Inha University

09:26-09:31

Introduction:

Biomolecule reconstruction by glycosylation platform

Prof. Jae Kyung Sohng

Sun Moon University

09:31-09:43

Mutagenetic studies for product specificity of YjiC towards α-mangostin monoglucoside

Dr. Tae-Su Kim, Hue Thi Nguyen, Kye Woon Cho, Jae Kyung Sohng*

Sun Moon University

09:43-09:55

Isolation and characterization of bioactive molecules from Nocardia

Dr. Dipesh Dhakal, Jae Kyung Sohng*

09:55-10:00

Sun Moon University

Biosynthesis and characterization of unusual furan type compound from marine *Streptomyces*

<u>Hue Thi Nguyen</u>, Chung Thanh Nguyen, Van Thuy Thi Pham, Jae-Kyung Sohng* Sun Moon University

10:00-10:05

Discovery of a new natural product in Streptomyces peucetius

<u>Van Thuy Thi Pham</u>, Chung Thanh Nguyen, Hue Thi Nguyen, Jae-Kyung Sohng* Sun Moon University

10:05-10:10

Activating cryptic biosynthetic pathway in *Streptomyces* sp. KTCT0041BP <u>Chung Thanh Nguyen</u>, Van Thuy Thi Pham, Hue Thi Nguyen, Jae-Kyung Sohng* Sun Moon University

10:10-10:15

Domain mutagenesis of herboxidiene gene cluster to generate novel analogues <u>Rubin Thapa Magar</u>, Jae-Kyung Sohng* Sun Moon University

10:15-10:30 Coffee Break

Session 7: Group Presentation from Korea (2) (Chair: Prof. Eung-Soo Kim) 10:30-10:35

Iron competition triggers antibiotic biosynthesis in Streptomyces coelicolor during

co-culture with *Myxococcus xanthus*

<u>Namil Lee</u>, Woori Kim, Jinkyoo Chung, Yongjae Lee, Suhyung Cho, Kyoung-Soon Jang, Sun Chang Kim, Bernhard Palsson, and Byung-Kwan Cho* Korea Advanced Institute of Science and Technology (KAIST)

10:35-10:40

Primary transcriptome and translatome analysis determines transcriptional and translational regulatory elements encoded in the *Streptomyces clavuligerus* genome <u>Soonkyu Hwang</u>, Namil Lee, Yujin jeong, Yongjae Lee, Woori Kim, Suhyung Cho, Bernald O Palsson, Byung-Kwan Cho* Korea Advanced Institute of Science and Technology (KAIST)

10:40-10:45

The transcription unit architecture of *Streptomyces lividans* TK24

<u>Yongjae Lee</u>, Namil Lee, Yujin Jeong, Soonkyu Hwang, Woori Kim, Suhyung Cho, Bernhard O. Palsson, and Byung-Kwan Cho* Korea Advanced Institute of Science and Technology (KAIST)

10:45-10:50

Biochemical identification of a neoagarooligosaccharides hydrolase in Streptomyces coelicolor A3 (2) Ju Won Seo, Soon-Kwang Hong* Myongji University

10:50-10:55

Regulation of agararses production in *Streptomyces coelicolor* A3 (2) <u>Maral Tsevelkhorloo</u>, Soon-Kwang Hong* Myongji University

Session 8: Group Presentation from China (2) (Chair: Prof. Linquan Bai)

10:55-11:15

Synthetic biology approaches for multi-copy chromosomal integration of natural product biosynthetic gene clusters in *Streptomyces*

<u>Prof. Yinhua Lu</u> Shanghai Normal University

11:15-11:30

A CRISPR/Cas12a-derived biosensing platform for the highly sensitive detection of diverse small molecules <u>Associate Prof. Gaoyi Tan</u> East China University of Science and Technology

11:30 Lunch (Bus to Canteen 2, SJTU)

Session 9: Group Presentation from China (3) (Chair: Prof. Linquan Bai)

13:00-13:20

Introduction:

Global regulation of crotonylation on metabolic development of *Streptomyces* through carbon catabolite repression

Prof. Xu-Ming Mao, Prof. Yongquan Li

Zhejiang University

13:20-13:25

Artificial construction of glycopeptide antibiotic A82846B biosynthetic pathway and mechanism study for high yield <u>Hui Qian</u>, Yongquan Li Zhejiang University

13:25-13:30

DNA methylation regulates the secondary metabolism of *Streptomyces roseosporus* <u>Jiao-Le Fang</u>, Yongquan Li Zhejiang University

13:30-13:50

Bio-formation and bio-control of cyclopropane ring in CC-1065 and yatakemycin biosynthesis <u>Prof. Gong-Li Tang</u> Shanghai Institute of Organic Chemistry, CAS

13:50-14:05

Anthraquinone-γ-pyrone ring formation in complex aromatic polyketide biosynthesis Associate Prof. Xian-Feng Hou

Shanghai Institute of Organic Chemistry, CAS

14:05-14:10

A self-resistance strategy existing in naphthyridinomycin biosynthesis

Wan-Hong Wen, Yue Zhang, and Gong-Li Tang*

Shanghai Institute of Organic ChemistryAS

14:10-14:15

Investigating divergent tailoring pathways of two series of aromatic polyketides generated through mining

<u>Qiu-Yue Nie</u>, Zhen-Yu Ji, Xian-Feng Hou, and Gong-Li Tang* Shanghai Institute of Organic Chemistry, CAS

14:15-14:35

Introduction:

Deciphering the late steps of rifamycin biosynthesis

<u>Prof. Wei Zhang</u>, Prof. Shengying Li Shandong University, China

14:35-14:40

Complete elucidation of the late steps of bafilomycin biosynthesis in *Streptomyces lohii* <u>Zhong Li</u>, Lei Du, Wei Zhang, XingWang Zhang, David H. Sherman and Shengying

Li*

Qingdao Institute of Bioenergy and Bioprocess Technology, CAS

14:40-14:45

Introduction

Prof. Yuemao Sheng

Shandong University

14:45-14:50

A single-component flavoenzyme catalyzed regioselective halogenation of pyrone in the biosynthesis of venemycins

<u>Rentai Song</u>, Haixia Shi, Jing Zhu, Haoxin Wang*, Yuemao Shen* Shandong University

14:50-14:55

Construction of a hybrid gene cluster to reveal coupled ring formationhydroxylation in the biosynthesis of HSAF and analogues from *Lysobacter enzymogenes*

Xue Li, Haoxin Wang, Yaoyao Li* and Liangcheng Du* Shandong University

14:55-15:00

Base editing in Streptomyces with Cas9-deaminase fusions

<u>Zhiyu Zhong</u>, Junhong Guo, Liang Deng, Li Chen, Jian Wang, Sicong Li, Wei Xu, Zixin Deng, Yuhui Sun* Wuhan University

15:00-15:20 Coffee Break

Session 10: Group Presentation from China (4) (Chair: Prof. Linquan Bai)

15:20-15:25

Introduction

Yan Feng

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

15:25 -15:30

Engineering promiscuous thioesterase TesA from *E. coli* for medium-chain fatty acids biosynthesis

Xi Deng, Liuqing Chen, Guangyu Yang*, Yan Feng*

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

15:30 - 15:35

Algorithm-based coevolution network identification reveals key functional sectors of the α/β hydrolase subfamilies

Zhiyun Wu, Lishi Xu, Hao Liu, Haifeng Chen*, Yan Feng*

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

15:35 -15:40

Introduction

Delin You

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

15:40-15:45

Tailoring enzymes involved in anisomycin biosynthesis

Qing Wang and Delin You*

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

15:45-15:50

Construction of the engineering strain for production of drug precursors demeclocycline and demecycline

Weinan Yang and Delin You*

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

15:50-16:05

Introduction:

Theoretical studies on the mechanisms of thioesterase-catalyzed polyketide chain release

Associate Prof. Ting Shi, Prof. Yilei Zhao

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

16:05-16:10

The Molecular recognition basis for phosphorothioated DNA diverge from normal DNA

Jiayi Li, Ting Shi and Yilei Zhao*

State Key Laboratory of Microbial Metabolism, School of Life Scienes and Biotechnology, Shanghai Jiao Tong University

16:10-16:15

Computational studies for mechanism of enantioselectivity in a zinc-dependent dehydrogenase

Shenggan Luo, Ting Shi and Yilei Zhao*

State Key Laboratory of Microbial Metabolism, School of Life Scienes and Biotechnology, Shanghai Jiao Tong University

16:15-16:20

An explanation for tautomycetin thioesterase preferring hydrolysis to macrocyclization

Lei Liu, Ting Shi and Yilei Zhao*

State Key Laboratory of Microbial Metabolism, School of Life Scienes and Biotechnology, Shanghai Jiao Tong University

16:20-16:25

Insights into substrate specificity and catalytic mechanism of nystatin/ amphotercin B thioesterase

Rufang Wang, Ting Shi and Yilei Zhao*

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

16:25-16:45

Introduction:

Structural and functional dissection of modular polyketide synthases

Prof. Jianting Zheng

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

16:45-16:50

Crystal structure of acetyl-CoA carboxylase β-subunit of *Streptomyces antibioticus* <u>Imtiaz Ali</u> and Jianting Zheng*

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

16:50-16:55

Crystal structure of FAD dependent halogenase ChlB4 and acyl-transferase ChlB3 in the biosynthesis of chlorothricin

Asad Ullah Saeed and Jianting Zheng*

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

16:55-17:00

Theoretical studies on the catalytic mechanism and substrate specificity of acyltransferase domains from salinomycin polyketide synthase

Huining Ji and Jianting Zheng*

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

17:00-17:05

Structural basis for transcription initiation by HrdB regulon in *Streptomyces* coelicolor

Guiyang Liu and Jianting Zheng*

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

17:05-17:10

Introduction

Prof. Linquan Bai

State Key Laboratory of Microbial Metabolism, School of Life Scienes and Biotechnology, Shanghai Jiao Tong University

17:10-17:20

Self-resistance of the ansamitocin producer *Ac. pretiosum* ATCC31280 Xinran Wang and Linquan Bai*

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

17:20-17:25

Structure-directed engineering of thioesterase domain toward improved chain release of unnatural polyene antibiotics

<u>Yucong Zhou,</u> Ting Shi, Zhen Qi, Qianjin Kang, Jianting Zheng, Yilei Zhao, Linquan Bai^{*}

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

17:25-17:30

ATP/ADP-dependent carbamoylations in the biosynthesis of ansamycins

<u>Jianhua Wei</u>, Yucong Zhou, Jianting Zheng, Qianjin Kang and Linquan Bai* State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

17:30-17:35

Comparative transcriptome-based mining of genes involved in polyether antibiotics export

Xian Liu, Jin Li, Xiaojie Zhang, Yuanting Wu and Linquan Bai*

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

17:35-17:38

Pathway redesign to produce rearranged acarbose analogs with improved activity in *Actinoplanes* spp.

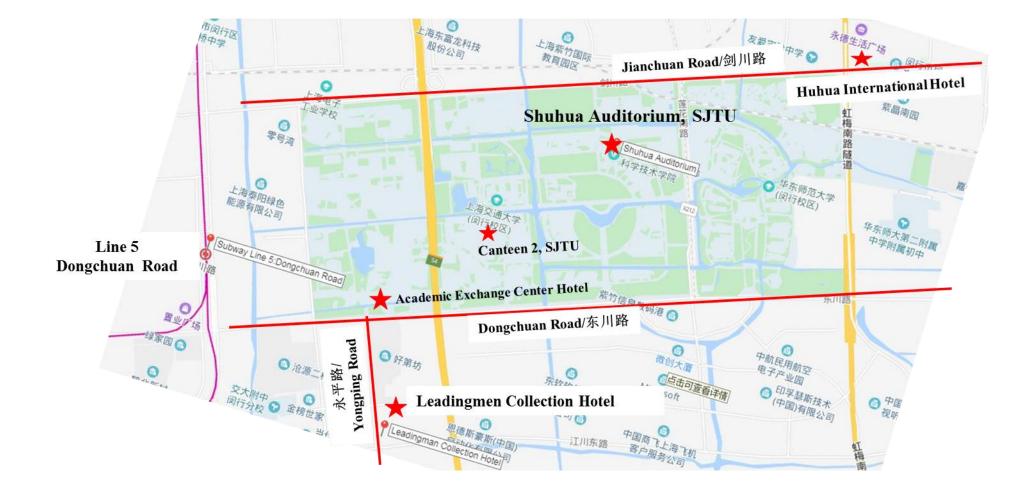
Xin Zhang, Huixin Xie, Qinqin Zhao and Linquan Bai*

State Key Laboratory of Microbial Metabolism, School of Life Sciences and Biotechnology, Shanghai Jiao Tong University

17:38-17:50 Panel Discussion

17:50-18:10 Awards and Closing Ceremony

18:30 Dinner (Bus to Academic Exchange Center Hotel, SJTU)





雷汀曼酒店 (上海交大店) 上海市闵行区永平南路178号,021-33882999 **Leadingmen Collection Hotel (SJTU)** No.178 South Yongping Road, Minhang District



沪华国际大酒店(上海吴泾店)

上海市闵行区剑川路368号(虹梅南路路口)021-64508999

Huhua International Hotel (Wujing)

No.368 Jianchuan Road, Minhang District

