

# Program

**14 December 2025**

Time	Session
12:00 PM – 01:00 PM	<b>Registration</b>
01:00 PM – 01:15 PM	<b>Welcome Speech by Skorn MONGKOLSUK</b>
01:15 PM – 02:00 PM	<b>Keynote: Peter C. DEDON</b> (Massachusetts Institute of Technology, USA) Revisiting the central dogma in the age of epigenomes and epitranscriptomes: Nucleic acid modifications driving health and disease
	<b>Session 1: Post-transcription</b> <b>Chair:</b> CHO Eun-Jung <b>Co-chair:</b> Mayuree FUANGTHONG
02:00 PM – 02:20 PM	<b>WU Cheng-Wen</b> (National Yang Ming Chiao Tung University, Taiwan) S1-01: Intron 2 variation of human <i>POU5F1</i> creates OCT4Borf1 microprotein that promotes tumorigenesis in lung squamous cell carcinoma
02:20 PM – 02:40 PM	<b>Mayuree FUANGTHONG</b> (Chulabhorn Research Institute, Thailand) S1-02: Functional and regulatory roles of tRNA modifications in the stress responses of <i>Pseudomonas aeruginosa</i>
02:40 PM – 03:00 PM	Refreshment
03:00 PM – 03:20 PM	<b>Reshma TANEJA</b> (National University of Singapore, Singapore) S1-03: Interrogating the epigenetic landscape in rhabdomyosarcoma
03:20 PM – 03:40 PM	<b>CHO Eun-Jung</b> (Sungkyunkwan University, Republic of Korea) S1-04: Epigenetic control of pericentromeric chromatin dynamics underlying myogenic differentiation
03:40 PM – 04:00 PM	<b>Narumon THONGDEE</b> (Chulabhorn Research Institute, Thailand) S1-05: Small RNA base pairing within coding sequences transiently slows translation elongation and alters target protein activity
04:00 PM – 04:40 PM	<b>2-min Flash Presentations</b>
04:40 PM – 08:00 PM	Welcome Drink and Dinner

## 15 December 2025

Time	Session
09:30 AM – 10:00 AM	<b>Plenary Lecture: Varodom CHAROENSAWAN</b> (Mahidol University, Thailand) Decoding transcriptional regulation through the lens of a systems biologist: from functional genomics to spatial transcriptomics
	<b>Session 2: Stress response, regulation and signaling</b> <b>Chair:</b> NAKASONE Kaoru <b>Co-chair:</b> Oranart MATANGKASOMBUT
10:00 AM – 10:20 AM	<b>NAKASONE Kaoru</b> (Kindai University, Japan) S2-01: Molecular mechanisms of phytotoxin tropolone in rice-seedling blight, <i>Burkholderia plantarii</i>
10:20 AM – 10:40 AM	Group Photo
10:40 AM – 11:00 AM	Refreshment
11:00 AM – 11:20 AM	<b>KATAOKA Masakazu</b> (Shinshu University, Japan) S2-02: pH homeostasis: exhaustive screening of the genes involved in pH regulation on <i>Escherichia coli</i>
11:20 AM – 11:55 AM	<b>2-min Flash Presentations</b>
11:55 AM – 01:30 PM	Lunch
01:30 PM – 01:50 PM	<b>Oranart MATANGKASOMBUT</b> (Chulalongkorn University, Thailand) S2-03: Exploring the roles of DNA-protein crosslink repair proteases in <i>Candida albicans</i> survival under oxidative stress
01:50 PM – 02:10 PM	<b>Piyajit WATCHARASIT</b> (Chulabhorn Research Institute, Thailand) S2-04: Arsenic impaired insulin and IGF-1 signaling and neural network: the protective roles of insulin and IGF-1, through the PI3K/Akt axis
02:10 PM – 02:30 PM	<b>Kusumawadee UTIS PAN</b> (Thammasat University, Thailand) S2-05: Nitric oxide and stress response in head and neck cancer
02:30 PM – 02:45 PM	<b>Biswanath CHATTERJEE</b> (Taipei Medical University, Taiwan) S2-06: Hypoxia-induced EMT and transcription of EMT-associated genes of cancer cells require DNA demethylation BY DNMT3A
02:45 PM – 04:15 PM	<b>Poster Presentations and Refreshment</b>

## 15 December 2025 (continued)

Time	Session
	<b>Session 3: Molecular insights into health, diseases, and treatment</b> <b>Chair:</b> Kalai MATHEE <b>Co-chair:</b> Nisa PATIKARNTHON
04:15 PM – 04:35 PM	<b>Kalai MATHEE</b> (Euleris LLC, USA) S3-01: Microbiolomics meets AI: revolutionizing the future of microbiology
04:35 PM – 04:55 PM	<b>Tanapat PALAGA</b> (Chulalongkorn University, Thailand) S3-02: The role of Ezh2 of polycomb repressive complex 2 in driving tumor-associated macrophages toward tumor promoting phenotypes in liver cancer model
04:55 PM – 05:15 PM	<b>Nisa PATIKARNTHON</b> (Mahidol University, Thailand) S3-03: Nanoparticle-mediated intracellular delivery of anti-envelope antibodies against dengue: a strategy to interfere with viral gene expression
05:15 PM – 05:35 PM	<b>Worasak KAEWKONG</b> (Naresuan University, Thailand) S3-04: Oncogenic editing of expression: targeting the splicing machinery for cancer suppression
05:35 PM – 05:55 PM	<b>YANG Jae-Hyun</b> (Korea Advanced Institute of Science and Technology, Republic of Korea) S3-05: Epigenetic strategies for reversing mammalian aging
06:00 PM – 08:00 PM	Dinner

## 16 December 2025

Time	Session
	<b>Session 4: Omics approaches to gene expression</b> <b>Chair:</b> WANG Gang <b>Co-chair:</b> Punyawee DULYAYANGKUL
09:00 AM – 09:20 AM	<b>Voraratt CHAMPATTANACHAI</b> (Chulabhorn Research Institute, Thailand) S4-01: Anticancer efficacy of the combination treatment of oxaliplatin and protein O-GlcNAcylation reduction in SW620 metastatic colorectal cancer cells
09:20 AM – 09:40 AM	<b>Naphat CHANTARAVISOOT</b> (Chulalongkorn University, Thailand) S4-02: Multi-omics insights into the mechanisms of brain cancer aggressiveness
09:40 AM – 10:00 AM	<b>Narisara CHANTRATITA</b> (Mahidol University, Thailand) S4-03: Host–pathogen transcriptomics in melioidosis: predicting outcomes and revealing resistance mechanisms
10:00 AM – 10:20 AM	Refreshment
10:20 AM – 10:40 AM	<b>Kriengsak LIRDPRAPAMONGKOL</b> (Chulabhorn Research Institute, Thailand) S4-04: ERK-mediated $\beta$ IVa-tubulin up-regulation confers metastasis-associated paclitaxel resistance in naïve lung cancer cells
10:40 AM – 11:00 AM	<b>Kanokwan SRIWATTANAPONG</b> (Chulalongkorn University, Thailand) S4-05: Novel therapeutic strategy to enhance bone regeneration in osteogenesis imperfecta
11:00 AM – 11:20 AM	<b>Phum TACHACHARTVANICH</b> (Chulabhorn Research Institute, Thailand) S4-06: Identification of emerging environmental contaminants as novel metabolism disruptors via PPAR $\gamma$ activation: insights from <i>in vitro</i> and transcriptome studies
11:20 AM – 12:20 PM	Lunch
12:20 PM – 09:00 PM	<b>Networking Excursion (Optional)</b>

## 17 December 2025

Time	Session
	<b>Session 5: Transcription machinery, regulation and diseases</b> <b>Chair:</b> KANG Changwon <b>Co-chair:</b> Narumon THONGDEE
09:00 AM – 09:20 AM	<b>KANG Changwon</b> (Korea Advanced Institute of Science and Technology, Republic of Korea) S5-01: Distinct preferences for termination mechanisms across RNA polymerases
09:20 AM – 09:40 AM	<b>SHIMAMOTO Nobuo</b> (National Institute of Genetics, Japan) S5-02: Biological significance of abortive initiation
09:40 AM – 10:00 AM	<b>HOHNG Sungchul</b> (Seoul National University, Republic of Korea) S5-03: Mechanisms of co-transcriptional R-loop formation at double-stranded breaks revealed by single-molecule fluorescence imaging
10:00 AM – 10:20 AM	Refreshment
10:20 AM – 10:40 AM	<b>Danaya PAKOTIPRAPHA</b> (Mahidol University, Thailand) S5-04: Versatility of MarR domain in the regulation of phenolic compound utilization
10:40 AM – 11:00 PM	<b>ZHANG Yan Jessie</b> (University of Texas, Austin, USA) S5-05: “Zipper” grammar of CTD governs the spatial programing of the transcription cycle of RNA polymerase II
11:00 PM – 11:20 PM	<b>WANG Gang</b> (Fudan University, China) S5-06: Nucleolar MED19 regulates the rRNA 2'-O-methylation in promoting tumorigenesis
11:20 PM – 11:40 PM	<b>Patompon WONGTRAKOONGATE</b> (Mahidol University, Thailand) S5-07: From RNA biology to mRNA therapeutics: from academics to startups
11:40 PM – 01:20 PM	Lunch / IAB, IOC, and ISC Meeting
01:20 PM – 01:40 PM	<b>KIM AeRi</b> (Pusan National University, Republic of Korea) S5-08: TAD-dependent sub-TAD is required for enhancer-promoter interaction enabling the $\beta$ -globin transcription
01:40 PM – 01:55 PM	<b>CHOI Jinmi</b> (Sungkyunkwan University, Republic of Korea) S5-09: Repeat-rich RNA guides genomic repeats into condensates for heterochromatin organization

## 17 December 2025 (continued)

Time	Session
01:55 PM – 02:10 PM	<b>HAN Sun</b> (Seoul National University, Republic of Korea) S5-10: Termination mechanisms of human RNA polymerase III revealed by single-molecule fluorescence imaging
02:10 PM – 03:40 PM	<b>Poster Presentations and Refreshment</b> <b>Session 6: AMR: genomic, mechanisms, and regulations</b> <b>Chair:</b> James M. DUBBS <b>Co-chair:</b> Nisanart CHAROENLAP
03:40 PM – 04:00 PM	<b>Suganya YOUNGKIETTRAKUL</b> (National Science and Technology Development Agency, Thailand) S6-01: From genomic insights to diagnostics: MassARRAY-Based identification and serotyping of foodborne and zoonotic bacteria
04:00 PM – 04:20 PM	<b>Adisak ROMSANG</b> (Mahidol University, Thailand) S6-02: Dual reactive chlorine species responsive sensor adaptation overcomes tolerance to disinfectants and host immunity in <i>Pseudomonas aeruginosa</i>
04:20 PM – 04:40 PM	<b>Anusak KERDSIN</b> (Kasetsart University, Thailand) S6-03: Genomic epidemiology of carbapenem-resistant <i>Enterobacterales</i> in Thailand
04:40 PM – 05:00 PM	<b>Punyawee DULYAYANGKUL</b> (Chulabhorn Research Institute, Thailand) S6-04: Molecular characterization of pandrug- and extensively drug-resistant <i>Stenotrophomonas maltophilia</i> strain SMC1291 and SMC3920
05:00 PM – 05:15 PM	<b>Poonyaporn KANAWONG</b> (Chulabhorn Graduate Institute, Thailand) S6-05: Mutation-induced structural alteration of SoxR enhances antibiotic resistance in <i>Stenotrophomonas maltophilia</i>
05:15 PM – 05:35 PM	Poster Awards Announcement
05:35 PM – 08:00 PM	Closing Ceremony and Conference Dinner