

Program

2 February 2026 (Arnoma Grand Hotel)

8:50	9:00	Opening Remarks by Assoc. Prof. Prasit Suwannalert, Dean, Faculty of Science, Mahidol University
9:00	9:10	Opening Remarks by Prof Igor Mokrousov, Saint Petersburg's Pasteur Institute
		Session chaired by Igor Mokrousov and Prasit Palittapongarnpim
9:10	9:40	Keynote – Taane G. Clark (London School of Hygiene and Tropical Medicine, UK) <i>What can 'omics tell us about tuberculosis?</i>
9:40	10:00	Christophe Sola (Université Paris-Saclay, Saint-Aubin, France) <i>Reconstructing a global evolutionary history of tuberculosis: what are the still unanswered questions ??</i>
10:00	10:20	Kiatichai Faksri (Faculty of Medicine, Khonkaen University, TH) <i>Application of Genomics and OMICS Approaches for the Diagnosis and Control of Mycobacterium and Related Pathogens</i>
10:20	10:50	Coffee Break
		Session chaired by Christophe Sola
10:50	11:10	Amador Goodridge (INDICASAT-AIP, City of Knowledge), Panama) <i>Endemic transmission of Mycobacterium tuberculosis Sublineage L2.2.M3 within Panama</i>
11:10	11:35	Igor Mokrousov (Saint Petersburg Pasteur Institute, Russia) <i>Genomic insight into Mycobacterium tuberculosis adaptation to external stress in the in vivo and in vitro models</i>
11:35	12:35	Presentations sponsored by Qiagen Speaker 1: Vanitha Palaeya (QIAGEN, Malaysia) <i>Transforming TB Genomic Surveillance and Resistance Profiling with QIAseq xHYB Mycobacterium tuberculosis Panel</i> Speaker 2: Zirwatul Adilah Bt Aziz , (National Public Health Surveillance Laboratory, Malaysia) <i>Culture-Free Whole Genome Sequencing: The NPHL Experience with QIAseq xHYB Mycobacterium tuberculosis Workflow</i>
12:35	13:30	Lunch
13:30	14:30	Poster Session 1: Presentation of posters with odd numbers
		Session chaired by Kiatichai Faksri
14:30	14:50	Danila Zimenkov (Engelhardt Institute of Molecular Biology, Russian Academy of Sciences, Moscow, Russia) <i>The hidden diversity of Mycobacterium</i>
14:50	15:10	Nawamin Pinpathomrat (Faculty of Medicine, Prince of Songkla University, Songkla, TH) <i>TB vaccine platforms and delivering system</i>
15:10	15:40	Coffee Break
15:40	16:55	Oral Presentations chaired by Sayera Banu Oren Tzfadia (Institute for Tropical Medicine, Belgium) <i>A next-generation Mycobacterial knowledge base</i> Mohammad Khaja Mafij Uddin (icddr, Bangladesh) <i>Genomic insights into drug resistance and transmission dynamics of multidrug-resistant tuberculosis strains in Bangladesh</i>

		<p>Dyshelly Nurkartika Pascapurnama (Universitas Padjadjaran, Indonesia) <i>Proportion of pathogenic non-tuberculous mycobacteria identified by partial-gene sequencing among presumptive tuberculosis patients in Indonesia</i></p> <p>John Carlo Macasaddug Malabed (Department of Science and Technology, Philippines) <i>A molecular epidemiologic analysis of drug-resistant Mycobacterium tuberculosis isolates derived from the Third Philippines TB Drug Resistance Survey 2018 and community-based study</i></p> <p>Pundharika Piboonsiri (Medical Life Sciences Institute, Department of Medical Sciences, MOPH, Thailand) <i>Whole genome sequencing of Mycobacterium tuberculosis reveals emerging bedaquiline resistance variants in Thailand</i></p> <p>Motunrayo Badejo (Stellenbosch University, South Africa) <i>Superparamagnetic iron oxide nanoparticles perturb mycobacterial homeostasis and unveil gene expression signatures of antimicrobial action</i></p>
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3 February 2026 (Arnoma Grand Hotel)

		Session chaired by Katsushi Tokunaga
9:00	9:30	Keynote – Qian Gao (Shanghai Medical College of Fudan University, China) <i>Molecular epidemiology of Mycobacterium tuberculosis in China</i>
9:30	10:00	Keynote – Katsushi Tokunaga (National Center for Global Health and Medicine, Tokyo, Japan) <i>Genomic approach to infectious diseases</i>
10:00	10:20	Surakameth Mahasirimongkol (Assistant Permanent Secretary, MOPH, TH) <i>Precision Control of TB: The future of ending TB by 2030.</i>
10:20	10:50	Coffee Break
		Session chaired by Richard Anthony
10:50	11:10	Taisei Mushiroda (RIKEN Center for Integrative Medical Sciences, Japan) <i>Toward Patient-Centered, Stratified Tuberculosis Treatment Through Pharmacogenomics</i>
11:10	11:30	Margarita Shleeva (Federal Research Centre “Fundamentals of Biotechnology” of the Russian Academy of Sciences, Russia) <i>A novel biochemical reaction in mycobacteria: coproporphyrin III tetramethyl ester synthesis and its adaptation significance</i>
11:30	11:50	Egor Shitikov (Lopukhin Federal Research and Clinical Center of Physical-Chemical Medicine, Moscow, Russia) <i>Mycobacteriophages: From Laboratory Research to Practical Applications in Medicine</i>
11:50	13:00	Lunch
13:00	14:00	Poster Session 2: Presentation of posters with even numbers
		Session chaired by Amador Goodridge
14:00	14:20	Marisa Ponpuak (Department of Microbiology, Faculty of Science, Mahidol University, TH) <i>Monocyte transcriptional responses to Mycobacterium tuberculosis associated with natural resistance to tuberculosis infection</i>
14:20	14:40	Alexander Apt (Central Institute of Tuberculosis, Moscow, Russia) <i>Tuberculosis infection control by the MHC-II in mice and the sequence of innate/adaptive immune responses to infection</i>
14:40	15:00	Tomasz Jagielski (University of Warsaw, Poland) <i>A water tale – new insights into the ecology of nontuberculous-mycobacteria</i>
15:00	15:20	David Couvin (Institut Pasteur de la Guadeloupe, France) <i>SITVIT++: Next-Generation Mycobacterium tuberculosis databases and biobanks for genomic surveillance, drug resistance monitoring, and predictive intelligence</i>

15:20	15:50	Break
15:50	17:00	<p>Oral Presentations chaired by Marisa Ponpuak</p> <p>Noppadon Nuntawong (National Electronics and Computer Technology (NECTEC), Thailand) <i>Engineering a robust SERS-based POC system for tuberculosis screening</i></p> <p>Saradee Warit (National Center for Genetic Engineering and Biotechnology (BIOTEC) Thailand) <i>Advancing tuberculosis diagnosis in macaques: A One Health approach</i></p> <p>Narisa Mohthong (Prince of Songkla University, Thailand) <i>Investigating immune profiles in healthy and latent tuberculosis-infected participants.</i></p> <p>Yoopie Setiawan (Universitas Katolik Parahyangan, Indonesia) <i>The interferon-gamma response to Latin American Mediterranean and modern Beijing sub-lineages of Mycobacterium tuberculosis in drug-sensitive and resistant tuberculosis</i></p> <p>Pyae Sone Oo (Prince of Songkla University, Thailand) <i>Differential immune responses to novel and established Mycobacterium tuberculosis antigens among tuberculosis patients in Southern Thailand</i></p>

4 February 2026 (Arnoma Grand Hotel)

		Session chaired by Qian Gao
9:00	9:30	<p>Keynote – Urvashi Singh (All India Institute of Medical Sciences, Delhi and Head of National TB program, India) <i>Mycobacterium tuberculosis genomics and transmission tracing in India</i></p>
9:30	9:50	<p>Virasakdi Chongsuvivatwong (Faculty of Medicine, Prince of Songkla University, Songkla, TH) <i>Geno-spatio-temporal analysis, interpretation and implications on Infectious diseases</i></p>
9:50	10:10	<p>Richard Anthony (National Institute for Public Health and the Environment, Netherland) <i>Micro-evolution in clustered cases of tuberculosis a useful signal or noise?</i></p>
10:10	10:40	Coffee Break
		Session chaired by Igor Mokrousov
10:40	11:00	<p>Sayera Banu (Head, Programme on Emerging Infections, Infectious Diseases Division, icddr, Bangladesh) <i>Targeted next-generation sequencing for detection of drug-resistant tuberculosis: Challenges and insights from high burden settings in Bangladesh</i></p>
11:00	11:30	<p>Pakorn Aiewsakun (Center for Microbial Genomics, Faculty of Science, Mahidol University, TH) <i>Whole genome sequence analysis to detect Mycobacterium tuberculosis transmission cluster</i></p>
11:30	12:00	<p>Robin Warren (Stellenbosch University, South Africa) <i>The genesis and transmission of drug-resistant TB.</i></p>
12:00	13:00	Lunch
13:00	14:00	<p>Oral Presentation chaired by Violeta Valcheva</p> <p>Tania Jim (National University of Singapore, Singapore) <i>The use of a novel conjugated oligoelectrolyte molecule to potentiate antibiotic efficacy against Non-Tuberculous Mycobacteria</i></p> <p>Kishan Kumar Parida (National Institute of Pharmaceutical Education and Research, India) <i>Structure based identification and validation of Benzothiazole derivatives as potent DprE1 inhibitors against Mycobacterium tuberculosis</i></p> <p>Hleziphi Violah Mpundu (Stellenbosch University, South Africa) <i>Targeting drug-tolerant tuberculosis through repurposing of approved drugs</i></p> <p>Arnab Roy (National Institute of Pharmaceutical Education and Research, India) <i>Overcoming metabolic redundancy in Mycobacterium tuberculosis via dual inhibition of terminal oxidases</i></p>

		Session chaired by Egor Shitikov
14:00	14:20	Mi Kaixia (Institute of Microbiology, Chinese Academy of Sciences, China) <i>Elucidating the dual role of mycobacterial MfpA in fluoroquinolone resistance and virulence regulation</i>
14:20	14:40	Violeta Valcheva (Department of Infectious Microbiology, Stephan Angeloff Institute of Microbiology, Bulgarian Academy of Sciences, Bulgaria) <i>Development and pharmacological evaluation of novel hydrazide - hydrazone derivatives as potential antituberculosis drug candidates</i>
14:40	15:00	Oleg Ogarkov (Scientific Center for Family Health and Human Reproduction Problems, Irkutsk, Russia) <i>The formation of a polybacterial community in caseous necrosis: a common and adverse consequence of tuberculosis</i>
15:00	15:20	Prasit Palittapongarnpim (Center for Microbial Genomics, Faculty of Science, Mahidol University, TH) <i>Structural variations of the pe/ppe gene family</i>
15:20	15:30	Awarding and Closing ceremony
15:30	16:30	Coffee Break

Abstracts (Poster presentations)

1. **P-01: Xiaoming Liu** (Beijing Chest Hospital, Capital Medical University; Beijing Tuberculosis and Thoracic Tumor Research Institute, China) | **Global burden of childhood tuberculosis (1990–2021): a systematic analysis from the Global Burden of Disease Study 2021**
2. **P-02: Zaidah Abdul Rahman** (School of Medical Sciences, Universiti Sains Malaysia, Malaysia) | **Latent tuberculosis infections (LTBI) among healthcare workers & laboratory staff: is screening mandatory?**
3. **P-03: Lapasrada Pattarapreeyakul** (Medical Life Sciences Institute, Department of Medical Sciences, MOPH, Thailand) | **Geographical distribution and characteristics of cutaneous non-tuberculous mycobacterial infections in Thailand from 2015 to 2024**.....
4. **P-04: Mahfuza Talukder Flowra** (Oslo Metropolitan University, Oslo, Norway) | **Barriers of multidrug-resistant tuberculosis (MDR- TB) detection from the community perspective in Bangladesh: A mixed method Study**
5. **P-05: Ayush Bhutada** (Indira Gandhi Government Medical College Nagpur, India) | **Open Negative Syndrome in tuberculosis: A rare diagnostic dilemma post-treatment**
6. **P-06: Ayush Bhutada** (Indira Gandhi Government Medical College Nagpur, India) | **Unveiling Gender and BMI influences on outcome of hospitalization in TB cases: Insights from a Central India hospital study**
7. **P-07: Rashmi Ratnam** (Microbiology, King George's Medical University, Lucknow, India) | **Influence of diabetes and socioeconomic variables on treatment response in pulmonary tuberculosis patients**
8. **P-08: Rashmi Ratnam** (Microbiology, King George's Medical University, Lucknow, India) | **Emerging phenotypic resistance to delamanid and pretomanid in MDR-TB: Co-resistance patterns from a North Indian Reference Laboratory**
9. **P-09: Nenekazi Masikantsi** (Department of Biomedical Sciences, Faculty of Medicine and Health Sciences, Stellenbosch University, South African) | **CRISPR interference-mediated discovery of antitubercular compounds from South African medicinal plants**.....
10. **P-10: Nuhu Ibrahim Tukur** (Department of Biomedical Sciences, Faculty of Medicine and Health Sciences, Stellenbosch University, South Africa) | **Discovery of potent DnaK-targeting Cyclomarin- A derived BacPROTACs as a potential anti-tuberculosis therapeutic modality**.....
11. **P-11: Onpreeya Kritwatcharas** (Department of Medical Technology, Faculty of Associated Medical Sciences, Chiang Mai University, Thailand) | **Investigation of propolis-loaded niosomes for tuberculosis: anti-mycobacterial activity and three-dimensional granuloma model**
12. **P-12: Sujata Sharma** (Department of Biophysics, All India Institute of Medical Sciences, India) | **Lactoferrin as a multifunctional modulator of oxidative stress and antimycobacterial immunity in tuberculosis**.....
13. **P-13: Nikhat Khan** (Symbiosis International University, India; Regional Medical Research Centre, India) | **Population genetic analysis of NAT2 Gene in Saharia Tribe: A particularly vulnerable tribal group of Central India**.....

14. **P-14: Pathida Prakongsup** (Department of Biochemistry, Faculty of Pharmacy, Mahidol University, Thailand) | **The association study between DNA methylation on *CYP2D6* and *CYP2E1* gene promoter and anti-tuberculosis drug-induced liver injury in tuberculosis patients in Thailand**
15. **P-15: Taratorn Kemthong** (Faculty of Science, Chulalongkorn University, Thailand) | **Evaluating the tuberculin skin test as a practical tool for staging *Mycobacterium tuberculosis* infection in cynomolgus macaques**
16. **P-16: Prapaporn Srilohasin** (Faculty of Medicine Siriraj Hospital, Mahidol University, Bangkok, Thailand) | ***Mycobacterium tuberculosis* Beijing Lineage from cynomolgus macaques compared with human isolates in Thailand**
17. **P-17: Zhenli Liang** (Guangxi Center for Disease Control and Prevention, China) | **The impact of Influenza on TB treatment outcomes: a retrospective matched cohort study in Guangxi, China, 2012-2024**
18. **P-18: Natapohn Saowaphong** (Faculty of Medicine, Prince of Songkla University, Thailand) | **Evaluation of stability and viability of BCG microneedle vaccine derived from three different conditions**
19. **P-19: Trinh Minh Phuong** (College of Medicine, Gyeongsang National University, Republic of Korea) | **Disentangling relapse and reinfection in *Mycobacterium intracellulare* using MIRU-VNTR genotyping**
20. **P-20: Minh Phuong Trinh** (College of Medicine, Gyeongsang National University, Republic of Korea) | **Gram-negative-driven TLR4 activation reshapes cytokine dynamics and host response during *Mycobacterium avium* infection**.....
21. **P-21: Sahasawat Suksan** (Department of Biomedical Science and Biomedical Engineering, Prince of Songkhla University, Thailand) | **Modeling human tonsil organoids to study adaptive immune responses against BCG**
22. **P-22: Suthirote Meesawat** (Department of Biology, Faculty of Science, Chulalongkorn University, Thailand) | **Rapid detection of antibody against *Mycobacterium tuberculosis* complex in synanthropic rhesus (*Macaca mulatta*) and long-tailed (*M. fascicularis*) macaques across Thailand**
23. **P-23: Anna Vyazovaya** (St. Petersburg Pasteur Institute, Russia) | **Overwhelming dominance of the Beijing genotype among pre-extensively drug-resistant *Mycobacterium tuberculosis* strains in western Siberia, Russia**
24. **P-24: Nik Mohd Noor Nik Zuraina** (Department of Medical Microbiology & Parasitology, School of Medical Sciences, Universiti Sains Malaysia, Malaysia) | **Rapid detection of *Mycobacterium tuberculosis* SIT745/EAI1-MYS Using a five-spacer multiplex PCR**.....
25. **P-25: Manu Singh** (Department of Respiratory Medicine, King George's Medical University, India) | **Discrepancies in molecular and culture-based detection of central nervous system tuberculosis and its drug resistance profile**.....

26. **P-26: Waritta Sawaengdee** (Department of Medical Sciences, MOPH, Thailand) | **Whole-genome sequencing reveals the co-dominance of *Mycobacterium tuberculosis* lineages 1 and 2 in Thailand**
27. **P-27: Pornpen Tantivitayakul** (Department of Oral Microbiology, Faculty of Dentistry, Mahidol University, Thailand) | **Interactions of human genomes and mycobacterial genomes contribute to severity and transmissibility of pulmonary tuberculosis**
28. **P-28: Manita Yimcharoen** (Department of Medical Technology, Faculty of Associated Medical Sciences, Chiang Mai University, Thailand) | **Adaptive genome remodeling and conserved virulence shape interactions between host and pathogen in drug-resistant *Mycobacterium tuberculosis***.....
29. **P-29: Zayar Phyo** (Department of Medical Technology, Faculty of Associated Medical Sciences, Chiang Mai University, Thailand) | **Comparative transcriptional signatures reveal strain-specific isoniazid responses in *Mycobacterium tuberculosis* under host-mimicked stress conditions** ...
30. **P-30: Grigorii Sergeev** (St. Petersburg Pasteur Institute, St. Petersburg, Russia) | **Development of bedaquiline resistance in *Mycobacterium tuberculosis* serial isolates recovered during treatment of MDR/pre-XDR tuberculosis patients**
31. **P-31: SM Rezvi** (Microbiology Department, Faculty of Medicine, Universitas Andalas, Indonesia) | **Development and validation of an RT-PCR Kit for rapid detection of *Mycobacterium tuberculosis* in clinical samples**.....
32. **P-32: Sushma Yadav** (King George's Medical University, India) | **Identification of Drug-resistant profile of multidrug-resistant patients receiving bedaquiline and delamanid based regimen**
33. **P-33: Charintip Yenyuvadee** (Medical Life Sciences Institute, Department of Medical Science, MOPH, Thailand) | **Comparative analysis of phenotypic and whole genome sequencing for drug susceptibility testing in *Mycobacterium tuberculosis* isolates**
34. **P-34: Ye Win Aung** (Department of Microbiology, Faculty of Medicine, Khon Kaen University, Thailand) | **Heteroresistance of *Mycobacterium tuberculosis* in the sputum detected by Droplet Digital PCR**.....
35. **P-35: Arkan Hidayat** (Stem Cell and Cancer Institute, Indonesia) | **Performance of Wide-Coverage Open-System RT-PCR to detect *Mycobacterium tuberculosis*, rifampicin, and isoniazid resistance in sputum samples from presumed pulmonary tuberculosis patients in Indonesia**.....
36. **P-36: Muhammad Mu'iz Ehsannudin bin Abu Bakar** (National Mycobacteria Reference Laboratory, Department of Laboratory Services, Ministry of Health, Brunei Darussalam) | **High prevalence of inducible macrolide resistance in *Mycobacterium abscessus* Complex revealed by nationwide implementation of the GenoType NTM-DR Assay in Brunei Darussalam** ...
37. **P-37: Selly Erwina** (Faculty of Medicine, Universitas Andalas, Indonesia) | **Potency of whole-genome sequencing for mapping lineage and drug resistance of *Mycobacterium tuberculosis* in Indonesia: a systematic review**