



Welcome to SCELSE's Open House, an opportunity to delve into the unseen world of microbes that shape our health and environment. As you explore our booths, you will embark on a journey through five themes that reflect the breadth of SCELSE's research. Meet The Invisible Architects, microbes building complex and resilient communities called biofilms, and find out how they can either support or disrupt our health and environment. Uncover how microbes become Host Hackers and explore how microbes interact with the human body in both harmful and beneficial ways, from evading immune defences during infection to supporting skin health. The Microbial Menu theme invites you to explore the good, the bad, and the edible sides of microbiology, from food spoilage to probiotics for gut health and sustainable food production. Dive into the theme of Hot Springs to High Seas: Exploring Microbial Frontiers, revealing the abundance and significance of microbial life in marine settings, hot springs, and wastewater treatment systems. Finally, discover how we decode genomic diversity in the theme From Genomes to Algorithms, where cutting-edge sequencing and computational tools unlock the secrets of microbial communities and their hosts or environments.

Each booth showcases how multidisciplinary science at SCELSE advances our knowledge of the microbial world. Start anywhere, chat with our scientists, and get ready to be amazed by the microbial world!

Programme

11.30 am Registration

12.15 pm Welcome address by Prof Alain Filloux,

SCELSE Centre Director

12.20 pm Path to PhD by Ms Sumathi Lechimanan.

Assistant Manager, Student & Academic Services

12.30 pm Fireside chat with SCELSE experts

1.20 pm Networking Lunch with Professors

2.40 pm Lab tours 1

3.40 pm Microbiology seminars: chaired by A/Prof Janelle Thompson

3.45 pm • Talk 1: Microbial solutions for economic & climate resilience in food production

- Dr Andrew Laloo, SCELSE-NUS Research Fellow • Talk 2: Microbiomes influencing ocean health
- Dr Jeslyn Poo, SCELSE-NTU Research Fellow
- Talk 3: Biofilms in chronic infection: A biophysical

Dr Binu Kundukad, SCELSE-NTU Senior Research Fellow

4.40 pm Lab tours 2 End

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Wheel of Microbial Fortune

Start your Open House experience with a spin! While you wait to register, try SCELSE's Wheel of Microbial Fortune. With so many booths to explore, this quick activity helps point you toward the booth you should visit first.



Registration

Register at SCELSE Level 1 to kickstart your Journey into the Microbial World! Pre-registered guests will receive a special redemption card, which can be used to collect an exclusive goodie bag at the SCELSE Level 1 redemption counter. Don't miss out, fun surprises and useful keepsakes



Overview of SCELSE

Take a minute to learn what SCELSE is all about and explore the fascinating world of biofilms and microbiomes. Check out our posters to see where biofilms and microbiomes can be found and why these microbial communities matter for health and the environment. A quick stop to set the stage for the rest of your visit.



THEME 1: THE INVISIBLE ARCHITECTS

Microbial biofilm - A visit to the House of Bacteria



Prof Staffan Kjelleberg Advisor, SCELSE
Director, Singapore National
Biofilm Consortium Emeritus Distinguished University Professor, NTU



Dr Wong Lan Li SCELSE-NTU Research Fellow



SCELSE-NTU Research Fellow

A microbial biofilm matrix is like a sticky, slimy layer made by microorganisms to house and protect themselves. Imagine a group of tiny organisms, like bacteria, creating a mixture of gooey substances including proteins, polysaccharides or nucleic acids that act like glue to hold them



This slimy matrix helps them stick to surfaces and stay close to each other, preventing them from being easily washed away. It's the basis for the plaque that forms on your teeth or the slimy coating on the inside of a water bottle if it's not cleaned regularly.

Visit us to learn more about how we study the composition of these matrices, combining knowledge from biology, chemistry, and biophysics, and how this understanding helps us to harness and control the microbial inhabitants.



Understanding bacterial cell envelope biology and its impact on intrinsic antimicrobial resistance



A/Prof Chng Shu Sin Associate Director, SCELSE-NUS PI, Biofilms & Health cluster



SCELSE-NUS Research Fellow



SCELSE-NUS PhD student

Many bacterial species build an additional layer of protective membrane. called the outer membrane, surrounding the cell. The outer membrane blocks the entry of antibiotics, thus conferring intrinsic antimicrobial resistance

Find out how we are using multiple scientific disciplines to unravel the fundamental biological basis of how bacteria build and maintain their outer membranes. A variety of approaches, including but not limited to biochemistry, genetics, structural biology, and chemical biology are allied to study the molecular mechanism of protein complexes that build and maintain the stability of the outer membrane in model organisms like Mycobacteria smeamatis and Escherichia coli.

The fundamental knowledge we generate fuels future development of antibiotics, for example, targeting the outer membrane directly or bypassing the outer membrane to kill the cells.

THEME 2: THE HOST HACKERS



Exploiting Pseudomonas aeruginosa to prevent biofilm formation and tackle antibiotic resistance



Prof Alain Filloux Centre Director, SCELSE Research Director, Biofilms Snr Research Fellow & Health cluster



Dr Sujatha Subramoni SCELSE-NTU



Dr Kay To SCELSE-NTU Research Fellow



Dr Casandra Tan SCELSE-NTU Research Fellow



SCELSE-NTU

Research Fellow

PhD student



SCELSE-NTŬ PhD student

Discover the secrets of biofilms at our booth! Biofilms are clusters of microbes that attach to surfaces and are shielded by an extracellular polymeric substance (EPS) matrix, a sticky and slimy layer composed of various macromolecules.

This extracellular matrix acts as a protective layer for bacteria against antimicrobials and environmental stress, making infections harder to treat. However, the mechanisms that drive matrix formation are complex, making it challenging to devise preventive strategies against the formation of this protective barrier. Visit our booth to explore how biofilms impact health, how bacteria gain an advantage within these communities, and the technologies being developed to tackle biofilm formation.

Host-E. faecalis interactions in wound infection and infective endocarditis



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Prof Kimberly Kline Visiting Academic Department of Microbiology & Molecular Medicine, Faculty of Medicine, University of Geneva



Dr Haris Antypas SCELSE-NTU Snr Research Fellow



Frederick **Reinhart Tanoto** SCELSE-NTU PhD student

Take a peek into the secret relationships of microbes and their hosts. Invading bacteria and host defenders are in a constant tug-of-war to win the battle of infection. Some bacteria are experts at hijacking the human body. What strategies do bacteria use to win this battle? And how can we ensure the victory of host defences instead?

Our group specialises in dissecting the role and mechanisms of bacterial virulence factors in establishing wound and heart infections. We employ in vivo infection models, as well as various in vitro cellular and molecular techniques (e.g., microscopy, next generation sequencing, and in vitro cell infection models) in our investigation. Join our tactical specialist team as we uncover bacterial virulence strategies and pave the way for host victory!

Exploring bacterial lifestyles



Prof Cynthia Whitchurch Research Director, Biofilm Biology cluster, SCELSE Professor, School of Biological Sciences, NTU



Dr Seetanshu Junnarkar SCELSE-NTU Snr Research Fellow



Dr Sharan Prakash SCELSE-NTU Research Fellow



Dr Micaela Mossop SCELSE-NTU Research Fellow



Alicia Pek SCELSE-NTU Visiting PhD student

Are you fascinated by microbial strategies like host evasion, biofilm formation, or phenotypic switching? Come dive into the hidden world of microbes and discover how they collaborate, adapt, and survive. Our poster showcases:

- · Biofilm Expansion powered by the versatile slime of extracellular DNA (eDNA)
- The stealthy persistence of gut pathobionts—typically harmless gut bacteria that can turn against us under certain conditions
- Microbial shapeshifting—is it a superpower or an Achilles heel?

Whether you're exploring new research angles or looking for your next project, this is your chance to engage with cutting-edge ideas and connect with fellow researchers. Stop by our booth and see how microbes are rewriting the rules of survival

SCELSE-L'Oréal Joint Laboratory



Prof Cynthia Whitchurch Biology cluster, SCELSE Professor, School of Biological Sciences, NTU



Dr Marcus Lee Research Fellow



Samantha Quah Research Director, Biofilm SCELSE-L'Oréal Joint Lab SCELSE-L'Oréal Joint Lab Research Associate /PhD Student



Karaiyagowder Govindarajan Deenadayalan SCELSE-L'Oréal Joint Lab Research Associate



Hashmath Fatimah SCELSE-L'Oréal Joint Lab Research Assistant

Let's explore exciting research at the SCELSE-L'Oréal Joint Laboratory, where we delve into the fascinating world of skin microbiomes and see how microorganisms contribute to playing a part in promoting skin health.

We utilise state-of-the-art laboratory-scale microbiome models, cutting-edge microscopy imaging techniques, and advanced molecular methods to understand the skin microbiome's role in promoting healthy skin and hair, and to create innovative and targeted solutions for treating skin and hair conditions.

If you are interested to learn more about what we do, come talk to our friendly team and we'll be happy to share more!

THEME 3: MICROBIAL MENU: THE GOOD, THE BAD & THE EDIBLE





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Prof Stefan Wuertz Deputy Centre Director (Education & Training) Research Director, Environmental Engineering cluster



Dr Ezequiel Santillan SCFLSF-NTU Snr Research Fellow



Dr Anika Cokro SCFLSF-NTU Research Fellow



Dr Anisa Cokro SCFLSF-NTU Research Fellow



Dr Woo Yissue SCELSE-NTU Research Fellow



Dr Soheil Neshat SCELSE-NTU Research Fellow

Revolutionise aquaculture with our innovative approach using microbial community-based protein to replace traditional fishmeal. Our research showcases the pivotal role of microbes in creating a sustainable, circular bioeconomy. By transforming food processing wastewater into high-quality single-cell protein (SCP), we're using microbes to turn waste into valuable resources, enhancing the efficiency of food production systems, and addressing critical challenges in protein sustainability and food security



Studying biofilms from "farm to fork" to reduce food waste



Asst Prof Laura Nolan Pl. Environmental Engineering cluster, SCELSE Assistant Professor, School of Biological Sciences, NTU



Dr Zhao Xue SCELSE-NTU Research Fellow



SCELSE-NTU Research Assistant



John Harold Nataño SCELSE-NTU PhD student



SCELSE-NŤU Visiting PhD student

Almost 10% of the world's population do not get enough to eat. Despite this, around 20% of all food produced for human consumption worldwide is wasted. One major contributor to this wastage is microbial food spoilage. These microbes grow on food and food chain surfaces in complex structures called biofilms. While we know which microbes cause food spoilage, we know very little about how they form biofilms in the food chain prior to spoilage occurring. We are studying spoilage microbe biofilm development from "farm to fork" using a range of cutting-edge approaches. We are then working to use this understanding to develop new ways to reduce food spoilage and extend food shelf life, which will contribute to a reduction in global food waste.

Synbiotic formulations for nutraceutical applications



Prof Joachim Loo PI, Environmental Engineering, SCELSE School of Materials Science & Engineering, NTU



Zhao Zhongtian SCELSE-NTU Visiting PhD student



Clive Ng Yi Jie SCELSE-NTU Visiting PhD student

Did you know that some bacteria can enhance gut health? Our team is developing a synbiotic system based on the Bacillus subtilis SOM8 bacterial strain isolated from sesame oil meal. This strain demonstrated potent antipathogenic activity, high tolerance to gastrointestinal stressors, and key probiotic traits including antioxidant and bile salt hydrolase activity. Safety evaluations confirmed low cytotoxicity, absence of mobile genetic elements, and intrinsic resistance only to streptomycin. Additionally, SOM8 produces EPS-SOM8, a levan-like exopolysaccharide with β -2,6 and β -2,1 linkage, resistant to digestion and selectively fermented by beneficial microbes such as Bifidobacterium longum DSM20219. In vitro gut models revealed that EPS-SOM8 modulates microbiota composition and enhances short chain fatty acids production. These findings highlight the dual probiotic and prebiotic potential of B. subtilis SOM8 for microbiome-targeted applications.

Functional foods for gut health modulation



Prof Patricia Conway

Visiting Professor, SCELSE

Edric Sim SCELSE-NTU PhD student



Lim Ye Joon SCELSE-NTU

You are what you eat! Join us to discover how your diet, among many factors, affects your gastrointestinal health and wellbeing!

Our group has harnessed the metabolic potential of microorganisms to develop several novel functional foods, including fermented oats beverages and black soldier fly larvae hydrolysate protein. We have shown that these foods have the potential to modulate the gut microbiome for the improvement of gastrointestinal health and wellbeing.

Find out how we collaborate with Singapore National Biofilm Consortium (SNBC) and industry partners such as Protenga, Adisseo, and Haleon to develop and investigate functional foods that modulate the microbiome for improved gut health.

Establishment of plastic-associated microbial community from superworm gut microbiome

Gut microbiomes of plastic-munching superworms offer promising avenues

for plastic biodegradation. Due to the complexity of worm maintenance.

direct application is impractical. Instead, we established stable plastic-

associated biofilm communities via a two-stage enrichment: feeding

Zophobas atratus worms with different types of plastics (HDPE, PP, and PS),

followed by in vitro incubation of their gut microbiomes. While plastic feeding

minimally affected bacterial diversity, it significantly altered community

composition and enriched potential degraders. In vitro incubation induced

more pronounced shifts, further enriching plastic-degrading taxa. Functional

inference and plastic characterization confirmed enhanced degradation

potential. Deterministic ecological processes, especially selection, drove

community succession. Our findings demonstrate that worm-derived

microbial communities can be harnessed to develop biotechnologies for



A/Prof Cao Bin PI, Biofilm Biology & Environmental Engineering clusters, SCELSE Associate Professor, School of Civil and Environmental Engineering, NTU

plastic waste processing.



Dr Sakcham Bairoliya SCELSE-NTU



Dr Liu Yinan SCELSE-NTU Visiting Research Fellow Visiting Research Fellow

Decoding a_cute diarrhoea in the community



A/Prof Yann Boucher Pl. Meta-o'mics & Microbiomes cluster, SCELSE Associate Professor, Saw Swee Hock School of Public Health, NUS



SCELSE-NUS Research Assistant



Su Shiqi SCELSE-NUS Research Assistant

A/Prof Janelle Thompson PI Environmental Engineering cluster, SCELSE Associate Professor, Asian School of Environment, NTU

MICROBIAL FRONTIERS

or public health



THEME 4: HOT SPRINGS TO HIGH SEAS: EXPLORING

Vastewater surveillance and extremophile monitoring

Dr Desmond Chua SCELSE-NTU Research Fellow



SCELSE-NTU Visiting PhD student

Always getting food poisoning or stomach flu? Or knows someone who does? Let's find out what is really going around!

Acute diarrhoea affects thousands in Singapore every year; many cases go unexplained, with the exact cause often unknown.

Here, we aim to address these knowledge gaps to uncover the real culprits behind food poisoning. By analysing stool samples, collecting lifestyle and dietary information, we can identify bacteria, viruses, and parasites that are responsible for diarrhoea.

Be part of an important discovery. Join this study and contribute your sample; you can help uncover the causes of illness in Singapore and contribute valuable insights for improved prevention and public health strategies.

Take the first step! Sign up today and help make a difference!'



Scan for more information

As we transition into the post-COVID-19 era, our team is constantly exploring how we can further utilise the wastewater surveillance used to detect local outbreaks, to monitor novel viruses. We employed qPCR and target enrichment sequencing, combined with both read- and assembly-level analyses, and identified a wide range of viruses on a university campus. We also detected unknown viruses as a proof of concept that wastewater surveillance can support Disease X monitoring.

We leverage genomic surveillance, from urban wastewater to extreme environments, to uncover microbial diversity, detect emerging viruses, and understand ecosystem responses.

Find out how our research is also investigating microbes in extreme environments such as hot springs to understand:

Which microbes are present?

What roles do they play in their community?

How do they respond to environmental disturbances?



Marine biofilms & microbiomes



A/Prof Rebecca Case PI, Biofilm Biology cluster, SCELSE Associate Professor School of Biological Sciences, NTU



Dr Clarence Sim SCELSE-NTU Research Fellow



Prasha Maithani SCELSE-NTU Research Assistant



Ananya Verma SCELSE-NTU



Microbes abound in marine settings, with marine biofilms:

- Playing pivotal roles in biogeochemical cycles,
- · Developing and maintaining ecosystems, and
- · Supporting marine life

Understanding the role of marine microbes (including bacteria, archaea, viruses, fungi and algae) in the formation of complex ecological networks is a focal point in marine biology, environmental microbiology, and industries that operate in marine environments.

SCELSE investigates these microorganisms and interactions, such as:

- Factors affecting seagrass function & health.
- · Production of coral probiotics to enhance resilience to stressors, and
- Molecule discovery in bacterial-algal interactions

THEME 5: BIOINFORMATICS (FROM GENOMES TO ALGORITHMS)



From humans to turtles: Exploring evolutionary history current diversity, and future implications spanning public nealth to conservation through whole-genome research.



A/Prof Kim Hie Lim PI, Meta-'omics & Microbiomes cluster, SCELSE, Asian School of the Environment, NTU



Dr Amit Gourav Ghosh SCELSE-NTU Research Fellow



Dr Regine Tiong ASE-NTU Research Fellow



Sim Chin Yee Faith SCELSE-NTU PhD student



1. GenomeAsia 100K

We're helping to uncover the genetic fabric of Asia through the GenomeAsia 100K project—sequencing thousands of diverse populations from across the continent to uncover deep demographic histories, understand present-day genetic diversity influencing health and immunity (through HLA immune genes), and pave the way for equitable precision medicine.

2. Turtle conservation genomics

Our conservation genomics project focuses on hawksbill sea turtles, a critically endangered species. Using non-invasive sampling (e.g., unsuccessful eggs and hatchlings, eggshells and sand), we sequence turtle genomes to study population structure, natal origins, and genetic diversity insights that are vital for conservation efforts in Southeast Asia.

8B The dynamics of airborne microbes



Prof Stephan Schuster Deputy Centre Director (Facilities & Capacities). Research Director, Meta-o'mics & Microbiomes cluster Professor, School of Biological Sciences, NTU President's Chair in Genomics, NTU Scientific Director, GenomeAsia 100K



Dr Anton Sadovov SCELSE-NTU Snr Research Fellow



Scan for more on Air Microbiome



Vineeth Kodengil SCELSE-NTLI Research Scientist



Santhi Puramadathil Sasi SCELSE-NTU SCELSE-NTU Research Associate Research Fellow



SCELSE-NTU Research Associate

Airborne microbes affect our health and the environment in ways we are only beginning to understand. Traditional methods cannot capture the full picture of these invisible communities, leaving us vulnerable to unknown risks and missing potential benefits. SCELSE's air microbiome team pioneers novel air sampling and analysis strategies. Using next-generation sequencing and highperformance computing, they developed a robust workflow for studying ultralow biomass air samples. Through these techniques, we are now able to understand the complex airborne microbial dynamics.

Uncovering diverse microbial mechanisms via computation



Asst Prof Anni Zhang Courtesy Appointment, SCELSE Assistant Professor School of Biological Sciences, NTU



Jeffry Gaston Computer Scientist



Qiwen Zheng SBS PhD student



Zhiqi Tanq SBS PhD student

In earlier research. Prof Anni Zhang's team employed computational methods to investigate the adaptation mechanisms and phage-host interactions within the human gut microbiome, motivated by the challenge that healthy microbes often struggle to survive in patients' gut environments after faecal microbiota transplantation. The analysis flagged limitations in existing tools for microbial sequence analysis, leading to the team's development of X-Mapper for sequence alignment and QuickVariants for variant identification. These tools are specifically tailored to microbial data to improve accuracy and efficiency.

Come to Asst Prof Zhang's booth to learn how her lab is now developing Al models to explore broader microbial mechanisms, including DNA-protein binding, genetic sweeps, regulatory networks, and host-microbe interactions.

SCELSE Business Development & Partnerships (BDP)



Dr Maria Yung Assistant Director, Business Development & Partnerships



Dr James Ho Snr Research Fellow

Bridging the gap between academic research and industry is critical to ensure that cutting-edge biofilm and microbiome research is translated into market facing products.

BDP facilitates academic-industry connections, offering expertise through:

Consultancies | Joint research initiatives | Collaborative agreements.

SCELSE welcomes industry members to partner with us for unparalleled access to knowledge and technical skills.

Contact:

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 Dr James Ho Snr Research Fellow Email: james.hcs@ntu.edu.sg

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Singapore National Biofilm Consortium (SNBC)



Prof Staffan Kjelleberg Director, Singapore National Biofilm Consortium



Dr Sheela Reuben Assistant Director & Programme Manager



Mr Frnest Tech Assistant Manager

The Singapore National Biofilm Consortium (SNBC) is a national platform that serves as a coordinating hub at the convergence of health, engineering, technology, and science, that integrates innovation and business in biofilm and microbiome technologies to address emerging challenges across diverse industries. SNBC focusses on deepening collaborations and connections to catalyse academic-industry partnerships that lead to relevant industry outcomes and economic impact. This initiative is based on a multidisciplinary research platform and skill base, with translational capacity at SCELSE and other institutions across Singapore.

Contact:

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