

ME Spring Seminar 07

Seeing the Invisible: Bridging Clinical Microbiome Cohorts and Single-Cell Engineering for Precision Medicine



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While large-scale clinical cohort studies have successfully identified various microbial signatures associated with human diseases-ranging from metabolic disorders to the gut - liver axis - the field currently faces a critical transition from association to causation. Conventional bulk-sequencing approaches (16S rRNA and Shotgun Metagenomics) provide a population-level snapshot but often obscure the functional heterogeneity and metabolic activity of individual microbes within complex ecosystems.

In this seminar, I will present my research group's work on deciphering the complex interplay between the gut microbiome and host health, with a particular focus on the gut-liver axis and immune modulation. I will discuss our approach to managing high-dimensional clinical cohort data, integrating shotgun metagenomics and metabolomics, and our efforts to validate these findings through *in vivo* and *in vitro* models. We will explore how deconfounding analysis and multi-omics integration can reveal hidden microbial signatures that drive disease pathogenesis.

Finally, I will briefly share some 'Bridging Questions' regarding the current technical bottlenecks in microbiome research. From a clinical researcher's perspective, I will touch upon the potential of emerging single-cell technologies to address challenges such as functional heterogeneity and real-time metabolic activity, which remain largely invisible through conventional sequencing alone. This session aims to foster an interdisciplinary dialogue on how precision measurement tools might be integrated into large-scale clinical frameworks.

Bldg.110 #N105

16:00 - 17:15

Wednesday, May 6



Host:
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