# Studying the Skin Microbiome in Singapore: Diversity, genetics and functions

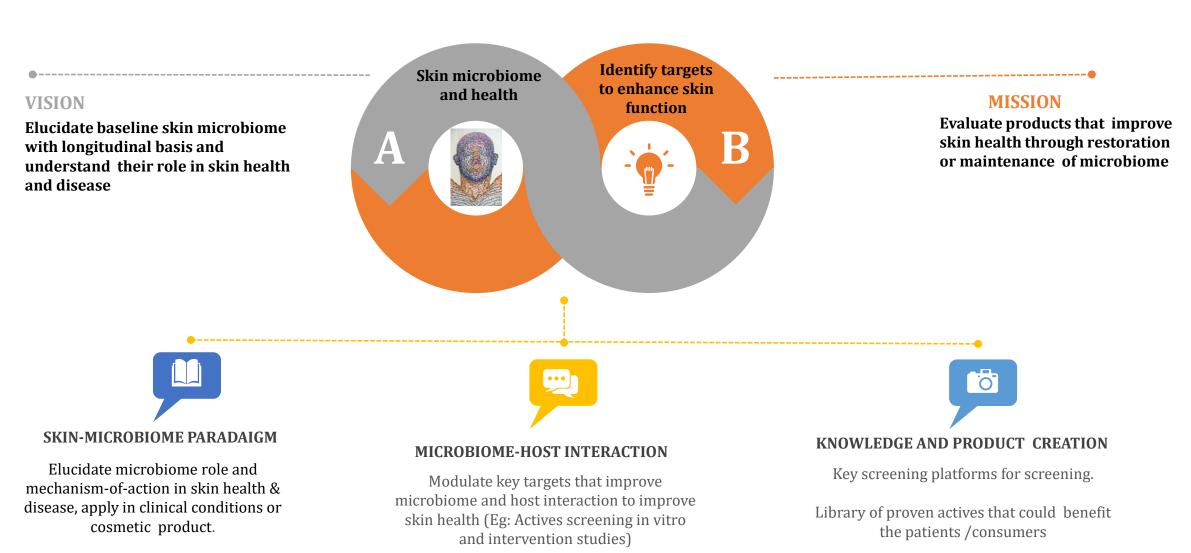
Thomas Dawson John Common Nirajan Nagarajan

Skin Research Institute of Singapore GIS A\*STAR



#### **Asian Skin Microbiome Program**







Skin Research Institute of Singapore

### Asian Skin Microbiome Program



**Comprehensive Multi-omics**, **Optimized sampling** Data integration and analysis from different body sites **Metagenomics** Microbiome, Transcriptome, **Transcriptomics** Lipidome and Metabolome **Metabolomics (Lipidomics) Skin Physiology** 10000+ HEALTHY ASIANS **Skin Phenotyping MULTI-COHORTS** AND BODY SITES Meta data analytics PUBERTY, MENOPAUSE, AGEING In vitro models **Developing test screening** platforms (eg:) pre and probiotic agents, small molecules Skin Questionnaires **Clinical and Consumer Endpoints** 



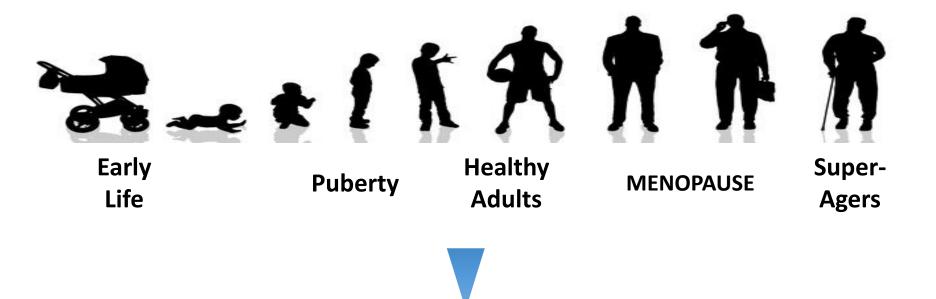
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## **Asian Skin Microbiome Program**

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#### **Global differentiation:**

Almost all current skin microbiome research is parallel group – we have shown longitudinal cohorts are superior.



Unique Singapore Cohorts: characterized, longitudinal, recallable



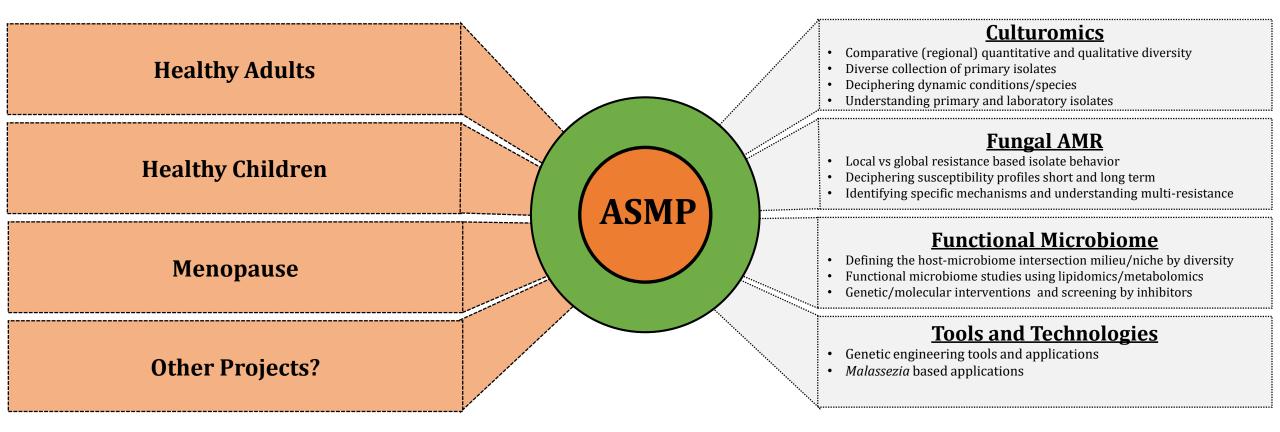
**Asian Skin Microbiome Program** 



Microbiome Program

### **Clinical Collaborations**

### Scientific Focus Areas



#### **Asian Skin Microbiome Program**

#### **Corresponding PIs (Sub-Programme Leads)**



Niranjan Nagarajan, GIS SP1: Tools and Technologies Microbiome genomics, big data analysis pipeline development



Thomas Dawson, SRIS SP2: Functional Skin Microbiome Skin microbiome, yeast biology, drug screening

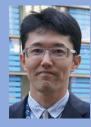
#### **Co-Pls**



John Common, IMB SP3: Skin Microbiome Models Skin biology, skin in vitro models, microbiome in skin disease



John Chambers, NTU Clinical Epidemiology, HELIOS cohort



Kenji Kabashima, SIgN/SRIS Skin immunology, host/microbe interactions, clinical dermatology



Chong Yap Seng, NUH/SICS Puberty and menopausal cohorts



Steven Thng, SRIS/NSC Clinical design and execution, clinical dermatology