

# The Interaction Between Gut Microbiome, Immune System and Allergy Outcomes

Professor Mimi Tang

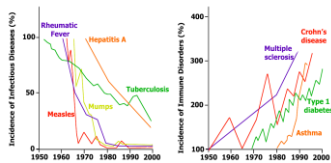
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## Presentation Outline

- DOHaD concept
- Establishment of the gut microbiota
- Gut microbiota and immune development
- Gut microbiota and allergic disease

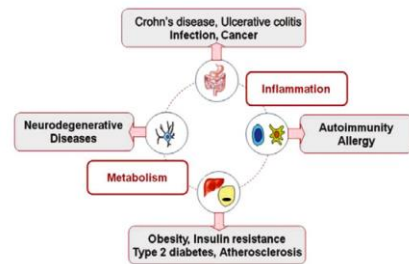
## Hygiene Hypothesis: Allergic and Autoimmune Diseases



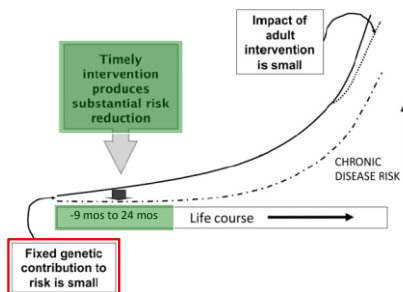
- Increased prevalence of immune mediated disorders correlated with reduced prevalence of infectious diseases
- Strachan first posed the Hygiene hypothesis in 1989
- Importance of overall microbial exposure rather than specific infections
  - Since early/mid 19<sup>th</sup> century... Reduced exposure to microorganisms which we have lived with since the paleolithic age → Concept of evolved adaptation

Strachan DP. BMJ 1989;299:1259-60;  
Bach JF. N Engl J Med 2002;347:911-20;  
Rook GAW. Clin Exp Immunology; 160:70-79.

## Auto-inflammatory Diseases: Common Pathways?



## Early Life Programming of Disease Risk: DOHaD



Hanson and Gluckman AJCN 2011

## Environmental Risk Factors

- Microbial exposures in early life
  - Hygiene hypothesis
  - Intestinal microbiota
- Infant diet
  - Type of food first year of life
  - Timing of exposure to food allergens
- Vitamin D / UV exposure
- Immunomodulatory dietary factors
  - Omega-3 fatty acids
  - Folate?
- Pollutants

## Intestinal Microbiota

The most abundant microbial exposure during life is establishment of the gut microbiota ... major impact on immune responses

100 trillion bacteria  
"Gut Microbiota"

60-70% of immune cells



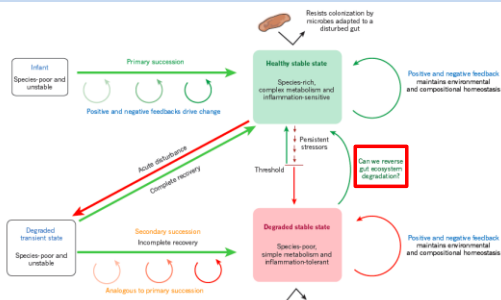
Surface of approximately 300m<sup>2</sup>

100 million neurons

## Establishment of the Intestinal Microbiota

- Gut microbiota evolves rapidly from birth → Relatively stable by 2 years
  - First bacterial communities (seeding bacteria) are acquired from the mother and the environment
  - Diversity increases over the first year and stabilises thereafter
- Microbiota composition is influenced by early life exposures
  - Mode of delivery (cesarean vs vaginal)
  - Maternal microbiota (stress/diet during pregnancy)
  - Breast milk vs formula feeding
  - Diet
  - Sanitation and level of cleanliness
  - Antibiotic therapy
- Stable microbiota is resilient to acute environmental changes
  - Compositional homeostasis

## Transitions in Gut Microbiota Composition



Lozupone et al. Diversity, Stability and Resilience of the Human Gut Microbiota Nature 2012;489:220

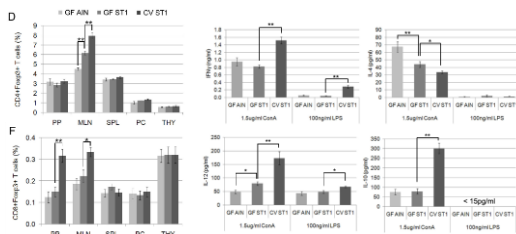
## Intestinal Microbiota and Immune Development

- The intestinal microbiota plays a crucial role in development of the mucosal and systemic immune system
- Mice bred in germ free conditions have abnormal lymphoid tissues<sup>1-4</sup>
  - Small underdeveloped Peyer's Patches, MLNs and spleen
- Mice raised in a germ free environment fail to develop oral tolerance<sup>5</sup>
  - Have persistent and exaggerated Th2 dependent responses to OVA
- These abnormalities of lymphoid tissue structure and oral tolerance can be corrected by seeding the intestine with *Bacteroides fragilis* but.... ONLY if this occurs in the neonatal period<sup>1,5</sup>

- Klaasen HL et al. Infect Immun 1993 61(1): 303-6.
- Umesaki Y et al. Microbiol Immunol 1995 39(8): 555-62.
- Mazmanian SK et al. Cell 2005;122:107-18.
- Hmrcir et al. BMC Immunology 2008, 9:65
- Sudo N et al. J Immunol 1997;159:1739-45.

## Intestinal Microbiota and Immune Regulation

- Germ free mice (that lack intestinal microbiota)<sup>1-3</sup>
  - Reduced numbers of FoxP3+ Treg in MLN +/- PP
  - Treg impaired suppressor function<sup>1,2</sup> and reduced IL10 and TGF $\beta$  production<sup>2</sup>
  - Reduced production of IL12, IFN $\gamma$ , IL10; and increased production of IL4<sup>3</sup>

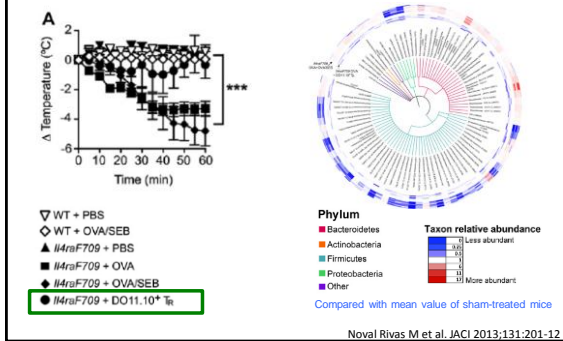


1. Ostman et al. Eur J Immunol 2006; 2. Ishikawa et al. Clin Exp Immunol 2008; 3. Hmrcir et al. BMC Immunology 2008

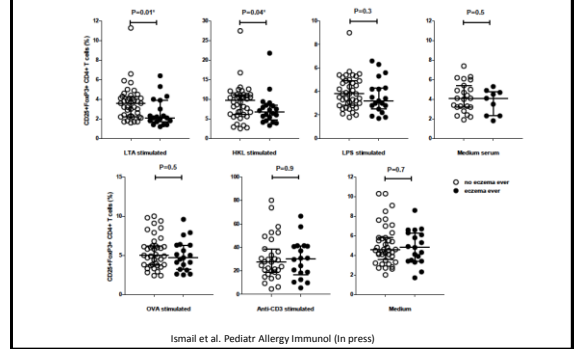
An altered microbiota is associated with increased risk for development of allergic disease



## Transfer of Treg cells can prevent food allergy and induces a tolerogenic microbial signature

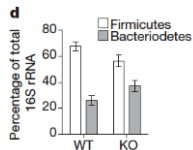


## Infants with eczema have reduced Treg capacity in response to TLR stimulation at birth



## iTreg can Influence Microbiota

- iTreg are important for control of unwanted allergic inflammatory responses at mucosal sites
  - Absence of iTreg in CNS1- mice was associated with intestinal and lung Th2 inflammation
- iTreg help maintain a 'normal' microbial community in the gut
  - Absence of iTreg in CNS1- mice resulted in altered gut microbial communities
  - enrichment of phylum TM7, genus Bacteroidetes Alistipes
  - overall decrease in the ratio of Firmicutes to Bacteroidetes



Percentage of total 16S rRNA gene sequences of the Firmicutes and Bacteroidetes phyla in stool from individually housed CNS1- (n=9) and WT (n=6) littermate mice.

Josefowicz et al. Nature 2012

## Summary

- The rapid rise in non-communicable diseases has occurred as a result of environmental changes associated with the modern lifestyle; in particular, reduced microbial exposures have led to altered intestinal microbial signatures that are associated with allergic disease risk
- The intestinal microbiota plays a critical role in immune development and early life immune programming
- Alterations in the intestinal microbiota in the first weeks of life are associated with development of allergic diseases in later life
- In mouse models, disease risk can be conferred by transfer of disease associated microbial signatures; and conversely, protection against disease can be conferred by favourable microbial signatures
- Infants who develop eczema by 1yr have reduced Treg response to TLR stimulation from the time of birth
- Modulation of the intestinal microbiota in early life may offer a novel approach to prevention or treatment of allergic disease; and may be most beneficial in infants at increased risk of allergic disease