

# CHIM: Opportunities in human immunology?

IABS 2<sup>nd</sup> Human Challenge Trials in Vaccine Development Conference

Beth Kirkpatrick, MD

University of Vermont, Vaccine Testing Center

September 29, 2017



# Overview



All models are wrong, but some are useful.

— *George E. P. Box* —

and design CHIMs  
our knowledge of

ie and

es in the field

**"I think you should be more explicit here in step two."**

CN  
COLLECTION

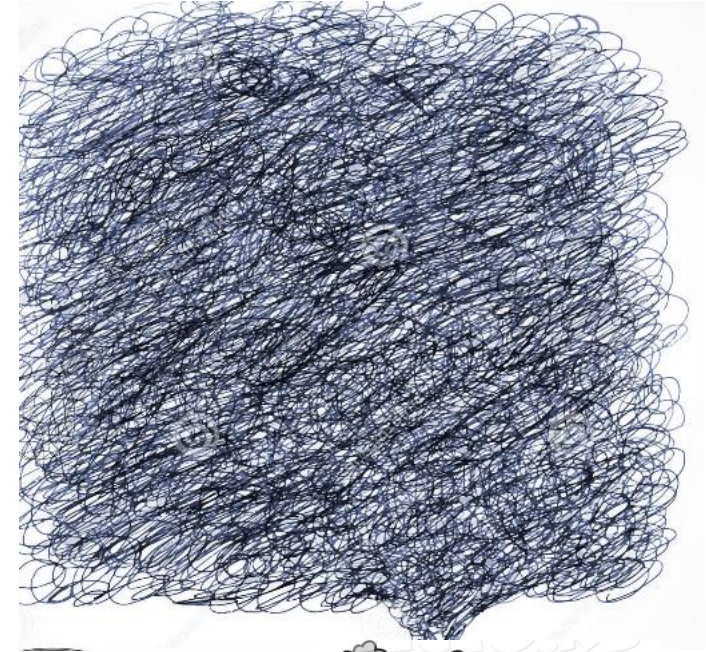
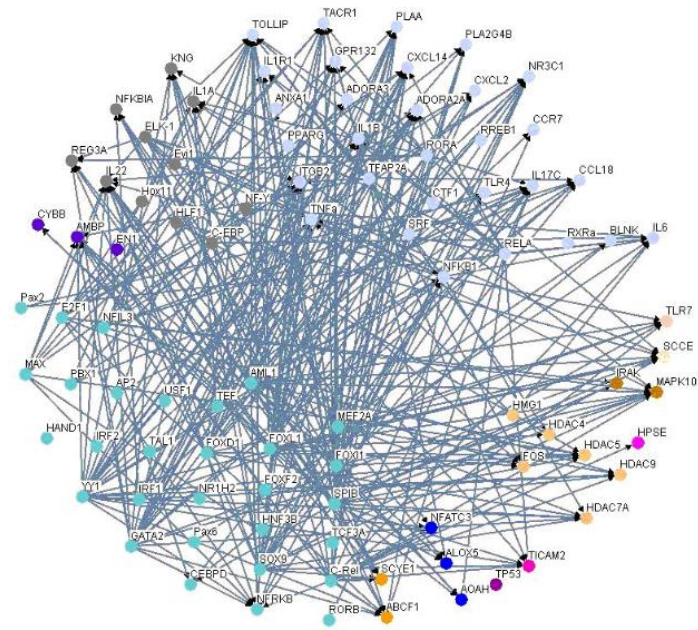
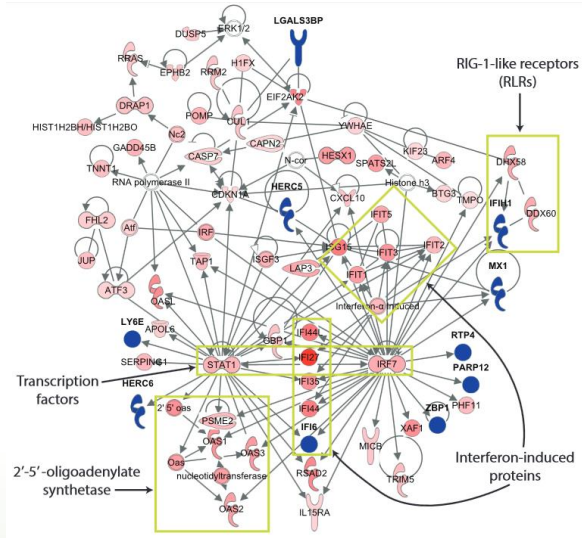
The University of Vermont





Mucosal vs. systemic  
 Infants vs. adults  
 Innate vs. adaptive

Homotypic vs. heterotypic  
 Isolated vs. cross-protective  
 Antibiotic truncated vs. not



Humoral vs. cellular  
 Sterilizing vs. shedding  
 Permanent vs. short lived

Endemic populations vs. naive  
 Co-infections?



# What type of immunology questions might be desired from CHIM?

- Confirmation of epidemiologic or animal models
- Descriptive and comparative immunology
- Functional or mechanistic understanding of immune components
- Antigen discovery
- Identification of monoclonal antibodies for diagnosis or treatment
- Selection and standardization of immunogenicity for future bridging
- Correlates of protection



What type of immunology information  
might be actually learned?  
What might be actually valuable toward  
improving human health?

**IT DEPENDS**

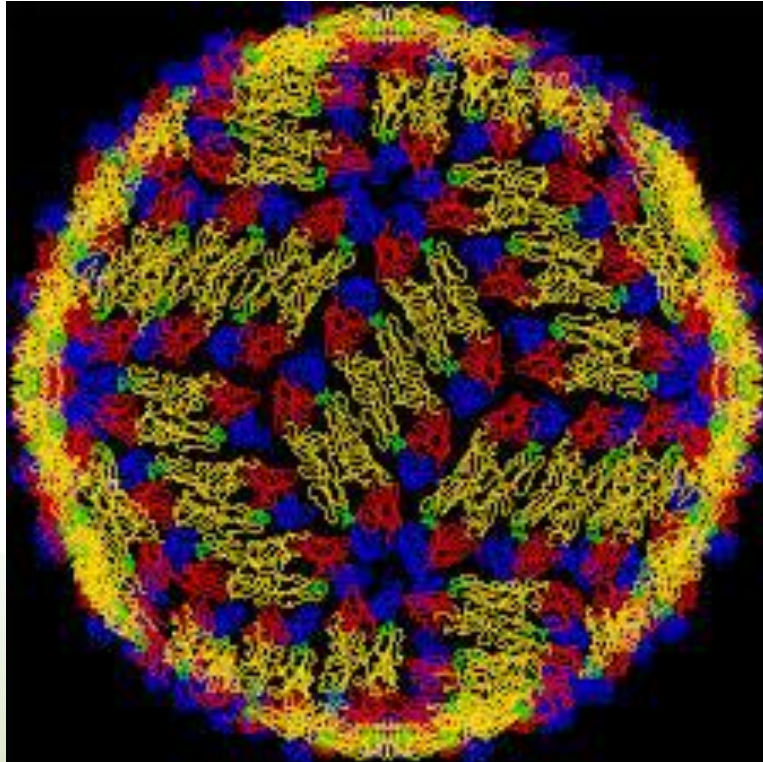


# Where are you starting?

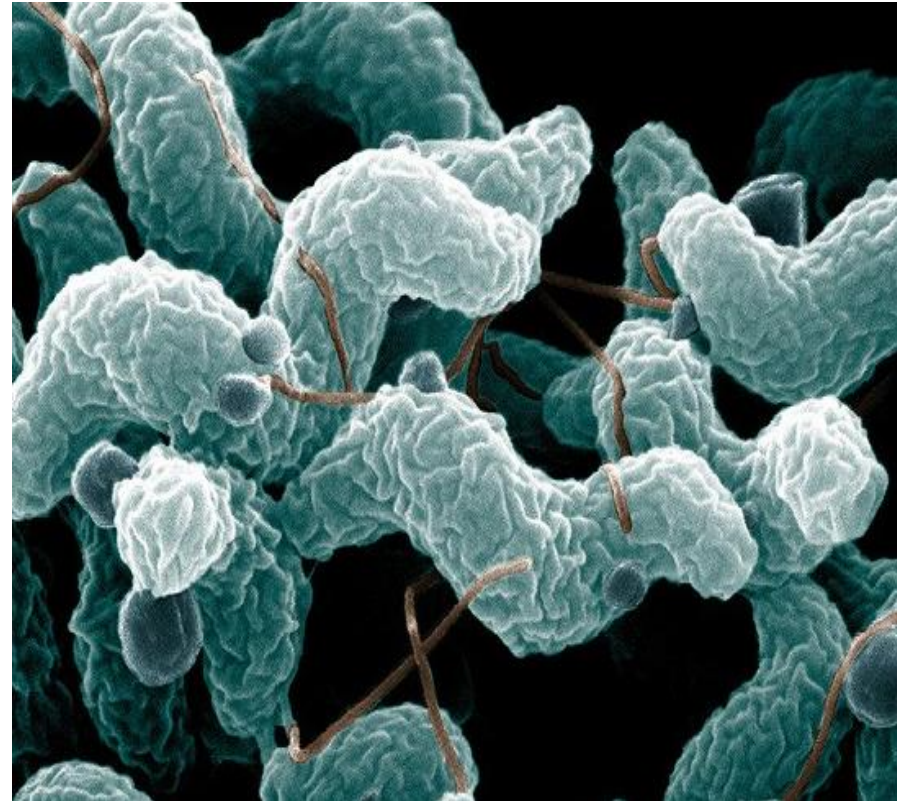
- What is known about the natural history of disease for your pathogen?
- Are robust epidemiologic data available? Populations/ages?
- Are animal models available? Do they reflect human disease?
- Does clinical immunity actually develop? Is it sterilizing/neutralizing? Does it last?
- What immune components are linked with clinical protection? Are immunodominant antigens known? Are there correlates/surrogates?
- Are immune assays with specific antigens established? Validated/Qualified?



# Let's look at two examples:



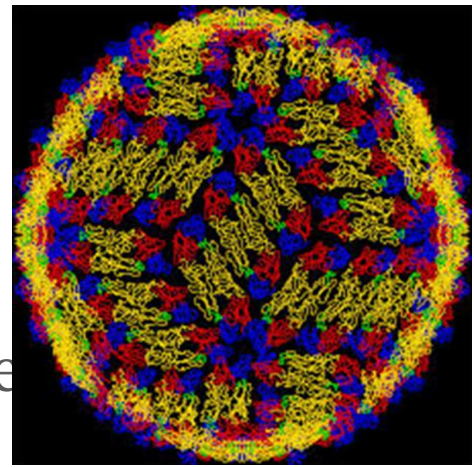
Dengue Viruses



*Campylobacter jejuni*



# Dengue



- Virus and biology well understood. Genetically stable. 4 serotypes
- Viral genetics worked out, manipulable
- Natural history of clinical disease well known, lots of epidemiologic studies
- Good non-human primate model mimics human disease
- Long lasting homotypic protection. Sterilizing immunity
- Correlate of protection is probably neutralizing antibodies
  - Established functional assays (NAB), validated (mostly), cut offs
- Vaccine robust. CHIM robust. Vaccine +CHIM robust.

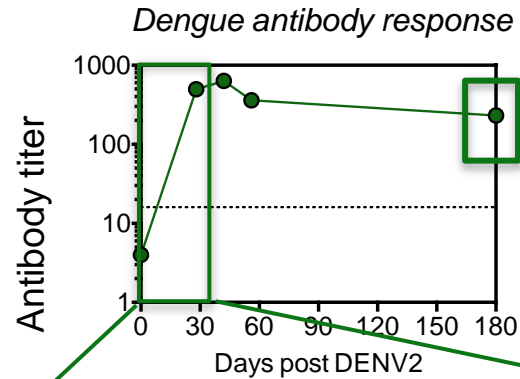
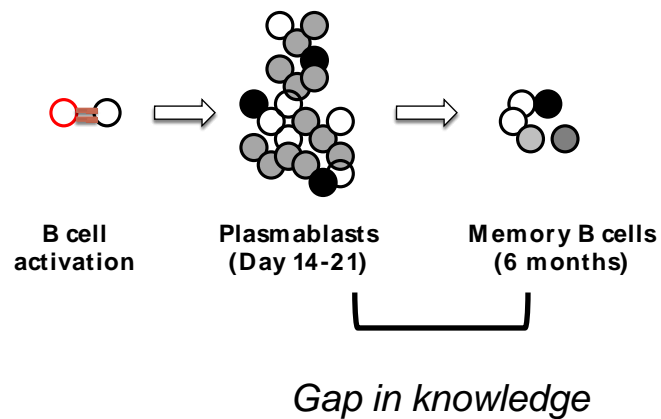


# Immunology advances: Dengue (infection) CHIM

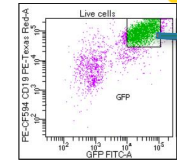
- Advanced descriptive immunology:
  - What is the relative role of CD4 vs. CD8 to structural vs. non-structural antigens?
  - How clonal is the dengue-specific B cell pool and are these antibodies all functional (neutralizing)?
  - Can we characterize memory B cells and their function?
- Identification of monoclonal antibodies with diagnostic or therapeutic potential.
- Study of dengue epitopes
- Confirmation or refinement of immune correlates of protection



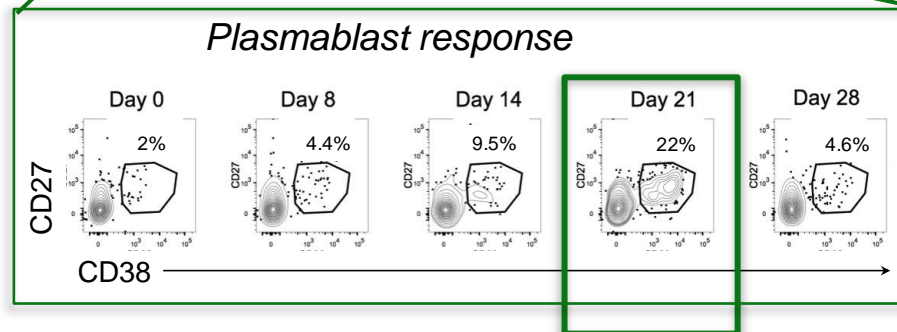
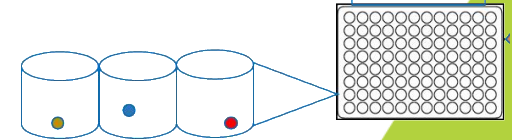
# Probing the B cell response to find new protective measures



High throughput memory B cell sorting



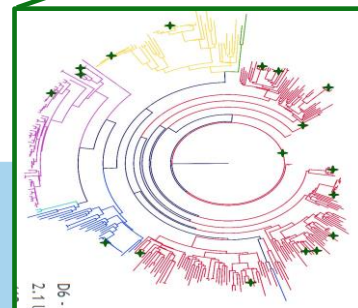
Immortalized MBC (1 cell per well)



Screen antibodies for antigen binding and activity



Next-gen sequencing  
Antibody phylogeny (Atreca)



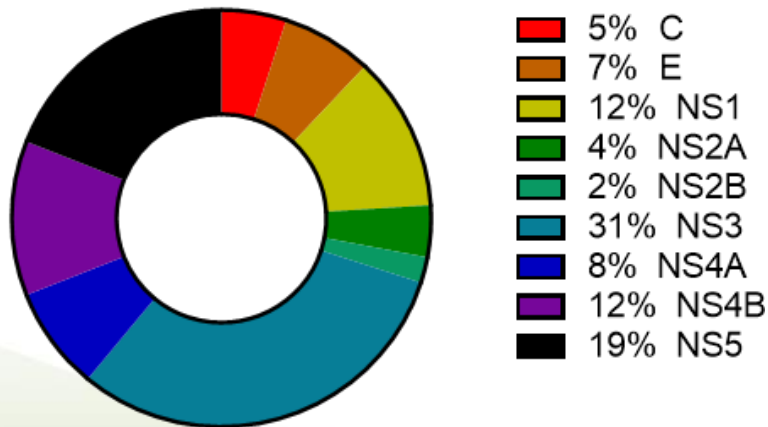
+ Select test Aff Mat Abs based on bioinformatics



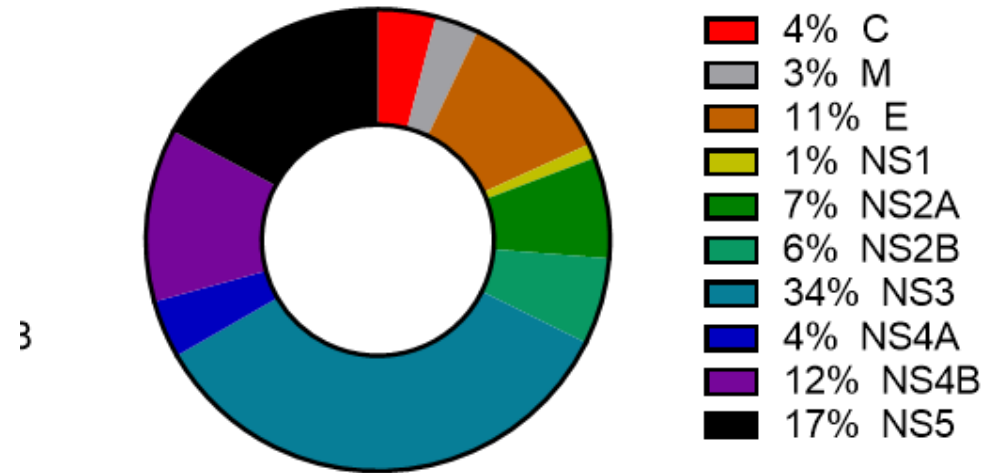
# Cellular responses to the DENV2 challenge strain?

Are the CD8 protein targets the same? Yes

**A Tonga CD8 T cell response**



**B NI DENV2 CD8 T cell response**



3



# DENGUE

- Incredibly rich opportunity to advance natural history “immunology” and compare vaccine to natural disease.
- Can move immune correlates of protection forward (easier if vaccine was less than 100% effective!)
- Opportunities for everyone!
- Four serotypes of dengue
- Difficult to tease out issue of cross-reactive components of immunity
- Vaccine works “too well”!
- “Correlates of Protection”...is a process not a slam dunk.
- Iterative improvements of standardized assays
  - PRNT→microneut (96 well MTP)



# *Campylobacter jejuni*



- Enteric microaerophilic enteric pathogen. Many subtypes. Large amount of capsular phase variation (antigenic changes)
- Clinical spectrum known, epidemiology strong.
- Epi studies show clinical protection (no diarrhea), but no sterilizing immunity. Short lived protection.
- Post-infectious sequelae, molecular mimicry.
- Mechanisms of protection unclear. Components of immune protection assumed from epi and immunocompromised patients.
- Unknown role of the microbiome
- No great animal models
- Human Immunologic assays mostly not functional.

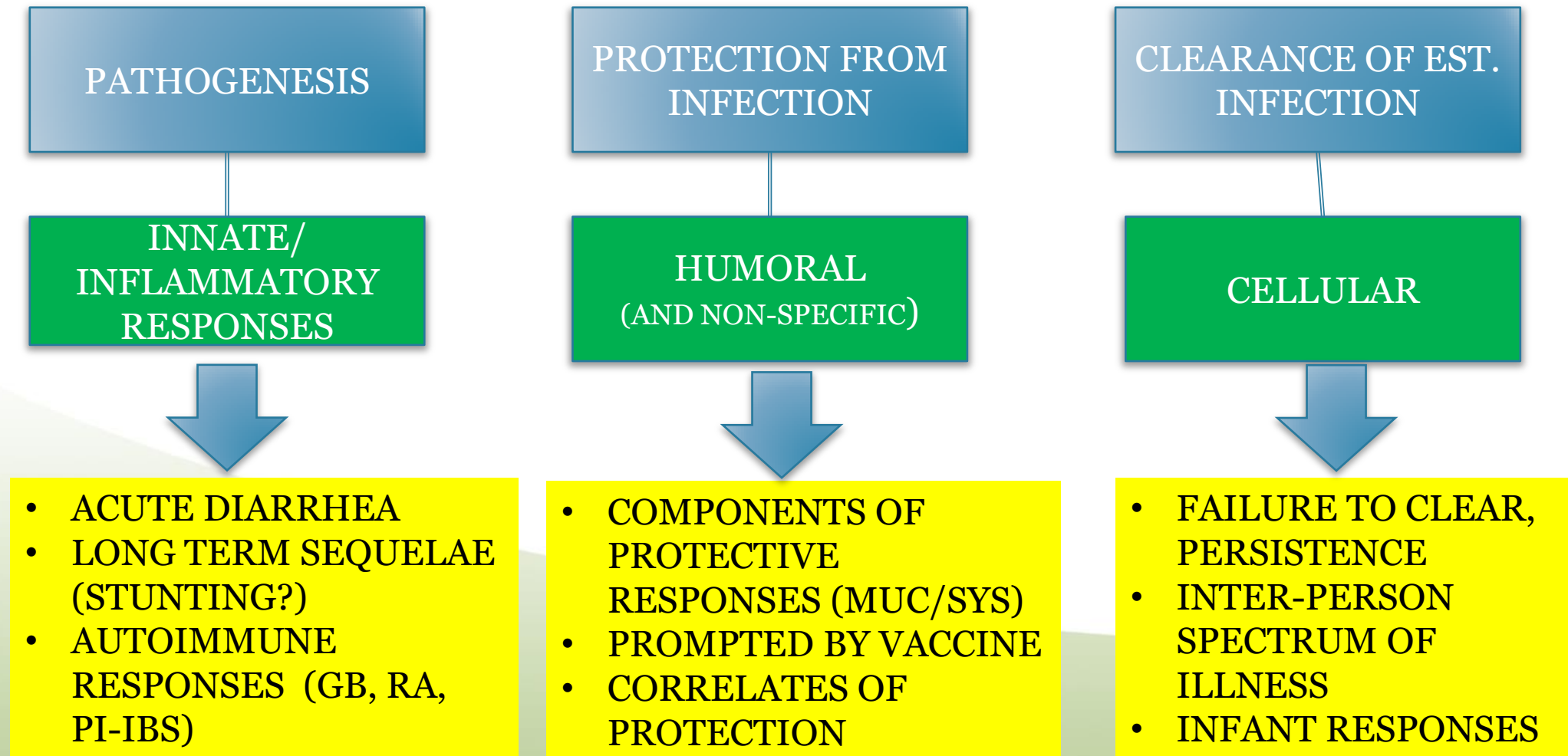


# CHIM: *Campylobacter* immunology

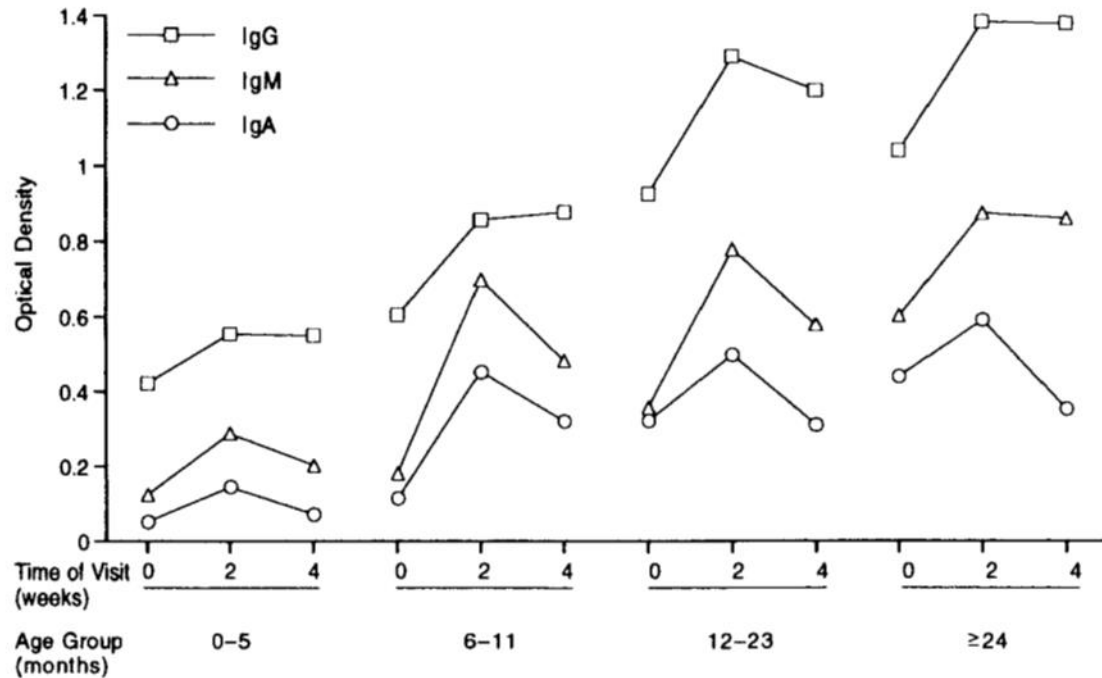
- Comparative immunology vs. epidemiologic observations/natural disease
  - Except issue of antibiotic truncation of full response?
- Descriptive immunity: components of the immune response, effector molecules, involvement of systemic and mucosal compartments.
- Kinetics of immune responses (vs. pre-exposure baselines)
- Confirmation of critical antigens and identification of new antigens
- Understanding composite pattern associated with clinical protection.



# Why do we need to understand $C_j$ human immunology? A simplified view.



# Protective immunity: Cohort Studies (developing world)



In conclusion, in a Thai population in which enteric infections are frequent, *Campylobacter* infections are hyperendemic. As children age, their *Campylobacter* infections become milder, they excrete fewer organisms, and *Campylobacter*-specific serum antibodies rise progressively. All of these findings are consistent with the development of immunity, which suggests that development of a vaccine against *Campylobacter* infections should be possible.

## *Campylobacter* Immunity and Quantitative Excretion Rates in Thai Children

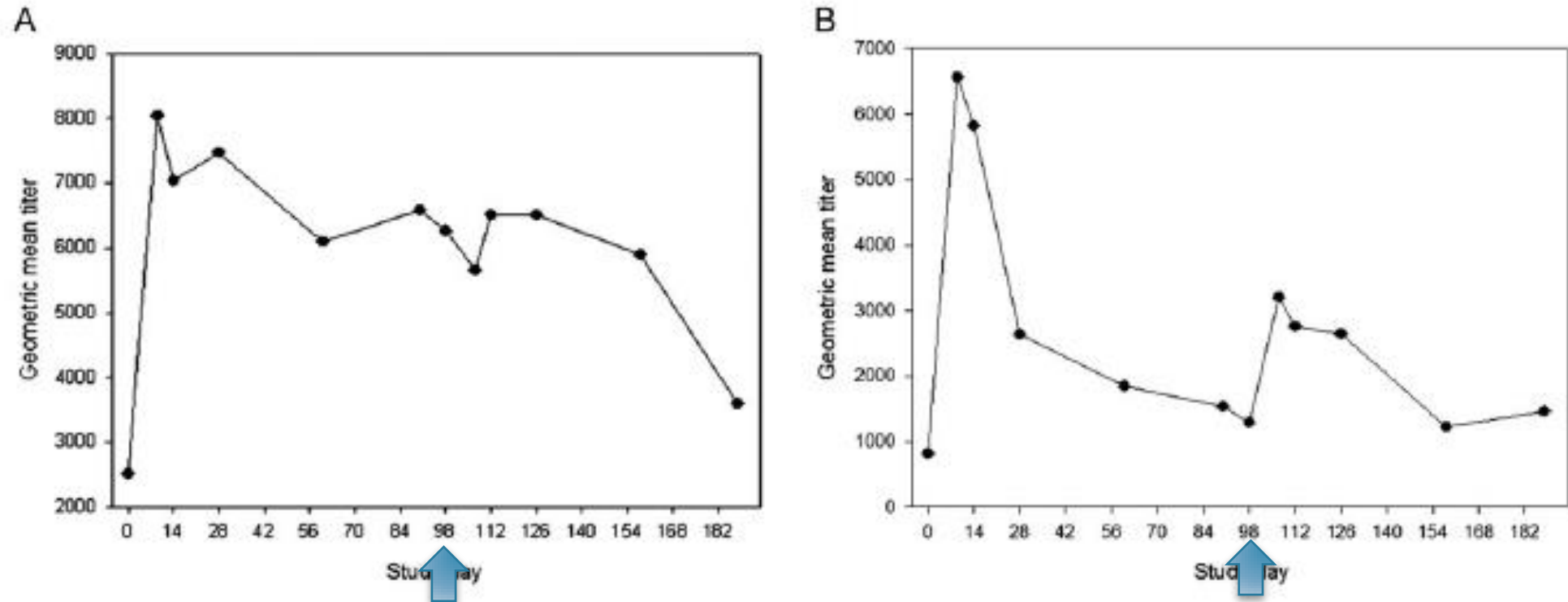
JID 1993;168 (September)

DAVID N. TAYLOR,<sup>1+\*</sup> PETER ECHEVERRIA,<sup>1</sup> CHITTIMA PITARANGSI,<sup>1</sup> JITVIMOL SERIWATANA,<sup>1</sup>  
LADAPORN BODHIDATTA,<sup>1</sup> AND MARTIN J. BLASER<sup>2</sup>

THE UNIVERSITY OF VERMONT



# Immunologic findings w/o protection (CG 8421, 3-month re-challenge)



**Figure 2** Serum immunoglobulin G (IgG), immunoglobulin A (IgA), and fecal IgA responses after challenge (day 0) and secondary challenge (day 98) with *Campylobacter jejuni* CG8421. *A*, Serum IgG responses after primary (day 0) and secondary (day 98) challenge with *C. jejuni* CG8421. *B*, Serum IgA responses after primary (day 0) and secondary (day 98) challenge with *C. jejuni* CG8421. *C*, Fecal IgA responses after primary (day 0) and secondary (day 98) challenge with *C. jejuni* CG8421. Hatched lines represent total fecal IgA. Solid lines represent *C. jejuni* antigen-specific fecal IgA. Abbreviation: IgA, immunoglobulin A.



# Immunogenicity ≠ Immunity

**Table 1. Clinical Outcomes of Subjects After Challenge With *Campylobacter jejuni* CG8421**

Outcome	All Naive	Veteran
No.	15	8
Met primary endpoint of campylobacteriosis, No. (%)	14 (93)	8 (100)
Any diarrhea, No. (%)	14 (93)	8 (100)
Severe diarrhea, No. (%)	8 (53)	7 (88)
Time in hours to campylobacteriosis, median (IQR)	52.3 (46.9–56.1)	53.7 (48.3–73.7)



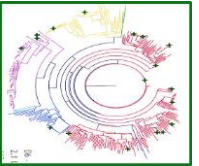
# Next steps for Campylobacter

- Need to understand the role of the microbiome (+/- inflammation, biofilms) in immunologic development.
- Back to the field?-more nuanced study of the development of protective/clinical immunity in children. Including the role of heterologous infections in protection and the “ideal” durability of protection
- Not yet ready for a correlate or surrogate of protection
- Need to understand the function of the immune components.
  - How are those antibodies working (opsonization?)
  - Assay standardization will help with if bridging needed
  - Hypothesis generating



# Overall: Context needed

- Overall, only rare pathogens have comprehensive “natural history” human immunology
  - Most are viruses (e.g. Yellow Fever)
  - The mucosal/gut pathogens are the most difficult (personal bias) due to complexity of the mucosal environment and difficulty evaluating these compartments.
- Trouble with data relevance:
  - Immunology assays are often not standardized or harmonized, making it difficult to build on historical data.
  - Data analysis plans for immunology require attention to correction for multiple variable.
  - CHIM power calculation may be insufficient for inter-personal ranges of immunologic parameters.
- Even without infection, the understanding of normal immune parameters is in its infancy
  - <http://10kimmunomes.org/>
  - Ten thousand immune gnomes?



# The 10,000 Immunomes

The 10,000 Immunomes Project is a reference dataset for human immunology, derived from over 10,000 control subjects in the NIAID ImmPort Database .

Available data include flow cytometry, CyTOF, multiplex ELISA, gene expression, HAI titers, clinical lab tests, HLA type, and others. Check out the [Visualize](#) page to view the available data. Visit the [Download](#) page to download a subset of the data or the entire dataset for your own use.

Select Data Type:

Flow Cytometry: PBMC

Select Analyte:

Effector\_memory\_CD4\_T\_cells

Normalized

Age Range:



Sex:

Female

Male

Ethnicities:

White

Black or African American

Asian

Other

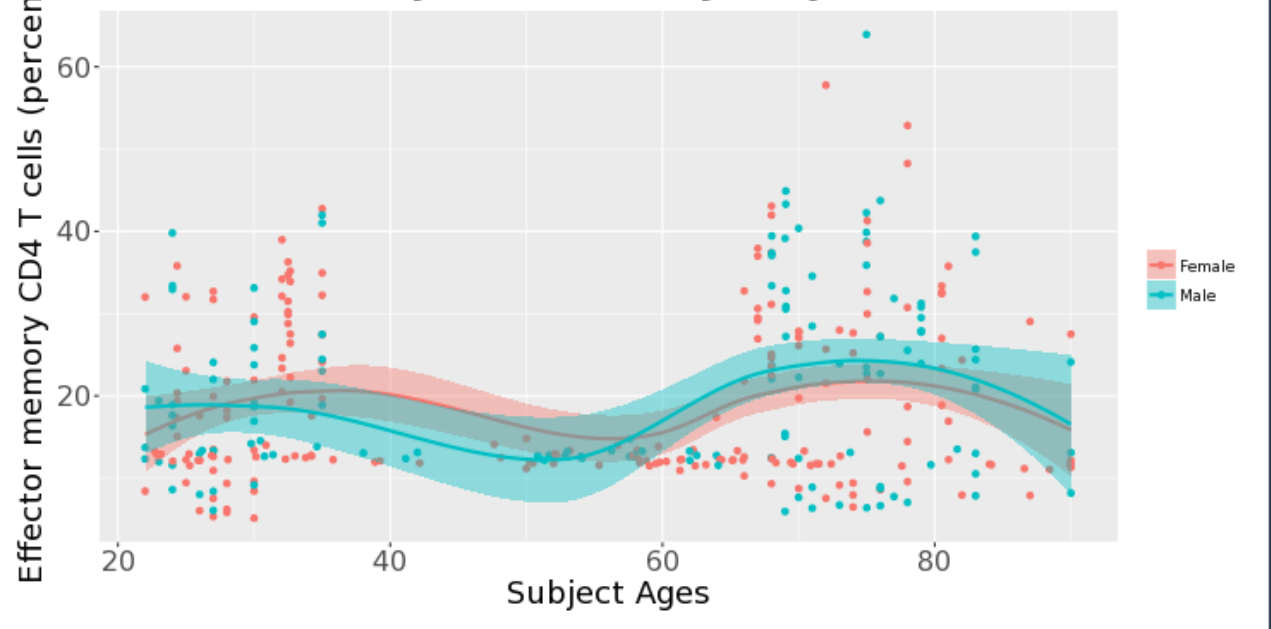
Plot By:

Age & Sex

Ethnicity

Study

### Effector memory CD4 T cells by Subject



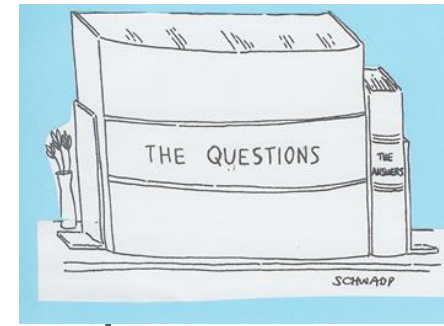
Download Plot

Download Data

Distinct Subjects: 330



# Conclusion



- CHIMS offer unparalleled opportunities to advance human immunology,
  - Especially if re-challenges are done.
  - We shouldn't be surprised if the data makes our life more difficult.
- We might consider as a group what immunologic data is needed from natural disease populations before CHIM is started.
- Minimal standardization of assays would permit use of historical controls.
- Functional immune assays
- Statistics, modeling, bio-informatics will be more important than ever and analytic packages must meet these challenges.

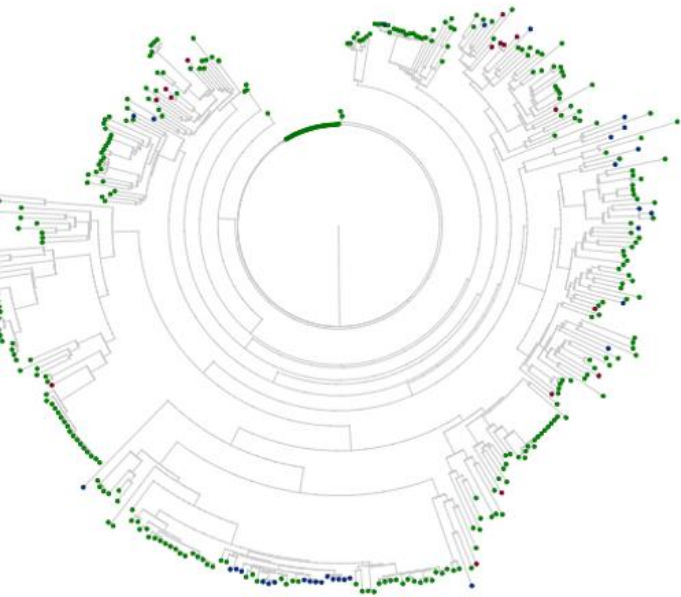


# Thank you





# Antibody Immune Repertoire



IgG Isotype

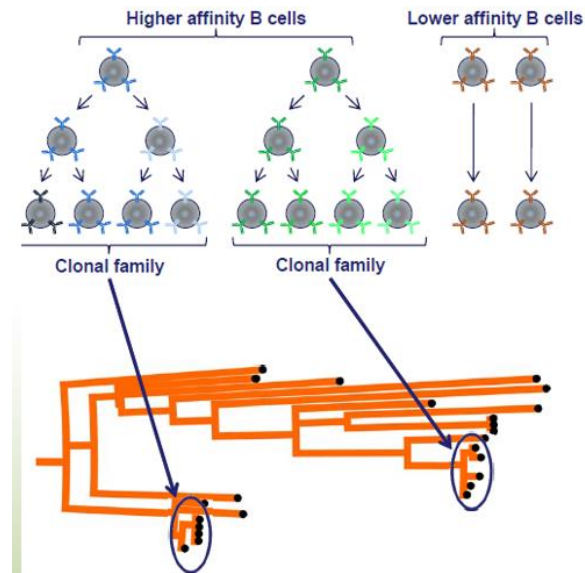
- G1
- G2
- G3

- Viremic D6 – D7 post-challenge
- 2.1 log<sub>10</sub> PFU/ml peak titer
- 412 pairs



Courtesy of Daniel Emerling, Atreca

## CLONAL FAMILY ANALYSIS IDENTIFIES AFFINITY MATURED ANTIBODIES



ATRECA, INC.

Affinity maturation of B cells in germinal centers leads to clonal expansion

### Clonal families

- Contain antibodies with shared HC V(D)J and LC VJ
- Identify high-affinity and functional antibodies

