Exploration of the microbiota in inflammatory diseases

Matthew Stoll MD Research Computing Day September 13, 2012

We're surrounded by bugs

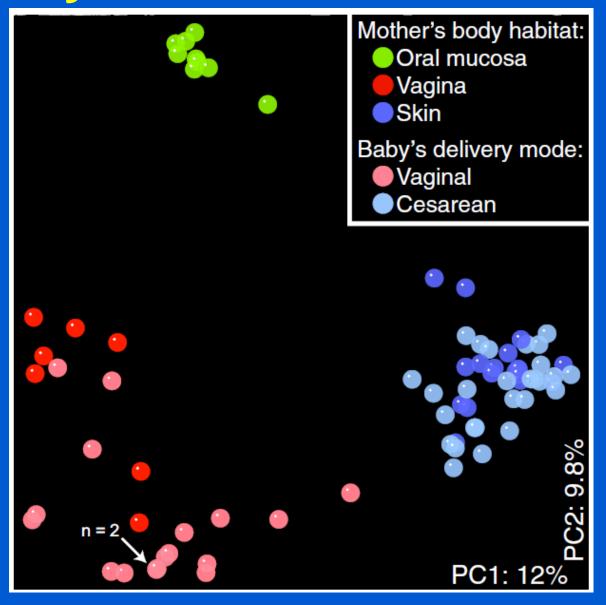
- Human body contains 100 trillion microbes
 - Out-number human cells 10:1
- Human gut alone:
 - -~ 99% of all of the microbiota
 - 3 pounds of bacteria
 - 50% of fecal volume
 - ~ 1000 different species
 - 3 million different genes (100x human host)

Scher, *Nat Rev Rheumatol* 2011;7:569 De Cruz, *Inflamm Bowel Dis* 2012;18:372

Influences of microbiota composition

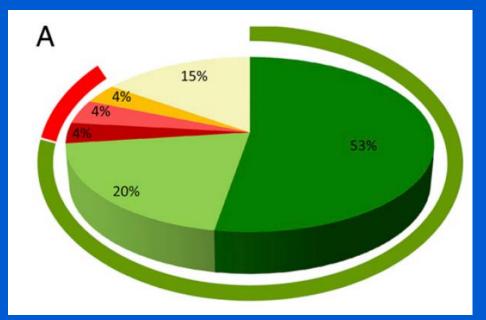
- Infant specific factors
 - Mode of delivery
 - Food source (breast vs bottle)
 - Maternal composition
- Diet
- Geography
- Antibiotic use

Delivery mode and neonatal flora



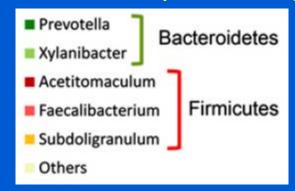
Domingues-Bello *PNAS* 2010;107:11971

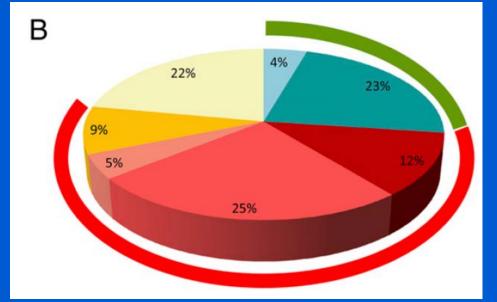
Microbiota and diet / geography



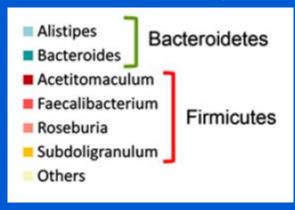
Children age 1 – 6 years

Burkina Faso (rural Africa)



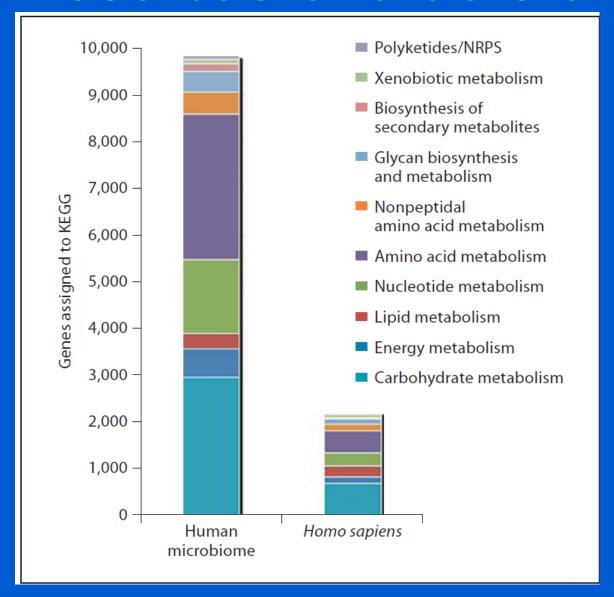


Florence, Italy



De Filippo, PNAS 2010;107:14691

These bacteria do stuff



Important metabolic functions

- Sugar / starches
- Drug metabolism
- Vitamin synthesis
- Bile acid deconjugation
 - Affects on triglycerides, cholesterol

Microflora appear to have a profound impact on a variety of diseases

Immunology and gut bacteria

- Mice raised in GF conditions show:
 - Defective T, B, and innate cells in mucosal lymphoid tissue
 - Smaller mucosal LNs
 - Fewer CD4 cells in peripheral LN
 - Th2 bias, with few Th17 cells
 - Decreased IgG and IgA levels

Hygeine hypothesis

- Recent increases in
 - Inflammatory bowel disease
 - Atopy (allergies, asthma)
 - Type I diabetes
- Attributed to cleaner environments

Germ-free humans, cont.

- Temporal: pre / post Industrial Revolution
- IBD, atopy risk increase with income
 - Across continents
 - Within countries
- Other markers of infectious burden
 - Day care attendance
 - Family size (increased # of sibs is protective)

Bach, Cell Immunol 2005;233:158
Blanchard, Am J Epidemiol 2001;154:328
Werner, Br Jrnl Dermatol 2002;147:95

Flora and animal models of disease

- Germ-free state results in altered / absent disease expression in several models of autoimmune diseases.
 - IBD
 - Spondyloarthritis with colitis
 - Rheumatoid arthritis
 - Type I Diabetes
 - Multiple sclerosis



Disease transfer: TRUC model

- Colitis develops in double k/o mice:
 - -RAG2
 - T-bet (transcription factor)
- Role for flora evidenced by
 - Response to broad-spectrum abx
 - Absence of disease in progeny of abx-Tx mice
 - Development of colitis in wt mice housed with TRUC mice



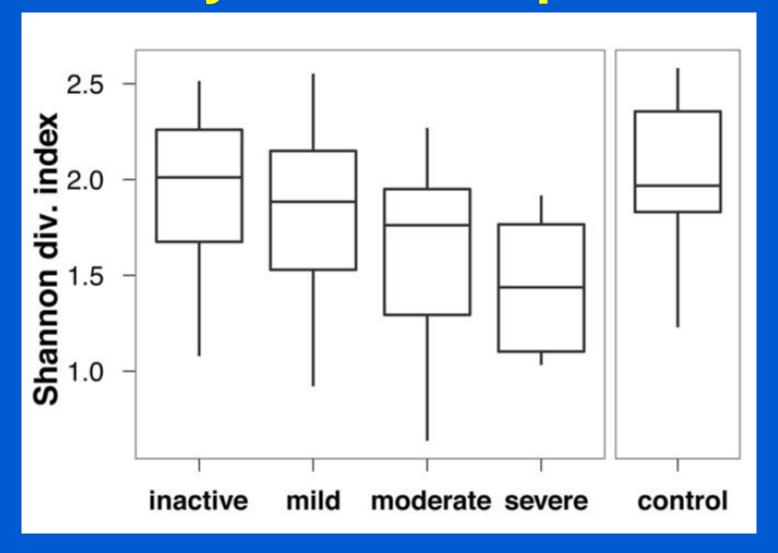
Microbiota and inflammatory diseases

- Inflammatory bowel disease (IBD)
- Insulin-dependent (Type I) diabetes
- Arthritis?

Microbiota and IBD: indirect evidence

- Altered flora in multiple studies
- Decreased diversity

Diversity of flora in peds IBD



Microbiota and IBD: therapeutic response to alterations in flora

- Antibiotics routinely used
- Probiotics may be effective
- Fecal microbiota transplantation
 - Used extensively in chronic C. diff
 - Case reports in IBD
- Gnotobiotic facility at UAB can permit further study of microbiota transfer

Microbiota and Type I DM

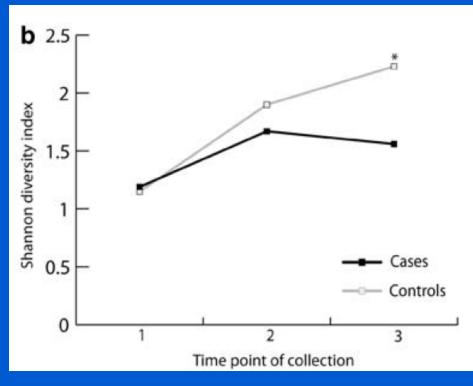
- Diabetes Prediction and Prevention Study
- Serial stool collections on high-risk infants
- Case defined as pos. autoantibodies
 - Ultimately dx with diabetes
- Matched controls

Microbiota changes precede onset of Type I diabetes

Increase in Bacteroides

50 ┌ Sacteroides-Firmicutes (% of total reads) 😢 40 30 -20 -10 Controls Time point of collection

Decreased diversity



Onset

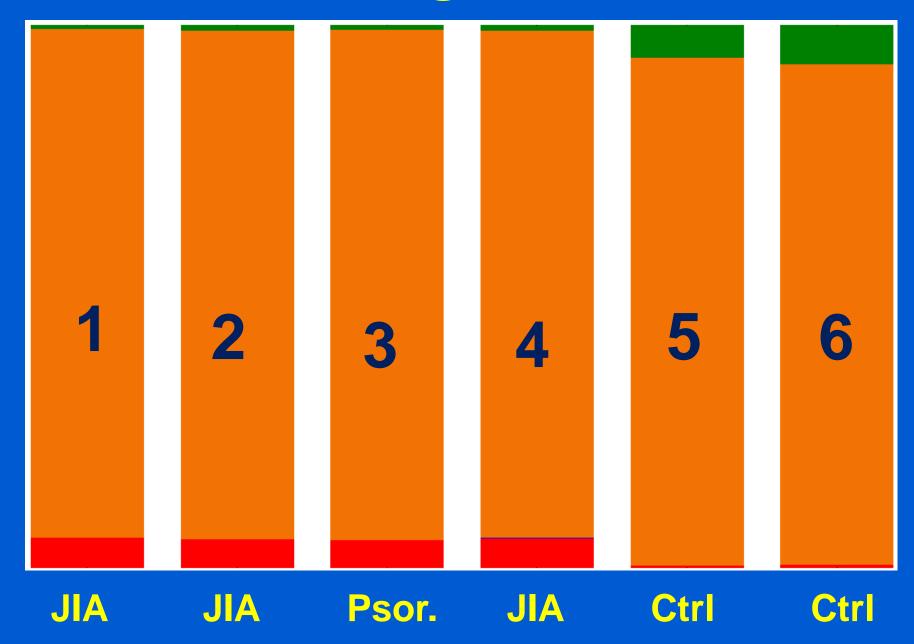
Onset

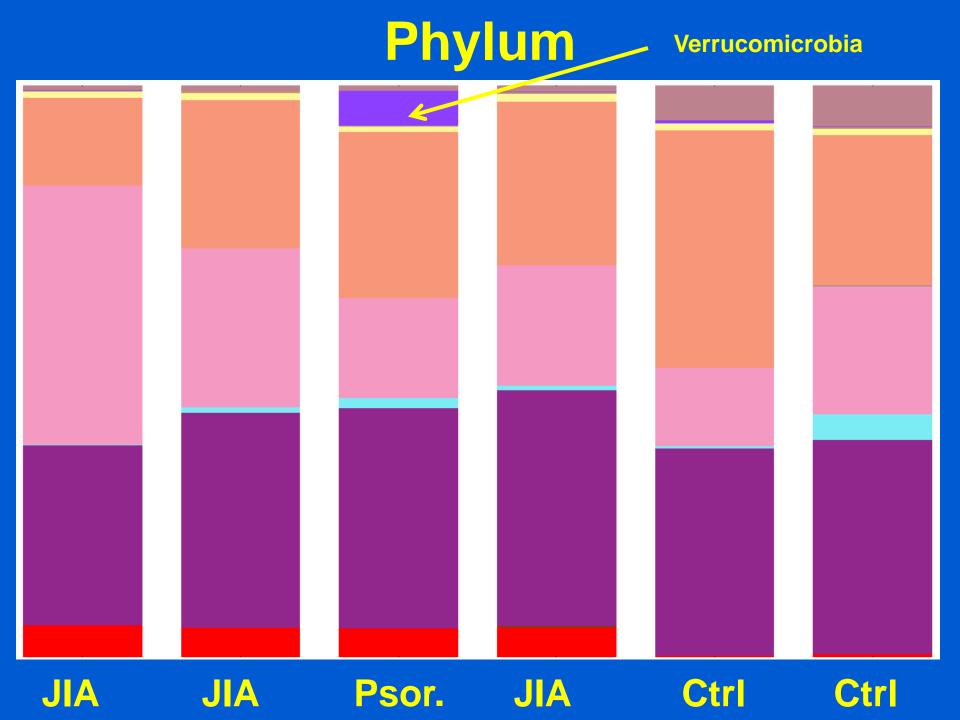
Juvenile arthritis: preliminary study

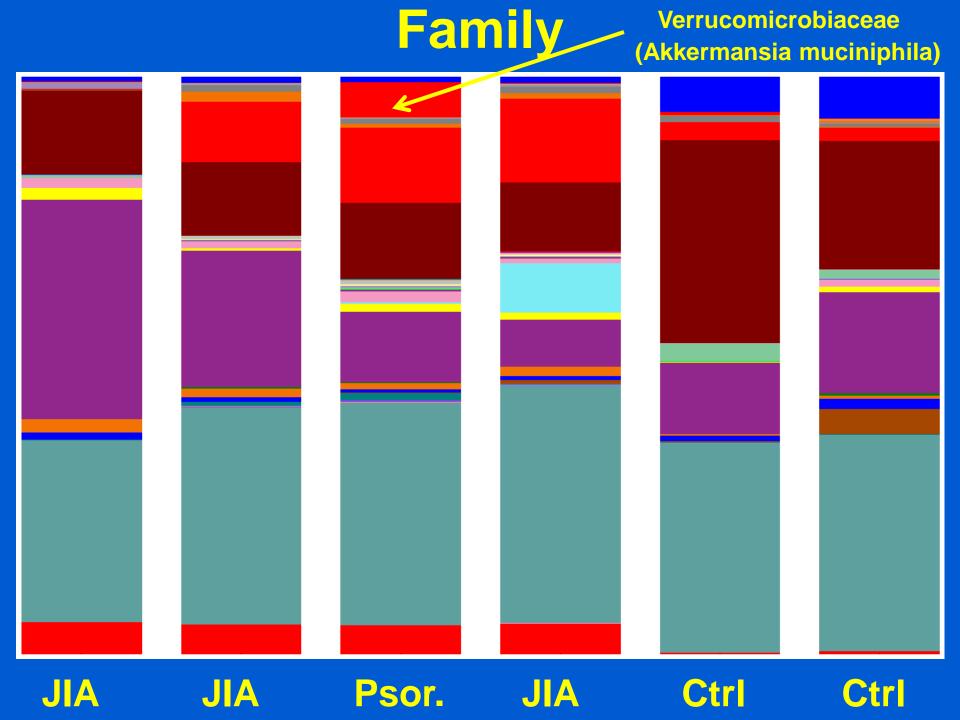
- Stool collected from
 - Three children with juvenile arthritis
 - One with psoriasis + joint pain
 - Two adult controls

16S sequencing performed

Kingdom







Start of a story...

- 6 yo female with psoriasis and joint pain
- Subsequently developed abdominal pain
 - Diagnosed with IBD
- A. muciniphila chews up protective mucin layer in intestines



Microbiota may impact noninflammatory diseases

- Obesity
- Metabolic syndrome

Obesity

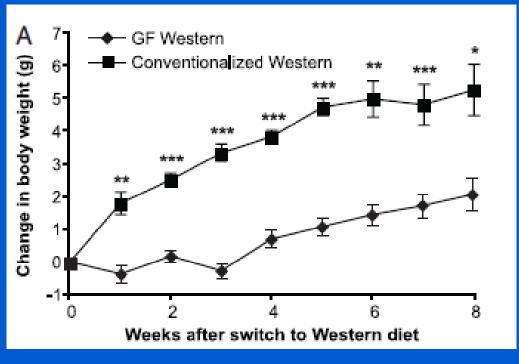
- Differences in obese vs lean microbiome
 - Not universally detected
- Even if so, cause or effect?
 - Weight loss results in altered microbiome
 - Bacteria adapt to our diet
 - Vegetarians enriched for bugs that can metabolize monosaccharides and disaccharides

However, in mice...

Germ-free mice showed less weight gain on

Western diet

Backhed, *PNAS* 2007;104:979



 Transplantation of microbiota from leptindeficient mice resulted in increased body fat compared to recipients of microbiota from wt

mice Turnbaugh, *Nature* 2006;444:1027

And in humans...

- Infant abx use is associated with increased BMI during childhood¹
- Infant microbiota is associated with future weight gain:
 - B fragilis correlates with BMI at age 3²
 - Low bidifobacter predicts overweight at age 7³
- Upregulation in obese subjects of genes involved in phosphotransferase, which is involved in putting sugars into cells⁴

¹Trasande, Int Jrnl Obesity 2012;ePub ²Vael, *Gut Pathogens* 2011;3:8 ³Kalliomaki, *Am J Clin Nutr* 2008;87:534 ⁴Greenblum, *PNAS* 2012;109:594

Moreover...

- 18 adults with metabolic syndrome underwent small bowel lavage, with either:
 - Allogenic microbiota transfer (lean donors)
 - Autologous microbiota transfer
- Results
 - Increased microbial diversity
 - Increased insulin sensitivity
 - Weight changes not reported

Summary

- New technology enables collection of previously unattainable amounts of data
- Analysis would not be possible without high-power computing platforms discussed throughout today
- Areas of research include:
 - Ontogeny of microflora, factors that influence it
 - Effect of flora on inflammatory diseases
 - How to alter the flora to improve human health