

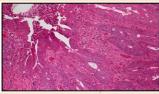
Jonathan Braun, MD, PhD

David Geffen School of Medicine at UCLA CCFA National Scientific Advisory Committee

Crohn's Disease

- · Chronic disease of the intestines
 - Sores (ulceration), perforation, scaring, strictures
 - All regions of intestine (especially junction of small and large intestine)
 - Abdominal pain, diarrhea, bleeding, malabsorption, abdominal infection, elevated risk of cancer







Normal

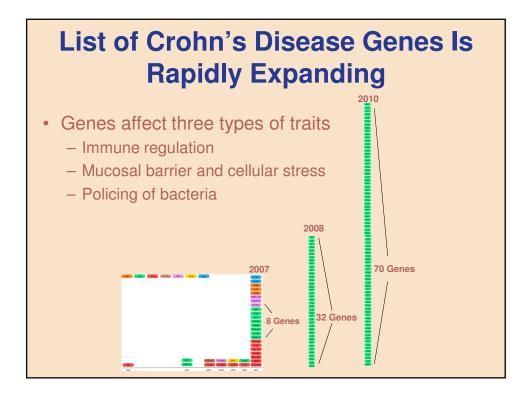
Active Disease

Treatment

Crohn's Disease

- Peak onset in teens
 - All ages affected
 - Growth and development problems in children
- · Immune-mediated
- Family (genetic) susceptibility
- Environment affects disease risk



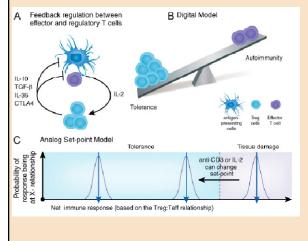


What Do CD Genes Teach Us?

- There will be many genes when the list is complete
 - Estimated > 200
 - Single patients may have only $\sim\!5\!-\!10$
 - By good fortune, unaffected siblings have slightly fewer
 - Significance: "fixing" only a few genes may be enough



Immune Regulation: Hormones Controlling the Balance of Inflammation



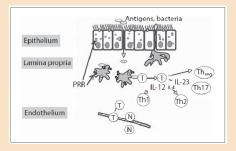
- Animal research discovers immune hormones that control colitis risk
- Examples
 - IL10 quiets inflammation
 - IL23 drives inflammation

Early Onset (<1 y/o) Aggressive Crohn's Disease Due to a Rare Mutation in the IL-10 Hormone Receptor



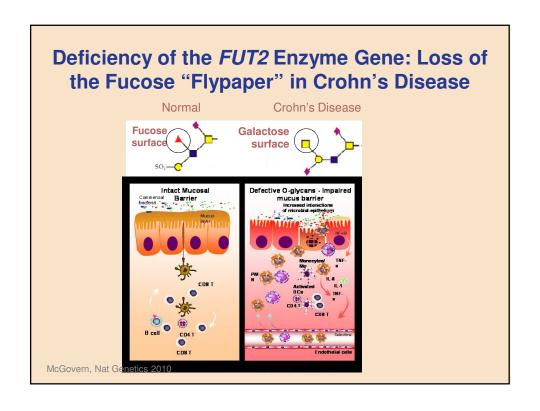
- Team Leader, Dr. Scott Snapper
 - CCFA Research Initiatives Chair
 - Glocker EO. N Engl J Med, 2009
- · Treatment implications
 - IL10 hormone won't correct
 - Stem cell replacement gave complete remission
 - Future: identify alternative hormone

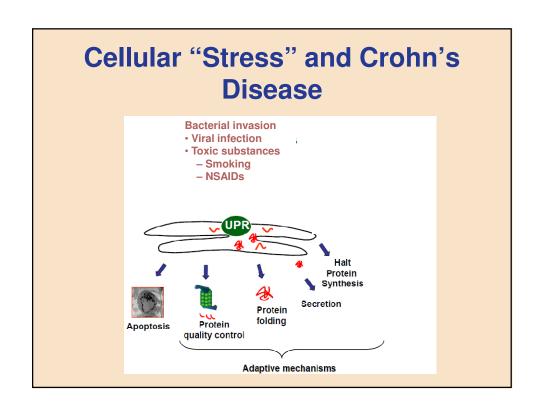
Targeting the IL-12/IL-23 Pathway in Crohn's Disease

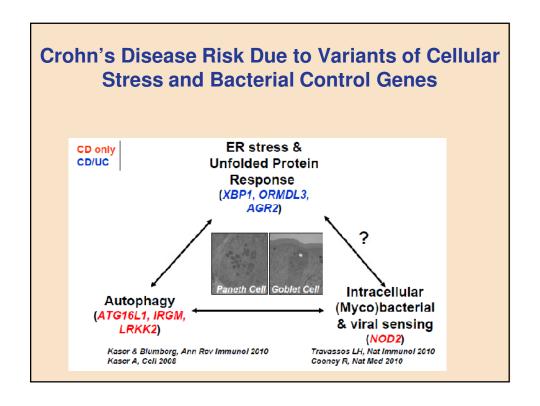


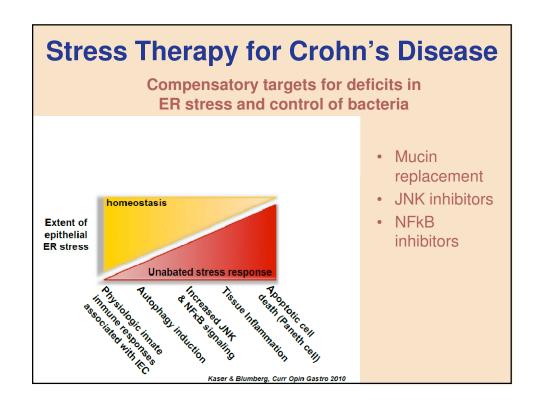


- Human genetics
 - An <u>overactive</u> IL23 receptor gene variant is present in <u>90%</u> of Crohn's patients
- Strategy: Block the IL23 receptor
 - IL12 and IL23 receptors both can be targeted via shared p40
- First success: phase 2 clinical trial (Mannon et al. N Engl J Med. 2004)
- · Ustekinumab phase 3 trials underway









CCFA Genetics Initiative

- First phase (inception, 2000)
 - Creation of first international team
 - DNA bank from patients
 - Discovery of original IBD genes
- Second phase (inception, 2011)
 - Create a gene testing toolkit for patients and doctors
 - Find the clinically most important genes:
 - · Affect response to treatment
 - · Determine disease severity
 - Identify genes suitable for treatment strategies

80 Agents in the Clinical Trial Pipeline

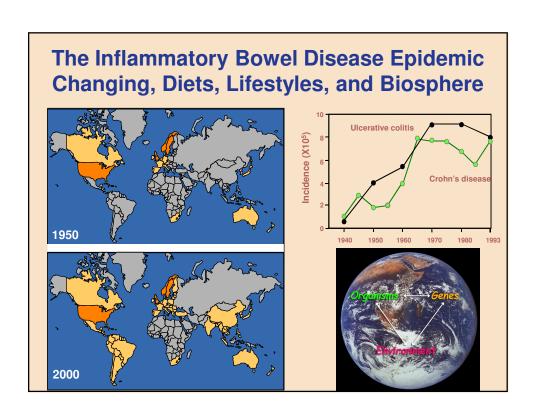
- Homing blockers
 - Natalizumab (approved, 2008)
 - Vedolizumab (phase III, 2009)
- IL12 and 23 blockers
 - Ustekinumab (phase III, 2009)
- Adult mesenchymal stem cells
 - Control inflammation, promote tissue repair, prevent scar formation
 - Prochymal (phase III, 2009)
- Combination of TNF blockers and methotrexate
 - More frequent response and better maintenance for fistulizing Crohn's
 - Concern: infection and cancer risk
 - CCFA Clinical Alliance trial to clarify best patients for combination therapy

PIANO

Pregnancy in Inflammatory Bowel Disease And Neonatal Outcomes

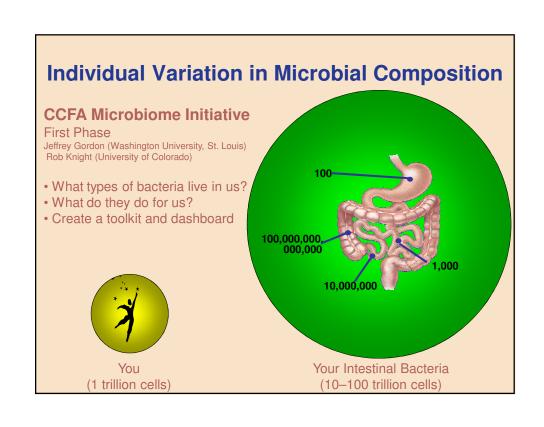
- CCFA-initiated clinical study
 - Leader: Uma Mahedevan, UCSF
- 413 patients divided into 4 groups
 - No immunosuppression; AZA/6MP; Biologics; Combination
- · Medication use not associated with increased risk of:
 - Any complication
 - Preterm birth, low birth weight
 - Cesarean section
 - Congenital anomalies: 17 anomalies/15 births
- Biologics: increased risk of NICU stay
- · Combination: increased risk of infection at 1 year of age

Join the registry: www.ccfa.org/trials (Search: PIANO)

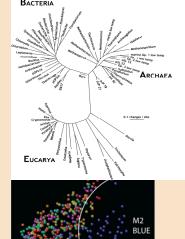


Injurious and Protective Bacteria at the Intestinal Surface Associated With Crohn's Disease Bad bacteria candidates Elevated in patients and in flares Products damage intestine Adherent/invasive E. coli Segmented filamentous bacteria Lachnospiraceae (CBir) SFB bacteria

Gut surface



CCFA Microbiome Initiative (First Phase)

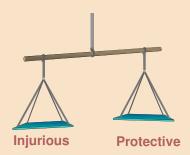


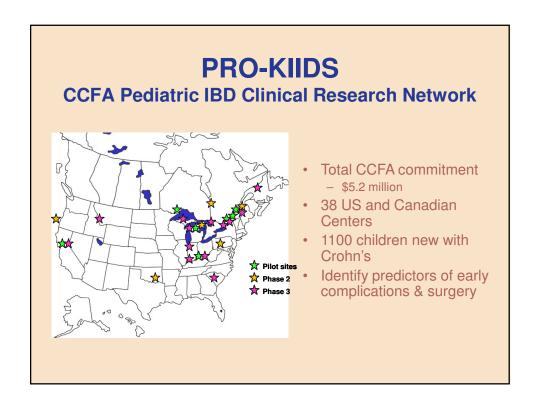
- 200 species per person
- Little species overlap between people
- Mother effect: species are shared by siblings
- However, a mosaic of functions are shared between people
- Toolkit: QIIME online (http://qiime.sourceforge.net)

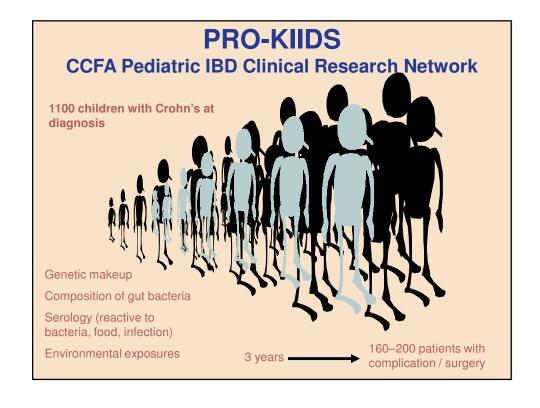
CCFA Microbiome Initiative (Second Phase)

- Determine full bacterial composition in individual IBD patients
- Alterations in bacterial functions in individual IBD patients
- Effect of IBD-related genes on intestinal bacteria
- Test strategies to alter intestinal bacteria
- Effect of dietary manipulations on bacterial microbial composition
- Creating a dashboard for patients to monitor and adjust their bacteria

- Selectively alter the balance
 - Diet and prebiotics
 - Probiotics
 - Antibiotics
 - Engineered bacteria (IL10, KGF2)







CCFA Partners

- A new program to enlist patient and family participation in activities to further increase our understanding of Inflammatory Bowel Diseases (IBD)
- A long-term patient registry to participate in IBD research
- Broaden participation and inform registry members of upcoming studies and trials
- Increased patient involvement for more rapid research progress by shortening the time required to compile sufficient research data to complete the project(s)
- The investigators of CCFA Partners are:
 - Lloyd Mayer, MD (Chair, National Scientific Advisory Committee)
 - Bruce Sands, MD, MS (Chair, Clinical Research Alliance)
 - James D. Lewis, MD, MSCE (Vice-Chair, Clinical Research Alliance)
 - Sunanda Kane, MD (Chair, Patient Education Committee)
- Contact
 - info@ccfa.org

Questions & Answers