

EFFECT OF EARLY-LIFE UNDERNUTRITION ON THE GUT MICROBIOTA

Geoffrey A. Preidis, M.D., Ph.D.

Assistant Professor of Pediatrics

Section of Gastroenterology, Hepatology, and Nutrition

Department of Pediatrics

Baylor College of Medicine

Texas Children's Hospital

THE #1 HEALTH PROBLEM PLAGUING CHILDREN TODAY



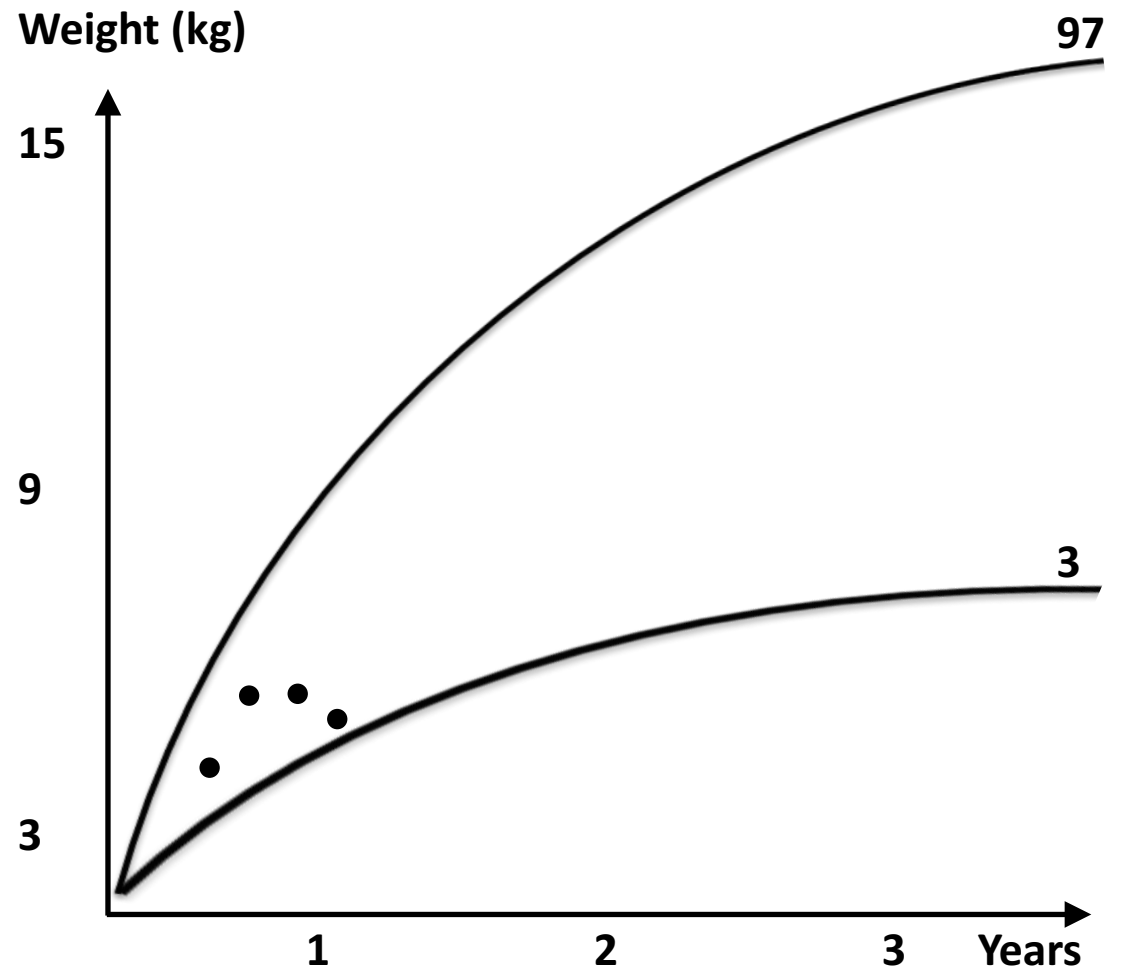
- Globally, undernutrition contributes to **3.1 million child deaths per year**.
- 96 million children (14%) are underweight and 159 million (24%) are stunted.

A “VICIOUS CYCLE” OF GI PATHOPHYSIOLOGIES



- Enteric dysfunction, malabsorption
→ increased fermentation substrates
- Low-protein diet, hypoalbuminemia
→ ascites
- Liver function anomalies
→ steatosis, hepatomegaly, ascites
→ decreased bile acid synthesis,
impaired weight gain
→ coagulopathy
- Gastrointestinal dysmotility
→ luminal stasis, fecal impaction,
poor appetite
- **Infection or microbial “dysbiosis”**
→ **inflammation, gas, bloating**
- “Thrifty phenotype” → increased risk of
obesity and metabolic diseases

CASE PRESENTATION



READY-TO-USE THERAPEUTIC FOOD (RUTF)

- Nut paste, sugar, vegetable oil, milk powder, vitamins & minerals
- Can be expensive, often must be imported
- Long-term outcomes with respect to child growth have yielded mixed results in meta-analyses

Ashworth. *Food Nutr Bull* 2006;27:S24-48.

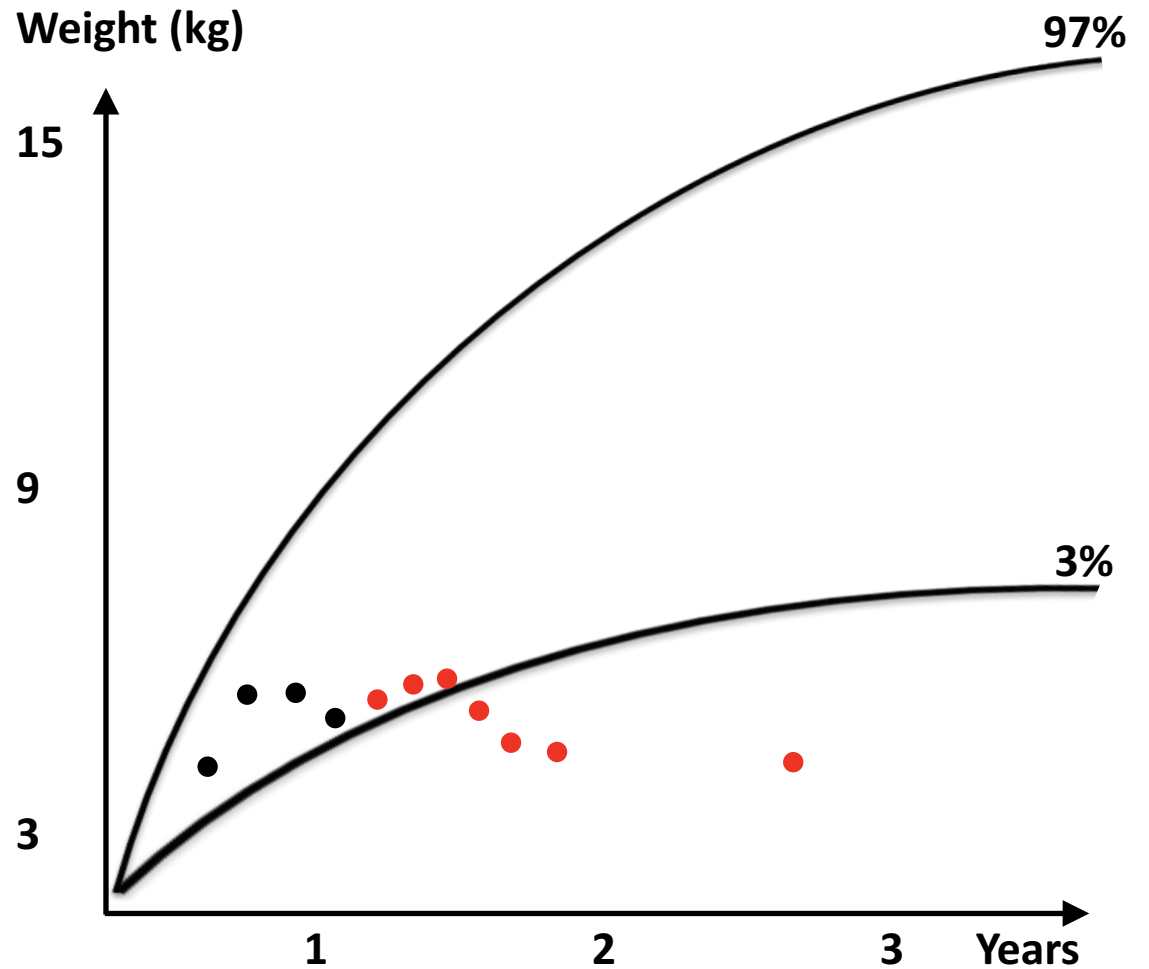
Lenters.. Bhutta. *BMC Public Health* 2013;13:1-15.

Schoones.. Volmink. *Cochrane Database Syst Rev* 2013;6:1-90.

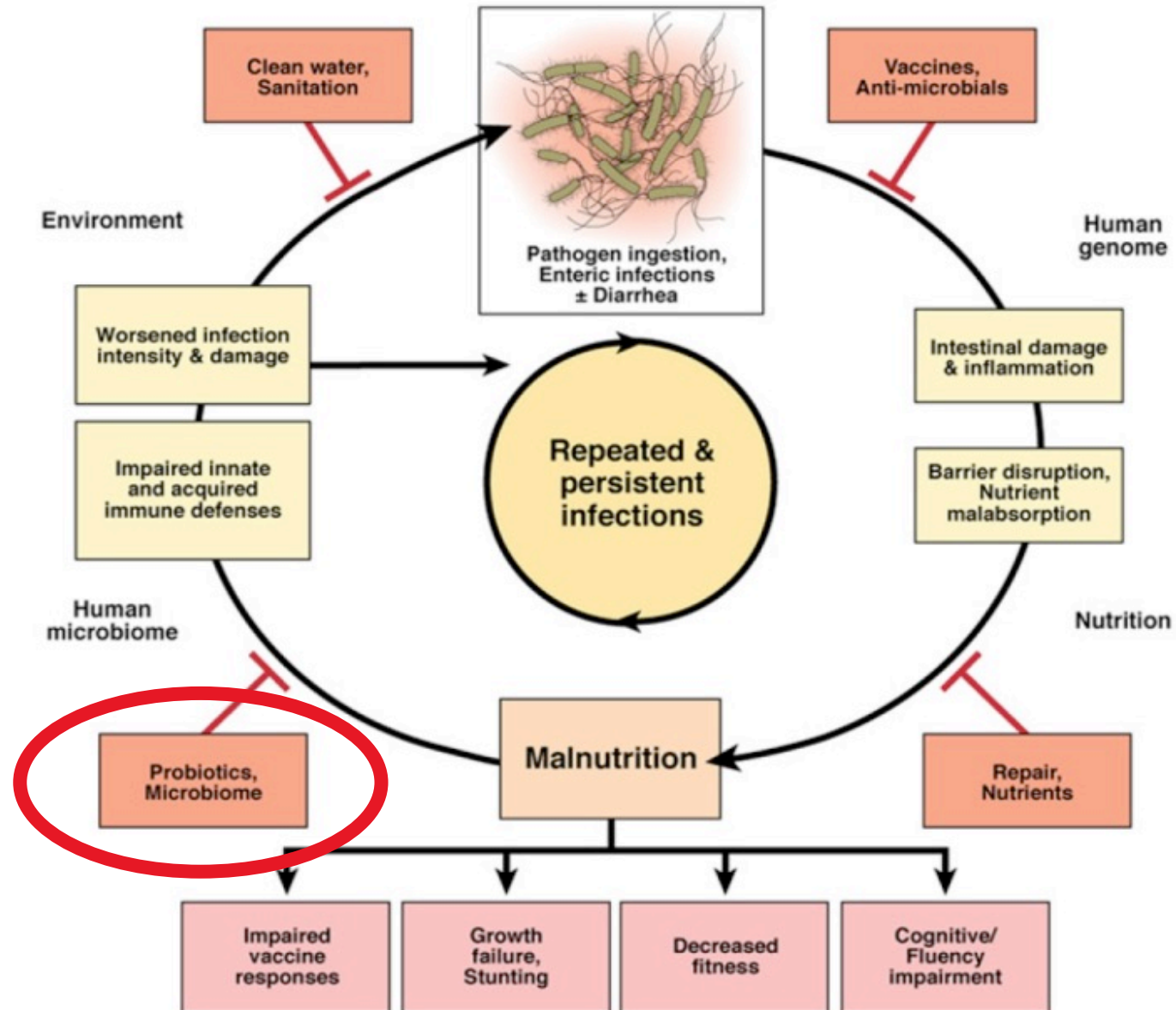


<http://www.thp.org/plumpynut-a-cure-for-malnutrition>

CASE PRESENTATION



THE VICIOUS CYCLE OF REPEATED INFECTIONS AND EARLY UNDERNUTRITION



OBJECTIVES

1. Recognize the distinct patterns of gut bacterial community configurations in undernourished children
2. List dietary, environmental, and host factors that shape the gut microbiome of undernutrition
3. Evaluate the clinical evidence supporting the use of microbiome-targeting therapies to enhance growth

CHANGES IN INTESTINAL BACTERIAL FLORA AND ROLE OF INFECTION IN KWASHIORKOR

P. M. SMYTHE
M.B. Cantab., M.R.C.P.

DEPARTMENT OF CHILD HEALTH, UNIVERSITY OF CAPE TOWN AND
GROOTE SCHUUR HOSPITAL

Gastric/duodenal bacterial overgrowth was subsequently reported in undernourished children from:

- **Guatemala**

Dammin GJ. *Bull World Health Organ* 1964;31:29-32.
Mata.. Viteri. *Am J Clin Nutr* 1972;25:118-26.

- **Aboriginal Australia**

Gracey & Stone. *Aust N Z J Med* 1972;2:215-9.

- **Indonesia**

Gracey & Stone. *Am J Clin Nutr* 1973;26:1170-4.

- **Brazil**

Maffei & Nobrega. *Gut* 1975;16:719-26.

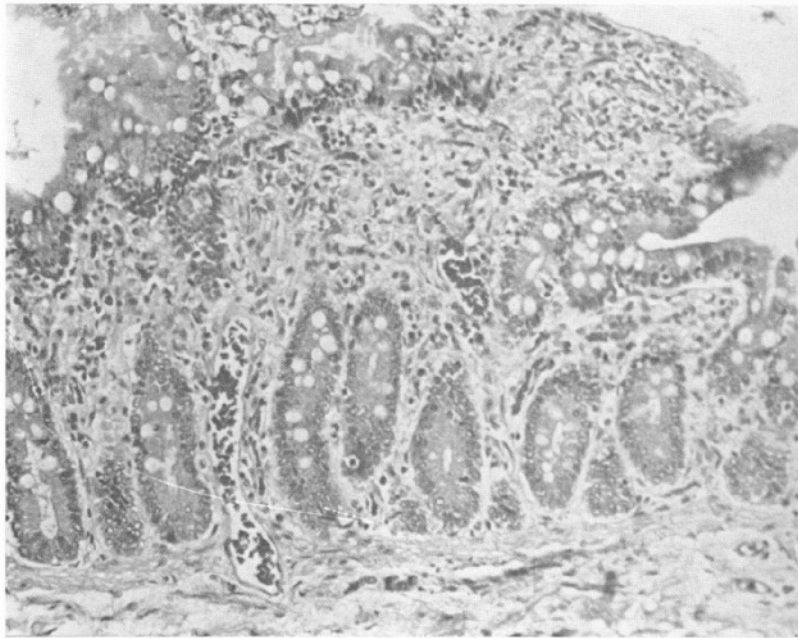
- **The Gambia**

Heyworth & Brown. *Arch Dis Child* 1975;50:27-33.

BACTERIAL GROWTH FROM GASTRIC JUICE, AND RECTAL SWAB, AND EFFECT OF THREE DAYS' ANTIBIOTIC TREATMENT ON THE INTESTINAL FLORA, IN 20 PATIENTS WITH KWASHIORKOR

Organisms	Gastric juice	Rectal swab	Both gastric juice and rectal swab	Rectal swab after 3 days' antibiotics
<i>Staph. aureus</i>	10	8	6	0
Coliforms	11	17	11	5
Salmonella	1	3	1	1
<i>Shigella flexneri</i>	0	2	0	0
Paracolon	2	5	1	4
Enterococci	0	4	0	7
<i>Proteus morgani</i>	0	14	0	4
<i>Ps. aeruginosa</i>	0	0	0	6
<i>Extent of bacterial growth:</i>				
None	3	0	..	5
Scanty	10	0	..	12
Moderate	6	0	..	3
Heavy	1	20	..	0

SEVERELY UNDERNOURISHED (BUT OTHERWISE HEALTHY) CHILDREN HAVE ABNORMAL INTESTINAL HISTOLOGY



1. Section of jejunum obtained from Guatemalan child with malnutrition but no diarrhoea. The structure of the villi is distorted and there is a reduced ratio of villus height to crypt depth. Note the sparse cellularity of the lamina propria and the complement of goblet cells.

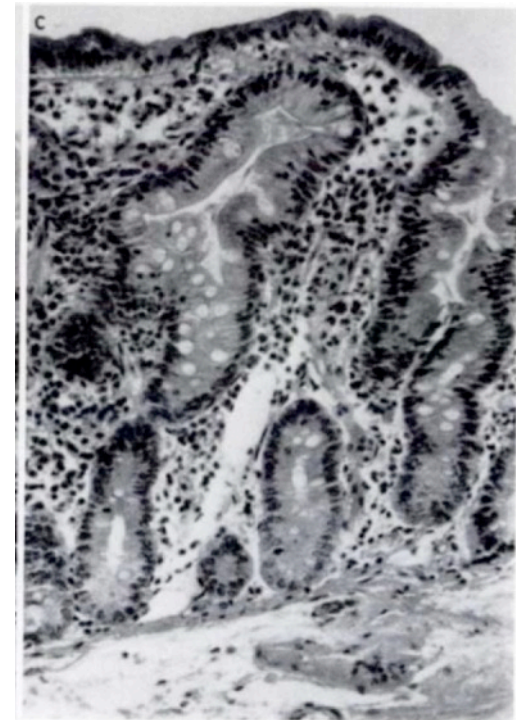
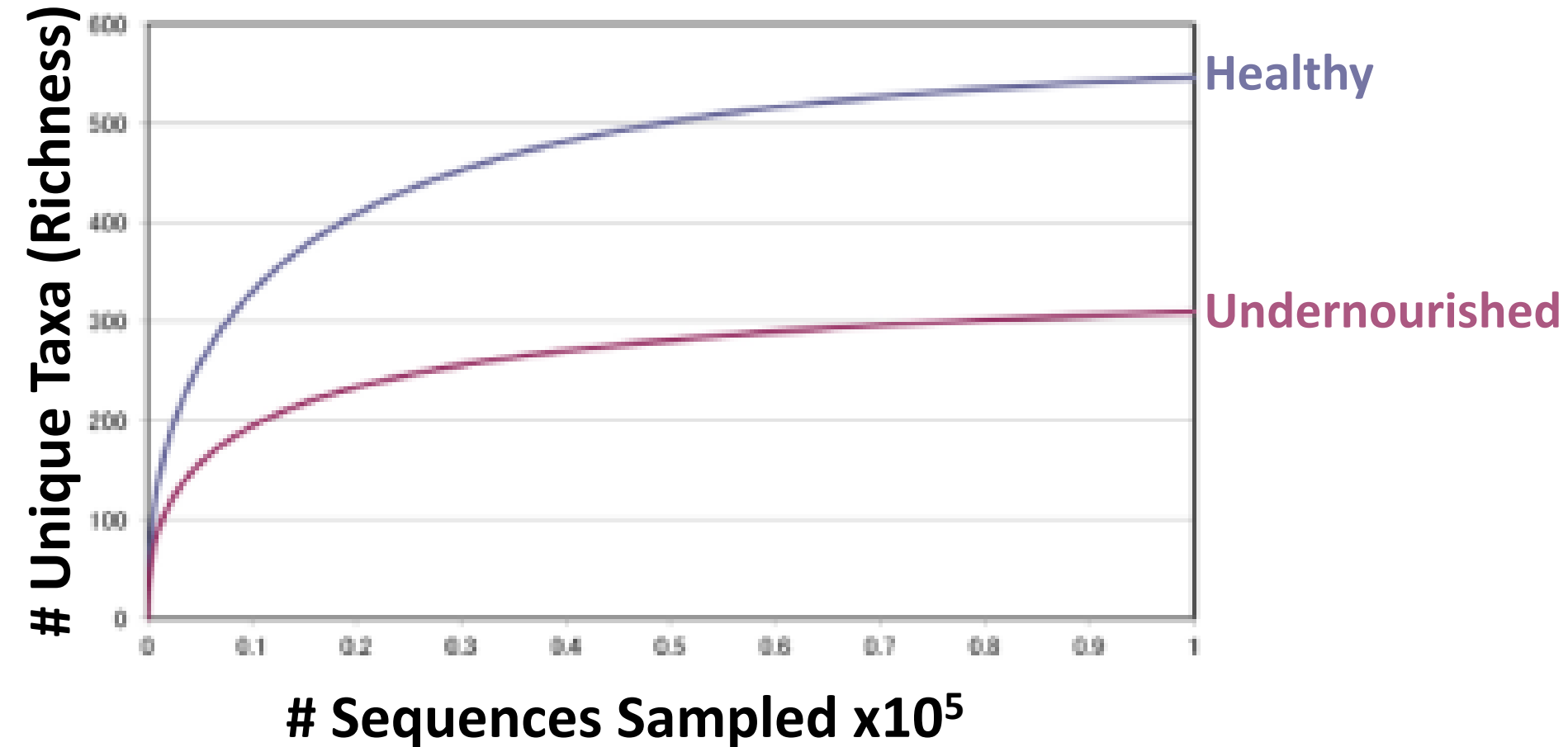
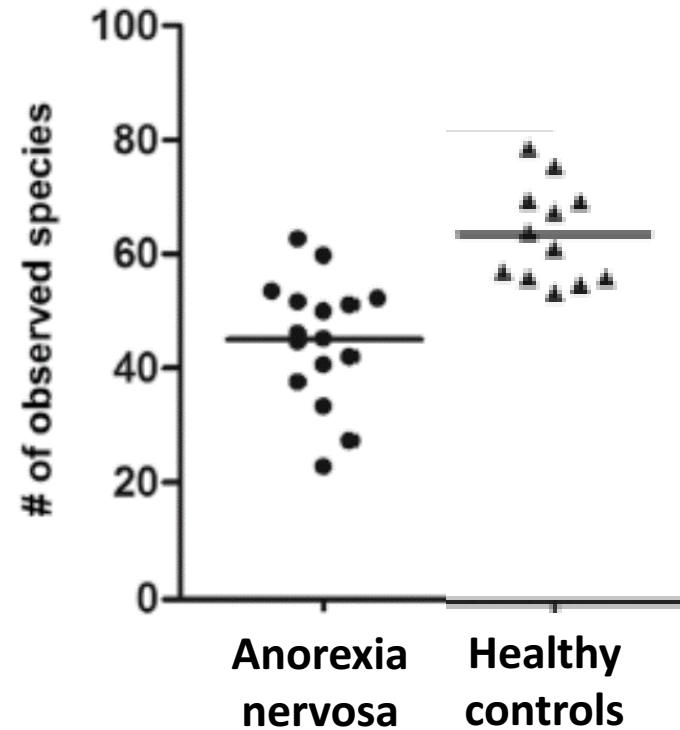
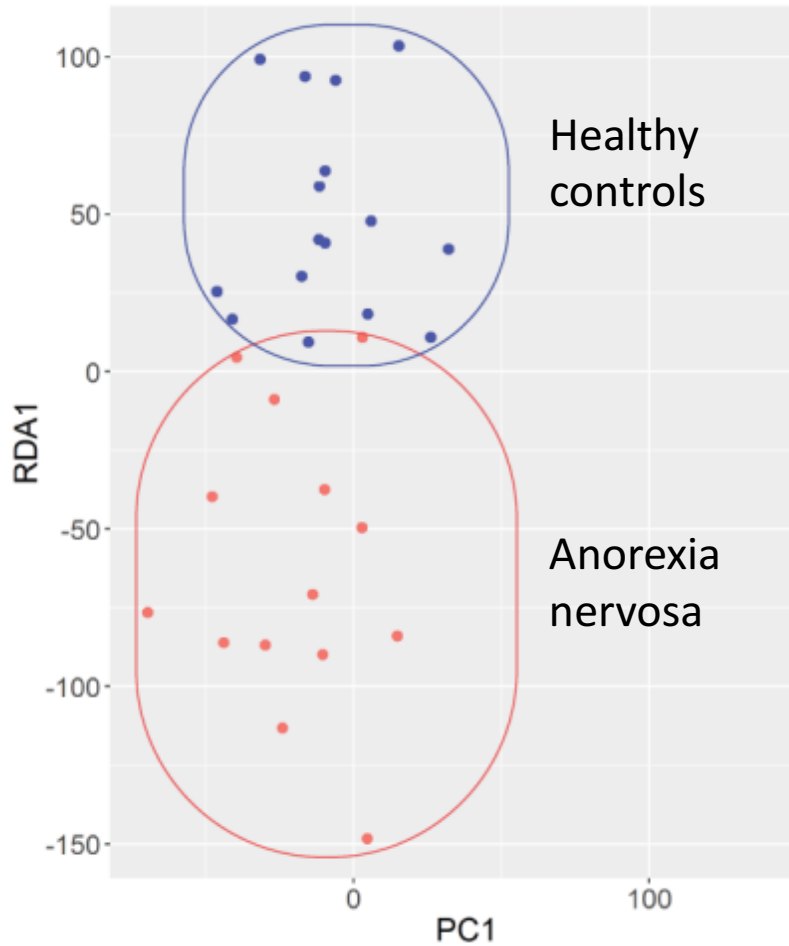


FIG. 2. Progressive fusion of the tips of two villi

STOOL FROM UNDERNOURISHED VS HEALTHY CHILDREN HAS DECREASED MICROBIOTA RICHNESS



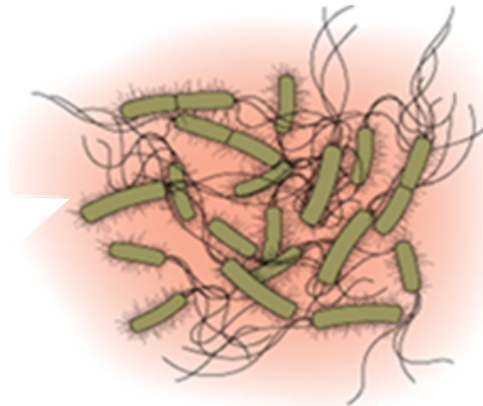
ANOREXIA NERVOSA PATIENTS ALSO HAVE “DYSBIOSIS” WITH DECREASED DIVERSITY



Proteobacteria, including Enterobacteriaceae, are overrepresented in anorexia vs healthy controls

BETA DIVERSITY IN UNDERNOURISHED VS HEALTHY CHILDREN

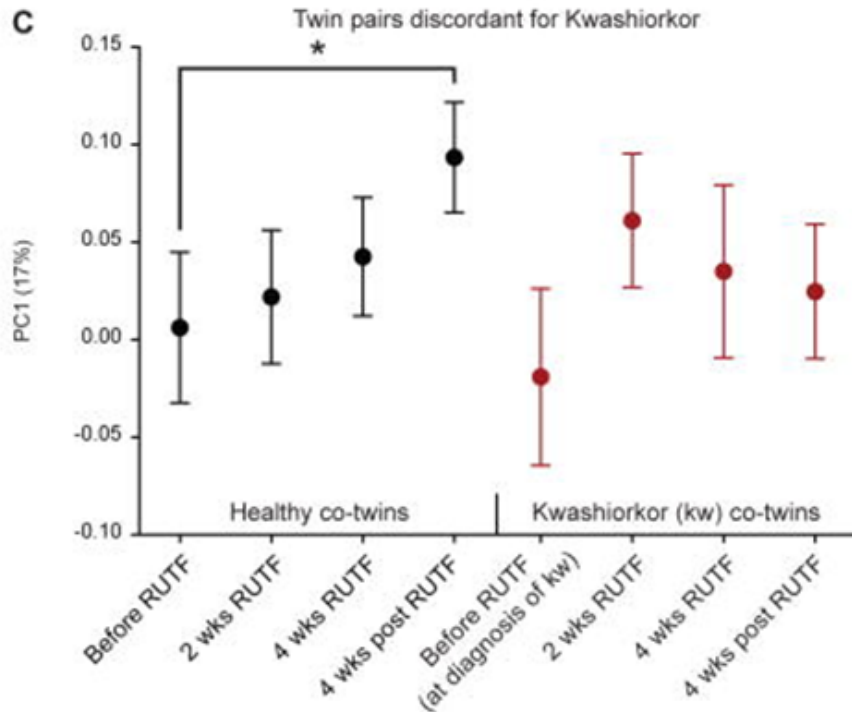
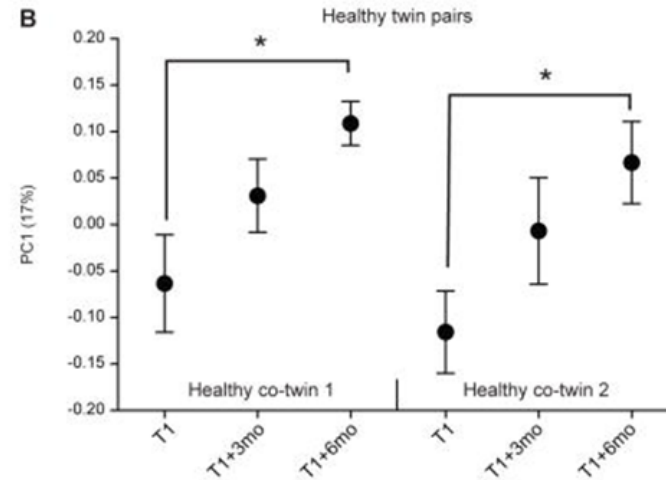
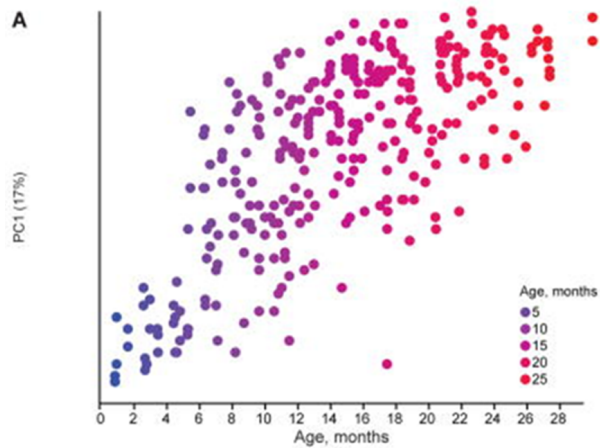
- **Increased** abundance of pathogenic genera within the phylum **Proteobacteria**, including *Enterobacter*, *Escherichia*, *Klebsiella*, and *Shigella*, even in the absence of diarrhea.



Microbial dysbiosis
±Enteropathogens

- **Decreased** abundance of genera with potentially **beneficial microbes**, including *Roseburia*, *Faecalibacterium*, *Butyrivibrio*, *Lactobacillus*, and *Bifidobacterium*.

GUT MICROBIOTA MATURITY IS IMPAIRED IN CHILD UNDERNUTRITION



KEY POINT 1

Objective 1: Recognize the distinct patterns of gut bacterial community configurations in undernourished children

- Children who are undernourished from a variety of causes have gut microbial community alterations (“**dysbiosis**”), characterized by:
 - Decreased richness (number of unique taxa)
 - Increased abundance of pathogens & potential pathogens
 - Decreased abundance of potentially beneficial microbes
 - Delayed microbiome maturation

OBJECTIVES

1. Recognize the distinct patterns of gut bacterial community configurations in undernourished children
2. List dietary, environmental, and host factors that shape the gut microbiome of undernutrition
3. Evaluate the clinical evidence supporting the use of microbiome-targeting therapies to enhance growth

Factors that may cause intestinal microbial dysbiosis in undernourished children

Maternal & perinatal factors

Dietary monotony, limited variation of nutrient-poor foods

↓ ratio of dietary: endogenous glycans available for metabolism

Decreased production of ileal antimicrobial peptides

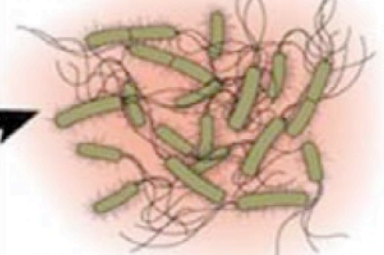
Enteropathogens (toxins, nutrient competition, mucus layer changes, altered motility)

Inflammation (antimicrobial peptides, disrupted O₂ gradient, reactive O₂ & N₂ species)

Impaired innate & acquired immunity exacerbating infection intensity & damage

Altered bile acid biosynthesis

Microbial "dysbiosis"



Decreased richness & commensal: pathogen ratio, Delayed maturation

Mechanisms by which intestinal microbial dysbiosis may impair weight gain

Toxins & other virulence factors

Enhanced pathogen colonization & invasion

Promotion of inflammation & enteric dysfunction

Less efficient energy harvest from non-digestible dietary components

Impaired *de novo* micronutrient biosynthesis

Differential metabolism of primary & conjugated bile acids

Altered transit time, nutrient malabsorption, decreased appetite, barrier disruption

Changes in systemic metabolites, hormones, and somatotrophic axis

Impaired weight gain

Persistent microbial dysbiosis

Undernutrition

Stunting, decreased fitness & earning potential

Impaired vaccine responses

Increased risk of obesity & co-morbidities

Altered brain metabolites, cognitive impairment

INFLAMMATION (A FEATURE OF ENVIRONMENTAL ENTEROPATHY) ALTERS THE GUT MICROBIOME

1. By triggering an immune response in which antimicrobial peptides released into the lumen innately defend against pathogens, but also target subsets of commensals

Sanchez de Medina.. Martinez-Augustin. *Inflamm Bowel Dis* 2014;20:2394-404

2. By disrupting the tightly regulated oxic-microoxic-anoxic zones in the lumen, influencing bacterial growth and transcriptional programs

Morris & Schmidt. *Nat Rev Microbiol* 2013;11:205-12.

Albenberg.. Wu. *Gastroenterol* 2014;147:1055-63.

Marteyn.. Tang. *Nature* 2010;465:355-61.

3. By generating reactive oxygen and nitrogen species, which shape microbial populations by facilitating respiration among certain bacteria

Winter.. Baumler. *Nature* 2010;467:426-9.

Winter.. Baumler. *Science* 2013;339:708-11.

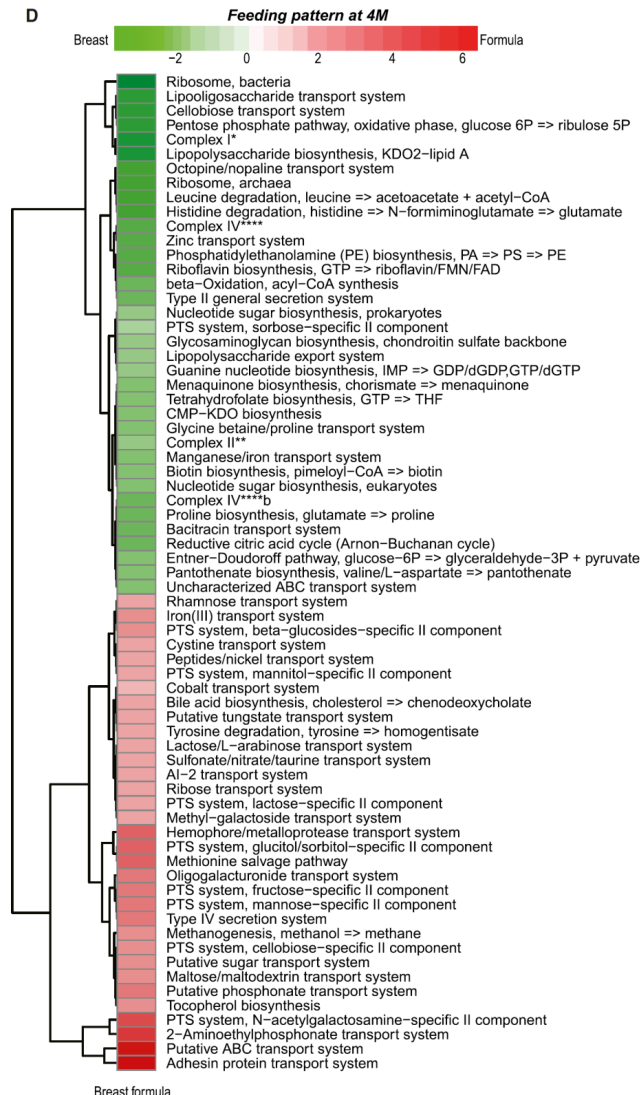
DIET (CARBOHYDRATE CONTENT) INFLUENCES A CHILD'S MICROBIOME

Compared to healthy Italian children, stool from healthy children in Burkina Faso was enriched with microbes (e.g., *Prevotella*, *Xylanibacter*) that harbor enzymes for metabolizing non-digestible dietary cellulose and xylans, key components of the Burkina Faso diet.



Fig. 1. Life in a rural village of Burkina Faso. (A) Village of Boulpon. (B) Traditional Mossi dwelling. (C) Map of Burkina Faso (modified from the United States CIA's World Factbook, 34). (D) Millet and sorghum (basic components of Mossi diet) grain and flour in typical bowls. (E) Millet and sorghum is ground into flour on a grinding stone to produce a thick porridge called Tô.

DIET (BREAST MILK VS FORMULA) INFLUENCES AN INFANT'S MICROBIOME



Compared to formula-fed infants, the gut microbiota of breastfed infants is *less diverse*, consistent with enrichment of genes required for the degradation of **human milk oligosaccharides (HMOs)** from breast milk.

* (NADH dehydrogenase), NADH dehydrogenase I

** (succinate dehydrogenase / fumarate reductase), succinate dehydrogenase

*** (Cytochrome bc1 complex)

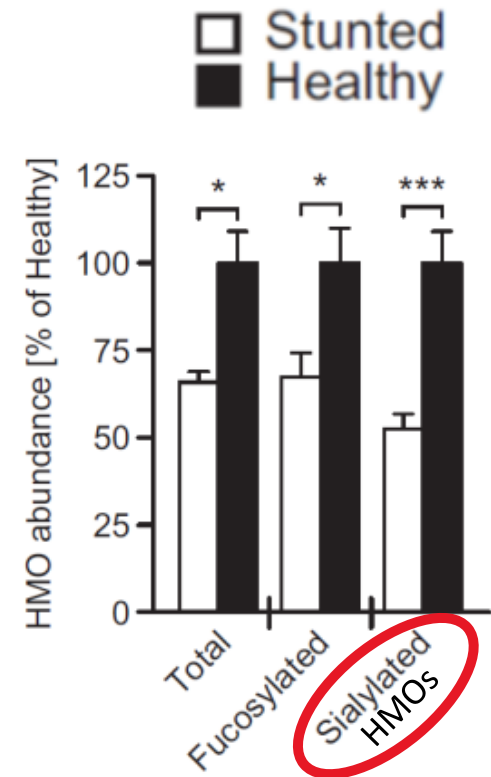
**** (Cytochrome c oxidase), cytochrome c oxidase, ccb3-type

****b (Cytochrome c oxidase), cytochrome c oxidase/ubiquinol oxidase/cytochrome c oxidase/quinol oxidase polypeptide

Backhed.. Wang. *Cell Host Microbe* 2015;17:690-703.

DIET INFLUENCES THE INFANT MICROBIOME

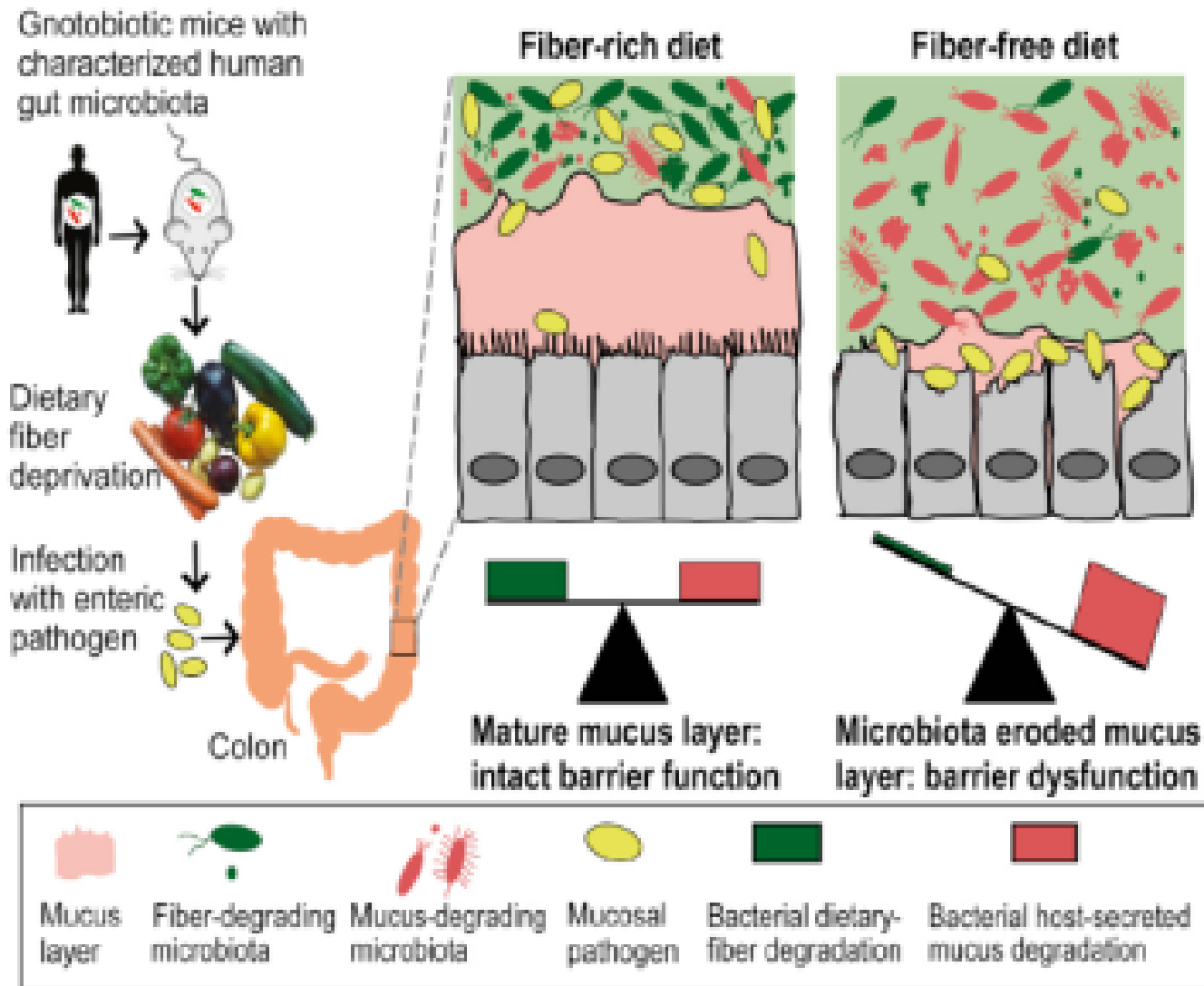
- Not all breast milk is equal
- Malawian mothers with severely stunted vs healthy infants produced decreased quantities of **human milk oligosaccharides (HMOs)**
- How might HMOs affect growth?



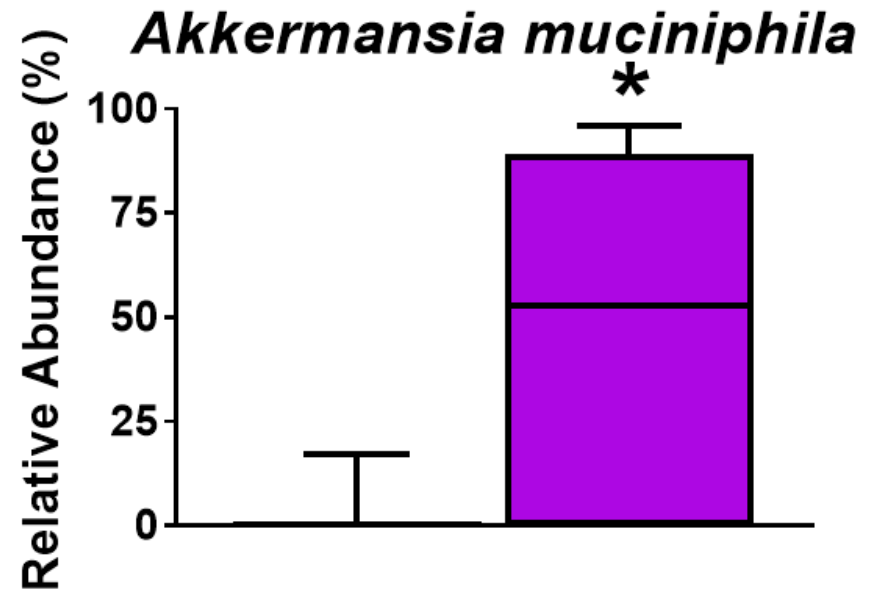
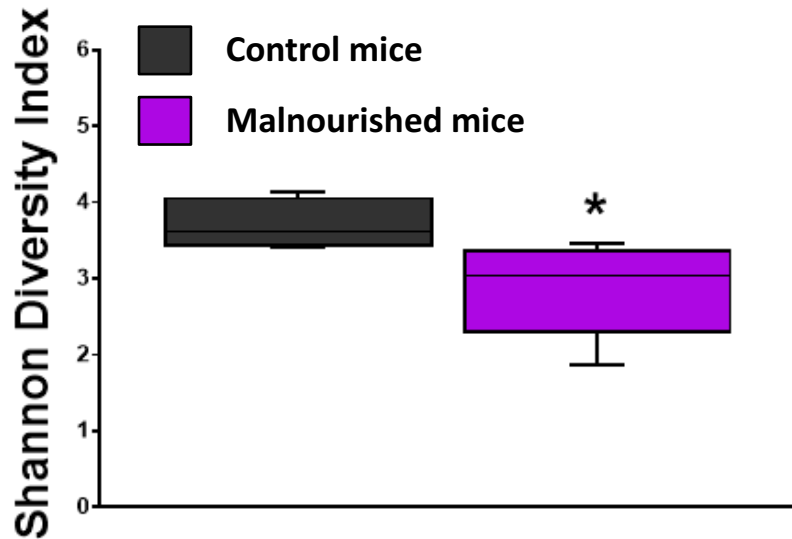
DIET INFLUENCES THE INFANT MICROBIOME



DIET-INDUCED GUT MICROBIAL “DYSBIOSIS” CAN CONFER SUSCEPTIBILITY TO INFECTION



UNDERNOURISHED MOUSE PUPS HAVE DECREASED FECAL MICROBIAL DIVERSITY, WITH INCREASED ABUNDANCE OF MUCOLYTIC BACTERIA



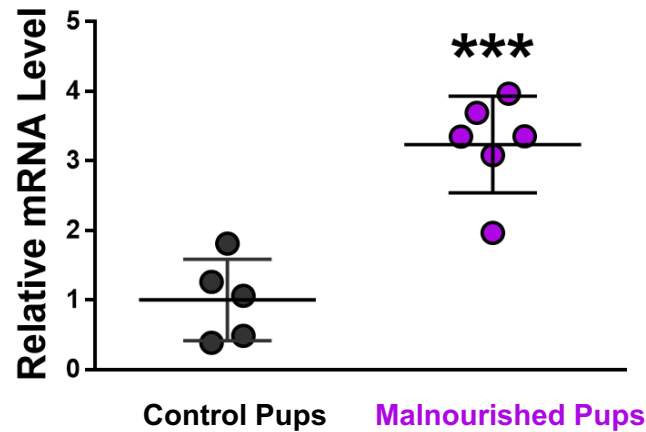
In humans, the relative abundance of *Akkermansia muciniphila* is inversely proportional to body mass index (BMI).

Santacruz.. Sanz. *Br J Nutr* 2010;104:83-92.

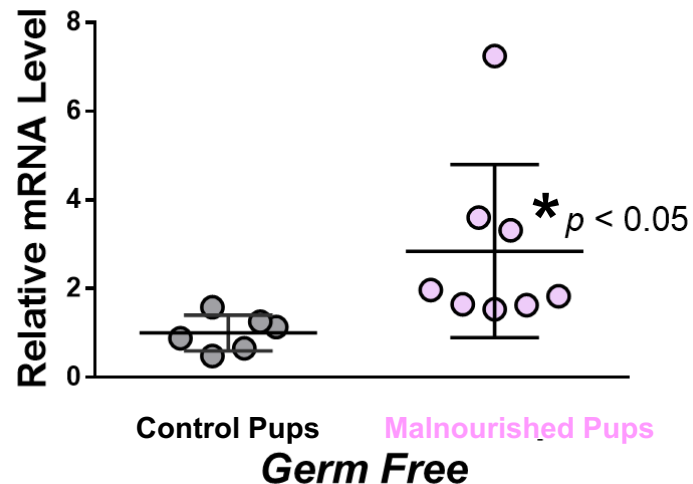
Karlsson.. Thorngren-Jerneck. *Obesity* 2012;20:2257-61.

THIS OVERABUNDANCE OF MUCOLYTIC MICROBES IS ASSOCIATED WITH INCREASED HOST MUCIN GENE EXPRESSION

Muc2



Conventional



THREE MOUSE MODELS OF EARLY-LIFE UNDERNUTRITION

- **Timed Separation (TmSep)**

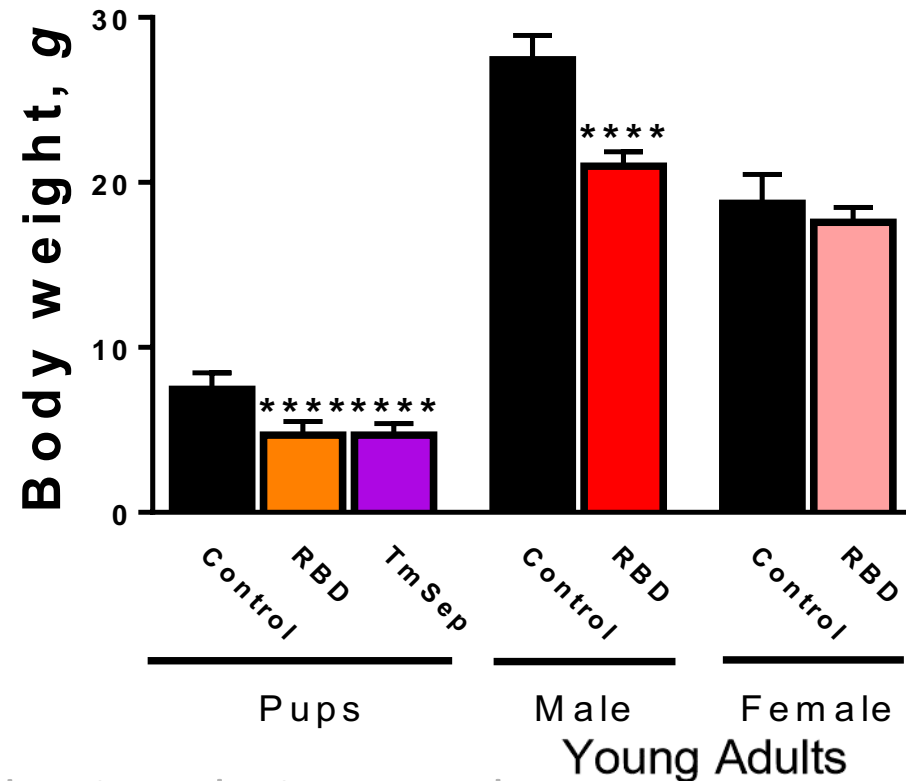
- TmSep: 12 hours/day away from mother
- Controls: litters of normal pups

- **Regional Basic Diet (RBD) Female**

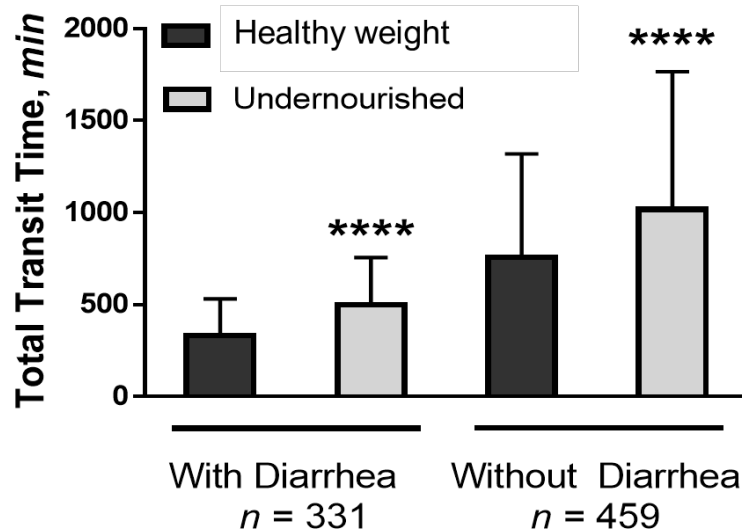
- RBD: Mothers fed 5% fat, 7% protein diet
- Controls: Mothers fed isocaloric diet

- **RBD Young Adults**

- RBD Pups weaned to RBD chow
- Controls: Control Pups weaned to isocaloric Control Diet



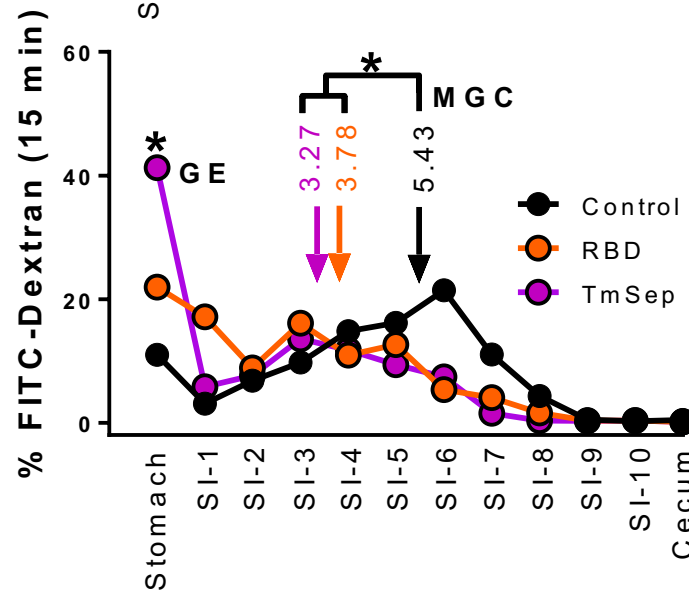
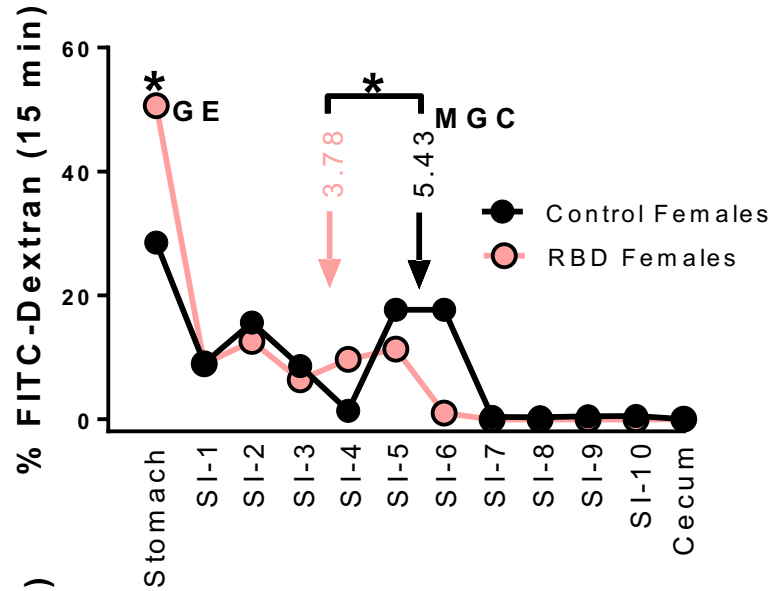
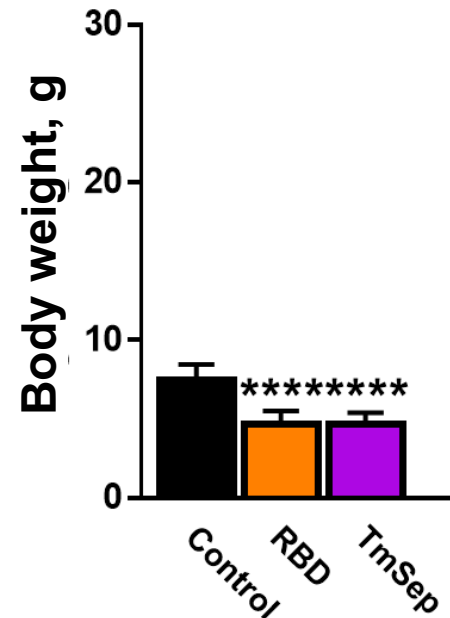
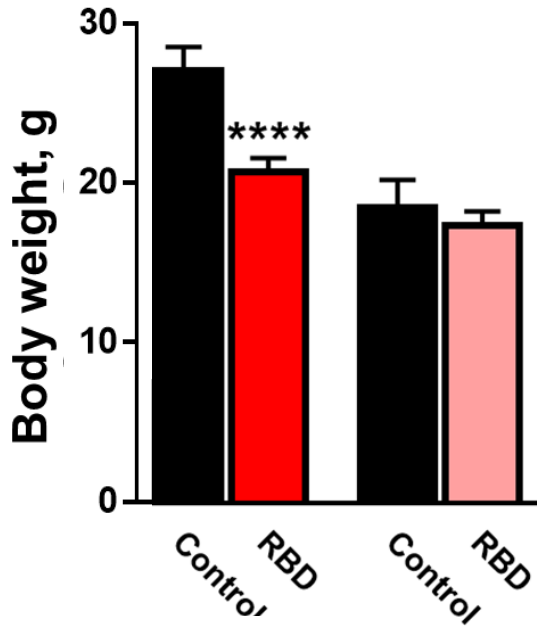
UNDERNOURISHED CHILDREN HAVE SLOW GASTROINTESTINAL MOTILITY



Viteri & Schneider. *Med Clin North Am* 1974;58: 1487-505.

Slow transit in undernutrition is linked to bacterial overgrowth, abdominal distention, constipation, and blunted appetite.

UNDERNUTRITION SLOWS GASTRIC AND SMALL BOWEL TRANSIT IN MULTIPLE MOUSE MODELS



GE = gastric emptying
MGC = mean geometric center of bolused dye

Transit time in undernourished young adult males was minimally affected.



Control TmSep

UNDERNOURISHED CHILDREN HAVE DECREASED INTESTINAL BILE ACIDS

Luminal events of lipid absorption in protein-calorie malnourished children; relationship with nutritional recovery and diarrhea.

II. Alterations in bile acid content of duodenal aspirates¹

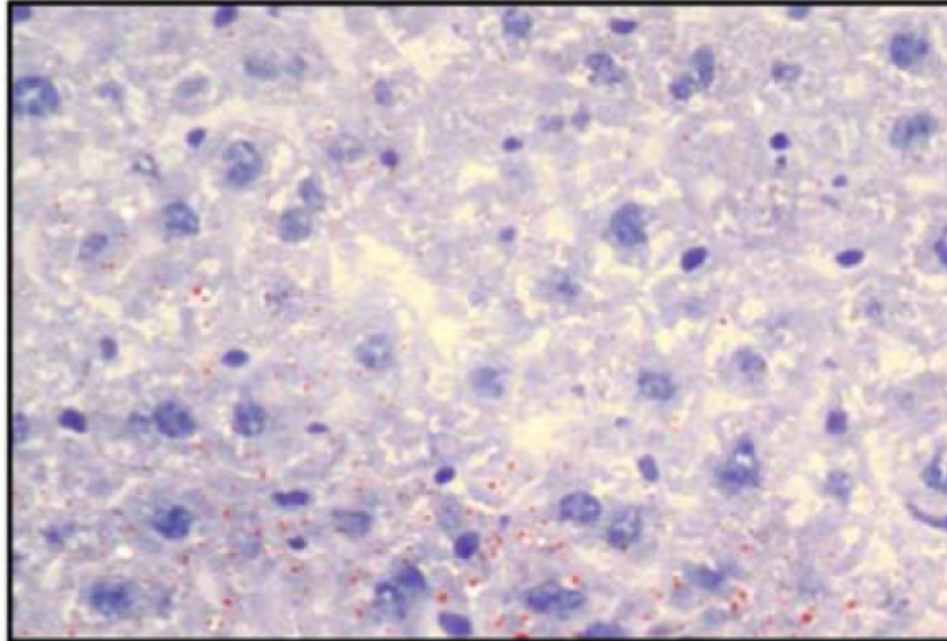
Roberto E. Schneider,² M.D. and Fernando E. Viteri,³ M.D., D.Sc.

ABSTRACT Duodenal aspirates from PCM patients and from children with diarrhea present quantitative and qualitative alterations in their content of bile acids, consisting mainly in decreased concentrations of conjugated bile acids (CBA) and increased amounts of primary and secondary free bile acids (FBA). In the absence of diarrhea, PCM was associated with relatively uniform decrements of CBA, increasing to normal levels with nutritional recovery; the amounts of primary and secondary FBA did not change significantly with recovery. The effect of diarrhea on the bile composition of the children varied depending on the patient's nutritional status: *a*) In recovered children with diarrhea, all CBA, especially the taurine conjugates, were decreased; the glycine/taurine (G/T) ratios were above 1:4 in most of these patients. Although all FBA increased with diarrhea, such an increase was most significant in cholic and lithocholic acids; these children also presented high FBA/CBA ratios. *b*) In PCM children, the degree of decrement of CBA during diarrhea was similar in taurine and glycine conjugates, the G/T ratio remaining within normal limits. The concentrations of primary and secondary FBA did not change significantly in PCM children with diarrhea when compared with the group without diarrhea. It is believed that the changes in bile acids observed in these children are due to the interaction of malnutrition, diarrhea, and an increased gastrointestinal flora. A theory of how the bile acid events described herein occur is also proposed. *Am. J. Clin. Nutr.* 27: 788-796, 1974.

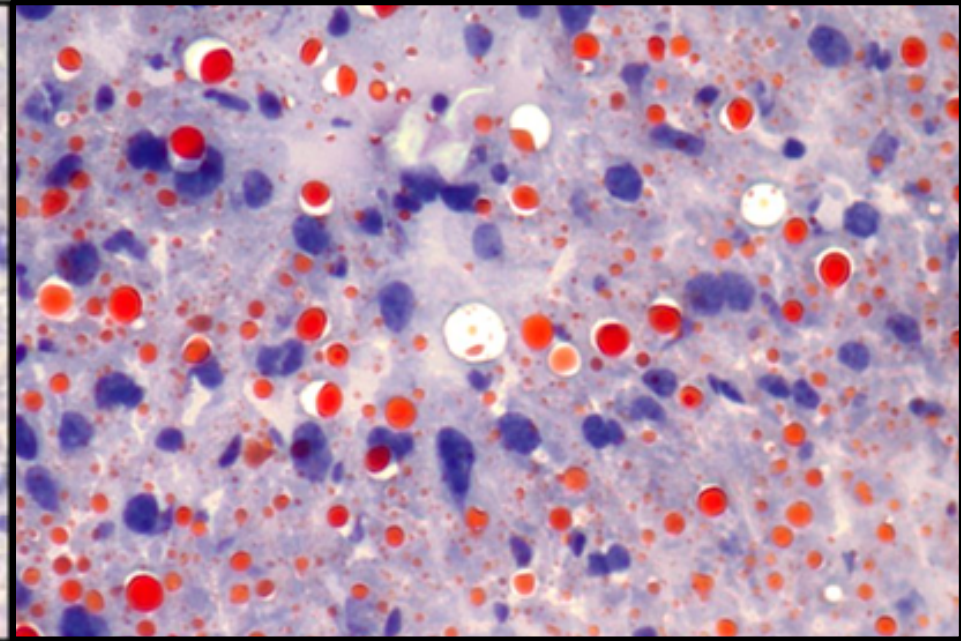
Schneider & Viteri. *Am J Clin Nutr* 1974;27: 788-96.

Decreased bile acids in undernutrition are linked to dietary fat malabsorption, fat-soluble vitamin deficiencies, bacterial overgrowth, and poor weight gain during refeeding.

UNDERNOURISHED MICE (AND CHILDREN) EXHIBIT MACROVESICULAR STEATOSIS



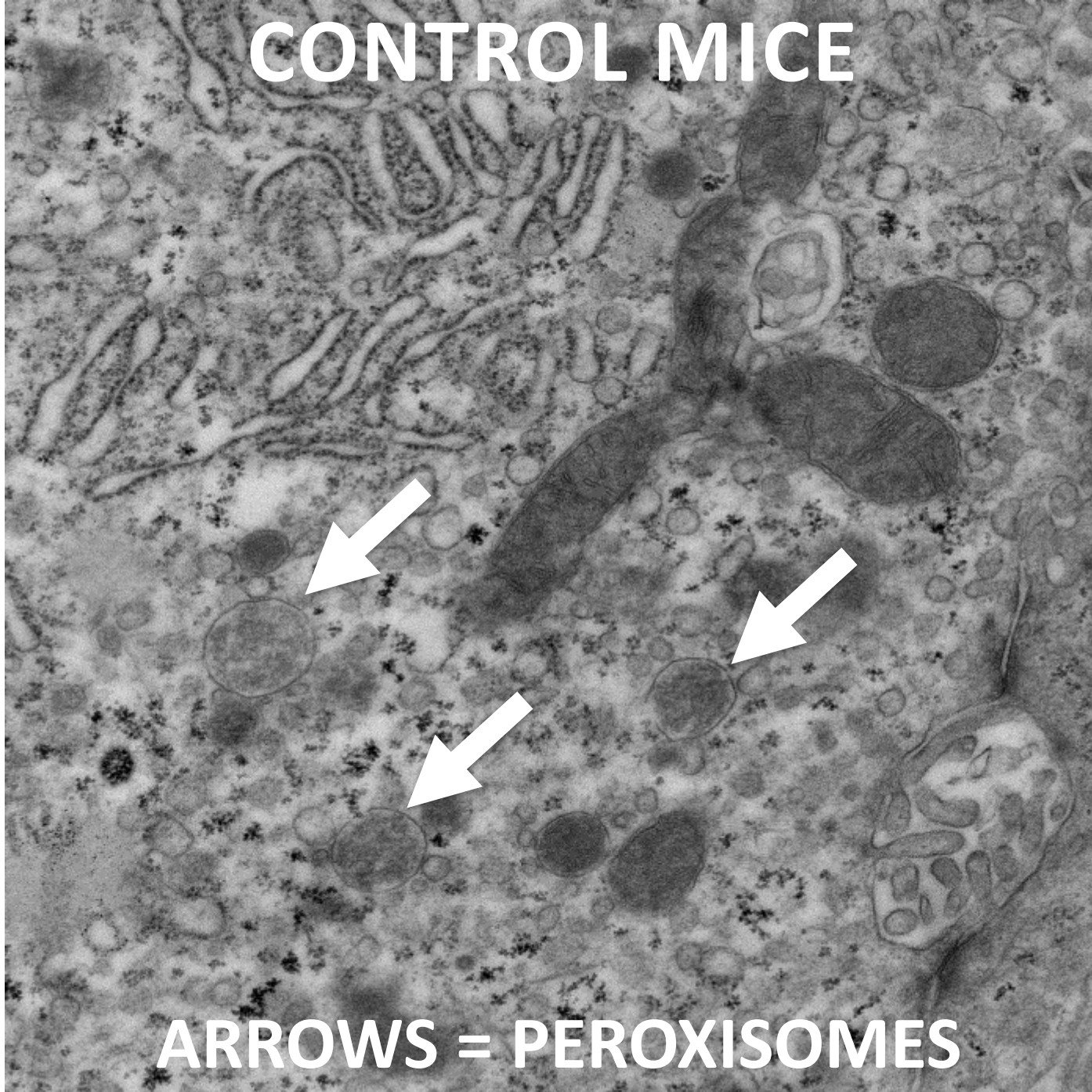
Control



RBD Young Adults

Oil red O stain, flash-frozen livers, x160

CONTROL MICE



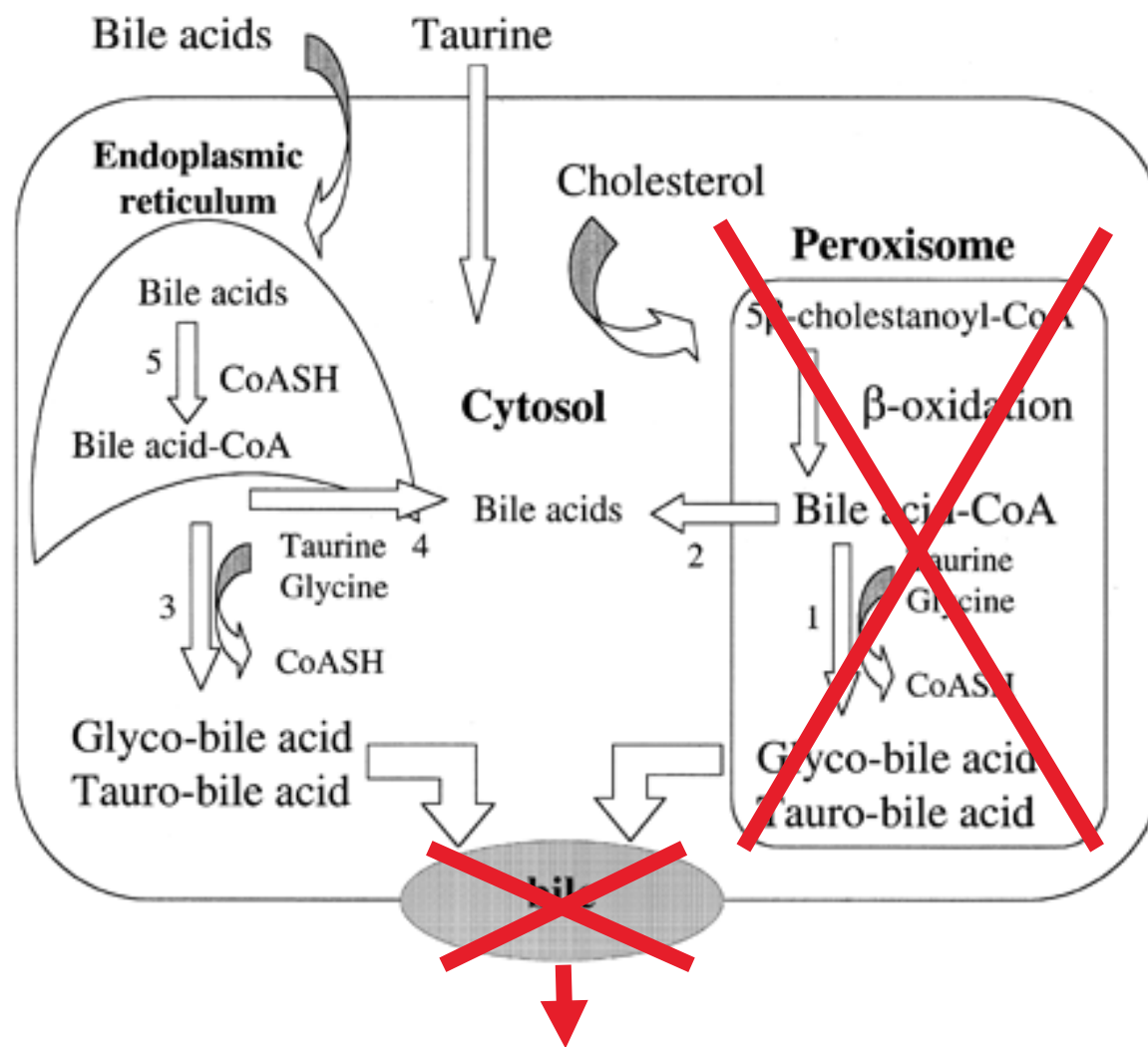
ARROWS = PEROXISOMES

—| 500 nm

UNDERNOURISHED MICE HAVE NO PEROXISOMES



1 um



Altered Intestinal Microbiota?

Factors that may cause intestinal microbial dysbiosis in undernourished children

Maternal & perinatal factors

Dietary monotony, limited variation of nutrient-poor foods

↓ ratio of dietary: endogenous glycans available for metabolism

Decreased production of ileal antimicrobial peptides

Enteropathogens (toxins, nutrient competition, mucus layer changes, altered motility)

Inflammation (antimicrobial peptides, disrupted O₂ gradient, reactive O₂ & N₂ species)

Impaired innate & acquired immunity exacerbating infection intensity & damage

Altered bile acid biosynthesis

Microbial "dysbiosis"



Decreased richness & commensal: pathogen ratio, Delayed maturation

Mechanisms by which intestinal microbial dysbiosis may impair weight gain

Toxins & other virulence factors

Enhanced pathogen colonization & invasion

Promotion of inflammation & enteric dysfunction

Less efficient energy harvest from non-digestible dietary components

Impaired *de novo* micronutrient biosynthesis

Differential metabolism of primary & conjugated bile acids

Altered transit time, nutrient malabsorption, decreased appetite, barrier disruption

Changes in systemic metabolites, hormones, and somatotrophic axis

Impaired weight gain

Persistent microbial dysbiosis

Undernutrition

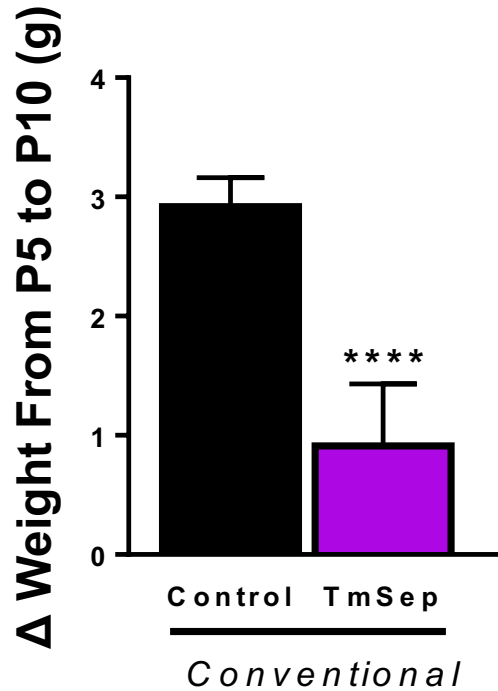
Stunting, decreased fitness & earning potential

Impaired vaccine responses

Increased risk of obesity & co-morbidities

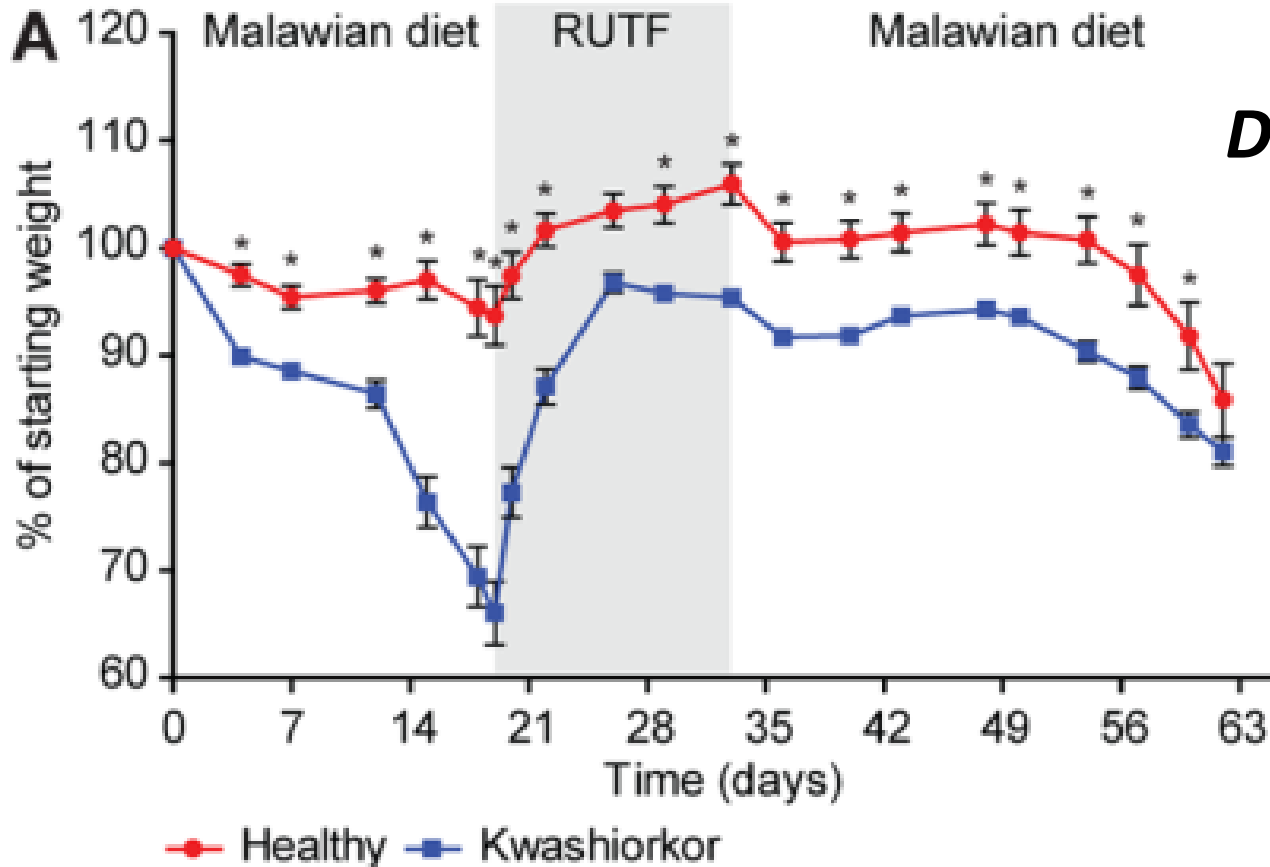
Altered brain metabolites, cognitive impairment

GERM-FREE MICE TOLERATE EARLY-LIFE UNDERNUTRITION BETTER THAN CONVENTIONAL MICE WITH INTESTINAL BACTERIA



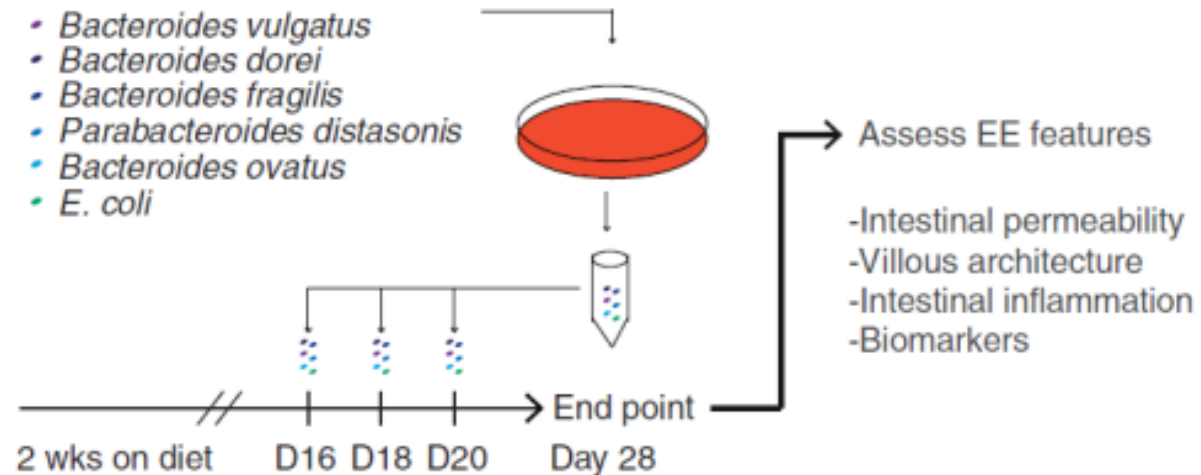
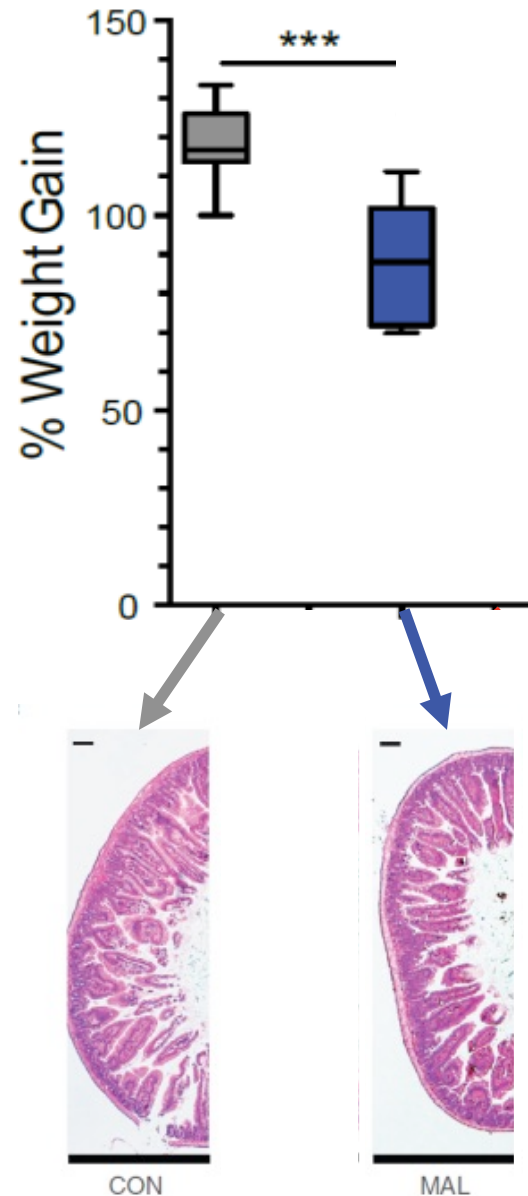
$n = 10-18$

FECAL MICROBES FROM UNDERNOURISHED CHILDREN CAN CAUSE UNDERNUTRITION IN GNOTOBIOTIC MICE (UNDER THE RIGHT CONDITIONS)



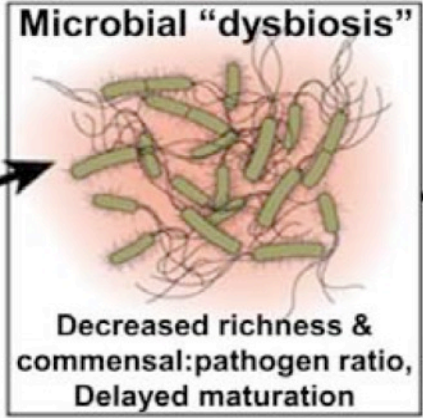
GUT BACTERIA (EVEN NON-PATHOGENS) CAN CAUSE INFLAMMATION AND GROWTH IMPAIRMENT

Consumption of a low-protein, low-fat diet, in combination with iterative exposure to 6 non-pathogenic gut microbes, produces inflammation and weight loss without overt diarrhea

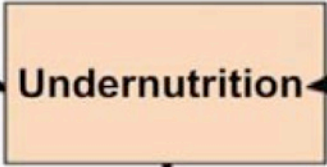
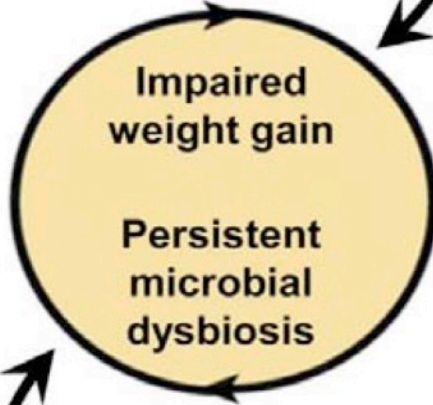


Factors that may cause intestinal microbial dysbiosis in undernourished children

- Maternal & perinatal factors
- Dietary monotony, limited variation of nutrient-poor foods
- ↓ ratio of dietary: endogenous glycans available for metabolism
- Decreased production of ileal antimicrobial peptides
- Enteropathogens (toxins, nutrient competition, mucus layer changes, altered motility)
- Inflammation (antimicrobial peptides, disrupted O₂ gradient, reactive O₂ & N₂ species)
- Impaired innate & acquired immunity exacerbating infection intensity & damage
- Altered bile acid biosynthesis



- Mechanisms by which intestinal microbial dysbiosis may impair weight gain**
- Toxins & other virulence factors
- Enhanced pathogen colonization & invasion
- Promotion of inflammation & enteric dysfunction
- Less efficient energy harvest from non-digestible dietary components
- Impaired *de novo* micronutrient biosynthesis
- Differential metabolism of primary & conjugated bile acids
- Altered transit time, nutrient malabsorption, decreased appetite, barrier disruption
- Changes in systemic metabolites, hormones, and somatotrophic axis



- Stunting, decreased fitness & earning potential
- Impaired vaccine responses
- Increased risk of obesity & co-morbidities
- Altered brain metabolites, cognitive impairment

KEY POINT 2

Objective 2: List dietary, environmental, and host factors that shape the gut microbiome of undernutrition

- “Dysbiosis” of undernutrition can be shaped by many factors, including:
 - Prenatal/perinatal factors, carbohydrate composition of breast milk or the post-wean diet, inflammation, presence of pathogens, host intestinal mucus profile...
- Mechanisms by which “dysbiosis” can impair weight gain are less clear, but might include:
 - Bacterial toxins, subclinical inflammation, decreased efficiency of energy harvest from diet, impaired micronutrient biosynthesis, altered gastrointestinal motility, bile acid pool changes...

OBJECTIVES

1. Recognize the distinct patterns of gut bacterial community configurations in undernourished children
2. List dietary, environmental, and host factors that shape the gut microbiome of undernutrition
3. Evaluate the clinical evidence supporting the use of microbiome-targeting therapies to enhance growth

A PROMISING TRIAL OF ANTIBIOTICS FOR CHILDREN WITH SEVERE ACUTE UNDERNUTRITION

- 2,767 Malawian children prescribed RUTF as outpatient treatment for severe acute undernutrition
- Children randomly assigned to twice-daily placebo vs amoxicillin (80-90 mg/kg/day) or cefdinir (14 mg/kg/day) for 7 days
- **Placebo** increased the relative risk of **treatment failure**:
 - RR 1.32 [1.04 – 1.68] vs amoxicillin
 - RR 1.64 [1.27 – 2.11] vs cefdinir
- **Placebo** increased the relative risk of **mortality**:
 - RR 1.55 [1.07 – 2.24] vs amoxicillin
 - RR 1.80 [1.22 – 2.64] vs cefdinir

TWO OTHER LARGE TRIALS FAILED TO SHOW BENEFIT OF ANTIBIOTICS FOR UNDERNOURISHED CHILDREN

- 2,412 children in Niger with severe acute undernutrition randomized to twice-daily placebo vs amoxicillin (80 mg/kg/day) x7 days
 - **No effect on nutritional recovery over 8 week follow-up**
- 1,778 children in Kenya who had recovered from severe acute undernutrition randomized to daily placebo vs co-trimoxazole (120 or 240 mg/day) x6 months
 - **No effect on mortality over 12 month follow-up**

THE PRONUT STUDY: TESTING A PREBIOTIC + PROBIOTIC IN CHILD UNDERNUTRITION

- Randomized, placebo controlled trial enrolled 795 Malawian children hospitalized for nutritional rehabilitation
- Children randomized to RUTF + placebo vs RUTF + **Synbiotic 2000 Forte**
 - 4 probiotics: *Pediococcus pentosaceus*, *Leuconostoc mesenteroides*, *Lactobacillus paracasei*, *Lactobacillus plantarum*
 - 4 prebiotics: oat bran, inulin, pectin, resistant starch
- Median 33 days of treatment

THE PRONUT STUDY: TESTING A PREBIOTIC + PROBIOTIC IN CHILD UNDERNUTRITION

- Result: No significant effect on nutritional cure or on any other nutritional outcome
- Reasons for this negative result?

Synbiotic 2000 Forte

- 4 probiotics: *Pediococcus pentosaceus*, *Leuconostoc mesenteroides*, *Lactobacillus paracasei*, *Lactobacillus plantarum*
- 4 prebiotics: oat bran, inulin, pectin, resistant starch

SUMMARY OF LARGE RANDOMIZED, PLACEBO-CONTROLLED CLINICAL TRIALS TO DATE

Table 1. Randomized controlled trials that evaluate microbiome-targeting therapies to improve nutritional status in undernourished children.

Intervention	Setting	Number of Study Participants	Result	Reference
Synbiotic 2000 Forte	Malawi	795	No significant effect on nutritional cure	104
Amoxicillin or cefdinir	Malawi	2767	Placebo increased risk of treatment failure (RR 1.32 [1.04–1.68] vs amoxicillin; RR 1.64 [1.27–2.11] vs cefdinir) and mortality (RR 1.55 [1.07–2.24] vs amoxicillin; RR 1.80 [1.22–2.64] vs cefdinir)	105
Amoxicillin	Niger	2412	No significant effect on nutritional recovery	106
Co-trimoxazole	Kenya	1778	No significant effect on mortality	107

Note. RR, relative risk followed by 95% confidence interval in brackets.

- None of these landmark studies assessed how treatment vs placebo affected the gut microbiome
- Would a beneficial effect of a broad-spectrum antibiotic be worth the risks?

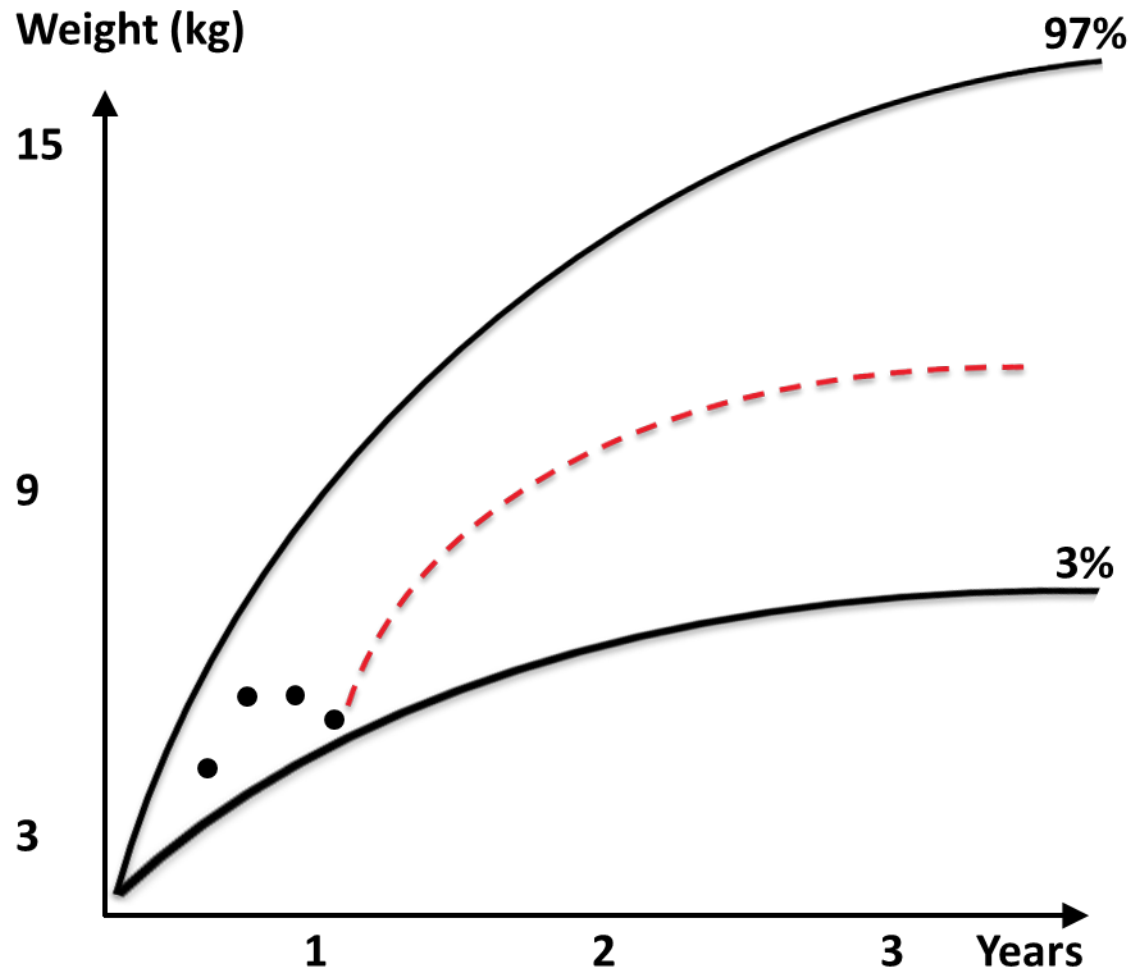
KEY POINT 3

Objective 3: Evaluate the clinical evidence supporting the use of microbiome-targeting therapies for undernutrition

- Although there is currently not enough clinical evidence to recommend microbiome-targeting therapies for undernourished children, promising preclinical models suggest that individualized therapies might one day allow clinicians to improve a child's growth trajectory

WHAT LIES AHEAD?

- Development of low-cost biomarkers to:*
- A) Identify children who would benefit from microbiome-targeting therapies*
 - B) Select the specific agents needed to address an individual's functional imbalances*



ACKNOWLEDGEMENTS

•Baylor College of Medicine

Pediatric GI, Hepatology & Nutrition Research Laboratories

- Tripti Halder
- Sanjiv Harpavat
- Swapna Krishnamoorthy
- Subapradha Narayanan
- Benjamin Shneider
- Krishnakant Soni
- Sundararajah Thevananther
- M. Elizabeth Tessier
- Jennifer Yeh

•Moore Laboratory

- Sungwoo Choi
- Kangho Kim
- David Moore

•Robert Britton

•Rui Chen

•Cristian Coarfa

•Margaret Conner

•Sridevi Devaraj

•Milton Finegold

•Robert Shulman

•Jennifer Spinler

•Lanlan Shen

•Arun Sreekumar

•James Versalovic

•Lisa White

Funding

- Young Investigator Grant for Probiotics Research, Global Probiotics Council
- Early Career Award, Thrasher Research Fund
- Pediatric GI Training Grant, NIH/NIDDK T32DK007664 (PI: Shulman)
- NASPGHAN Foundation / Nestlé Nutrition Research Young Investigator Development Award
- Pilot/Feasibility Award, Texas Medical Center Digestive Disease Center (PHS grant P30DK56338)
- Functional and Mechanistic Award, Alkek Center for Metagenomics and Microbiome Research
- Research Grant, American Neurogastroenterology and Motility Society
- AGA-Rome Foundation Functional GI & Motility Disorders Pilot Research Award
- Chao Physician-Scientist Award
- NIH/NIDDK K08DK113114

