Definition

The concept of the human microbiome was first suggested by Joshua Lederberg, who coined the term "microbiome, to signify the ecological community of commensal, symbiotic, and pathogenic microorganisms that literally share our body space".

Microbiome = the community of microorganisms that shares our body space

Population 100 Trillion: Microbes outnumber human cells 10:1.

Humans have 23,000 genes. Microbiome has 8 million genes.

Microbiome

"2nd human genome"

"Newly discovered organ"

"Bacterial inner rainforest"



Evolutionary Timeline

• 200 million years ago: Mammals

• 60 million years ago: Primates

• 2.5 million years ago: Genus Homo

(including humans)

Microbes appeared 3.5 billion years ago



Welcome to Chicago Population: 2.6 million



Welcome to Gut Population: 100 trillion

Microbiomes are quite dominant

- GI tract is sterile at birth
- GI tract colonized by bacteria within a few hours
- Microbial flora established within 3-4 weeks
- More than 1000 species of microbes take up house

Gut bacteria must resist removal

- 1-3 billion cells per hour are shed by the small intestine. 100-300 million cells per hour are shed by the colon. (Xu J. Gordon, JI, PNAS 2003. 100.)
- Entrenched "resident" bacteria are able to establish themselves by embedding in biofilms. (Sonnenburg, JL, Angenent, LT, Gordon, Jl. Nature Immunology 2004. 5.)

High throughput "omic" technologies used to *characterize*

- DNA (genomics)
- RNA (transcriptomics)
- Small molecules (metabolomics)



• Proportional abundance of various microbes in a bio-specimen and inference of their functions

Weight of Microbiome 3 LBS!

What shapes our microbiome?

Birth

Antibiotics

Probiotics

Where we live

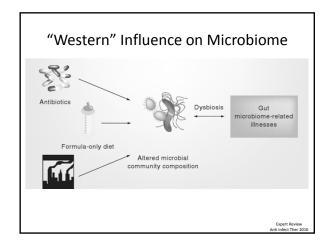
Travel

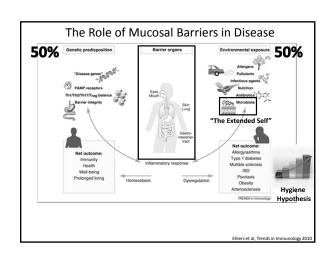
Diet

Exercise

PPIs

Malnutrition





A Seminal Article and a Thoughtful Editorial

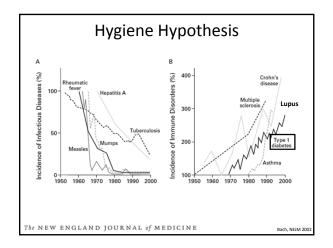
Review Article:

The Effect of Infections on Susceptibility to Autoimmune and Allergic Diseases

JF Bach, M.D., D.Sc NEJM 347(12), 2002

Editorial:

Eat Dirt – The Hygiene Hypothesis and Allergic Diseases ST Weiss, M.D. NEJM 347(12), 2002



Major Ancient Pandemics

- The Plague of Athens (430-427 BC)
- The Great Plague of Milan (1629-1631)
- The Third Pandemic (1855-1905)

The Third Pandemic (1855-1905)

- Began in Yunnan Province in 1855
- Spread to all inhabited continents
- Killed more than 12 million in China and India alone

Bubonic Plague or Black Death Yesinia pestis

The Big Question

FOE or FRIEND?

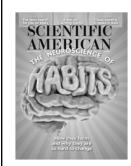
Microbiome and Obesity

- Studies in mice as well as in humans show that gut microbiota differ in composition between obese and lean subjects
- Notable difference in ratio of Bacteroidetes and Firmicutes, with a decrease in Bacteroidetes and a corresponding increase in Firmicutes

Obesity: Twin Studies Shed Light

- Four pairs of identical twins. One lean and the other obese in each pair.
- Genetically identical baby mice had their guts populated with intestinal microbes from either obese women or their lean twin sister.
- Mice that received bacteria from obese twin gained weight and had more body fat. They also had a less diverse community of gut microbes.

Gut Microbiota from Twins Discordant for Obesity Modulate Metabolism in Mice Ridaura, VK et al. Science, 6 September 2013: Vol. 341 no. 6150



How Gut Bacteria Help Make Us Fat and Thin By Claudia Wallis

Increased acetate production could be the culprit

- Acetate stimulates secretion of insulin by the pancreas in rodents
- Acetate infusion into the brain triggers insulin secretion by activating the parasympathetic nervous system
- Acetate also stimulates secretion of gastrin and ghrelin, leading to increased food intake

Continued: Gut Microbiota a Key Player?

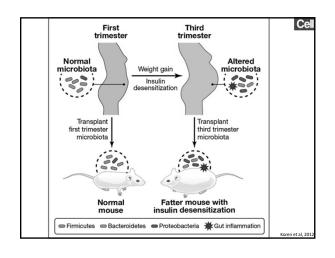
 Transfusion of fecal material from one group of rodents to another leads to changes in gut microbiota, acetate levels, and insulin secretion

R.J. Perry et al. Nature 534: 215-217, 2016.

Pregnancy, Weight Gain, and Altered Microbiota

- Gut microbiota changed dramatically from first to third trimesters, with vast expansion of diversity between mothers, an overall increase in Proteobacteria and Actinobacteria, and reduced richness.
- When transferred to germ-free mice, third trimester microbiota induced greater adiposity and insulin insensitivity compared to first trimester.

Host remodeling of the gut microbiome and metabolic changes during pregnancy Koren et al., Cell, 2012



Long Term Infant Outcomes are Influenced by Mode of Delivery

- Children born by Cesarean section are more likely to develop:
 - · Type 1 Diabetes
 - Celiac disease
 - · Hospitalization for Gastroenteritis
 - Asthma
 - · Allergic rhinitis

Neu, J and J Rushing. Cesarean versus Vaginal Delivery: Long term infant outcomes and the Hygiene Hypothesis. *Clin Perinatol*. 2001 June: 38(2): 321-331.

What is this condition?





Necrotizing Enterocolitis

- · A study of 1000 premature newborns
- Up to 6% incidence in premature newborns
- 36% mortality rate





Gut Microbes Linked to Necrotizing Enterocolitis in Premature Newborns

- Premature newborns who survive the first two weeks have a much higher risk of dying from Necrotizing Enterocolitis
- Gut microbiota in these newborns are different more Gram negative, less anaerobes
- This dysbiosis observed before any clinical event
- · Potential window for intervention

Warner, B.B. *et al.* Gut bacterial dysbiosis and necrotizing enterocolitis in very low birth weight infants: a prospective case control study. Lancet, March 2016.

The Microbiome affects Obesity and T2DM

- Obesity is associated with changes in the intestinal microbiota.
- The obese microbiome seems to be more efficient in harvesting energy from the diet.
- Differences in gut microbiota composition might function as early diagnostic markers for the development of T2DM.
- Butyrate, a product of intestinal microbes, may induce beneficial metabolic effects through enhancement of mitochondrial activity, prevention of metabolic endotoxemia, and activation of intestinal gluconeogenesis.

nsights into the Role of the Microbiome in Obesity and Type 2 Diabetes. Hartstra et al. Diabetes Care 2015, 38(1).

Bariatric Surgery Affects Gut Microbiome Composition

- Gut microbiomes different in patients after gastric bypass surgery compared to obese controls
- E. coli, Klebsiella, Pseudomonas, all more common
- Stool from gastric bypass patients transferred to germ free mice:
 - Mice had improved fat oxidation
 - Microbe trasplanted mice gained 43% less body fat compared to mice that received stool from gastroplasty patients

Tremaroli, V. et al. Cell Metabolism, 22, 2015.

Diets Modulate Gut Microbiome and Improve Insulin Sensitivity

- 20 obese men with CAD. Randomized to Mediterranean diet (Med) or low-fat, high-complex carbohydrate diet (LFHCC) for one year
- Bacterial composition and relationship with fecal and plasma metabolome evaluated
- Both diets shown to exert a predictive effect on the development of T2DM. Med diet increased Roseburia genus and LFHCC diet increased F. prausnitizii.

Haro, C et al. Endocrinol Metab 101:233-242, 2016

Kids Get Arthritis

- 300,000 kids in USA have some form of arthritis -Juvenile Rheumatoid Arthritis (JRA) is most common
- Kids exposed to antibiotics have high risk of getting IRA
- Risk highest when exposed to multiple courses of antibiotics
- Risk not seen with exposure to antivirals or antifungals

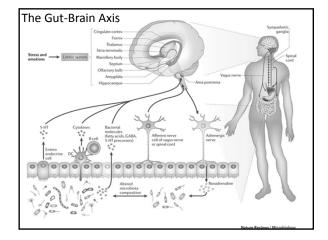
Rheumatoid Arthritis: Expansion of rare lineage gut microbes

- Expansion of Eggerthella and Collinsella
- Decrease of beneficial genera- Faecalibacterium
- Collinsella increases gut permeability and induces production of (IL) 17A, a proinflammatory cytokine
 - Chen J et al. An expansion of rare lineage intestinal microbes characterizes
 rheumatoid arthritis. Genome Medicine 2016 8:43

Is there a Gut-Brain connection?

- The immune system is active in myelin destruction.
- Alteration in gut flora in mice leads to a MS-like disease, called Experimental Autoimmune Encephalomyelitis (EAE)
- Preliminary work suggests that the gut in MS patients contained bugs that drive inflammation and are low in the types of the bacteria that control inflammation.*

*Consistent with work in IBD and RA



"The E. coli made me do it!"

- May 2000, heavy rains pummel Walkertown, Ontario
- Epidemic of *E. coli* ensues. Half the residents get ill and seven individuals die.
- Two years later, question posed: "Since the outbreak, have you developed depression, anxiety, panic disorder, or PTSD?"
- · Exposed residents more likely to answer YES!

"The E. coli made me do it!"

- Gut organisms capable of making neurotransmitters such as norepinephrine and serotonin.
- Mice fed Lactobacillus alters swimming mobility (shorter times suggesting less hopelessness).
- If the vagus nerve is severed, the mice become hopeless again.

An Important Reference

Ingestion of *Lactobacillus* strain regulates emotional behavior and central GABA receptor expression in a mouse via the vagus nerve Bravo *et al.*, *PNAS*, 2011, 108(38).

There is cross-talk between Gut Microbiota and Brain Function

- In germ free mice, restraint leads to exaggerated ACTH response.
- Above response partially reversed by colonization with fecal material from controls.
- Normal mice subjected to psychosocial stress have a decrease in Bacteroides and increase in Clostridia in the cecum.
- In humans, probiotics may modulate brain activity
- Bariatic surgery followed by profound changes in microbiota and improvement in memory.

Study Protocol

- 20 obese patients, 19 lean subjects
- Obese patients had decrease in bacterial biodiversity
- Subjects with the highest gut microbial diversity had changes in hypothalamus, hippocampus, and cordate nucleus
- Subjects with greater abundance of Actinobacteria had better motor speed and attention

Conclusions

The human Gut Microbiota profile is significantly associated with Brain microstructure and function.

Fernandez-Real, JM *et al.*, Gut Microbiota Interacts with Brain Microstructure and Function. J Clin Endocrinol Metab 100: 4505-4513. 2015.

Microbes can play games with the mind

- Bacteria in our guts may decide who gets anxiety and depression
- Pilot study in 22 men, subtle but definite brain benefits by taking capsules brimming with bacteria.
- Reported at annual meeting for Society of Neuroscience
- Different bacteria make neural messengers such as dopamine, norepinephrine, GABA, serotonin, acetylcholine

Do gut microbes influence severity of **stroke**?

- · Two colonies of mice:
 - Group A: gut bacteria resistant to antibiotics
 - Group B: gut bacteria vulnerable to antibiotics
- Cerebral arteries of the mice occluded, resulting in ischemic stroke
- Brain damage 60% smaller in Group B mice

Do gut microbes influence severity of **stroke**?

- Fecal transplant from mice that had reduced brain damage given to naïve mice who now developed an altered gut microbiome (were not given antibiotics)
- Cerebral arteries occluded, leading to ischemic stroke again, significant protection against brain damage seen
- Composition of immune cells "good" regulatory T cells and "bad" Gamma Delta T Cells altered
- "These cells determine what kind of inflammatory immune response the brain experiences after a stroke."

Constantino ladecola

Benakis, C. *et al.* Commensal microbiota affects ischemic stroke outcome by regulating intestinal $\gamma\delta$ T cells. Nat. Med. 2016.

Caveat: This is a study in mice.

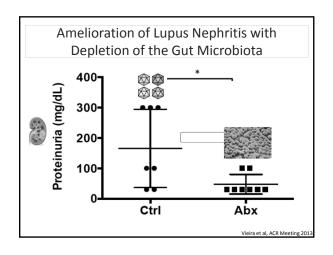
Gut Dysbiosis Impairs Recovery after Spinal Cord Injury (SCI)

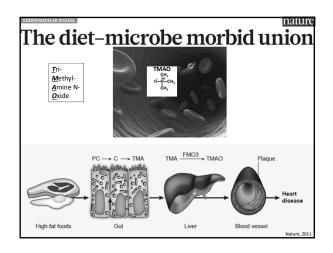
- SCI increases intestinal permeability and bacterial translocation from the gut.
- These changes associated with immune cell activation.
- Changes in gut microbiota persist for at least one month and predict the magnitude of locomotor impairment.

Gut Dysbiosis Impairs Recovery after Spinal Cord Injury (SCI) Part Two:

- Experimental of gut dysbiosis in naïve mice before SCI worsens neurological impairment and spinal cord pathology.
- Feeding SCI mice commercial probiotics triggers a predictive immune response and confers neuroprotection.

Kigeri K A. et al Journal of Experimental Medicine October 2016.





TMAO, Microbiome, and CAD/Stroke!

- TMAO is produced when intestinal bacteria digest the nutrient Lecithin.
- Human subjects, after eating two hard-boiled eggs and a capsule of labelled Lecithin, have an increase in TMAO levels.
- However, when subjects are given broad-spectrum antibiotics, their TMAO levels are suppressed.
- High TMAO blood levels are associated with higher risk of heart attack or stroke, independent of other risk factors and other blood test results.





TMAO Important Predictor of Atherosclerosis

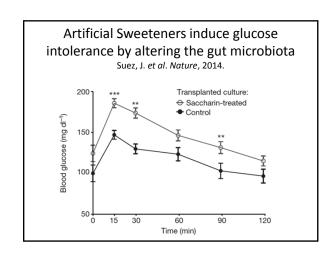
- Dietary meat is a major source of TMAO in humans
- TMAO levels independent risk factor for atherosclerosis in humans
- In Apo E deficient mice, TMAO levels correlate with atheroma burden
- When Apo E deficient mice are treated with antibiotics, there is a significant decrease in atheroma burden

Diet is destiny!

Fate is shaped by genome and microbiome.

An Important Reference

Intestinal Microbial Metabolism of Phosphatidylcholine and Cardiovascular Risk. Wilson Tang *et al.*, *NEJM* 2013, 368.



"Dis-moi ce que tu manges, je te dirai ce que tu es."

Tell me what you eat and I will tell you what you are.

Anthelme Brillat-Savarin, 1826

Preliminary work shows...

- Vegan diet
- · High fiber diet
- Highly fermented foods (Kimchi)
- · Breast milk

All have a favorable influence on the composition of the gut microbiome.

Gut Microbiota and Autism Spectrum Disorder

- Pregnant mice injected with artificially created virus-like DNA
- Offspring display less socialization, greater sense of being startled by sounds, fewer vocalizations
- Serum of ASD mice contain more than 45x the amount of 4-ethylphenyl sulfate (4-EPS) (a metabolite of gut bacteria)

Gut Microbiota and Autism Spectrum Disorder

- Children with ASD have high concentrations of a similar compound p-cresol in their urine
- Healthy mice injected with 4-EPS have a leaky gut and display ASD symptoms
- Probiotic treatment with B. fragilis in ASD mice restored intestinal permeability and 4-EPS levels returned to normal

Hsiao, E.Y. *et al.*, Presentation at Gut Microbiota for Health. World Summit. Barcelona, Spain. 2015.

Gut Microbes Are Essential

- · Not free-loaders
- · Digest food
- Produce anti-inflammatory chemicals and compounds
- Guide the Immune System to distinguish friend from foe

Basic Definitions

- <u>Prebiotics</u> are selectively fermented products that confer changes in the composition and/or activity of the GI tract microflora and confer health benefits
- <u>Probiotics</u> are ingested microorganisms that are associated with health benefits

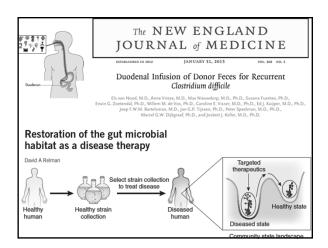
Global sales of probiotics estimated to exceed **\$31 billion** in 2015

More than a Century ago...

"The dependence of the intestinal microbes on the food makes it possible to adopt measures to modify the flora in our bodies and to replace the harmful microbes by useful microbes."

Elie Metchnikoff (1907) Nobel Laureate

Therapeutic Implications



Frozen Poopsicle!

- 14 of 20 with recurrent C. diff diarrhea resolved
- 4 of 6 non-responders' diarrhea resolved with retreatment
- No adverse effects noted in this small study

Oral, Capsulized, Frozen Fecal Microbiota Transplantation for Relapsing *Clostridium difficile* Infection. Youngster *et al.*, *JAMA*, 2014, 212(17)

Treatment of C.diff

Conclusions Fecal microbiota transplantation using colonoscopy to infuse feces was significantly more effective than vancomycin regimen for the treatment of recurrent *C. difficile* infection. The delivery of donor feces via colonoscopy has the potential to optimize the treatment strategy in patients with pseudomembranous colitis.

Cammarota G. et al Aliment Pharmacol Ther. 2015;41(9):835-843.

The No. 1 treatment for C.diff may well be No. 2

A Case Study in Humans

- 32 year old woman with recurrent CDI receives FMT.
- Baseline weight was 136 lbs (BMI 26).
- Patient receives FMT from 16 year old daughter who weighed 140 lbs (later it increased to 170 lbs).
- 36 months after FMT, the patient's weight increased to 177 lbs (BMI 34.5).

Weight Gain After Fecal Microbiota Transplantation.

Alang, N, and CR Kelly. OFID, 2015.

Probiotics Beneficial in Hepatic Encephalopathy (HE)

- Daily intake of a probiotic, VSL #3, over 6 months significantly reduced the risk of hospitalization for HE in cirrhotics
- Patients given VSL #3 had improvement in Childs Pugh and MELD scores

R.K. Dhiman et al., Gastroenterology 2014; 147: 1327-1337.

The Gut Microbiota and NAFLD

- Microbiota in addition to regulating body fat gain and insulin resistance:
 - Change gene expression
 - Increase energy harvest from diet
 - Produce ethanol
 - · Affect inflammation and immunity

E Lau et al. Gut Microbiota: Association with NAFLD and Metabolic Disturbances. Biomedical Research International. 2015.

Let's Summarize

- 1. Microbes appeared 3.5 billion years ago.
- 2. There's a cross talk! They've been doing it for billions of years. Yet to be silent!
- 3. Gut microbiota affect:

Obesity Diabetes
Cardiovascular Disease Colon Cancer

Neurological Disease IBD Arthritis IBS

Necrotizing Enterocolitis Longevity and more...

Let's Summarize (continued)

- 4. Composition of the microbiota is in flux and altered by a diverse array of factors
- 5. Microbiota-directed therapies include:
 - Fecal transplants (C. difficile colitis)
 - Hepatic Encephalopathy
 - And more...

Have you heard of Biological Dark Matter?

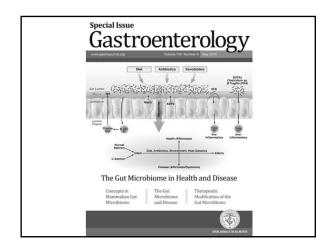
- It is uncategorized genetic material found in humans that does not fall under the three existing domains of life: bacteria, archaea, and eukaryotes.
- It may well be a fourth domain of life yet to be discovered.
- Biological dark matter accounts for:
 - 40-50% of the genetic material in the human gut
 - 20% of the genetic material in the nose
 - 1% of genetic material in sterile blood

Worth listening to...

"What's left to explore?"

Nathan Wolfe, biologist and explorer gave a terrific TED talk in 2015 about biological dark matter.

http://www.ted.com/talks/nathan wolfe what s left to explore





An African Proverb





We have only seen the head of the hippo.