

# **Factors that Modify the Microbiome in Perinatal and Neonatal Periods**

Josef Neu, M.D.

University of Florida

# Agenda

- **The Past**

- Human Microbiome Project and Intestinal Microbe Host Interactions

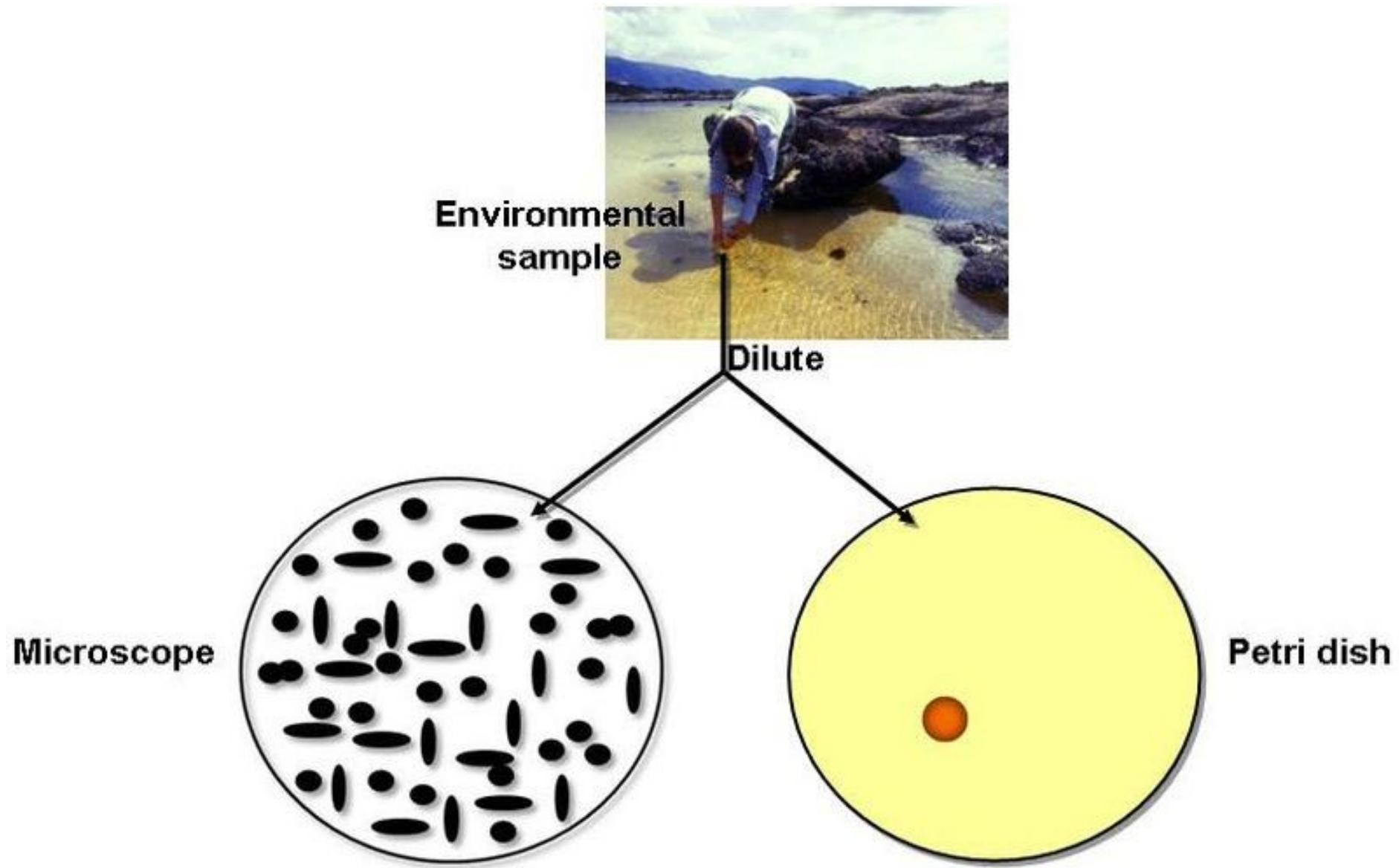
- **Recent Studies**

- Preterm Birth.
- Mode of Delivery Effects
- Environmental Effects:
  - Antibiotics
- Microbes in Human Milk

- **The Future**

- Multi-omics
- Host Microbial Interactions

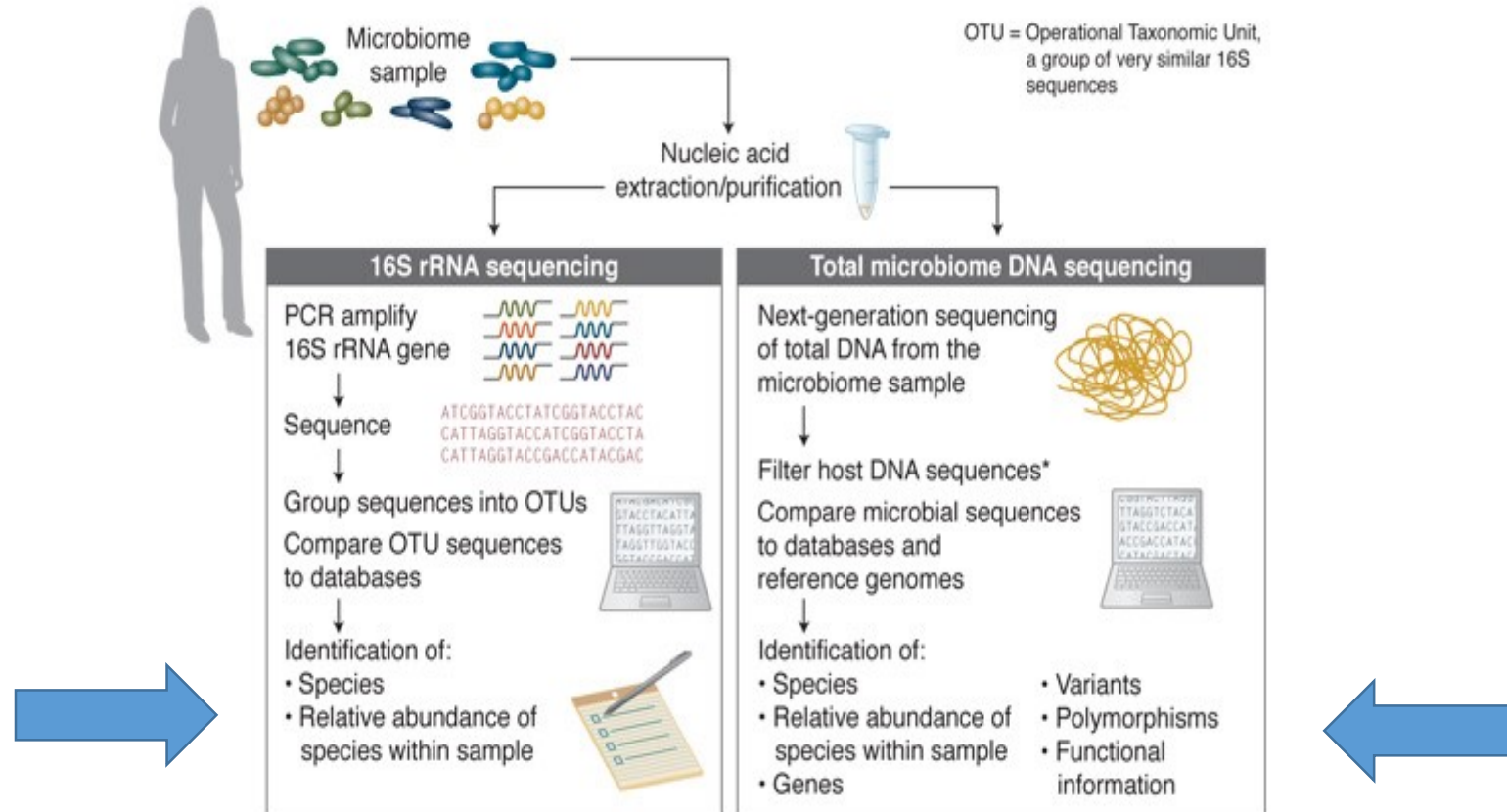




**“THE GREAT PLATE COUNT ANOMALY”**

**~100 times > cells than colonies, 99% unculturable**

# Major Sequencing Technologies



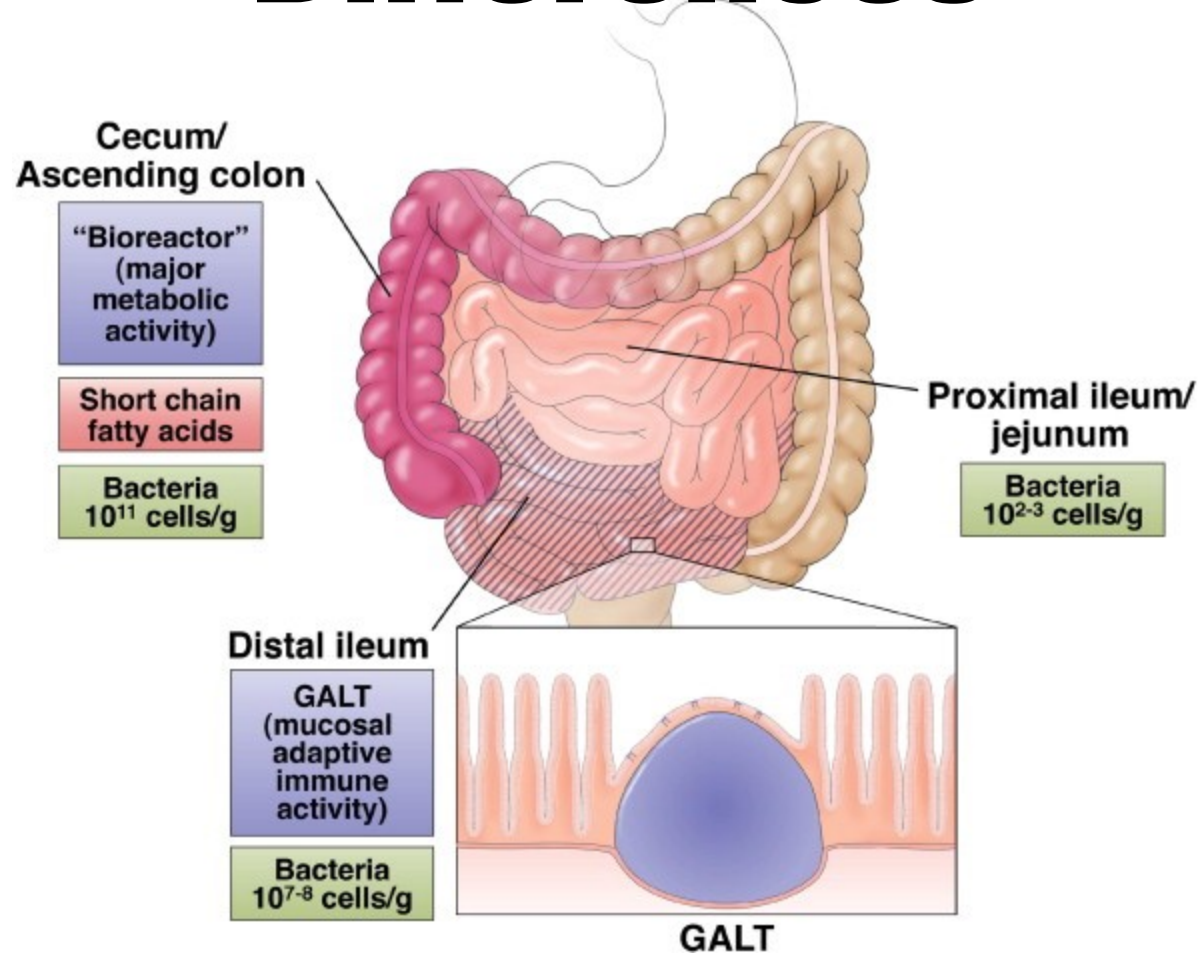
Adapted from : <https://www.neb.com/tools-and-resources/feature-articles/addressing-challenges-in-microbiome-dna-analysis>

# Holobiont

A “super or meta -organism” consisting of an assemblage of different species that form ecological units. It adjusts and transforms itself according to environmental changes causing evolution of the entire entity.

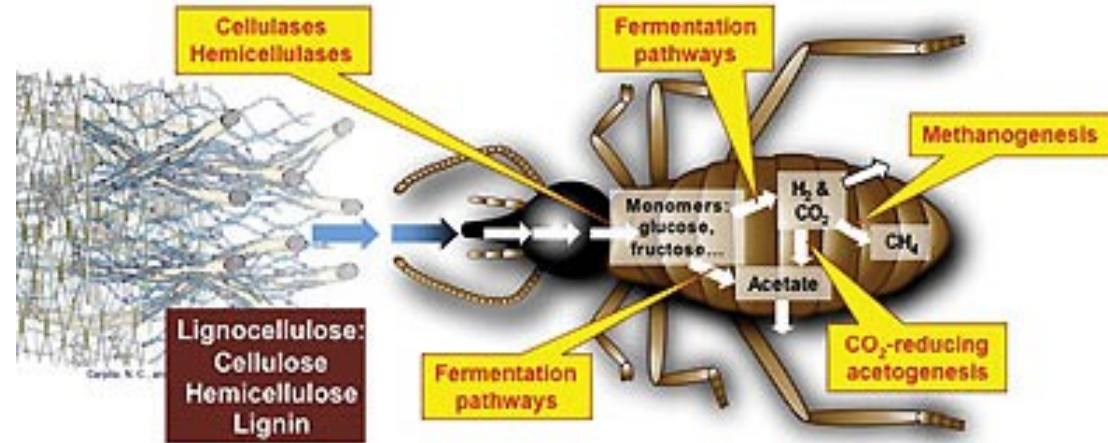


# The Gut: Regional Microbial Differences



Neish, A. Gastroenterology vol. 136, No. 1, 2009

# Mechanisms: Metabolomic (Bioreactor) Role



WWW.APPLE.COM/GETAMAC/



Gut bacteria and obesity

“Holy shit!”

Nov 12th 2009

From *The Economist* print edition

A new way of finding out how diet affects gut microbes

Roy C. Nutr. Clin Practice 21;351:2006

Turnbaugh, PJ Cell Host Microbe Volume 3, Issue 4, 17 April 2008, Pages

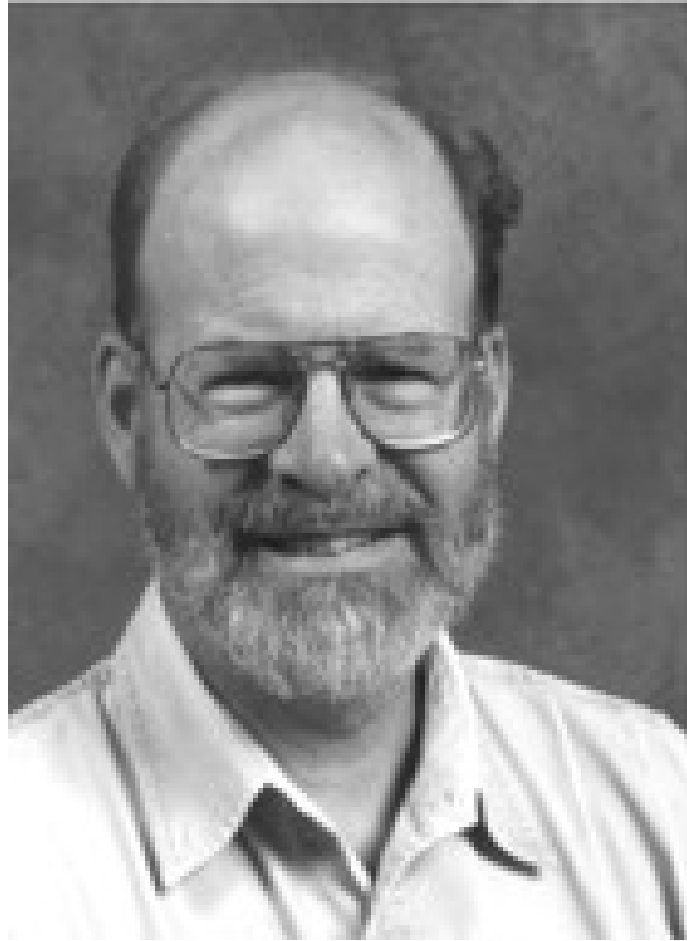
213-223

# “TOLL” Proteins-1985



Christiane Nüsslein-Volhard

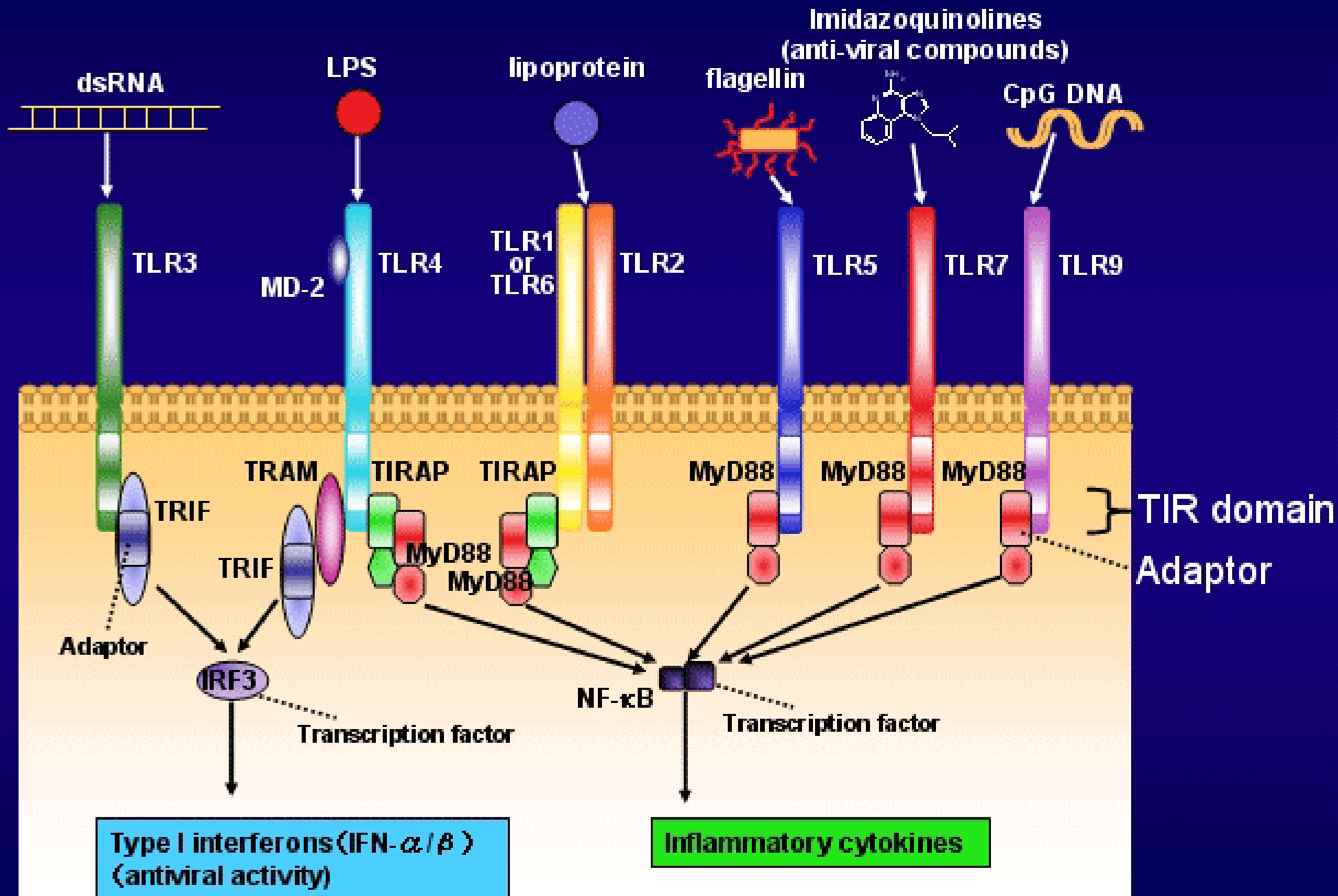
# “Toll Receptors”-1996



© 1996 MIT/Johns Hopkins

**"Father of Innate Immunity," Charles Janeway**

# Toll-like receptors: ligands and signaling pathways



# Recognition of Commensal Microflora by Toll-Like Receptors Is Required for Intestinal Homeostasis

Seth Rakoff-Nahoum,<sup>1</sup> Justin Paglino,<sup>2</sup>  
Fatima Eslami-Varzaneh,<sup>3</sup> Stephen Edberg,<sup>2</sup>  
and Ruslan Medzhitov<sup>1,\*</sup>

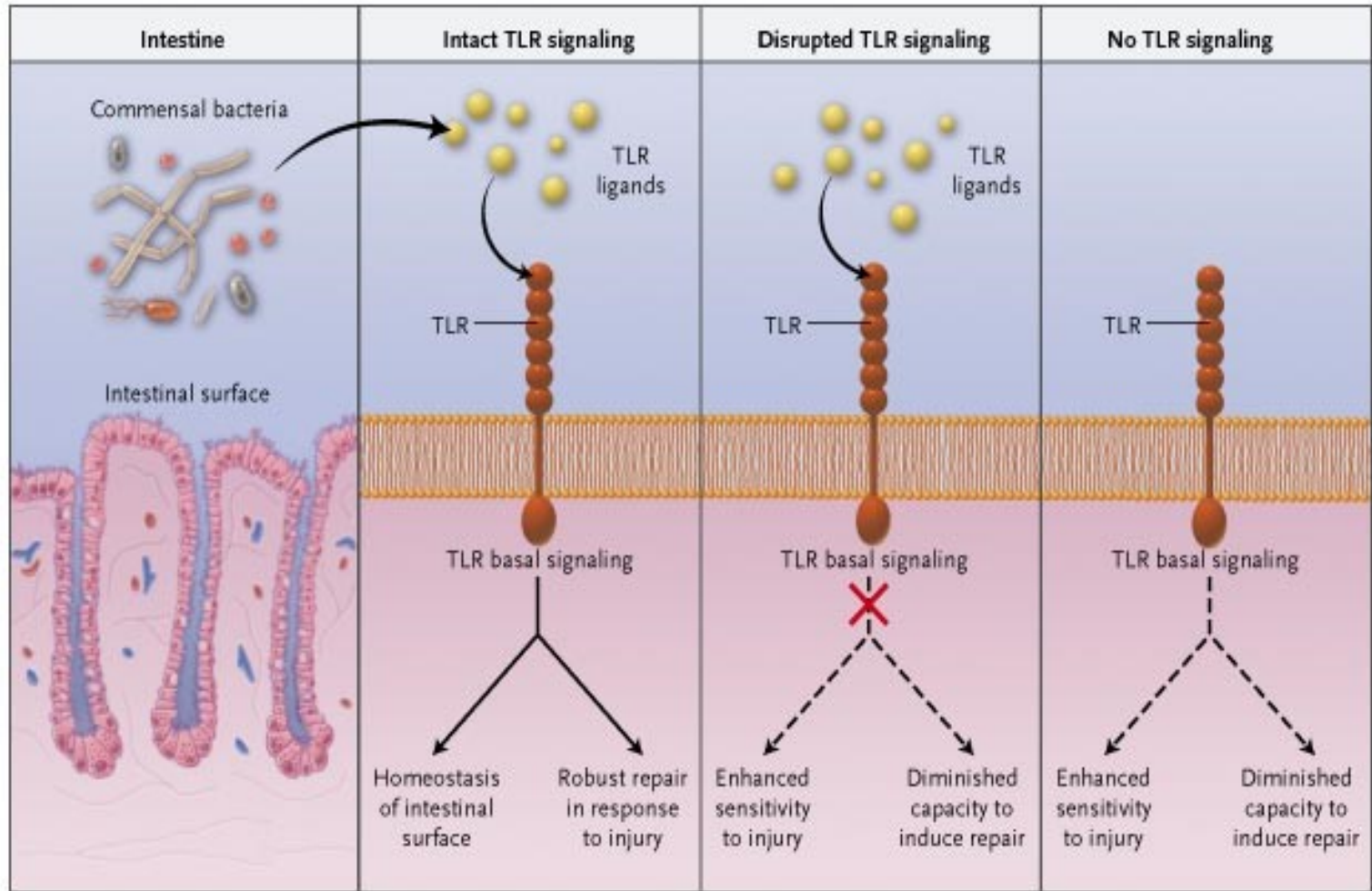
<sup>1</sup>Howard Hughes Medical Institute  
and Section of Immunobiology

<sup>2</sup>Department of Laboratory Medicine

<sup>3</sup>Department of Pathology  
Yale University School of Medicine  
New Haven, Connecticut 06510

by commensal bacteria (Farrell and LaMont, 2002; Sartor, 2000). The nature and regulation of the host-commensal interactions in the gut is thus an area of intense scientific and clinical interest.

TLRs comprise a family of pattern-recognition receptors that detect conserved molecular products of microorganisms, such as lipopolysaccharide (LPS) and lipoteichoic acid (LTA), recognized by TLR4 and TLR2, respectively (Takeda et al., 2003). TLRs function as sen-

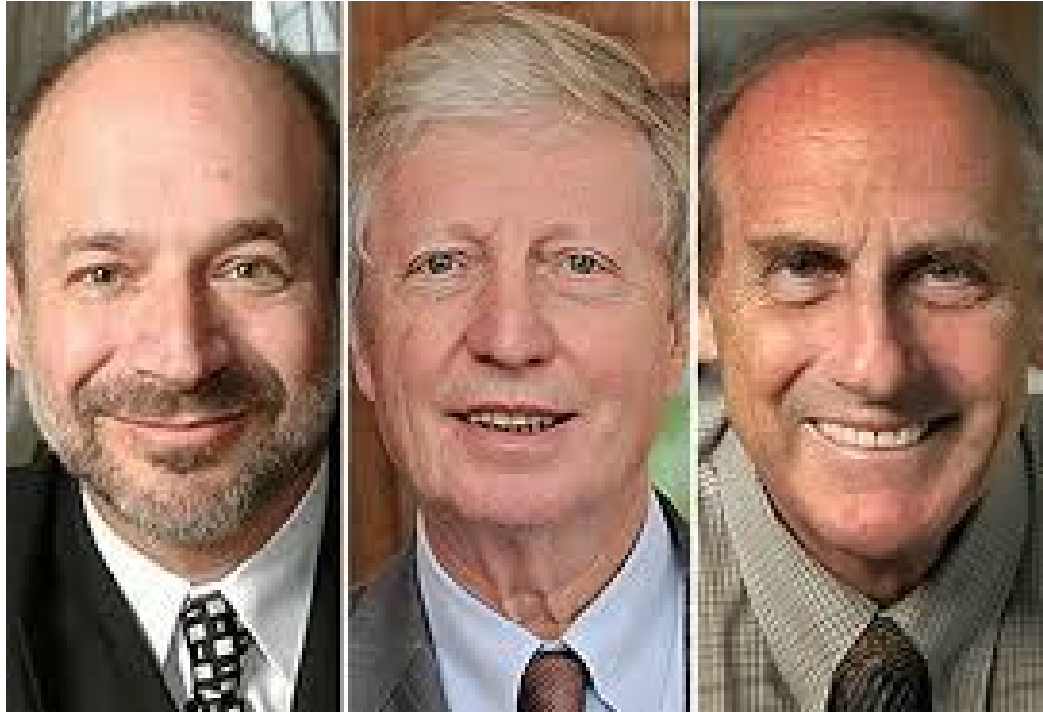


From Madara J. Building an intestine – architectural contributions of commensal bacteria. *New Engl. J. Med.* 2004; 351: 1685-86.

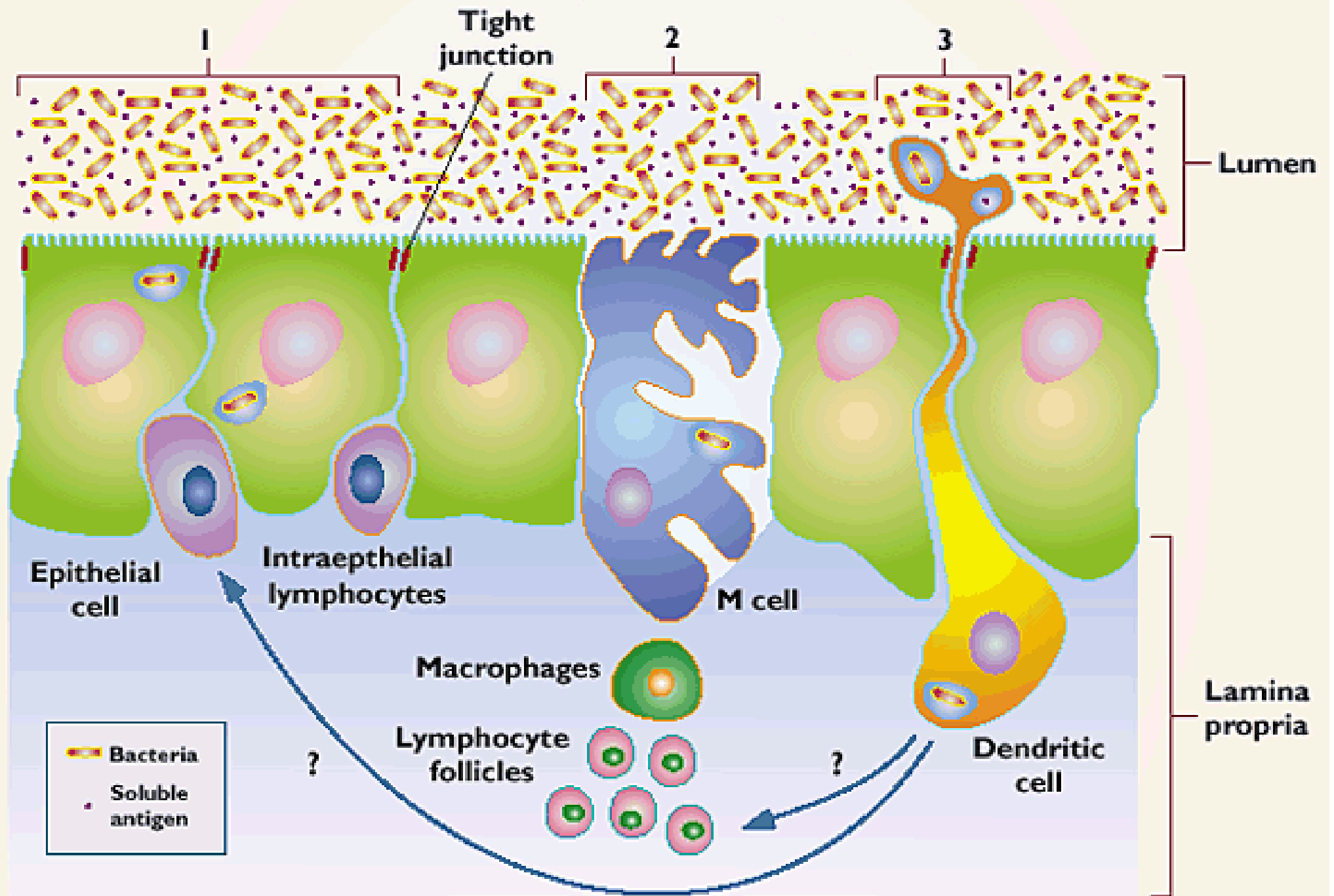
# Lesson

Low grade stimulation (“tickling”) of toll receptors can prevent high grade inflammation and intestinal damage and promotes intestinal homeostasis.

# The Dendritic Cell and its Role in Adaptive Immunity



Bruce Beutler,  
Jules Hoffman and  
Ralph Steinman



# Questions

## **Amniotic fluid is sterile?**

- True
- False

## **Placenta is sterile?**

- True
- False

## **Meconium is sterile?**

- True
- False

## **Human Milk is sterile?**

- True
- False



Most infants first  
come into contact with  
microbes during birth —  
or so researchers have  
assumed.

A “Scientific Knife Fight”

Is it all “Fake News”?

# Baby’s first bacteria

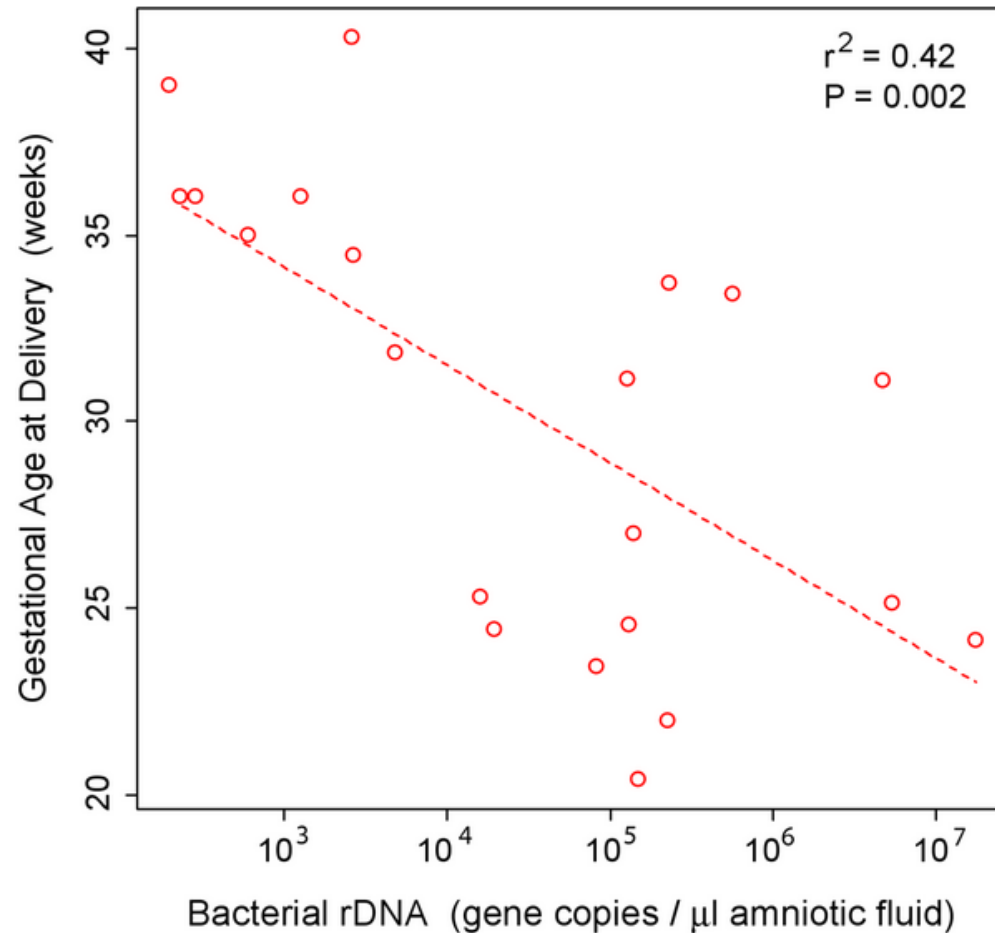
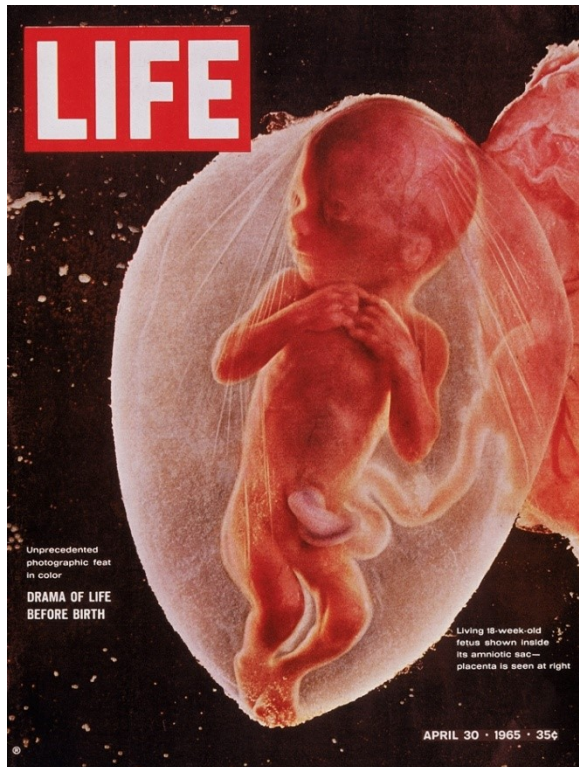
THE WOMB WAS THOUGHT TO BE STERILE. SOME SCIENTISTS  
ARGUE IT’S WHERE THE MICROBIOME BEGINS.

Soon after conception, a human embryo begins to assemble a remarkable organ crucial to its survival. The placenta is both a lifeline and a guardian: it shuttles oxygen, nutrients and immune molecules from the mother’s blood *By Cassandra Willyard* nearly 200 placentas collected from women giving birth at a hospital in St Louis, Missouri. When the researchers examined the samples under a microscope, they found bacteria in nearly one-third of them! “They were actually inside cells there?”

DOO/REUTERS

Willyard. C. Nature: Jan 18, 2018

# Is there a Fetal Microbiome? Might this Relate to Prematurity



MICROBIOME

# The Placenta Harbors a Unique Microbiome

Kjersti Aagaard,<sup>1,2,3\*</sup> Jun Ma,<sup>1,2</sup> Kathleen M. Antony,<sup>1</sup> Radhika Ganu,<sup>1</sup> Joseph Petrosino,<sup>4</sup> James Versalovic<sup>5</sup>

Humans and their microbiomes have coevolved as a physiologic community composed of distinct body site niches with metabolic and antigenic diversity. The placental microbiome has not been robustly interrogated, despite recent demonstrations of intracellular bacteria with diverse metabolic and immune regulatory functions. A population-based cohort of placental specimens collected under sterile conditions from 320 subjects with extensive clinical data was established for comparative 16S ribosomal DNA-based and whole-genome shotgun (WGS) metagenomic studies. Identified taxa and their gene carriage patterns were compared to other human body site niches, including the oral, skin, airway (nasal), vaginal, and gut microbiomes from nonpregnant controls. We characterized a unique placental microbiome niche, composed of nonpathogenic taxa distinct from those found in other human body sites.

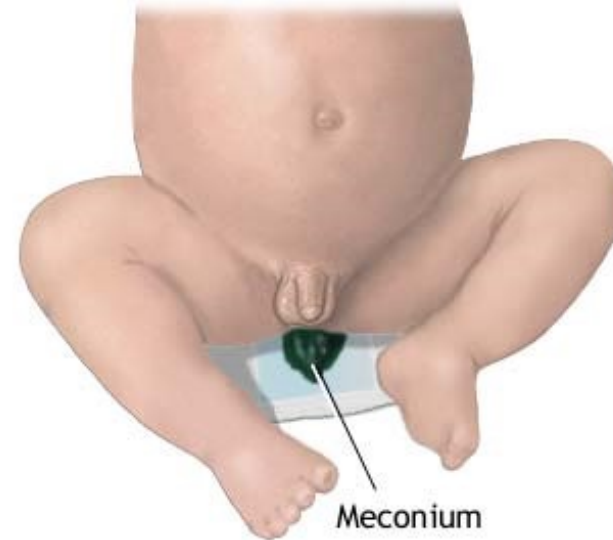
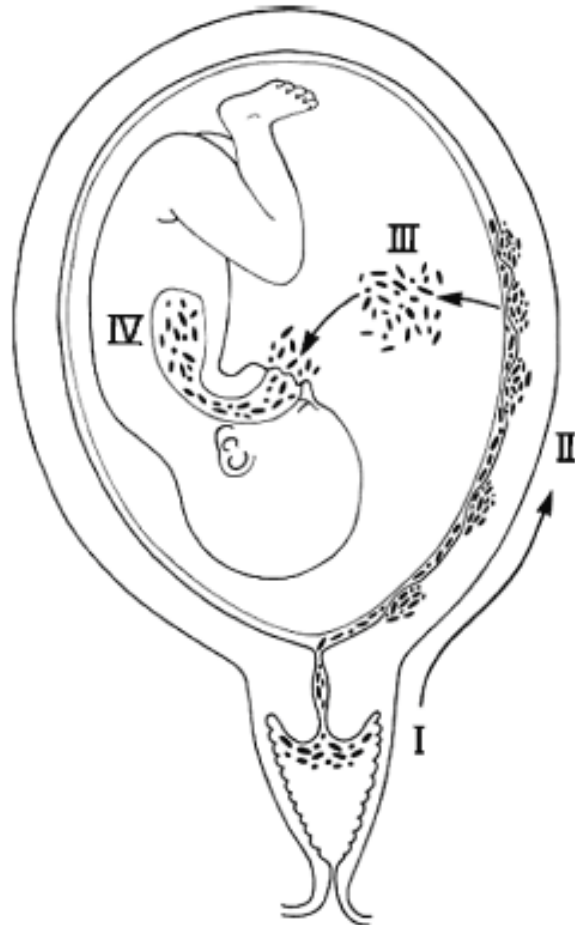
These findings reveal associations of the placental microbiome with a remote site, the oral cavity, and suggest a role for the placental microbiome in fetal development. This study provides the first comprehensive characterization of the placental microbiome and its relationship to other human body site niches.

## INTRODUCTION

# How Do the microbes get into the fetus? What are the implications?

Medscape®

www.medscape.com



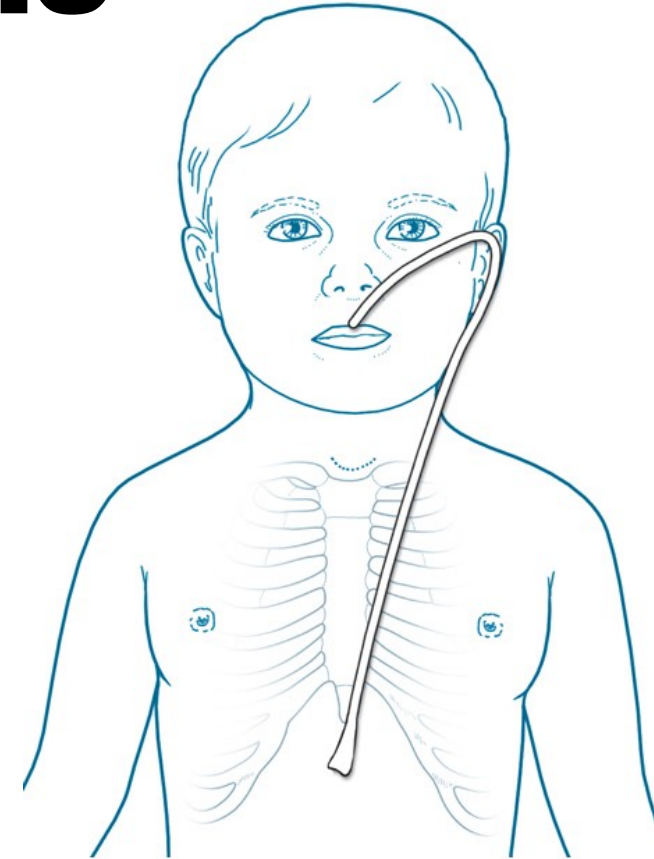
Over 50% of the microbes in meconium are also found in Amniotic Fluid suggesting fetal swallowing of colonized amniotic fluid.

Source: Semin Reprod Med © 2007 Thieme Medical Publishers

Ardissone, et al. PLOS One 2014

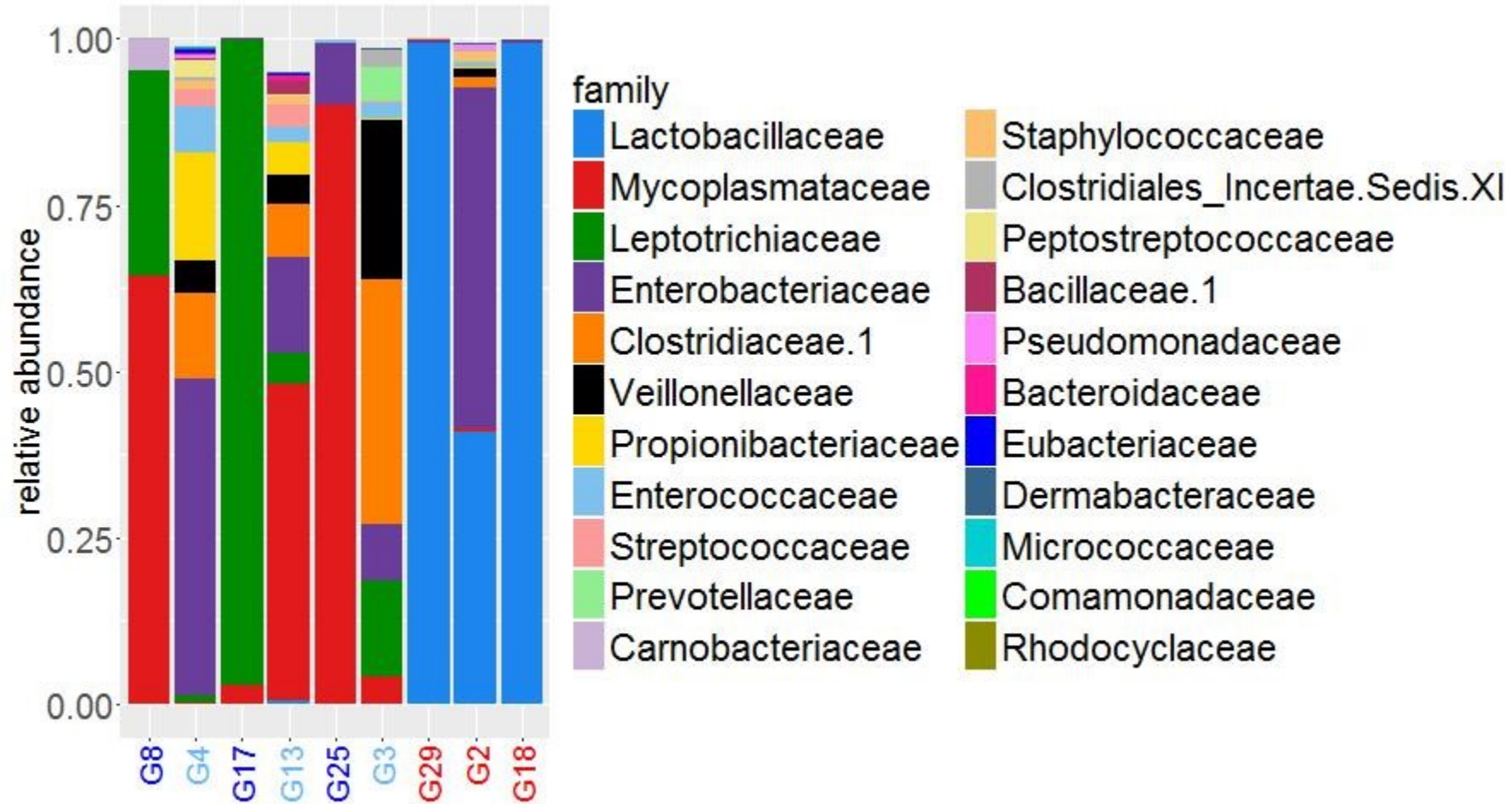
# First Gastric Aspirate Microbiome

- Do first gastric aspirates as a reflection of amniotic fluid differ in term versus preterm infants?
- How do these compare to microbes present in other niches such as vagina or GI tract?
- Could these microbes relate to inflammation of the GI tract and preterm birth?



Source: Goodman DM, Green TP, Unti SM, Powell EC: *Current Procedures: Pediatrics*: [www.accesspediatrics.com](http://www.accesspediatrics.com)  
Copyright © The McGraw-Hill Companies, Inc. All rights reserved.

# Gastric Aspirates: Term versus Preterm



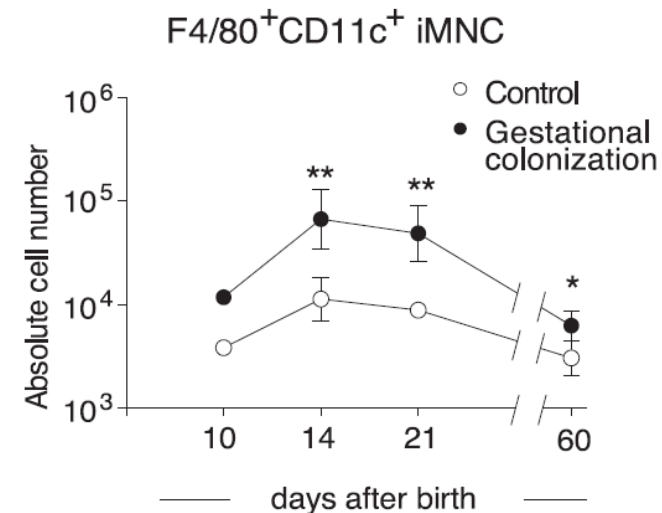
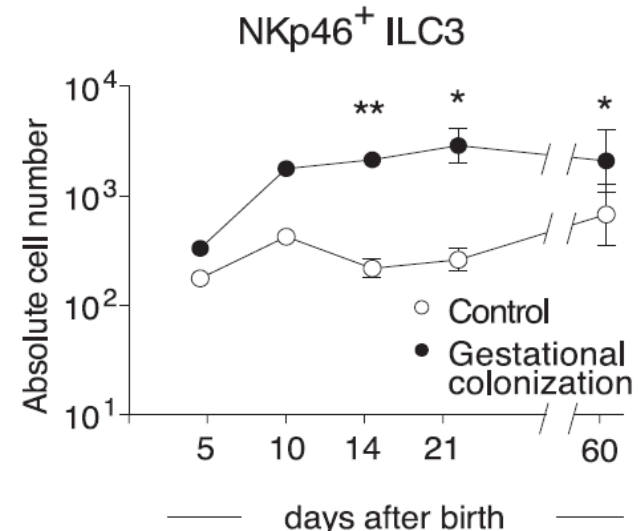
Bajorek, S. et al. Under Revision

# **Sterile Womb Hypotheses: the Controversy**

- Are the microbes found in the womb simply a “kitome”? U. Penn Group found from 6 placentas that using strict controls from the extraction kits that the microbes present in their studies were derived from the kits( Lauder, et al. Microbiome, 2016).
- Sheep fetuses have a placental and intestinal microbiome (where microbes can be cultured) and the placental appears to act as a conduit during times of stress that allow microbes to reach the brain (Zarate, M.A. Sci Rep. 2017 Jul 25;7(1)).
- “If humans do not have an in-utero microbiome, we would be the only species that doesn’t” (Susan Lynch, UCSF).
- Roberto Romero from NIH and other colleagues are planning to answer this question more definitively.

# Mouse Study: Maternal Intestinal Microbiota Drives Early Postnatal Innate Immune Development

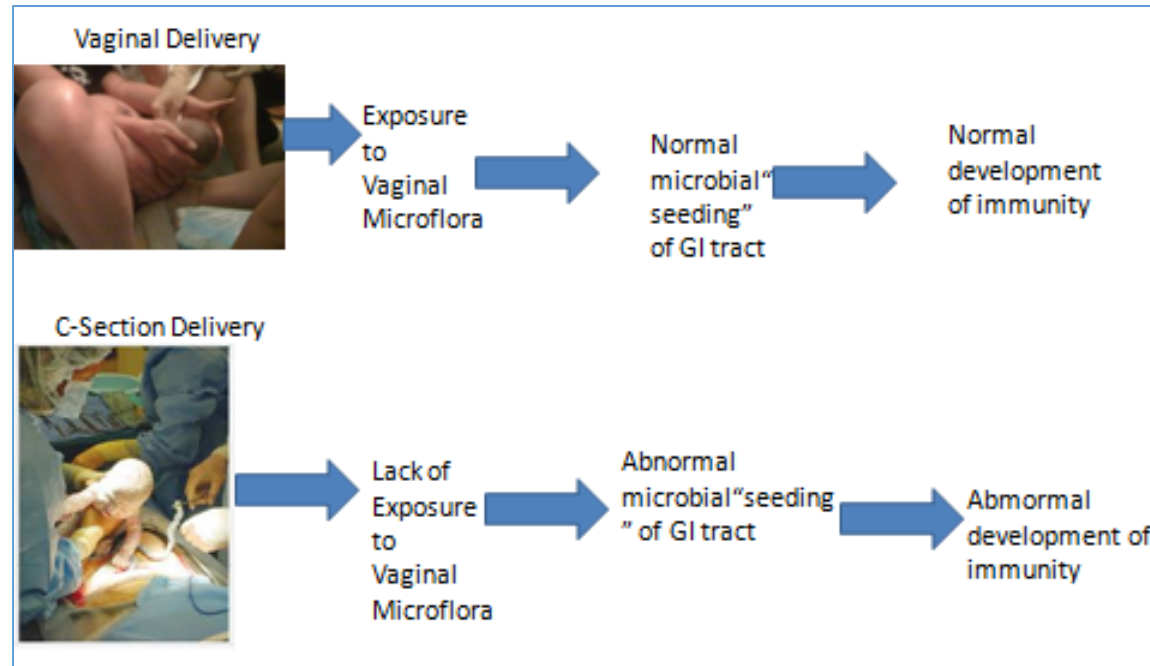
- Pregnant germ free mice colonized with *E. coli* HA 107 (strain does not persist in the intestine).
- Compared inoculated to non-inoculated offspring intestinal innate immune systems.



# Epidemiology Studies of C-section vs. Vaginal Delivery

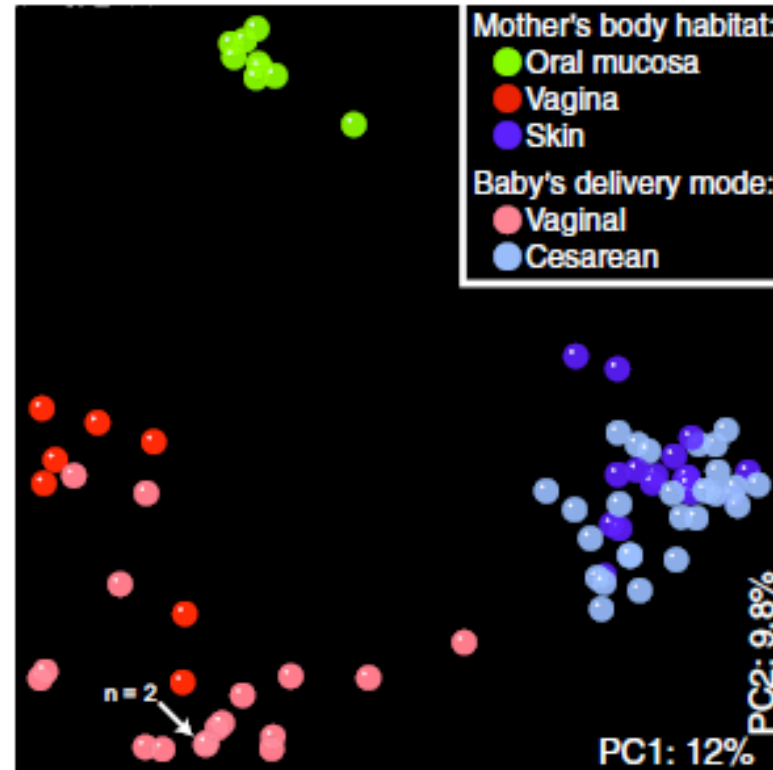
Higher odds of:

- Allergic Rhinitis
- Asthma
- Celiac Disease
- Type 1 Diabetes
- Gastroenteritis
- Allergies
- Obesity



\*Neu J, Rushing, J.  
Clinics in Perinatology 2011

# Vaginal vs. C-section Delivery: Non-Culture Technique



Meconium obtained  
<24 hours after  
Delivery.

Vaginally delivered infants acquired bacterial communities resembling their own mother's vaginal microbiota and C-section infants harbored bacterial communities similar to those found on the skin surface. (Dominguez-Bellow, MG. Proc Natl Acad Sci U S A. 2010 Jun 29;107(26):11971-5)



## Partial restoration of the microbiota of cesarean-born infants via vaginal microbial transfer

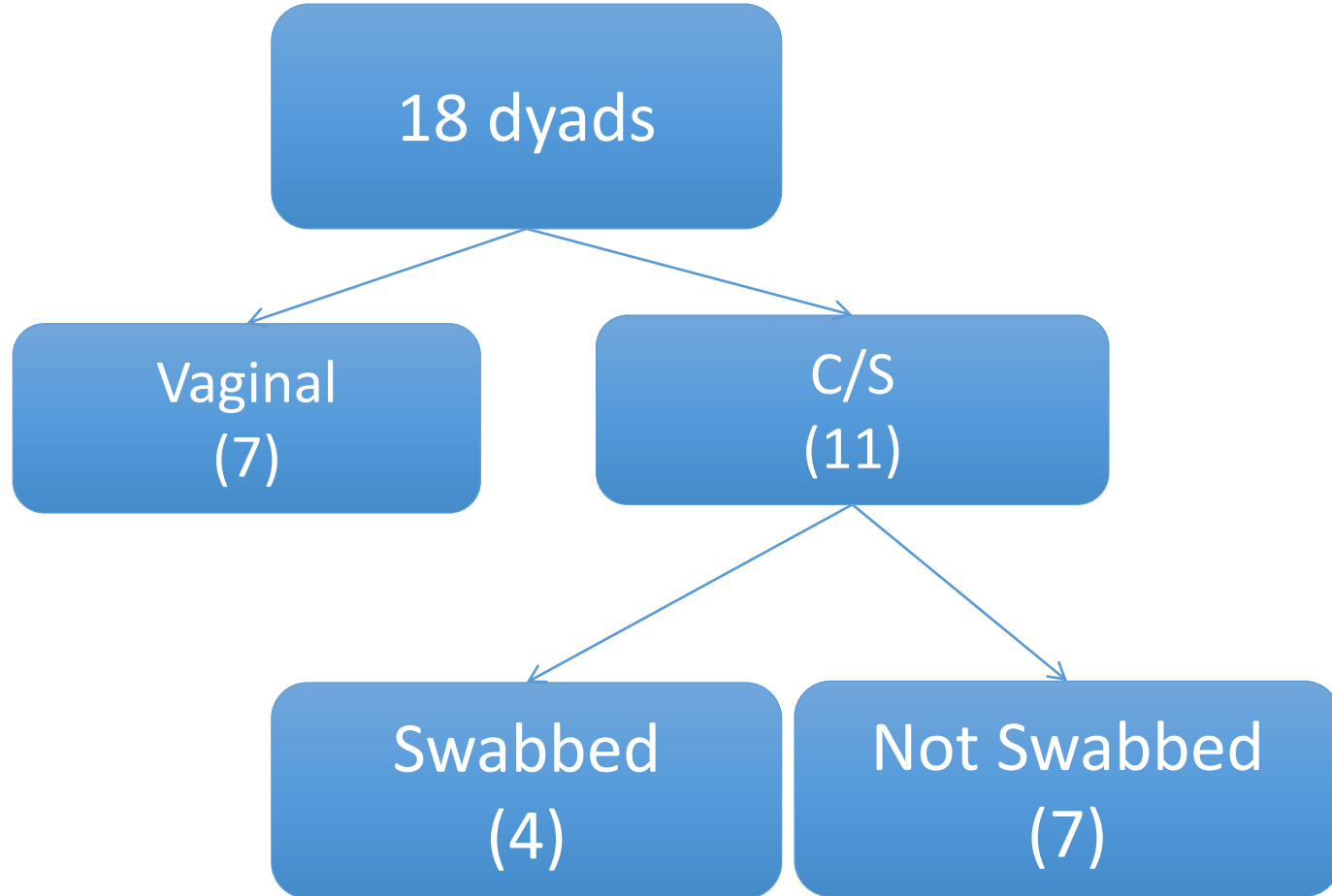
[Maria G Dominguez-Bello](#), [Kassandra M De Jesus-Laboy](#), [Nan Shen](#), [Laura M Cox](#), [Amnon Amir](#), [Antonio Gonzalez](#), [Nicholas A Bokulich](#), [Se Jin Song](#), [Marina Hoashi](#), [Juana I Rivera-Vinas](#), [Keimari Mendez](#), [Rob Knight](#) & [Jose C Clemente](#)

[Affiliations](#) | [Contributions](#) | [Corresponding authors](#)

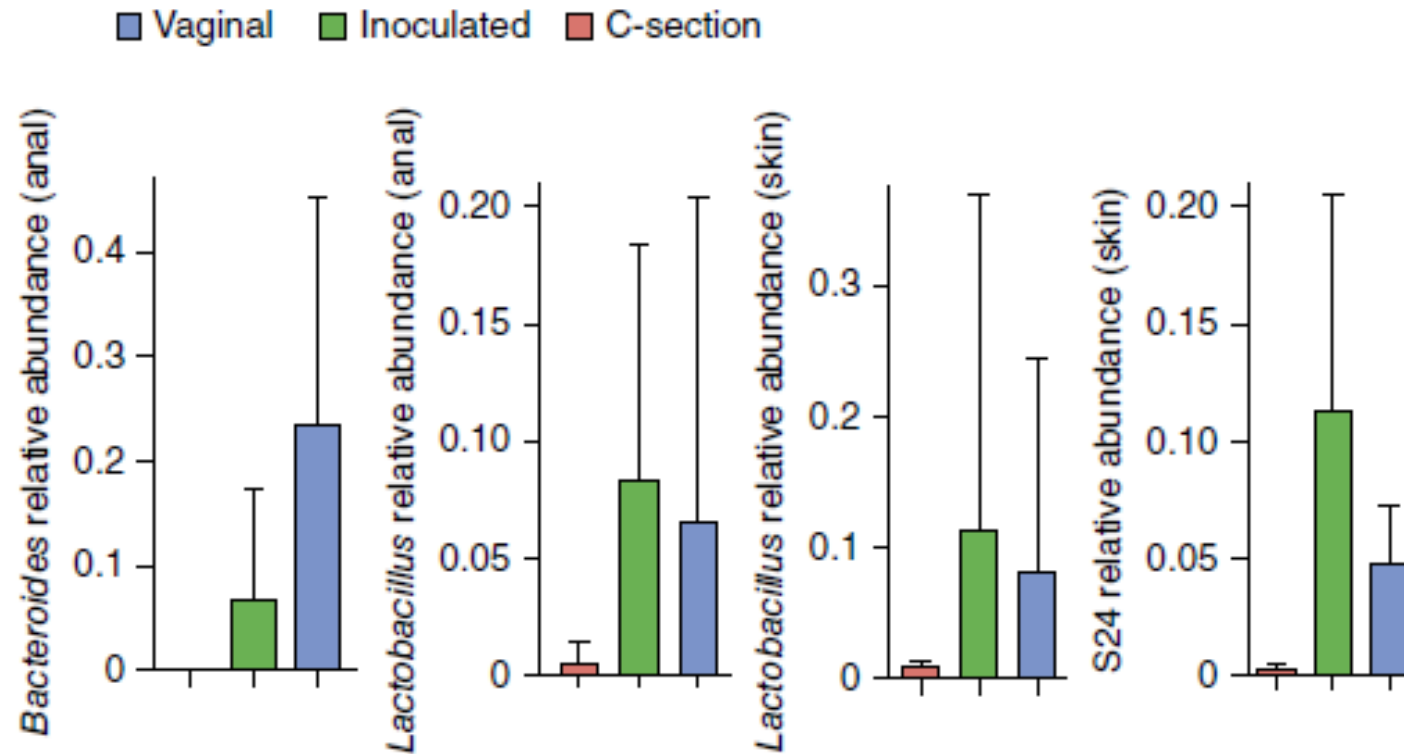
*Nature Medicine* (2016) | doi:10.1038/nm.4039

Received 03 July 2015 | Accepted 22 December 2015 | Published online 01 February 2016

# Groups



# Restitution of Vaginal Microbes



Dominguez-Bello, Nature Medicine, Feb. 2016

# Practice Advisory: American College of Ob/Gyn Nov. 8, 2016

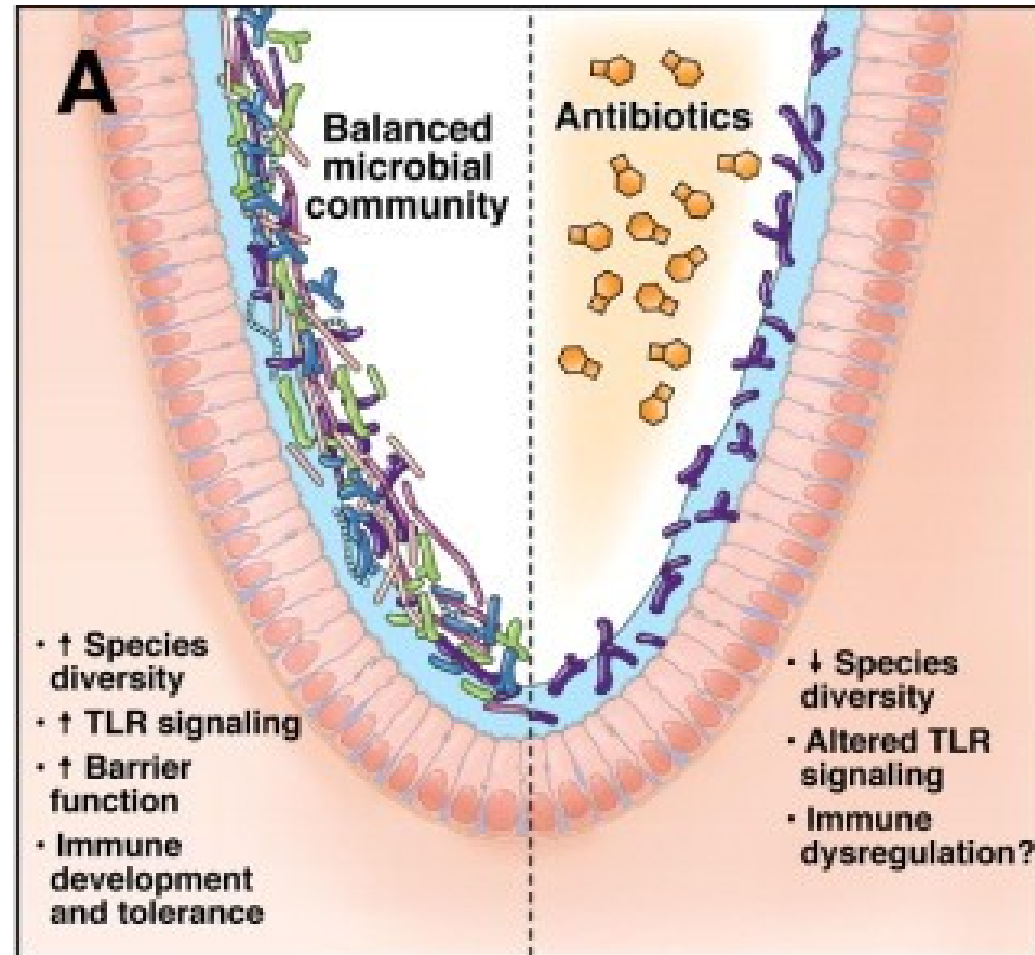
“With the increasing attention given to this concept in public media, providers are being faced with incorporating vaginal seeding into clinical practices. At this time, based on the lack of data about the safety and harms, and the potential for harm from transfer of pathogens associated with adverse neonatal outcomes, **we recommend against the practice of vaginal seeding until better data are available establishing the safety and benefits of this practice.**”

# Antibiotic Use, Dysbiosis and Adverse Outcomes

- Intrapartum antibiotic prophylaxis (IAP) is provided in 30% of Pregnancies.
- Majority of preterms less than 33 weeks gestation are treated with ampicillin and gentamicin for at least 48 hours.
- Average length of treatment of “standard of care” practice in preterm neonates is between 5 to 7 days.
- How do these practices affect the developing microbiome, microbial metabolites, resistance factors and neonatal disease.

# Antibiotic Induced Dysbiosis

**Early  
Antibiotic  
Use: What  
are we  
really  
doing?**



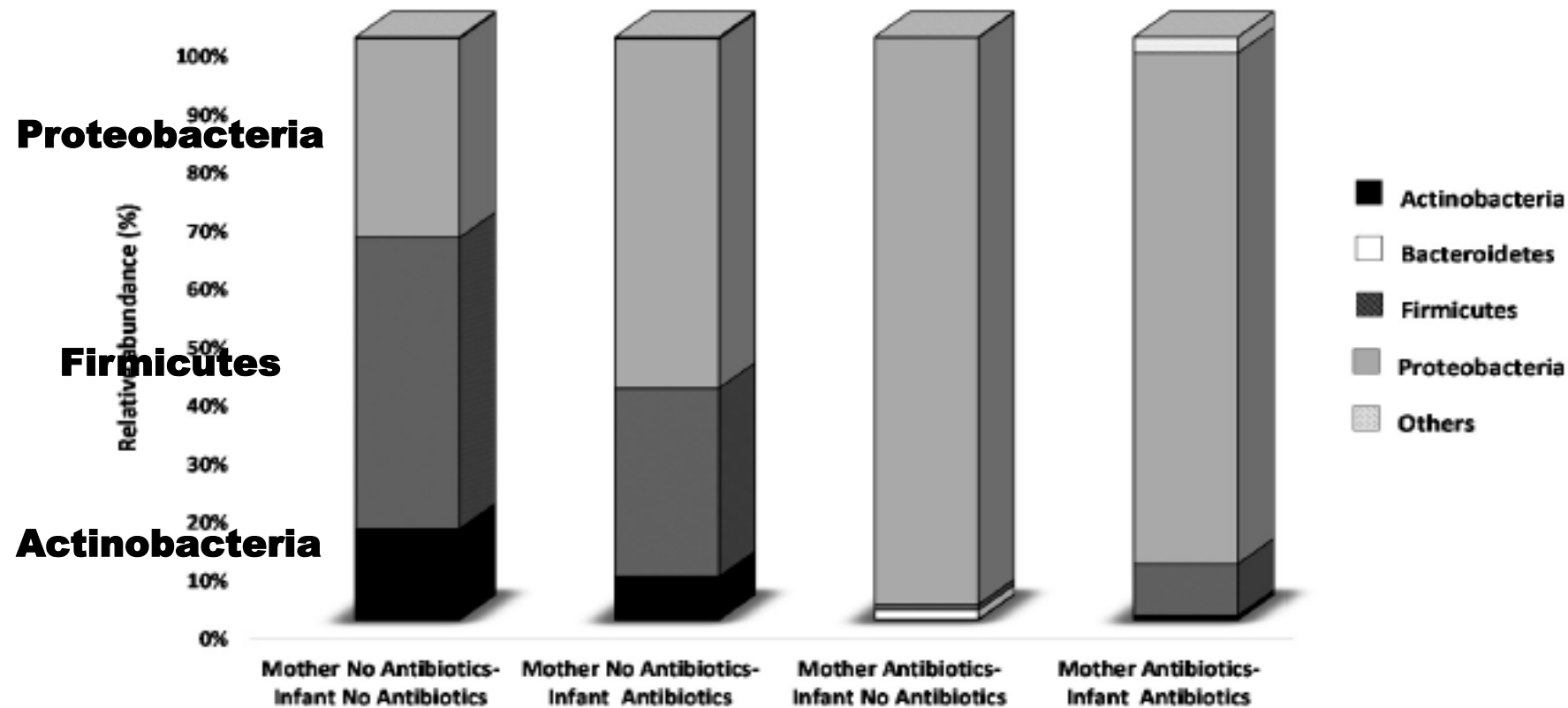
# Variation in Duration of Initial Exposure

- Retrospective review of 50,261 neonates across 127 NICUs from California showed a forty-fold variation in antibiotic prescribing practice with similar burden of infection and mortality.
- What does this mean?
- “Forty-fold variation in NICU antibiotic prescribing practice across 127 NICUs with similar burdens of proven infection, NEC, surgical volume, and mortality indicates that a considerable portion of antibiotic use lacks clear warrant; in some NICUs, antibiotics are overused.”

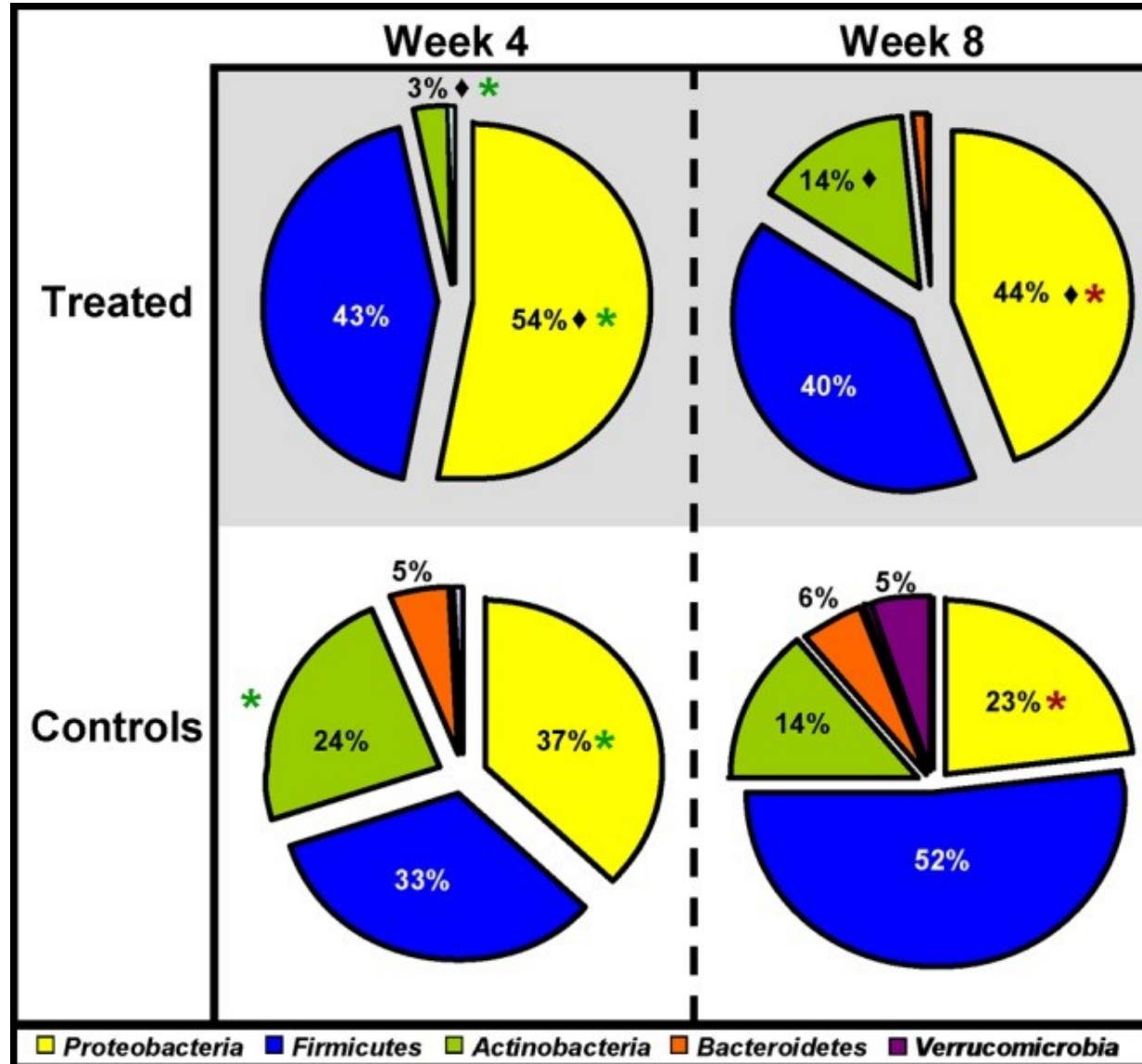
“ So I give a couple days antibiotics to my preterm patients, what does that matter if I change the microbes in the GI tract since I could potentially be saving the baby’s life by treating unrecognized early onset sepsis” ---  
Anonymous Neonatologist.

# Do Antenatal and/or Postnatal Antibiotics Affect Neonatal Microbiota?

Phylum Level Distributions at 30 days depending on Antibiotic Use

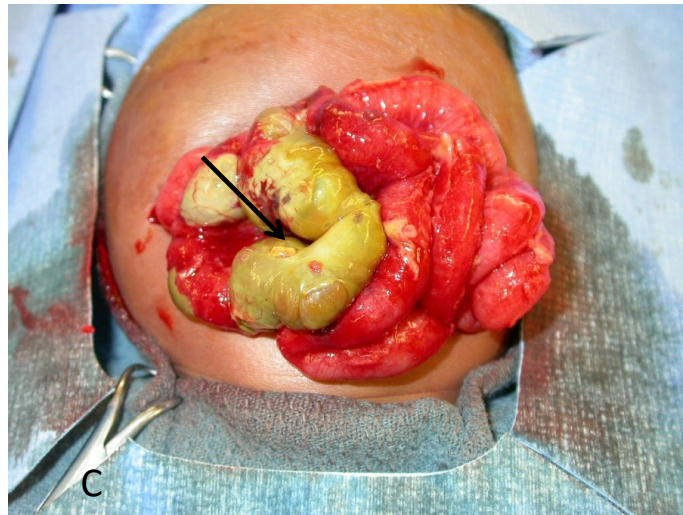
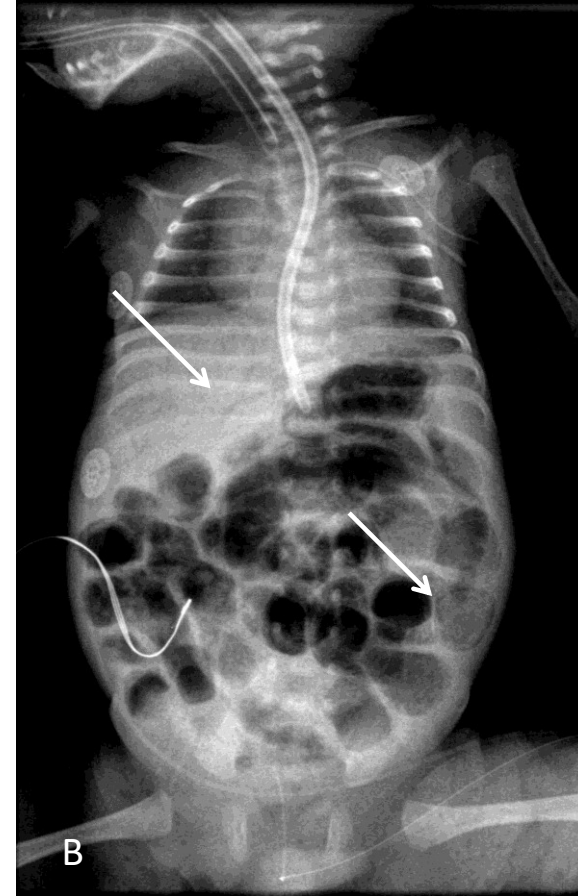


# Effects of Ampicillin and Gentamicin within 48 hours after birth

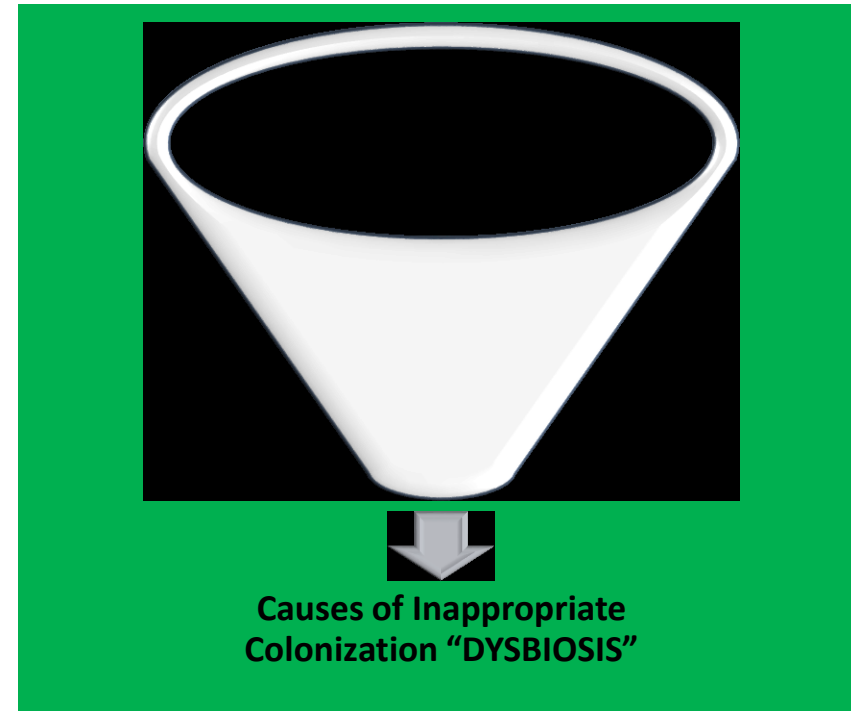
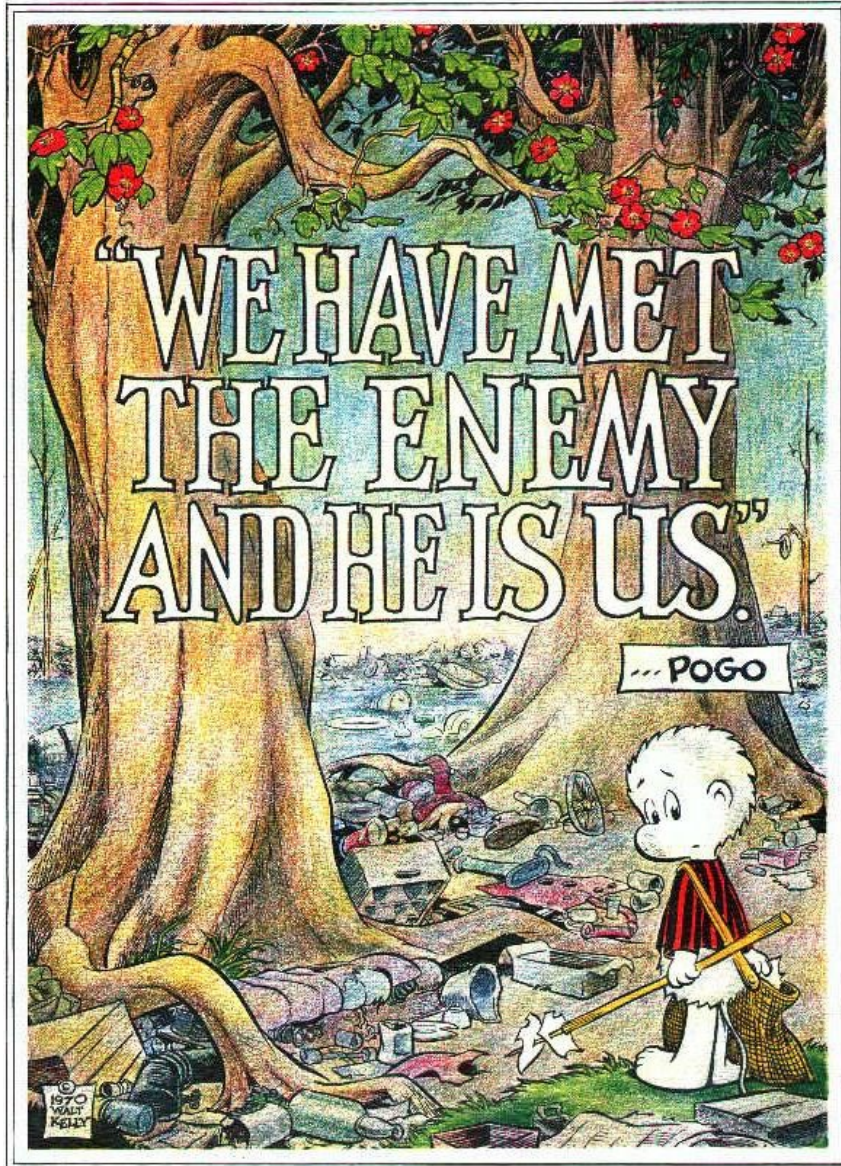


Fouhy, R. et al. Antimicrobial Agents and Chemotherapy. Nov. 2012, vol 56., Number 11.

# “Classic” NEC

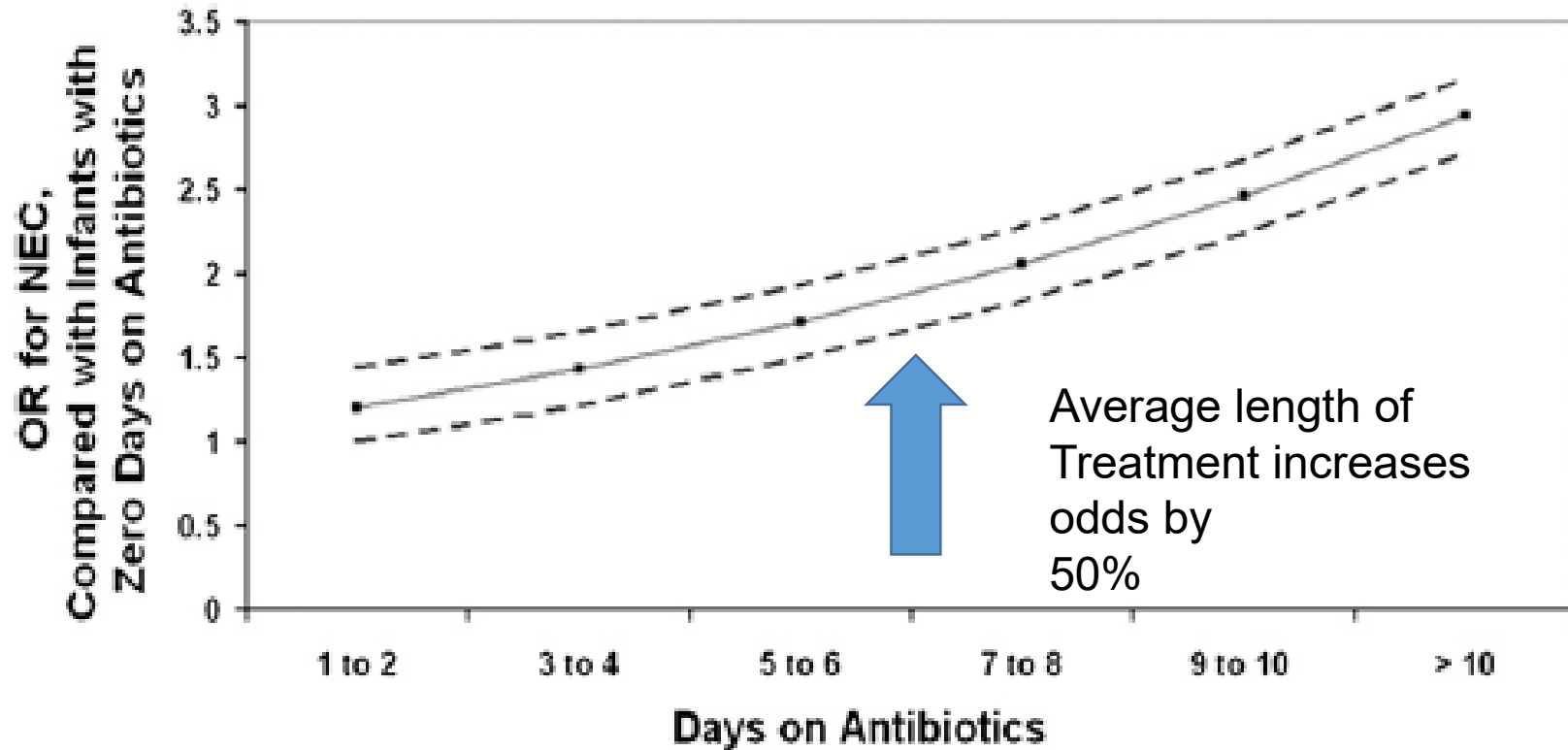


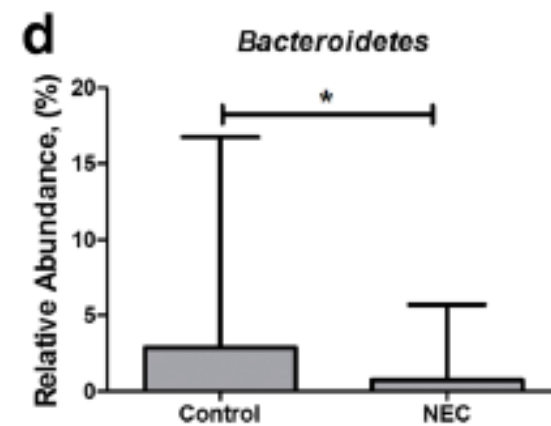
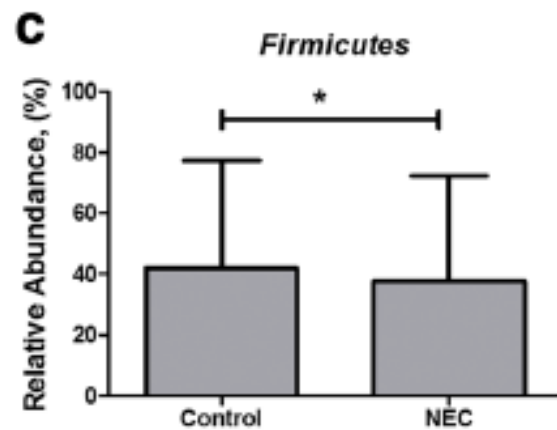
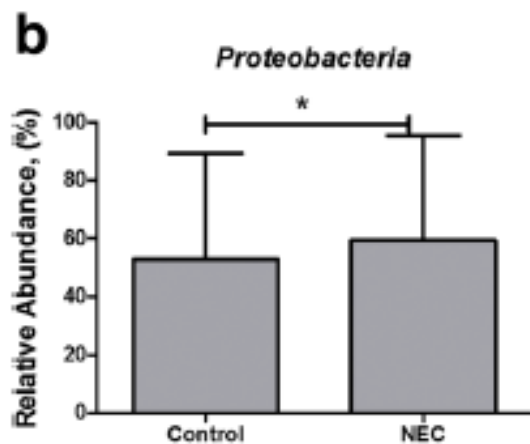
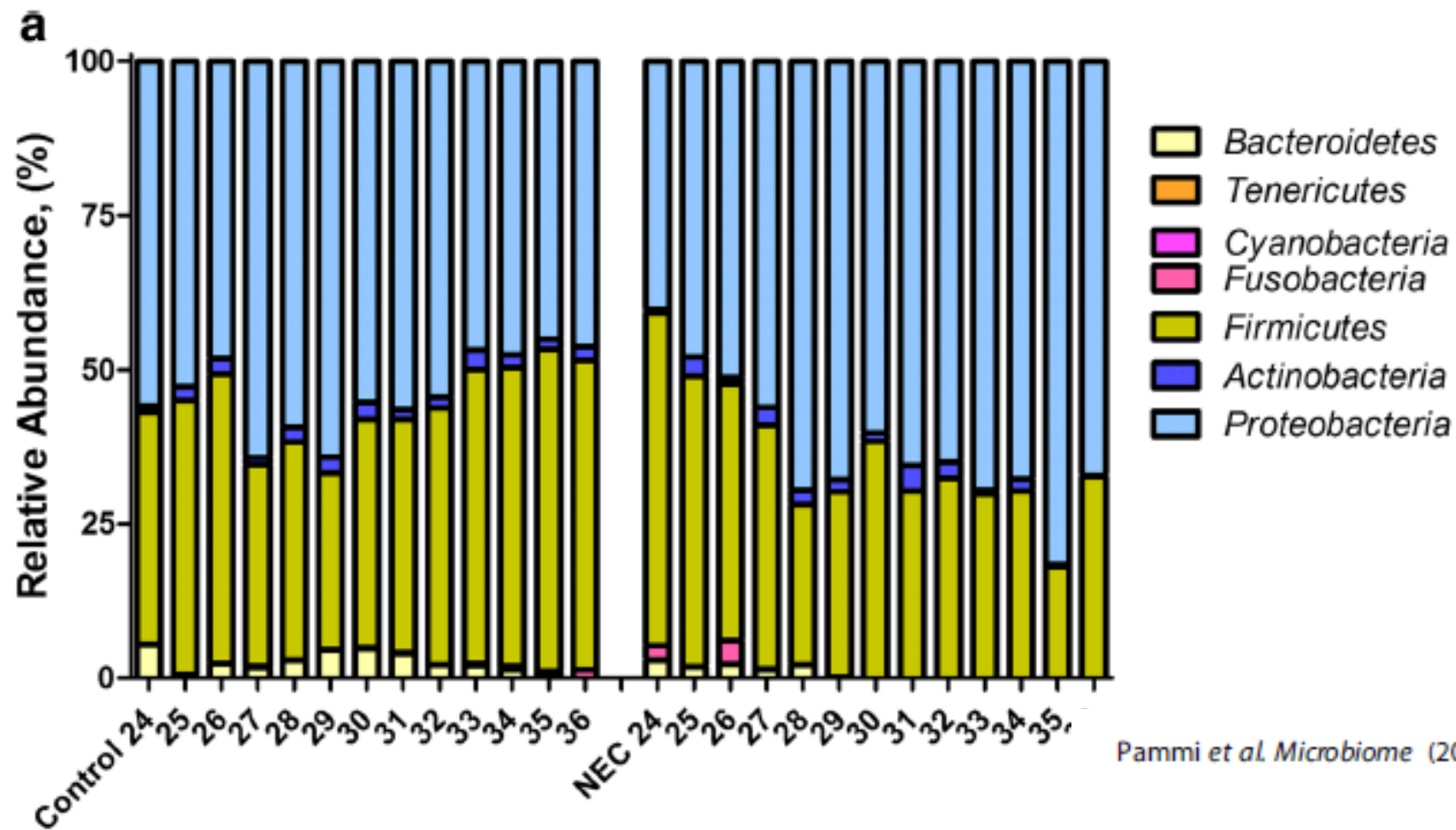
# Do Common Neonatal Practices Cause NEC?



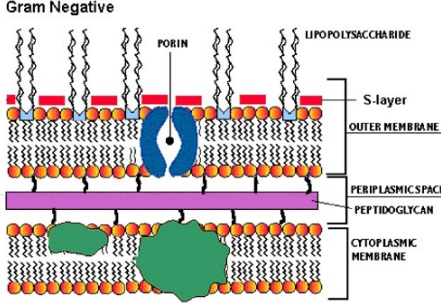
# Odds Ratio of NEC with Increased Days on Antibiotics

Alexander, V.N. J. Pediatrics, Sept. 2011

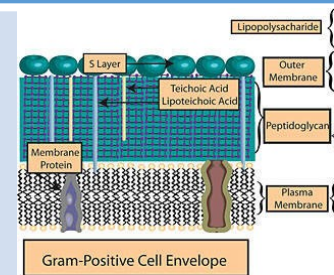




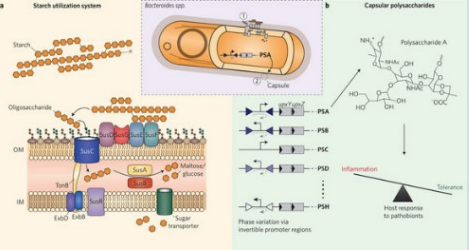
# Comparison of three major phyla: Proteobacteria, Firmicutes and Bacteroidetes

| Phylum         | Gram Staining | Functional Relationship                                                                                                                                                                                                                    | Comment                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            |
|----------------|---------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Proteobacteria | Gram negative | <u>High Lipopolysaccharide (LPS) content in cell wall.</u> Abundance of Proteobacteria increased prior to exacerbations of inflammatory bowel disease. Strong stimulator of TLR4. E. Coli, Klebsiella and Pseudomonas are representatives. |  <p>The diagram illustrates the cell wall structure of a Gram-negative bacterium. It shows an outer membrane (OM) composed of phospholipids with lipopolysaccharide (LPS) chains on the surface. A porin is embedded in the OM. Below the OM is the periplasmic space (PS), which contains a thin layer of peptidoglycan. The innermost layer is the cytoplasmic membrane (CM), which is a phospholipid bilayer. The space between the OM and PS is labeled as the S-layer.</p> |
|                |               |                                                                                                                                                                                                                                            |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |
|                |               |                                                                                                                                                                                                                                            |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |

# Comparison of three major phyla: Proteobacteria, Firmicutes and Bacteroidetes

| Phylum     | Gram Staining | Functional Relationship                                                                                                                                                                                                                                                                                         | Comment                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      |
|------------|---------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Firmicutes | Gram positive | Lactobacilli are a common class of the Firmicutes phylum. Have high <u>lipoteichoic acid</u> in the cell wall, but <u>low LPS</u> . Have excellent capacity for energy harvest. Produce butyrate in high quantities. Butyrate is a major fuel for colonocytes and important for maintenance of tight junctions. |  <p>The diagram illustrates the Gram-Positive Cell Envelope. It shows an outer layer labeled 'S Layer' composed of blue spheres. Below this is the 'Outer Membrane' containing 'Lipopolysaccharide' (LPS) and 'Teichoic Acid Lipoteichoic Acid'. A thick 'Peptidoglycan' layer is situated between the outer membrane and the 'Plasma Membrane'. The plasma membrane is a phospholipid bilayer with 'Membrane Protein' embedded in it. The entire structure is labeled 'Gram-Positive Cell Envelope'.</p> |

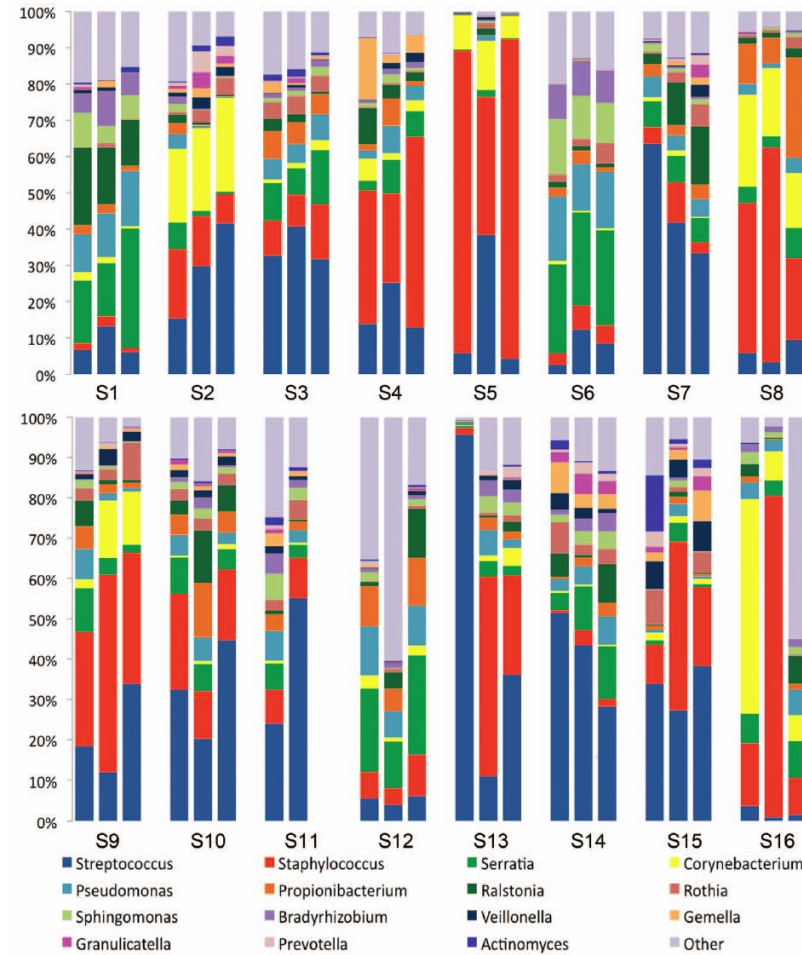
# Comparison of three major phyla: Proteobacteria, Firmicutes and Bacteroidetes

| Phylum        | Gram Staining                                  | Functional Relationship                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         | Comment                                                                             |
|---------------|------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------|
| Bacteroidetes | Gram negative, anaerobic, rod shaped bacteria. | Involvement in fermentation of carbohydrates (propionate and acetate producers), utilization of nitrogenous substances, and biotransformation of bile acids. <i>Bacteroides fragilis</i> is a representative. <u>The immunomodulatory molecule, polysaccharide A (PSA), of <i>B. fragilis</i> mediates the conversion of CD4<sup>+</sup> T cells into Foxp3<sup>+</sup> Treg cells that produce IL-10 during commensal colonization. PSA is not only able to prevent, but also cure experimental colitis in animals. Propionic acid is also a strong inducer of the Foxp3<sup>+</sup> R regulatory pathway.</u> |  |

## Breast milk microbes Over time



Hunt, et al. PlosOne  
2011



**Figure 1.** The community composition of the 15 most abundant bacterial genera in each of 3 milk samples from 16 subjects was diverse. The communities observed were found to be reasonably complex, and while consistent in composition over time for some subjects, a great deal of variation was observed over time in the samples of others.  
doi:10.1371/journal.pone.0021313.g001

# American Academy of Pediatrics Guidelines 2012

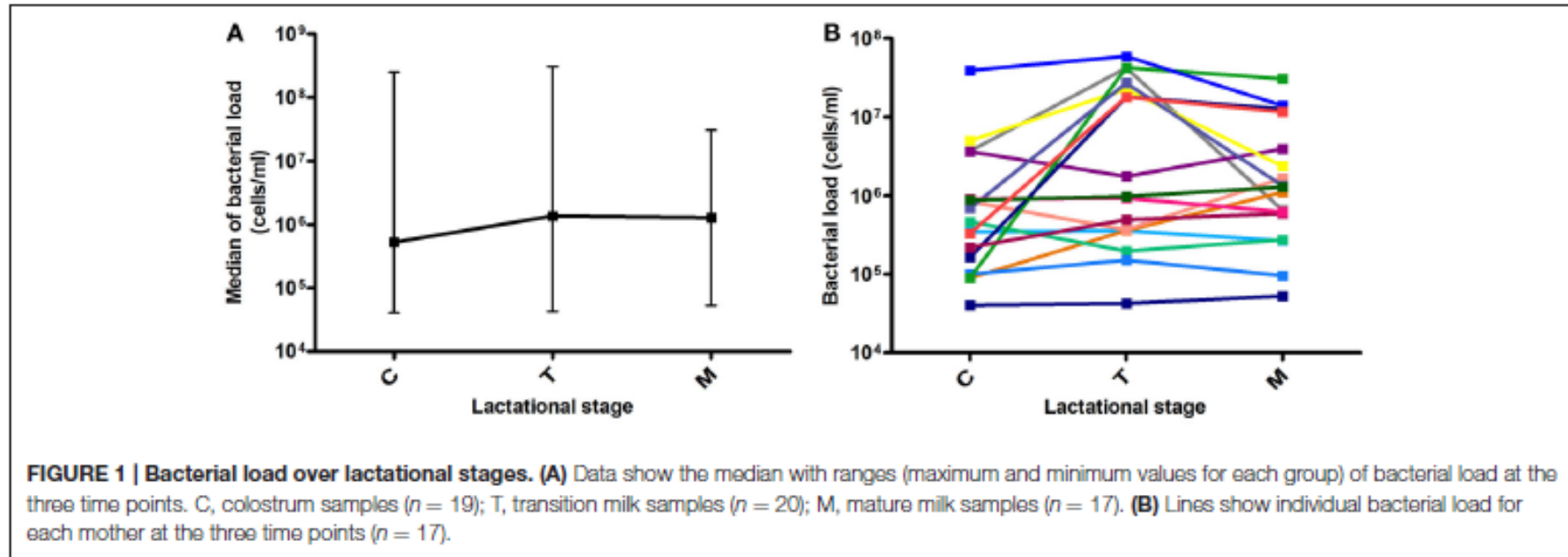
“The potent benefits of human milk are such that all preterm infants should receive human milk). Mother’s own milk, fresh or frozen, should be the primary diet, and it should be fortified appropriately for the infant born weighing less than 1.5 kg. If mother’s own milk is unavailable despite significant lactation support, pasteurized donor milk should be used.

# Donor Human Milk after Holder Pasteurization

|                                    |                    |
|------------------------------------|--------------------|
| Total Lipid                        | NA or ↓ 3.5-5.5%   |
| Free fatty acids                   | ↑ 83%              |
| Lactose and oligosaccharides       | NA                 |
| Total protein                      | NA or 4% reduction |
| Alkaline phosphatase (ALP)         | ↓ 99%              |
| Bile salt-stimulated lipase (BSSL) | Abolished          |
| Lactoferrin (LF)                   | NA or ↓ 44-91%     |
| LF-iron-binding capacity           | ↓ 71%              |
| IgA                                | NA or ↓ 20-60%     |
| IgM                                | Abolished          |
| IgG                                | ↓ 34%              |
| Lactoperoxidase (LP)               | ↓ 82%              |
| LP activity                        | ↓ 88%              |
| Lysozyme                           | ↓ 24-60%           |
| Lysozyme activity                  | NA or ↓ 65-85%     |
| TGF-β                              | NA                 |
| IGF-I                              | ↓ 39%              |
| IGF-II                             | ↓ 10%              |
| IGFBPs                             | ↓ 7-19%            |
| EGF                                | NA                 |



# Bacterial Load over Lactational Stages: FusA Gene PCR



Boix-Amaros, A. *Frontiers in Microbiology*. 20 April, 2016

# Microbial Dose from Human Milk

- Assume intake of 800 ml/day
- Assume  $10^{5-6}$  bacterial cells/ml
- This will provide  $10^{7-8}$  bacterial cells (personalized?) daily, close to the dose in most probiotic studies.

Edited by: Christophe Lacroix, ETH Zurich, Switzerland  
Reviewed by:

ORIGINAL RESEARCH  
published: 03 August 2017 doi: 10.3389/fmicb.2017.01470

# Personalization of the Microbiota of Donor Human Milk with Mother's Own Milk

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The American Academy of Pediatrics recommends that extremely preterm infants receive mother's own milk (MOM) when available or pasteurized donor breast milk (DBM) when MOM is unavailable. The goal of this study was to determine whether DBM could be inoculated with MOM from mothers of preterm infants to restore the live microbiota (RM). Culture dependent and culture independent methods were used to analyze the fluctuations in the overall population and microbiome, respectively, of DBM, MOM, and RM samples over time. Using MOM at time 0 (T0) as the target for the restoration

# Re-Faunation Experiments

- Donor Milk Refaunation: Adding mother's own milk to pasteurized donor breast milk (DBM) to add back potentially beneficial microbes: 10-30% dilution will expand a large part of microbiome after 4-8 hours.



# Take Home Messages

- Studies of the Microbiome are opening new avenues in our understanding of Health and Disease.
- The Fetal Maternal Unit most likely is not “sterile” and prenatal exposure to microbes may be very important in subsequent development.
- Early life pre, peri and postnatal exposures such as antibiotic use and diet are particularly important in that they can result in life-long changes in health and disease.
- Future studies will utilize multi-omic approaches to better delineate how microbes interact with the environment and the host in the cause and/or prevention of disease.