

#### 16SrRNA Intermediate Bioinformatics Online Course: Int\_BT\_2019

#### **Module 2:**

#### Introduction to the microbiome – why 16S?

**Part 2.2** 









Gestational duration
Prenatal antibiotic treatment
Maternal prenatal stress
Maternal diabetes status

Mode of delivery

Antibiotics?

Solid foods

Antibiotics?

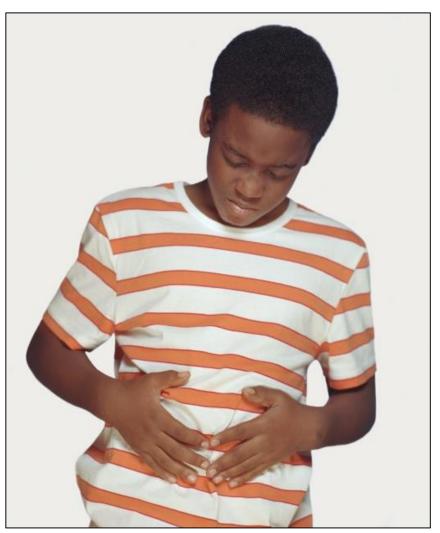












Decker et al. (2011). Gut Microbes. 2:91-98, Marcobal et al. (2010). J Agric Food Chem. 58:5334-40; Renz-Polster et al. (2005) Clin Exp Allergy. 35: 1466-72





















Published in final edited form as:

Nat Med. 2016 March; 22(3): 250-253. doi:10.1038/nm.4039.

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

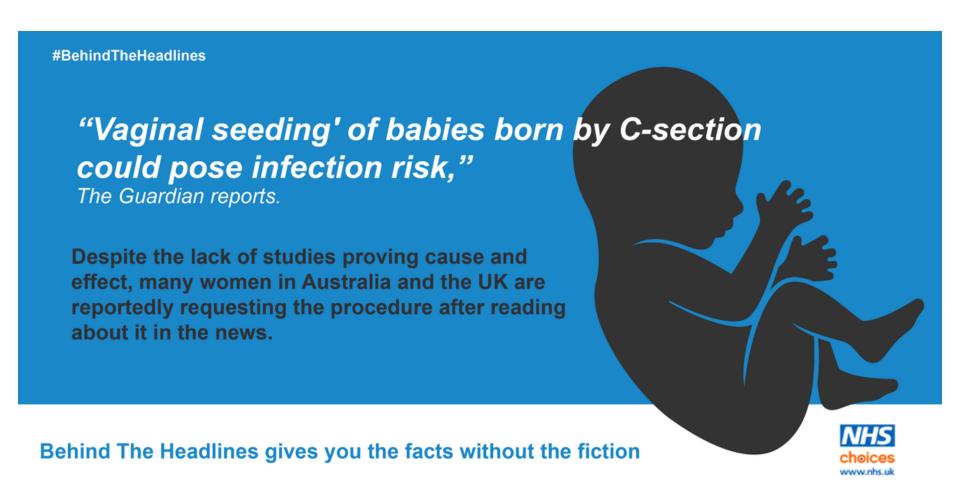
Maria G. Dominguez-Bello<sup>1,2,\*</sup>, Kassandra M. De Jesus-Laboy<sup>2</sup>, Nan Shen<sup>8</sup>, Laura M. Cox<sup>1</sup>, Amnon Amir<sup>3,7</sup>, Antonio Gonzalez<sup>3,7</sup>, Nicholas A. Bokulich<sup>1</sup>, Se Jin Song<sup>3,4</sup>, Marina Hoashi<sup>5</sup>, Juana I. Rivera-Vina<sup>6</sup>, Keimari Mendez<sup>6</sup>, Rob Knight<sup>3,7</sup>, and Jose C. Clemente<sup>8,9,\*</sup>



















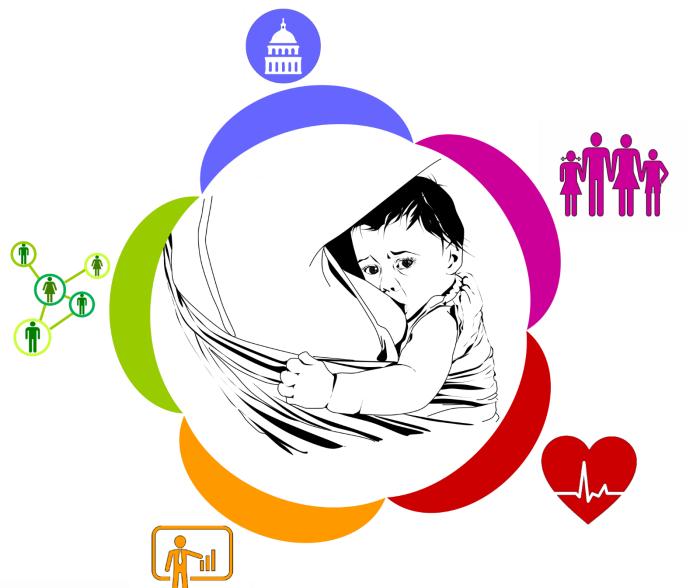




















### Randomized controlled trial on the impact of early-life intervention with bifidobacteria on the healthy infant fecal microbiota and metabolome

Monika Bazanella, <sup>1</sup> Tanja V Maier, <sup>4</sup> Thomas Clavel, <sup>2</sup> Ilias Lagkouvardos, <sup>2</sup> Marianna Lucio, <sup>4</sup> Maria X Maldonado-Gòmez, <sup>5</sup> Chloe Autran, <sup>7</sup> Jens Walter, <sup>6</sup> Lars Bode, <sup>7</sup> Philippe Schmitt-Kopplin, <sup>3,4</sup> and Dirk Haller <sup>1,2</sup>

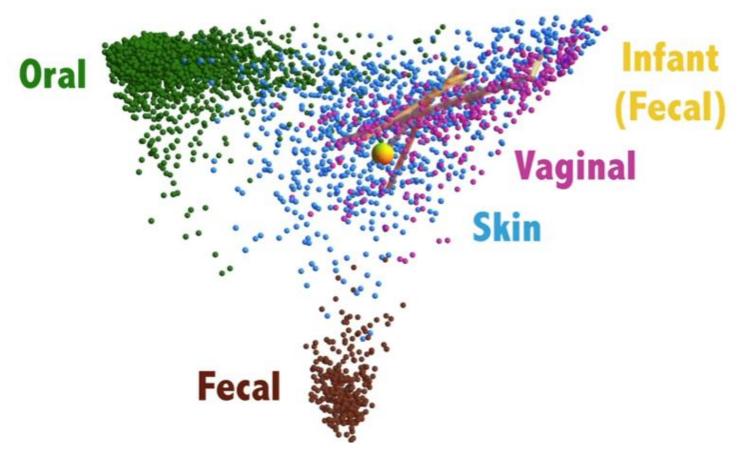
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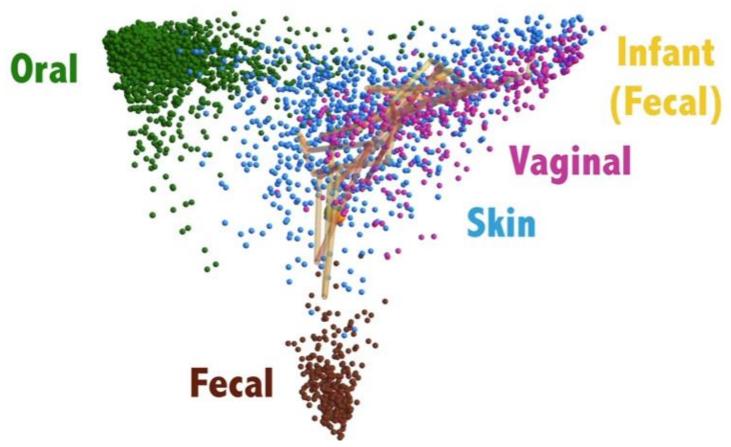










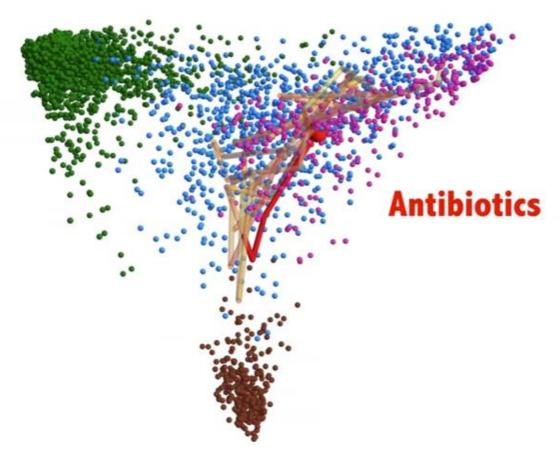










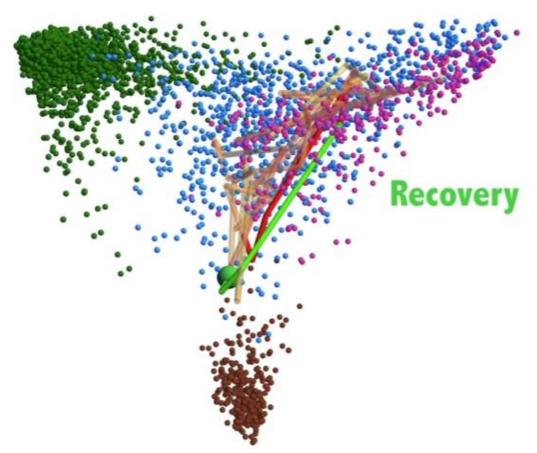










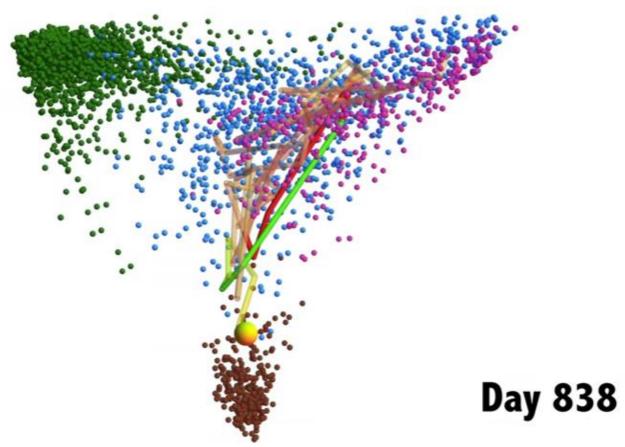






















RESEARCH ARTICLE Host-Microbe Biology



Antibiotic-Induced Alterations of the Gut Microbiota Alter Secondary Bile Acid Production and Allow for *Clostridium* difficile Spore Germination and Outgrowth in the Large Intestine

Casey M. Theriot, a,b Alison A. Bowman, b Vincent B. Youngb,c











Therapeutic Advances in Gastroenterology

Review

## Fecal microbiota transplantation: in perspective

Shaan Gupta, Emma Allen-Vercoe and Elaine O. Petrof

Abstract: There has been increasing interest in understanding the role of the human gut microbiome to elucidate the therapeutic potential of its manipulation. Fecal microbiota transplantation (FMT) is the administration of a solution of fecal matter from a donor into the intestinal tract of a recipient in order to directly change the recipient's gut microbial composition and confer a health benefit. FMT has been used to successfully treat recurrent Clostridium difficile infection. There are preliminary indications to suggest that it may also carry therapeutic potential for other conditions such as inflammatory bowel disease, obesity, metabolic syndrome, and functional gastrointestinal disorders.

Ther Adv Gastroenterol

2016, Vol. 9(2) 229-239

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Creating crapsules: is faeces in a pill the cure for our

ills?



https://www.smh.com.au/lifestyle/health-and-wellness/creating-crapsules-is-faeces-in-a-pill-the-cure-for-our-ills-20180319-p4z53z.html











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References, detailed credits and more in the description











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Table 1. Role of Pathogenic Gut Microbiota in Gastrointestinal Diseases

Risk factor	Microbial change	Possible mechanisms	Ref.
ID			
Genetics (NIrp6 deficient)	Prevotellaceae ↑, TM7 ↑	IL-18↓, CCL5↑, and innate and adaptive immune cell recruitment	14, 15, 16
Genetics (IL-10, IL-2 deficient)	Escherichia coli or Enterococcus faecalis (monocolonization)	IL-12, IFN-γ↑	23
Genetics (HLA-B27)	Bacteroides fragilis (monocolonization)	Unknown	24
Diet (high fat derived from milk)	Firmicutes ↓, Bilophila wadsworthia ↑	Immune system (Th1) disruption	26, 27
Diet (high protein)	Desulfovibrio spp. ↑, Desulfuromonas spp. ↑	Genotoxic ↑, DNA damage ↑, inflammation ↑	28, 29
Diet (high fat, high beef)	Erysipelotrichaceae ↑, Bacteroides fragilis ↑	Unknown	30, 31
Smoking	Anaerostipes ↓	Butyrate ↓	35
Antibiotics (ciprofloxacin, metronidazole)	Dorea ↓, Butyricicoccus ↓, Coriobacteriaceae ↓	Organic acid \$\formic (e.g., formic acid, butyrate)	40, 41
Antibiotics	Clostridium scindens ↓, Clostridium difficile ↑	DCA ↓	47
Unknown	Faecalibacterium prausnitzii ↓	Anti-inflammatory effect ↓	50, 51
Unknown	pks+ Escherichia coli ↑	Colibactin ↑, DNA damage ↑	64

Nagao-Kitamoto et al. (2016) Intest Res. 14: 127-138

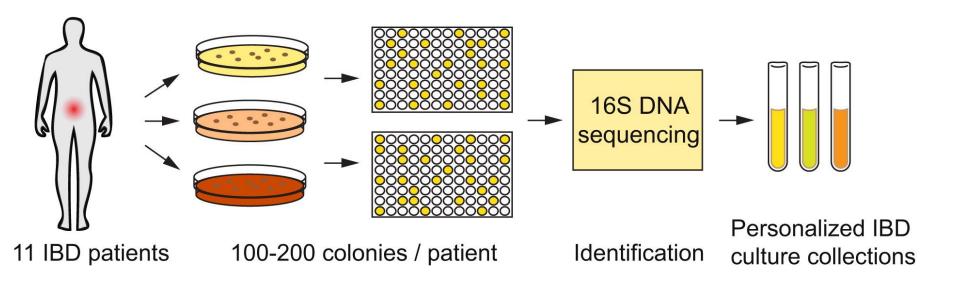








#### Irritable bowel disease: confirmed using mouse models



Selected individual bacterial isolates comprising of IgA+ and IgA- bacteria and colonized germ free mice.

High IgA coating are thought to mark colitogenic bacteria in inflammatory bowel disease

Palm et al. (2014) Cell 158: 1000-1010

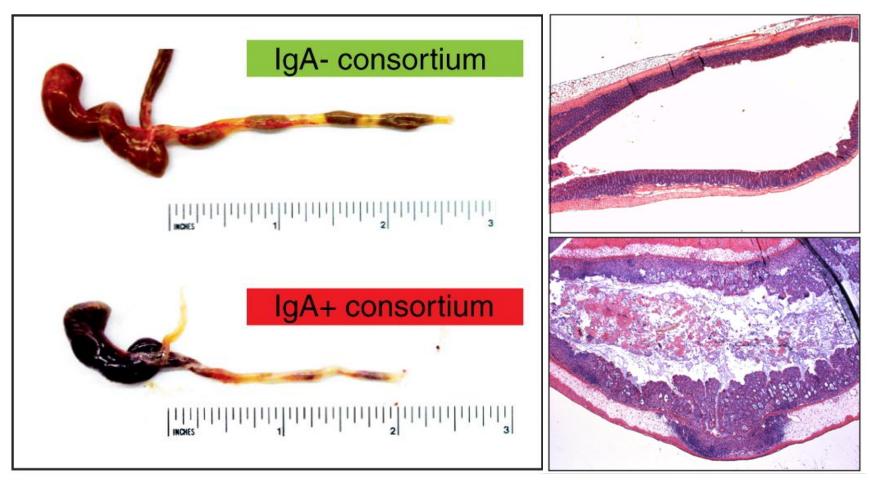








#### Irritable bowel disease: confirmed using mouse models













#### In summary:

- Our GIT microbiome is a "plastic entity" which is modulated by a number of exposures throughout our lives.
- A large number of 16S studies have contributed to our current knowledge of the GIT microbiome – which has led to a number of potential interventions for disease states.
- To date, the majority of 16S studies have focussed on the GIT microbiome.
- This research, however, is still very new and more well designed studies are needed to better understand not only "what's there", but also "what they're doing".



