

IUNS 21st ICN

International Congress of Nutrition

“From Sciences to Nutrition Security”



INTERNATIONAL UNION OF
NUTRITIONAL SCIENCES

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Technical Secretariat:
Fase 20
Fase 20 Argentina
Calle 20 Avenida 843. Piso 2 departamentos 7 RETIRO
1061 Buenos Aires Capital Federal (Argentina)
www.fase20.com - info@fase20.com

Conflict of Interest Disclosure

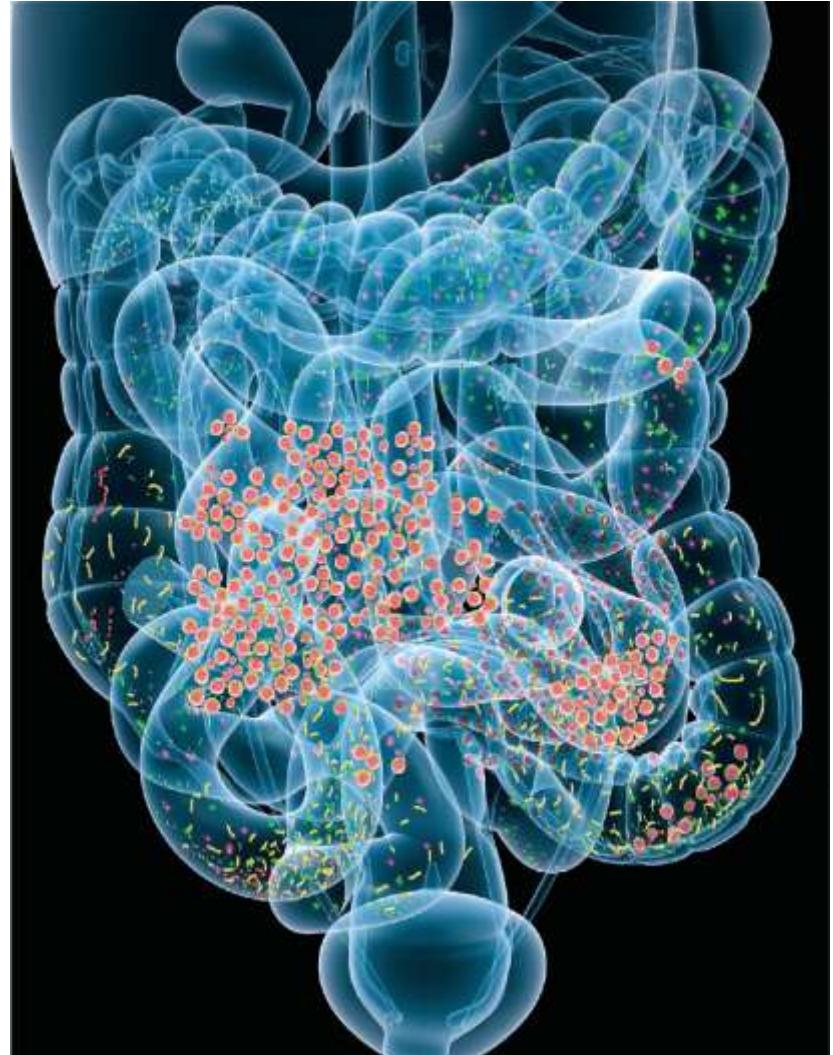
I have no conflict of interest to report in relation to
this presentation.

Sharing Our Bodies: The Symbiosis of Humans and Our Microbiota

- ▶ Christian Hoffmann, PhD
- ▶ School of Pharmaceutical Sciences
- ▶ Dept. of Food Science and Experimental Nutrition
- ▶ 10/2017
- ▶ c.hoffmann@usp.br

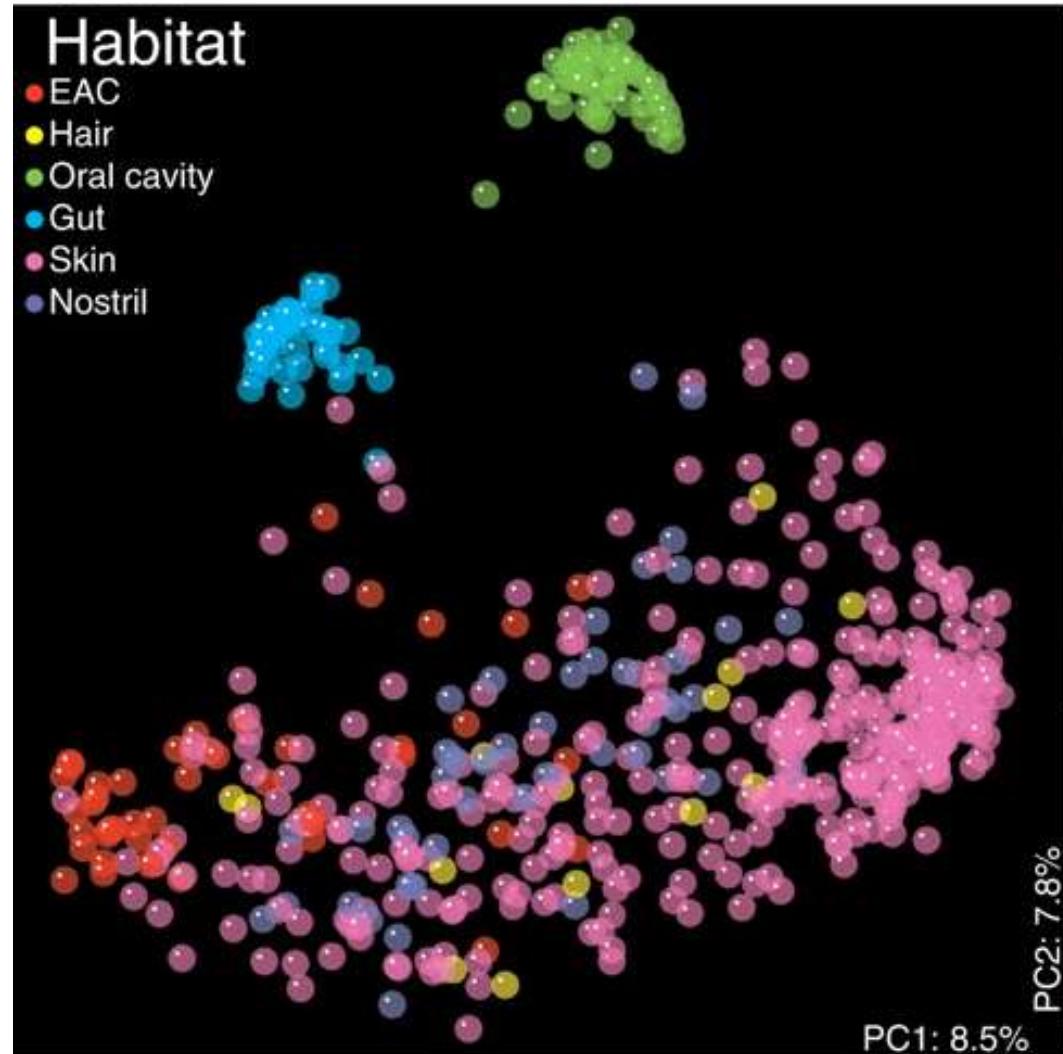
The Human Microbiome

- ▶ Data overload <- NGS(sequences)
- ▶ Composed of: Bacteria, Archaea, Eukaryotes and Virus
- ▶ Density: 10^{11} a 10^{12} cells/mL (colon)
- ▶ High number of species present
- ▶ One microbiome to each body part (nose, mouth, intestine, skin etc.)

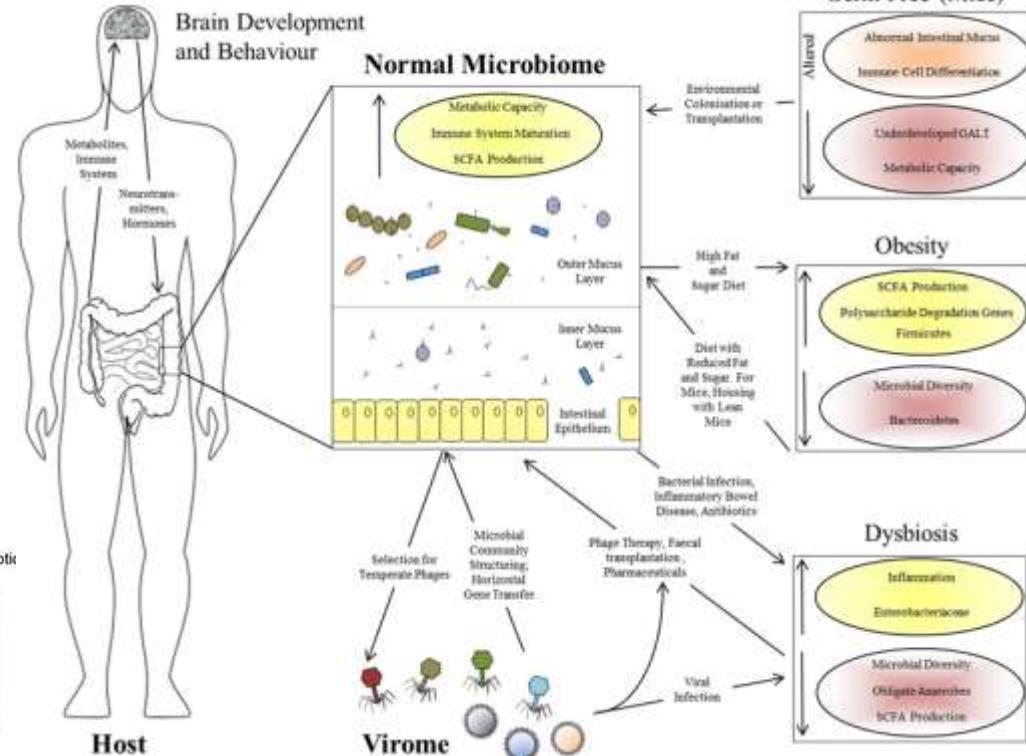
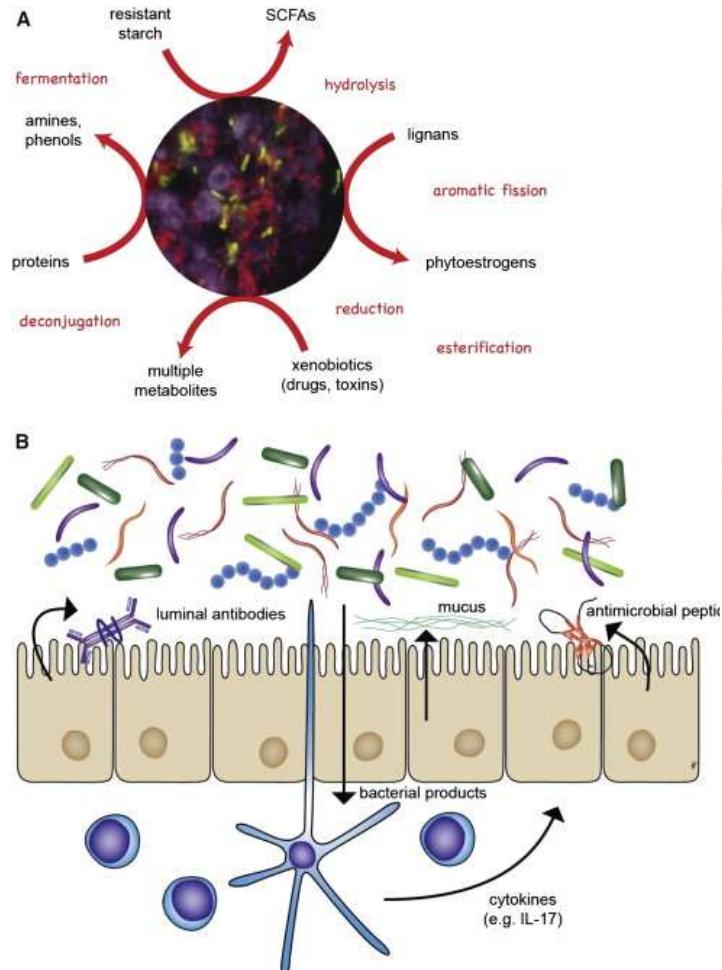


The Human Microbiome

- ▶ Composed of :
Bacteria, Archaea,
Eukaryotes e Viruses
- ▶ Density: 10^{11} a 10^{12}
cells/mL (colon)
- ▶ High number of
species present
- ▶ Distinct
compositions for
each body site
(nose, mouth,
intestine, skin etc.)

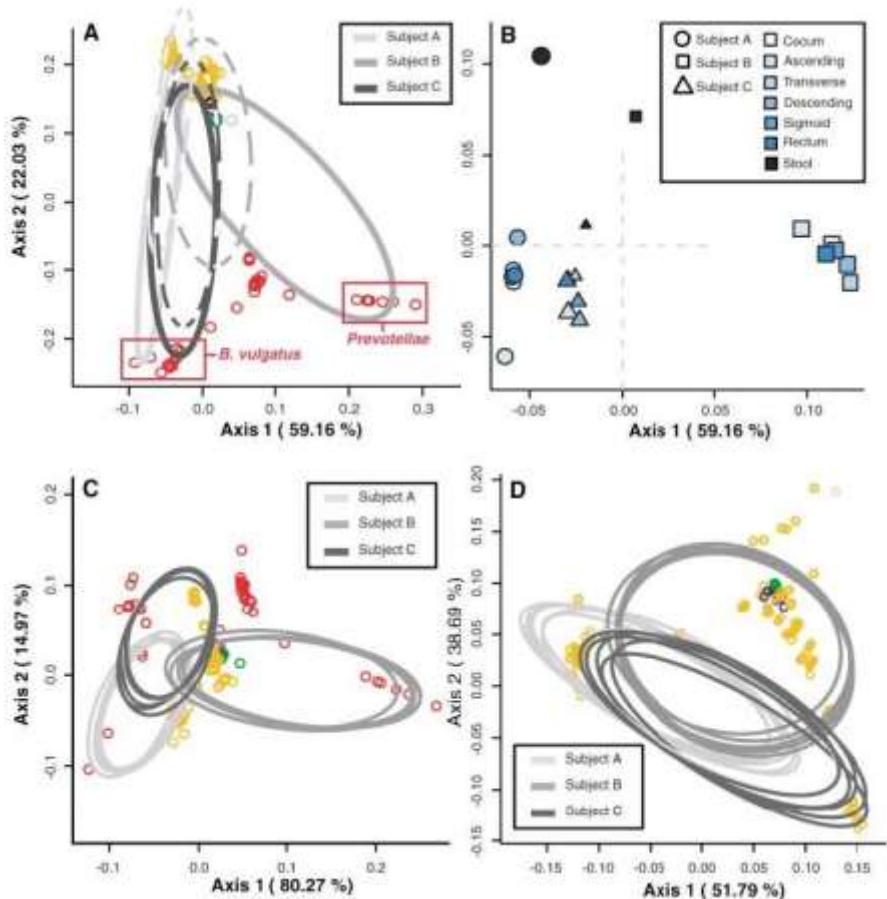


Symbiosis and Dysbiosis

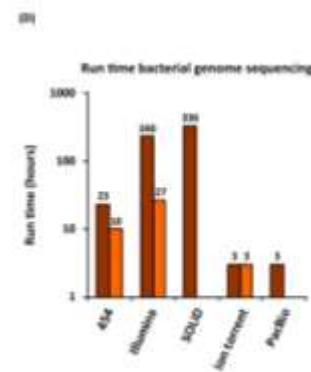
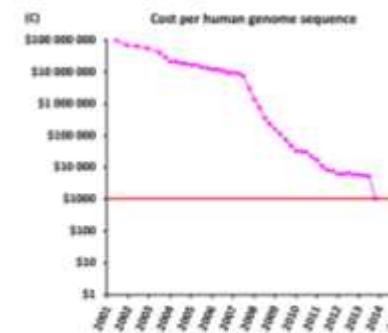
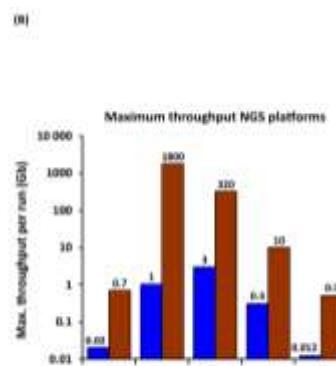
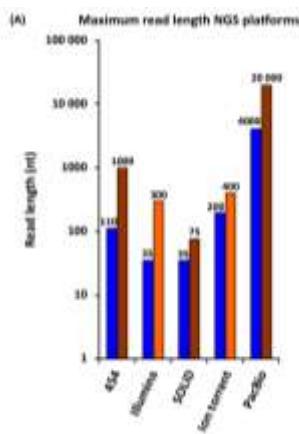


In the beginning:

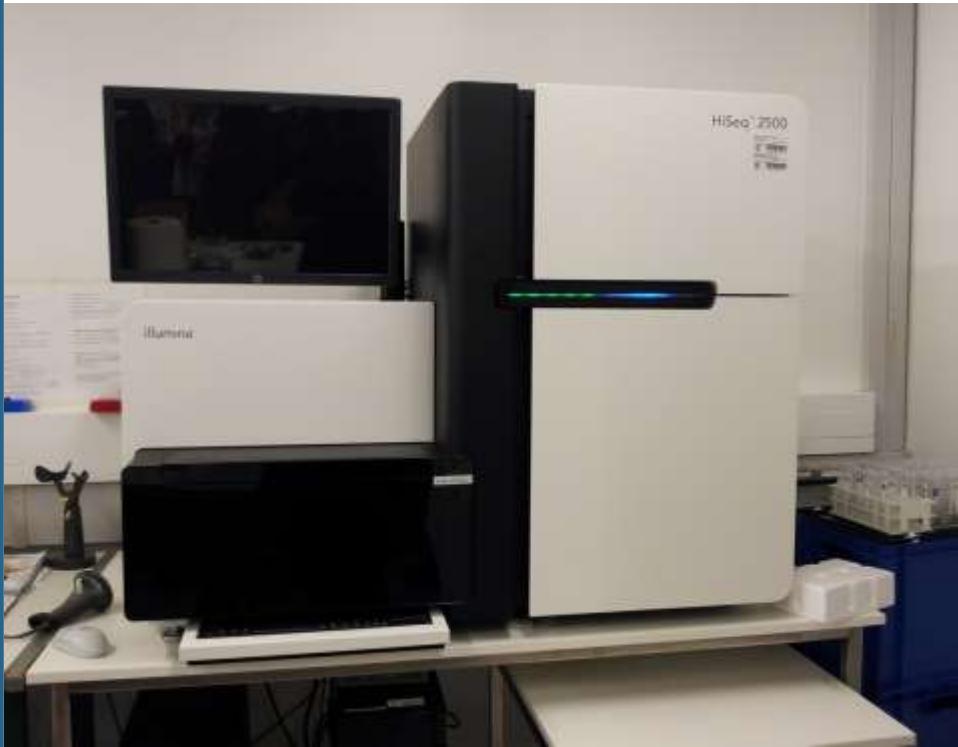
- ▶ In this issue of *Science*, we have:
 - ▶ 3 volunteers
 - ▶ Sanger
 - ▶ 11831 seqs: bacteria
 - ▶ 1524 seqs: archaea
 - ▶ “*Diversity of the Human Intestinal Microbial Flora*”



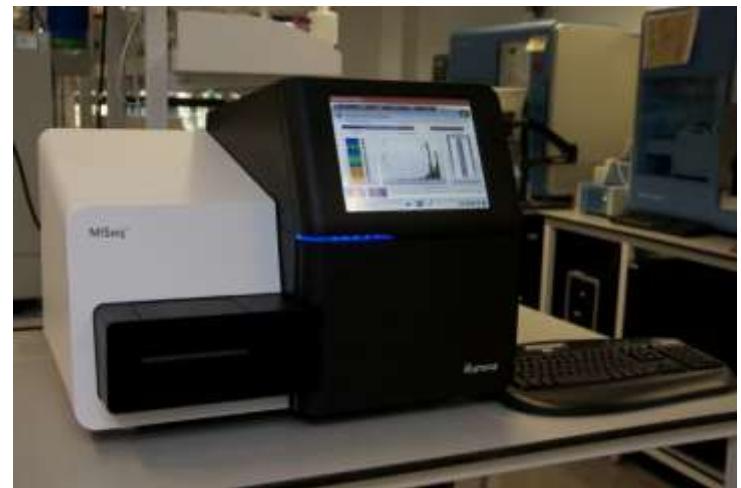
With the advent of NGS



Now

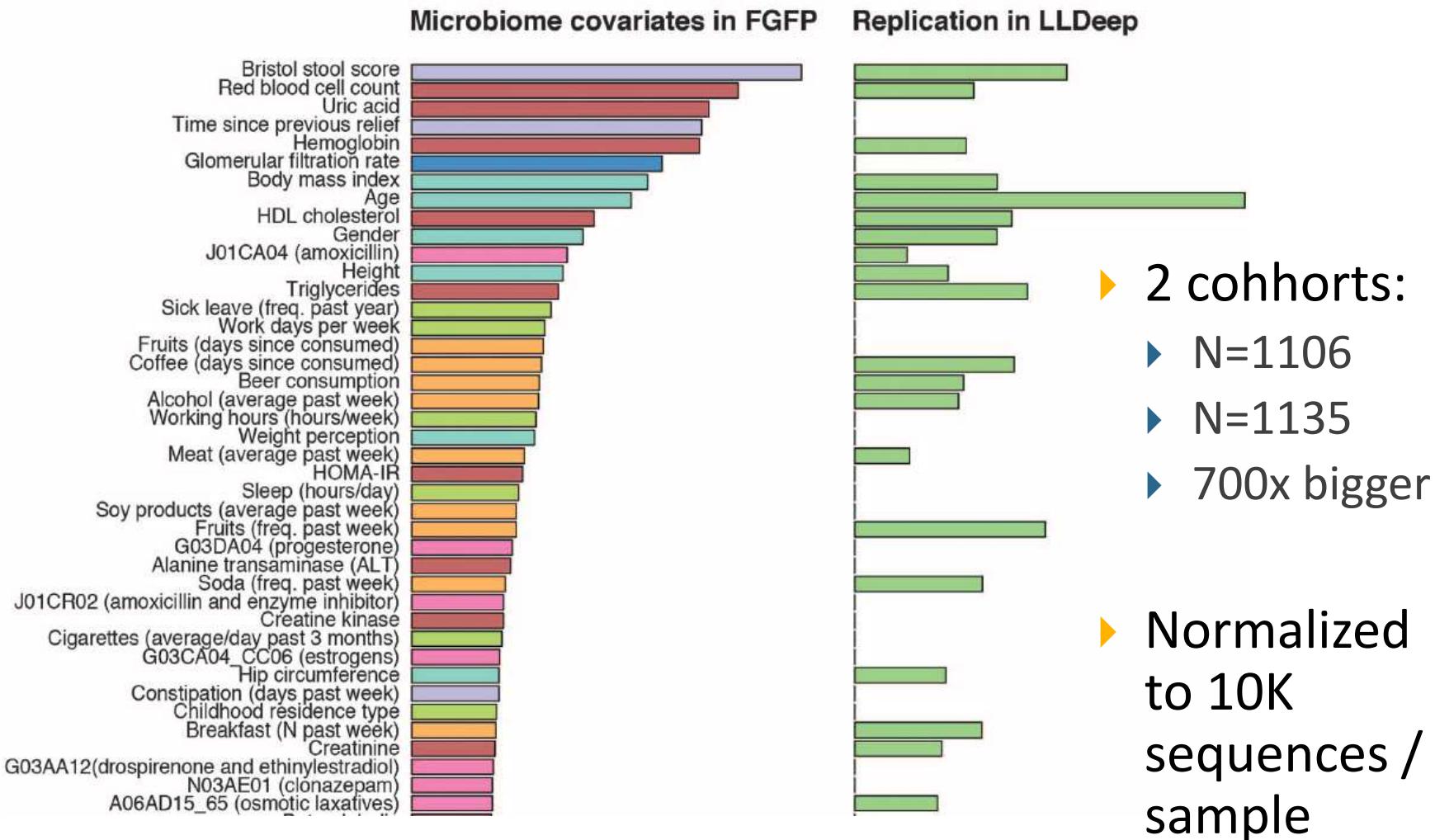


► 3 to 6 billion *reads*



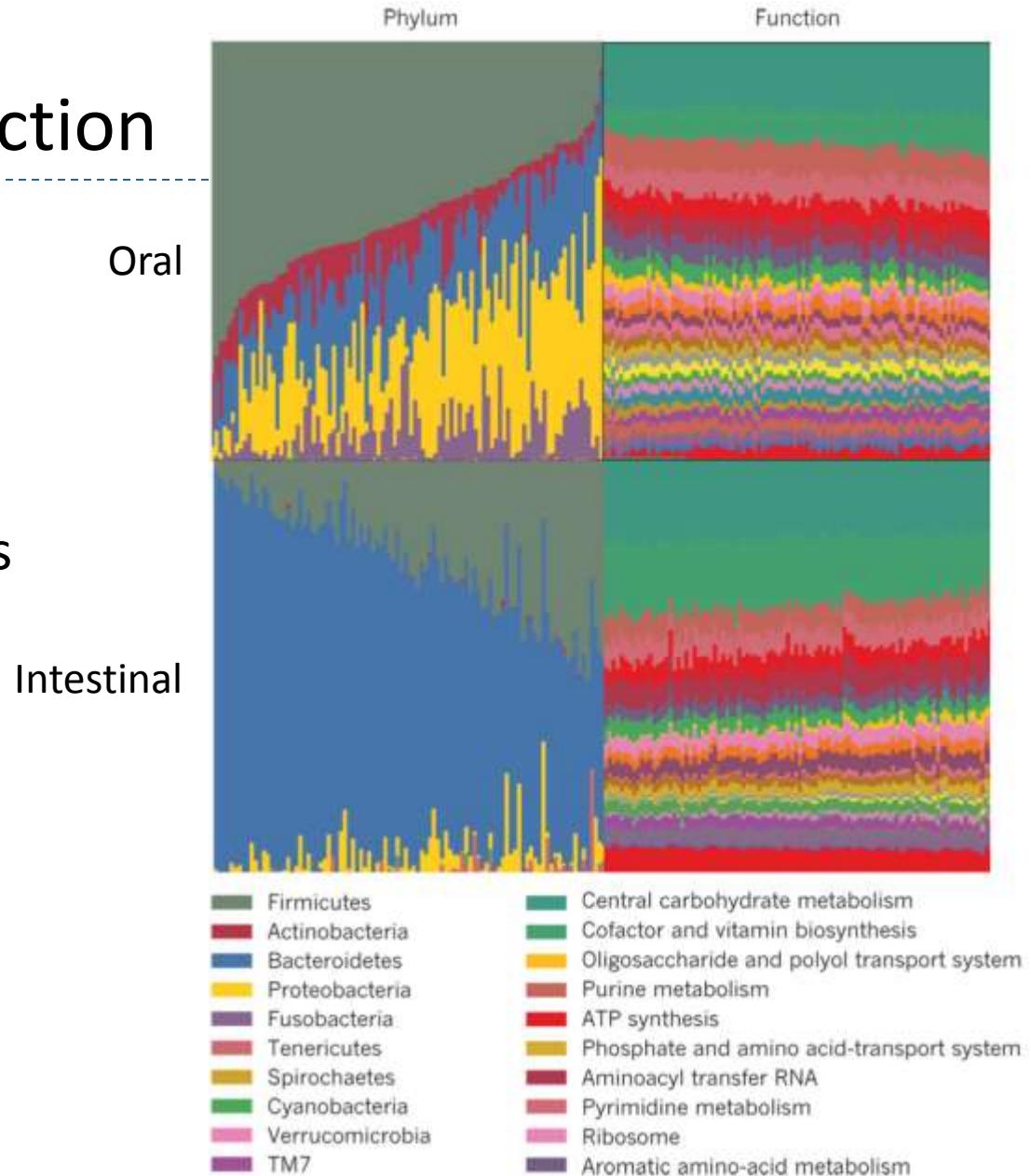
► 25 million *reads*

Co-variation with health/diet/lifestyle



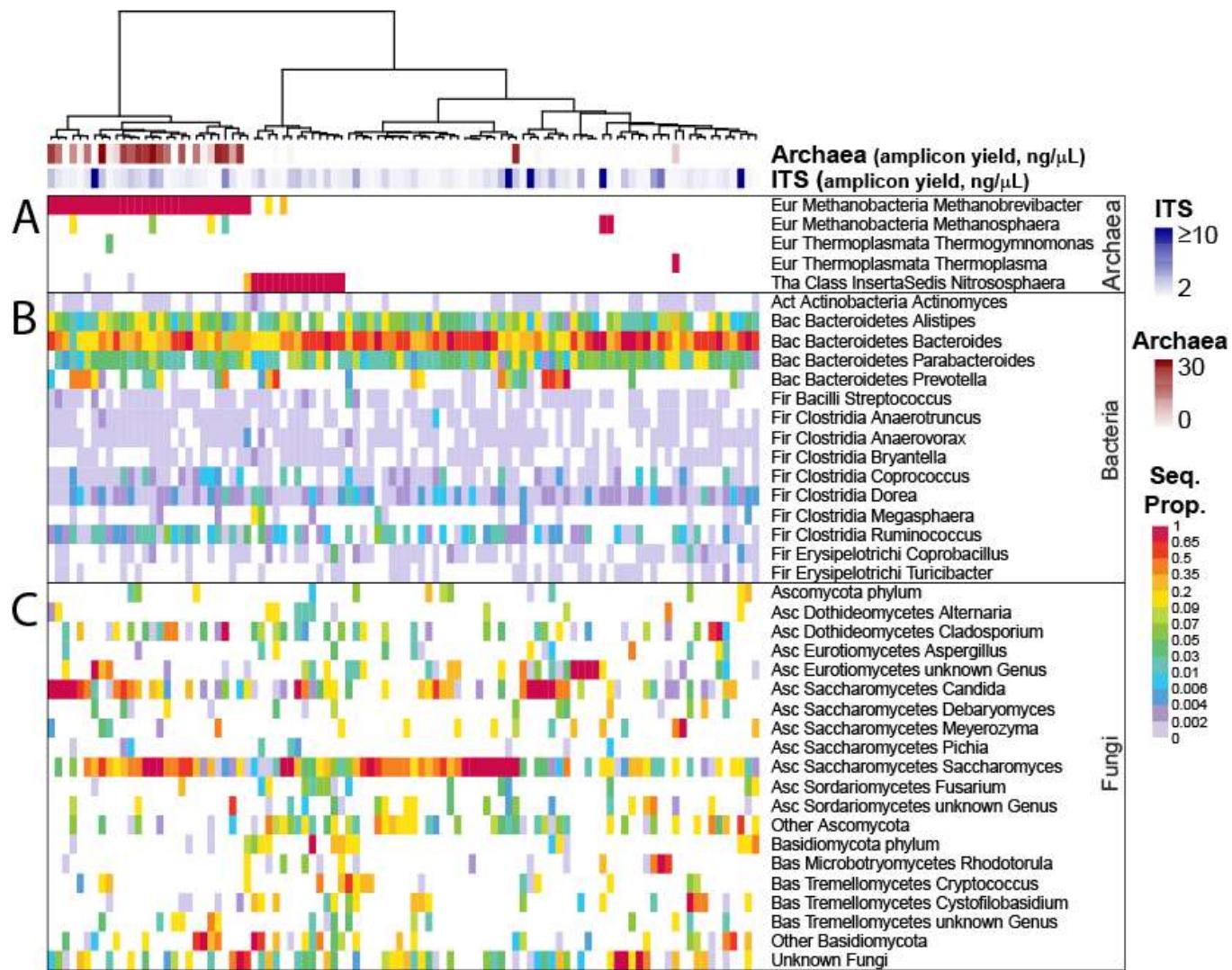
Diversity and Function

- ▶ Bacteria
- ▶ Differences in species composition
- ▶ Similarities in metabolic functions
- ▶ Phage?



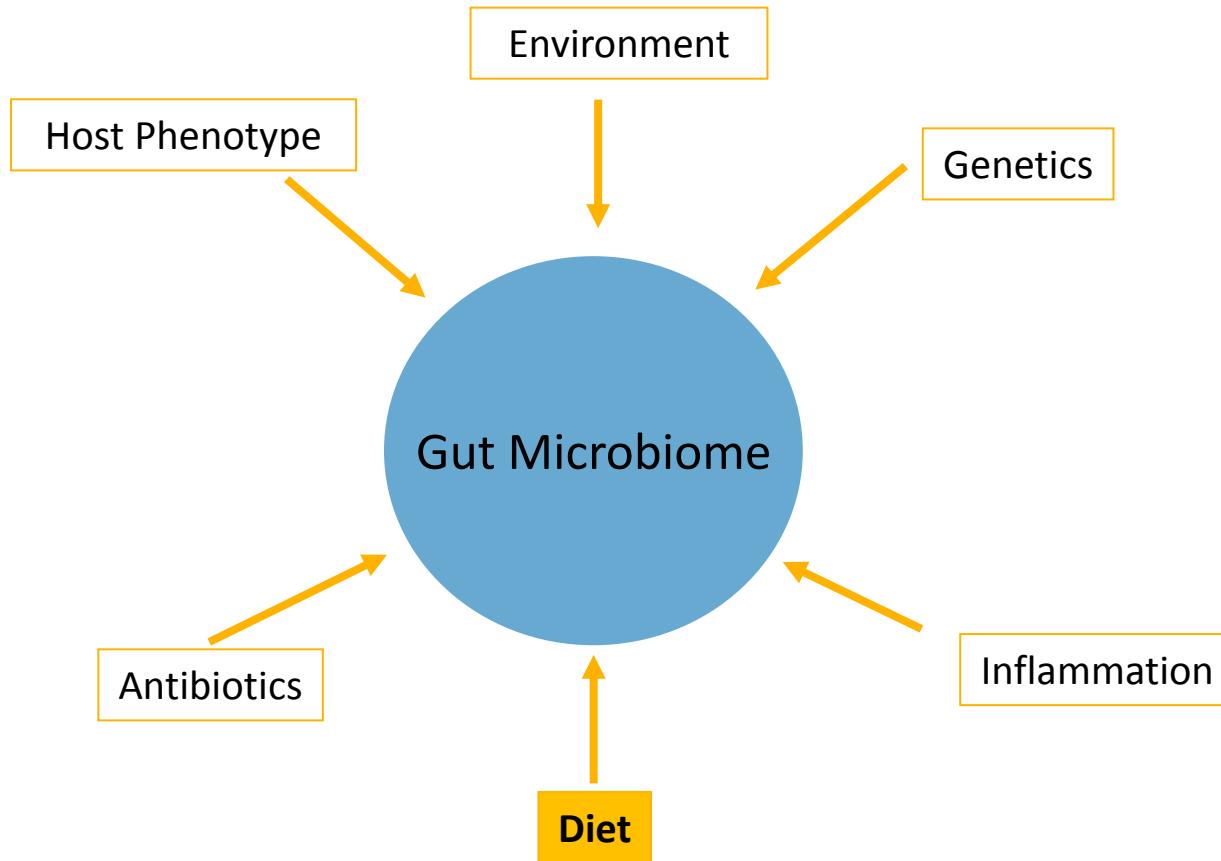
Diversity: Fungi and Archaea

- ▶ 5 archaea genera
- ▶ 66 fungi genera

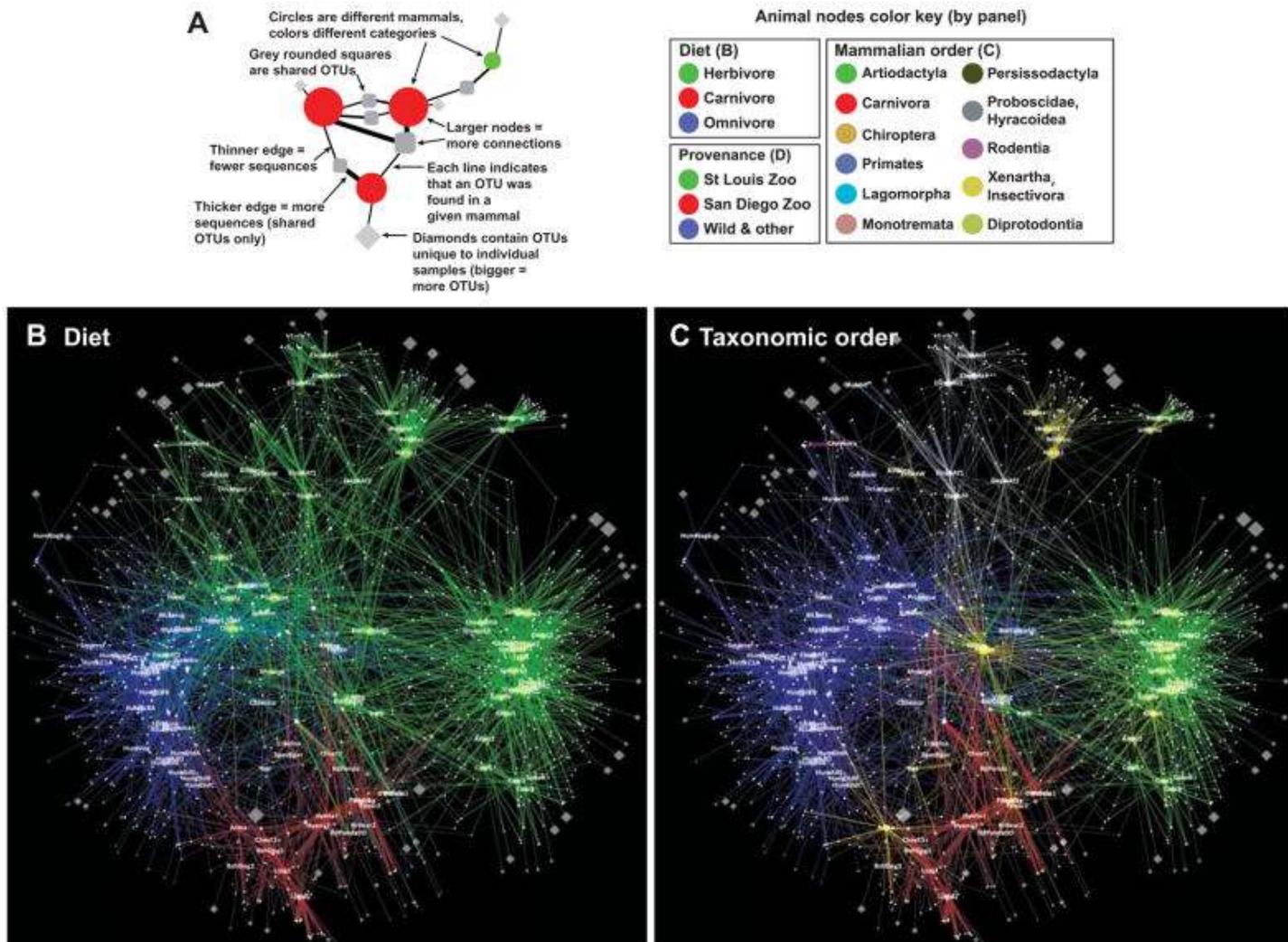


What Influences the Gut Microbiome?

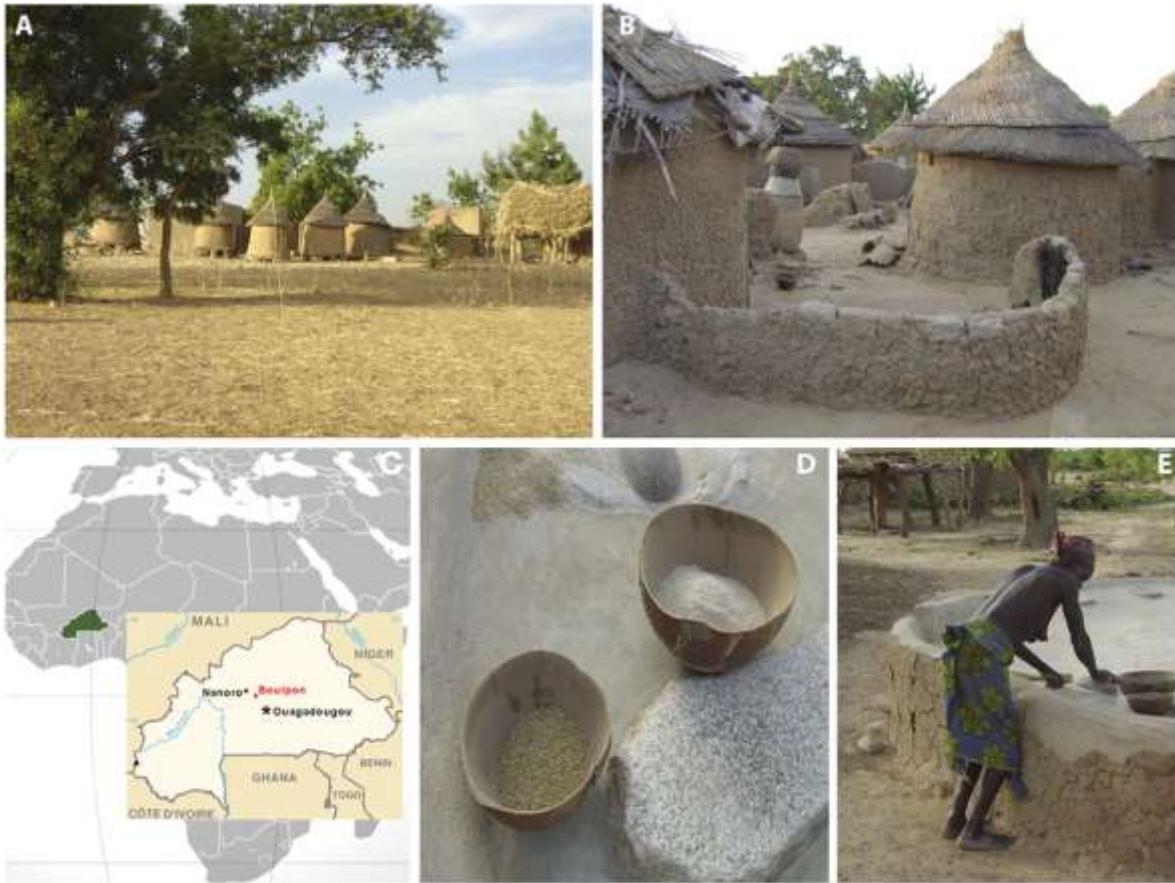
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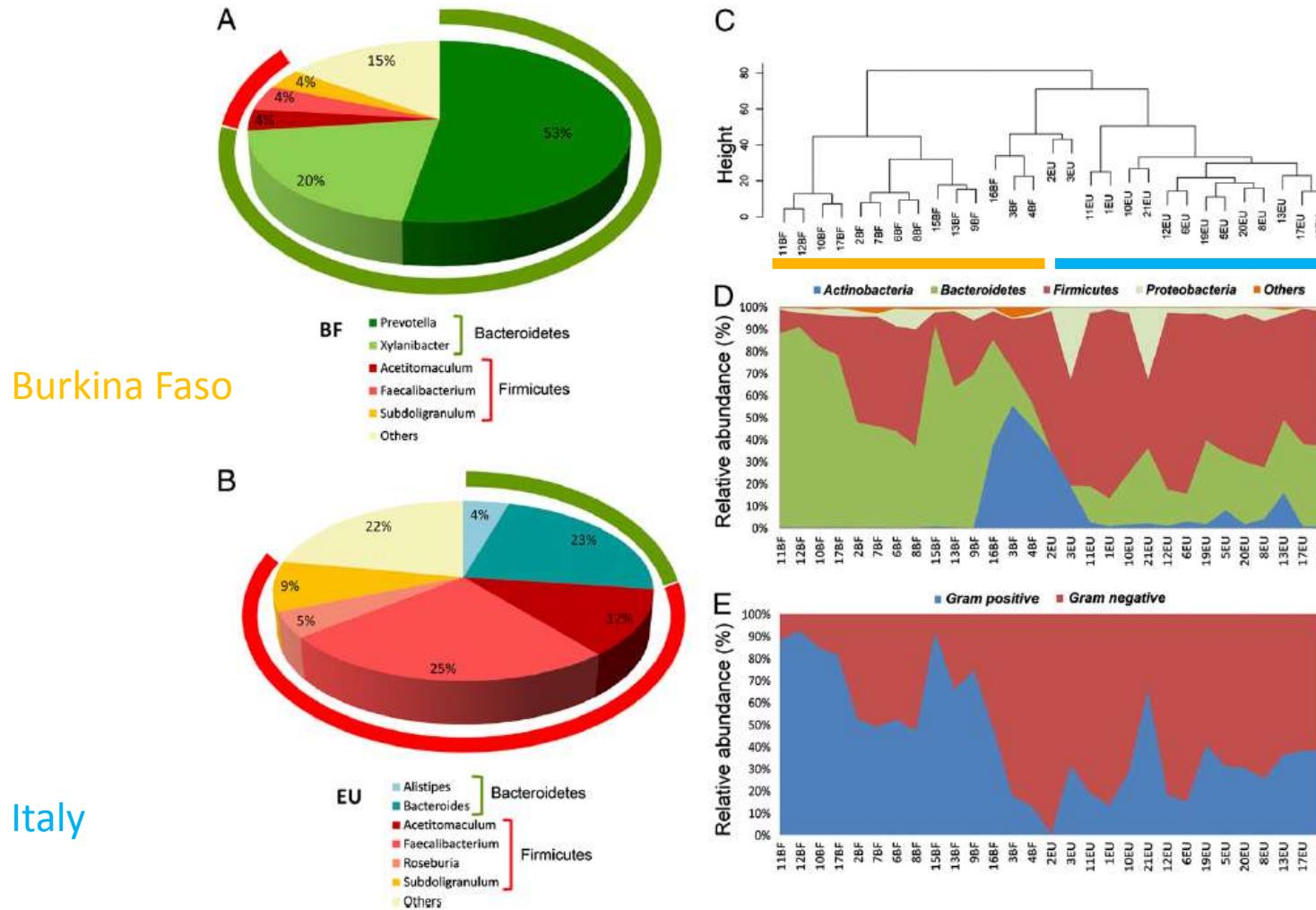
Microbiome, Diet e Co-evolution



Diet and the Microbiome: the case of Burkina Faso.

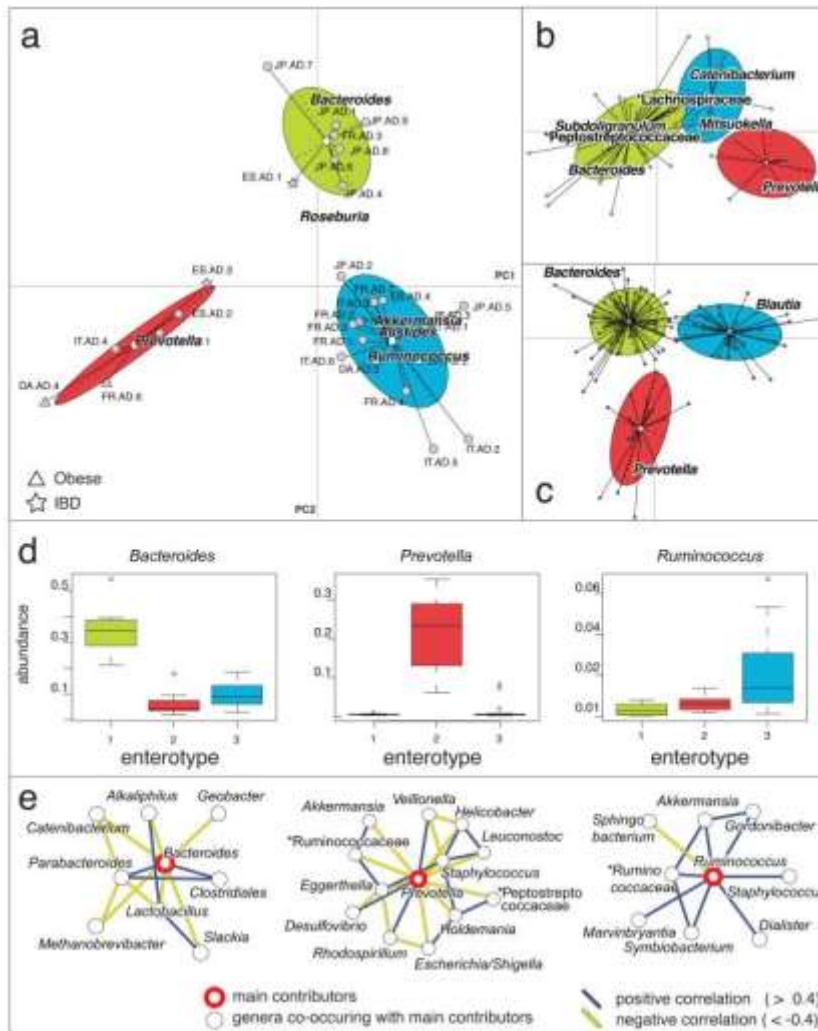


Diet and the Microbiome: the case of Burkina Faso.

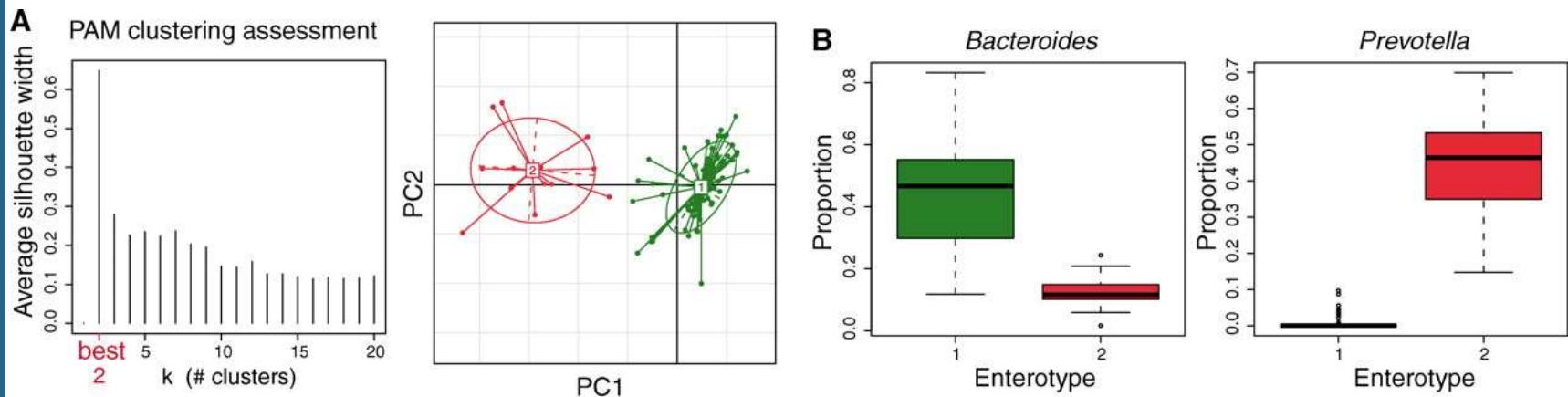


Italy

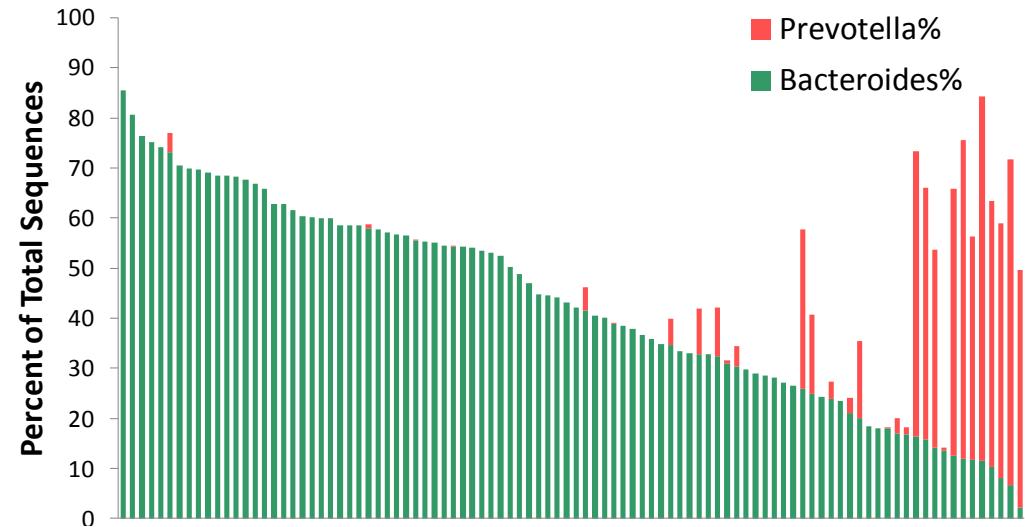
Microbiome variation: Enterotypes



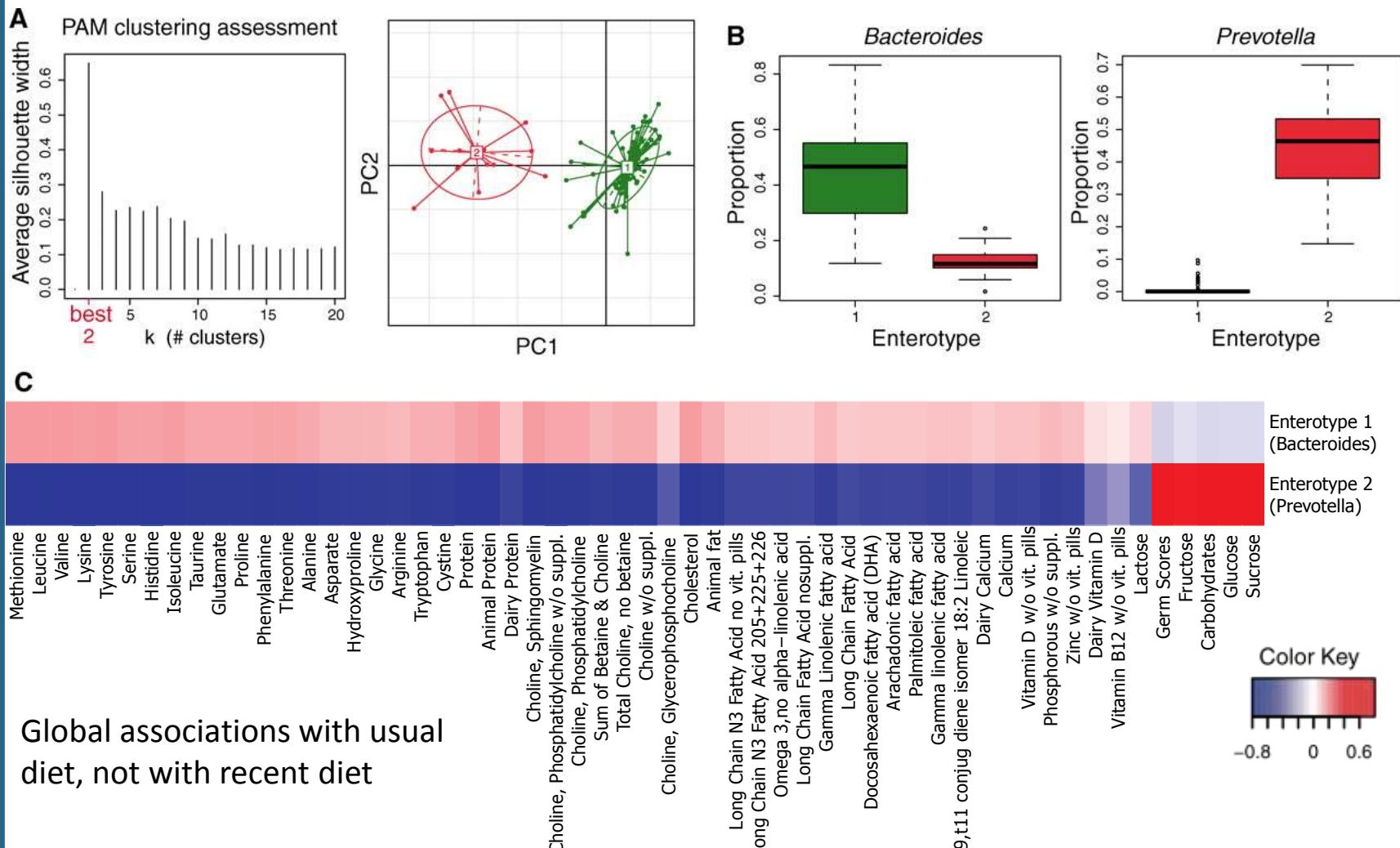
Microbiome variation: Enterotypes



► A balance between
Bacteroides e
Prevotella

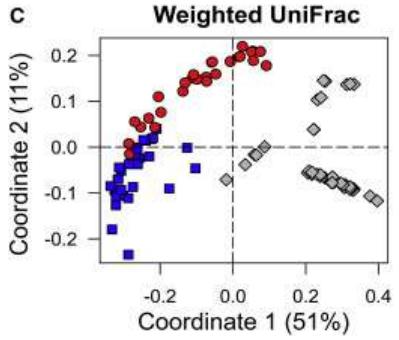
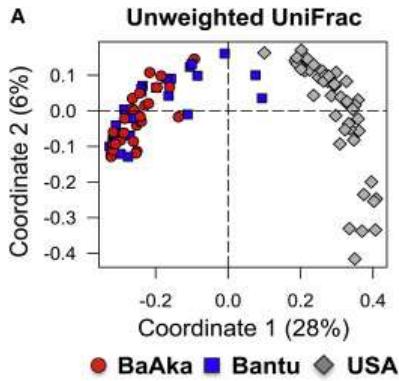


Enterotypes and human diet

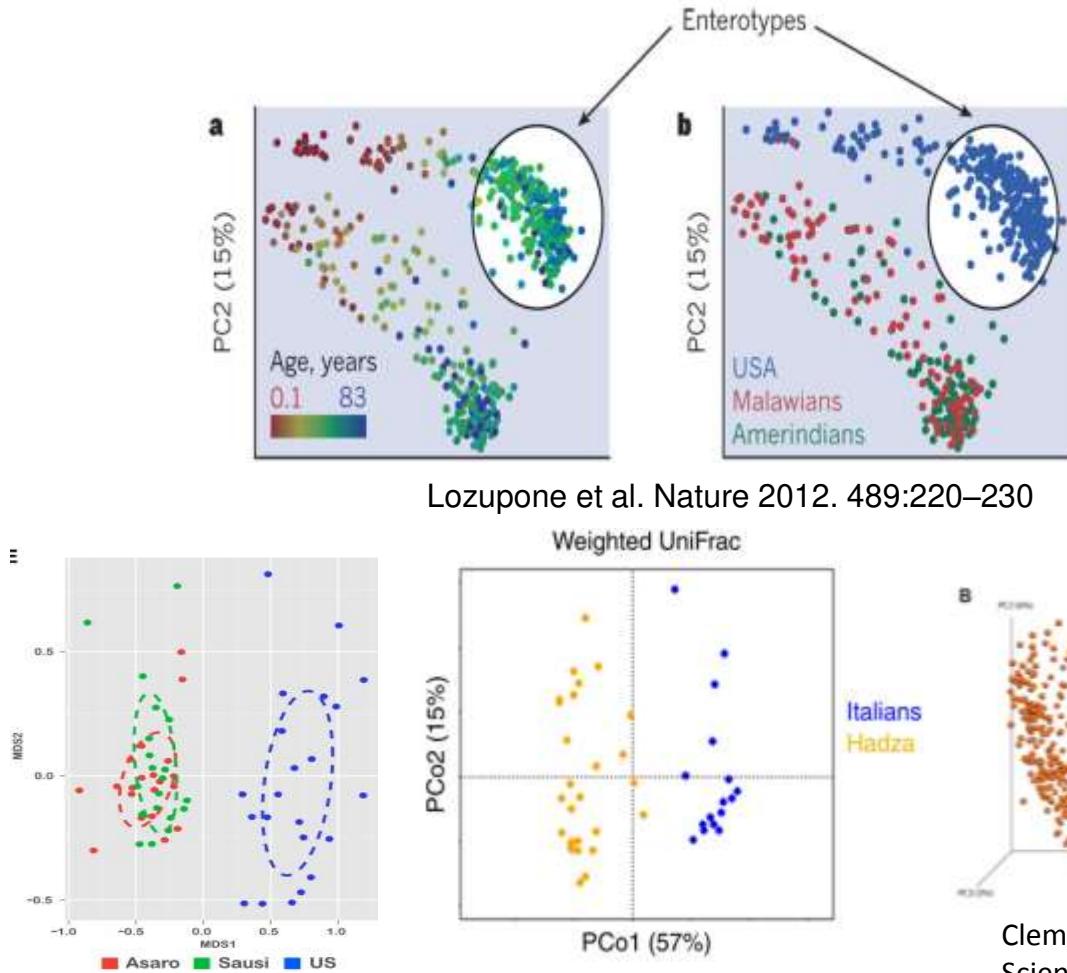


Global associations with usual diet, not with recent diet

Population variation



Gomez et al. 2016. Cell Reports 14(9):2142-2153



Martinez et al. 2015 Cell Reports 11(4):527–538

Schnor et al. 2014. Nature Communications 5:3654

Clemente et al. 2015. Science Advances 1(3):e1500183

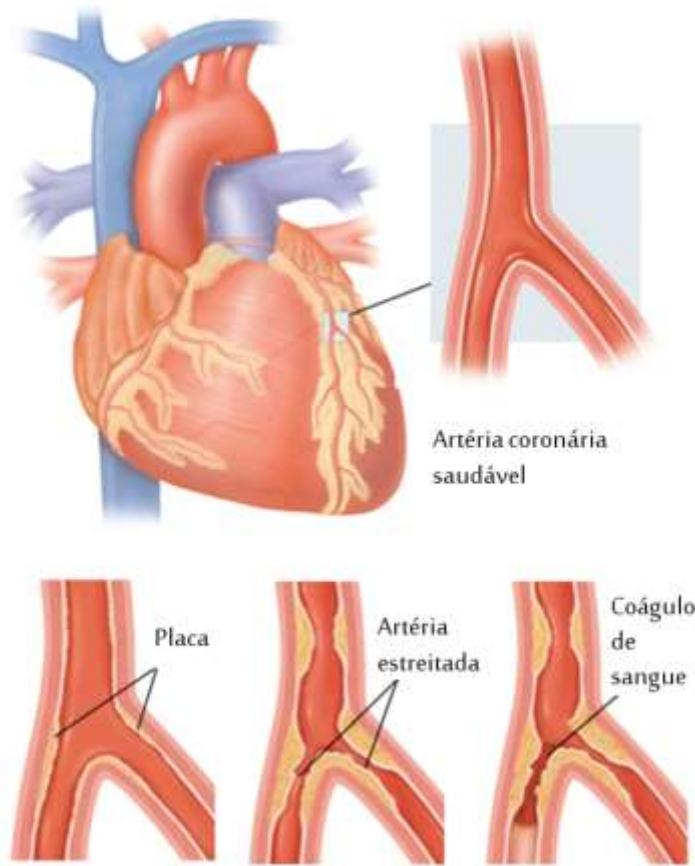
Cardiovascular disease: Atherosclerosis

- ▶ An example of where we are now



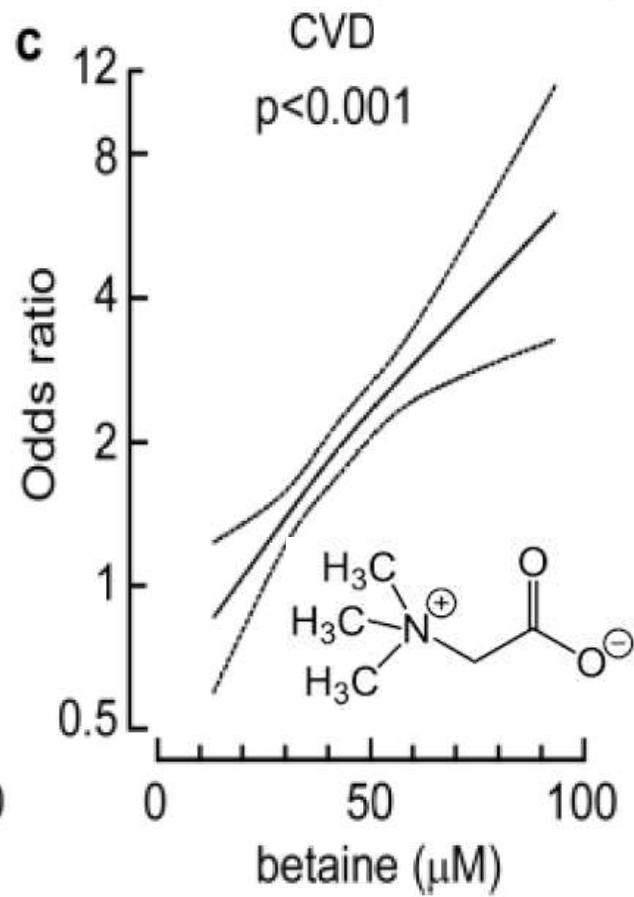
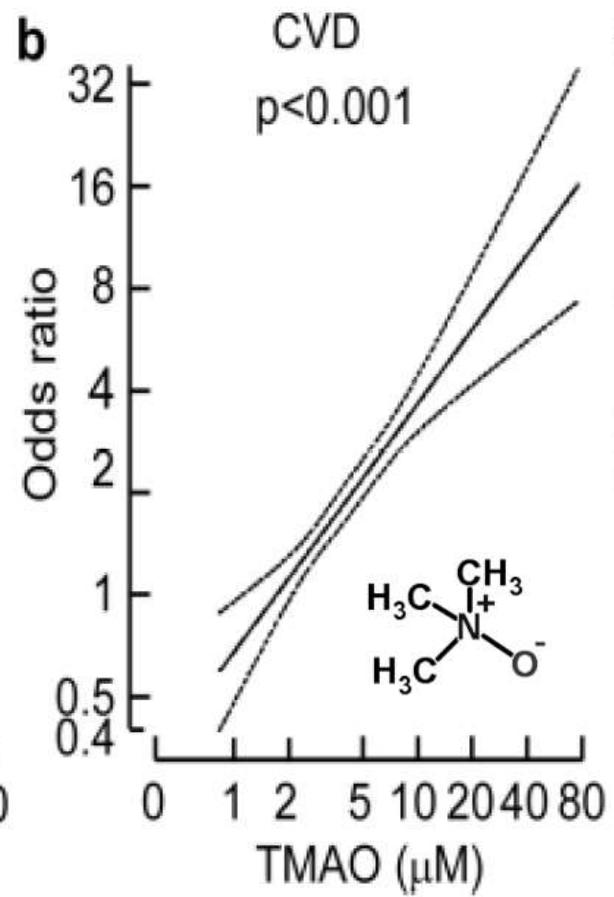
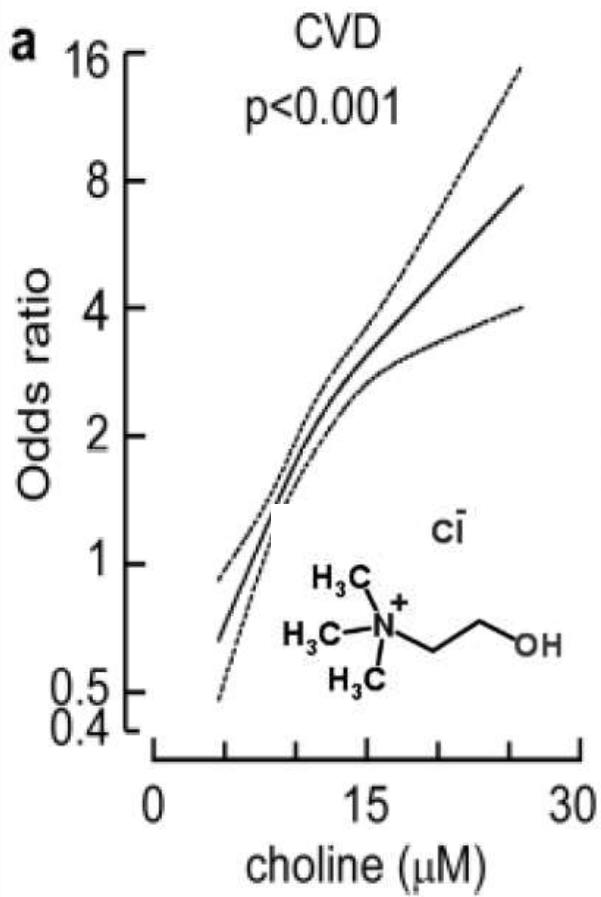
Cardiovascular disease: Atherosclerosis

- ▶ Risk factors:
 - ▶ High cholesterol
 - ▶ Obesity
 - ▶ Diabetes
 - ▶ Etc



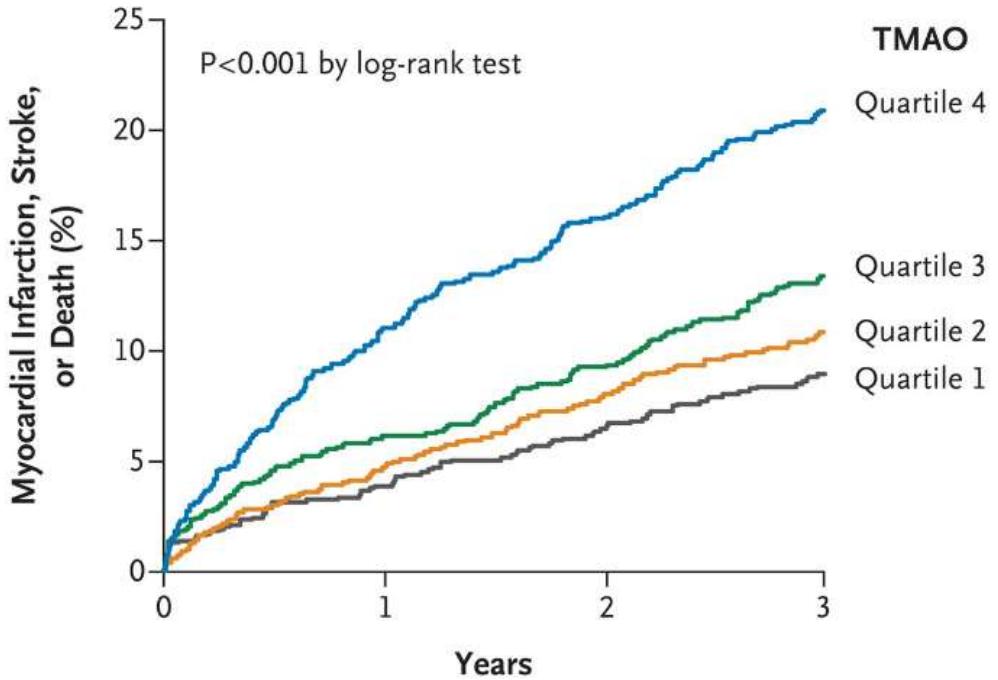
CVD risk

► N = 1865



CVD risk

- ▶ TMAO
- ▶ N=4007
- ▶ 3 year followup
- ▶ What is TMAO?

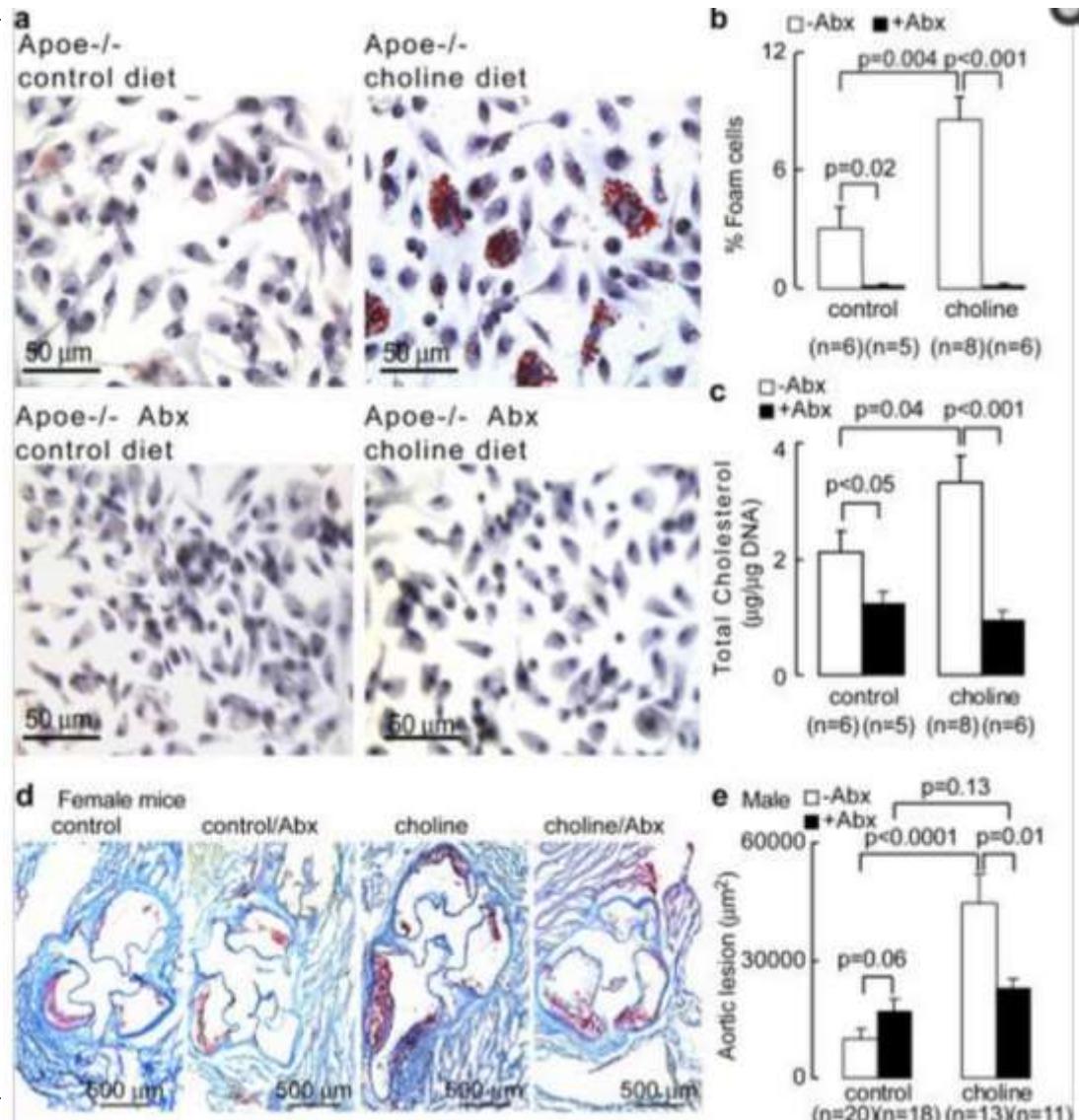


No. at Risk

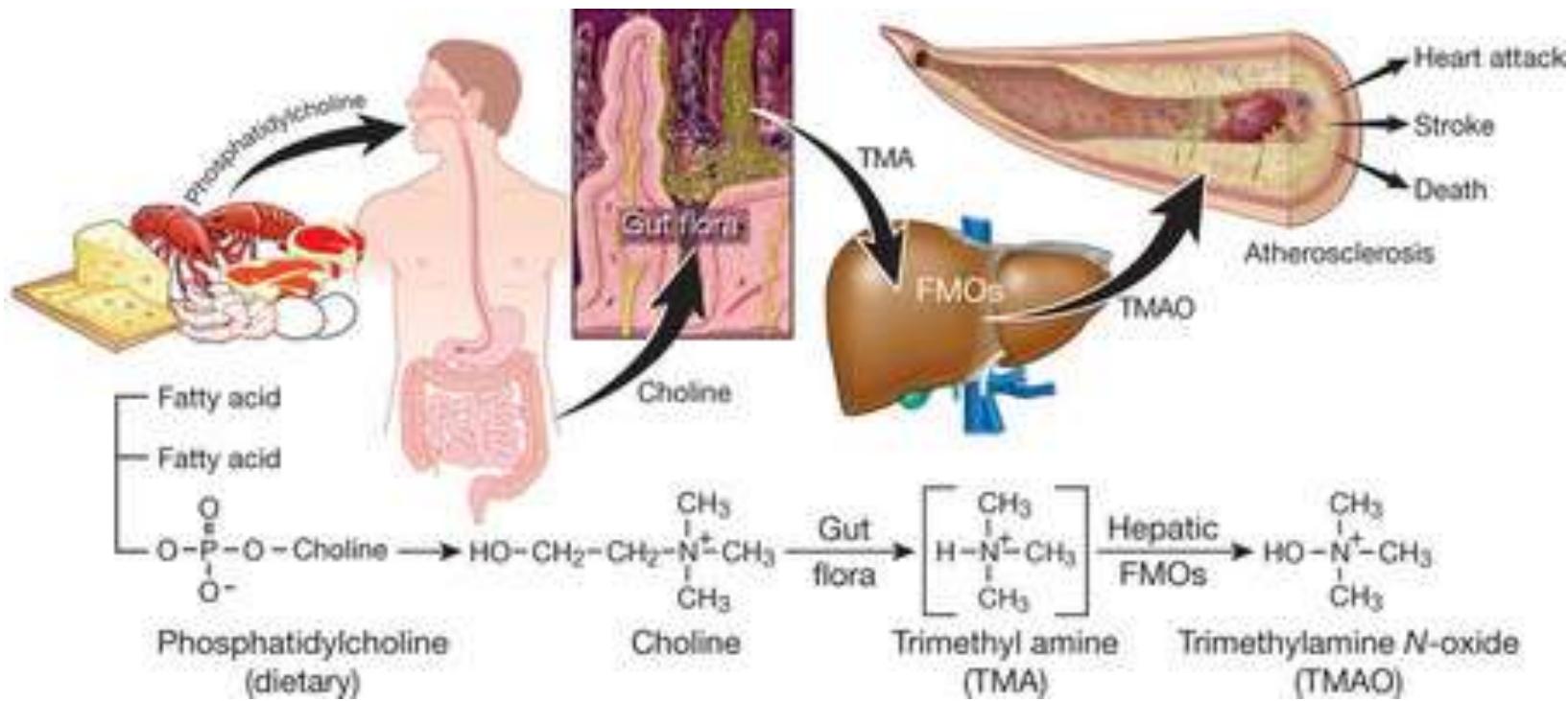
Quartile 1	1001	933	869	827
Quartile 2	998	940	884	843
Quartile 3	1003	938	888	835
Quartile 4	1005	913	849	791

Atherosclerosis

- ▶ Animal models:
 - ▶ Choline
 - ▶ Foam cells (macrophage),
 - ▶ With and without ABX



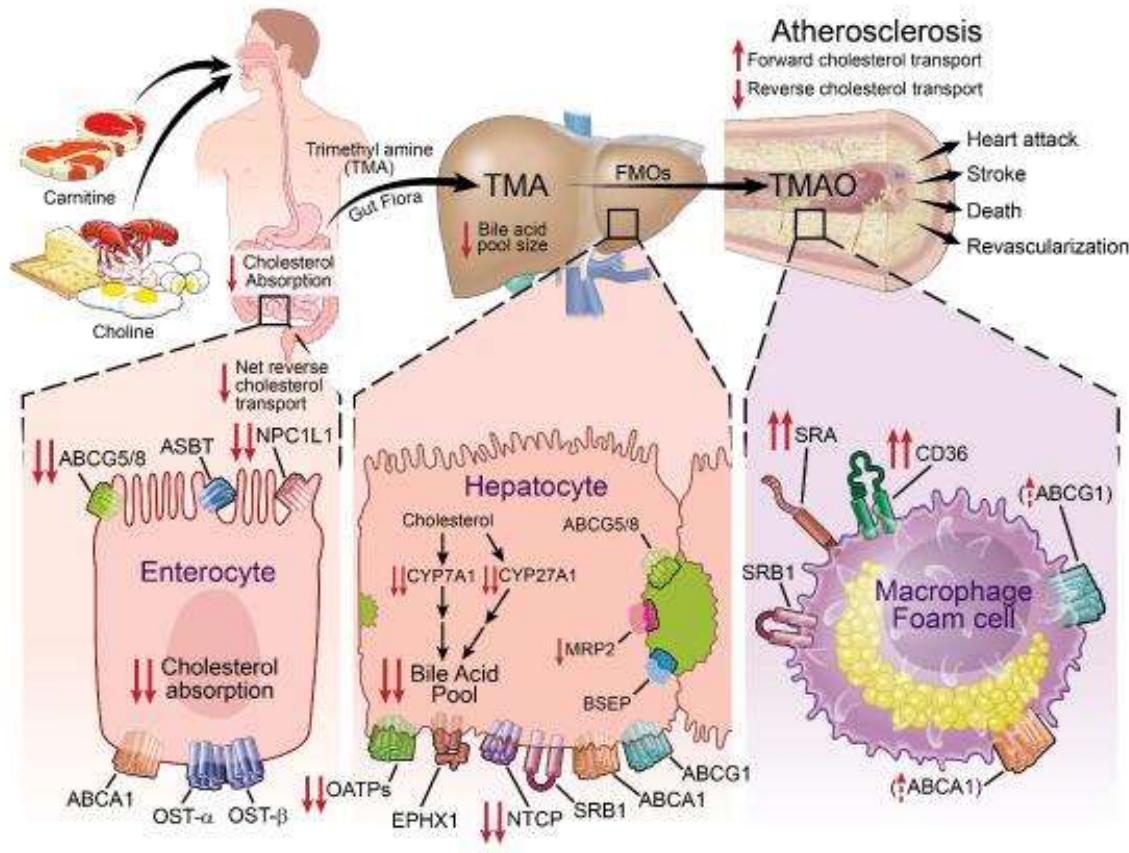
Atherosclerosis



Atherosclerosis

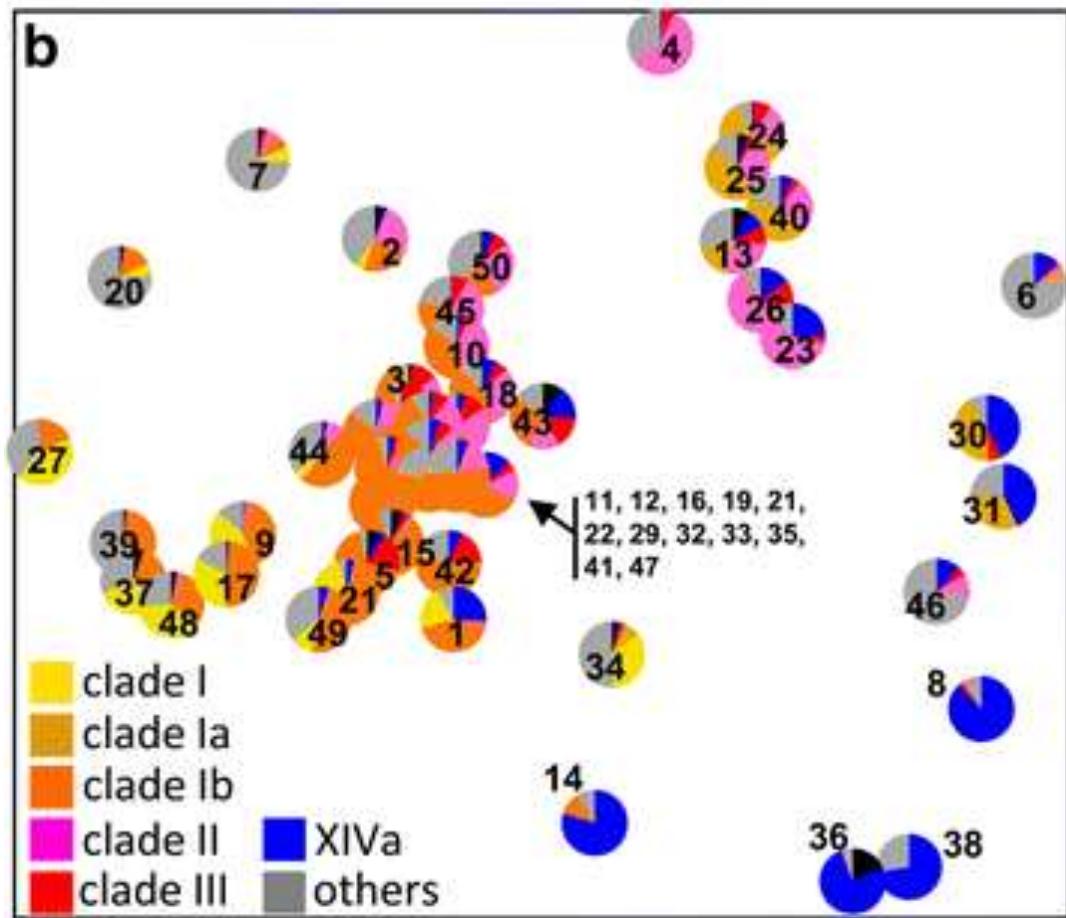
► Microbiome metabolism influences:

- Example: Choline/Carnitine → TMA→ TMAO → Atherosclerosis



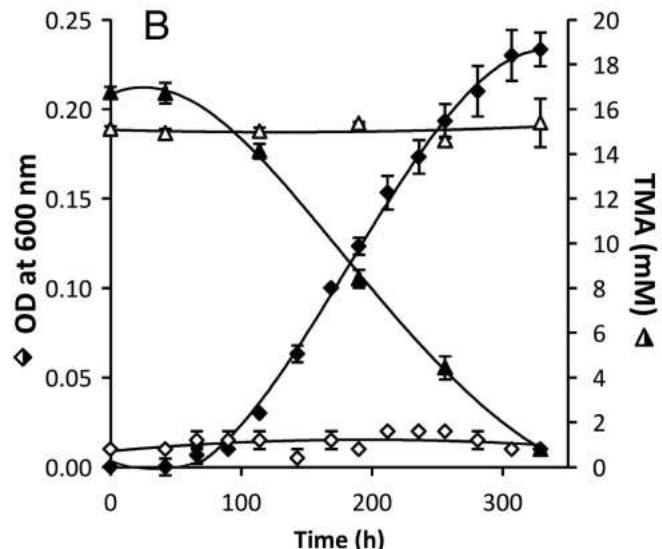
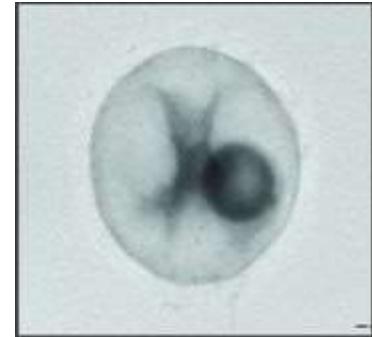
Atherosclerosis

- ▶ Many bacterial genes with the same function



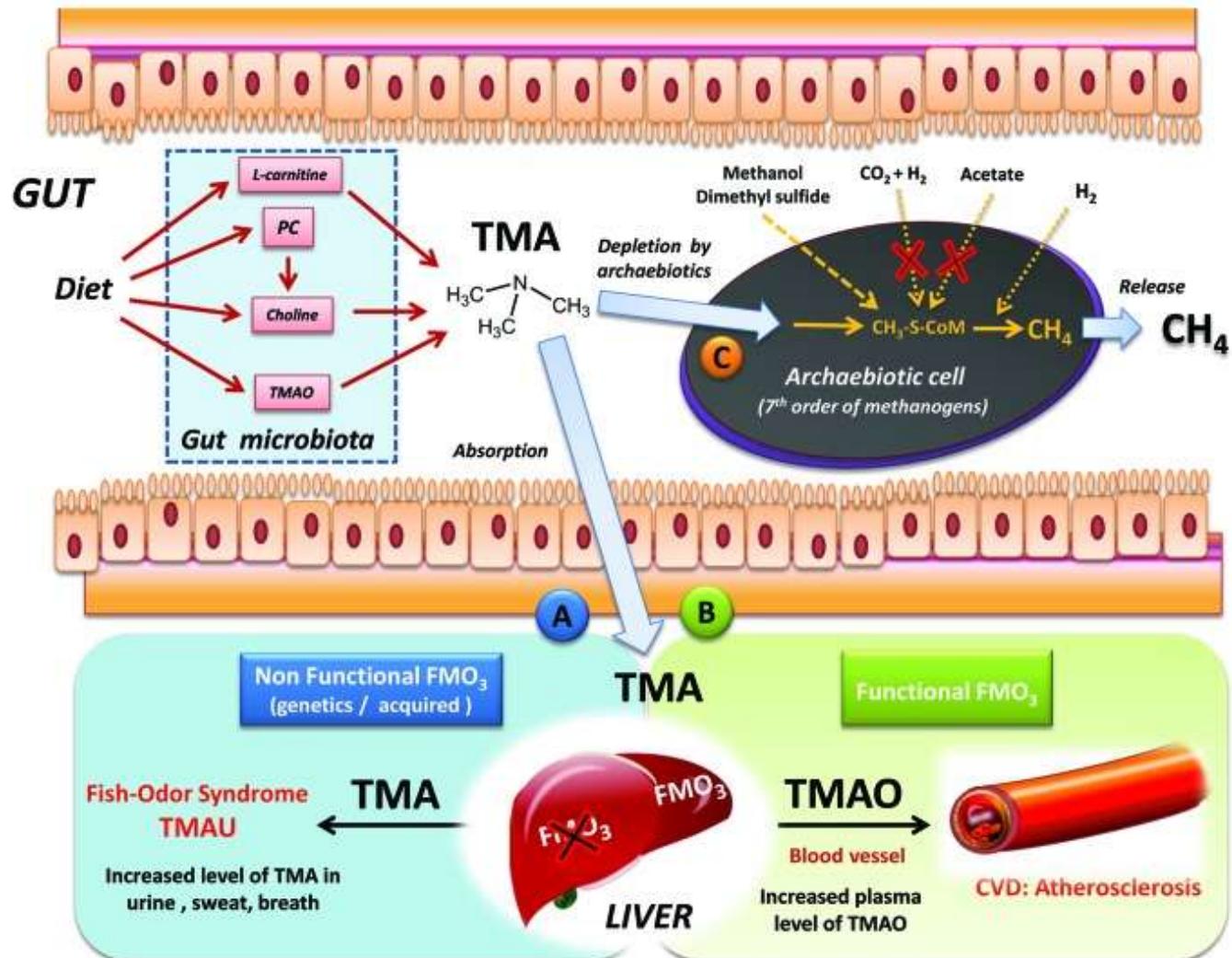
Archaeobiotic X Atherosclerosis

- ▶ *Methanomassiliicoccus luminyensis*
- ▶ Methanogenic
- ▶ Intestinal isolate
- ▶ Can grow using TMA producing methane



- ▲ [TMA] - TMA with H₂
- △ [TMA] - TMA w/out H₂
- ◆ *M.lum.* Growth with TMA / H₂
- ◇ *M.lum.* Growth w/out TMA / H₂

Archaeobiotic X Atherosclerosis



- ▶ Thanks!
 - ▶ c.hoffmann@usp.br
- ▶ Funding

