# Personalized Gut Mucosal Colonization Resistance to Empiric Probiotics Is Associated with Unique Host and Microbiome Features (*Cell*, September 6, 2018)

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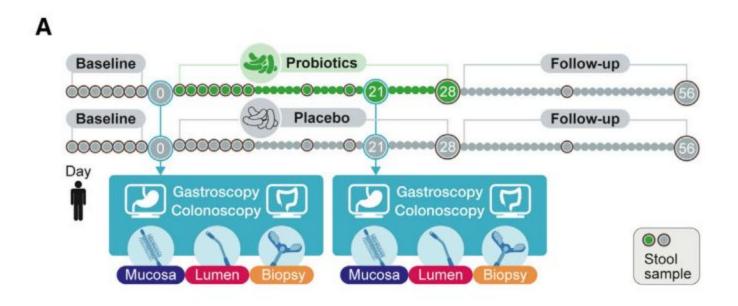
<u>Motivation</u>: Evidence of **probiotic gut mucosal colonization efficacy** remains sparse and controversial.

<u>Experiment:</u> Assessed murine and 15 human GI microbiome (2 endoscopy procedures along with fecal sampling) before and after 11-strain probiotic treatment.

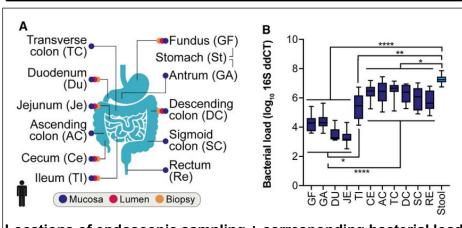
Conclusions:

- 1. Metagenomically characterized the murine and human mucosal-associated gastrointestinal microbiome and found it to **only partially correlate with stool microbiome**.
- 2. A sequential invasive multi-omics measurement at baseline and during consumption of an 11-strain probiotic combination or placebo demonstrated that probiotics remain viable upon gastrointestinal passage.
- Probiotics induced a transient, individualized impact on mucosal community structure and gut transcriptome. Collectively, empiric probiotics supplementation may be limited in universally and persistently impacting the gut mucosa, meriting development of new personalized probiotic approaches.

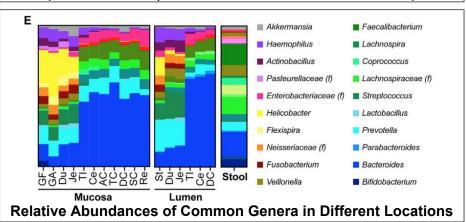
### **Experimental Outline in Humans**

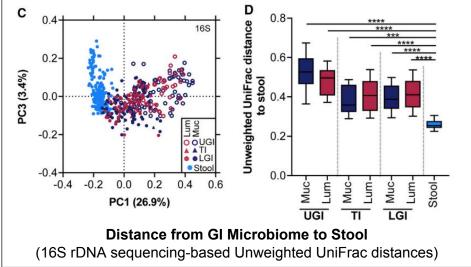


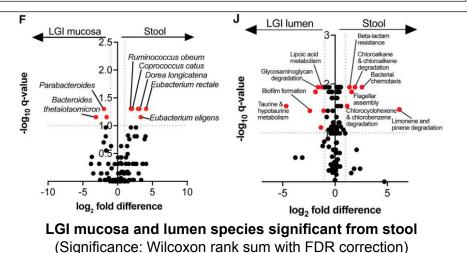
## Human Stool ≠ GI Bacteria



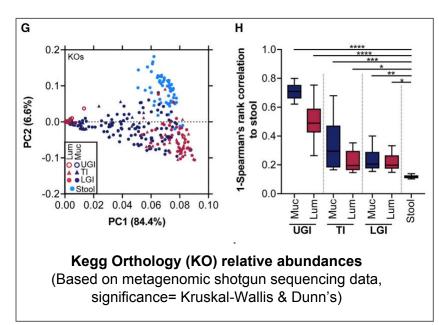
Locations of endoscopic sampling + corresponding bacterial load (Quantified with qPCR normalized to detection threshold)

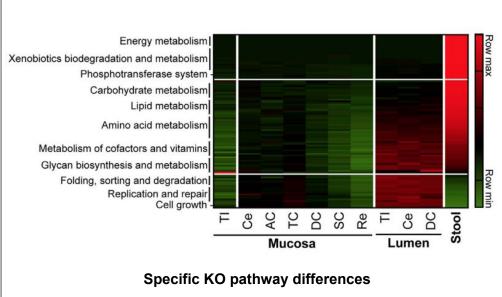




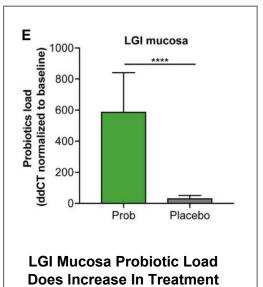


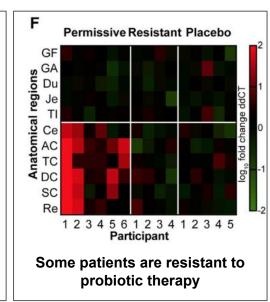
## Human Stool ≠ GI Bacteria: Functional Orthologs

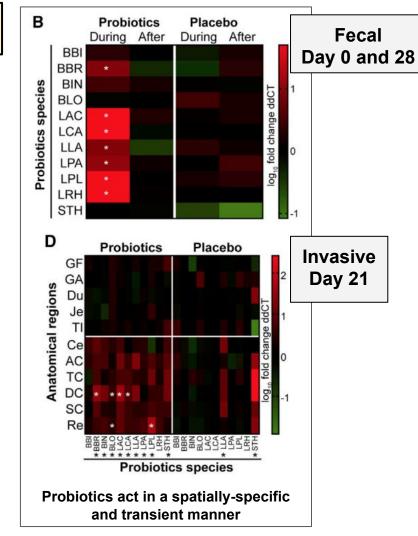




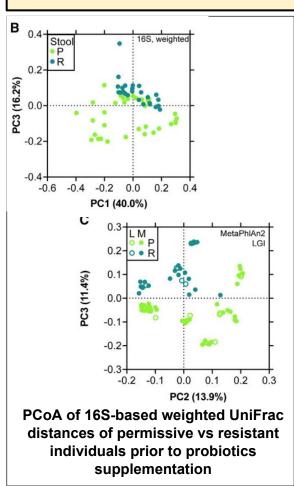
#### **Probiotic Treatment in Humans**

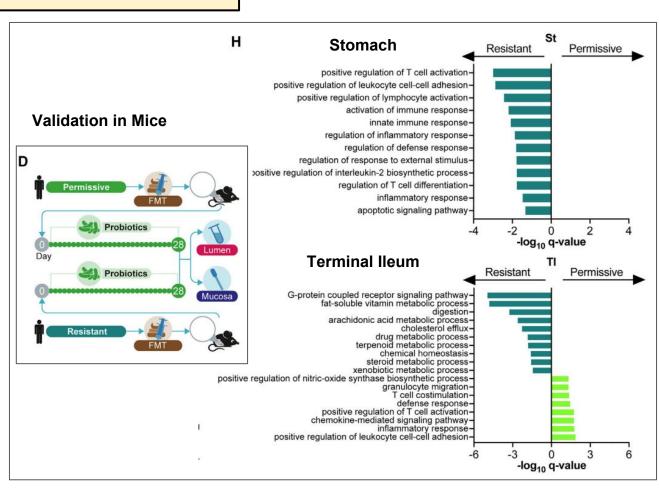






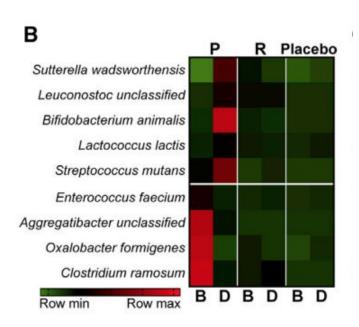
#### Permissive vs. Resistant



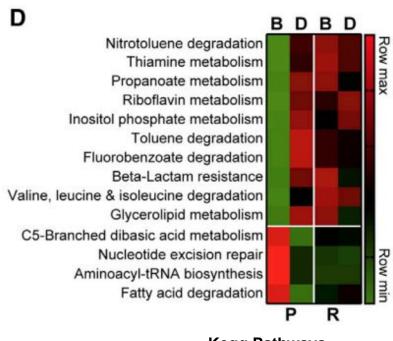


#### Permissive vs. Resistant Stool

**B** = Baseline and **D** = During Treatment



16S-based genera that bloomed or diminished in probiotics, but not in placebo



**Kegg Pathways**