PERSONALIZING PREBIOTIC REGIMENS for ASTRONAUTS

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MICROBIOME

• All animals are colonized by microbial communities



MICROBIOME

- All animals are colonized
 by microbial communities
- In humans:
 - ~100 trillion bacteria colonize epithelial surfaces
 - 100 billion bacteria per gram of fecal matter in the colon



GUT MICROBIOME IS DYNAMIC



David et al., *Genome Biol* 2014

DIET CAN BE A WAY TO MANIPULATE



Peter Turnbaugh







• Prebiotics are **indigestible dietary carbohydrates** known to stimulate gut microbiota

 Prebiotics are indigestible dietary carbohydrates known to stimulate gut microbiota

Body Site	Lag (days)	Host factor	Representative OTUs (#)	
	0	Stool:Hardness	Eggerthella/Clostridium(11)	
	0	Stool:TimeOfDay	Eggerthella/Clostridium(11)	0.27
	1	Nutrition:Fiber	Clostridium(6)	-0.38
	1	Nutrition:Fiber	Ruminococcaceae/F.prausnitzii(4)	-0.44
	1	Nutrition:Fiber	Eggerthella/Clostridium(11)	-0.39
	1	Nutrition:Fiber	Ruminococcus/R.gnavus/Clostridium(4)	-0.51
	1	Nutrition:Fiber	Ruminococcus/R.gnavus/Clostridium(5)	-0.51
	1	Nutrition:Fiber	Blautia(3)	-0.38
Subject A Gut	1	Nutrition:Fiber	Bifidobacteriales(13)	0.36
	1	Nutrition:Fiber	Coprococcus(8)	0.44
	1	Nutrition:Fiber	Clostridium(1)	-0.42
	1	Nutrition:Fiber	Ruminococcus/R.gnavus/Clostridium(6)	-0.44
	1	Nutrition:Fiber	Roseburia/E.rectale(30)	0.37
	1	Food:OrangeJuice	Clostridium(1)	0.28
	1	Food:BreakfastBar	Ruminococcus/R.gnavus/Clostridium(4)	-0.27
	1	Food:BreakfastBar	Ruminococcus/R.gnavus/Clostridium(5)	-0.40
	1	Food:BreakfastBar	Bifidobacteriales(13)	0.27
	1	Food:BreakfastBar	Clostridium(1)	-0.43
	1	Food:Yogurt	Bifidobacteriales(2)	0.45
	1	Food:Fruits:Fresh	Clostridiales(4)	-0.27
	1	Food:Fruits:Citrus	Ruminococcaceae/F.prausnitzii(4)	0.36
	1	Food:Soup	Clostridiales(1)	-0.25
	1	Food:Soup	Blautia(21)	-0.26
	1	Food:Soup:Other	Clostridiales(1)	-0.27
	1	Food:Soup:Other	Blautia(21)	-0.28

 Prebiotics are indigestible dietary carbohydrates known to stimulate gut microbiota

Body Site	Lag (days)	Host factor	Representative OTUs (#)	ρ	
0		Stool:Hardness	Eggerthella/Clostridium(11)	-0.30	
	0	Stool:TimeOfDoy Eggorthollo/Clootridium(11)		0.07	
	1	Nutrition:Fiber	Clostridium(6)	-0.38	
	1	Nutrition:Fiber	Ruminococcaceae/F.prausnitzii(4)	-0.44	1
	1	Nutrition:Fiber	Eggerthella/Clostridium(11)	-0.39	ĺ
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	1	Nutrition:Fiber	Ruminococcus/R.gnavus/Clostridium(5)	-0.51	
	1	Nutrition:Fiber	Blautia(3)	-0.38	
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	1	Nutrition:Fiber	Ruminococcus/R.gnavus/Clostridium(6)	-0.44	
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	1	Food:OrangeJuice	Clostridium(1)	0.28	
	1	Food:BreakfastBar	Ruminococcus/R.gnavus/Clostridium(4)	-0.27	
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	1	Food:BreakfastBar	Bifidobacteriales(13)	0.27	
	1	Food:BreakfastBar	Clostridium(1)	-0.43	
			0.15		
	1	Food:Fruits:Fresh	Clostridiales(4)	-0.27	
	1	Food:Fruits:Citrus	Ruminococcaceae/F.prausnitzii(4)	0.36	
	1	Food:Soup	Clostridiales(1)	-0.25	
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	1	Food:Soup:Other	Clostridiales(1)	-0.27	
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- Of interest because gut microbial fermentation leads to short-chain fatty acids like **butyrate**, a primary energy source for colonic epithelial cells

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- Of interest because gut microbial fermentation leads to short-chain fatty acids like **butyrate**, a primary energy source for colonic epithelial cells
- Also of relevance to spaceflight because shelfstable and tend to originate in plants

• **Dozens** of dietary fiber species exist

Prebiotics: Oligo- and Polysaccharides					
Amylopectin (corn)	Galactomannan	Lichenin	Pullulan		
Amylopectin (potato)	Glucomannan	Lignin	Rhamnogalacturonan I		
Arabinan	Glycogen	Oat spelt xylan	Starch (rice)		
Arabinogalactan	Gum (guar)	Pectic galactan (lupin)	Starch (corn)		
Arabinoxylan (wheat)	Gum (rosin)	Pectic galactan (potato)	Starch (wheat)		
Cellobiose	Heparin	Pectin (apple)	Starch (potato)		
Chitin	Hyaluronan	Pectin (citrus fruit)	Xyloglucan		
Chondroitin sulfate	Inulin	Pectin (citrus peel)	a-cellulose		
Dextran	Laminarin	Polygalacturonate	α-mannan		

- **Dozens** of dietary fiber species exist
- Each may vary in effect

	Butyrate			
Group ^{<i>b</i>}	Before	During	Change (%)	P value
Accessible starch (<i>n</i> = 39)	13 ± 6.1	15 ± 8.3	+13	0.18
Hi-Maize $(n = 43)$	9.3 ± 4.1	9.7 ± 5.6	+5	0.81
Potato $(n = 43)$	13 ± 6.0	16 ± 7.5	+29	<0.001
Inulin (<i>n</i> = 49)	11 ± 6.0	13 ± 7.0	+17	0.14

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- Individuals also vary in response



Venkataraman et al., Microbiome 2016

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How do we avoid provisioning astronauts with prebiotics that don't work?

GOALS OF OUR WORK

Answer three questions for astronauts:

- Which prebiotics should they take?
- Who should take them?
- And, when?

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CHALLENGES TO SCREENING PREBIOTICS

- Potentially a large number of prebiotics to screen
- Human studies can be logistically demanding
- In vivo physiological variation (e.g. SCFA absorption) may mask microbial response

A SIMPLE HOST-FREE PREBIOTIC ASSAY



Zack Holmes

PUMP: Prebiotic Utilization and Metabolite Production



Holmes et al., mBio 2020

OBSERVED SIGNIFICANT VARIATION BETWEEN PREBIOTICS



n=17 stool donors p < 0.0001, two-way ANOVA

Holmes et al., *<u>mBio</u>* 2020

YET, VARIATION BETWEEN DONORS LARGER

Grouped by **DONOR**





Average Total SCFA Concentration / Control

Holmes et al., mBio 2020

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Culture can be **hard to scale**:

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capture

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Culture can be hard to scale:

- Manual isolation can be tedious and time-consuming
- Automation may require expensive and/or bulky robotics that need to be enclosed anaerobically





90% of gut bacteria

Colony Picking Robot

Anaerobic Chamber

MICROFLUIDIC BACTERIAL CULTURE



Rachael Bloom

MICROFLUIDIC BACTERIAL CULTURE



Rachael Bloom



MICROFLUIDIC BACTERIAL CULTURE



Rachael Bloom





Max Villa





Max Villa











Phylum Firmicutes Bacteroidetes Actinobacteria Proteobacteria

EVERYONE HOSTS PREBIOTIC UTILIZERS



Villa*, Bloom*, et al., mSystems 2020

EVERYONE HOSTS PREBIOTIC UTILIZERS *But, people vary in abundance of utilizers*

across prebiotics



Villa*, Bloom*, et al., mSystems 2020

EVERYONE HOSTS PREBIOTIC UTILIZERS

But, people vary in abundance of utilizers across prebiotics



Villa*, Bloom*, et al., mSystems 2020

STUDY GOAL

 Test the hypothesis that individuals will vary in prebiotic metabolic potential, regardless of prebiotic

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- Test the hypothesis that individuals will vary in prebiotic metabolic potential, regardless of prebiotic
- Requires a *unique study design* where all individuals consume the same battery of prebiotics

CHoosing the OptiMal Prebiotic (CHOMP)





Zack Holmes

Heather Durand



Figure 2: Design for healthy human prebiotic trial. Study is uniform and balanced in period and frequency

Pre A: Inulin Pre B: Dextrin Pre C: Galactooligosaccharides n = 28 healthy
volunteers completed

ClinicalTrials.gov ID: NCT03595306

Holmes, et al., In prep

AGAIN, SIGNIFICANT VARIATION BETWEEN PREBIOTICS



IMPORTANTLY, INDIVIDUAL VARIATION CORRELATED ACROSS PREBIOTICS



WHAT DRIVES **INDIVIDUAL VARIATION?**

WHAT DRIVES INDIVIDUAL VARIATION?



WHAT DRIVES INDIVIDUAL VARIATION?

Can recent fiber intake affect prebiotic response?



GOALS OF OUR WORK

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HYPOTHESIS

Exposure to **prior doses** of a prebiotic will affect microbiota response **over time**



Silverman, Durand, et al, Microbiome 2018















IS A SINGLE DOSE SUFFICIENT IN PEOPLE?



Jeff Letourneau

n = 40 healthy volunteers

HOW DOES THIS WORK?



Within 6 hrs of prebiotic exposure, human gut bacteria activate polysaccharide utilization loci



• Prebiotics vary in effect on gut microbiota

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- But, stronger individual variation in response
- Variation can be linked to gut microbial function, which in turn is linked to prior diet
- Even initial exposure to prebiotics will alter response to subsequent doses

Three questions:

1. Which prebiotics should they take?

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 - Might not matter too much.

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- 1. Which prebiotics should they take?
 - Might not matter too much.
- 2. Who should take them?
 - Greatest benefit among those who normally eat the least fiber
- 3. And, when?
 - Poor responders may start shortly before spaceflight

ONGOING WORK

What are the effects of prebiotics on cognition and mental performance?





Ken Racicot



Mathias Basner

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