Microbiome Analysis Overview

Josh Granek

Housekeeping

- Me
- Names
- Test Wifi and IBIEM Computing Environment
- Bootcamp Daily Groups

Bootcamp Overview

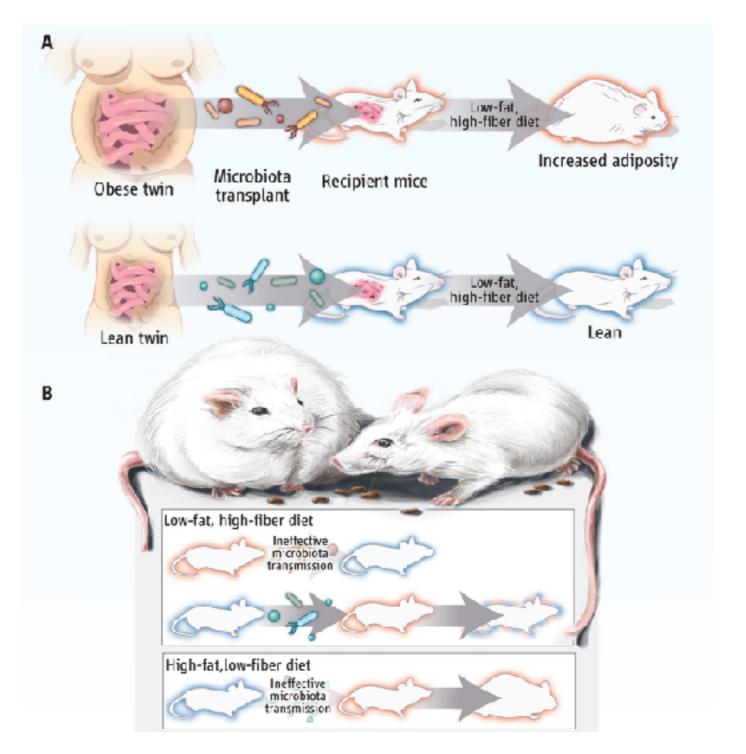
Microbiome Analysis Overview

Josh Granek

Microbiomes in Human Health

- Acne
- Asthma/allergies
- Autism
- Autoimmune diseases
- Cancer
- Dental cavities
- Depression and anxiety
- Diabetes
- Eczema
- Gastric ulcers
- Hardening of the arteries
- Inflammatory bowel diseases
- Malnutrition
- Obesity
- Parkinson's Disease
- Drug Metabolism
- Vaccine Effectiveness
- ...

Causation



Metagenomics

	What	Information	Analogy	Target Size	Cost
Amplicon	Marker Gene	Who is Present	Name	100bp - 1kb	Low
Shotgun Metagenome	Genomes	What Genes are Present	CV	100kb - 100Mb	High
Shotgun Metatranscriptome	All RNA	What Genes are Expressed	Twitter Feed	100kb - 100Mb	High

Fungal genome and mating system transitions facilitated by chromosomal translocations involving intercentromeric recombination.

Sun S, Yadav V, Billmyre RB, Cuomo CA, Nowrousian M, Wang L, Souciet JL, Boekhout T, Porcel B, Wincker P, **Granek JA**, Sanyal K, Heitman J. PLoS Biol. 2017 Aug 11;15(8):e2002527. doi: 10.1371/journal.pbio.2002527. eCollection 2017 Aug.

PMID: 28800596 Free PMC Article

Similar articles

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2.

Elucidation of the calcineurin-Crz1 stress response transcriptional network in the human fungal pathogen Cryptococcus neoformans.

Chow EW, Clancey SA, Billmyre RB, Averette AF, Granek JA, Mieczkowski P, Cardenas ME, Heitman J.

PLoS Genet. 2017 Apr 4;13(4):e1006667. doi: 10.1371/journal.pgen.1006667. eCollection 2017 Apr.

PMID: 28376087 Free PMC Article

Similar articles

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3.

Evidence for distinct brain networks in the control of rule-based motor behavior.

Granek JA, Sergio LE.

J Neurophysiol. 2015 Aug;114(2):1298-309. doi: 10.1152/jn.00233.2014. Epub 2015 Jul 1.

PMID: 26133796 Free PMC Article

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4.

Rapid mapping of insertional mutations to probe cell wall regulation in Cryptococcus neoformans.

Esher SK, **Granek JA**, Alspaugh JA.

Fungal Genet Biol. 2015 Sep;82:9-21. doi: 10.1016/j.fgb.2015.06.003. Epub 2015 Jun 23.

PMID: 26112692 Free PMC Article

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5.

Integrating chemical mutagenesis and whole-genome sequencing as a platform for forward and reverse genetic analysis of Chlamydia.

Kokes M, Dunn JD, **Granek JA**, Nguyen BD, Barker JR, Valdivia RH, Bastidas RJ. Cell Host Microbe. 2015 May 13;17(5):716-25. doi: 10.1016/j.chom.2015.03.014. Epub 2015 Apr 23.

PMID: 25920978 Free PMC Article

Similar articles

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6.

Antifungal drug resistance evoked via RNAi-dependent epimutations.

Calo S, Shertz-Wall C, Lee SC, Bastidas RJ, Nicolás FE, **Granek JA**, Mieczkowski P, Torres-Martínez S, Ruiz-Vázquez RM, Cardenas ME, Heitman J.

Nature. 2014 Sep 25;513(7519):555-8. doi: 10.1038/nature13575. Epub 2014 Jul 27.

PMID: 25079329 Free PMC Article

Similar articles 1 comment Select item 24392035

7.

Decoupled visually-guided reaching in optic ataxia: differences in motor control between canonical and non-canonical orientations in space.

Granek JA, Pisella L, Stemberger J, Vighetto A, Rossetti Y, Sergio LE.

PLoS One. 2013 Dec 31;8(12):e86138. doi: 10.1371/journal.pone.0086138. eCollection 2013.

PMID: 24392035 Free PMC Article

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Q

Amplicon Sequencing

PCR amplify and sequence a marker gene

	Marker Gene		
Bacteria	16s rRNA		
Fungi	18s or ITS rRNA		
Archaea	16s rRNA		
Protozoa	18s rRNA		
Viruses	?????		

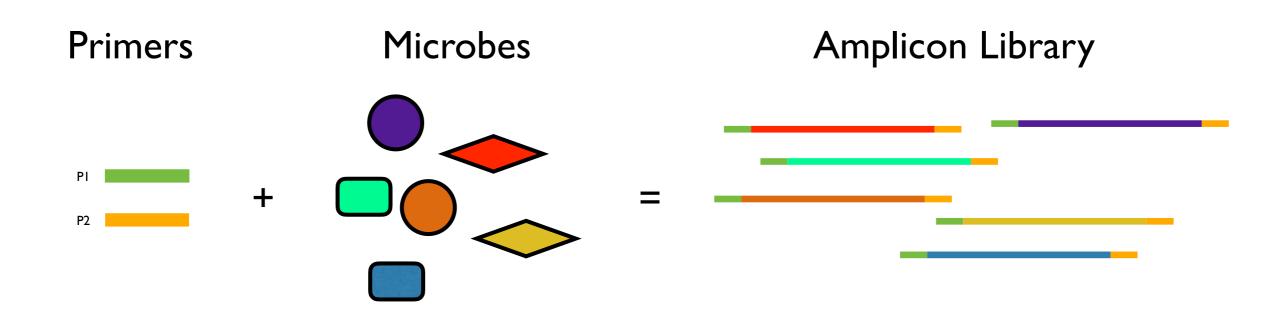
Metagenomics

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Metagenomics

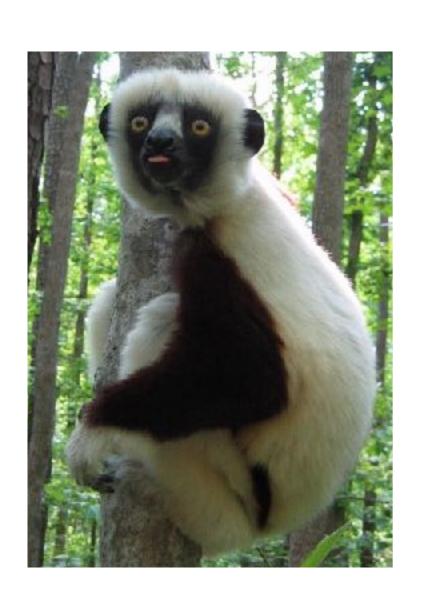
	What	Information	Analogy	Target Size	Cost	Discovery?
Amplicon	Marker Gene	Who is Present	Name	100bp - 1kb	Low	+/-
Shotgun Metagenome	Genomes	What Genes are Present	CV	100kb - 100Mb	High	++
Shotgun Metatranscriptome	All RNA	What Genes are Expressed	Twitter Feed	100kb - 100Mb	High	++

Amplicon Sequencing



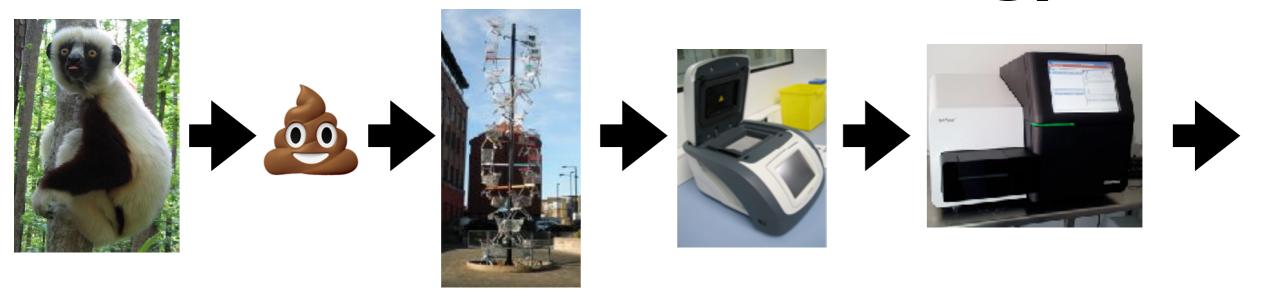
I 6s Amplicon Analysis Overview

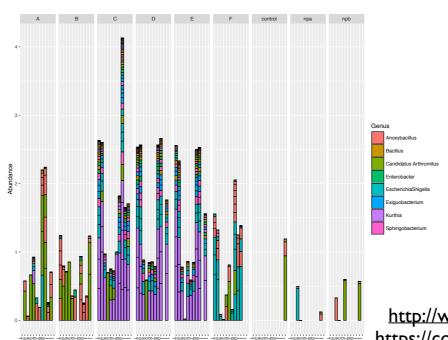
Big Picture



- I. What is present?
- 2. How much?
- 3. Are there differences between treatments, host species, ...?
- 4. What are the differences?

Molecular Biology





http://www.geograph.org.uk/photo/2847164 https://commons.wikimedia.org/wiki/File:Pcr.jpg

Bioinformatic Analysis





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ABBBABBBAFFFGGGGGGGGGGGGGGGCG2GF3FFGHHHHHHGGFGHEHHGGGEHHHHAGGHHGHHHFFDHFHHHGEGGGG@F@H?GHH/GBEFGGG @M00698:36:000000000-AFBEL:1:1101:16483:1412 1:N:0:0

CTGCCAGTTGAACGACGGCGAGCAGTTATAAGCCAGCAGTTTGCCCGGATATTTCGCGTGGATAGCTTGTGCAAAGCGACGCCAGTTCCAGATCCGGCG

GTAAAGTCCTGAGTGATACCGGCAACTTTTACCCCCAGTCCCACTTTCGAACCGGCAAACATATCGGCAAAAGAGGCCGTGCCTGATTTAAAGCCGTAGGT

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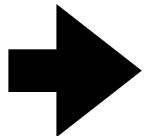


	Sample 1	Sample 2	 Sample N
Bacteria 1			
Bacteria 2			
Bacteria N			

Statistical Analysis

	Sample 1	Sample 2	 Sample N
Bacteria 1			
Bacteria 2			
Bacteria N			

- I. What is present?
- 2. How much?



- 3. Are there differences between treatments, host species, ...?
- 4. What are the differences?

Caveat





Bioinformatic Analysis





@M00698:36:000000000-AFBEL:1:1101:14738:1412 1:N:0:0

TTACGCTAACAGGCGGTAGCCTGGCAGGGTCAGGAAATCAATTAACTCATCGGAAGTGGTGATCTGTTCCATCAAGCGTGCGGCATCGTCAAAACGCCC

ABBBABBBAFFFGGGGGGGGGGGGGGGCG2GF3FFGHHHHHHGGFGHEHHGGGEHHHHAGGHHGHHHFFDHFHHHGEGGGG@F@H?GHH/GBEFGGG @M00698:36:000000000-AFBEL:1:1101:16483:1412 1:N:0:0

CTGCCAGTTGAACGACGGCGAGCAGTTATAAGCCAGCAGTTTGCCCGGATATTTCGCGTGGATAGCTTGTGCAAAGCGACGCCAGTTCCAGATCCGGCG

GTAAAGTCCTGAGTGATACCGGCAACTTTTACCCCCAGTCCCACTTTCGAACCGGCAAACATATCGGCAAAAGAGGCCGTGCCTGATTTAAAGCCGTAGGT

+



	Sample 1	Sample 2	 Sample N
Bacteria 1			
Bacteria 2			
Bacteria N			

The End