

# Microbiome Overview

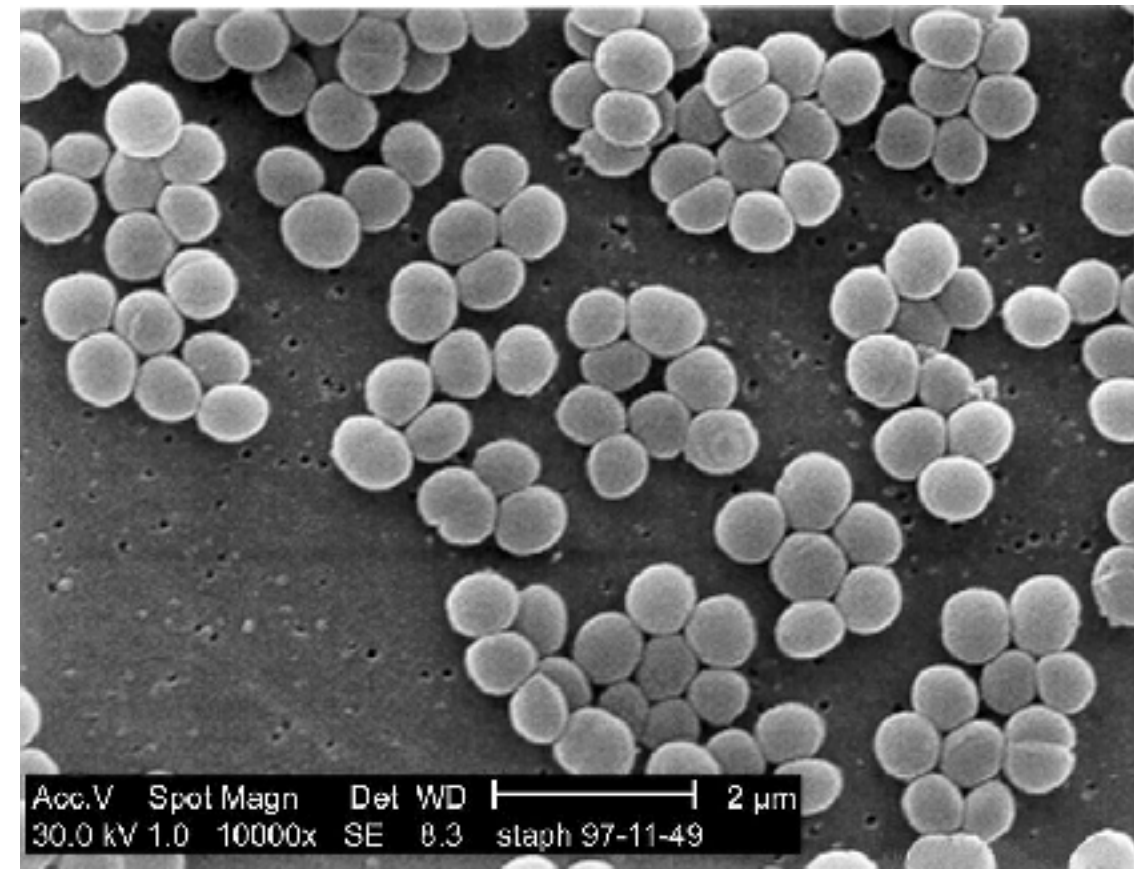
Josh Granek

# Microbiome

- The collection of microorganisms (microbes) living in an environment

# Microorganism

- microscopic organisms



# Microbes: Complexity

**Humans** Supercomputer

**Fungi\***

Tablet

Not as powerful or complex as a supercomputer, but able to do lots of stuff on its own

**Bacteria**

Cell Phone

Less complicated than Fungi, but still able to do stuff on its own

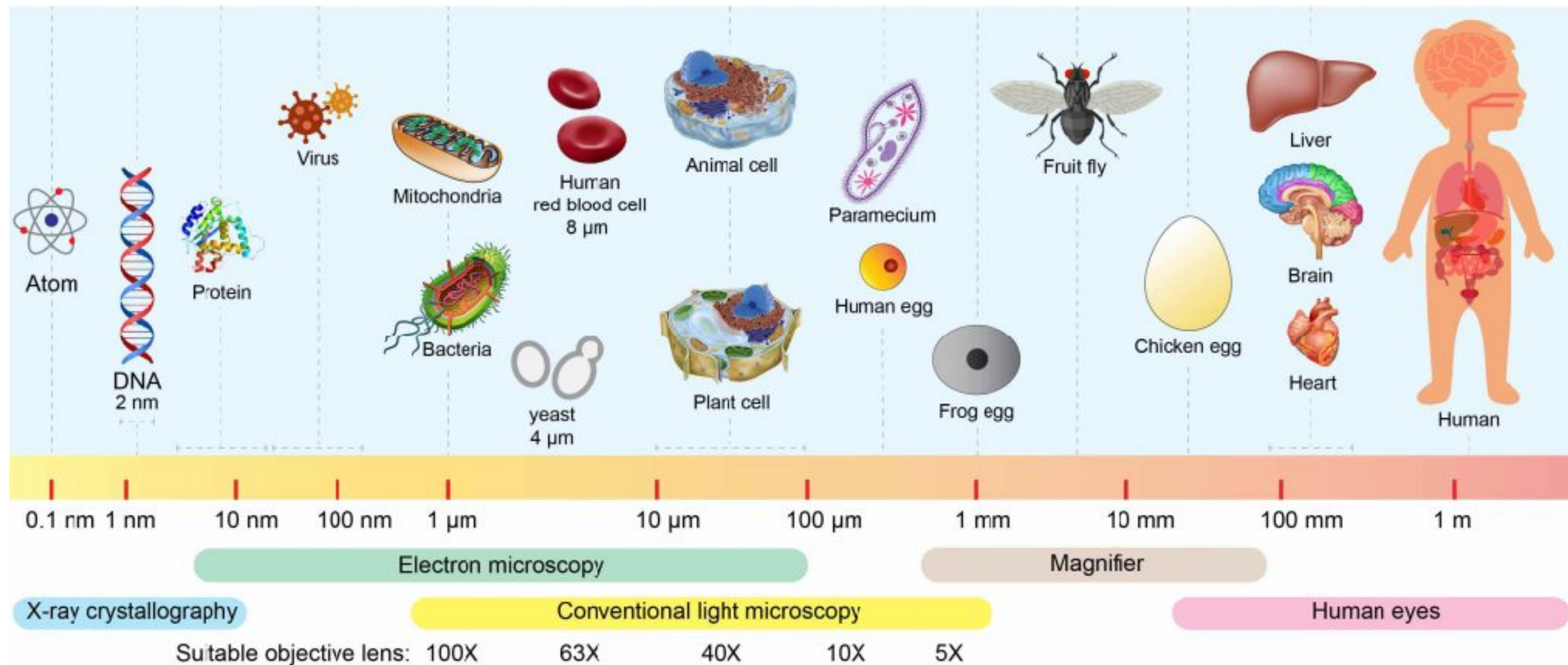
**Viruses**

USB drive

Can't do anything on its own, depends on a computer (see: Humans) to do anything

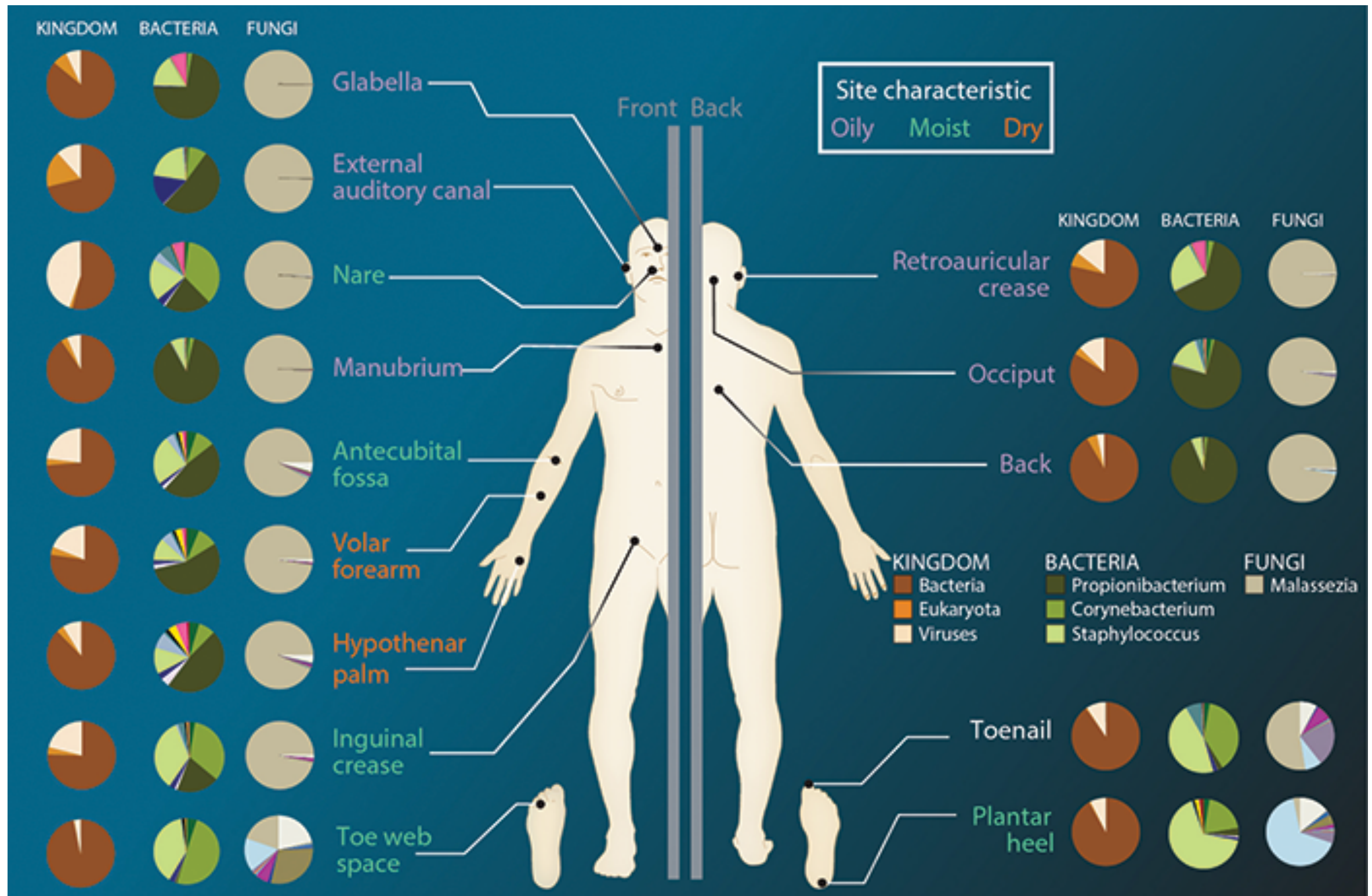
\* fungal microbes, there are also multicellular fungi

# Microbes: Scale





# Microbiomes: Where



# Microbiomes: Where

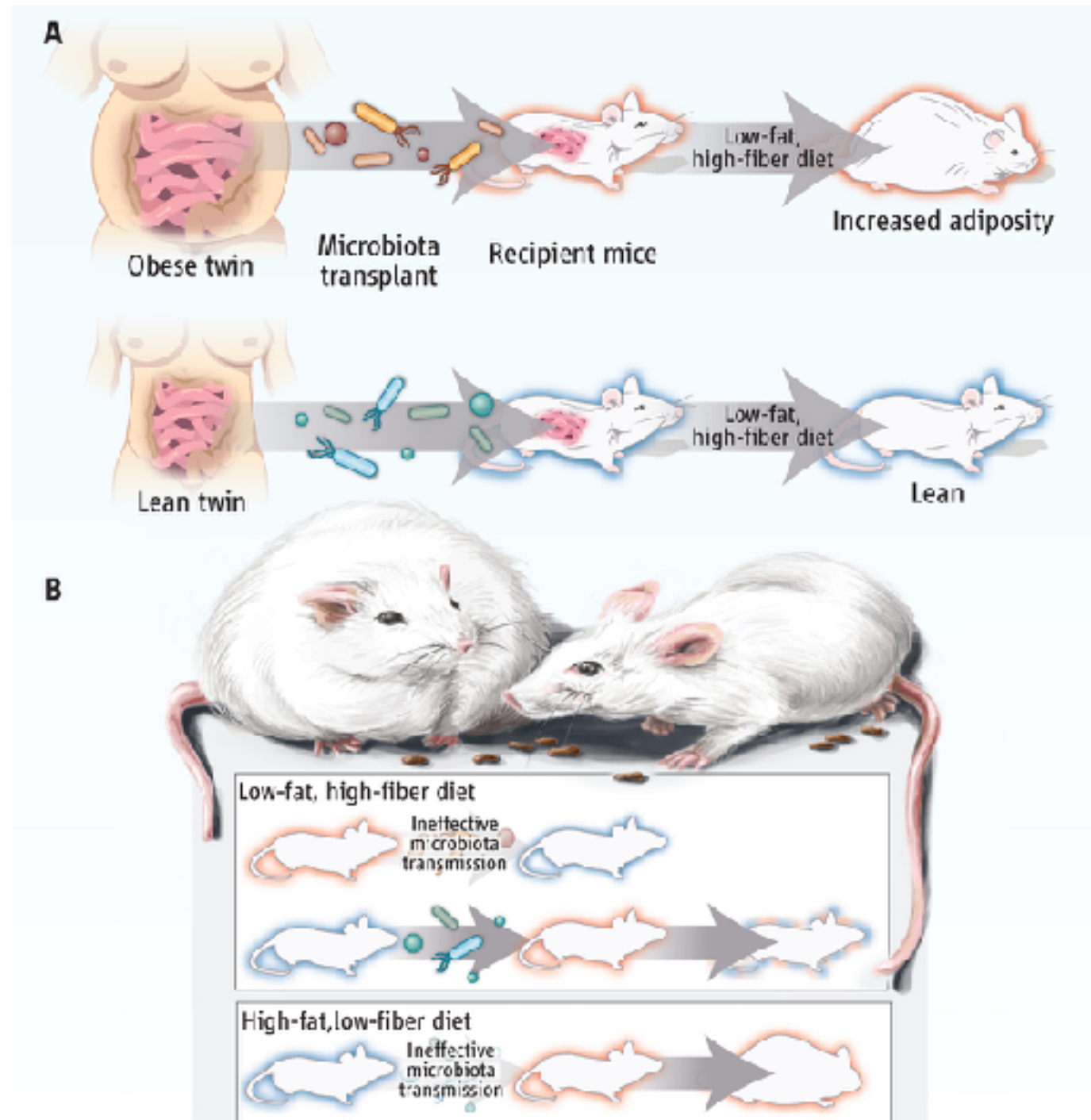


# Microbiomes in Human Health

- 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



Walker, et al. Science 2013. <https://doi.org/10.1126/science.1243787>

Ridaura I, et al. Science 2013. <https://doi.org/10.1126/science.1241214>

# Metagenomics

	What	Information	Analogy	Target Size	Cost
<b>Amplicon</b>	Marker Gene	Who is Present	Name	100bp - 1kb	Low
<b>Shotgun Metagenome</b>	Genomes	What Genes are Present	CV	100kb - 100Mb	High
<b>Shotgun Metatranscriptome</b>	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](#)

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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](#)

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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.

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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](#)

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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](#)

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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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# Amplicon Sequencing

PCR amplify and sequence a marker gene

	Marker Gene
<b>Bacteria</b>	16s rRNA
<b>Fungi</b>	18s or ITS rRNA
<b>Archaea</b>	16s rRNA
<b>Protozoa</b>	18s rRNA
<b>Viruses</b>	?????

# Metagenomics

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



# Metagenomics

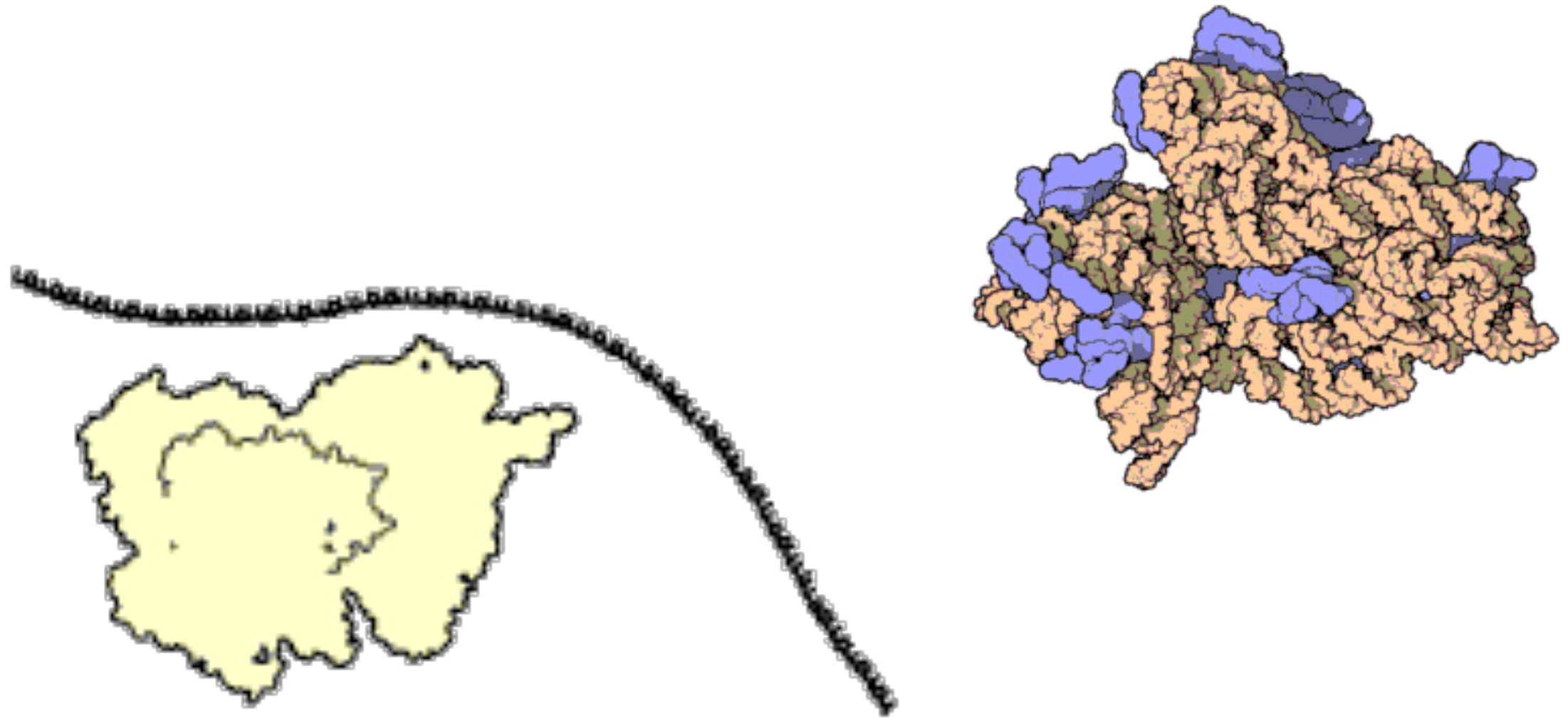
	What	Information	Analogy	Target Size	Cost	Discovery?
<b>Amplicon</b>	Marker Gene	Who is Present	Name	100bp - 1kb	Low	+/-
<b>Shotgun Metagenome</b>	Genomes	What Genes are Present	CV	100kb - 100Mb	High	++
<b>Shotgun Metatranscriptome</b>	All RNA	What Genes are Expressed	Twitter Feed	100kb - 100Mb	High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

# A Whale of an Analogy

- Amplicon: 250 characters (“These reflections just here are occasioned by the circumstance that after we were all seated at the table, and I was preparing to hear some good stories about whaling; to my no small surprise, nearly every man maintained a profound silence. And not o”)
- Whole book:  $1.2 \times 10^6$  characters

# The 16S Amplicon

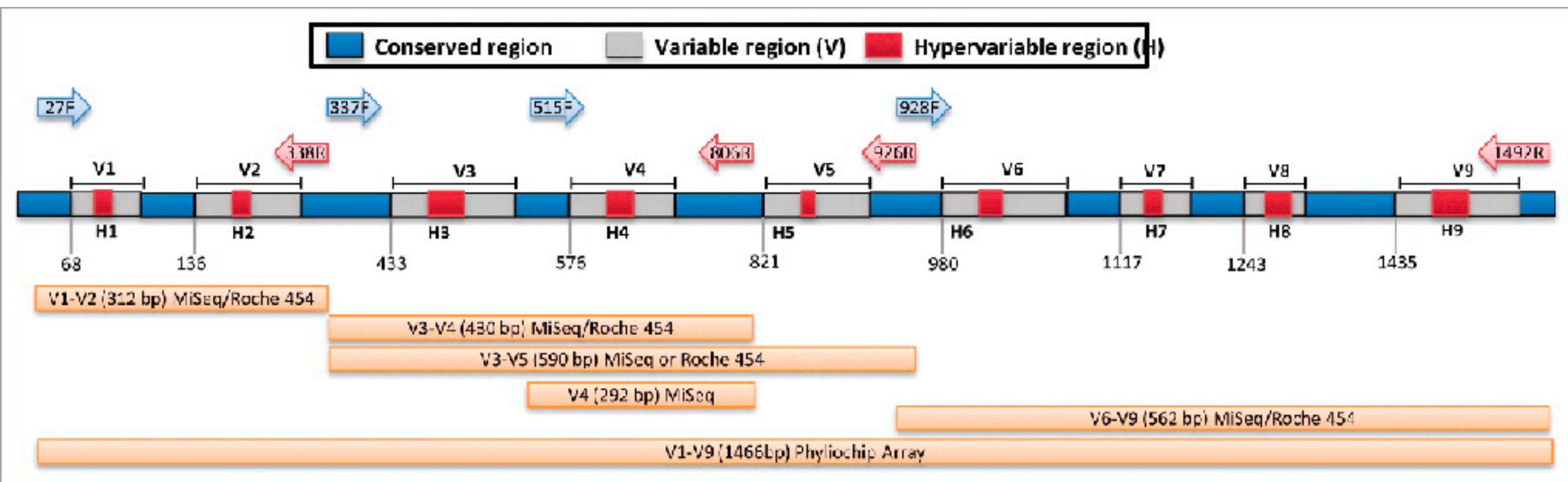
# The Ribosome



[https://upload.wikimedia.org/wikipedia/commons/9/94/Protein\\_translation.gif](https://upload.wikimedia.org/wikipedia/commons/9/94/Protein_translation.gif)

[https://upload.wikimedia.org/wikipedia/commons/9/90/010\\_small\\_subunit-1FKA.gif](https://upload.wikimedia.org/wikipedia/commons/9/90/010_small_subunit-1FKA.gif)

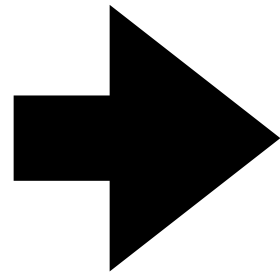
# 16S rRNA Gene





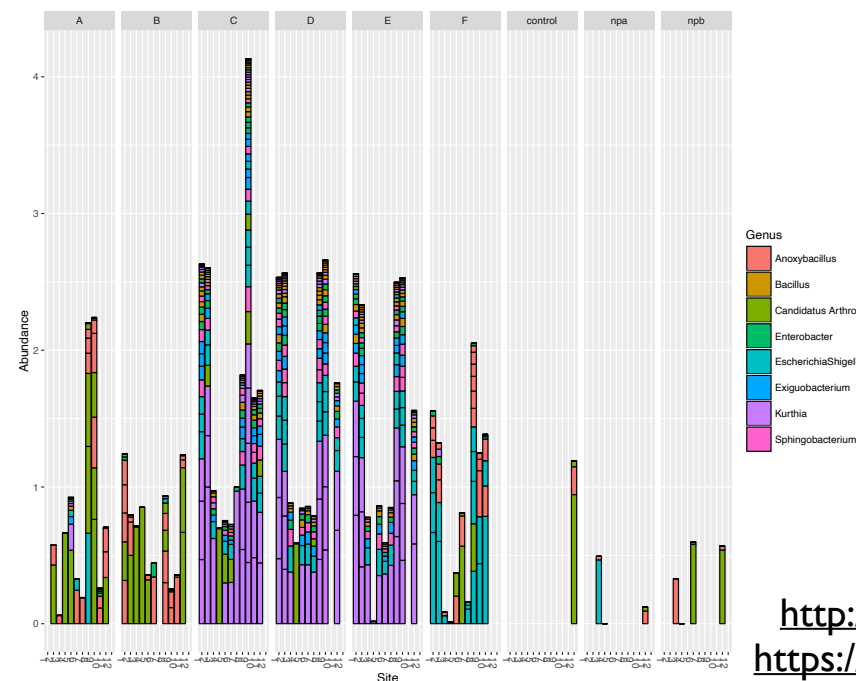
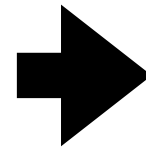
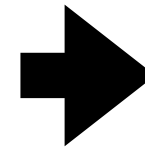
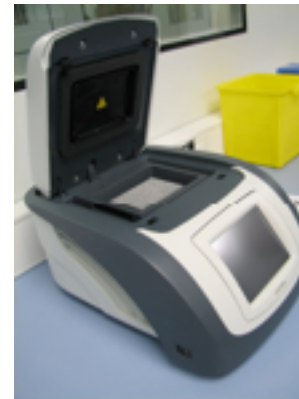
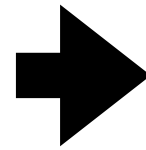
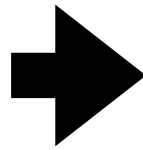
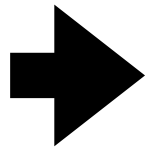
# Amplicon Analysis

# Big Picture



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

# Molecular Biology

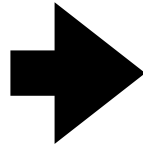


<http://www.geograph.org.uk/photo/2847164>

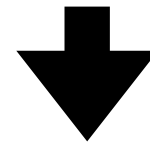
<https://commons.wikimedia.org/wiki/File:Pcr.jpg>

[https://commons.wikimedia.org/wiki/File:Illumina\\_MiSeq\\_sequencer.jpg](https://commons.wikimedia.org/wiki/File:Illumina_MiSeq_sequencer.jpg)

# Bioinformatic Analysis



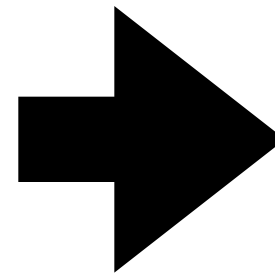
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+
ABBBABBBBAFFGGGGGGGGGGHGGHGGGCG2GF3FFGHHHHHHGGFGHEHHGGGEHHHHAGGHHGHHHFFDHFHHHGGGGG@F@H?GHH/GBEFGGG
@M00698:36:000000000-AFBEL:1:1101:16483:1412 1:N:0:0
CTGCCAGTTGAACGACGGCGAGCAGTTATAAGCCAGCAGTTTGCCCGGATATTTCCGCTGGATAGCTTGTCAAAGCGACGCGCCAGTTCCAGATCCGGCG
+
AAABBFBBBBFFGGGGGGGGGGHHHHHHHHHGHGHHHHHHHHGGHHGGGGGGHHHHGGGGGGHHHHHFFHHHHHHHGGGGGGGGHHHHHHHHHHHHGGG
@M00698:36:000000000-AFBEL:1:1101:15928:1413 1:N:0:0
GTAAAGTCCTGAGTGATACCGGCACTTTTACCCCCAGTCCCACTTTTGAACCGGCAACATATCGGCAAAAGAGCCGTGCCTGATTTAAAGCCGTAGGT
+
```



	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				



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



# Caveat



**The End**