

# Microbiome Overview

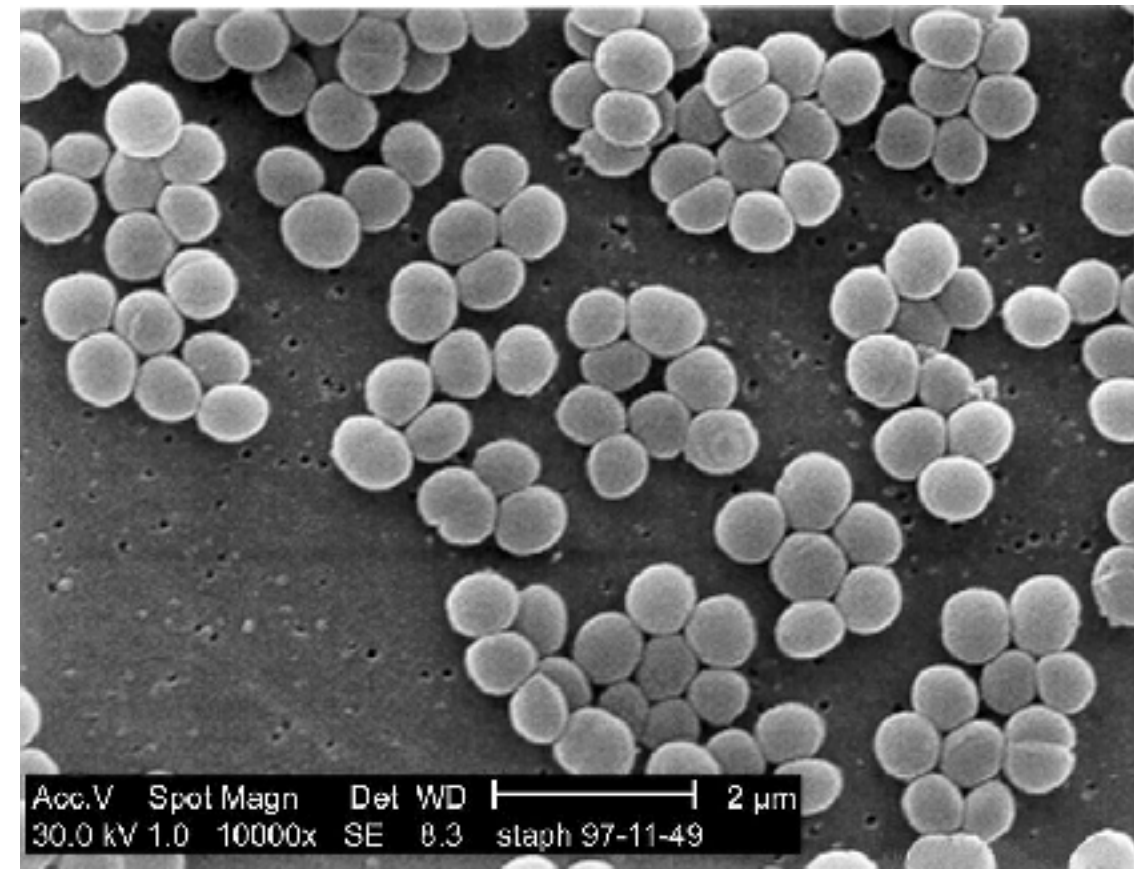
Josh Granek

# 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

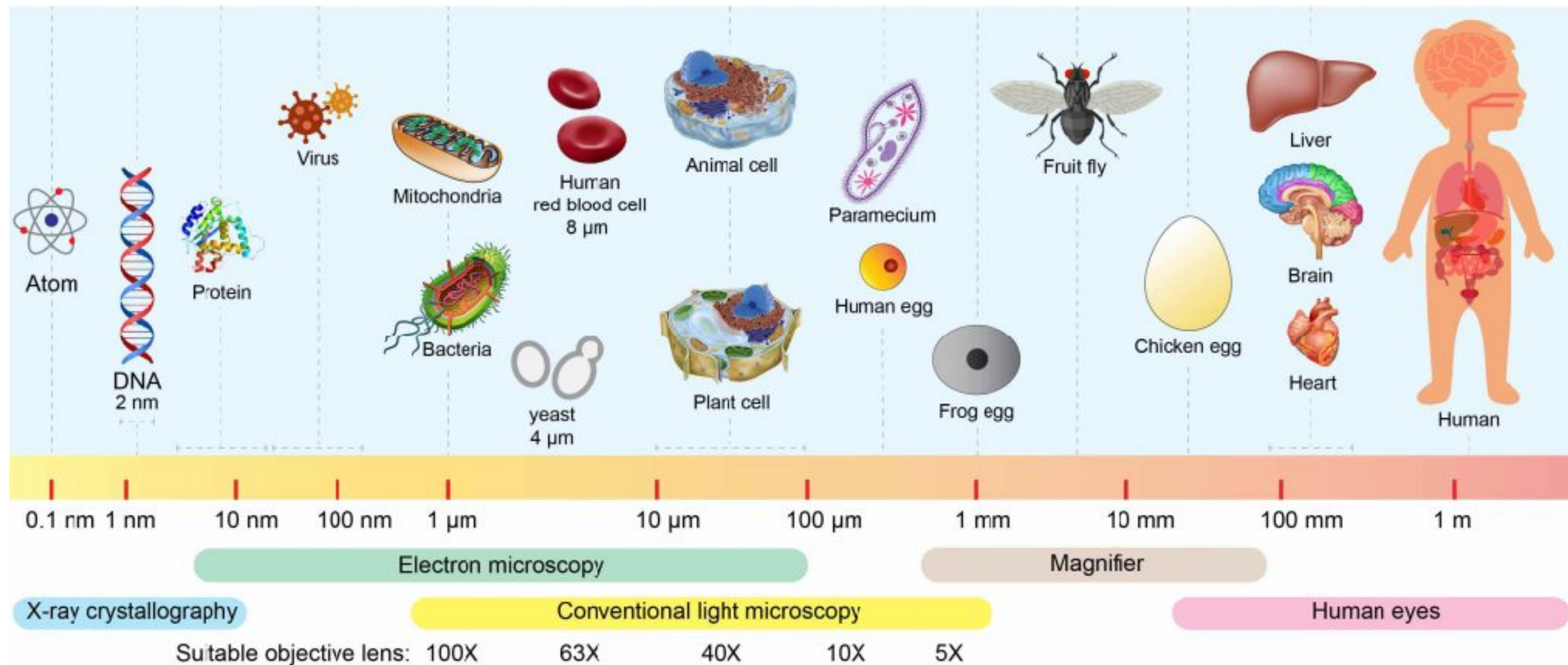
**Viruses**

USB drive

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

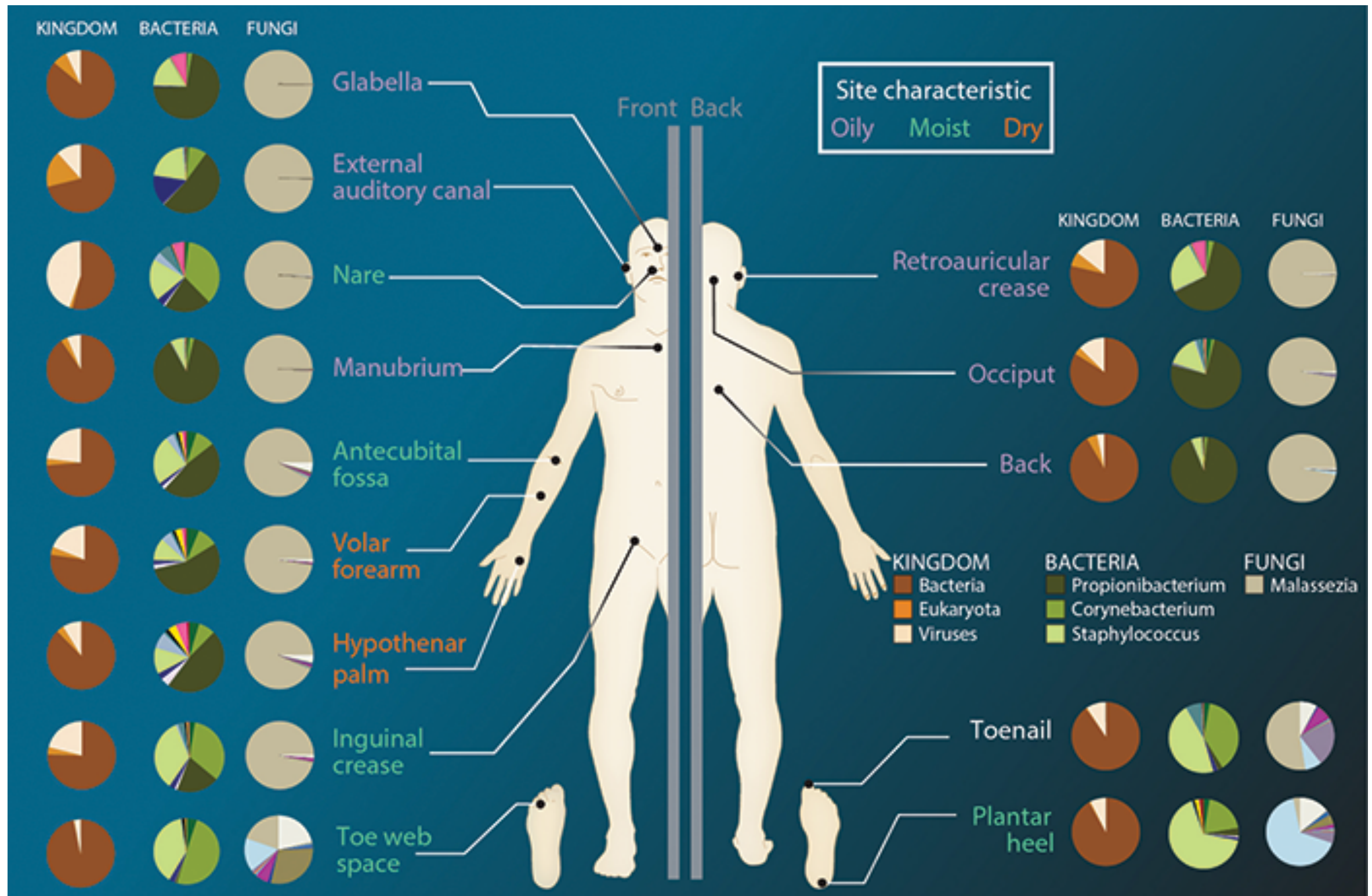
\* fungal microbes, there are also multicellular fungi

# Microbes: Scale

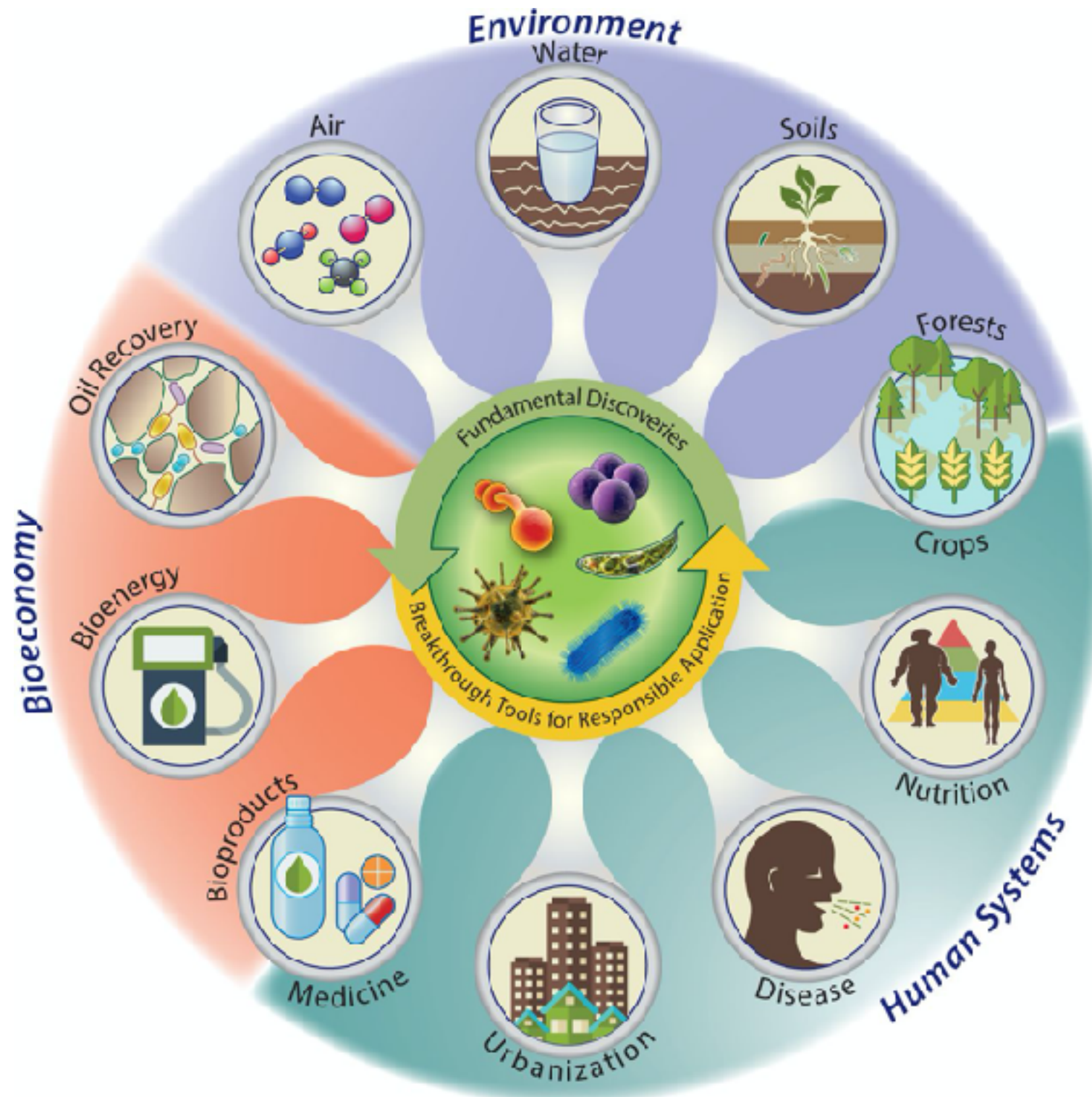




# Microbiomes: Where



# Microbiomes: Where

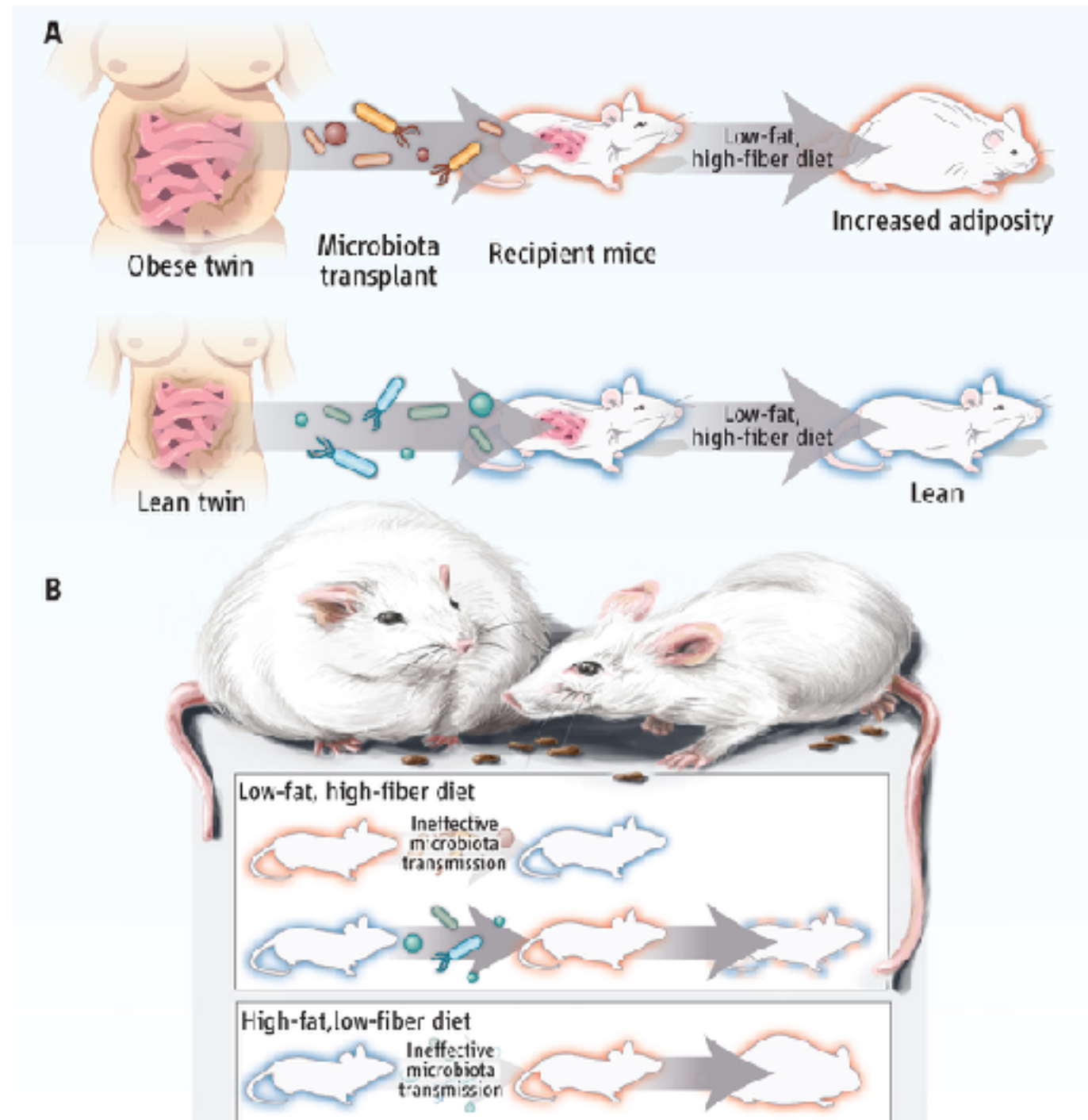


# Microbiomes in Human Health

- 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



# Metagenomics

# Metagenomics



**Amplicon**

**Shotgun  
Metagenome**

**Shotgun  
Metatranscriptome**

# Metagenomics

	What
<b>Amplicon</b>	Marker Gene
<b>Shotgun Metagenome</b>	Genomes
<b>Shotgun Metatranscriptome</b>	All RNA

# Metagenomics

	What	Information
<b>Amplicon</b>	Marker Gene	Who is Present
<b>Shotgun Metagenome</b>	Genomes	What Genes are Present
<b>Shotgun Metatranscriptome</b>	All RNA	What Genes are Expressed



# Metagenomics

	What	Information	Analogy
<b>Amplicon</b>	Marker Gene	Who is Present	Name
<b>Shotgun Metagenome</b>	Genomes	What Genes are Present	CV
<b>Shotgun Metatranscriptome</b>	All RNA	What Genes are Expressed	Twitter Feed

# Metagenomics

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

# 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
<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

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# Amplicon Sequencing

PCR amplify and sequence a marker gene

# 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



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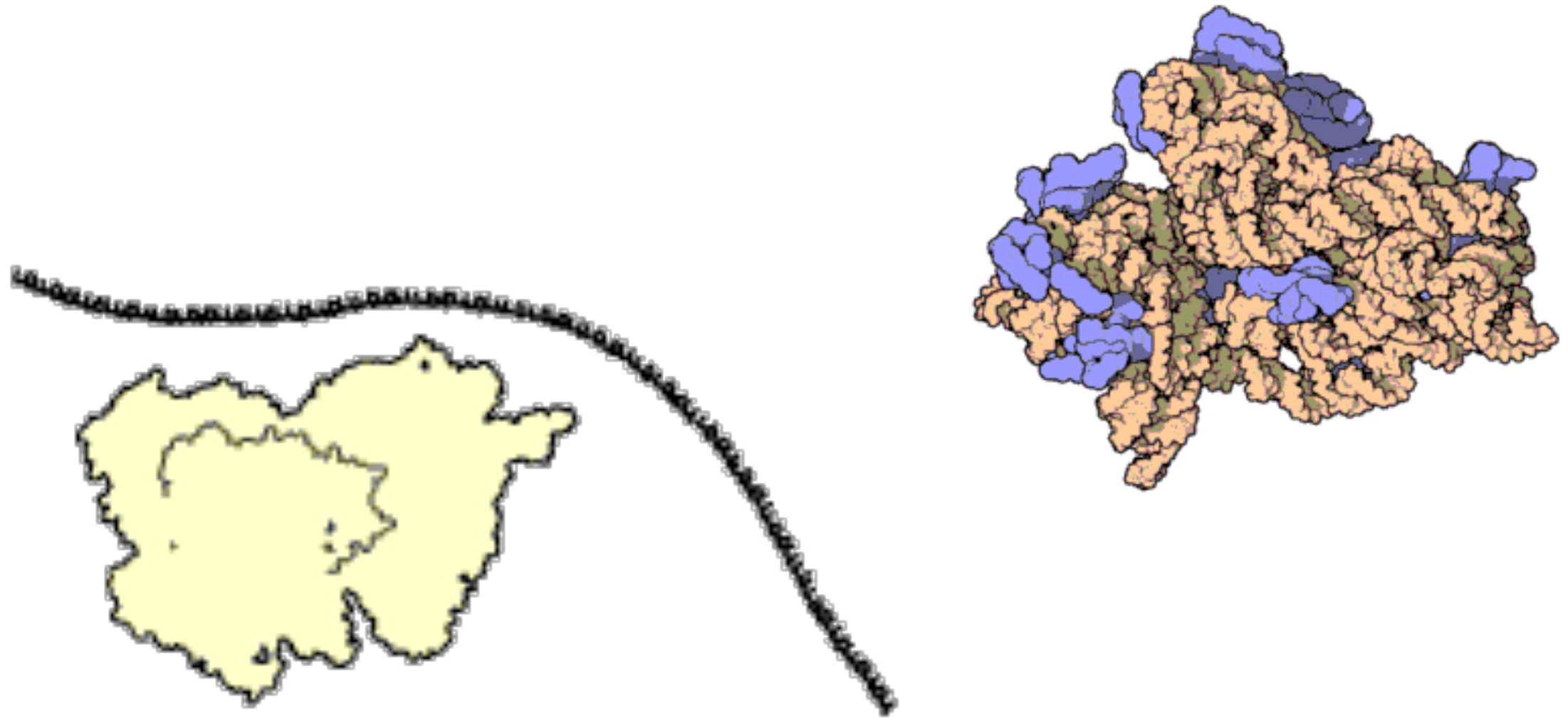
	What	Information	Analogy	Target Size	Cost	Discovery?
<b>Amplicon</b>	Marker Gene	Who is Present	Name	100bp - 1kb	Low	+/-
<b>Shotgun Metagenome</b>	Genomes	What Genes are Present	CV	100kb - 100Mb	High	++
<b>Shotgun Metatranscriptome</b>	All RNA	What Genes are Expressed	Twitter Feed	100kb - 100Mb	High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

# 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

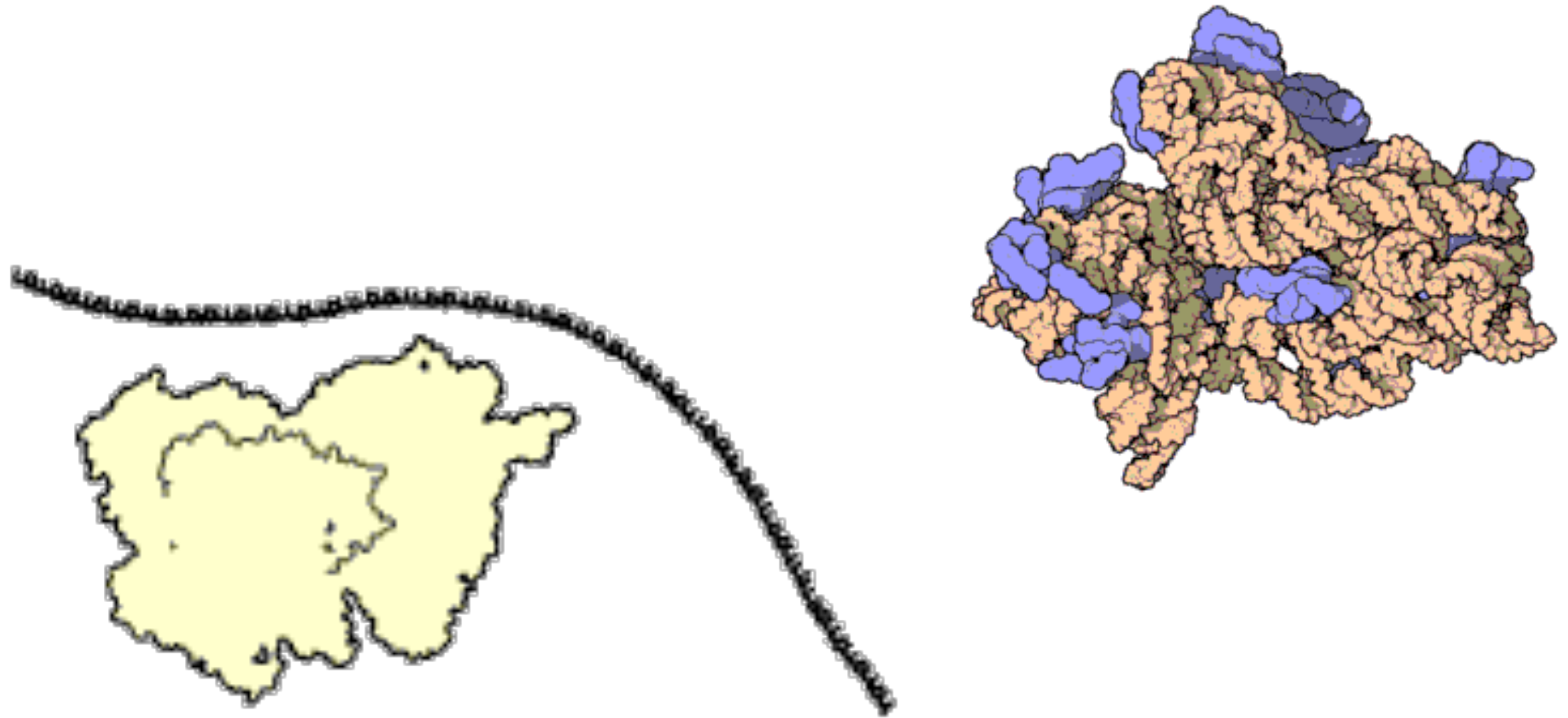
# The Ribosome



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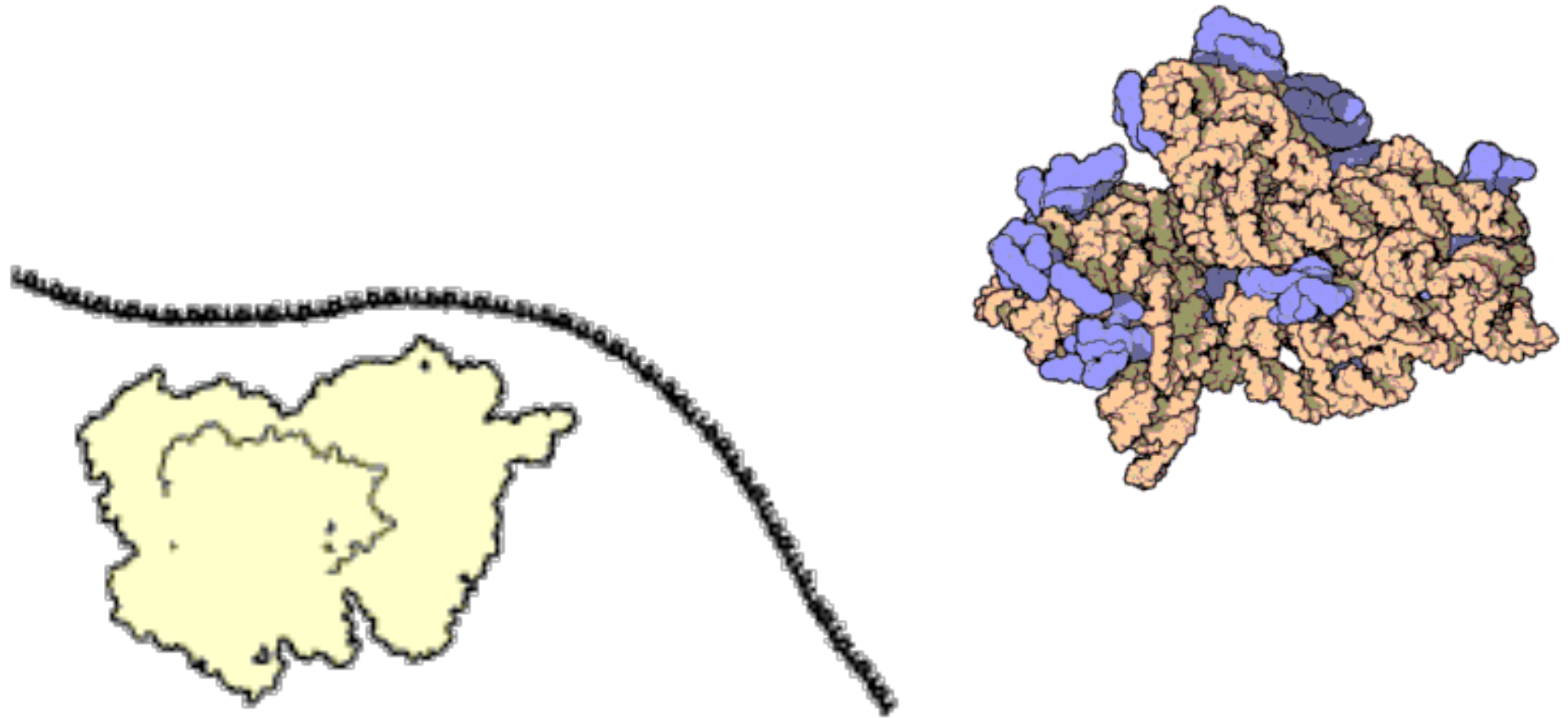


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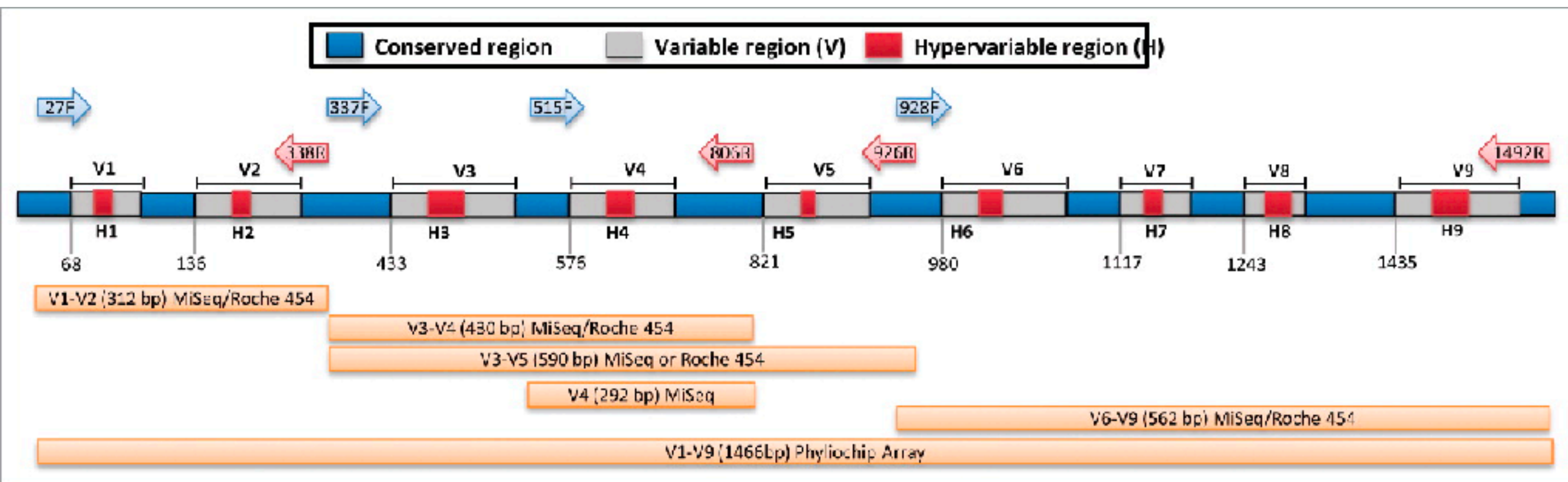
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# 16S rRNA Gene



# Sequencing Details

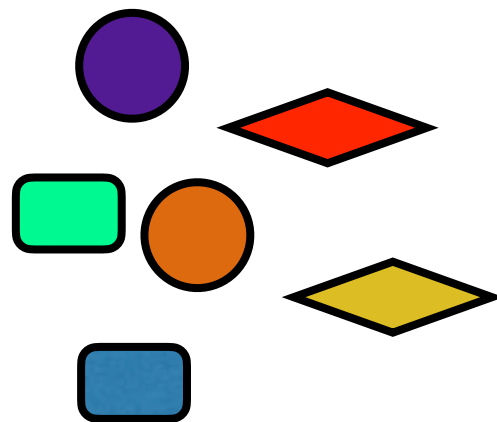
# Amplicon Sequencing

Primers

P1   
P2 

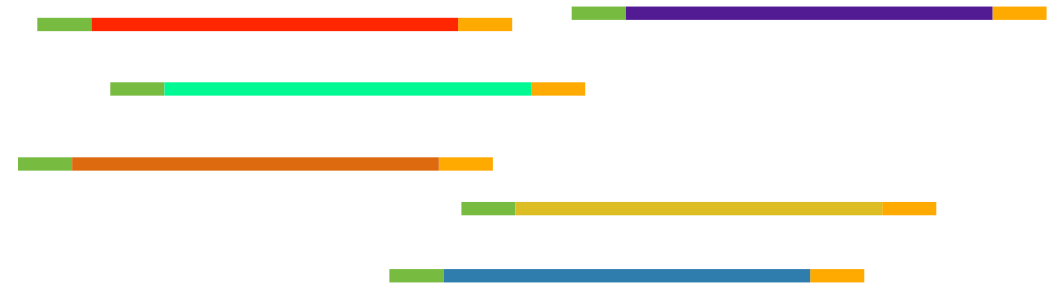
+

Microbes



=

Amplicon Library

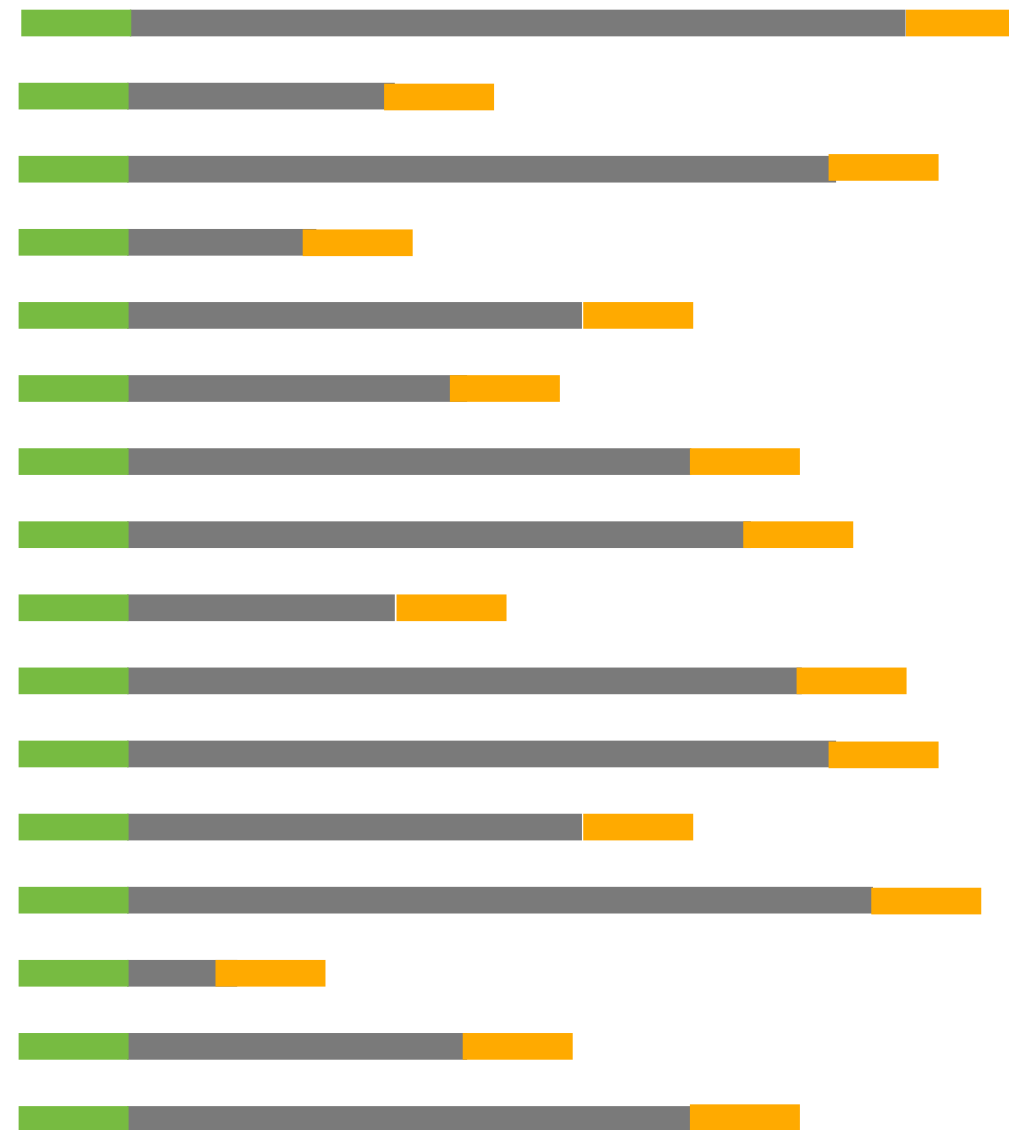


# Sequencing Library

## Amplicon Library



## Shotgun Library



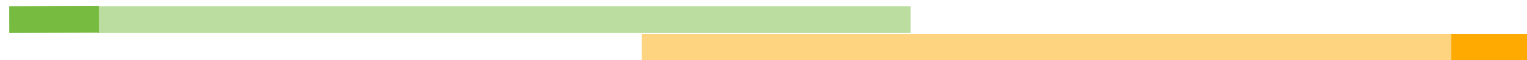
# Read Length

Read  
Length

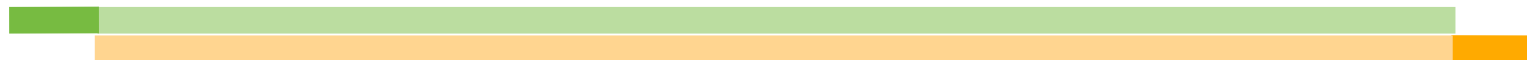
100bp



150bp



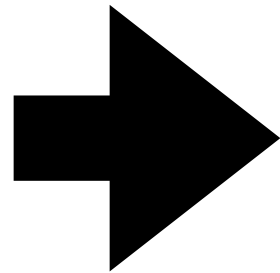
250bp



250bp Amplicon

# Amplicon Analysis

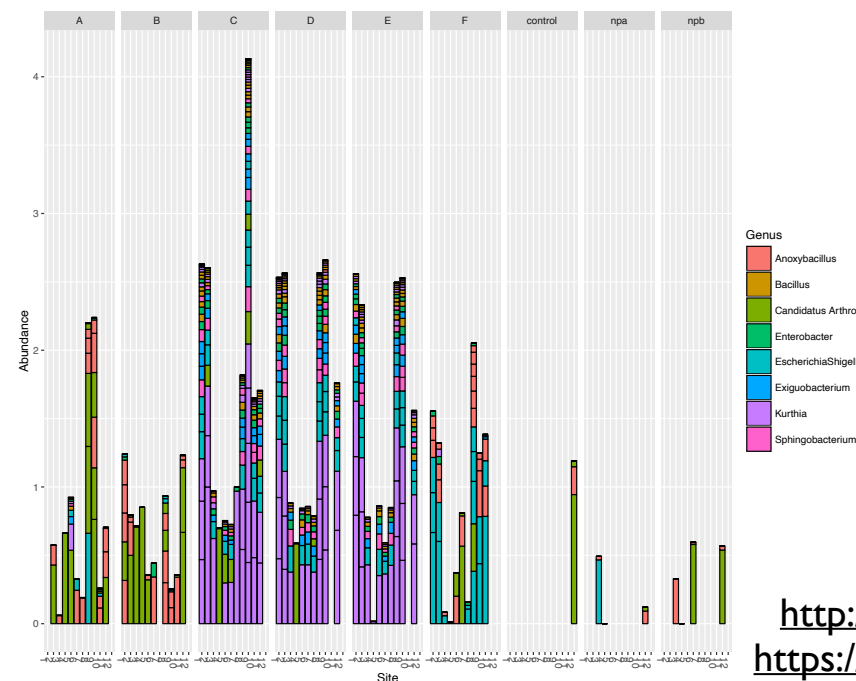
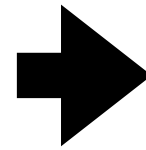
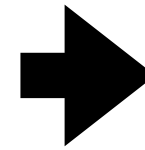
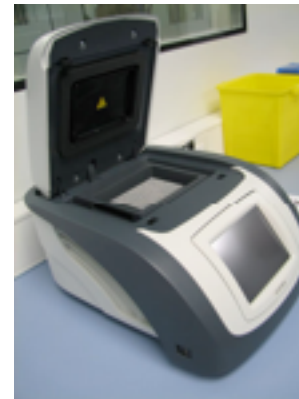
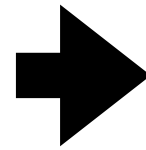
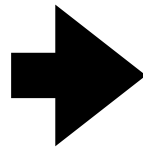
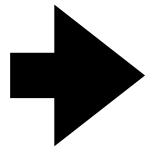
# Big Picture



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



# Molecular Biology

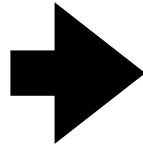


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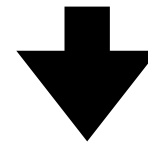
<https://commons.wikimedia.org/wiki/File:Pcr.jpg>

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

# Bioinformatic Analysis



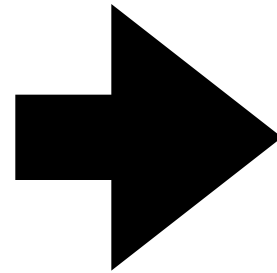
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+
ABBBABBBBAFFGGGGGGGGGGHGGHGGGCG2GF3FFGHHHHHHGGFGHEHHGGGEHHHHAGGHHGHHHFFDHFHHHGEgggg@F@H?GHH/GBEFGGG
@M00698:36:000000000-AFBEL:1:1101:16483:1412 1:N:0:0
CTGCCAGTTGAACGACGGCGAGCAGTTATAAGCCAGCAGTTTGCCCGGATATTTCCGCTGGATAGCTTGTCAAAGCGACGCGCCAGTTCCAGATCCGGCG
+
AAABBBFFFFFGGGGGGGGGGGHHHHHHHHHGHGHHHHHHHHGGHHGGGGGGHHHHGGGGGGHHHHHFFHHHHHHGHGGGGGGGGHHHHHHHHHHHHGGG
@M00698:36:000000000-AFBEL:1:1101:15928:1413 1:N:0:0
GTAAAGTCCTGAGTGATACCGGCAACTTTTACCCCCAGTCCCACTTTGCAACCGGCAACATATCGGCAAAAGAGCCGTGCCTGATTTAAAGCCGTAGGT
+
```



	Sample 1	Sample 2	...	Sample N
Bacteria 1	1	10		
Bacteria 2	100	0		
...				
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**