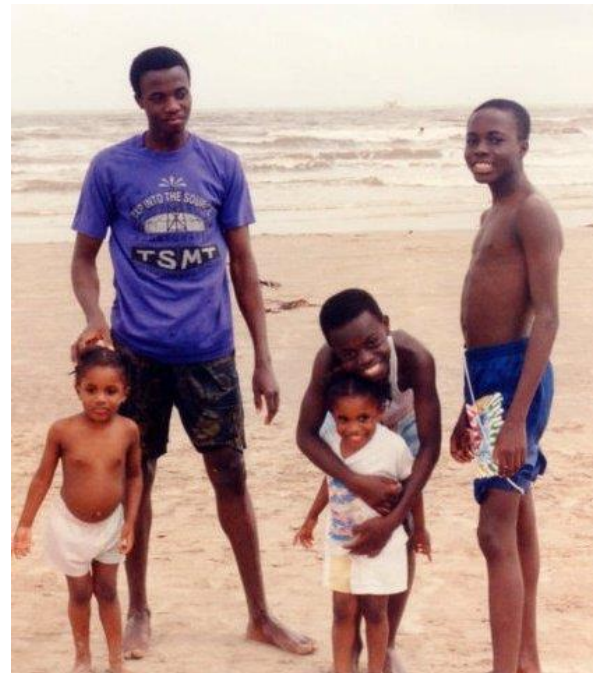
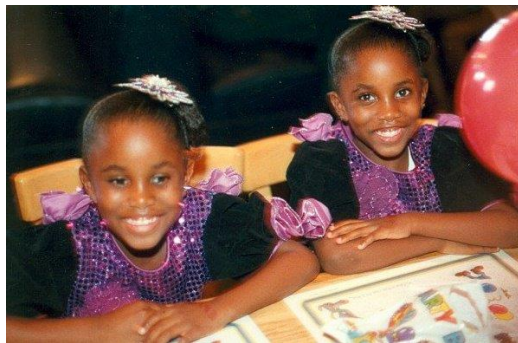
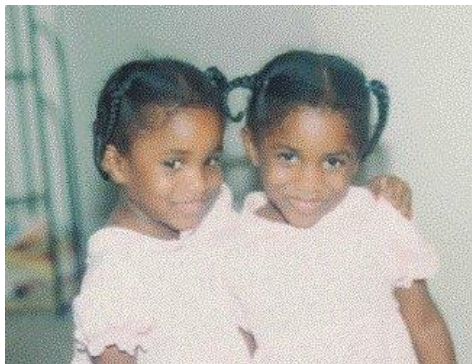


TEEN SCIENCE CAFÉ

Why and how I became a scientist

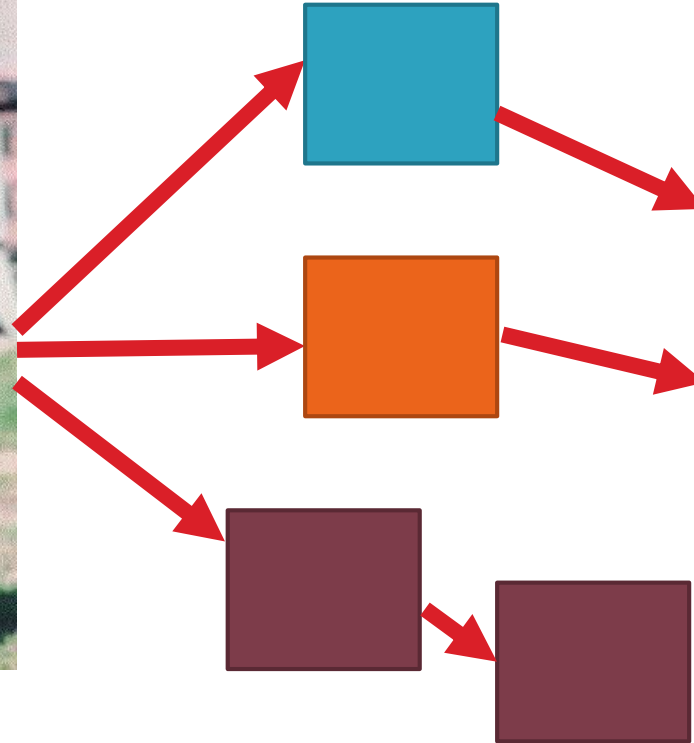
My Beginning.....



My Initial Dreams



My Reality





How was I going to make my family proud?

Dream School



RICE

Unconventional Wisdom



Middle School– High School– College



What should I major in?



RICE

Unconventional Wisdom



Amazing Advisors



Exploring Research: *Dictyostelium discoideum*



3 published papers as
an undergraduate

Exploring Research



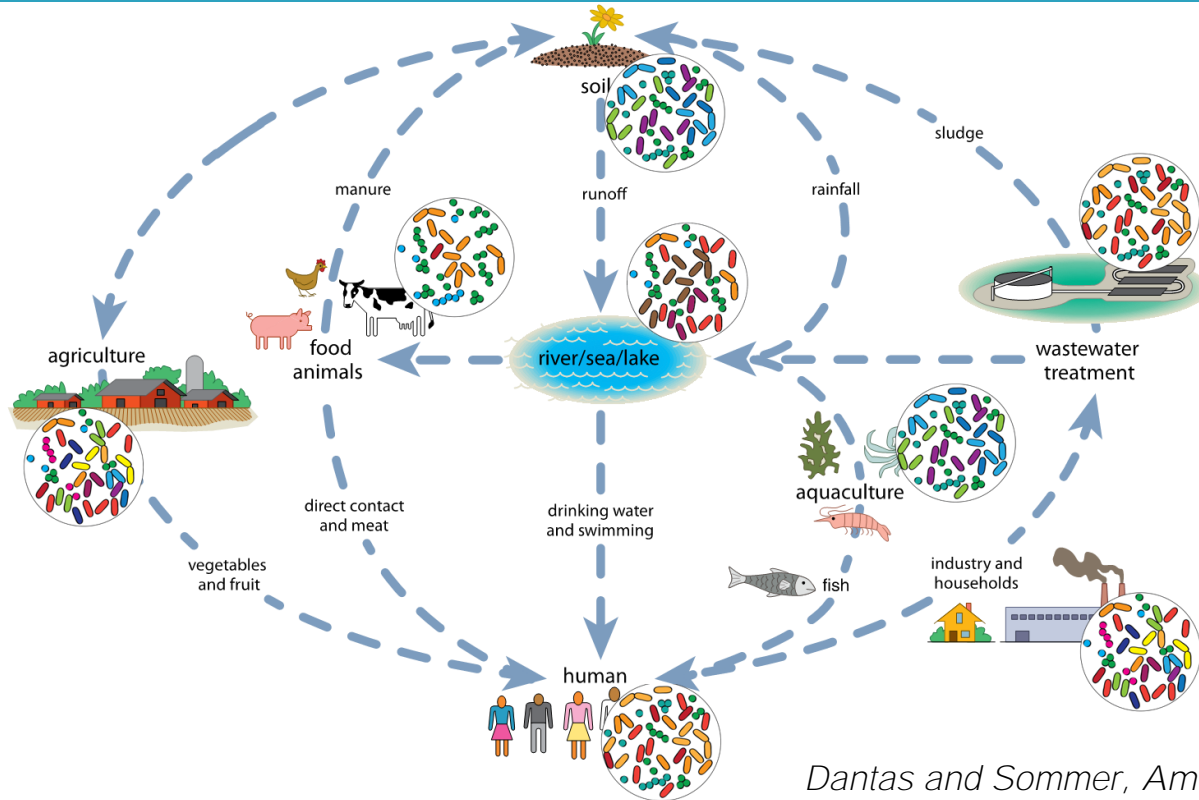
Exploring Research



Microbial communities are highly diverse, abundant and serve important functions in almost every environment



Highly connected network of microbial communities across habitats

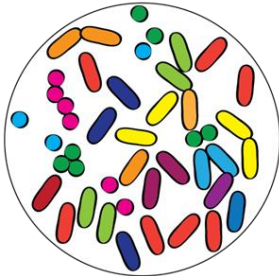


*adapted from:
Dantas and Sommer, American Scientist (2014)*

Microbes are traditionally studied by culture-dependent methods



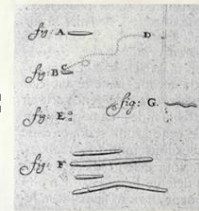
99 - 99.9% of bacteria in most habitats are not easily cultured!



+



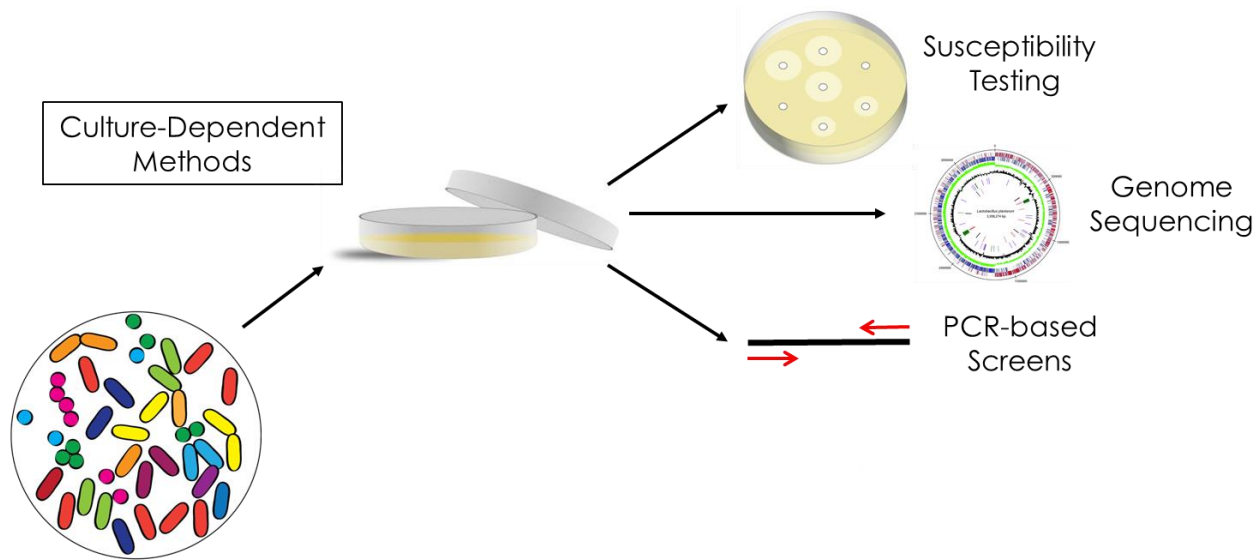
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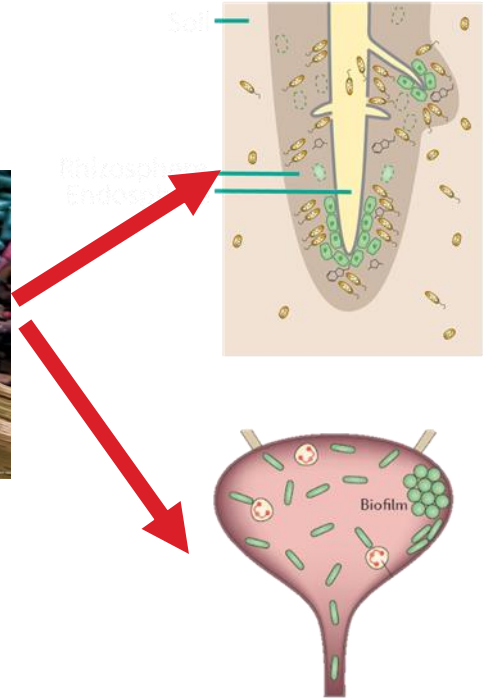
“...animalcules were in such enormous numbers, that all the water...seemed to be alive.” — van Leeuwenhoek (1683)



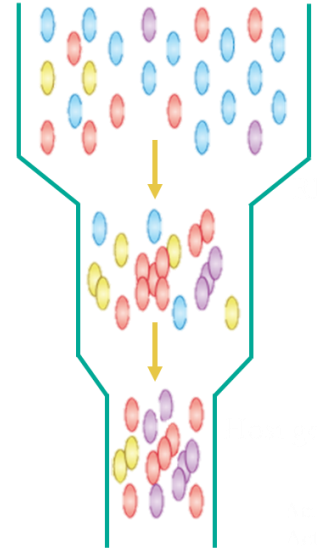
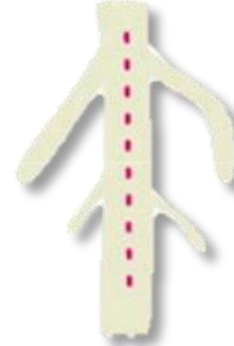
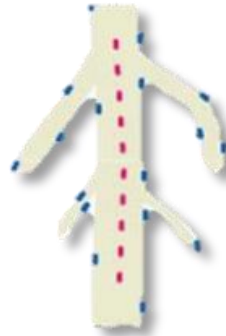
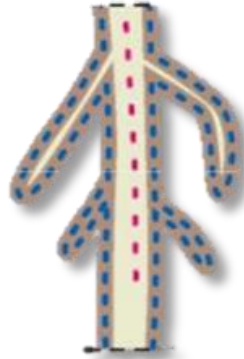
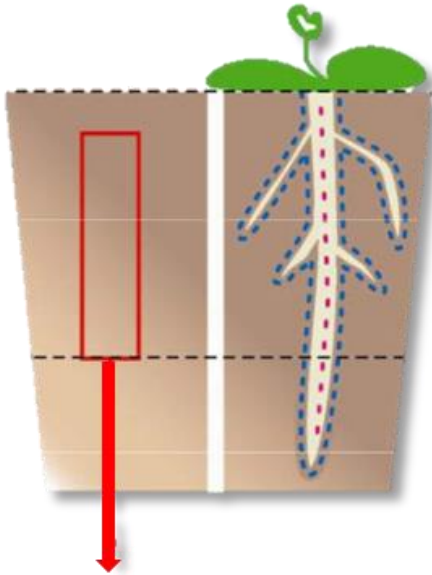
Metagenomics: study microbial communities through direct DNA sequencing



Current Research



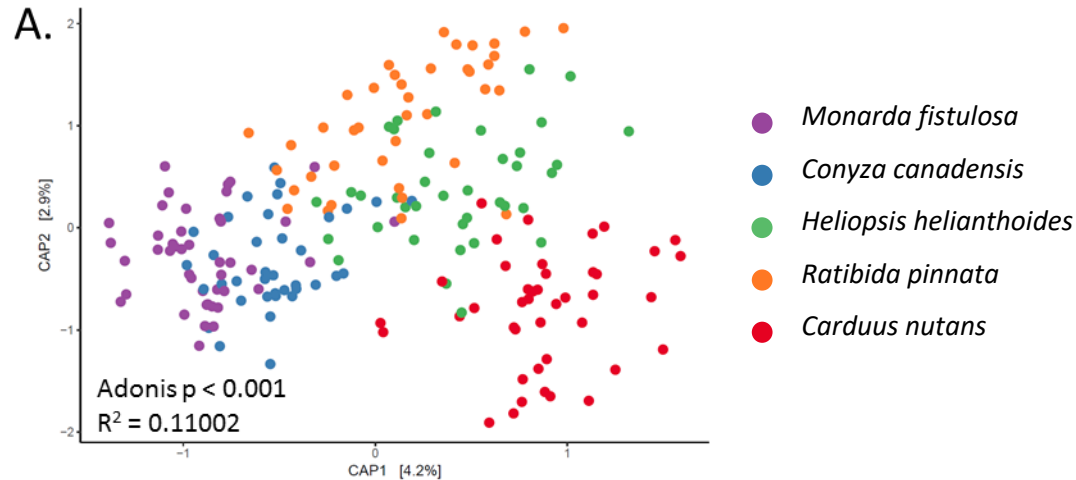
Root Microbiome Study



Root Microbiome Study



Root Microbiome Study



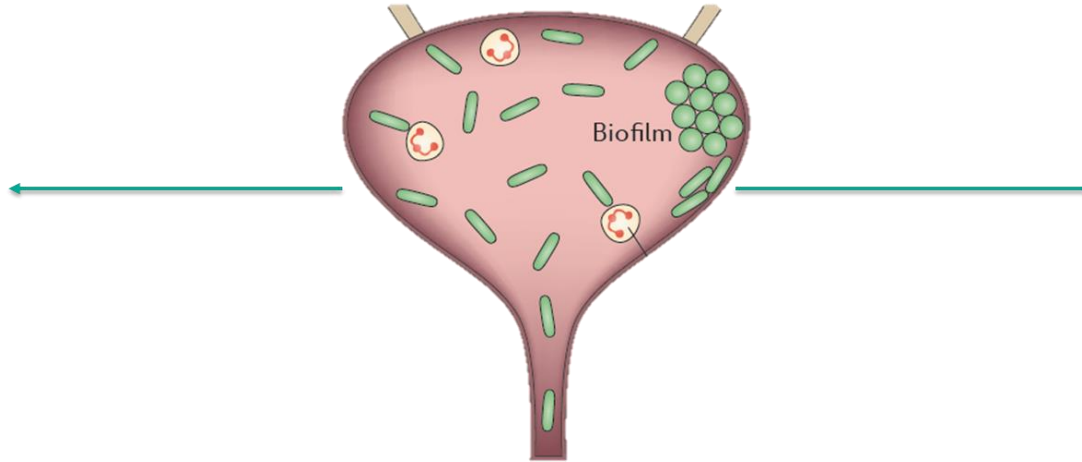
Plants are selecting bacteria from the soil

Urine Microbiome Study

Healthy Urine



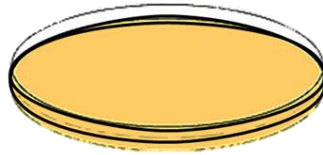
Infected Urine



Urinary Tract Infection = Culture growing a uropathogenic bacteria in the presence of symptoms or signs compatible with a Urinary Tract Infection

Urine Microbiome Study

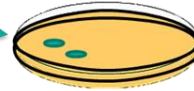
Healthy Urine



Positive Culture



Contaminated



Insignificant

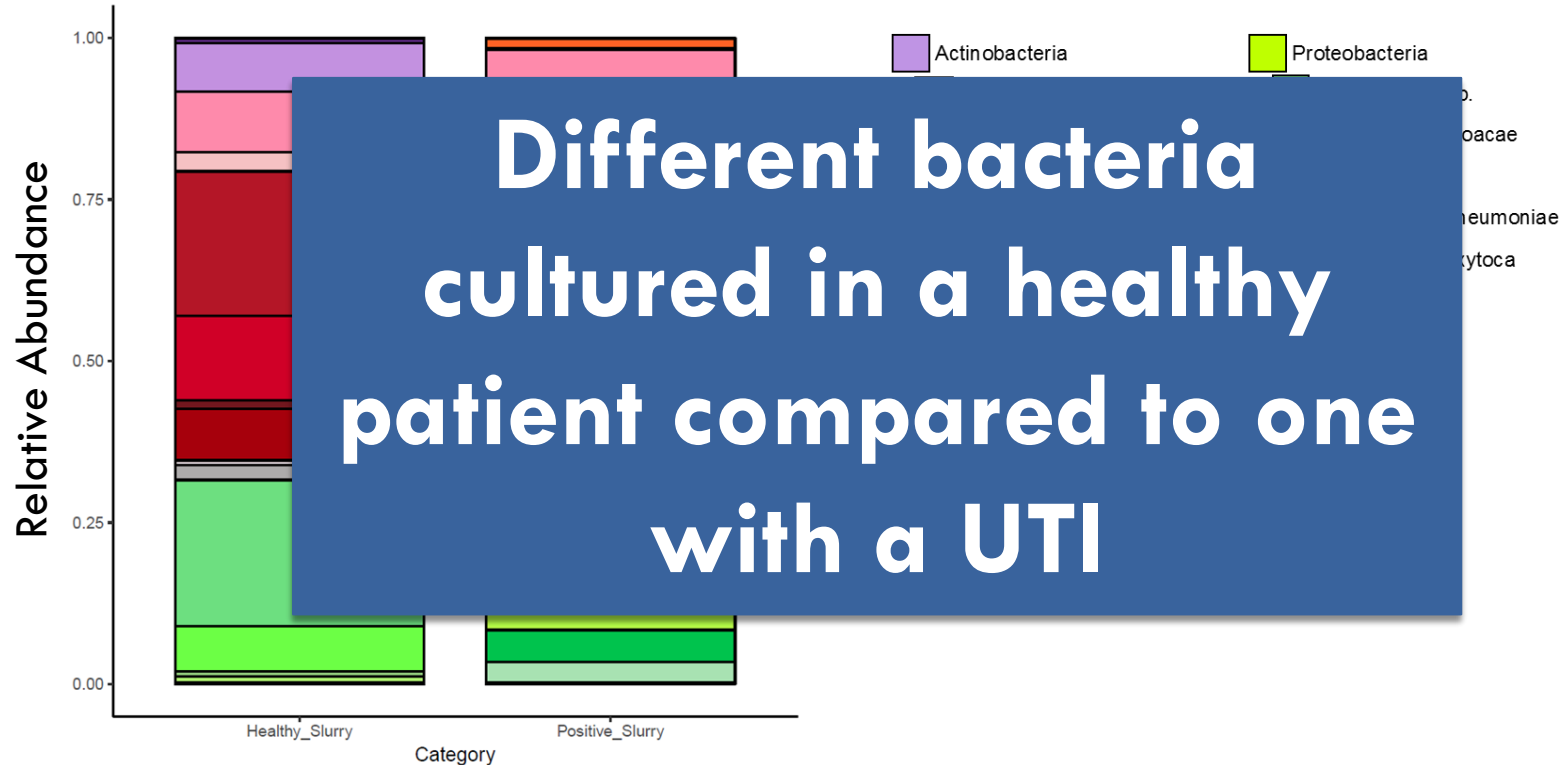


No-Growth

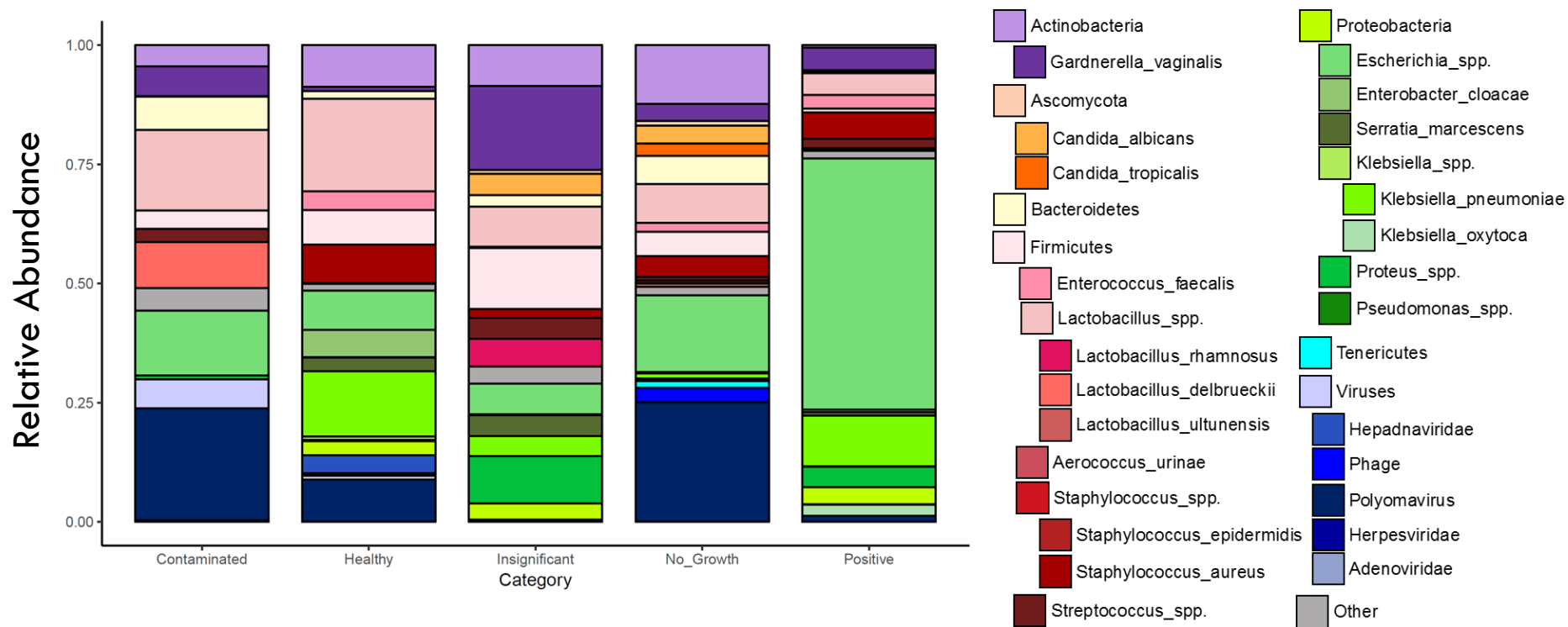
Infected Urine



Cultured Urine Microbiome



Direct Sequencing of the Urine Microbiome



Thank you!! Questions?



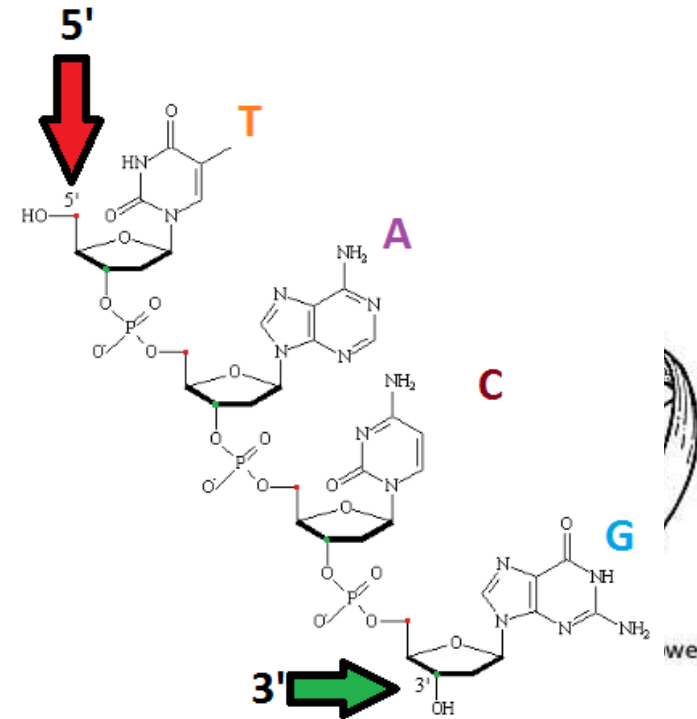


Activity Time!

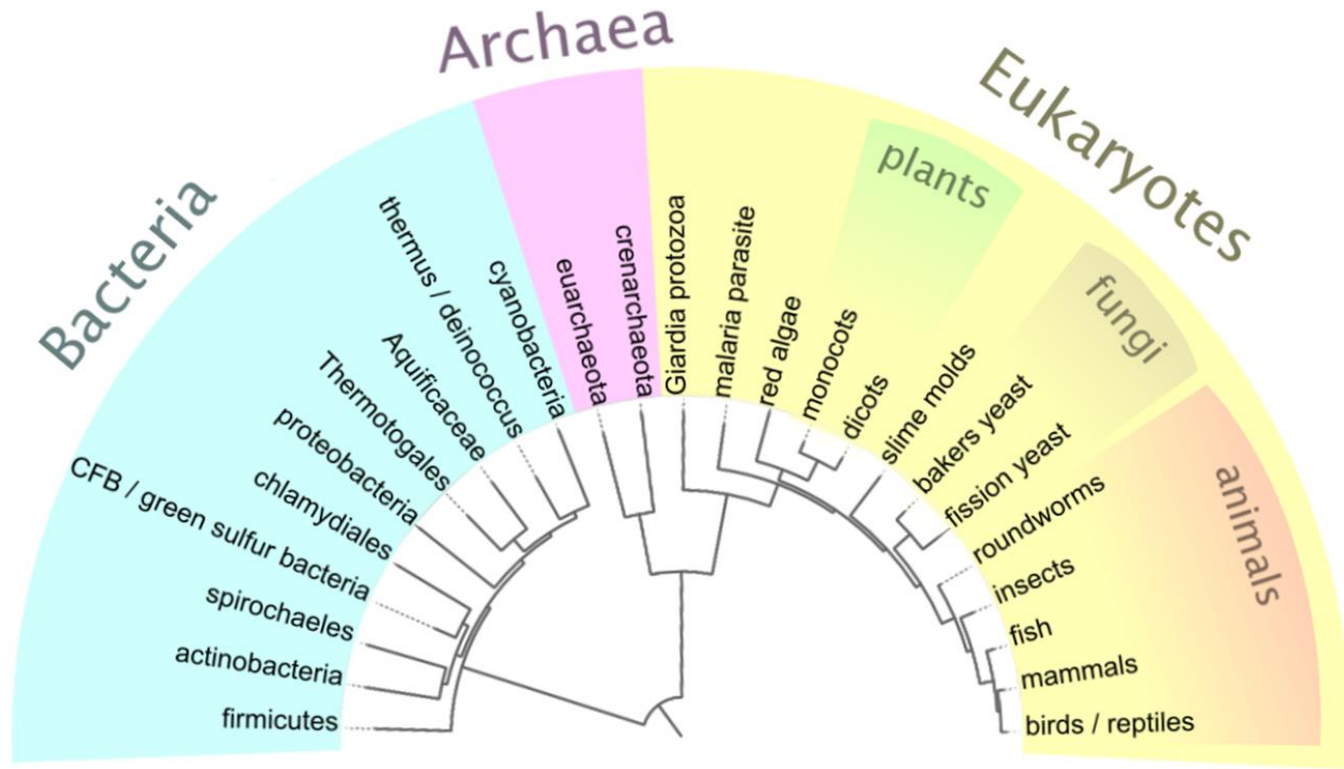
What environment did we sequence?

DNA is the building block of all living organisms

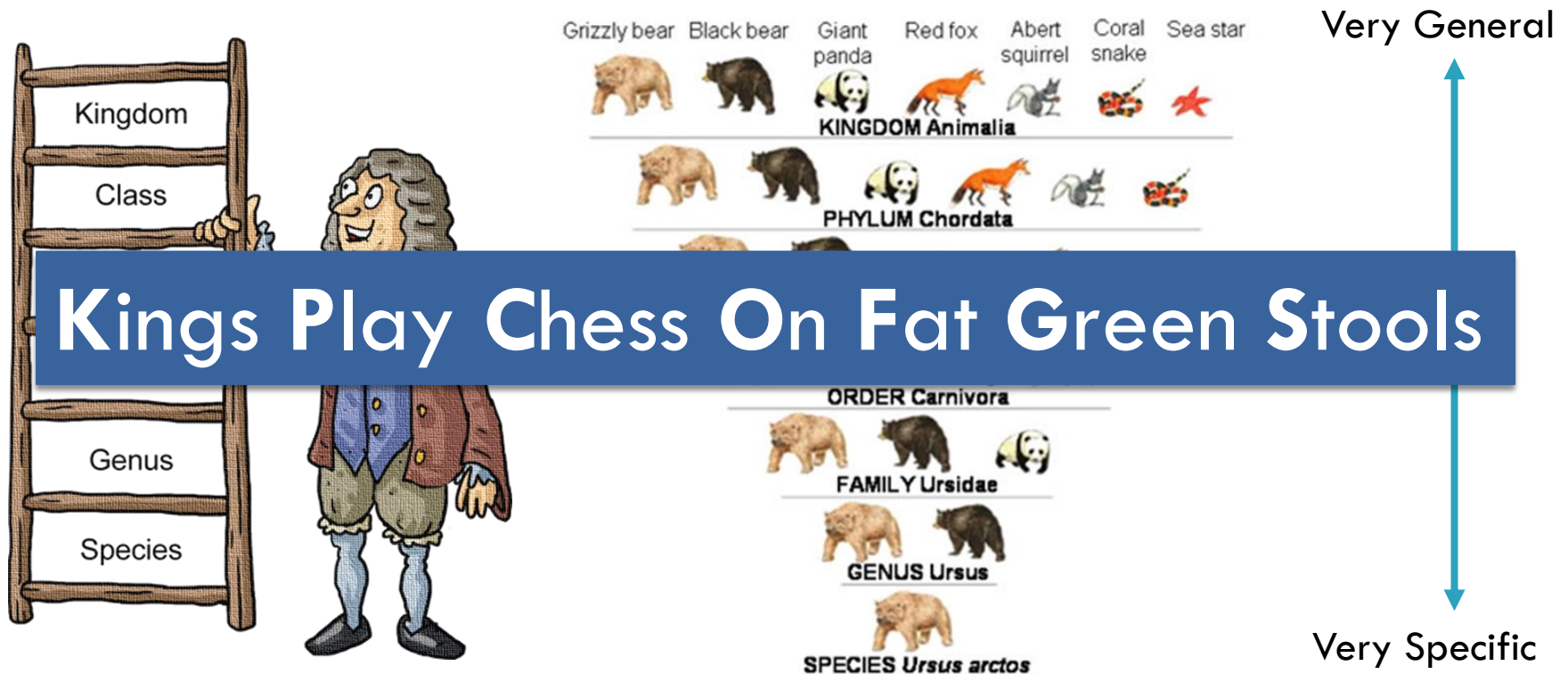
- Adenine (A) - Thymine (T)
- Guanine (G) - Cytosine (C)
- Forward sequence 5' – 3'
- Reverse sequence 3' – 5'
- For example:
 - ▣ 5' – AGGACGT – 3'
 - ▣ 3' – TCCTGCA – 5'
 - ▣ Reverse complement - ACGTCCT



Living things on our planet are classified into three domains

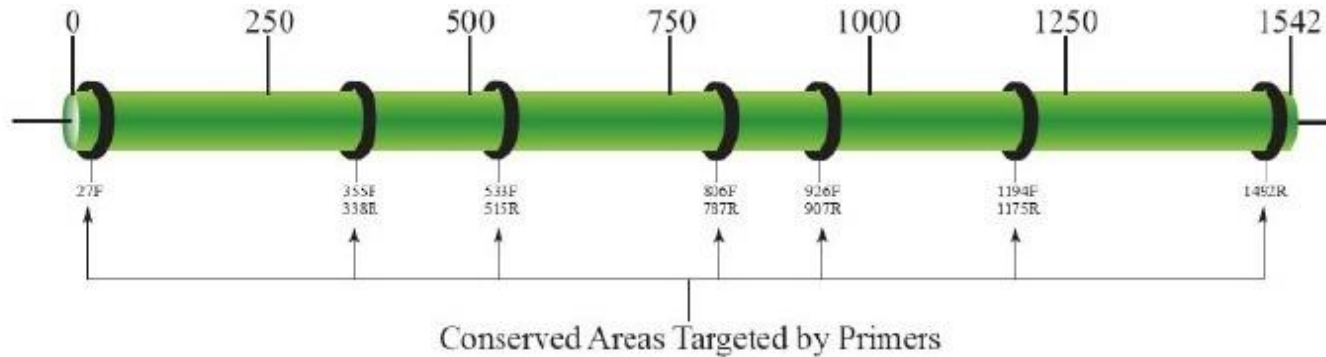


How do we classify living organisms?



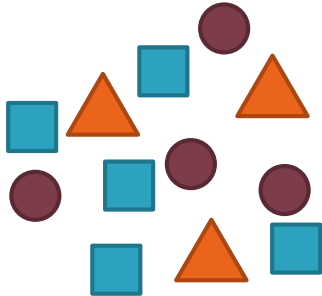
How do we classify bacteria?

16S ribosomal RNA gene



How to calculate relative abundance

Relative abundance is the percent composition of an organism of a particular kind *relative* to the total number of organisms in the area.




How many circles? 4
How many squares? 5
How many triangles? 3
Total number of individuals? 12

Circles RA? 33%
Squares RA? 41.6%
Triangles RA? 25%


$$\text{Relative abundance} = \frac{\text{\# of individuals}}{\text{total number of individuals}} \times 100$$



Wrapping up



Why is it important to study all of the
species in the environment?



Why is it important to study bacteria that
live on this planet?



How will global warming affect
bacteria?



Why is sequencing bacteria better than
culturing?

Why is culturing bacteria better than
sequencing?