



# Welcome to the M<sup>3</sup> Taxonomic Identification Workshop!



Before we begin, please:

1. Check in and let us know if you need a parking voucher
2. Log onto your machine using the username and password on the right hand corner of your computer
3. Log onto the openlab server using the username and password found on the back of your name tag: Go to <https://github.com/shahnidhi/m3-taxonomy-workshop> for detailed directions. It might be helpful to create a file on your Desktop to store name and password

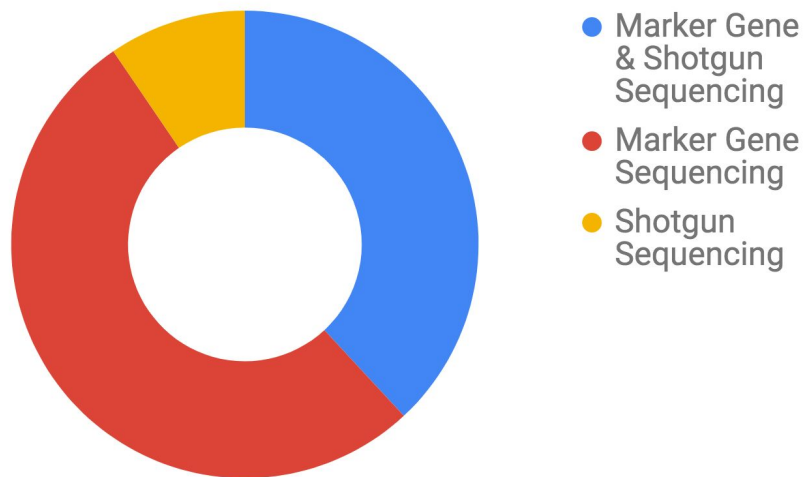
ssh USERNAME@openclass.umiacs.umd.edu

# Icebreaker: Introduce your neighbor!

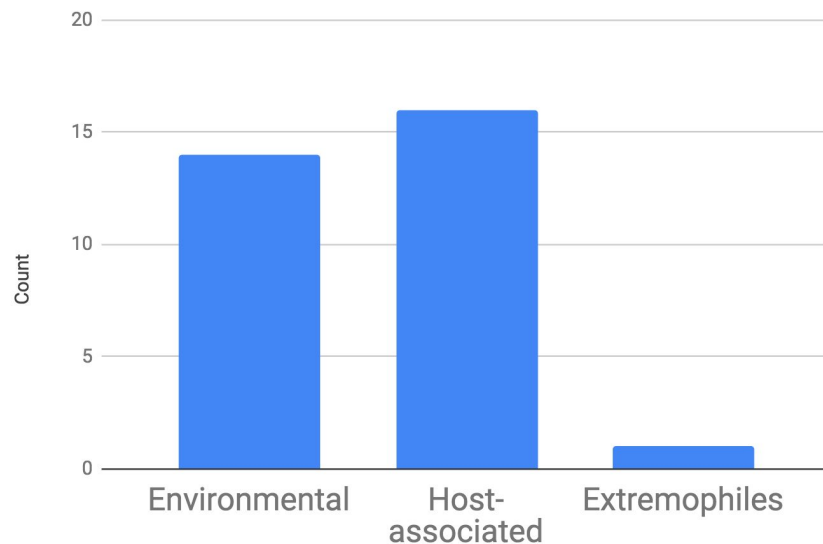
- Name
- Position and Institution
- Ten words or less -- an interesting fact about them:
  - What do they study?
  - Why do they hope to learn from the workshop?
  - What is their favorite microbe?

# Results from the Pre-Workshop Survey

What kind of microbial data do you typically analyze?

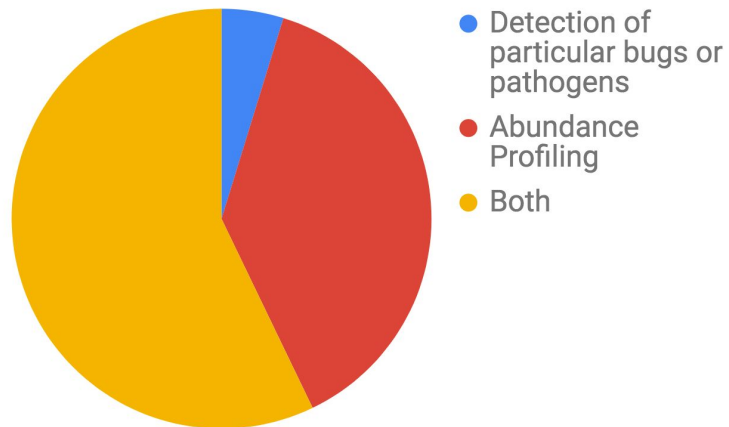


What systems do you typically study?

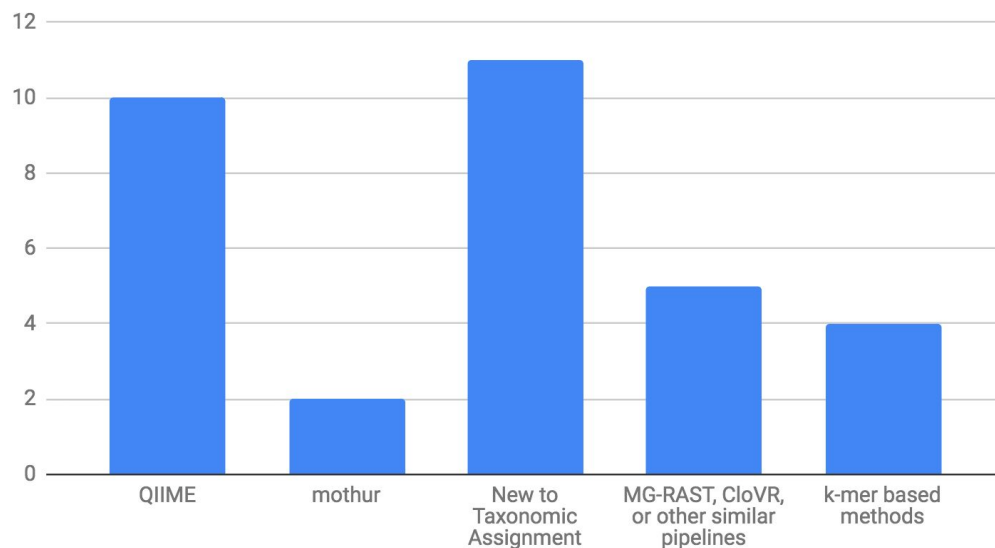


# Results from the Pre-Workshop Survey

When you perform taxonomic assignment, what are your primary goals?



What tools have you used for taxonomic assignment?



# Results from the Pre-Workshop Survey

On a scale from 1 to 10, how comfortable are you with using the command line?

23 responses

