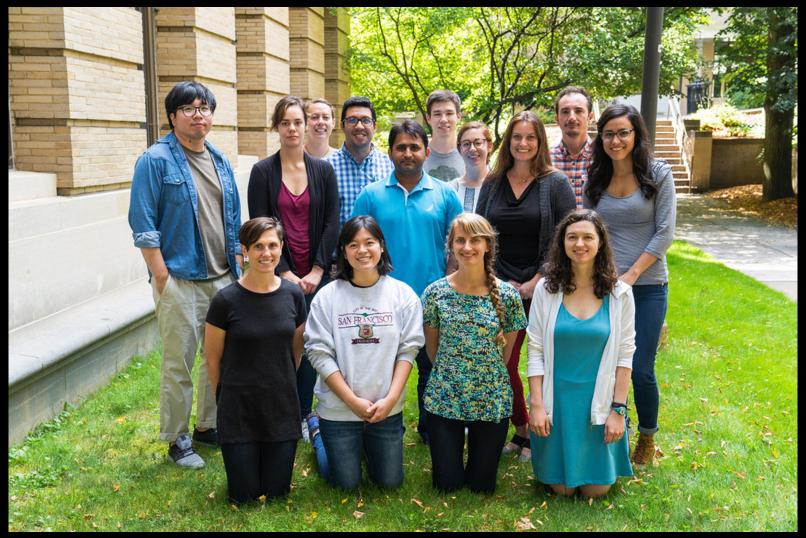
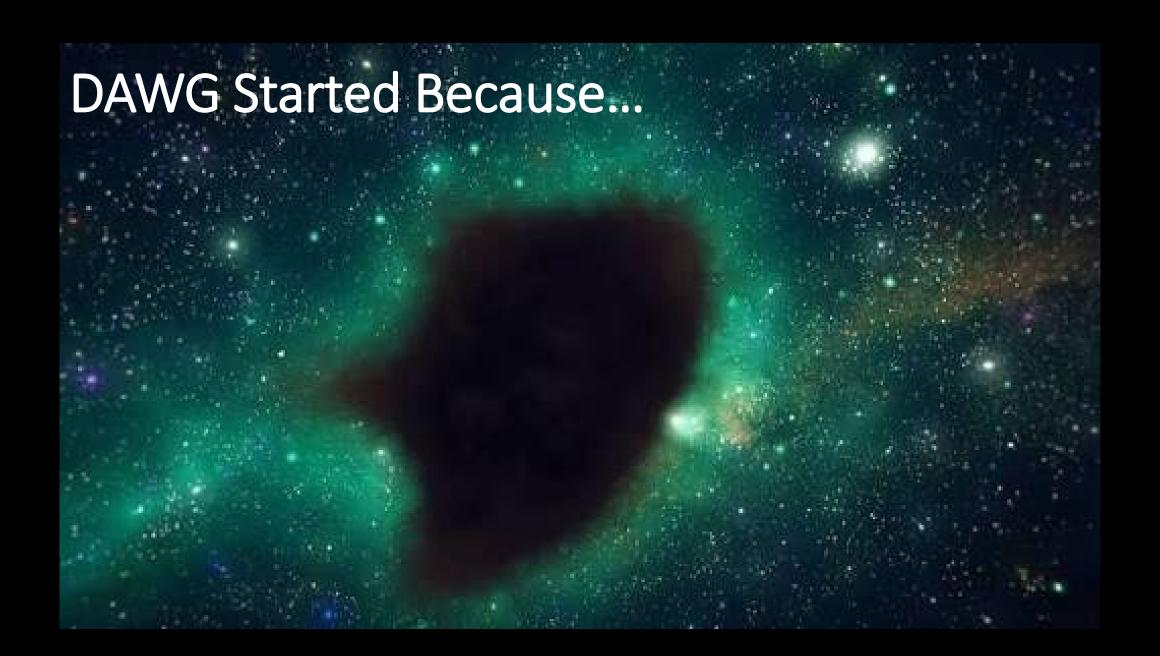
Microbiome
Center's DAWG
(data analysis
working group)



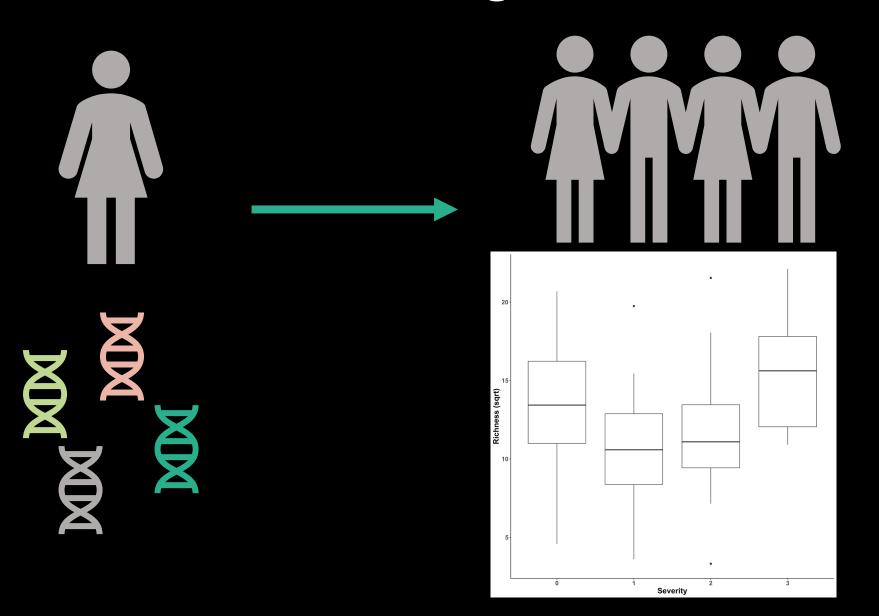
Who is DAWG?!?!

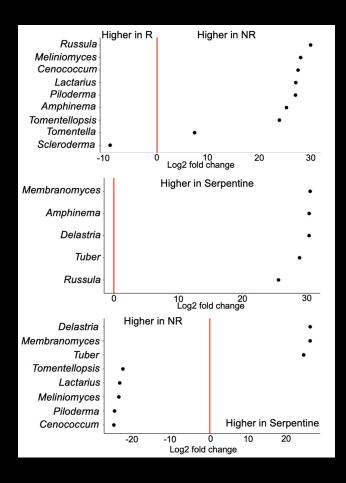


https://news.psu.edu/story/588593/2019/09/17/academics/penn-state-microbiome-centers-dawg-chews-microbiome-data-sets



What is DAWG Doing?





DAWGs Series of Workshops

- Cohort building
- Career development (publications/workshops)

- Training
- Stay up to date on latest packages/analyses



DAWGs Goals Are to:

- Use the skills of our Microbiome Center members to help a researcher at Penn State process data and analyze results, culminating in a publication or potentially being used as preliminary data for a grant application,
- Provide a unique opportunity for students/post-docs to further develop transferable skills as group leaders of DAWG,
- and help build collaborations between the members already associated with the Microbiome Center and Penn State researchers who are interested in microbiome research.
- But, the main goal of the workshops is to train graduate students and post-doctoral scholars how to process/analyze microbiome datasets

What has DAWG done so far?

- Started in Spring 2019
- Hosted 4 workshops on processing/analyzing a 454 pyrosequencing dataset for Dr. Bert Eardly from Penn State Berk's
- Ectomycorrhizal fungi from pitch pine roots along a metals gradient associated with a Zinc smelter
- DAWG leaders + Bert presented our findings to the Microbiome Center
- Bert presented our findings at the annual ASM meeting



Average number of attendees per workshop was 14

What has DAWG done so far? (ctd)

- Fall 2019/Spring 2020
- Held 4 workshops each semester
- Working with Dr. Judie Howrylak on analyzing 16S amplicon sequences from patients that were admitted into the ICU at Penn State Hershey Medical Center
- Contains ~ 230 samples from ~100 patients
- Currently writing manuscript for stool samples

Average number of attendees was 15 (until COVID)

Topics we have covered

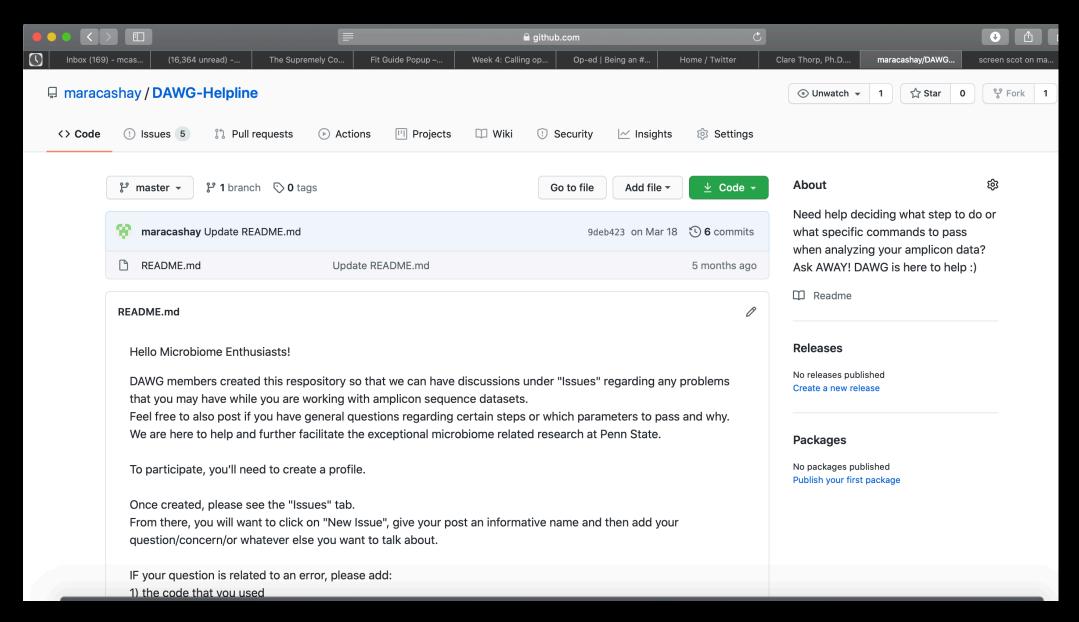
- ASV versus OTU
- Rarefying data
- Which reference data base to use
- How to incorporate ancillary data with amplicon sequencing data
- PICRUSt analysis/interpretation
- Network analyses
- Machine learning

DAWGs Plan for Fall 2020

- METAGENOMICS!
- Have 5-6, 1 hour long presentations (including Kbase tutorial)
- MAGs (reconstruction and annotation of genomes)
- Marker Gene Analysis (Nitrogen, carbon, phosphorous cycling, antibiotic resistance genes, etc.)

- First session: Aug 28th at 1 pm Dr. Steven Higgins, post-doctoral scholar for Dan Buckley and now works with Michelle Heck at Cornell
- 4 PSU students and Dr. Luis Orellana

DAWGs Helpline: Github.com/maracashay



DAWGs Future

- We need you!!!!
- Participants
- Workshop leaders
- Any level of expertise ©
- Data sets

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