

## **MICB 475 Meeting November 25th**

### **1. Progress Updates - issues and troubleshooting**

- a. ANCOM-BC
- b. ALDEx2
- c. Picrust
- d. Random forest

### **2. Updated powerpoint slides**

- i. [MICB 475 - Team 10 - Meeting 10- 11/26/2024 - Google Slides](#)
- ii. Choosing figures for manuscript - For actual paper?
- iii. Appendix
- b. Aim 1
  - i. Conclusion: Looking at alpha diversity between 2 groups, there are no statistically significant differences
  - ii. MANUSCRIPT: Observed, Chao1 and Shannon = typical observed ones in a microbiome paper (Figure 1)
  - iii. PMS should change to pMS
  - iv. Beta Diversity: add circles around groups
- c. Aim 2
  - i. Only include Jaccard / Bray-Curtis
  - ii. Table for supplemental pairwise analysis (supplemental and table for presentation)
  - iii. Taxa bar plot
    - 1. Make labels bigger
    - 2. Mostly pMS making an impact
    - 3. Crop out only to include main colours
    - 4. Look into Verrucomicrobiae
- d. Aim 3
  - i. Core Microbiome
    - 1. Make taxa bar plots → look at top 80%, or look at bottom 20%
    - 2. Colour bars in a different colour
    - 3. Use “healthy” consistently in labels
    - 4. Two venn diagrams from first slideshow
    - 5. Include those instead for figure 3: do the age one
    - 6. Can add in results that we did age by but have no significance
    - 7. Losing some species with age but then the smoking is increasing other species
  - ii. ALDeX2
    - 1. Instead: do old smokers with oms/healthy. Young smokers pms/healthy. Young non smokers - pms/healthy

2. Young more susceptible to smoking
  - i. Influential, pMS is an additional factor
  - ii. Control
  - iii. Supplemental
  - iv. Four panel:
    - i. Young non smokers PMS v. healthy
    - ii. Young smokers PMS v. healthy
    - iii. Old non smokers PMS v. healthy
    - iv. Old smokers PMS v. healthy
  - v. Annotate the species in the graphs
  - vi. Stick to volcanos
  - vii. Be transparent that most not significant
- iii. Picrust
  1. Make MS comparisons
  2. Upregulated vs down regulated
  3. Table if many pathways
  4. Figure if few pathways
- e. Aim 4
  - i. Refine model. Do 1 prediction for young and 1 for old
  - ii. Test old model on young to verify prediction differences

#### Action items

- Listed in minutes