## **Meeting October 1st**

## Research question

- What species of microbes are changing in MS patients with an increase in age and severity? Aims are as follow:
- Do for one city to narrow down the samples
  - 1. Correlation between age and severity
    - Group between male and female patients
  - Correct for TD2 and smoking to get rid of factors that may impact microbial diversity (could be 2nd aim to eliminate confounding variables)
  - 3. Look at alpha and beta diversity (PCA plot, group by age colour severity) for each age group and different severity
  - 4. Microbes researched can then be correlated to inflammation
    - Inflammatory microbes to tie into literature
    - Do the inflammatory microbes vary between patients with other variables
  - 5. Random forest if time

## To start off:

- 1. Trim down manifest before qiime2
- 2. Figure how many samples relevant (ideal sample size in the hundreds)
  - Make a revised manifest
- 3. All figures are needed for the proposal
  - Chris can look at some of the files at meetings

## **Action items**

- 1. Generating a new manifest files after trimming data
  - Pick a city with more than 100 samples and filter other factors as discussed
- 2. Start writing proposal (due the 20th)
  - Divide the aims for all 5 members to write
  - MS severity and different microbial differences
    - Eg: age component and lifestyle (smoking)
    - Does smoking result in a change in microbiomes independent of age in MS patients?
- 3. More formal agenda for next meeting