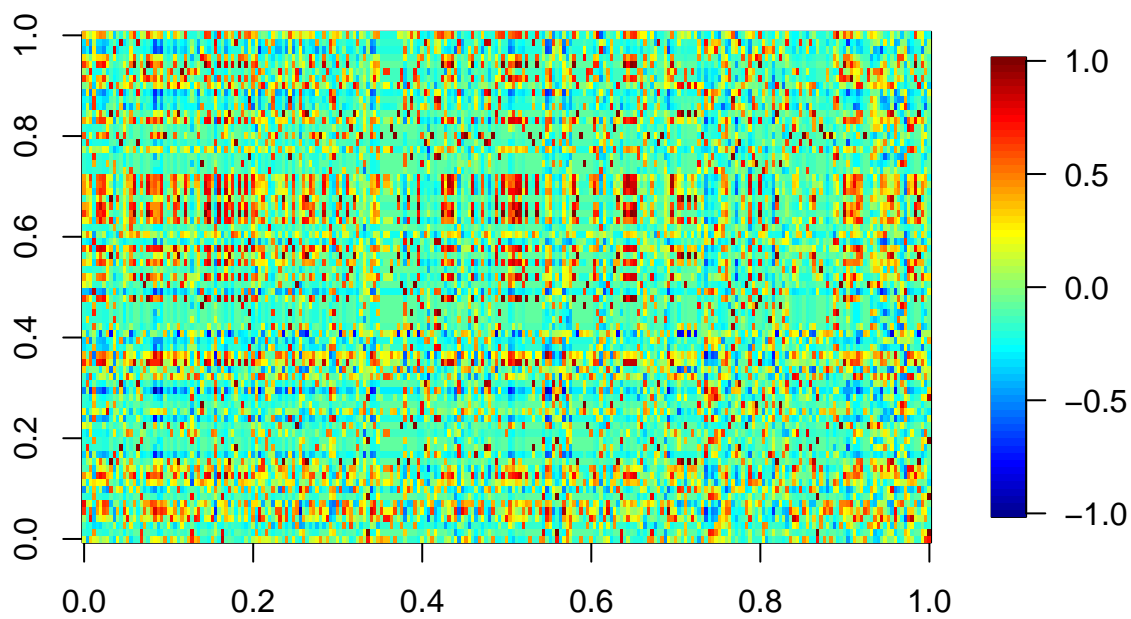


Correlations

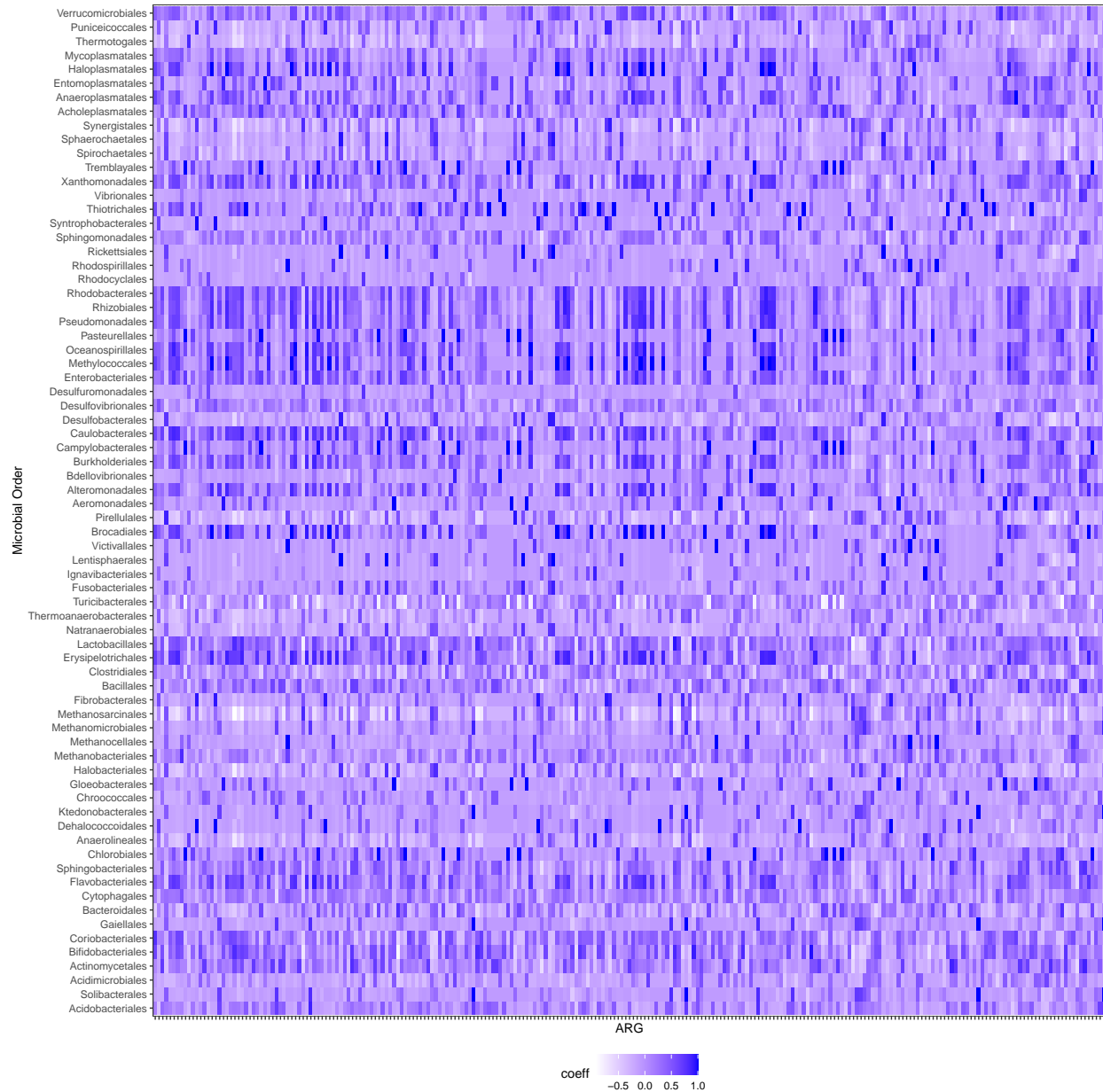
Daniel Flores Orozco

08/02/2022



```
## Joining, by = "microbe"
```

```
## Joining, by = "gene_id"
```



```
## Warning: Setting row names on a tibble is deprecated.
```

```
## Warning: Setting row names on a tibble is deprecated.
```

```
## Warning in brewer.pal(ColourCount1, "Paired"): n too large, allowed maximum for palette Paired is 12
## Returning the palette you asked for with that many colors
```

