

Step2_RNA_Analysis

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Load all the packages used in the analysis

```
library("ranacapa")
library("phyloseq")
library("ggplot2")
library("stringr")
library("plyr")
library("reshape2")
library("reshape")
library("dplyr")
library("tidyr")
library("doBy")
library("plyr")
library("microbiome")
library("ggpubr")
library("vegan")
library("tidyverse")
library("magrittr")
library("cowplot")
library("dendextend")
library("WGCNA")
library("metagenomeSeq")
library("decontam")
library("RColorBrewer")
library("ampvis2")
```