BIOST 546 Final Project

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```r
# set global options for code chunks
knitr::opts_chunk$set(message = FALSE, warning = FALSE, collapse = TRUE)
knitr::opts_knit$set(root.dir = rprojroot::find_rstudio_root_file())

library(dplyr)
library(knitr)
library(ggplot2)
```

```r
# load data
load("./dataset/ADProj.RData")
str(ADProj, max.level = 1)
## List of 3
##  $ X_train: tibble [400 x 360] (S3: tbl_df/tbl/data.frame)
##  $ y_train: tibble [400 x 1] (S3: tbl_df/tbl/data.frame)
##  $ X_test : tibble [400 x 360] (S3: tbl_df/tbl/data.frame)
```