Problem 1:

create a simple R package called forloops that reproduces some Base R functions using For-loops.

Part a:

• Create a folder called "forloop", set the working directory to this folder



Part b:

• In the R sub-directory create a R script called practice



Part c:

• Build a function inside of R script practice called col_means() that will take as input a data frame and return a vector of column means. We will not use the colMeans() function from {base} package

```
col_means <- function(df) {
  means <- numeric(ncol(df))
  for (i in seq_along(df)) {
    means[i] <- mean(df[[i]], na.rm = TRUE)
  }
  return(means)
}</pre>
```

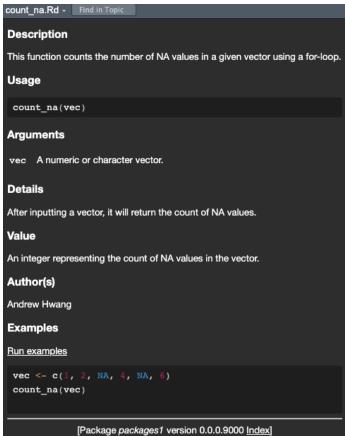
Part d:

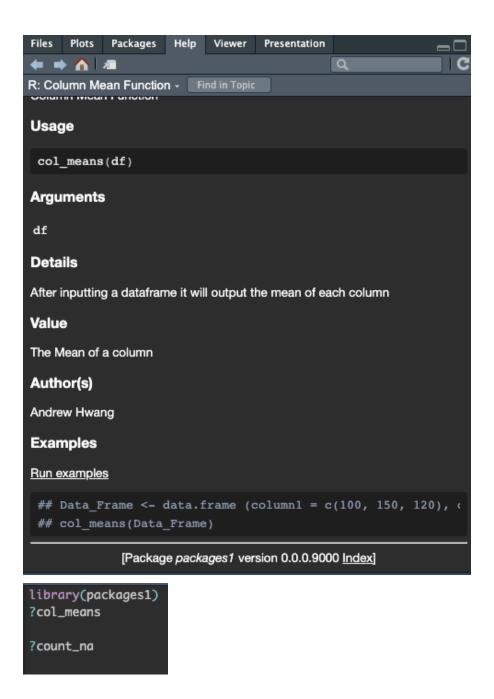
• Build a function inside of R script practice called count_na() that will use a for-loop to count how many NA's there are in a vector.

```
count_na <- function(vec) {
  count <- 0
  for (i in seq_along(vec)) {
    if (is.na(vec[i])) {
      count <- count + 1
    }
  }
  return(count)</pre>
```

Part e:

• Create documentation for each functions that you have in your R script. Then load the package and type ?col_means and then ?count_na.





Part f:

• In your R studio move to Terminal and type Is, it will show the list of all files and folders that you have in your working directory forloops.

```
andrews-mbp-2:forloops andrewhwang$ ls

DESCRIPTION LICENSE LICENSE.md NAMESPACE R man
andrews-mbp-2:forloops andrewhwang$ ls
```