## Question1.R

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```
library(tidyverse)
## -- Attaching packages ----- tidyverse 1.2.1 --
## <U+221A> ggplot2 3.1.0
                             <U+221A> purrr
                                              0.2.5
## <U+221A> tibble 1.4.2
                              <U+221A> dplyr
## <U+221A> tidyr
                   0.8.1
                              <U+221A> stringr 1.3.1
## <U+221A> readr
                   1.1.1
                             <U+221A> forcats 0.3.0
## Warning: package 'dplyr' was built under R version 3.5.1
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                  masks stats::lag()
# Q1. Run the below command and the parts:
   mtcars %>%
        group_by(cyl) %>%
        summarize_at(.vars=c('mpg', 'drat'), funs(mean))
## # A tibble: 3 x 3
      cyl mpg drat
     <dbl> <dbl> <dbl>
##
        4 26.7 4.07
## 1
## 2
        6 19.7 3.59
## 3
        8 15.1 3.23
    #1. Easy - Modify the "funs" command and see if you can calculate mean,
                    # sum and count, don't change the summarise_at command.
    #2. Hard - Can you use quantile instead of mean and see what happens?
                    # What error you got? Fix it and run for quantile
    #3. Easy - Find mean, count, median, sum for all the continous data in mtcars,
                    # grouped across cyl. Please avoid descrete column values.
# Q2. Easy - Use iris data and find the summary stats( count , sum , mean, std. deviation)
                    # for all the numeric columns grouped by Species
# Q3. Easy - Run a gaplot for the density plot for all the unique Species types from iris
                    # dataset for Sepal.Length column
# Q4. Easy - Plot a histogram across Each "cut" for depth from diamonds data
                   # (use library(ggplot2) to access diamonds dataset)
# Q5. Medium
   df \leftarrow list(df1 = data.frame(x = 1:10, y = letters[1:10]),
                    df2 = data.frame(x = 11:20, y = LETTERS[11:20]))
    \# Use the above "df" and append the list item to make one table
    # Make column y in the appended column as all uppercase
    # Use lapply to fetch 1st column and 2nd column separately
# Q6. Hard - split the superhero names with their ages in the given vector:
    vector <- c("Superman1000", "Batman35", "Wonderwoman240")</pre>
# Q7. Easy - Return a copy of hflights that contains the four columns related to delay
                    \# ( ActualElapsedTime , AirTime , ArrDelay , DepDelay )
```

```
## x1 X2 X3 X4 X5
## a 1 2 3 4 5
## b 1 2 3 NA NA
```