STATS 506 Problem Set #3

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Vision

a. Read data and merge data

```
library(haven)
vix <- read_xpt('./VIX_D.XPT')
demo <- read_xpt('./DEMO_D.XPT')
df <- merge(vix, demo, by='SEQN')
cat('Sample size:', nrow(df))</pre>
```

Sample size: 6980

b. The max age is 85, so there will only be 9 age brackets.

Age Group	Proportion
10-19	32.09
20-29	32.66
30-39	35.87
40-49	37.00
50-59	55.01

Age Group	Proportion
60-69	62.22
70-79	66.89
80-89	66.88

c. Here are fitted models and their summary.

```
#' Create Summary Table for Logistic Regression
# '
#' @param model a fitted logistic regression model
#' @return a table with required stats
summary_table <- function(model) {</pre>
  odds_ratios <- as.data.frame(t(exp(coef(model))))</pre>
  res <- data.frame(</pre>
  'Sample Size' = nobs(model),
  'Pseudo R2' = 1 - model$deviance / model$null.deviance,
  'AIC' = AIC(model)
  res <- cbind(odds_ratios, res)</pre>
  return(t(res))
# data cleaning
df mod <- subset(df, select=c(VIQ220, RIAGENDR, RIDAGEYR, RIDRETH1, INDFMPIR))
df_mod <- na.omit(df_mod)</pre>
df_mod$VIQ220 <- as.factor(df_mod$VIQ220)</pre>
df_mod$RIAGENDR <- as.factor(df_mod$RIAGENDR)</pre>
df_mod$RIDRETH1 <- as.factor(df_mod$RIDRETH1)</pre>
# model fitting
mod1 <- glm(VIQ220 ~ RIDAGEYR, data = df_mod,</pre>
             family = binomial(link = 'logit'))
knitr::kable(summary_table(mod1))
```

(Intercept)	3.4170427
RIDAGEYR	0.9760682
Sample.Size	6247.0000000
Pseudo.R2	0.0473330
AIC	8119.8714676

6.0521094
0.9779121
0.8560069
0.5277604
0.7737545
0.5310998
0.6053170
6247.0000000
0.0685555
7949.0753329

(Intercept)	7.5094311
RIDAGEYR	0.9780560
RIDRETH12	0.8904552
RIDRETH13	0.6056040
RIDRETH14	0.8127071
RIDRETH15	0.5870020
RIAGENDR2	0.5967415
INDFMPIR	0.8926169
Sample.Size	6247.0000000
Pseudo.R2	0.0733995
AIC	7909.8082208

d. From previous part, we have the odds ratio for women is 0.5967415. This can be interpreted as the value of female odds divided by male odds. From the summary of model 3, the coefficient is significant, thus implying that female odds differs from male odds for being a glass wearer is statistically significant. Since the positive class of glm is the last level of the factor, 2 in this case, the positive class for the model is actually 'not a glass wearer'. Thus, we need to invert our interpretation. Having an odds ratio less than 1 from the model actually should imply that the odds of females wearing glasses/contacts for distance vision is higher than male.

summary(mod3)

```
Call:
glm(formula = VIQ220 ~ RIDAGEYR + RIDRETH1 + RIAGENDR + INDFMPIR,
   family = binomial(link = "logit"), data = df_mod)
Coefficients:
            Estimate Std. Error z value Pr(>|z|)
                      0.087788 22.966 < 2e-16 ***
(Intercept) 2.016160
RIDAGEYR
           -0.022188
                      0.001295 -17.135 < 2e-16 ***
RIDRETH12
           -0.116023
                     0.168265 -0.690 0.490495
           RIDRETH13
RIDRETH14
           -0.207385
                      0.079217 -2.618 0.008847 **
           -0.532727
                      0.140152 -3.801 0.000144 ***
RIDRETH15
           -0.516271
                      0.054305 -9.507 < 2e-16 ***
RIAGENDR2
INDFMPIR
           -0.113598
                      0.017707 -6.415 1.41e-10 ***
___
              0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Signif. codes:
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 8519.1 on 6246 degrees of freedom
Residual deviance: 7893.8 on 6239 degrees of freedom
AIC: 7909.8
Number of Fisher Scoring iterations: 4
```

As shown by the two sample proportion test, the p-value is extremely small. Thus, we can reject the null hypothesis and conclude that the proportion of wearers of glasses/contact lenses for distance vision differs between men and women. We can even say that male proportion is less than female proportion. (according to the fact that the confidence interval is below 0)

```
tab <- table(df_mod$RIAGENDR, df_mod$VIQ220)
prop.test(tab[,1], rowSums(tab))</pre>
```

2-sample test for equality of proportions with continuity correction

```
data: tab[, 1] out of rowSums(tab)
X-squared = 69.683, df = 1, p-value < 2.2e-16</pre>
```

```
alternative hypothesis: two.sided

95 percent confidence interval:

-0.12945505 -0.08007986

sample estimates:

prop 1 prop 2

0.3714379 0.4762054
```

Sakila

a. It appears that all movies are from 2006, which is the earliest release year from this database.

```
library(DBI)
sakila <- dbConnect(RSQLite::SQLite(), './sakila_master.db')
dbGetQuery(sakila,'
    SELECT film_id, title, release_year
    FROM film
    ORDER BY release_year ASC
')</pre>
```

b. Here's the R approach, note that the min value is unique.

```
film_cat <- dbGetQuery(sakila, 'SELECT * FROM film_category')
category <- dbGetQuery(sakila, 'SELECT * FROM category')
cat_count <- table(film_cat$category_id)
min_cat <- which.min(cat_count)
cat(category$name[category$category_id == min_cat], cat_count[min_cat])</pre>
```

Music 51

Here's the SQL approach

```
dbGetQuery(sakila, '
   SELECT c.name, COUNT(*) AS count
FROM film_category AS fc
   JOIN category AS c
   ON fc.category_id = c.category_id
   GROUP BY c.category_id
   ORDER BY count ASC
   LIMIT 1
')
```

```
name count
1 Music 51
```

c. Here's the R approach

```
customer <- dbGetQuery(sakila, 'SELECT * FROM customer')
address <- dbGetQuery(sakila, 'SELECT * FROM address')
city <- dbGetQuery(sakila, 'SELECT * FROM city')
country <- dbGetQuery(sakila, 'SELECT * FROM country')
merged_df <- merge(customer, address, by='address_id')
merged_df <- merge(merged_df, city, by='city_id')
merged_df <- merge(merged_df, country, by='country_id')
res <- table(merged_df$country)
res[res == 13]</pre>
```

```
Argentina Nigeria
13 13
```

Here's the SQL approach.

```
dbGetQuery(sakila, '
   SELECT country.country, COUNT(*) AS count
FROM customer, address, city, country
WHERE customer.address_id = address.address_id AND
   address.city_id = city.city_id AND
   city.country_id = country.country_id
GROUP BY country.country_id
HAVING count = 13
')
```

```
country count
1 Argentina 13
2 Nigeria 13
```

US Records

a. Here's the proportion of TLD with ".com'

```
df <- read.csv('./us-500.csv')
cat('Proportion of .com:', mean(grepl('\\.com$', df$email)))</pre>
```

Proportion of .com: 0.732

b. Here's the proportion of email with at least one non alphanumeric character in them. Since it's possible to have ".' in the username, we need to seperate the username and domain, then check each part separately.

```
emails <- strsplit(df$email, '@')
usernames <- lapply(emails, '[[', 1)
domains <- lapply(emails, '[[', 2)
domains <- gsub('\\.[a-z]{3}', '', domains)
mean(grepl('[^a-zA-Z0-9]+', usernames) | grepl('[^a-zA-Z0-9]+', domains))</pre>
```

[1] 0.506

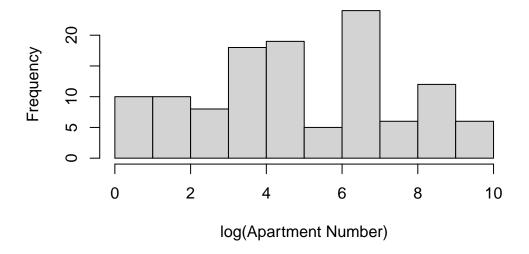
c. Here's the top 5 area code. Notice that there is no ties in top 5, so I can directly use the top 5 element.

```
res <- table(c(substr(df$phone1, 1, 3), substr(df$phone2, 1, 3)))
sort(res, decreasing=TRUE)[1:5]</pre>
```

```
973 212 215 410 201
36 28 28 28 24
```

d. Here's the frequency histogram of the apartment numbers.

Frequency of Apartment Numbers



e. The data appears to be synthetic, as the first digit does not have a decreasing trend.

```
first_digit <- as.numeric(substring(apt_nbrs, 1, 1))
hist(first_digit, main = 'First Digit Distribution', xlab = '', freq = FALSE)</pre>
```

First Digit Distribution

