The following libraries are necessary and need it to be downloaded and up to date, in order to develop the analysis required.

FB1 is a named assigned to the data provided, it can be renamed if the sponsor chooses to do so.

```
# Load in library
library(rpart)
library(rpart.plot)
library(readr)
library(dplyr)
library(foreign)
library(ggplot2)
library(class)
library(useful)
library(tidyverse)
library(caret)
library(gbm)
library(ISLR)
library(skimr)
library(party)
library(tree)
library(DAAG)
library(mlbench)
library(pROC)
library(rattle)
options(scipen = 999)
FB1 <- read_csv("FB1.csv")
fb1<- FB1[-1]
```

As stated on the code the following variables were eliminated because there were no values at all. Therefore, the variables are meaningless to our analysis.

Next, the variables containing only dates related to customers were assigned to a variable just to make the regression tree more visible. Dates do not affect our analysis, but the variables were not eliminated, rather they were assigned just in case they are required for a specific analysis.

After the variables containing dates were saved and assigned to a Datfb1 variable, then those variables are eliminated from our data set.

Variables characterized as characters need to be converted to factors because R does not handle characters as values, rather as factors. Notice the variable Client ID contains numbers but for our analysis the Client ID variable will helps us identify a specific client. Thus, it will be considered a categorical variable but needs to be converted to factor.

The next function in R Studio is creating a new column our data frame fb1 named "Client Preferred Agency", and populating it with binary values (either 1 or 0) based on whether the agency name "Interfaith Food Bank" appears in the original "Client Preferred Agency" column.fb1\$'Client Preferred Agency': This accesses the "Client Preferred Agency" column in the fb1 data frame using the \$ operator.

Here's a step-by-step breakdown of what's happening in the code:

- 1. <- with(fb1, ...): This specifies that the subsequent code will be executed within the context of the fb1 data frame.
- 2. ifelse(grepl("Interfaith Food Bank", Client Preferred Agency), 1, 0): This is a conditional statement that checks whether the string "Interfaith Food Bank" is present in the "Client Preferred Agency" column of fb1. If it is, the function returns 1. If it isn't, the function returns 0. The ifelse function in R allows for vectorized if-else statements, which means it can operate on entire columns of data at once.
- 3. The result of step 3 is then assigned to the new column "Client Preferred Agency" in fb1.

In summary, this code is creating a new binary column that indicates whether or not each row in the original "Client Preferred Agency" column contains the string "Interfaith Food Bank".

In addition "Interfaith Food Bank" was chosen because diving through the Client Preferred Agency database we observe the agency Interfaith Food Bank had the most visitors among the rest of the agencies. Notice that this can be adjusted and if wanted to test the regression trees, all is needed is to change the variable name.

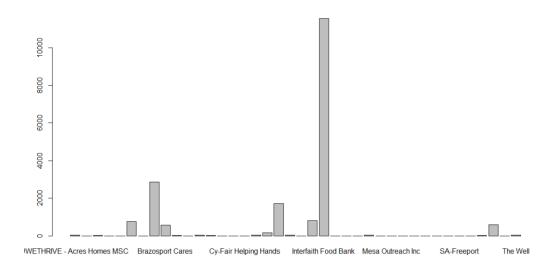
```
> table(fb15'Client Preferred Agency')

#WETHRIVE - Acres Homes MGC
#WETHRIVE - Fifth Ward MGC
24
0
0
A Place For Grace Foundation

Angleton Jr High
8 Brazoria County Dream Center
2869
8 Brazosport Cares
8 Brazosport Cares
8 Brazosport Cares
0
Compton Memorial Church of God in Christ
27
Covenant with Christ Heritage Pentecostal Church
10
Covenant with Christ Heritage Pentecostal Church
11
Covenant With Christ Heritage Pentecostal Church
11
1 Covenant With Christ Heritage Pentecostal
11
1 Food For Change
173
```

If necessary the Client Preferred Agency can be plotted to confirm our decision, using the following command:

plot(fb1\$`Client Preferred Agency`)



This function in R Studio is creating a new factor variable in the **fb1** data frame called "Client Marital Status", based on the values in the existing "Client Marital Status" column.

Here's a step-by-step breakdown of what's happening in the code:

- 1. **fb1\$'Client Marital Status'**: This accesses the "Client Marital Status" column in the **fb1** data frame using the \$ operator.
- factor(fb1\$'Client Marital Status', ...): This function creates a new factor variable based on the
 values in the "Client Marital Status" column of fb1. Factors are a special data type in R used for
 categorical variables.
- 3. levels = c("0", "common_law", "declined_to_answer", "did_not_ask", "divorced",
 "do_not_know", "married", "separated", "single", "widowed"): This specifies the levels of the
 new factor variable. In this case, the factor levels are a set of character strings that correspond to
 the possible values in the original "Client Marital Status" column.
- 4. **labels** = **c(0,1,2,3,4,5,6,7,8,9)**: This specifies the labels to be assigned to each factor level. In this case, the function is assigning numeric values to each factor level, with "0" being assigned the label "0", "common law" being assigned the label "1", and so on.
- 5. The result of step 2 is then assigned to the new column "Client Marital Status" in **fb1**

Using the following function, we can extract all the characteristics on the Client Marital Status:

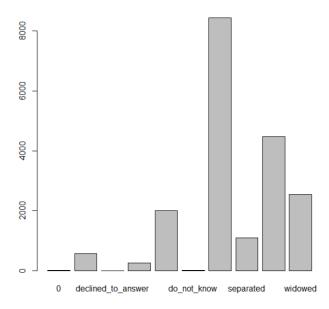
```
table(fb1$`Client Marital Status`)
```

```
> table(fb1$`Client Marital Status`)
```

```
0
                     common_law declined_to_answer
         19
                            566
did_not_ask
                       divorced
                                       do_not_know
                           2009
                                                 15
        265
                      separated
                                             single
    married
       8440
                           1097
                                               4468
    widowed
```

In addition, the function can be plot to visualize the characteristics of the variable, using the following function:

```
plot(fb1$`Client Marital Status`)
```



The same procedure was applied to the rest of the variables.

The following code we just rename the variables again to better fit the regression tree.

names(fb1): will print out all the variable names in the R console

attach(fb1): attaching the names will help you obtained the information needed in the dataset

sum(is.na(fb1)) : count total missing values in entire data frame

apply(X = is.na(fb1), MARGIN = 2, FUN = sum): prints the number of N/A values for each variable as a list

fb1 <- fb1[complete.cases(fb1),]: deletes all the missing values in the dataset without affecting the analysis.

sum(is.na(fb1)): prints the number of missing values after using the complete case function abovedim(fb1): prints out the numbers of rows, and columns. For our dataset the following is true: 19,428101

Setting a seed guarantees the same output no matter how many times you run the program.

The partition variable above is creating a partition of the data in the **fb1** data frame by randomly selecting a subset of rows for training a machine learning model.

Here's a step-by-step breakdown of what's happening in the code:

- 1. **fb1\$'MothHoldinc'**: This accesses the "MothHoldinc" column in the **fb1** data frame using the \$ operator. This column likely contains the target variable for the machine learning model.
- createDataPartition(y = fb1\$'MothHoldInc', p = 0.8, list = FALSE): This function is part of the caret package in R, which provides tools for training and evaluating machine learning models.
- 3. **y = fb1\$'MothHoldInc'**: This specifies the target variable for the machine learning model.
- 4. **p = 0.8**: This specifies the proportion of the data to be included in the training set. In this case, 80% of the data will be used for training.
- 5. **list = FALSE**: This specifies that the function should return a vector of row indices rather than a list of data frames.
- 6. The result of step 2 is then assigned to the **partition** variable.

The following functions split the data into two parts, one as a training set and the rest as a test.

```
cs_train <- fb1[partition, ]
cs_test <- fb1[-partition, ]</pre>
```

This function in R Studio is building a decision tree regression model on the **cs_train** data frame, using the Recursive Partitioning and Regression Trees (rpart) algorithm, to predict the target variable **MothHoldinc**.

Here's a step-by-step breakdown of what's happening in the code:

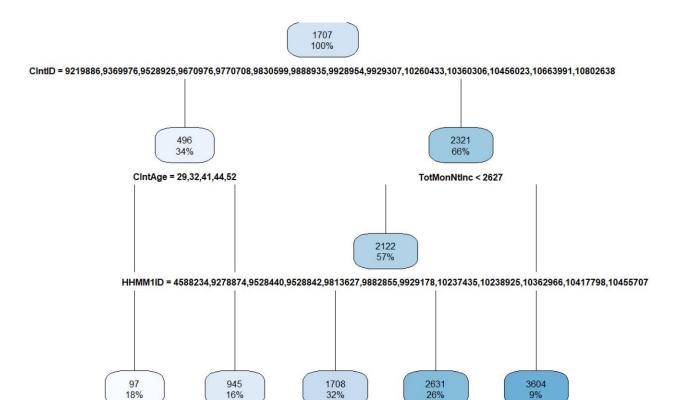
- 1. **rpart**(MothHoldInc ~ ., **cs_train**, **method** = **"anova"**): This function is using the **rpart** algorithm to build a decision tree model, where **MothHoldInc** is the target variable to be predicted and . represents all the other variables in the **cs_train** data frame.
- 2. **cs train**: This is the training data frame that the model will be fit to.
- 3. **method = "anova"**: This specifies the method used for splitting the data at each node of the decision tree. In this case, the **anova** method is used to perform an analysis of variance to determine the best variable to split the data.
- 4. The resulting decision tree model is then assigned to the cs_mdl_cart_full variable.

The function **print(cs_mdl_cart_full)** will allow us to print the information that will be used to generate the regression tree. Notice, Client ID, Client Age, Total Monthly Income, HH Member 1 ID, and Total Monthly Net Income, were the most important variables in our analysis.

```
> print(cs_mdl_cart_full)
n = 101
node), split, n, deviance, yval
      * denotes terminal node
1) root 101 126419100.0 1706.80900
  2) clntID=9219886,9369976,9528925,9670976,9770708,9830599,9888935,9928954,9929307,1
0260433,10360306,10456023,10663991,10802638 34
                                               7883529.0
                                                           496.29410
     4) ClntAge=29,32,41,44,52 18
                                   342361.1
                                               97.22222 *
     5) ClntAge=34,43,45,57,59,65 16
                                     1449535.0 945.25000 *
   3) clntiD=9194925,9278821,9279341,9369237,9445162,9528410,9528575,9548852,9623330,9
813593,9875184,9882821,9928956,9987172,10165425,10238777,10353058,10402869,10417842,10
455646,10591060,10714125,10714481,10783423 67 43431190.0 2321.10000
     6) TotMonNtInc< 2626.5 58 16030630.0 2121.99400
     12) HHMM1ID=4588234,9278874,9528440,9528842,9813627,9882855,9929178,10237435,102
38925,10362966,10417798,10455707 32 1533488.0 1708.12500 *
     13) HHMM1ID=2334784,9369383,9875235,9928961,9943116,9987183,10165436,10402956,10
591104,10714214,10714538,10783555 26
                                     2269819.0 2631.37200 *
     7) TotMonNtInc>=2626.5 9 10283640.0 3604.22200 *
```

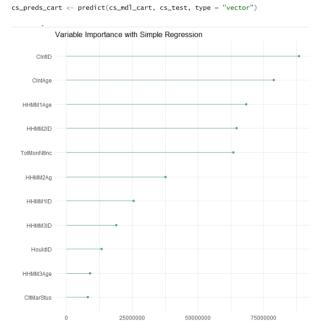
The following function will plot the tree for a better visualization of the results.

```
rpart.plot(cs_mdl_cart_full, yesno = TRUE)
```



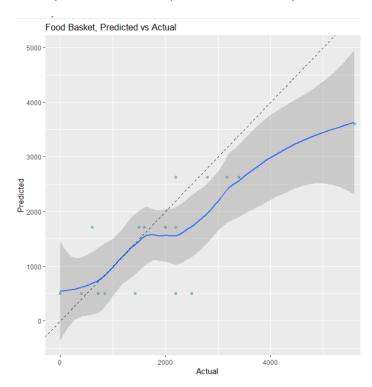
The following function will plot the most important variables:

```
cs_mdl_cart$variable.importance %%
  data.frame() %%
  rownames_to_column(var = "Feature") %%
  rename(overall = '.') %%
  ggplot(aes(x = fct_reorder(Feature, Overall), y = Overall)) +
  geom_pointrange(aes(ymin = 0, ymax = Overall), color = "cadetblue", size = .3) +
  theme_minimal() +
  coord_flip() +
  labs(x = "", y = "", title = "Variable Importance with Simple Regression")
```



There are less than 2,000 possible predicted values binning the observations.

```
data.frame(Predicted = cs_preds_cart, Actual = cs_test$MothHoldInc) %>%
    ggplot(aes(x = Actual, y = Predicted)) +
    geom_point(alpha = 0.6, color = "cadetblue") +
    geom_smooth() +
    geom_abline(intercept = 0, slope = 1, linetype = 2) +
    labs(title = "Food Basket, Predicted vs Actual")
```



The following functions helps us predict the accuracy of our model:

```
p <- predict(cs_mdl_cart_full, cs_train)

sqrt(mean(cs_train$`MothHoldInc`-p)^2)

#sually, the larger the R2, the better the regression model fits
#your observations, r square, 0.826503
(cor(cs_train$`MothHoldInc`, p))^2</pre>
```

Our model turns out to have an 83% of accuracy.