

Predicting Total Wealth: A Predictive Analysis Using the 1991 SIPP Data

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Introduction

Loading and Inspecting the Data

Let's take a look at the first 6 rows of the data.

```
data <- read.table('data_tr.txt', head = T)[-1]
head(data)
```

```
##      tw  ira e401  nifa  inc hmort  hval hequity educ male twoearn nohs hs
## 1  53550    0    0   100 28146 60150  69000   8850   12    0      0    0    1
## 2 124635    0    0 61010 32634 20000  78000  58000   16    0      0    0    0
## 3 192949 1800    0   7549 52206 15900 200000 184100   11    1      1    1    0
## 4   -513    0    0   2487 45252    0      0      0   15    0      1    0    0
## 5 212087    0    0 10625 33126 90000 300000 210000   12    0      0    0    1
## 6  24400    0    0   9000 76860 99600 120000  20400   15    0      1    0    0
##  smcol col age fsize marr
## 1     0  0  31     5     1
## 2     0  1  52     5     0
## 3     0  0  50     3     1
## 4     1  0  28     4     1
## 5     0  0  42     3     0
## 6     1  0  49     6     1
```

The variables in this dataset is defined as follows:

- tw: Total wealth (in US \$), which is defined as “net financial assets, including Individual Retirement Account (IRA) and 401(k) assets, plus housing equity plus the value of business, property, and motor vehicles.”
- ira: individual retirement account (IRA) balance (in US \$).
- e401: Binary variable, where 1 indicates eligibility for a 401(k)-retirement plan, and 0 indicates otherwise.
- nifa: Non-401k financial assets (in US \$).
- inc: Income (in US \$).
- hmort: Home mortgage (in US \$).
- hval: Home value (in US \$).
- hequity: Home value minus home mortgage.
- educ: Education (in years).
- male: Binary variable, where 1 indicates male and 0 indicates otherwise.
- twoearn: Binary variable, where 1 indicates two earners in the household, and 0 indicates otherwise.

- nohs, hs, smcol, col: Dummy variables for education levels - no high school, high school, some college, college.
- age: Age.
- fsize: Family size.
- marr: Binary variable, where 1 indicates married and 0 indicates otherwise.

```
colSums(is.na(data))
```

```
##      tw      ira    e401     nifa      inc    hmort      hval hequity      educ      male
##      0       0       0       0       0       0       0       0       0       0
## twoearn    nohs      hs    smcol      col      age    fsize      marr
##      0       0       0       0       0       0       0       0
```

```
any(duplicated(data))
```

```
## [1] FALSE
```

We can see that the data is in good shape, where categorical variables are already transformed into dummy variables. We can also see that there exists multi-collinearity in education levels (**nohs**, **hs**, **smcol**, **col**) and home-ownership-related variables (**hmort**, **hval**, and **hequity**).

```
summary(data)
```

```
##      tw      ira      e401      nifa
## Min.   :-502302 Min.    : 0 Min.   :0.0000 Min.    : 0
## 1st Qu.: 3246   1st Qu.: 0 1st Qu.:0.0000 1st Qu.: 200
## Median : 25225  Median : 0 Median :0.0000 Median : 1687
## Mean   : 63629  Mean   : 3471 Mean   :0.3714 Mean   : 13611
## 3rd Qu.: 82173  3rd Qu.: 0 3rd Qu.:1.0000 3rd Qu.: 8875
## Max.   :1887115 Max.   :100000 Max.   :1.0000 Max.   :1425115
##      inc      hmort      hval      hequity
## Min.    : -9   Min.    : 0 Min.    : 0 Min.    : -40000
## 1st Qu.: 19413 1st Qu.: 0 1st Qu.: 0 1st Qu.: 0
## Median : 31575 Median : 8000 Median : 50000 Median : 10000
## Mean   : 37177 Mean   : 30207 Mean   : 63965 Mean   : 33757
## 3rd Qu.: 48615 3rd Qu.: 52000 3rd Qu.: 95000 3rd Qu.: 48000
## Max.   :242124 Max.   :150000 Max.   :300000 Max.   :300000
##      educ      male      twoearn      nohs
## Min.    : 1.0   Min.    :0.0000 Min.    :0.0000 Min.    :0.0000
## 1st Qu.:12.0   1st Qu.:0.0000 1st Qu.:0.0000 1st Qu.:0.0000
## Median :12.0   Median :0.0000 Median :0.0000 Median :0.0000
## Mean   :13.2   Mean    :0.2018 Mean    :0.3808 Mean    :0.1277
## 3rd Qu.:15.0   3rd Qu.:0.0000 3rd Qu.:1.0000 3rd Qu.:0.0000
## Max.   :18.0   Max.    :1.0000 Max.    :1.0000 Max.    :1.0000
##      hs      smcol      col      age
## Min.    :0.0000 Min.    :0.0000 Min.    :0.0000 Min.    :25.00
## 1st Qu.:0.0000 1st Qu.:0.0000 1st Qu.:0.0000 1st Qu.:32.00
## Median :0.0000 Median :0.0000 Median :0.0000 Median :40.00
## Mean   :0.3819 Mean    :0.2422 Mean    :0.2482 Mean    :41.08
## 3rd Qu.:1.0000 3rd Qu.:0.0000 3rd Qu.:0.0000 3rd Qu.:48.00
## Max.   :1.0000 Max.    :1.0000 Max.    :1.0000 Max.    :64.00
##      fsize      marr
```

```
## Min.   : 1.00   Min.   :0.0000
## 1st Qu.: 2.00   1st Qu.:0.0000
## Median : 3.00   Median :1.0000
## Mean   : 2.87   Mean   :0.6075
## 3rd Qu.: 4.00   3rd Qu.:1.0000
## Max.   :13.00   Max.   :1.0000
```

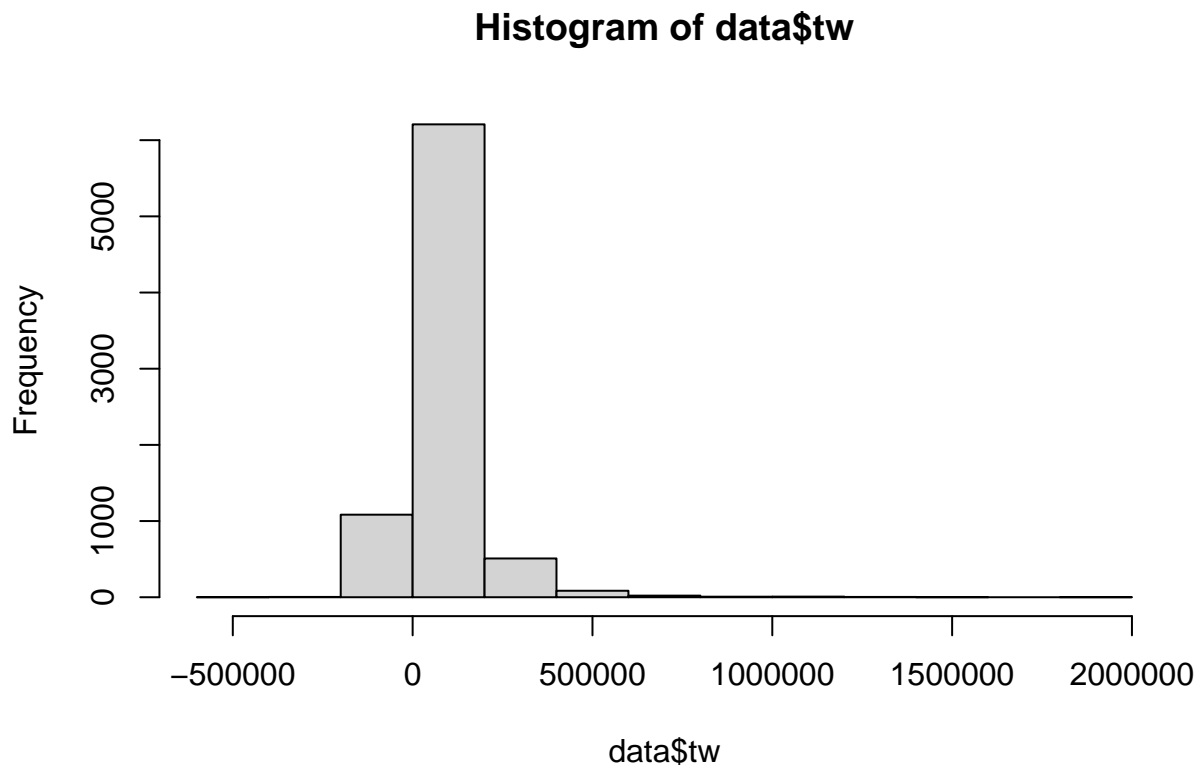
While there exist observations where total wealth is negative, it should be noted that the variable includes home equity, which can be negative, so it does not necessarily indicate that there are incorrect data entries.

The variables **ira**, **nohs**, **smcol**, **col**, and **male** exhibited a value of 0 at the 3rd quantile. They are probably a significant number of data points taking on the value of 0. Since **male** is on the list, it should also be noted that most observations are associated with female participants.

Also, the variable **tw**, **nifa**, **hmort**, and **hequity** have means that are much greater than medians, showing signs of large outliers.

In the histogram below, we can visualize the existence of outliers with enormous wealth.

```
hist(data$tw)
```



Using the graph, we can determine that removing the outliers with **tw** above \$1,000,000 would be appropriate.

```
data = subset(data, data$tw < 1000000)
```

Testing and Removing Multi-collinearity

Let's test whether removing different educational level predictors affect my model's performance, gauged by (MSPE). For simplicity sake, I did not use k-fold cross validation.

```
k <- 10
set.seed(123)
rand <- sample(nrow(data), floor(nrow(data)/k))
train <- setdiff(c(1:nrow(data)), rand)
y_rand <- data$tw[rand]

regnohs <- lm(tw ~ 1 + hs + smcol + col, data = data[train,])
reghs <- lm(tw ~ 1 + nohs + smcol + col, data = data[train,])
regsmcol <- lm(tw ~ 1 + nohs + hs + col, data = data[train,])
regcol <- lm(tw ~ 1 + nohs + hs + smcol, data = data[train,])

prnohs <- predict(regnohs, newdata = data[rand,])
prhs <- predict(reghs, newdata = data[rand,])
prsmcol <- predict(regsmcol, newdata = data[rand,])
prcol <- predict(regcol, newdata = data[rand,])

MSEnohs <- mean((y_rand-prnohs)^2)
MSEhs <- mean((y_rand-prhs)^2)
MSEsmcol <- mean((y_rand-prsmcol)^2)
MSEcol <- mean((y_rand-prcol)^2)

c(MSEnohs, MSEhs, MSEsmcol, MSEcol)
```

```
## [1] 9119936474 9119936474 9119936474 9119936474
```

No difference in performance is found between removing different terms for multi-collinearity. For interpretability, we choose to remove **hs** for education level.

More Feature Selections

Since **hequity** represents home value minus home mortgage, it is intuitively a better predictor of total wealth than **hval** or **hmort** itself. Hence, choosing **hequity** over **hval** and **hmort** is the more sensible choice.

Including years of education (**educ**) along with education levels is redundant. Considering that diplomas are usually much more important than years of education, prioritizing education level over years of education is appropriate.

```
data <- data[, !(names(data) %in% c("hs", "hval", "hmort", "educ"))]
```

Creating a Linear Baseline Model

Using Lasso and Forward/Backward Stepwise Selection

We will strive to create a linear baseline model. This will serve as a baseline to compare to when we later add nonlinear transformations and interaction terms.

For this approach, we are going to include all the features in the dataset. We will let the feature selection algorithms, Lasso and Stepwise Selection, to select the features for us.

For better accuracy, I employed 10-fold cross validation. Leave-one-out cross validation would yield a even more accurate result, but doing so on a dataset containing 7919 observations would take too much computational power.

```
source("KfoldCVFunctions.R")

mean_mspe_lasso <- compute_lasso_mspe(data, response_var = "tw")
mean_mspe_forward <- compute_forward_stepwise_mspe(data, response_var = "tw")
mean_mspe_backward <- compute_backward_stepwise_mspe(data, response_var = "tw")

# Print results
print(c(Lasso = mean_mspe_lasso,
        Forward_Stepwise = mean_mspe_forward,
        Backward_Stepwise = mean_mspe_backward))
```

```
##           Lasso  Forward_Stepwise  Backward_Stepwise
##      1427662119          1427849850          1427849850
```

As shown in the results above, Lasso yielded a lower MSPE than forward or backward stepwise selection.

Let's now inspect the coefficients that Lasso chose and their associated p-values.

Since Lasso performs both variable selection and shrinkage, leading to biased coefficient estimates, traditional significance tests for coefficients (like p-values) are not straightforwardly available. Therefore, we use the *hdi* (High Dimensional Inference) package to approximate the p-values.

```
library(hdi)

response_var <- "tw"
y <- data[[response_var]]
X <- as.matrix(data[, !(names(data) %in% response_var)])

lasso_cv <- cv.glmnet(X, y, alpha = 1)
best_lambda <- lasso_cv$lambda.min

lasso_model <- glmnet(X, y, lambda = best_lambda, alpha = 1)

lasso_inference <- hdi::lasso.proj(X, y)

print(coef(lasso_model))
```

```
## 14 x 1 sparse Matrix of class "dgCMatrix"
##              s0
## (Intercept) -1.576371e+04
## ira         1.606418e+00
## e401        7.935739e+03
## nifa        1.102051e+00
## inc         2.500733e-01
## hequity     1.081333e+00
## male        3.169643e+03
## twoearn     -5.386380e+03
```

```
## nohs      .
## smcol     1.006324e+03
## col       .
## age       2.470886e+02
## fsize     .
## marr      1.473061e+03
```

As shown above, Lasso has selected all of the features except nohs, col, and fsize. The coefficients are shown above.

```
print(lasso_inference$pval)
```

```
##          ira          e401          nifa          inc          hequity
## 5.152649e-198 1.807363e-17 0.000000e+00 1.982564e-23 0.000000e+00
##          male          twoearn          nohs          smcol          col
## 1.683754e-03 1.802943e-07 8.852471e-01 2.187133e-01 8.723161e-01
##          age          fsize          marr
## 1.172631e-07 7.921987e-01 5.334928e-02
```

Above are the approximated p-values the coefficients. We can gauge how strong of a predictor each feature is. As expected, ira and e401 have a really small p-value as they are literally a part of **tw**.

- tw: Total wealth (in US \$), which is defined as “net financial assets, including Individual Retirement Account (IRA) and 401(k) assets, plus housing equity plus the value of business, property, and motor vehicles.”

Let's also compute the MSPE for a simple OLS regression model and for a ridge regression model with the selected features for comparison.

```
data_subset <- data[, !(names(data) %in% c("nohs", "col", "fsize"))]

source("KfoldCVFunctions.R")
mean_mspe_ols <- compute_ols_mspe(data_subset, response_var = "tw")
mean_mspe_ridge <- compute_ridge_mspe(data, response_var = "tw")

print(c(
  OLS = mean_mspe_ols,
  Ridge = mean_mspe_ridge,
  Lasso = mean_mspe_lasso,
  Forward_Stepwise = mean_mspe_forward,
  Backward_Stepwise = mean_mspe_backward
))
```

```
##          OLS          Ridge          Lasso  Forward_Stepwise
## 1427051036 1450228711 1427662119 1427849850
## Backward_Stepwise
## 1427849850
```

Surprisingly, a simple OLS regression on selected features yielded better results than ridge regression, Lasso, forward/backward stepwise selection.

The mean MSPE of our OLS regression will then serve as our benchmark, which we will strive to improve upon when fitting nonlinear transformations.

Finding Nonlinear Relationships and Applying Transformations

Inspecting the Relationships between `tw` and Features

After creating an appropriate linear model, the appropriate next step to improve predictive performance is through applying nonlinear transformations.

Let's first inspect the relationships between `tw` and all the quantitative (non-binary) features in the dataset. Nonlinear transformations are inappropriate for binary features because they only have two distinct values, making such transformations ineffective and potentially meaningless.

The below function compiles all the scatterplots into a list of *ggplot* objects.

```
library(tidyverse)
library(gridExtra)

plot_scatterplots <- function(data, response_var, feature_names) {

  plot_list <- list()

  for (feature in feature_names) {
    p <- ggplot(data = data) +
      geom_point(mapping = aes(x = !!sym(feature), y = !!sym(response_var)), alpha = 0.5) +
      labs(x = feature, y = response_var) +
      theme_minimal() +
      theme(aspect.ratio = 1)
    plot_list[[feature]] <- p
  }

  return(plot_list)
}
```

We can then call the function and use the *gridExtra* package to arrange them.

```
plots <- plot_scatterplots(data_subset, response_var = "tw", c("ira", "nifa", "inc", "hequity", "age"))
do.call(grid.arrange, c(plots, ncol = 3))
```

