

Human Freedom Index and Suicides Project

Part 2: Data Exploration with HFI Dataset

Doug Cady

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Contents

Load dataset into R	1
Explore HFI dataset	1

R version 4.1.1 “Kick Things”

```
library(readr)
library(dplyr)
library(tidyr)
library(stringr)
library(ggplot2)
library(GGally)
```

Load dataset into R

```
hfi <- read_csv("../data/hfi_cc_2018.csv")
```

Explore HFI dataset

Questions to answer:

- Are the freedom score, number of suicides per 100k, and gdp per capita related?
 - Find correlations, make 3 scatter pair plots
- Does population size impact freedom?
- Does female freedom scores relate to female suicide rates?
- What are important factors that contribute to human freedom?
 - Linear regression model

Hypotheses:

1. Countries with high female freedom scores will have less than average female suicide rates

Ho: mean suicide rates lo female freedom \geq average female suicide rate

Ha: mean suicide rates hi female freedom $<$ average female suicide rate

- pf_ss_women_fgm - Female genital mutilation
- pf_ss_women_inheritance - Equal inheritance rights for widows and daughters
- pf_movement_women - Freedom of movement for women
- pf_identity_sex_female - Female to female relationships
- pf_identity_divorce - Divorce
- Perhaps educational opportunity or abortion access should be included here as well?

2. Countries with smaller populations will have more freedom

Ho: mean freedom small population \leq mean freedom large population

Ha: mean freedom small population $>$ mean freedom large population

- `hf_score` - Overall Human Freedom Index score
- `population` - (suicides dataset) country population

Column Descriptions

There are many, many columns in this dataset, so let us focus on a few columns

- 1 time-series variable
 - `year`
- 3 categorical variables
 - `country`
 - `ISO_code` (country abbreviation)
 - `region`
- 119 numeric (narrowed down to only a handful)
 - Freedom indicators and their aggregate category scores, like
 - * Female genital mutilation and inheritance rights (widows, daughters) make up 3 columns and are aggregated into one category score - Female security and safety

```
categ_cols <- c(
  "year",
  "country",
  "region"
)

focus_cols <- c(
  "pf_ss_women_fgm",
  "pf_ss_women_inheritance",
  "pf_movement_women",
  "pf_identity_sex_female",
  "pf_identity_divorce",
  "pf_score",
  "ef_score",
  "hf_score"
)

# Look at focus subset of columns and update categoricals to factor type
hfi_focus <- rename(hfi, country = countries) %>%
  mutate(country = factor(country),
         region = factor(region))
hfi_focus <- hfi_focus[, str_c(c(categ_cols, focus_cols))]
write_csv(hfi_focus, "../data/clean_hfi_2018.csv")

print(str(hfi_focus))

## tibble [1,458 x 11] (S3: tbl_df/tbl/data.frame)
##   $ year                : num [1:1458] 2016 2016 2016 2016 2016 ...
##   $ country              : Factor w/ 162 levels "Albania","Algeria",...: 1 2 3 4 5 6 7 8 9 10 ...
##   $ region               : Factor w/ 10 levels "Caucasus & Central Asia",...: 3 5 9 4 1 7 10 1 4 5 .
##   $ pf_ss_women_fgm      : num [1:1458] 10 10 10 10 10 10 10 10 10 NA 10 ...
##   $ pf_ss_women_inheritance: num [1:1458] 5 0 5 10 10 10 10 7.5 NA 0 ...
```

```
## $ pf_movement_women      : num [1:1458] 5 5 10 10 10 10 10 5 NA 5 ...
## $ pf_identity_sex_female : num [1:1458] 10 0 0 10 10 10 10 10 10 10 ...
## $ pf_identity_divorce    : num [1:1458] 5 0 10 10 5 10 10 5 NA 0 ...
## $ pf_score               : num [1:1458] 7.6 5.28 6.11 8.1 6.91 ...
## $ ef_score               : num [1:1458] 7.54 4.99 5.17 4.84 7.57 7.98 7.58 6.49 7.34 7.56 ...
## $ hf_score               : num [1:1458] 7.57 5.14 5.64 6.47 7.24 ...
## NULL
```

Head / Tail of Data

Nothing out of the ordinary here. It seems like we read in the whole file and do not need to skip any header or footer miscellaneous data.

```
print(head(hfi_focus))
```

```
## # A tibble: 6 x 11
##   year country region      pf_ss_women_fgm pf_ss_women_inhe~ pf_movement_wom~
##   <dbl> <fct>   <fct>          <dbl>          <dbl>          <dbl>
## 1  2016 Albania Eastern Eur~         10             5             5
## 2  2016 Algeria Middle East~         10             0             5
## 3  2016 Angola Sub-Saharan~         10             5            10
## 4  2016 Argenti~ Latin Ameri~         10            10            10
## 5  2016 Armenia Caucasus & ~         10            10            10
## 6  2016 Austral~ Oceania          10            10            10
## # ... with 5 more variables: pf_identity_sex_female <dbl>,
## #   pf_identity_divorce <dbl>, pf_score <dbl>, ef_score <dbl>, hf_score <dbl>
```

```
print(tail(hfi_focus))
```

```
## # A tibble: 6 x 11
##   year country region      pf_ss_women_fgm pf_ss_women_inhe~ pf_movement_wom~
##   <dbl> <fct>   <fct>          <dbl>          <dbl>          <dbl>
## 1  2008 Uruguay Latin Ameri~         10            10            10
## 2  2008 Venezue~ Latin Ameri~         10            10            10
## 3  2008 Vietnam South Asia          10            10            10
## 4  2008 Yemen, ~ Middle East~         NA            NA            NA
## 5  2008 Zambia Sub-Saharan~         10             0            10
## 6  2008 Zimbabwe Sub-Saharan~         9.5             5             5
## # ... with 5 more variables: pf_identity_sex_female <dbl>,
## #   pf_identity_divorce <dbl>, pf_score <dbl>, ef_score <dbl>, hf_score <dbl>
```

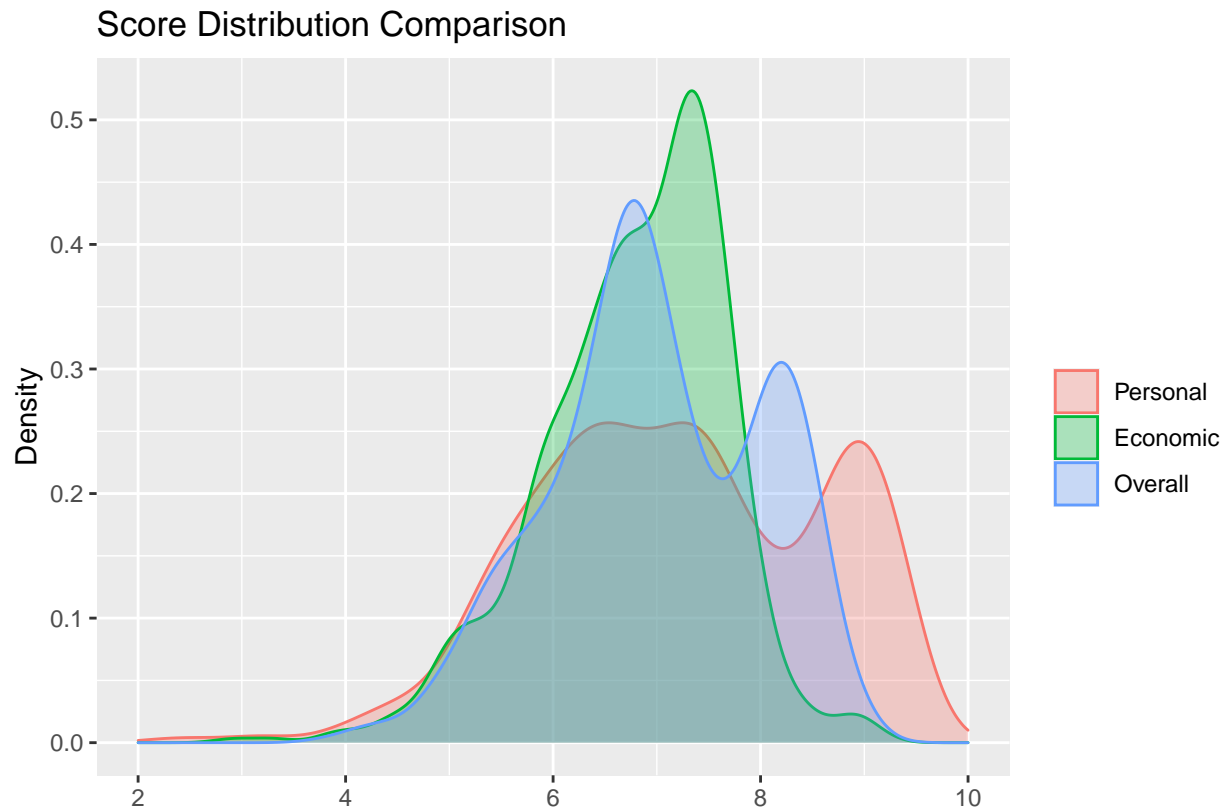
Distribution of Scores - Personal, Economic, Overall Human Freedom

```
score_cols <- c("pf_score", "ef_score", "hf_score")
scores <- hfi_focus %>%
  pivot_longer(cols = score_cols, names_to = "freedom", values_to = "score")
scores <- scores[, str_c(c(categ_cols, "freedom", "score"))]

scores$freedom <- recode_factor(scores$freedom, "pf_score" = "Personal",
                                "ef_score" = "Economic",
                                "hf_score" = "Overall")

# Freedom distribution
ggplot(scores) +
  geom_density(aes(x = score, color = freedom, fill = freedom), alpha = 0.3) +
```

```
labs(x = '', y = 'Density', title = 'Score Distribution Comparison', color = '', fill = '') +
scale_x_continuous(limits = c(2, 10))
```



Personal freedom scores are more spread out than economic or overall freedom scores. Very few countries have scores below 4. The majority of economic scores fall between 6 and 8, lower than many countries' personal freedom scores.

Women Freedom

```
women_freedom_cols <- c(
  "pf_ss_women_fgm",
  "pf_ss_women_inheritance",
  "pf_movement_women",
  "pf_identity_sex_female",
  "pf_identity_divorce")

women_freedom_init <- hfi_focus[, str_c(c(categ_cols, women_freedom_cols))]
summary(women_freedom_init)
```

##	year	country	region
##	Min. :2008	Albania : 9	Sub-Saharan Africa :378
##	1st Qu.:2010	Algeria : 9	Latin America & the Caribbean:234
##	Median :2012	Angola : 9	Eastern Europe :198
##	Mean :2012	Argentina: 9	Middle East & North Africa :171
##	3rd Qu.:2014	Armenia : 9	Western Europe :162
##	Max. :2016	Australia: 9	South Asia :153
##		(Other) :1404	(Other) :162
##	pf_ss_women_fgm pf_ss_women_inheritance pf_movement_women		

```
## Min. : 0.40 Min. : 0.00 Min. : 0.00
## 1st Qu.: 9.60 1st Qu.: 5.00 1st Qu.: 5.00
## Median :10.00 Median : 5.00 Median :10.00
## Mean : 9.24 Mean : 6.64 Mean : 8.04
## 3rd Qu.:10.00 3rd Qu.:10.00 3rd Qu.:10.00
## Max. :10.00 Max. :10.00 Max. :10.00
## NA's :172 NA's :119 NA's :141
## pf_identity_sex_female pf_identity_divorce
## Min. : 0.00 Min. : 0.0
## 1st Qu.:10.00 1st Qu.: 5.0
## Median :10.00 Median :10.0
## Mean : 7.94 Mean : 7.5
## 3rd Qu.:10.00 3rd Qu.:10.0
## Max. :10.00 Max. :10.0
## NA's :80 NA's :873
```

Missing values

The 5 women freedom variables have at least 5% missing values with Divorce having almost 60%:

- pf_ss_women_fgm - Female genital mutilation - 11.8% missing
- pf_ss_women_inheritance - Female inheritance - 8.2%
- pf_movement_women - Women movement - 9.7%
- pf_identity_sex_female - Female to female relationships - 5.5%
- pf_identity_divorce - Divorce - 59.9%

```
sapply(women_freedom_init,
       function(x) paste0(round(sum(is.na(x)) / nrow(women_freedom_init) * 100, 1), "%"))
```

```
##          year          country          region
##          "0%"          "0%"          "0%"
## pf_ss_women_fgm pf_ss_women_inheritance pf_movement_women
##          "11.8%"          "8.2%"          "9.7%"
## pf_identity_sex_female pf_identity_divorce
##          "5.5%"          "59.9%"
```

```
# What to do about Divorce missing values? Impute? Interpolate? Remove?
# Instead, we can average the 5 women freedom columns together and then remove the columns
# with NAs - equal to the minimum NA% from the individual columns (5.5%)
women_freedom_init$avg_women_score <- rowMeans(select(women_freedom_init, women_freedom_cols),
                                              na.rm = TRUE)

women_freedom <- women_freedom_init[!is.na(women_freedom_init$avg_women_score),
                                   c("year", "country", "region", "avg_women_score")]

ggplot(women_freedom) + geom_histogram(aes(x = avg_women_score))
```

