## Replication: The Structure of Inequality and the Politics of Redistribution

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## Design declaration

First we start by loading in the DeclareDesign package and defining the elements of the design.

- declare\_population refers to the sample size of the study. The study concerns country-year units. In this case, there are 858 observations.
- delcare\_potential\_oucomes refers to

```
library('DeclareDesign')

## Loading required package: randomizr

## Loading required package: fabricatr

## Loading required package: estimatr

design <-
    declare_population(N = 858) +
    declare_potential_outcomes() +
    declare_estimand() +
    declare_assignment() +
    declare_estimator()</pre>
```

## Replication

First we open the dataset with the haven package, which allows us to open .dta files.

```
library('haven')
directory <- "/Users/juliangerez/Google Drive/Semester_Fall 2018/Political Economy of Development/Lupu-
data <- read_dta(paste0(directory, "LupPon_APSR.dta"))</pre>
```

Next, the authors define invert disproportionality measures, disp\_gall as such:

```
data$disp_gall <- data$disp_gall*-1
```

Then the variables female participation, fempar, and annual net union density, union so that they are rescaled:

```
data$fempar <- data$fempar*100
data$union <- data$union*100</pre>
```

Then the variables pjoint and disp\_gall, which are partial and disproportionality, respectively are standardized from [0,1]. To do so, we are defining a function, range01, which standardizes the range of a variable such that it takes on values from 0 to 1.

```
range01 <- function(x){(x-min(x))/(max(x)-min(x))}
```

```
data$stdpjoint <- range01(data$pjoint)</pre>
data$stdpdisp_gall <- range01(data$disp_gall)</pre>
Next, we interpolate missing values:
library('zoo')
##
## Attaching package: 'zoo'
## The following objects are masked from 'package:base':
##
##
       as.Date, as.Date.numeric
data$pratio9050 <- NA
data_countries <- lapply(unique(data$country), function(x)</pre>
  subset(data, data$country==x)
for (i in 1:length(data_countries)){
data_countries[[i]][,24] <- na.approx(data_countries[[i]][,5], x = index(data_countries[[i]][,3], data_
}
data <- do.call("rbind", data_countries)</pre>
data$pratio5010 <- NA
data_countries <- lapply(unique(data$country), function(x)</pre>
  subset(data, data$country==x)
for (i in 1:length(data_countries)){
data_countries[[i]][,25] <- na.approx(data_countries[[i]][,6], x = index(data_countries[[i]][,3], data_
data <- do.call("rbind", data_countries)</pre>
data$pratio9050s <- NA
data_countries <- lapply(unique(data$country), function(x)</pre>
  subset(data, data$country==x)
for (i in 1:length(data_countries)){
data_countries[[i]][,26] <- na.approx(data_countries[[i]][,7], x = index(data_countries[[i]][,3], data_
data <- do.call("rbind", data_countries)</pre>
data$pratio5010s <- NA
data_countries <- lapply(unique(data$country), function(x)</pre>
  subset(data, data$country==x)
)
```

```
for (i in 1:length(data_countries)){
data_countries[[i]][,27] <- na.approx(data_countries[[i]][,8], x = index(data_countries[[i]][,3], data_
data <- do.call("rbind", data_countries)</pre>
data$pforeign <- NA
data_countries <- lapply(unique(data$country), function(x)</pre>
  subset(data, data$country==x)
for (i in 1:length(data_countries)){
data_countries[[i]][,28] <- na.approx(data_countries[[i]][,16], x = index(data_countries[[i]][,3], data
data <- do.call("rbind", data_countries)</pre>
data$pvoc <- NA
data_countries <- lapply(unique(data$country), function(x)</pre>
  subset(data, data$country==x)
)
for (i in 1:length(data_countries)){
data_countries[[i]][,29] <- na.approx(data_countries[[i]][,19], x = index(data_countries[[i]][,3], data
}
data <- do.call("rbind", data_countries)</pre>
We generate an immigration measure, pforeign which reflects the percentage of the population that is
foreign-born:
data$pforeign <- data$pforeign*1000</pre>
data$fpop <- data$pforeign/data$pop</pre>
Generate inequality measures
data$ratio9010 <- data$pratio9050*data$pratio5010
data$ratio9010s <- data$pratio9050s*data$pratio5010s</pre>
data$skew <- data$pratio9050/data$pratio5010</pre>
```

data\$skews <- data\$pratio9050s/data\$pratio5010s</pre>

Data cleaning

Redistribution models

Social spending models

Immigration

Partisanship

Redistribution and social spending with partisanship

Robustness checks via design modification

Extension