### Sample\_Project

Code ▼

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This document is a minimal, reproducible sample of my empirical workflow Full code and additional projects: **github.com/tuffyli** (https://github.com/tuffyli)

### 1 Objectives

The main objective of this sample work is to explore the expansion of Brazil's alimony rights to women in stable unions and its effects on education, particularly among young women who became newly entitled to this right. This analysis is inspired by Rangel's (2006) seminal work on the policy's expansion. The main difference in this study is the application of an additional Propensity Score Matching (PSM) method, with a particular focus on young mothers and their educational decisions.

#### 2 Data

In the repository (https://github.com/Tuffyli/Personal\_Projects/tree/main/PNAD-exercise), I include supplementary code detailing the selection of key variables.

Due to the large size of the PNAD dataset, it is not possible to make the full data publicly available. Therefore, I provide a filtered version of the dataset containing only the variables relevant to the analysis.

```
# ----- #
# 1. Data Extraction ----

data <- readRDS("C:/Users/tuffy/Documents/Trabalhos/Ava_Pol/Bases/final_filtered_9295.rds")

#1.1 Summary ----
#' In the data manipulation code I created a summary table. Here I present it

summary <- readRDS("C:/Users/tuffy/Documents/Trabalhos/Ava_Pol/Bases/summary.rds")
print(summary)
```

##		Mean	SD	Min	Max	
##	Treatment	0.45	0.50	0	1	
##	Female = 1	1.00	0.00	1	1	
##	<pre>Race (White or Asian = 1)</pre>	0.55	0.50	0	1	
##	Highest Education	4.22	0.56	0	9	
##	Age	21.04	2.41	15	24	
##	Years of education	6.61	3.37	1	17	
##	Last grade concluded	4.47	2.00	1	9	
##	Course enrollment	2.48	1.65	0	9	
##	Household per capita wage	173221.66	269756.48	0	9549387	
##	Pension (yes $=$ 1)	0.00	0.05	0	1	
##	CBO Group	4.34	2.58	0	8	
##	Male child (house)	0.57	0.72	0	12	
##	Female child (house)	0.55	0.71	0	6	
##	Total childs	1.17	1.00	0	13	

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rm(summary)

Treated individuals are women aged 15–24 in non-civil unions (consensual or religious-only) who are household heads or spouses; controls are otherwise similar women in legally recognized unions (civil or civil + religious).

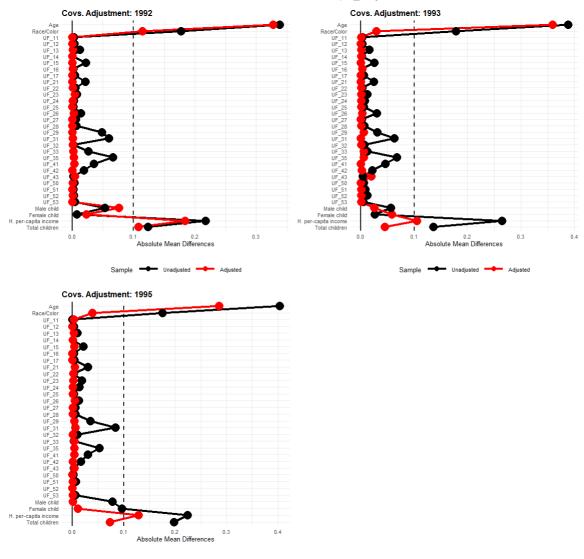
# 3 Propensity Score Matching

In this next step I will execute the propensity score matching between treated and control females.

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```
# 2. PSM balance
# -----
plots <- list()</pre>
for (year in c(1992, 1993, 1995)) {
  df_psm <- data %>%
    filter(ano == year) %>%
   select(
      treatment, uf, age, cor, anos_estudo, peso_pessoa,
      ultima_serie_concluida, tipo_curso_frequenta, grupo_cbo,
      trabalhou_ultimo_ano, renda_dom_per_capita,
      dummy_filhos_homens_dom, dummy_filhos_mulheres_dom,
      pensao_dummy, total_filhos
    ) %>%
   mutate(uf = as.factor(uf)) %>%
   filter(
      !is.na(treatment), !is.na(uf), !is.na(cor), !is.na(peso_pessoa),
      !is.na(dummy_filhos_homens_dom), !is.na(dummy_filhos_mulheres_dom),
      !is.na(renda_dom_per_capita), !is.na(total_filhos)
    )
  match_model <- matchit(</pre>
   treatment ~ age + cor + uf + dummy_filhos_homens_dom +
      dummy_filhos_mulheres_dom + renda_dom_per_capita + total_filhos,
   data = df_psm,
    s.weights = df_psm$peso_pessoa,
   caliper = 0.05,
   method = "nearest",
   replace = TRUE
  # Show the match summary counts in the document
  sum <- summary(match_model)</pre>
  latex df <- as.data.frame(sum$nn)</pre>
  latex_table <- kable(latex_df,</pre>
                       format = "latex",
                       booktabs = TRUE,
                       caption = "Estatísticas descritivas ponderadas",
                       align = "lccc")
  #The files are availabre in the repository
  writeLines(latex_table,
             paste0("C:/Users/tuffy/Documents/Trabalhos/", year," match summary.tex"))
  # Love plot (INLINE)
  custom labels <- c(</pre>
    distance_weighted
                              = "Distance (weighted)",
   uf
                              = "UF",
                              = "Race/Color",
    cor
    dummy_filhos_homens_dom
                             = "Male child",
    dummy_filhos_mulheres_dom = "Female child",
                              = "Age",
    age
                              = "H. per-capita income",
    renda_dom_per_capita
```

```
= "Total children"
    total_filhos
  )
  love <- love.plot(</pre>
    match_model,
                  = "mean.diffs",
    stats
    abs
                  = TRUE,
                 = 0.10,
    threshold
    colors
                  = c("black", "red"),
    line
                  = TRUE,
                  = custom_labels,
    var.names
    drop.distance = TRUE
  ) +
    labs(
      title = paste0("Covs. Adjustment: ",year)
   theme_minimal(base_size = 5) +
    theme(plot.title = element_text(face = "bold"),
                   legend.position = "bottom")
  #Grouping the plots
  plots[[as.character(year)]] <- love</pre>
  matched_data <- match.data(match_model) %>%
    mutate(ano = year)
  assign(paste0("df_matched_", year), matched_data)
  rm(year, custom_labels, sum, ess_table, latex_df, latex_table, love)
}
       <- cowplot::plot_grid(plots[["1992"]], plots[["1993"]],</pre>
top
                              ncol = 2, align = "hv", rel_widths = c(1, 1))
bottom <- cowplot::plot_grid(plots[["1995"]],</pre>
                              ncol = 2, align = "hv", rel_widths = c(1, 1)) # legend stays he
         re
combined <- cowplot::plot_grid(top, bottom, ncol = 1, rel_heights = c(1, 1.05))</pre>
combined
```



# 4 Regression

To assess treatment effects, we fit progressively richer models (no FE, UF×year FE, and FE + controls) using the matched sample.

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```
# ----- #
# 3. Regression -----
# ------ #
# Bind matched samples (1992, 1993, 1995) and prepare identifiers
df_psm <- bind_rows(df_matched_1992, df_matched_1993, df_matched_1995)</pre>
df psm$ano
           <- factor(df_psm$ano)
                                      # ensure year is a factor
           <- relevel(df_psm$ano, ref = "1992")</pre>
df psm$ano
                                              # reference year = 1992
df_psm$uf_ano <- interaction(df_psm$uf, df_psm$ano, sep = "_") # UF x year FE key</pre>
# --- Model 1: no fixed effects (year-to-year comparison via i())
model_nc <- feols(</pre>
 anos_estudo ~ treatment + i(ano, treatment, ref = "1992"),
 data = df_psm, weights = df_psm$weights, vcov = "hetero"
# --- Model 2: add UF×year fixed effects (absorbs geography-by-year shocks)
model_fe <- feols(</pre>
 anos_estudo ~ treatment + i(ano, treatment, ref = "1992") | uf_ano,
 data = df_psm, weights = df_psm$weights, vcov = "hetero"
# --- Model 3: UF×year FE + basic controls
model_cc <- feols(</pre>
 anos_estudo ~ treatment + i(ano, treatment, ref = "1992") +
   total_filhos + cor + age | uf_ano,
 data = df_psm, weights = df_psm$weights, vcov = "hetero"
)
```

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```
tab <- fixest::etable(</pre>
  model_nc, model_fe, model_cc,
                                          # pass models directly
  drop
            = "Constant",
  dict = c(
    "anos_estudo"
                                     = "Years of education",
    "treatment$"
                                     = "Treatment (1992)",
    "i\\(ano, treatment.*\\)::1993" = "Treatment × 1993",
    "i\\(ano, treatment.*\\)::1995" = "Treatment \times 1995",
    "age"
                                     = "Age",
    "cor"
                                     = "Race/Color",
    "total filhos"
                                    = "Total children",
    "treatment x ano = 1993"
                                    = "Treatment \times 1993",
    "treatment x ano = 1995"
                                    = "Treatment × 1995"
  ),
  se.below = TRUE,
  signif.code = "letters",
 digits
           = 3,
 fitstat = c("n","r2","rmse"),
           = "ATT by Year (ref = 1992)",
 title
  notes = "SEs are heteroskedastic-robust. Weights from matching sample."
  # <- DO NOT set tex=TRUE here (default is HTML/markdown-friendly)</pre>
knitr::kable(tab, format = "html", escape = FALSE) |>
  kableExtra::kable_styling(full_width = FALSE)
```

	model_nc	model_fe	model_cc
Dependent Var.:	Years of education	Years of education	Years of education
treatment	-1.77a	-1.66a	-1.25a
	(0.077)	(0.100)	(0.095)
treatment x ano = 1993	0.307a	0.230	0.112
	(0.080)	(0.146)	(0.136)
treatment x ano = 1995	0.452a	0.176	0.055
	(0.077)	(0.147)	(0.137)
Total children			-0.915a
			(0.028)
Race/Color			0.803a
			(0.057)
Age			0.260a
			(0.011)

	model_nc	model_fe	model_cc
Fixed-Effects:			
uf_ano	No	Yes	Yes
S.E. type	Heteroskedasrob.	Heteroskedasrob.	Heteroskedasrob.
Observations	17,779	17,779	17,779
R2	0.05238	0.11600	0.21156

There is no statistically significant effect of alimony eligibility on the years of education among young women. However, the low R² values indicate limited explanatory power, these specifications explain only a small share of the variation in schooling outcomes, so the results should be interpreted with caution.