Colombian trans wellbeing

Code and analyses

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Description

This document contains all code, and step by step explanations for all analyses, figures and tables (including supplementary figures and tables) for:

Reyes-Rodríguez, M. F., & Leongómez, J. D. (in prep). Colombian trans wellbeing

Data are available on the Open Science Framework (OSF): https://doi.org/10.XXXX/OSF.IO/XXXXX. The analyses were designed by Maria Fernanda Reyes-Rodríguez and Juan David Leongómez. This document and its underlying code were created in R Markdown by Juan David Leongómez using R and LaTeX, ensuring full reproducibility.

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1 Preliminaries

1.1 Load packages

This file was created using knitr (Xie, 2014), mostly using tidyverse (Wickham et al., 2019) syntax. As such, data wrangling was mainly done using packages such as dplyr (Wickham et al., 2023), and most figures were created or modified using ggplot2 (Wickham, 2016). Tables were created using knitr::kable and kableExtra (Zhu, 2021).

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Multi-model inference and model averaging was achieved using MuMIn (Bartoń, 2024), and model assumptions were performed using performance (Lüdecke et al., 2021).

All packages used in this file can be directly installed from the Comprehensive R Archive Network (CRAN). For a complete list of packages used to create this file, and their versions, see section 2, at the end of the document.

```
library(ltm)
library(psych)  # For statistical functions (e.g., Cronbach's alpha)
library(MuMIn)  # For model selection and averaging
library(performance)  # For model performance metrics
library(readr)  # For reading data files
library(scales)  # For percent formatting
library(knitr)
library(kableExtra)
library(tidyverse)  # For data manipulation and piping
```

1.2 Load data

Load raw CSV data

```
data_RAW <- read_csv("data/data.csv")
```

1.2.1 Define PANAS Subscales (Positive & Negative Affect)

XXXXXXX

1.3 Internal consistency

1.3.1 Calculate Cronbach's Alpha for Different Scales

To measure the internal consistency of these tests, we used standardized Cronbach's alpha (α or Tau-equivalent reliability: ρ_T) coefficients, using the function cronbach.alpha from the package ltm (Rizopoulos, 2006).

```
# Compute Cronbach's alpha for the Self-Efficacy (EAG) scale
alpha_EAG <- data_RAW |>
    mutate(across(where(is.numeric), ~ na_if(., 99))) |> # Replace 99 with NA (missing values)
    select(starts_with("EAG_")) |> # Select all columns starting with "EAG_"
    drop_na() |>
        cronbach.alpha(CI = TRUE, standardized = TRUE) # Compute Cronbach's alpha

# Compute Cronbach's alpha for the Life-Satisfaction (SWLS) scale
alpha_SWLS <- data_RAW |>
    mutate(across(where(is.numeric), ~ na_if(., 99))) |> # Replace 99 with NA
    select(starts_with("SWLS_")) |> # Select all columns starting with "SWLS_"
    drop_na() |>
        cronbach.alpha(CI = TRUE, standardized = TRUE)

# Compute Cronbach's alpha for the Resilience (EBR) scale
alpha_EBR <- data_RAW |>
```

```
mutate(across(where(is.numeric), ~ na_if(., 99))) |> # Replace 99 with NA
 select(starts_with("EBR_")) |> # Select all columns starting with "EBR_"
 drop_na() |>
  cronbach.alpha(CI = TRUE, standardized = TRUE)
alpha EBD <- data RAW |>
 mutate(across(where(is.numeric), ~ na_if(., 99))) |> # Replace 99 with NA
 select(starts_with("EBD_")) |> # Select all columns starting with "EBD_"
 mutate(across(everything(), ~ ifelse(is.na(.x), NA, .x - 1))) |> # Adjust values
 drop_na() |>
 cronbach.alpha(CI = TRUE, standardized = TRUE)
alpha_MOS2 <- data_RAW |>
 mutate(across(where(is.numeric), ~ na_if(., 99))) |> # Replace 99 with NA
 select(starts_with("MOS2_")) |> # Select all columns starting with "MOS2_"
 drop na() |>
 cronbach.alpha(CI = TRUE, standardized = TRUE)
alpha_PANAS_P <- data_RAW |>
 mutate(across(where(is.numeric), ~ na_if(., 99))) |> # Replace 99 with NA
 select(all_of(PANAS_P)) |> # Select PANAS_P variables
 drop_na() |>
 cronbach.alpha(CI = TRUE, standardized = TRUE)
alpha_PANAS_N <- data_RAW |>
 mutate(across(where(is.numeric), ~ na_if(., 99))) |> # Replace 99 with NA
 select(all_of(PANAS_N)) |> # Select PANAS_N variables
 drop na() |>
 cronbach.alpha(CI = TRUE, standardized = TRUE)
# Compute Cronbach's alpha for Community Cohesion (PCPS3) scale
alpha PCPS3 <- data RAW |>
 mutate(across(where(is.numeric), ~ na_if(., 99))) |> # Replace 99 with NA
 select(starts_with("PCPS3_")) |> # Select all columns starting with "PCPS3_"
 drop_na() |>
 cronbach.alpha(CI = TRUE, standardized = TRUE)
```

1.3.2 Table S1. Internal consistency of measured scales

The internal consistency of the measured scales was generally strong, with Cronbach's α values ranging from 0.767 to 0.977. In particular, the Social Support (MOS2) and Self-Efficacy (EAG) scales exhibited excellent internal consistency, while the Depression (EBD) and Community Cohesion (PCPS3) scales had acceptable reliability, suggesting a slightly lower but still adequate level of internal consistency.

```
"Community Cohesion$^x$"),
p = c(alpha_EAG$p, alpha_SWLS$p, alpha_EBR$p, alpha_EBD$p, alpha_MOS2$p,
      alpha_PANAS_P$p, alpha_PANAS_N$p, alpha_PCPS3$p),
n = c(alpha_EAG$n, alpha_SWLS$n, alpha_EBR$n, alpha_EBD$n, alpha_MOS2$n,
      alpha_PANAS_P$n, alpha_PANAS_N$n, alpha_PCPS3$n),
alpha = c(alpha_EAG$alpha, alpha_SWLS$alpha, alpha_EBR$alpha, alpha_EBD$alpha,
          alpha MOS2$alpha, alpha PANAS P$alpha, alpha PANAS N$alpha, alpha PCPS3$alpha),
ci2.5 = c(alpha_EAG$ci[1], alpha_SWLS$ci[1], alpha_EBR$ci[1], alpha_EBD$ci[1],
          alpha_MOS2$ci[1], alpha_PANAS_P$ci[1], alpha_PANAS_N$ci[1], alpha_PCPS3$ci[1]),
ci97.5 = c(alpha_EAG$ci[2], alpha_SWLS$ci[2], alpha_EBR$ci[2], alpha_EBD$ci[2],
           alpha_MOS2$ci[2], alpha_PANAS_P$ci[2], alpha_PANAS_N$ci[2], alpha_PCPS3$ci[2])) |>
mutate(across(starts_with("ci"), round, 3)) |>
unite(col = "CI", ci2.5:ci97.5, sep = " - ") |>
kable(digits = 3,
      booktabs = TRUE,
      linesep = "",
      align = c("l", rep("c", 4)),
      caption = "Internal consistency of measured scales",
      col.names = c("Variable", "Items", "$n$", "$\\alpha$", "$95\\% CI$"),
      escape = FALSE) |>
kable_styling(latex_options = "HOLD_position") |>
footnote(
  general = "95\\\% confidence intervals were calculated with 1,000 bootstrap samples.
         Standardized Cronbach's alpha ($\\\alpha$) coefficients were computed.
         $^1$\\\\cite{EAG};
         $^2$\\\cite{EBR};
         $^3$\\\\cite{EBD};
         $^4$\\\cite{MOS};
         $^5$\\\\cite{PANAS};
         $^6$\\\\cite{PANAS}",
  threeparttable = TRUE, footnote_as_chunk = TRUE, escape = FALSE
```

Table S1. Internal consistency of measured scales

Variable	Items	n	α	95%CI
Self-Efficacy ¹	10	223	0.905	0.875 - 0.926
Life-Satisfaction ²	5	253	0.869	0.841 - 0.895
$Resilience^3$	4	278	0.861	0.821 - 0.892
$Depression^4$	7	223	0.767	0.71 - 0.813
Social Support ⁵	19	195	0.977	0.971 - 0.982
PANAS Positive ⁶	10	285	0.884	0.854 - 0.907
PANAS Negative ⁶	10	282	0.827	0.793 - 0.867
Community Cohesion x	3	281	0.769	0.699 - 0.826

Note: 95% confidence intervals were calculated with 1,000 bootstrap samples. Standardized Cronbach's alpha (α) coefficients were computed. ¹Baessler and Schwarzer, 1996; ²Sinclair and Wallston, 2004; ³Andresen et al., 1994; ⁴Sherbourne and Stewart, 1991; ⁵Watson et al., 1988; ⁶Watson et al., 1988

2 Session info (for reproducibility)

```
# Display session information for reproducibility
# - Uses `pander()` for better formatting
# - `locale = FALSE` to exclude locale-specific info (reduces clutter)
library(pander)
pander(sessionInfo(), locale = FALSE)
```

R version 4.4.3 (2025-02-28)

Platform: x86 64-pc-linux-gnu

attached base packages: stats, graphics, grDevices, utils, datasets, methods and base

other attached packages: pander(v.0.6.6), lubridate(v.1.9.4), forcats(v.1.0.0), stringr(v.1.5.1), dplyr(v.1.1.4), purrr(v.1.0.4), tidyr(v.1.3.1), tibble(v.3.2.1), ggplot2(v.3.5.1), tidyverse(v.2.0.0), kableExtra(v.1.4.0), scales(v.1.3.0), readr(v.2.1.5), performance(v.0.13.0), MuMIn(v.1.48.4), psych(v.2.4.12), ltm(v.1.2-0), polycor(v.0.8-1), msm(v.1.8.2), MASS(v.7.3-64) and knitr(v.1.49)

loaded via a namespace (and not attached): gtable(v.0.3.6), xfun(v.0.51), insight(v.1.1.0), lattice(v.0.22-6), tzdb(v.0.4.0), vctrs(v.0.6.5), tools(v.4.4.3), generics(v.0.1.3), stats4(v.4.4.3), parallel(v.4.4.3), pkgconfig(v.2.0.3), Matrix(v.1.7-2), lifecycle(v.1.0.4), compiler(v.4.4.3), munsell(v.0.5.1), mnormt(v.2.1.1), htmltools(v.0.5.8.1), yaml(v.2.3.10), crayon(v.1.5.3), pillar(v.1.10.1), admisc(v.0.37), nlme(v.3.1-167), tidyselect(v.1.2.1), digest(v.0.6.37), mvtnorm(v.1.3-3), stringi(v.1.8.4), bookdown(v.0.42), splines(v.4.4.3), fastmap(v.1.2.0), grid(v.4.4.3), archive(v.1.1.11), colorspace(v.2.1-1), expm(v.1.0-0), cli(v.3.6.3), magrittr(v.2.0.3), survival(v.3.8-3), withr(v.3.0.2), bit64(v.4.6.0-1), timechange(v.0.3.0), rmarkdown(v.2.29), bit(v.4.5.0.1), hms(v.1.1.3), evaluate(v.1.0.3), viridisLite(v.0.4.2), rlang(v.1.1.5), Rcpp(v.1.0.14), glue(v.1.8.0), xml2(v.1.3.6), svglite(v.2.1.3), rstudioapi(v.0.17.1), vroom(v.1.6.5), R6(v.2.5.1) and systemfonts(v.1.2.1)

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