This folder contains all the materials necessary for replicating all the results in the paper, including both its main text and online appendix.

The data set used for the analyses is **"Data_full.dta"**. The syntax files are **"1_analysis.do"** and **"2_plotting.R",** explained below.

File "1_analysis.do" is the Stata do-file that produces all the results in all tables in the submitted manuscript and online appendix. The complete log of this do-file is **"statacapacity.txt".** All the tables produced by "1_analysis.do" are stored in tables.xlsx.

File "2_plotting.R" is the R file that produces all the figures in the manuscript and online appendix. All the files with names beginning with "fg_" are the data files for making figures, which are used in "2_plotting.R". The codes that produced these figure data files are also in the syntax files "1_analysis.do" and "2_plotting.R", but they are commented out in order for the full replication to be as easy as possible. Any user who wants to regenerate these figure data files simply needs to uncomment these codes within the two syntax files.

Folder "output" is empty, and once the user executes "1_analysis.do", tmp files created while the code is being run will go to this folder.