GTAPSSP: SSPs for GTAP Framework

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1 Introduction

This tutorial demonstrates the utilization of the gtapssp package in R for data processing. It covers various steps such as reading, transforming, and analyzing data, making it suitable for both beginners and advanced users.

The package provides optimized and user-friendly functions to download SSP data, interpolate data using spline and beers methods. The qtapssp functions is accompanied by detailed manual, you can also access this manual by running ?gtapssp in the R console or pressing F1 on the function name in RStudio.

2 Installation

To use the *qtapssp* package, it's necessary to have R installed on your computer, which can be downloaded from here. Additionally, we recommend downloading RStudio, available at here, which provides a user-friendly interface to work with R.

? R install details

R is a versatile programming language, with a focus on statistical computing. It is a big part of academic research in the social sciences. R is free and open-source and runs on Windows, Mac OS X, and Linux.

- 1. Download the R installer from the Comprehensive R Archive Network (CRAN)
- Choose the appropriate installer for your operating system and computer architecture (32-bit or 64-bit).

- If on Mac, you will need to know if you are using an Intel or Apple Silicon (M1) processor.
- 2. Run the installer and follow the instructions.
- 3. We recommend install **R Tools**. Many of the packages we will be using in this course require R Tools to be installed.

If you are on Windows:

- Download the latest version of the software from the R Tools for Windows page.
- Run the installer and follow the instructions. If you are on Mac:
- First, install the Xcode Command Line Tools. Go to the R Tools for Mac page and follow the instructions. Note: the precise instructions will vary according to the version of macOS you are using.
- Install the gfortran compiler, as also indicated on the R Tools for Mac page.

? IDE install details

We recommend RStudio as IDE for R.

RStudio is by far the most popular IDE used by R programmers. It is free and open-source and comes with a console and syntax-highlighting editor that supports direct code execution, as well as tools for plotting, history, debugging, and workspace management. Alternative: If you are already familiar with Visual Studio Code (VS Code), say because you have already used it for Python, you can also use it for R programming. You will need to do a bit of configuration to get it to work, though. If you choose to use VS Code, download and install the R Extension for Visual Studio Code and remember to follow the instructions on the Getting Started section of the extension page.

You can install the development version of gtapssp from GitHub with:

```
# If the devtools package is not already installed, please run the disabled line below.
# install.packages("devtools")
devtools::install_github("tsimonato/gtapssp")
```

3 Quick Start: Run the Entire Pipeline

If you'd like to execute the entire pipeline described in this tutorial with minimal effort, you can use the gtapssp::iiasa_gtap() function. This function runs all steps, including data

aggregation, interpolation, expansion, label merging, and optional file exports.

Here's an example:

```
# Run the entire pipeline and save the output as a .HAR file
gtapssp::iiasa_gtap(outFile = "gtap_ssp.har")

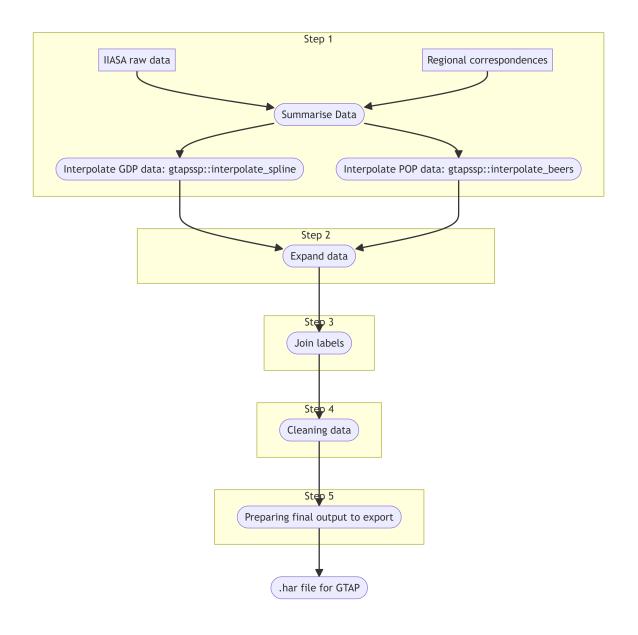
# Or save as a .CSV file
gtapssp::iiasa_gtap(outFile = "gtap_ssp.csv")

# To simply process the data without saving it
final_data <- gtapssp::iiasa_gtap()</pre>
```

This function provides flexibility to either save the output in .har or .csv formats or return the processed dataset for further use in R.

4 Workflow Overview

The pipeline involves the following steps:



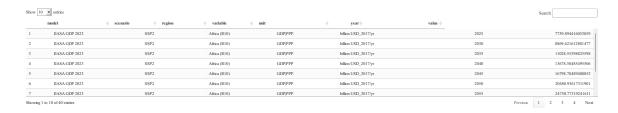
5 Data Source

The pipeline uses projections from the Shared Socioeconomic Pathways (SSPs) developed by IIASA (version 3.0.1, March 2024). These projections include **GDP** and **Population** data and are publicly available under a license allowing reuse by other research communities. For more details, visit the official IIASA SSP database.

Below is a preview of the default IIASA dataset used in the gtapssp package. This dataset can be updated with the gtapssp::updateData function if newer data or custom updates are

required.

gtapssp::iiasa_raw\$data



6 Step 1: Preprocessing and Aggregation

Aggregate the raw data using gtapssp::aggData. This function groups the dataset by specified columns and sums the values within each group, ensuring compatibility with the GTAP regional structure.

```
# Define grouping columns
group_cols <- c("model", "scenario", "reg_iso3", "variable", "unit")
# Aggregate raw data
agg_iiasa <- gtapssp::aggData(
    iiasa_raw = gtapssp::iiasa_raw,
    group_cols = group_cols
)</pre>
```