

EMBRACE-II Data Processing Pipeline

Overview

The `embraceR` package provides a comprehensive data processing pipeline for the EMBRACE-II study data. This article explains the data flow from raw input files to the final clean dataset ready for analysis.

Data Processing Pipeline

```
# Example of getting the final clean dataset  
clean_data <- emii_get_clean_data()  
  
# With Excel export  
clean_data <- emii_get_clean_data(save_excel = TRUE)
```

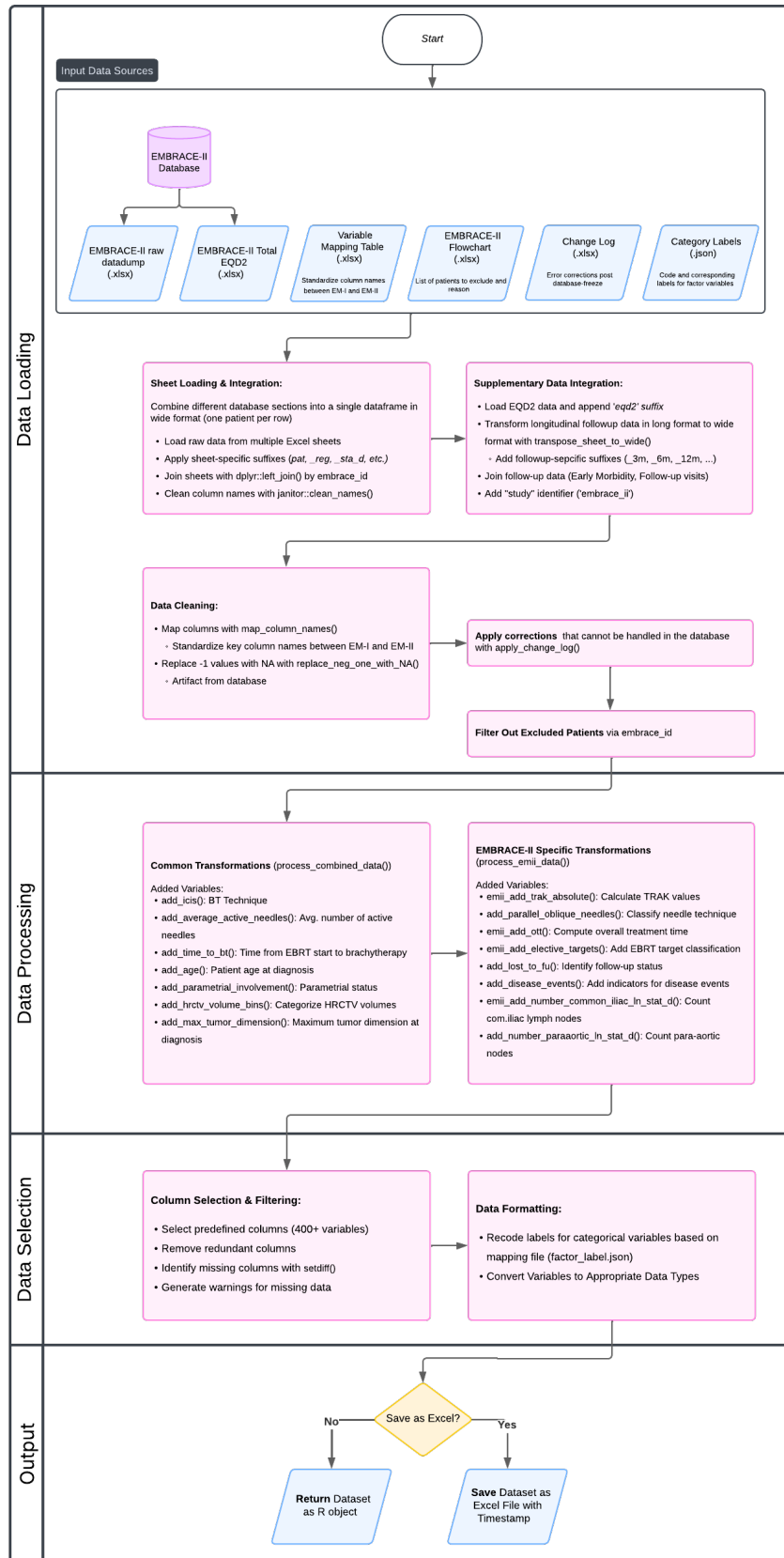


Figure 1: EMBRACE-II Data Processing Pipeline