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Statistics
Korea



KOSTAT-UNFPA Summer Seminar on Population

Workshop 1. Demography in R

Day 6: Data Prep for Processing and visualizing South Korean fertility microdata

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1 Summary

This markdown file steps through the data prep of the microdata. Here I have the simple task of converting the (very nicely formatted) `.csv` files provided by KOSTAT into a fixed width format. I want to do this because very often when we get microdata from official statistics offices, it is delivered in fixed-width files, often public-use files where we need to know column positions and widths in order to read the data into R. That is, here we take a step *backwards*! The metadata about which column is located in which position, and how wide it is is best delivered in a spreadsheet, but sometimes it is given in `pdf` files, which can be laborious to prepare.

2 examine file

```
library(tidyverse)
library(readr)
folder_pieces <- c("Data",
```

```

      "2000-2020_birth_annual data (for W1)_20220706",
      "2000-2020_birth_annual data", "Excel")

folder <- paste(folder_pieces, collapse="/")
files <- dir(folder)

A <- read_csv(file.path(folder, files[1]),
              show_col_types = FALSE)
# View(A)

```

2.1 Improvise way to write fwf

Solution inspired by: Hao Zhu's GitHub gist: https://gist.github.com/haozhu233/28d1309b58431f4929f782430541f58/raw/3c888574eba9bdde41b5f9081de7c9533e84f20d/write_fwf.R I have modified the function a bit to get the result we want: no headers given, this information is delivered with the column positions in a metadata file.

In real life, you will NEVER need to do what I'm doing here.

```

write_fwf = function(dt, file, width,
                     justify = "l", replace_na = "NA") {
  fct_col = which(sapply(dt, is.factor))
  if (length(fct_col) > 0) {
    for (i in fct_col) {
      dt[,i] <- as.character(dt[,i])
    }
  }
  dt[is.na(dt)] = replace_na
  n_col = ncol(dt)
  justify = unlist(strsplit(justify, ""))
  justify = as.character(factor(justify, c("l", "r"), c("-", "")))
  if (n_col != 1) {
    if (length(width) == 1) width = rep(width, n_col)
    if (length(justify) == 1) justify = rep(justify, n_col)
  }
  sptf_fmt = paste0(
    paste0("%", justify, width, "s"), collapse = ""
  )
  tbl_content = do.call(sprintf, c(fmt = sptf_fmt, dt))
  writeLines(tbl_content, file)
}

```

2.2 Write out as flat files

We now apply this function in a loop, sending the resulting text files to a folder called Korea_births_fwf inside Data.

```

# based on eyeballing the data; it's OK if these are bigger than needed; not
# OK if they are smaller than needed.
widths <- c(7,7,4,7,1,7,2,5,5,3,6)

years <- 2000:2020

```

```

# create our metadata and save a copy for sharing
positions_names <- tibble(colname = colnames(A),
                          width = widths)
write_csv(positions_names, "Data/Korea_births_fwf_metadata.csv")

if (!file.exists("Data/Korea_births_fwf")){
  dir.create("Data/Korea_births_fwf")
}
# a loop! i is an index counter of the elements of `years`
for (i in 1:length(years)){
  file_in      <- read_csv(file.path(folder, files[i]),
                          # suppresses messages when reading in
                          show_col_types = FALSE)

  # create name for the outgoing file
  file_out_name <- paste0("kb",years[i],".txt")

  # write the file using the custom function
  write_fwf(file_in,
            file = file.path("Data/Korea_births_fwf",file_out_name),
            width = widths)
  # remove biggish file from workspace, clear memory of it
  rm(file_in);gc()
}

```

2.3 Read back in?

```

library(vroom)

specs <- fwf_widths(widths = positions_names$width,
                    col_names = positions_names$colname)

files <- file.path("Data/Korea_births_fwf",
                  paste0("kb",2000:2020,".txt"))
# vroom reads them all in and rbinds them together nicely :-)
A      <- vroom_fwf(files,
                    col_positions = specs)

## Rows: 9369102 Columns: 11
## -- Column specification -----
##
## chr  (1): Birth weight
## dbl (10): Report year, Year, Report month, Report date, Sex, Year of birth, ...
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
# In Terminal, I typed:
# cd workspace/KOSTAT_Workshop1/Data
# zip -r Korea_births_fwf.zip Korea_births_fwf
# Also possible from R, but I didn't bother

```

2.4 Test unzip

We can unzip in session, no problem

```
unzip("Data/Korea_births_fwf.zip",  
  
      # in case this isn't the first time  
      overwrite = TRUE,  
  
      # which directory to stash the result in  
      exdir = "Data")
```

Objective achieved!