## FAO Crops Data 1961-2007

## Md. Tanvir Rahman

2024-09-22

```
# import packages
library(dplyr)
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
##
      filter, lag
## The following objects are masked from 'package:base':
##
       intersect, setdiff, setequal, union
library(tidyverse)
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v forcats 1.0.0 v readr
                                   2.1.5
## v ggplot2 3.5.1 v stringr 1.5.1
## v lubridate 1.9.3 v tibble
                                    3.2.1
## v purrr 1.0.2
                        v tidyr
                                    1.3.1
## -- Conflicts -----
                                          ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                    masks stats::lag()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
#import data of csv files in R programming
getwd() #fuction to know the current working directory(WD)
## [1] "D:/faodata"
setwd("D:\\faodata")#function to reset the current WD
#import csv files
faocrops <-read.csv("fao_data_crops.csv")</pre>
```

```
# Data Explore
View(faocrops)
colnames(faocrops)
## [1] "country_or_area" "element_code"
                                           "element"
                                                              "vear"
## [5] "unit"
                                           "value_footnotes" "category"
                         "value"
# Get unique country or area of the 'country_or_area' column
unique_country_or_area <- unique(faocrops$country_or_area)</pre>
# Remove empty string entries
unique_country_or_area <- unique_country_or_area[unique_country_or_area != ""]
# Print the filtered unique elements
#There are 259 unique elements in the country_or_area column
head(unique_country_or_area)
## [1] "Americas +"
                           "Asia +"
                                               "Caribbean +"
## [4] "Central America +" "Colombia"
                                               "Cuba"
tail(unique_country_or_area)
## [1] "Namibia"
                          "Marshall Islands" "Nauru"
                                                                 "Saint Helena"
## [5] "Liechtenstein" "Faroe Islands"
# Get unique elements of the 'element' column
unique_element <- unique(faocrops$element)</pre>
# Remove empty string entries
unique_element <- unique_element[unique_element != ""]</pre>
# Print the filtered unique elements
print(unique_element) # There are 10 unique elements in the element column
## [1] "Area Harvested"
## [2] "Yield"
## [3] "Production Quantity"
## [4] "Seed"
## [5] "Gross Production 1999-2001 (1000 I$)"
## [6] "Net Production 1999-2001 (1000 I$)"
   [7] "Gross PIN (base 1999-2001)"
## [8] "Grs per capita PIN (base 1999-2001)"
## [9] "Net PIN (base 1999-2001)"
## [10] "Net per capita PIN (base 1999-2001)"
# Get unique element code of the 'element_code' column
unique_element_code <- unique(faocrops$element_code)</pre>
print(unique_element_code) # There are some unwanted elements in the unique elements code
## [1] "31"
## [2] "41"
## [3] "51"
## [4] "Footnote"
```

```
## [5] "Calculated Data"
## [6] "May include official, semi-official or estimated data"
## [7] "Not reported by country"
## [8] "FAO Estimate"
## [9] "Unofficial figure"
## [10] "111"
## [11] "152"
## [12] "154"
## [13] "432"
## [14] "434"
## [15] "436"
## [16] "438"
# Define the elements to be removed
elements_to_remove <- c("Footnote", "Calculated Data", "May include official, semi-official or estimate
                        "Not reported by country", "FAO Estimate", "Unofficial figure")
# Remove the specified elements
unique_element_code <- setdiff(unique_element_code, elements_to_remove)</pre>
# Print the filtered unique elements
print(unique_element_code)
## [1] "31" "41" "51" "111" "152" "154" "432" "434" "436" "438"
# Get unique elements of the 'year' column
unique_year <- unique(faocrops$year)</pre>
# Remove NA values
unique_year <- unique_year[!is.na(unique_year)]</pre>
# Print the filtered unique elements
print(unique_year)
## [1] 2007 2006 2005 2004 2003 2002 2001 2000 1999 1998 1997 1996 1995 1994 1993
## [16] 1992 1991 1990 1989 1988 1987 1986 1985 1984 1983 1982 1981 1980 1979 1978
## [31] 1977 1976 1975 1974 1973 1972 1971 1970 1969 1968 1967 1966 1965 1964 1963
## [46] 1962 1961
# Print the filtered unique elements
print(unique_year)
## [1] 2007 2006 2005 2004 2003 2002 2001 2000 1999 1998 1997 1996 1995 1994 1993
## [16] 1992 1991 1990 1989 1988 1987 1986 1985 1984 1983 1982 1981 1980 1979 1978
## [31] 1977 1976 1975 1974 1973 1972 1971 1970 1969 1968 1967 1966 1965 1964 1963
## [46] 1962 1961
head(unique_year)
## [1] 2007 2006 2005 2004 2003 2002
tail(unique_year)
## [1] 1966 1965 1964 1963 1962 1961
```

```
# The dataset contain information of 1961-2007
# Get unique elements of the 'unit' column
unique_unit <- unique(faocrops$unit)</pre>
# Remove NA values
unique_unit <- unique_unit[unique_unit != ""]</pre>
# Print the filtered unique elements
print(unique_unit) # There are 5 unique elements in the elements column
## [1] "Ha"
                     "Hg/Ha"
                                   "tonnes"
                                                 "1000 Int. $" "Int. $"
# Get unique elements of the 'category' column
unique_category <- unique(faocrops$category) # There are 172 unique elements in the category column
head(unique_category)
## [1] "agave_fibres_nes"
                                    "almonds_with_shell"
## [3] "anise_badian_fennel_corian" "apples"
## [5] "apricots"
                                    "arecanuts"
tail(unique_category)
## [1] "vetches"
                            "walnuts_with_shell" "watermelons"
## [4] "wheat"
                            "yams"
                                                 "yautia_cocoyam"
```