# Population prescribed drugs for anxiety or depression or psychosis

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14 November, 2023, 20:31

```
# tidyverse includes dplyr and ggplot2 so I don't need to load them separately
library(tidyverse)
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr 1.1.3
                       v readr
                                    2.1.4
## v forcats 1.0.0
                                    1.5.0
                        v stringr
## v ggplot2 3.4.4
                        v tibble
                                    3.2.1
## v lubridate 1.9.3
                        v tidyr
                                    1.3.0
## v purrr
              1.0.2
## -- 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
library(here)
## here() starts at C:/Users/Fionnuala/OneDrive - University of Aberdeen/PU5063 Intro to HDS/Assessment
library(viridis)
## Loading required package: viridisLite
library(plotly)
##
## Attaching package: 'plotly'
## The following object is masked from 'package:ggplot2':
##
##
      last_plot
##
## The following object is masked from 'package:stats':
##
##
      filter
## The following object is masked from 'package:graphics':
##
##
      layout
```

### Question

What are the regional trends for the number of people prescribed drugs for anxiety, depression and psychosis in Scotland over the last ten years? What might these mean for employers' allocation of support resources? The next sections follow the Health Data Science Workflow to address these questions.

## **Data Acquisition**

The data was downloaded from https://scotland.shinyapps.io/ScotPHO\_profiles\_tool/ on 05/11/23 for the item "population prescribed drugs for anxiety/depression/psychosis" for all available years and all health boards. The downloaded file was called timetrend data.csv

```
#reading in the data and checking its columns:
adp data <- read csv(here("Inputs/timetrend data.csv"))</pre>
## Rows: 180 Columns: 12
## -- Column specification -----
## Delimiter: ","
## chr (7): indicator, area_name, area_code, area_type, period, definition, dat...
## dbl (5): year, numerator, measure, lower confidence interval, upper confiden...
## 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.
glimpse(adp_data)
## Rows: 180
## Columns: 12
## $ indicator
                               <chr> "Population prescribed drugs for anxiety/dep~
                               <chr> "Scotland", "NHS Ayrshire & Arran", "NHS Bor~
## $ area_name
                               <chr> "S00000001", "S08000015", "S08000016", "S080~
## $ area code
## $ area_type
                               <chr> "Scotland", "Health board", "Health board", ~
## $ year
                               <dbl> 2010, 2010, 2010, 2010, 2010, 2010, 2010, 20~
## $ period
                               <chr> "2010/11 financial year", "2010/11 financial~
                               <dbl> 787040, 60822, 17226, 22280, 55334, 43976, 7~
## $ numerator
## $ measure
                               <dbl> 14.96, 16.31, 15.15, 14.75, 15.26, 14.86, 12~
## $ lower_confidence_interval <dbl> 14.93, 16.20, 14.94, 14.57, 15.14, 14.73, 12~
## $ upper_confidence_interval <dbl> 14.99, 16.43, 15.36, 14.92, 15.38, 14.98, 12~
                               <chr> "Percentage", "Percentage", "Percentage", "P~
## $ definition
## $ data_source
                              <chr> "Public Health Scotland (Prescribing Informa~
#Prepare/Clean Data
# There are no missing values
# This chunk is for selecting and renaming columns, removing the rows for the
# whole of Scotland and removing the NHS prefix.
# The mutate line was suggested by chatgpt when I gave it the preceding lines
# in this chunk and asked it what to add to strip out the "NHS " prefix.
clean_data <- adp_data %>%
```

```
select('area_name','year','numerator', 'measure') %>%
  rename(number = 'numerator', NHS = 'area_name', percentage = 'measure') %>%
  filter(NHS !='Scotland') %>%
  mutate(NHS = sub("^NHS ","", NHS))
head(clean_data)
## # A tibble: 6 x 4
##
    NHS
                         year number percentage
##
     <chr>>
                         <dbl> <dbl>
                                           <dbl>
                          2010 60822
                                            16.3
## 1 Ayrshire & Arran
## 2 Borders
                          2010 17226
                                            15.2
## 3 Dumfries & Galloway 2010 22280
                                            14.8
                          2010 55334
## 4 Fife
                                            15.3
                          2010 43976
## 5 Forth Valley
                                            14.9
## 6 Grampian
                          2010 70337
                                            12.4
# The below code plots the data as-is. It's here so anyone can plot it and see
# for themselves that 14 is too many categories, but it's commented out so it
# doesn't get confused with the intended plot.
# options(scipen = 999)
# clean_data %>%
   ggplot() +
   geom\_area(aes(x = year, y = number, fill = NHS))
```

#### #Analyse

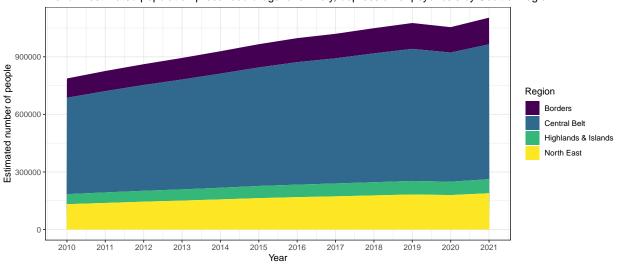
14 Health Boards are too many to plot in the same visualisation; the audience would be overwhelmed. So, I will create a new column, classifying neighbouring NHS boards into Central Belt, Borders, Highlands and Islands and North East. Then, I have to sum the old NHS Board percentages for each year into a single value for the Region for that year.

```
#I wanted to use functions from the course but when I looked up how to recategorise a categorical varia
group_data <- clean_data %>%
  mutate(Region = case_when(
   NHS %in% c("Ayrshire & Arran" , "Borders" , "Dumfries & Galloway")
    ~ "Borders",
   NHS %in% c("Fife" , "Forth Valley" , "Greater Glasgow & Clyde" ,
               "Lanarkshire", "Lothian")
    ~ "Central Belt",
   NHS %in% c("Grampian" , "Tayside")
   ~ "North East",
   NHS %in% c("Highland", "Western Isles", "Orkney", "Shetland")
    ~ "Highlands & Islands"))
#Now to re-calculate the regional percentages
summed_data <- group_data %>%
  group_by(Region, year) %>%
  summarise(total_people = sum(number))
```

```
## `summarise()` has grouped output by 'Region'. You can override using the
## `.groups` argument.
```

```
options(scipen = 999)
summed_data %>%
    ggplot(aes(x = year, y = total_people, fill = Region)) +
    geom_area()+
    xlab("Year")+
    ylab("Estimated number of people")+
    ggtitle("Trend in estimated population prescribed drugs for anxiety, depression or psychosis by Scott
    scale_fill_viridis(discrete=TRUE)+
    scale_x_continuous(breaks = unique(summed_data$year))+
    theme_bw()
```

### Trend in estimated population prescribed drugs for anxiety, depression or psychosis by Scottish region



```
# If switching the output to html, this line shows values when hovered over. # It's commented out here because the output is PDF. # ggplotly(tooltip = c("x", "y"))
```