

# Intro to Data Science - HW 5

Copyright Jeffrey Stanton, Jeffrey Saltz, and Jasmina Tacheva

```
# Enter your name here: Benjamin Tisinger
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## Attribution statement: (choose only one and delete the rest)

```
# 1. I did this homework by myself, with help from the book and the professor.
```

**This module: Data visualization** is important because many people can make sense of data more easily when it is presented in graphic form. As a data scientist, you will have to present complex data to decision makers in a form that makes the data interpretable for them. From your experience with Excel and other tools, you know that there are a variety of **common data visualizations** (e.g., pie charts). How many of them can you name?

The most powerful tool for data visualization in R is called **ggplot**. Written by computer/data scientist **Hadley Wickham**, this “**graphics grammar**” tool builds visualizations in layers. This method provides immense flexibility, but takes a bit of practice to master.

## Step 1: Make a copy of the data

- A. Read the **who** dataset from this URL: <https://intro-datascience.s3.us-east-2.amazonaws.com/who.csv>  
(<https://intro-datascience.s3.us-east-2.amazonaws.com/who.csv>)  
into a new dataframe called **tb**.

Your new dataframe, **tb**, contains a so-called **multivariate time series**: a sequence of measurements on 23 Tuberculosis-related (TB) variables captured repeatedly over time (1980-2013). Familiarize yourself with the nature of the 23 variables by consulting the dataset’s codebook which can be found here: [https://intro-datascience.s3.us-east-2.amazonaws.com/TB\\_data\\_dictionary\\_2021-02-06.csv](https://intro-datascience.s3.us-east-2.amazonaws.com/TB_data_dictionary_2021-02-06.csv) ([https://intro-datascience.s3.us-east-2.amazonaws.com/TB\\_data\\_dictionary\\_2021-02-06.csv](https://intro-datascience.s3.us-east-2.amazonaws.com/TB_data_dictionary_2021-02-06.csv)).

```
library(dbplyr)
library(tidyverse)
```

```
## — Attaching packages — tidyverse 1.3.2 —
## ✓ ggplot2 3.3.6      ✓ purrr   0.3.5
## ✓ tibble  3.1.8      ✓ dplyr   1.0.10
## ✓ tidyr   1.2.1      ✓ stringr 1.4.1
## ✓ readr   2.1.3      ✓ forcats 0.5.2
## — Conflicts — tidyverse_conflicts() —
## X dplyr::filter() masks stats::filter()
## X dplyr::ident()  masks dbplyr::ident()
## X dplyr::lag()    masks stats::lag()
## X dplyr::sql()    masks dbplyr::sql()
```

```
tb <- read.csv('https://intro-datascience.s3.us-east-2.amazonaws.com/who.csv')
head(tb,15)
```

##	iso2	year	new_sp	new_sp_m04	new_sp_m514	new_sp_m014	new_sp_m1524	
## 1	AD	1989	NA	NA	NA	NA	NA	
## 2	AD	1990	NA	NA	NA	NA	NA	
## 3	AD	1991	NA	NA	NA	NA	NA	
## 4	AD	1992	NA	NA	NA	NA	NA	
## 5	AD	1993	15	NA	NA	NA	NA	
## 6	AD	1994	24	NA	NA	NA	NA	
## 7	AD	1996	8	NA	NA	0	0	
## 8	AD	1997	17	NA	NA	0	0	
## 9	AD	1998	1	NA	NA	0	0	
## 10	AD	1999	4	NA	NA	0	0	
## 11	AD	2000	1	NA	NA	0	0	
## 12	AD	2001	3	NA	NA	0	NA	
## 13	AD	2002	2	NA	NA	0	0	
## 14	AD	2003	7	NA	NA	0	0	
## 15	AD	2004	3	NA	NA	0	0	
##			new_sp_m2534	new_sp_m3544	new_sp_m4554	new_sp_m5564	new_sp_m65	new_sp_mu
## 1			NA	NA	NA	NA	NA	NA
## 2			NA	NA	NA	NA	NA	NA
## 3			NA	NA	NA	NA	NA	NA
## 4			NA	NA	NA	NA	NA	NA
## 5			NA	NA	NA	NA	NA	NA
## 6			NA	NA	NA	NA	NA	NA
## 7			0	4	1	0	0	NA
## 8			1	2	2	1	6	NA
## 9			0	1	0	0	0	NA
## 10			0	1	1	0	0	NA
## 11			1	0	0	0	0	NA
## 12			NA	2	1	NA	NA	NA
## 13			0	1	0	0	0	NA
## 14			0	1	2	0	0	NA
## 15			0	1	1	0	0	NA
##			new_sp_f04	new_sp_f514	new_sp_f014	new_sp_f1524	new_sp_f2534	new_sp_f3544
## 1			NA	NA	NA	NA	NA	NA
## 2			NA	NA	NA	NA	NA	NA
## 3			NA	NA	NA	NA	NA	NA
## 4			NA	NA	NA	NA	NA	NA
## 5			NA	NA	NA	NA	NA	NA
## 6			NA	NA	NA	NA	NA	NA
## 7			NA	NA	0	1	1	0
## 8			NA	NA	0	1	2	3
## 9			NA	NA	NA	NA	NA	NA
## 10			NA	NA	0	0	0	1
## 11			NA	NA	NA	NA	NA	NA
## 12			NA	NA	NA	NA	NA	NA
## 13			NA	NA	0	1	0	0
## 14			NA	NA	0	1	1	1
## 15			NA	NA	0	0	1	0
##			new_sp_f4554	new_sp_f5564	new_sp_f65	new_sp_fu		
## 1			NA	NA	NA	NA		
## 2			NA	NA	NA	NA		
## 3			NA	NA	NA	NA		

## 4	NA	NA	NA	NA
## 5	NA	NA	NA	NA
## 6	NA	NA	NA	NA
## 7	0	1	0	NA
## 8	0	0	1	NA
## 9	NA	NA	NA	NA
## 10	0	0	0	NA
## 11	NA	NA	NA	NA
## 12	NA	NA	NA	NA
## 13	0	0	0	NA
## 14	0	0	0	NA
## 15	0	0	0	NA

B. How often were these measurements taken (in other words, at what frequency were the variables measured)? Put your answer in a comment.

```
min(tb$year)
```

```
## [1] 1980
```

```
max(tb$year)
```

```
## [1] 2008
```

```
#The measurements are taken at a frequency of every year starting at 1980 until 2008.
```

## Step 2: Clean-up the NAs and create a subset

A. Let's clean up the iso2 attribute in **tb**

Hint: use *is.na()* – well use *!is.na()*

```
tb <- tb[!is.na(tb$iso2),]
head(tb,5)
```

```
##   iso2 year new_sp new_sp_m04 new_sp_m514 new_sp_m014 new_sp_m1524 new_sp_m2534
## 1  AD 1989     NA         NA         NA         NA         NA         NA
## 2  AD 1990     NA         NA         NA         NA         NA         NA
## 3  AD 1991     NA         NA         NA         NA         NA         NA
## 4  AD 1992     NA         NA         NA         NA         NA         NA
## 5  AD 1993     15         NA         NA         NA         NA         NA
##   new_sp_m3544 new_sp_m4554 new_sp_m5564 new_sp_m65 new_sp_mu new_sp_f04
## 1           NA         NA         NA         NA         NA         NA
## 2           NA         NA         NA         NA         NA         NA
## 3           NA         NA         NA         NA         NA         NA
## 4           NA         NA         NA         NA         NA         NA
## 5           NA         NA         NA         NA         NA         NA
##   new_sp_f514 new_sp_f014 new_sp_f1524 new_sp_f2534 new_sp_f3544 new_sp_f4554
## 1           NA         NA         NA         NA         NA         NA
## 2           NA         NA         NA         NA         NA         NA
## 3           NA         NA         NA         NA         NA         NA
## 4           NA         NA         NA         NA         NA         NA
## 5           NA         NA         NA         NA         NA         NA
##   new_sp_f5564 new_sp_f65 new_sp_fu
## 1           NA         NA         NA
## 2           NA         NA         NA
## 3           NA         NA         NA
## 4           NA         NA         NA
## 5           NA         NA         NA
```

B. Create a subset of **tb** containing **only the records for Canada (“CA” in the iso2 variable)**. Save it in a new dataframe called **tbCan**. Make sure this new df has **29 observations and 23 variables**.

```
tbCan <- subset(tb, tb$iso2 == 'CA')
head(tbCan,5)
```

```
##      iso2 year new_sp new_sp_m04 new_sp_m514 new_sp_m014 new_sp_m1524
## 872   CA 1980   951         NA         NA         12         54
## 873   CA 1981   803         NA         NA         8         49
## 874   CA 1982   812         NA         NA         6         52
## 875   CA 1983   771         NA         NA         9         47
## 876   CA 1984   811         NA         NA         3         44
##      new_sp_m2534 new_sp_m3544 new_sp_m4554 new_sp_m5564 new_sp_m65 new_sp_mu
## 872              75           83          100          108          186         NA
## 873              61           64           87          103          141         NA
## 874              66           69           90           91          150         NA
## 875              63           62           90           92          123         NA
## 876              75           58           68           83          169         NA
##      new_sp_f04 new_sp_f514 new_sp_f014 new_sp_f1524 new_sp_f2534 new_sp_f3544
## 872           NA           NA           18           62           51           34
## 873           NA           NA           6           46           57           26
## 874           NA           NA           7           51           57           30
## 875           NA           NA          11           50           50           29
## 876           NA           NA           9           51           59           28
##      new_sp_f4554 new_sp_f5564 new_sp_f65 new_sp_fu
## 872              31           33          104          NA
## 873              28           35           92          NA
## 874              25           38           80          NA
## 875              24           35           86          NA
## 876              28           36          100          NA
```

C. A simple method for dealing with small amounts of **missing data** in a numeric variable is to **substitute the mean of the variable in place of each missing datum**.

This expression locates (and reports to the console) all the missing data elements in the variable measuring the **number of positive pulmonary smear tests for male children 0-4 years old** (there are 26 data points missing)

```
tbCan$new_sp_m04[is.na(tbCan$new_sp_m04)]
```

```
## [1] NA NA NA NA NA NA NA NA NA NA NA NA NA NA NA NA NA NA NA NA NA NA NA
## [26] NA
```

```
Error in eval(expr, envir, enclos): object 'tbCan' not found
Traceback:
```

D. Write a comment describing how that statement works.

```
#This expressions is finding all the missing data located in tbCan and is filling it with NA so
the database looks and behaves cleaner
```

E. Write 4 more statements to check if there is missing data for the number of positive pulmonary smear tests for: **male and female** children 0-14 years old (**new\_sp\_m014** and **new\_sp\_f014**), and **male and female** citizens **65 years of age and older**, respectively. What does empty output suggest about the number of missing observations?

```
youngdudes <- tbCan$new_sp_m014[is.na(tbCan$new_sp_m014)]
youngfemales <- tbCan$new_sp_f014[is.na(tbCan$new_sp_f014)]

olddudes <- tbCan$new_sp_m65[is.na(tbCan$new_sp_m65)]
oldfemales <- tbCan$new_sp_f65[is.na(tbCan$new_sp_f65)]

head(youngdudes,5)
```

```
## integer(0)
```

```
head(youngfemales,5)
```

```
## integer(0)
```

```
head(olddudes,5)
```

```
## integer(0)
```

```
head(oldfemales,5)
```

```
## integer(0)
```

*#An output of integer(0) simply means there is No NA/Missing data in these sets*

There is an R package called **imputeTS** specifically designed to repair missing values in time series data. We will use this instead of mean substitution.

The **na\_interpolation()** function in this package takes advantage of a unique characteristic of time series data: **neighboring points in time can be used to “guess” about a missing value in between.**

- F. Install the **imputeTS** package (if needed) and use **na\_interpolation()** on the variable from part C. Don't forget that you need to save the results back to the **tbCan** dataframe. Also update any attribute discussed in part E (if needed).

```
library('imputeTS')
```

```
## Warning: package 'imputeTS' was built under R version 4.2.2
```

```
## Registered S3 method overwritten by 'quantmod':
##   method           from
##   as.zoo.data.frame zoo
```

```
tbCan$new_sp_m04 <- na_interpolation(tbCan$new_sp_m04)

tbCan$new_sp_m014 <- na_interpolation(tbCan$new_sp_m014)
tbCan$new_sp_f014 <- na_interpolation(tbCan$new_sp_f014)

tbCan$new_sp_m65 <- na_interpolation(tbCan$new_sp_m65)
tbCan$new_sp_f65 <- na_interpolation(tbCan$new_sp_f65)
```

G. Rerun the code from C and E above to check that all missing data have been fixed.

```
youngdudes <- tbCan$new_sp_m014[is.na(tbCan$new_sp_m014)]
youngfemales <- tbCan$new_sp_f014[is.na(tbCan$new_sp_f014)]

olddudes <- tbCan$new_sp_m65[is.na(tbCan$new_sp_m65)]
oldfemales <- tbCan$new_sp_f65[is.na(tbCan$new_sp_f65)]

head(youngdudes,5)
```

```
## integer(0)
```

```
head(youngfemales,5)
```

```
## integer(0)
```

```
head(olddudes,5)
```

```
## integer(0)
```

```
head(oldfemales,5)
```

```
## integer(0)
```

## Step 3: Use ggplot to explore the distribution of each variable

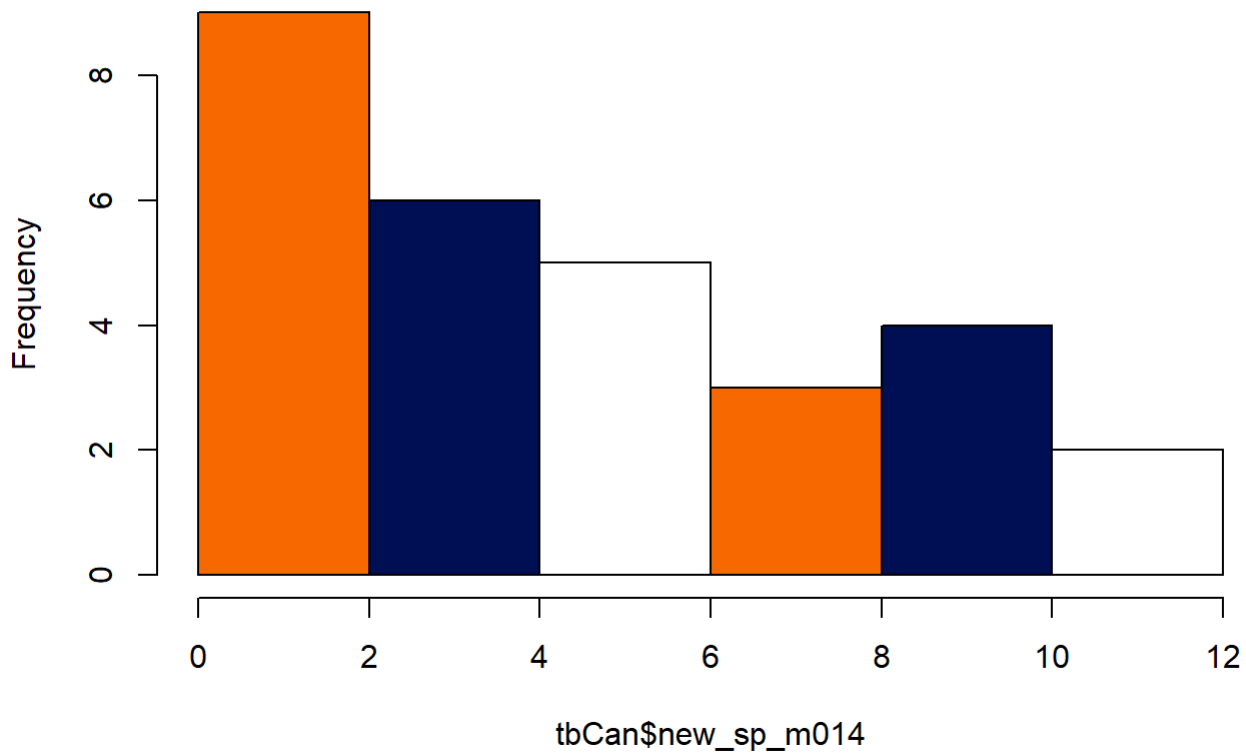
**Don't forget to install and library the ggplot2 package.** Then:

H. Create a histogram for **new\_sp\_m014**. Be sure to add a title and briefly describe what the histogram means in a comment.

```
library(ggplot2)

hist(tbCan$new_sp_m014, main="Histogram Males 0-14 with Positive Cases",
     col = c("#F76900", "#000E54", "#FFFFFF"))
```

## Histogram Males 0-14 with Positive Cases



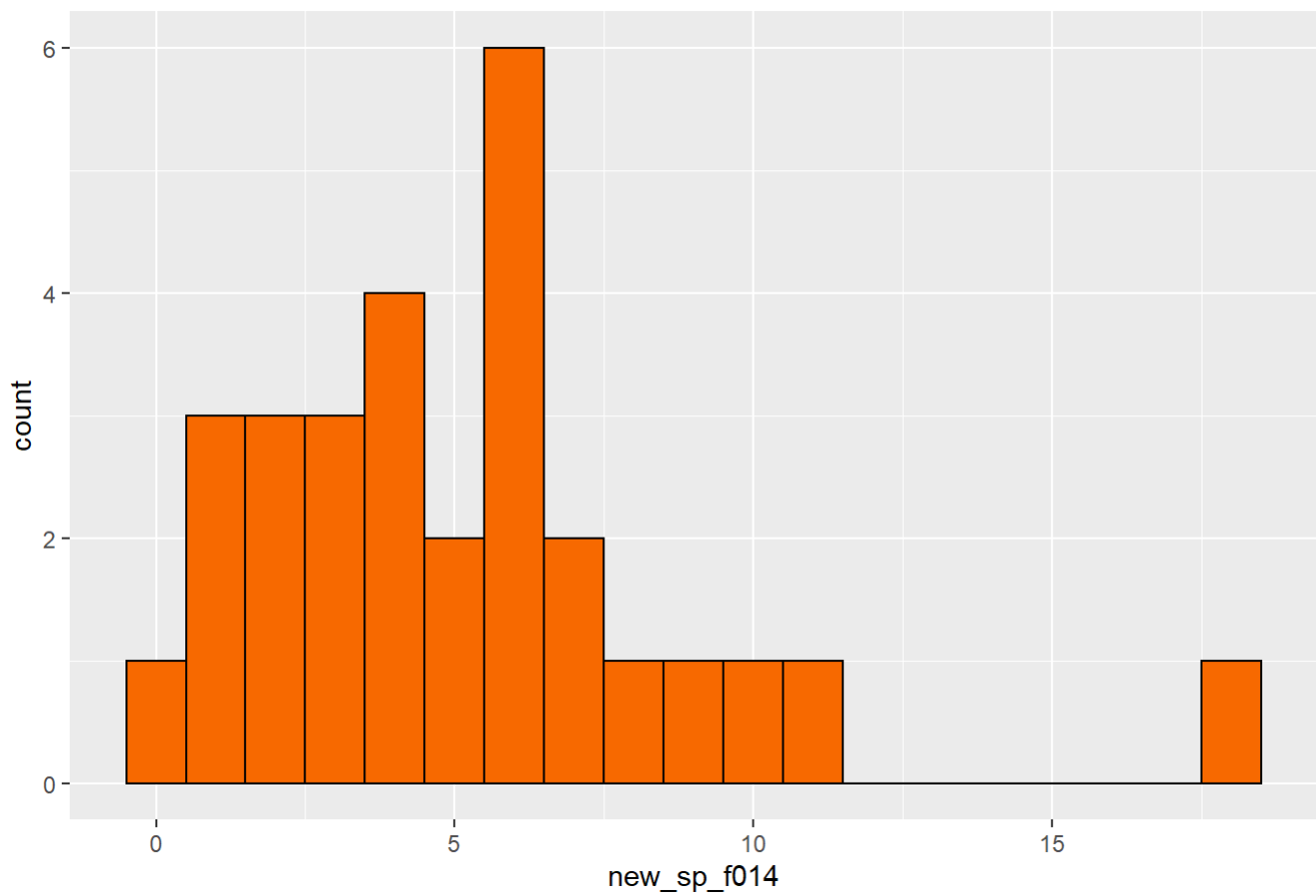
*#histogram of Male positive cases. This set of data is not very good for measurement*

- I. Create histograms (using ggplot) of each of the other three variables from E with ggplot( ).  
Which parameter do you need to adjust to make the other histograms look right?

```
tbCan %>% ggplot() +  
  geom_histogram(binwidth = 1,  
                 fill="#F76900",  
                 color="black",  
                 aes(x=new_sp_f014)) +  
  ggtitle('Histogram for Females in the 0-14 Age Range with a Postive Case')
```

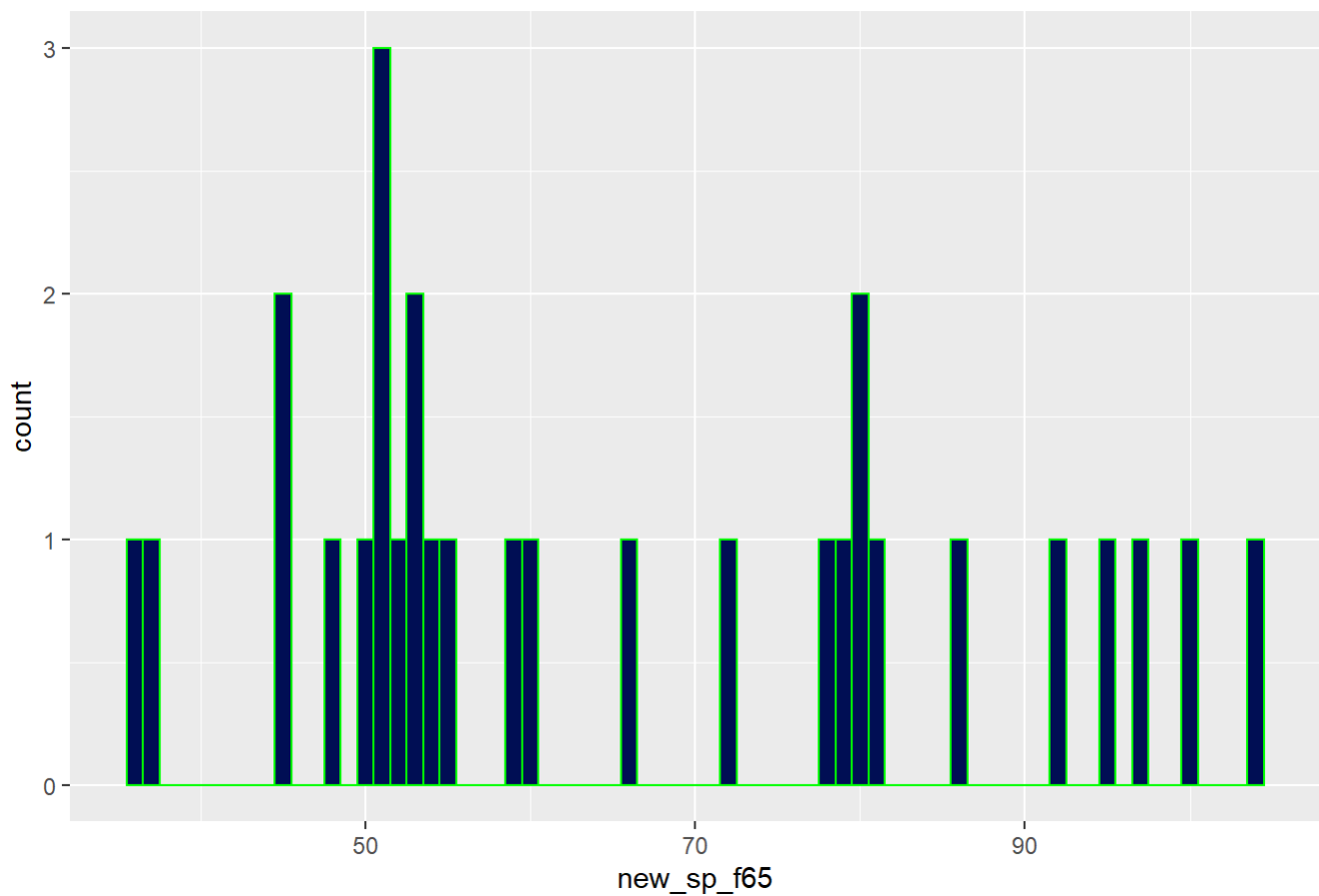


Histogram for Females in the 0-14 Age Range with a Postive Case



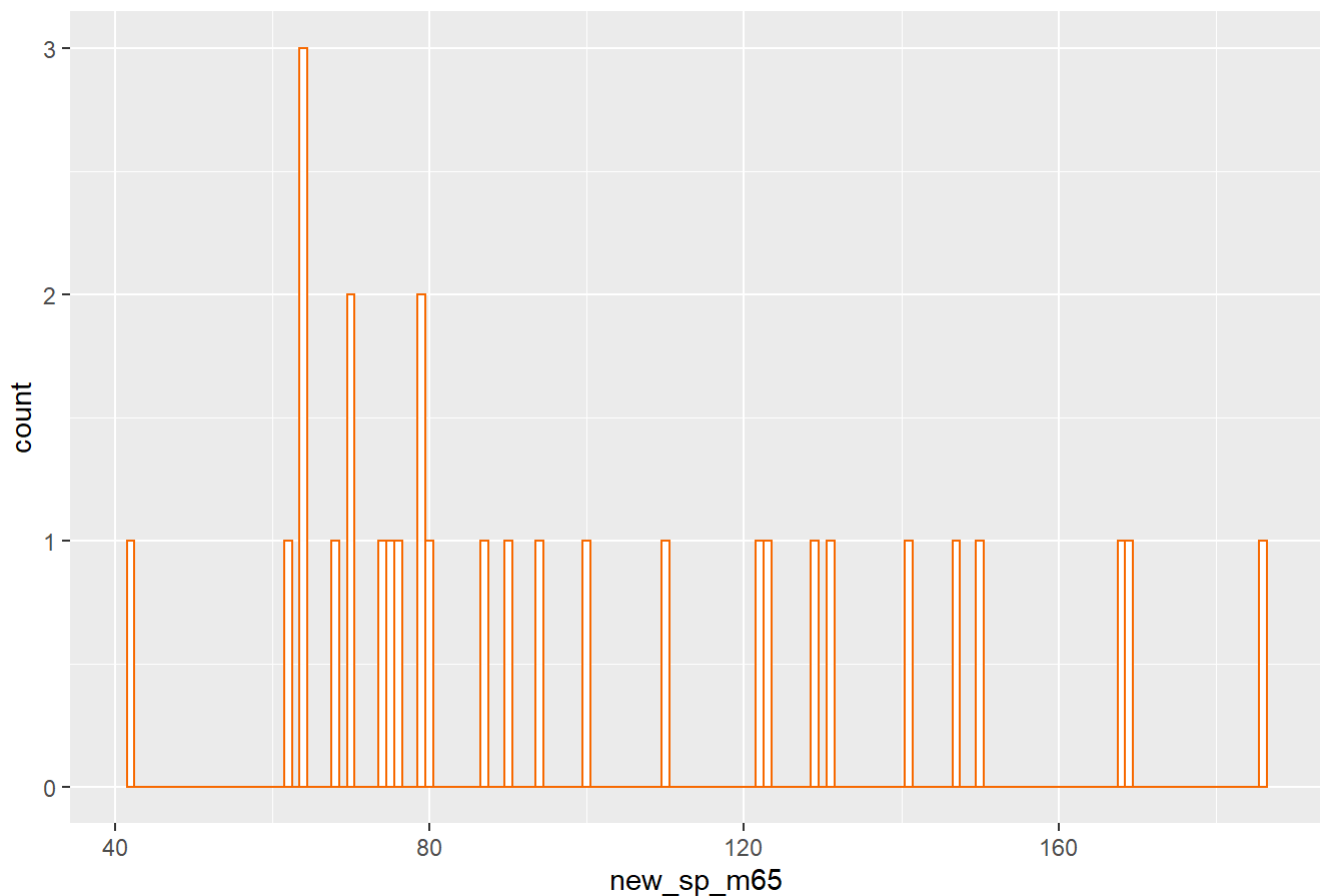
```
tbCan %>% ggplot() +  
  geom_histogram(binwidth = 1,  
                 fill="#000E54",  
                 color="green",  
                 aes(x=new_sp_f65)) +  
  ggtitle('Histogram for Females in the 65+ Age Range with a Postive Case')
```

Histogram for Females in the 65+ Age Range with a Postive Case



```
tbCan %>% ggplot() +  
  geom_histogram(binwidth = 1,  
    fill="#FFFFFF",  
    color="#F76900",  
    aes(x=new_sp_m65)) +  
  ggtitle('Histogram for Males in the 65+ Age Range with a Postive Case')
```

Histogram for Males in the 65+ Age Range with a Postive Case

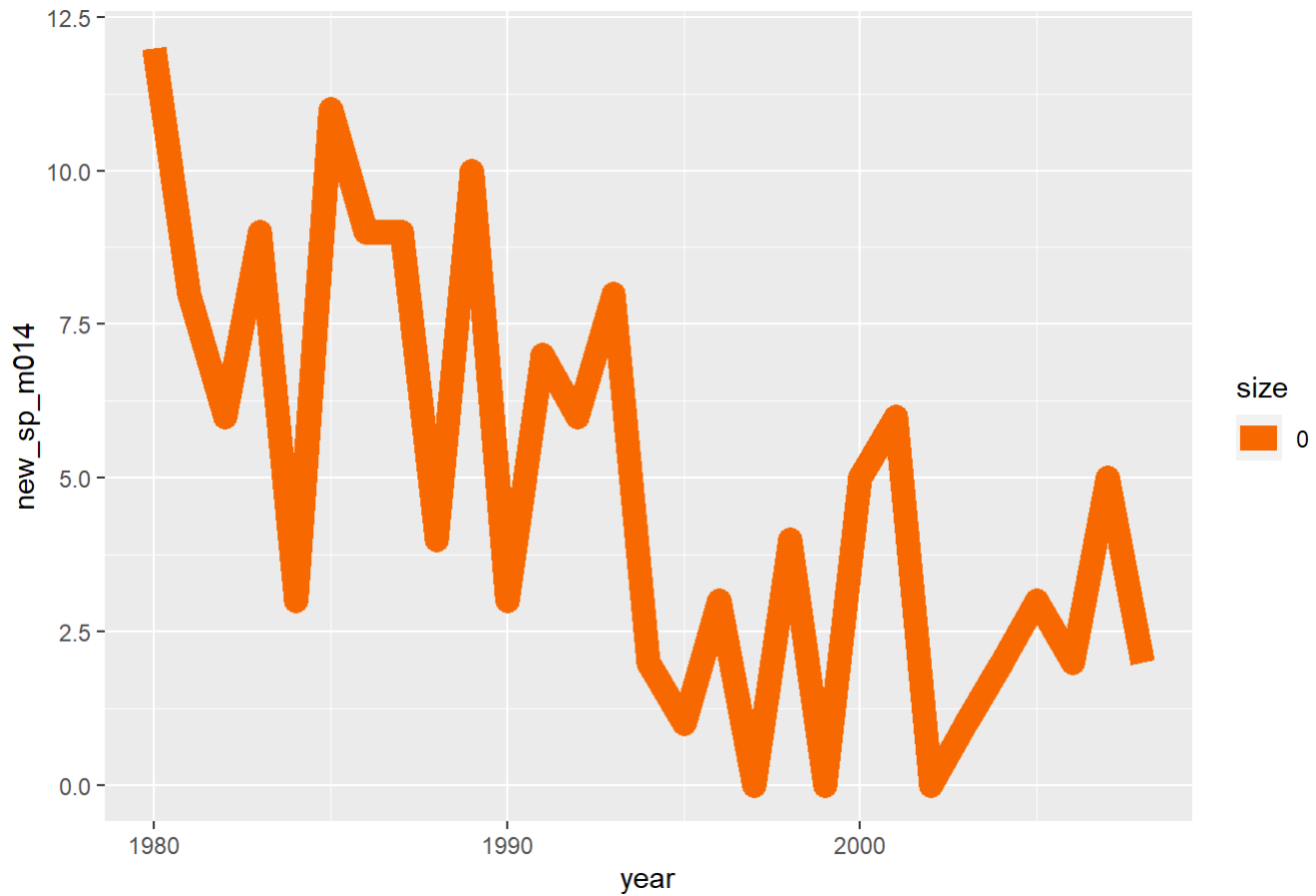


## Step 4: Explore how the data changes over time

J. These data were collected in a period of several decades (1980-2013). You can thus observe changes over time with the help of a line chart. Create a **line chart**, with **year** on the X-axis and **new\_sp\_m014** on the Y-axis.

```
tbCan %>% ggplot() +  
  geom_line(color = '#F76900',  
    aes(x=year, y=new_sp_m014, size=0 )) +  
  ggtitle('Histogram Males 0-14 with Positive Cases')
```

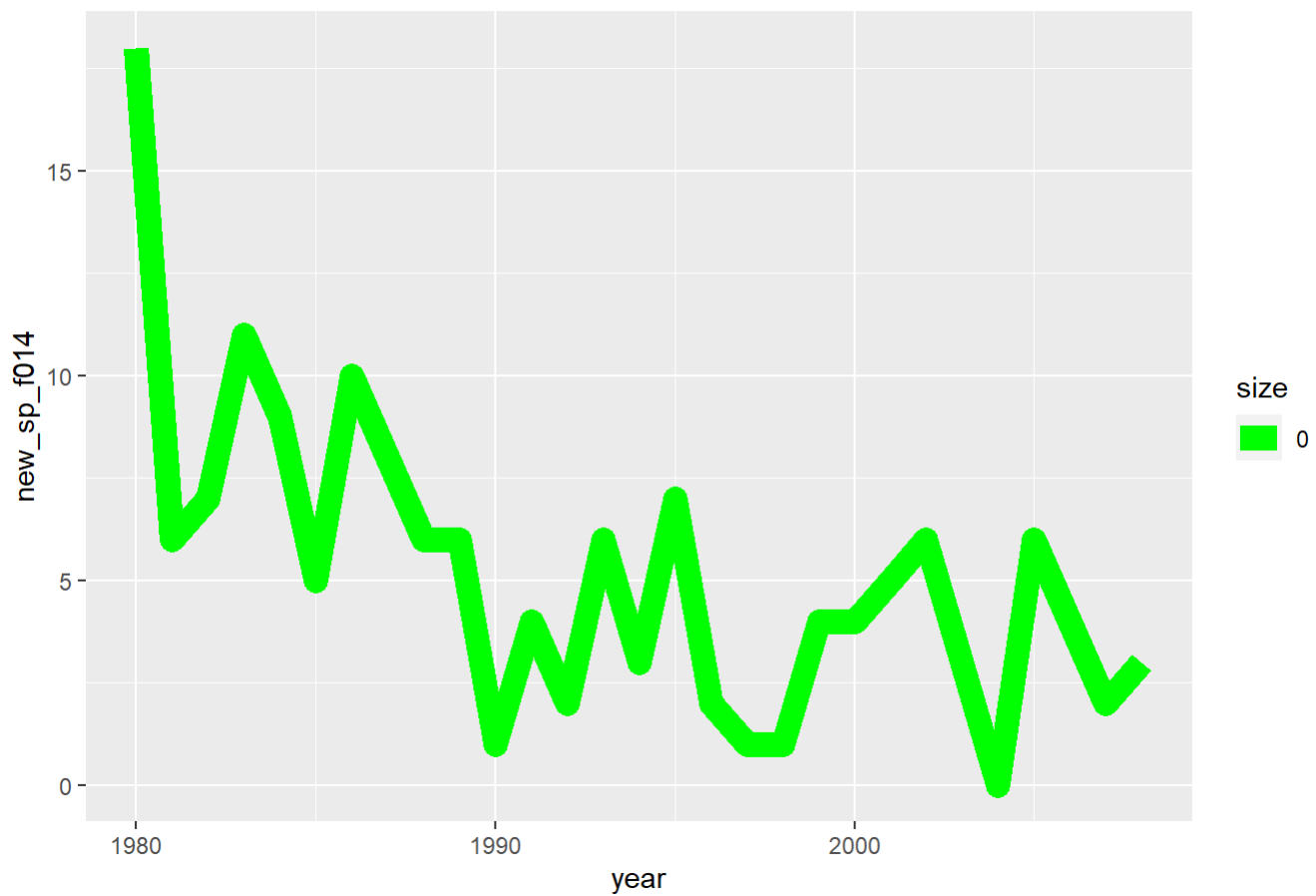
Histogram Males 0-14 with Positive Cases



K. Next, create similar graphs for each of the other three variables. Change the **color** of the line plots (any color you want).

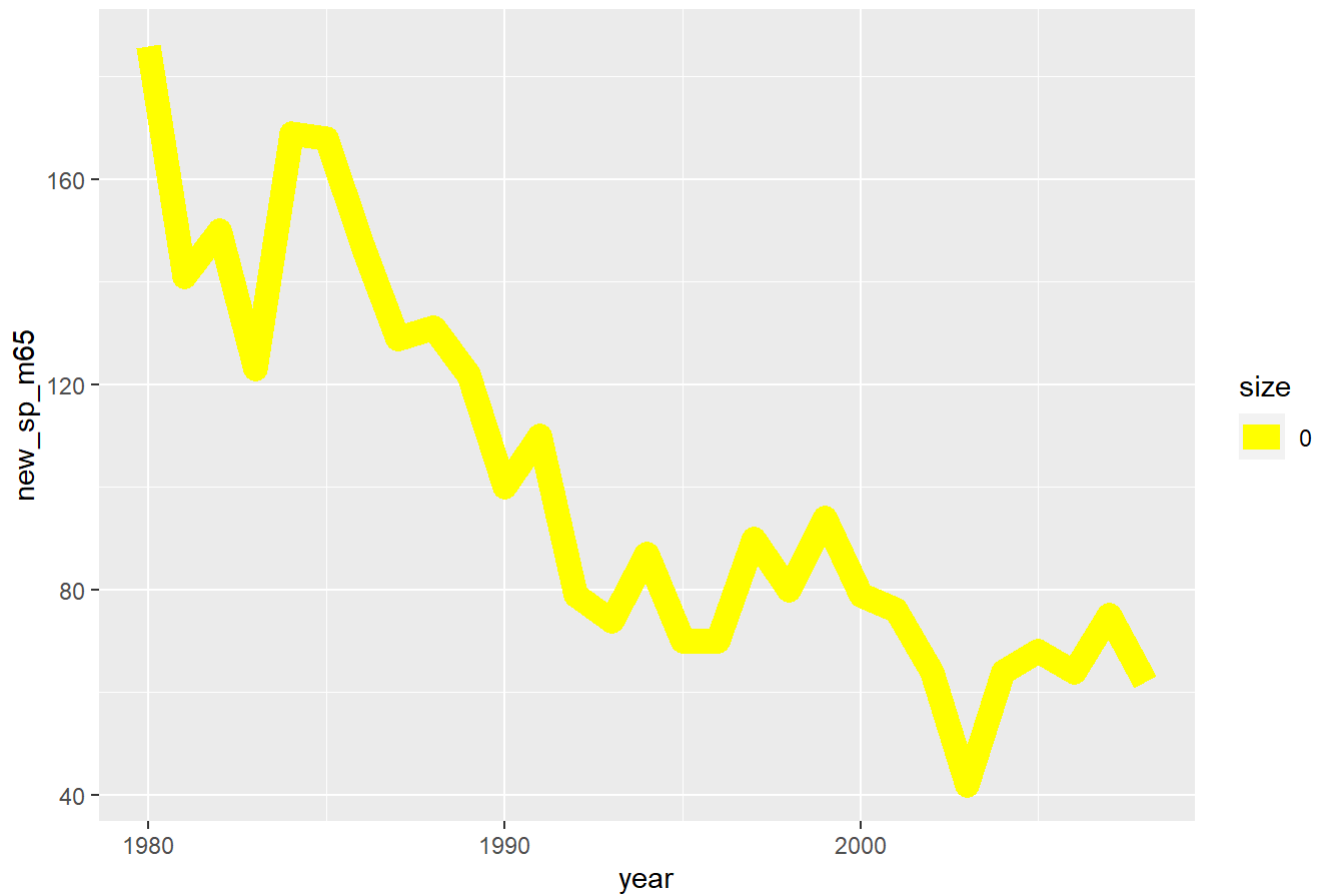
```
tbCan %>% ggplot() +  
  geom_line(color = 'green',  
    aes(x=year, y=new_sp_f014, size=0 )) +  
  ggtitle('Histogram Female 0-14 with Positive Cases')
```

Histogram Female 0-14 with Positive Cases



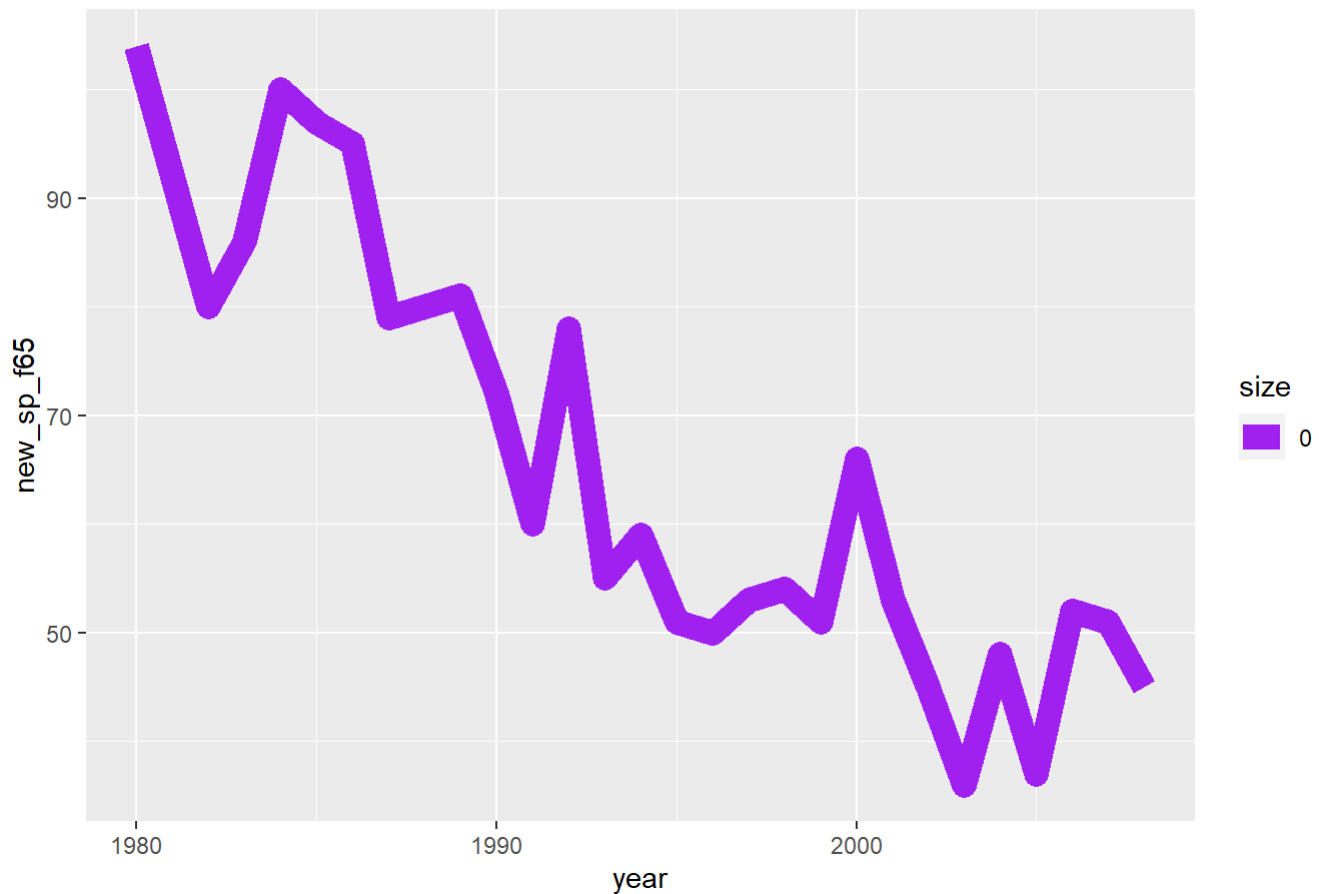
```
tbCan %>% ggplot() +  
  geom_line(color = 'yellow',  
    aes(x=year, y=new_sp_m65, size=0 )) +  
  ggtitle('Histogram Male 65+ with Positive Cases')
```

Histogram Male 65+ with Positive Cases



```
tbCan %>% ggplot() +  
  geom_line(color = 'purple',  
    aes(x=year, y=new_sp_f65, size=0 )) +  
  ggtitle('Histogram Female 65+ with Positive Cases')
```

Histogram Female 65+ with Positive Cases



- L. Using vector math, create a new variable by combining the numbers from **new\_sp\_m014** and **new\_sp\_f014**. Save the resulting vector as a new variable in the **tbCan** df called **new\_sp\_combined014**. This new variable represents the number of positive pulmonary smear tests for male AND female children between the ages of 0 and 14 years of age. Do the same for SP tests among citizens 65 years of age and older and save the resulting vector in the **tbCan** variable called **new\_sp\_combined65**.

```
tbCan$new_sp_combined014 <- (tbCan$new_sp_m014 + tbCan$new_sp_f014)
tbCan$new_sp_combined65 <- (tbCan$new_sp_m65 + tbCan$new_sp_f65)

show(tbCan$new_sp_combined014)
```

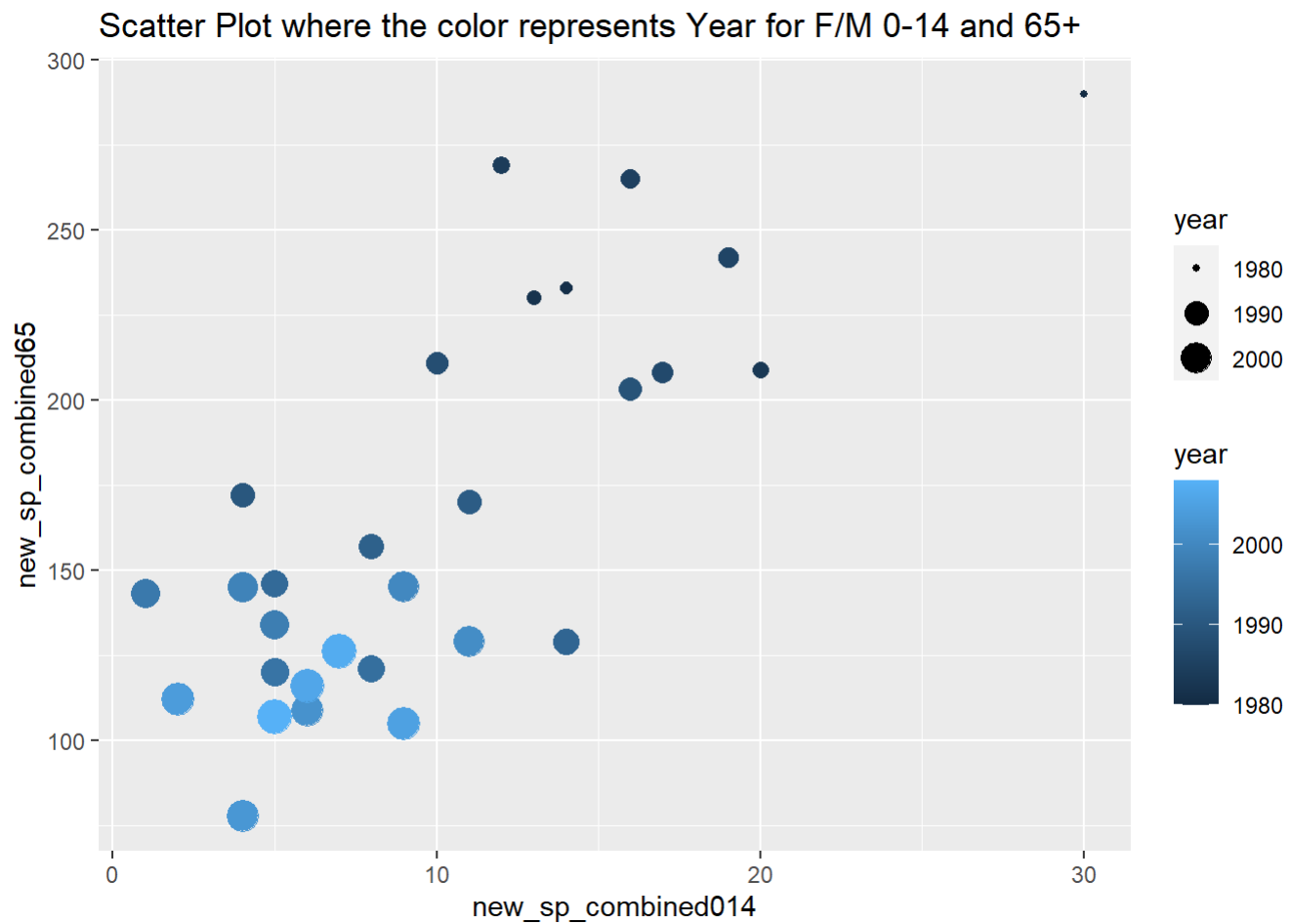
```
## [1] 30 14 13 20 12 16 19 17 10 16 4 11 8 14 5 8 5 1 5 4 9 11 6 4 2
## [26] 9 6 7 5
```

```
show(tbCan$new_sp_combined65)
```

```
## [1] 290 233 230 209 269 265 242 208 211 203 172 170 157 129 146 121 120 143 134
## [20] 145 145 129 109 78 112 105 116 126 107
```

- M. Finally, create a **scatter plot**, showing **new\_sp\_combined014** on the x axis, **new\_sp\_combined65** on the y axis, and having the **color and size** of the point represent **year**.

```
tbCan %>% ggplot() +
  geom_point() +
  aes(x=new_sp_combined014, y=new_sp_combined65, size=year, color=year ) +
  ggtitle('Scatter Plot where the color represents Year for F/M 0-14 and 65+')
```



N. Interpret this visualization – what insight does it provide?

*# The data shows that the Older People had more tests done, but the test count droppped as time increased.*