cm012 Exercises: Factors

Motivating the need for factors in R

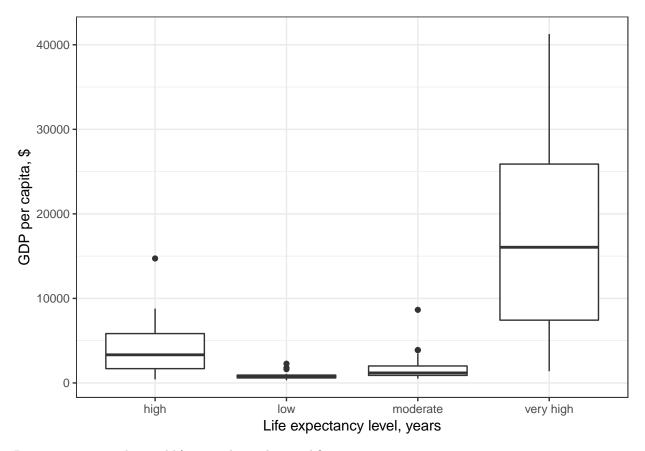
Activity 1: Using Factors for plotting

1.1 Let's look again into gapminder dataset and create a new cloumn, life_level, that contains five categories ("very high", "high", "moderate", "low" and "very low") based on life expectancy in 1997. Assign categories according to the table below:

Criteria	life_level
less than 23	very low
between 23 and 48	low
between 48 and 59	moderate
between 59 and 70	high
more than 70	very high

Function case_when() is a tidier way to vectorise multiple if_else() statements. you can read more about this function here.

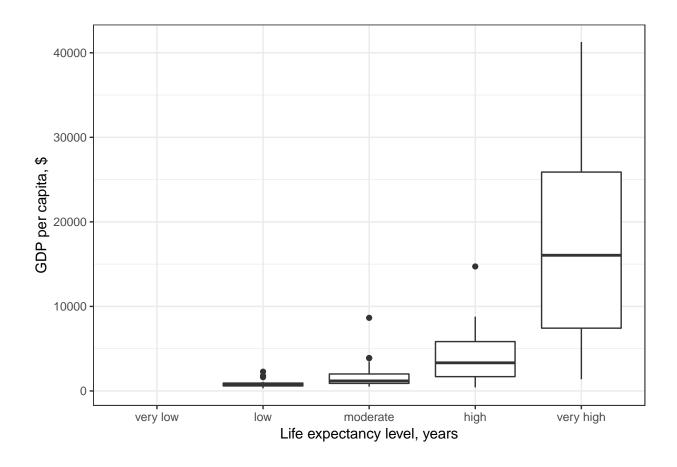
```
library(tidyverse)
## -- Attaching packages ----- tidyverse 1.2.1 --
## v ggplot2 3.0.0
                     v purrr
                                0.3.2
## v tibble 2.1.3
                      v dplyr
                                0.8.3
## v tidyr
           1.0.0
                      v stringr 1.3.1
## v readr
           1.3.1
                     v forcats 0.4.0
## Warning: package 'ggplot2' was built under R version 3.4.4
## Warning: package 'readr' was built under R version 3.4.4
## Warning: package 'purrr' was built under R version 3.4.4
## Warning: package 'stringr' was built under R version 3.4.4
## Warning: package 'forcats' was built under R version 3.4.4
## -- Conflicts ----- tidyverse conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                    masks stats::lag()
library(gapminder)
gapminder %>%
 filter(year == 1997) %>%
 mutate(life_level = case_when(lifeExp < 23 ~ "very low",</pre>
                               lifeExp < 48 ~ "low",</pre>
                               lifeExp < 59 ~ "moderate",</pre>
                               lifeExp < 70 ~ "high",
                               TRUE ~ "very high")) %>%
 ggplot() + geom_boxplot(aes(x = life_level, y = gdpPercap)) +
 labs(y = "GDP per capita, $", x= "Life expectancy level, years") +
 theme_bw()
```



Do you notice anything odd/wrong about the graph?

We can make a few observations:

- It seems that none of the countries had a "very low" life-expectancy in 1997.
- However, since it was an option in our analysis it should be included in our plot. Right?
- Notice also how levels on x-axis are placed in the "wrong" order.
- 1.2 You can correct these issues by explicitly setting the levels parameter in the call to factor(). Use, drop= FALSE to tell the plot not to drop unused levels



Inspecting factors (activity 2)

In Activity 1, we created our own factors, so now let's explore what categorical variables that we have in the gapminder dataset.

Exploring gapminder\$continent (activity 2.1)

Use functions such as str(), levels(), nlevels() and class() to answer the following questions:

- what class is continent(a factor or charecter)?
- How many levels? What are they?
- What integer is used to represent factor "Asia"?

```
gapminder$continent %>%
  class()
## [1] "factor"
gapminder$continent %>%
 levels()
                   "Americas" "Asia"
## [1] "Africa"
                                          "Europe"
                                                     "Oceania"
gapminder$continent %>%
  nlevels()
```

```
gapminder$continent %>%
str()
```

```
## Factor w/ 5 levels "Africa", "Americas", ...: 3 3 3 3 3 3 3 3 3 3 ...
```

Exploring gapminder\$country (activity 2.2)

Let's explore what else we can do with factors:

Answer the following questions:

- How many levels are there in country?
- Filter gapminder dataset by 5 countries of your choice. How many levels are in your filtered dataset?

```
h_countries <- c("Canada", "Brazil", "France", "Bolivia", "Russia")
gap <- gapminder %%
filter(country %in% h_countries)</pre>
```

Dropping unused levels

What if we want to get rid of some levels that are "unused" - how do we do that?

The function droplevels() operates on all the factors in a data frame or on a single factor. The function forcats::fct_drop() operates on a factor.

```
h_gap_dropped <- gap %>%
  droplevels()
h_gap_dropped %>%
  nlevels()
```

[1] 0

Changing the order of levels

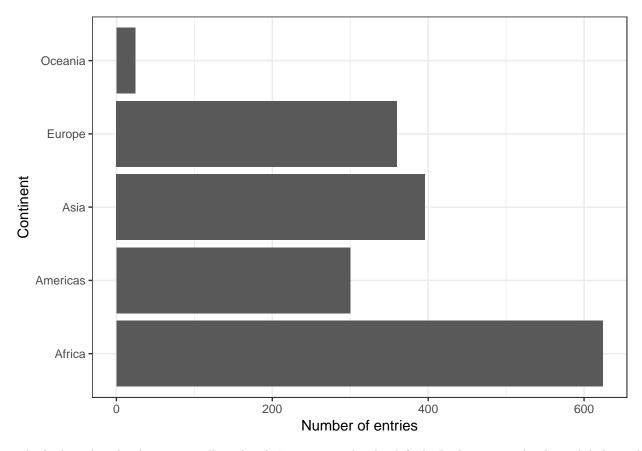
Let's say we wanted to re-order the levels of a factor using a new metric - say, count().

We should first produce a frequency table as a tibble using dplyr::count():

```
gapcount <- gapminder %>%
count(continent)
```

The table is nice, but it would be better to visualize the data. Factors are most useful/helpful when plotting data. So let's first plot this:

```
gapminder %>%
  ggplot() +
  geom_bar(aes(continent)) +
  coord_flip() +
  theme_bw() +
  ylab("Number of entries") + xlab("Continent")
```

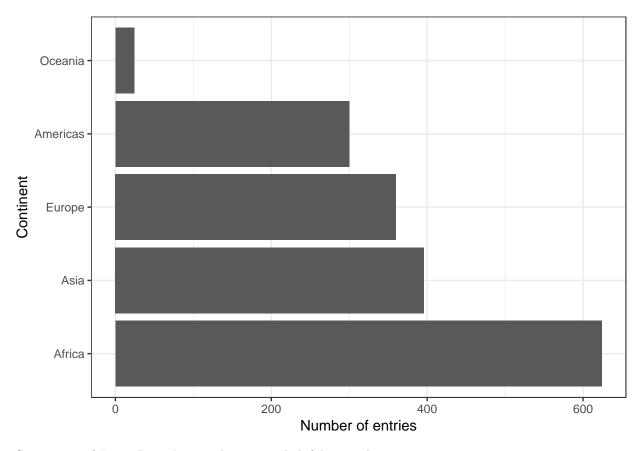


Think about how levels are normally ordered. It turns out that by default, R always sorts levels in alphabetical order. However, it is preferable to order the levels according to some principle:

- 1. Frequency/count.
- Make the most common level the first and so on. Function fct_infreq() might be useful.
- The function fct_rev() will sort them in the opposite order.

For instance, '

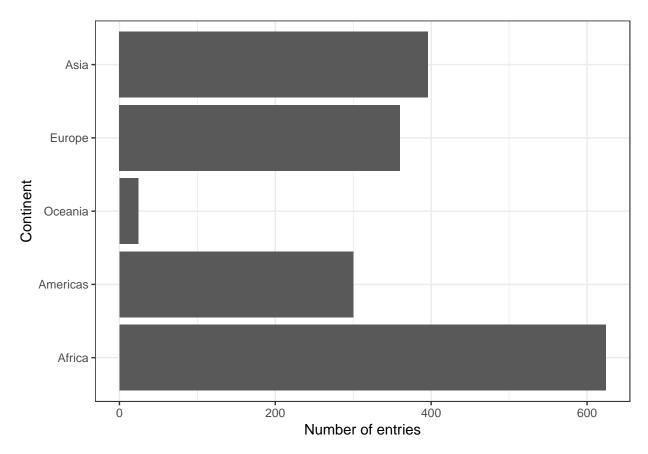
```
gapminder %%
ggplot() +
geom_bar(aes(fct_infreq(continent))) +
coord_flip()+
theme_bw() +
ylab("Number of entries") + xlab("Continent")
```



Section 9.6 of Jenny Bryan's notes have some helpful examples.

- 2. Another variable.
- For example, if we wanted to bring back our example of ordering gapminder countries by life expectancy, we can visualize the results using fct_reorder().

```
## default summarizing function is median()
gapminder %>%
    ggplot() +
    geom_bar(aes(fct_reorder(continent, lifeExp, max))) +
    coord_flip()+
    theme_bw() +
    xlab("Continent") + ylab("Number of entries")
```



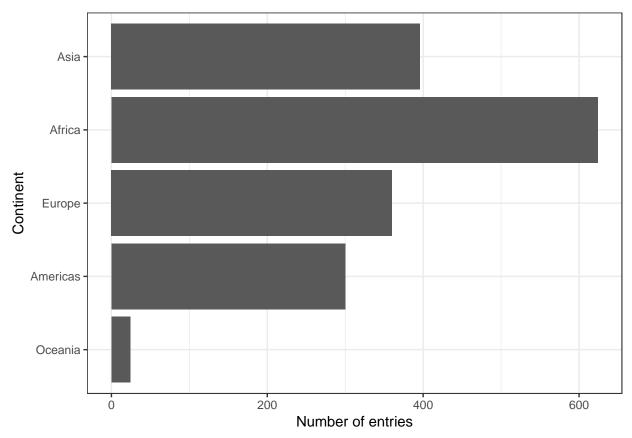
Use fct_reorder2() when you have a line chart of a quantitative x against another quantitative y and your factor provides the color.

 $\begin{tabular}{ll} ## Error in ggplot(h_gap, aes(x = year, y = lifeExp, color = FILL_IN_THIS)): object 'h_gap' not found in the property of the property o$

Change order of the levels manually

This might be useful if you are preparing a report for say, the state of affairs in Africa.

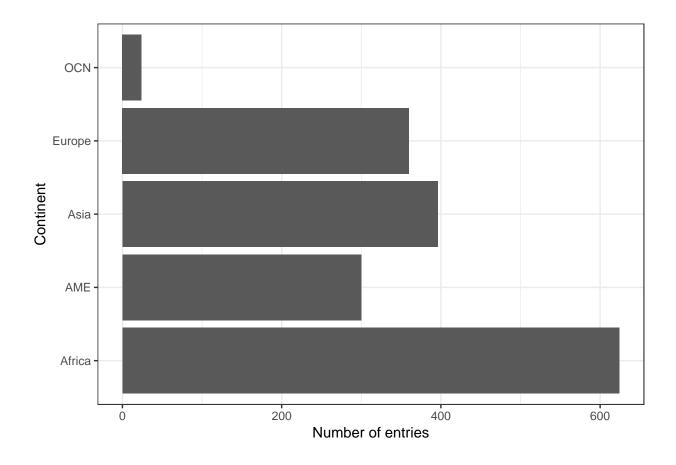
```
gapminder %>%
  ggplot() +
  geom_bar(aes(fct_relevel(continent, "Oceania", "Americas", "Europe"))) +
  coord_flip()+
  theme_bw() +
  xlab("Continent") + ylab("Number of entries")
```



More details on reordering factor levels by hand can be found [here]https://forcats.tidyverse.org/reference/fct_relevel.html

Recoding factors

Sometimes you want to specify what the levels of a factor should be. For instance, if you had levels called "blk" and "brwn", you would rather they be called "Black" and "Brown" - this is called recoding. Lets recode Oceania and the Americas in the graph above as abbreviations OCN and AME respectively using the function fct_recode().



Grow a factor (OPTIONAL)

Let's create two data frames,df1 and df2 each with data from two countries, dropping unused factor levels. The country factors in df1 and df2 have different levels. Can we just combine them?

The country factors in df1 and df2 have different levels. Can you just combine them using c()?

Explore how different forms of row binding work behave here, in terms of the country variable in the result.