

gapminder-exploration-phase2

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```
suppressPackageStartupMessages(library(tidyverse)) # The tidyverse contains ggplot2!
suppressPackageStartupMessages(library(gapminder))
knitr::opts_chunk$set(fig.width=8, fig.height=6)
```

Smell test of data

Here are some ways to explore some characteristics of the gapminder dataset.

Using the following functions, I have been able to determine some of the gapminder dataset characteristics.

```
class(gapminder)
```

```
## [1] "tbl_df"      "tbl"        "data.frame"
```

```
typeof(gapminder)
```

```
## [1] "list"
```

```
str(gapminder)
```

```
## Classes 'tbl_df', 'tbl' and 'data.frame':   1704 obs. of  6 variables:
## $ country   : Factor w/ 142 levels "Afghanistan",...: 1 1 1 1 1 1 1 1 1 ...
## $ continent: Factor w/ 5 levels "Africa","Americas",...: 3 3 3 3 3 3 3 3 3 ...
## $ year      : int   1952 1957 1962 1967 1972 1977 1982 1987 1992 1997 ...
## $ lifeExp   : num   28.8 30.3 32 34 36.1 ...
## $ pop       : int  8425333 9240934 10267083 11537966 13079460 14880372 12881816 13867957 16317921 22...
## $ gdpPercap: num   779 821 853 836 740 ...
```

I found the Help documentation very useful to understand better the concept of an object in R coding.

```
?class
?tbl_df
?tbl
?data.frame
?typeof
?str
```

Therefore, I now understand that gapminder is a tibble. It seems like using tibbles over classic data.frame() has several advantages, like avoiding problems such as accidental conversion of string as factor. I read this interesting wiki page on the topic.

The gapminder data set has 3 classes

tbl_df

tbl

**data.frame*

I found class() to be the most informative function for me

It revealed the specific data type of each variable!

country : Factor

continent: Factor

year : int

```
lifeExp : num
pop : int
gdpPercap : num
```

The class function told me that there were 6 variables (columns) and 1704 observations (rows) in gapminder dataset, but I could also use these:

```
ncol(gapminder)
```

```
## [1] 6
```

```
nrow(gapminder)
```

```
## [1] 1704
```

Here are some other useful functions to learn more about the dataset. If you have a specific question about the dataset, you can use more specific functions as well, like `dim()` which provides the dimension of the data tibble.

```
dim(gapminder)
```

```
## [1] 1704    6
```

Explore individual variables

To ensure I address all the requirements of this assignment, I have decided to include the STAT545 homework 2 questions above my answers in bold.

Q? What are possible values (or range, whichever is appropriate) of each variable?

I have used the `summary()` function to know the possible values and range of each variable.

```
overview <- summary(gapminder)
overview
```

```
##      country      continent      year      lifeExp
## Afghanistan: 12 Africa :624  Min.   :1952  Min.   :23.60
## Albania      : 12 Americas:300 1st Qu.:1966 1st Qu.:48.20
## Algeria       : 12 Asia    :396 Median :1980 Median :60.71
## Angola        : 12 Europe  :360 Mean    :1980 Mean   :59.47
## Argentina     : 12 Oceania : 24 3rd Qu.:1993 3rd Qu.:70.85
## Australia     : 12                Max.   :2007 Max.   :82.60
## (Other)       :1632
##      pop      gdpPercap
## Min.   :6.001e+04  Min.    : 241.2
## 1st Qu.:2.794e+06  1st Qu.: 1202.1
## Median :7.024e+06  Median : 3531.8
## Mean    :2.960e+07  Mean    : 7215.3
## 3rd Qu.:1.959e+07  3rd Qu.: 9325.5
## Max.    :1.319e+09  Max.    :113523.1
##
```

Or the prettier version with `knitr::kable`

```
knitr::kable(overview,
  format = "pandoc",
  digits = 2,
  caption = "Summary Stats - Gapminder dataset",
  align = c("l"),
```

```
longtable = TRUE,
padding = 5)
```

Table 1: Summary Stats - Gapminder dataset

country	continent	year	lifeExp	pop	gdpPercap
Afghanistan: 12	Africa :624	Min. :1952	Min. :23.60	Min. :6.001e+04	Min. : 241.2
Albania : 12	Americas:300	1st Qu.:1966	1st Qu.:48.20	1st Qu.:2.794e+06	1st Qu.: 1202.1
Algeria : 12	Asia :396	Median :1980	Median :60.71	Median :7.024e+06	Median : 3531.8
Angola : 12	Europe :360	Mean :1980	Mean :59.47	Mean :2.960e+07	Mean : 7215.3
Argentina : 12	Oceania : 24	3rd Qu.:1993	3rd Qu.:70.85	3rd Qu.:1.959e+07	3rd Qu.: 9325.5
Australia : 12	NA	Max. :2007	Max. :82.60	Max. :1.319e+09	Max. :113523.1
(Other) :1632	NA	NA	NA	NA	NA

Another way is to use individual functions to obtain the information. Below, these functions respectively informs on the minimum value, maximum value, range of value (minimum and maximum) and all possible distinct values of specific columns.

```
min(gapminder$lifeExp)
```

```
## [1] 23.599
```

```
max(gapminder$lifeExp)
```

```
## [1] 82.603
```

```
range(gapminder$gdpPercap)
```

```
## [1] 241.1659 113523.1329
```

```
distinct(gapminder, country)
```

```
## # A tibble: 142 x 1
```

```
##   country
```

```
##   <fctr>
```

```
## 1 Afghanistan
```

```
## 2 Albania
```

```
## 3 Algeria
```

```
## 4 Angola
```

```
## 5 Argentina
```

```
## 6 Australia
```

```
## 7 Austria
```

```
## 8 Bahrain
```

```
## 9 Bangladesh
```

```
## 10 Belgium
```

```
## # ... with 132 more rows
```

I also found this website useful.

Q? What values are typical? What's the spread? What's the distribution? Etc., tailored to the variable at hand. Feel free to use summary stats, tables, figures. We're NOT expecting high production value (yet).

If we group data according to their column, another useful function is summarise() to get some statistics on a variable, but data needs to be grouped before doing so. For example, if we group by continent, we can use the function to know the mean year of entry, lifeExp, pop and gdpPercap for each continent.

```
gapminder %>%
  group_by(continent) %>%
  summarise_if(is.numeric, mean, na.rm=TRUE)
```

```
## # A tibble: 5 x 5
##   continent  year  lifeExp      pop gdpPercap
##   <fctr>    <dbl>    <dbl>    <dbl>    <dbl>
## 1 Africa  1979.5  48.86533  9916003  2193.755
## 2 Americas 1979.5  64.65874 24504795  7136.110
## 3 Asia    1979.5  60.06490 77038722  7902.150
## 4 Europe  1979.5  71.90369 17169765 14469.476
## 5 Oceania 1979.5  74.32621  8874672 18621.609
```

Again, I find that the most useful function is `summary()`, as it provides the range (minimum, maximum), and summary statistics such as 1st quartile, mean and 3rd quartile (see above for data).

```
gapminder %>%
  group_by(country) %>%
  summarise_if(is.numeric, median)
```

```
## # A tibble: 142 x 5
##   country  year  lifeExp      pop gdpPercap
##   <fctr>    <dbl>    <dbl>    <dbl>    <dbl>
## 1 Afghanistan 1979.5  39.1460 13473708.5   803.4832
## 2 Albania    1979.5  69.6750  2644572.5  3253.2384
## 3 Algeria    1979.5  59.6910 18593278.5  4853.8559
## 4 Angola     1979.5  39.6945  6589529.5  3264.6288
## 5 Argentina  1979.5  69.2115 28162601.0  9068.7844
## 6 Australia  1979.5  74.1150 14629150.0 18905.6034
## 7 Austria    1979.5  72.6750  7571521.5 20673.2530
## 8 Bahrain    1979.5  67.3225  337688.5 18779.8016
## 9 Bangladesh 1979.5  48.4660 86751356.0   703.7638
## 10 Belgium   1979.5  73.3650  9839051.5 20048.9102
## # ... with 132 more rows
```

Explore various plot types

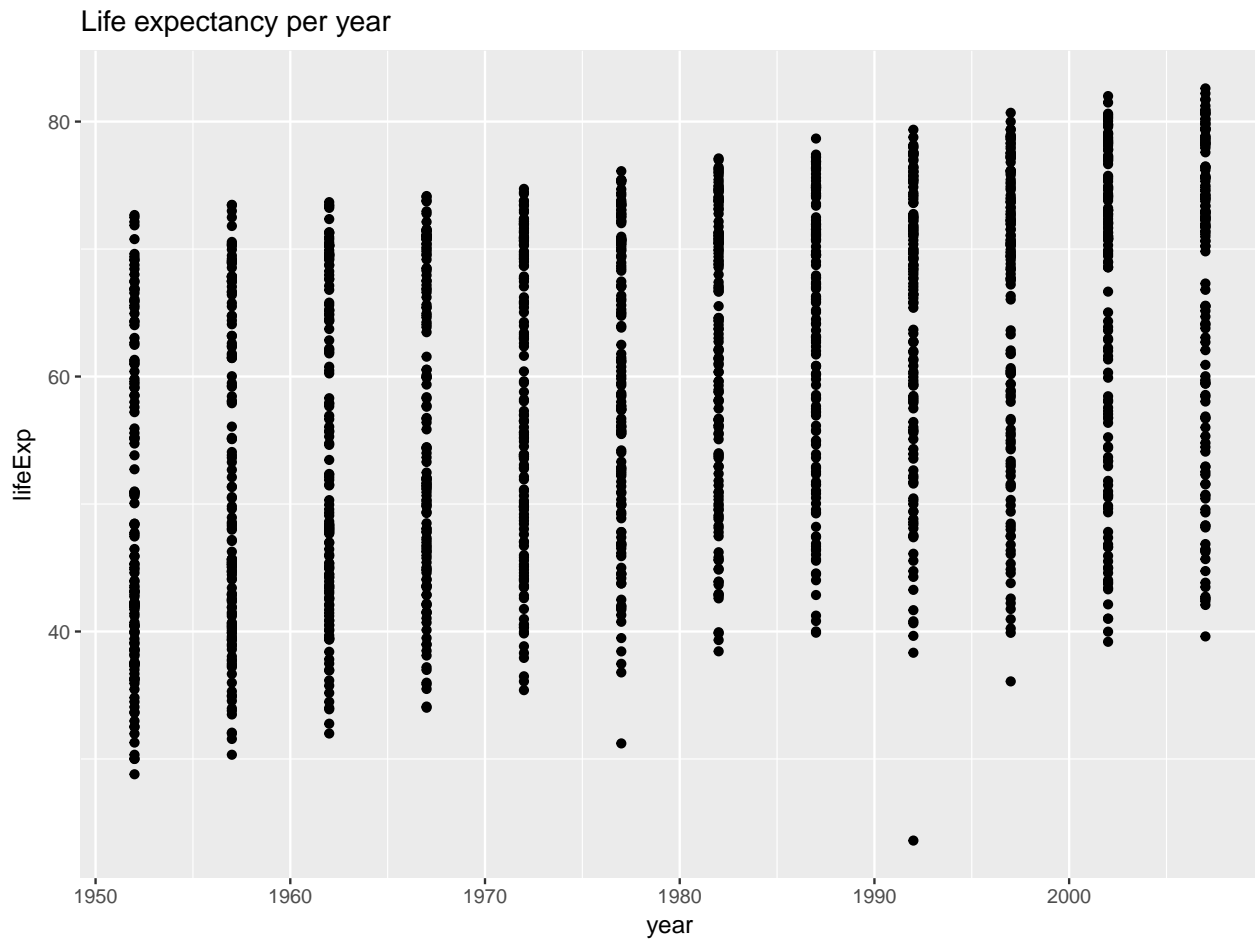
NOTE: The main resource used is the [stat545 github repository](#) of Jenny Bryan

POINT PLOTS

Now, I will explore diverse plot types.

Let's start with a basic data point plot of the life expectancy relative to the year of assesment for each country.

```
p <- gapminder %>%
  group_by(country) %>%
  ggplot(aes(x=year, y=lifeExp))
p + geom_point() + labs(title = "Life expectancy per year")
```



Overall, it seems that life expectancy has improved with years for the majority of countries, but I wonder what are the outliers at the bottom of the plot.

Let us see what is the minimum life expectancy for each year to find out.

```
t <- gapminder %>%
  group_by(country, year, lifeExp, pop, gdpPercap) %>%
  summarise_if(is.numeric, min) %>%
  arrange(lifeExp)
t
```

```
## # A tibble: 1,704 x 5
## # Groups:   country, year, lifeExp, pop [1,704]
##   country year lifeExp pop gdpPercap
##   <fctr> <int> <dbl> <int> <dbl>
## 1 Rwanda 1992 23.599 7290203 737.0686
## 2 Afghanistan 1952 28.801 8425333 779.4453
## 3 Gambia 1952 30.000 284320 485.2307
## 4 Angola 1952 30.015 4232095 3520.6103
## 5 Sierra Leone 1952 30.331 2143249 879.7877
## 6 Afghanistan 1957 30.332 9240934 820.8530
## 7 Cambodia 1977 31.220 6978607 524.9722
## 8 Mozambique 1952 31.286 6446316 468.5260
## 9 Sierra Leone 1957 31.570 2295678 1004.4844
## 10 Burkina Faso 1952 31.975 4469979 543.2552
```

```
## # ... with 1,694 more rows
```

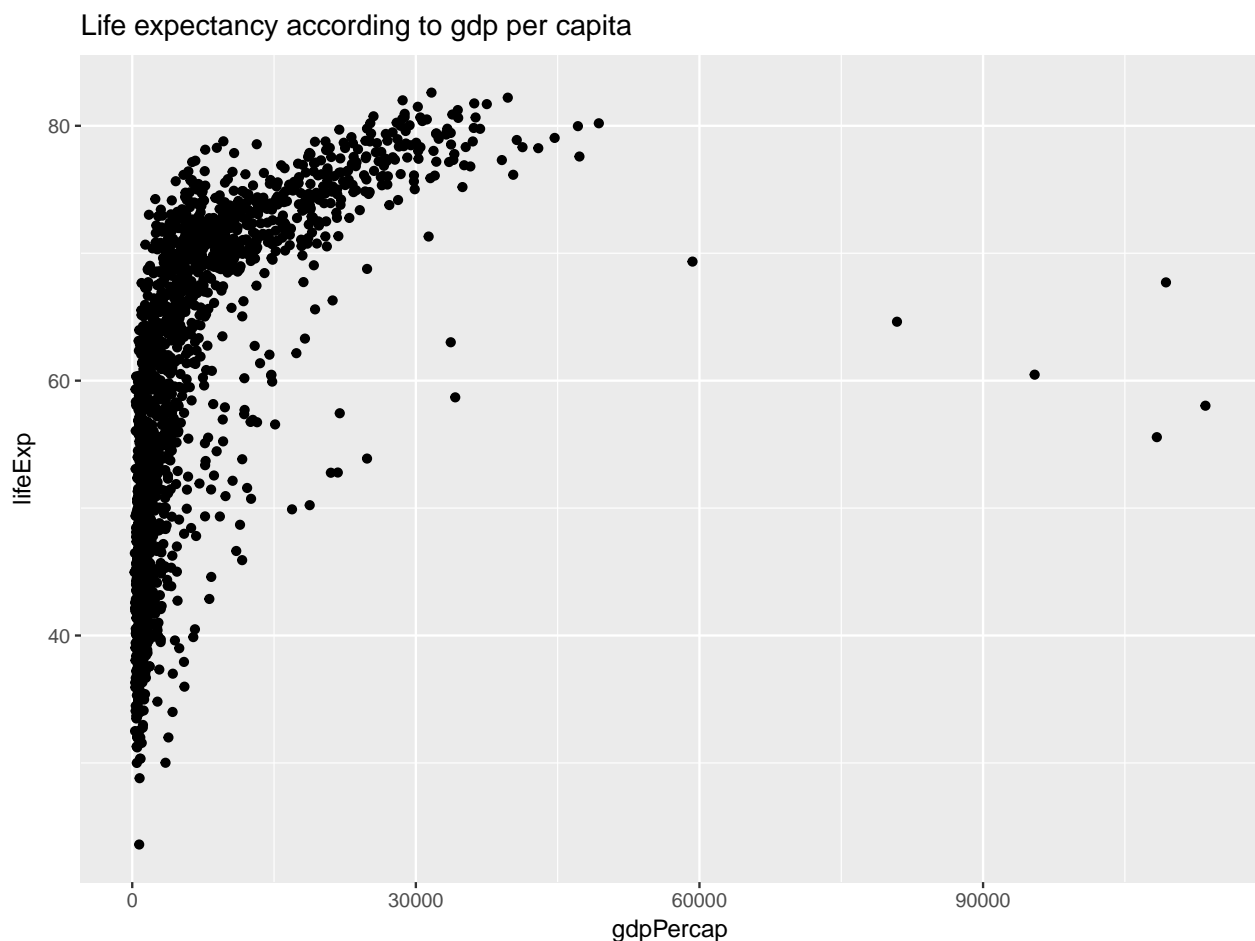
The lowest life expectancy recorder in the gapminder dataset is 23.5999 years in Rwanda in 1992, which corresponds to the tragedy of the Rwandan genocide.

Another way to find this outlier was thought to me by a student of STAT545 (thanks Alistair Barton!):

```
outlier_lifeExp <- filter(gapminder, lifeExp == min(lifeExp))
View(outlier_lifeExp)
```

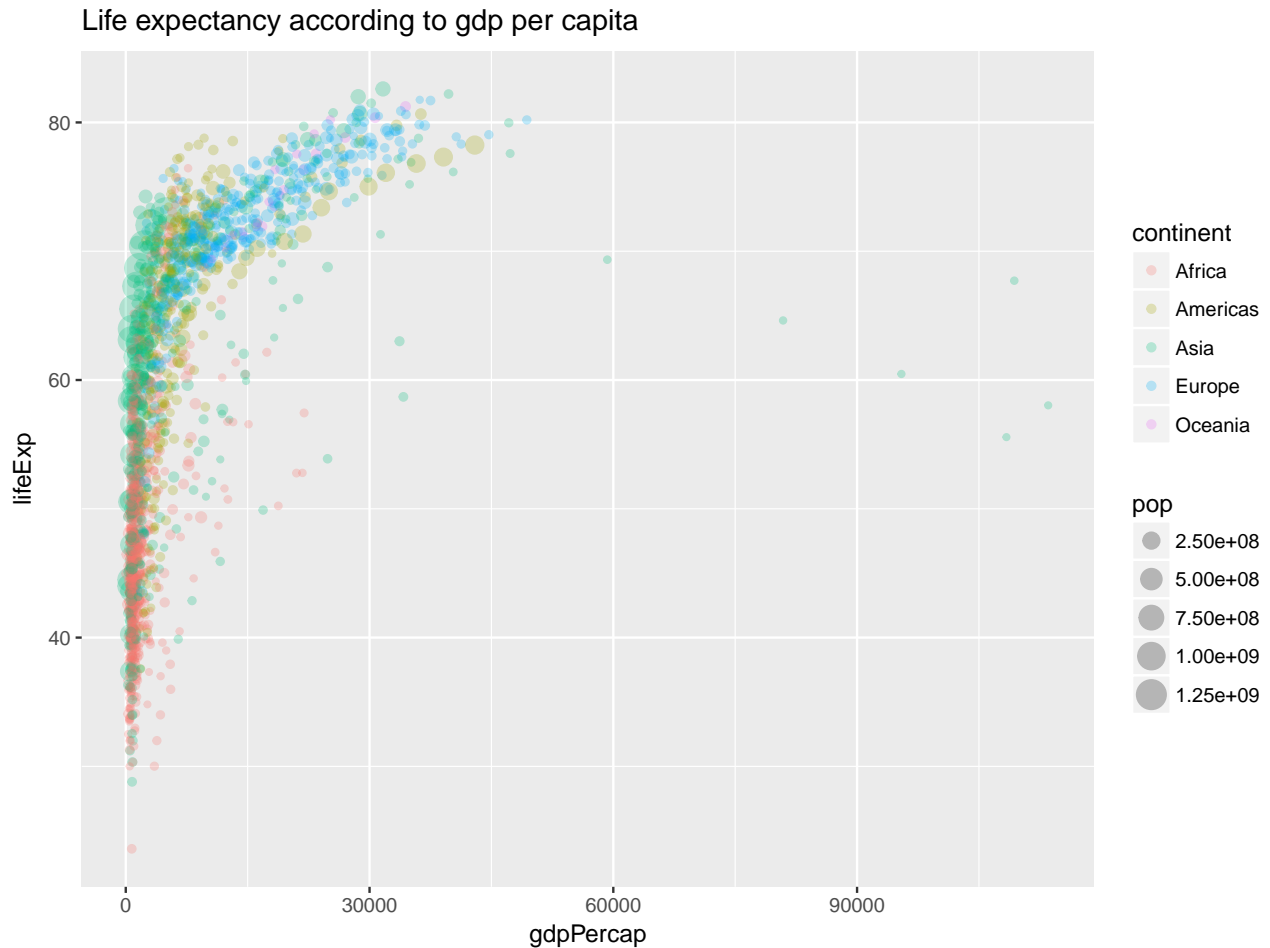
Let us carry on in our analysis. Of course, one may suspect there will be a relationship between the lifeExp and gdpPercap variables, so let's plot these.

```
p1 <- gapminder %>%
  group_by(country) %>%
  ggplot(aes(x=gdpPercap, y=lifeExp))+
  labs(title = "Life expectancy according to gdp per capita")
p1 + geom_point()
```



However, since I am still wondering if the size of the population has an effect on the life expectancy, I will plot the size of each dots according to the population, and I would also like to see if some continents segregate from each other, so I will use colours for this. I will also use some transparencies in order to see the dots when they layer on top of each other.

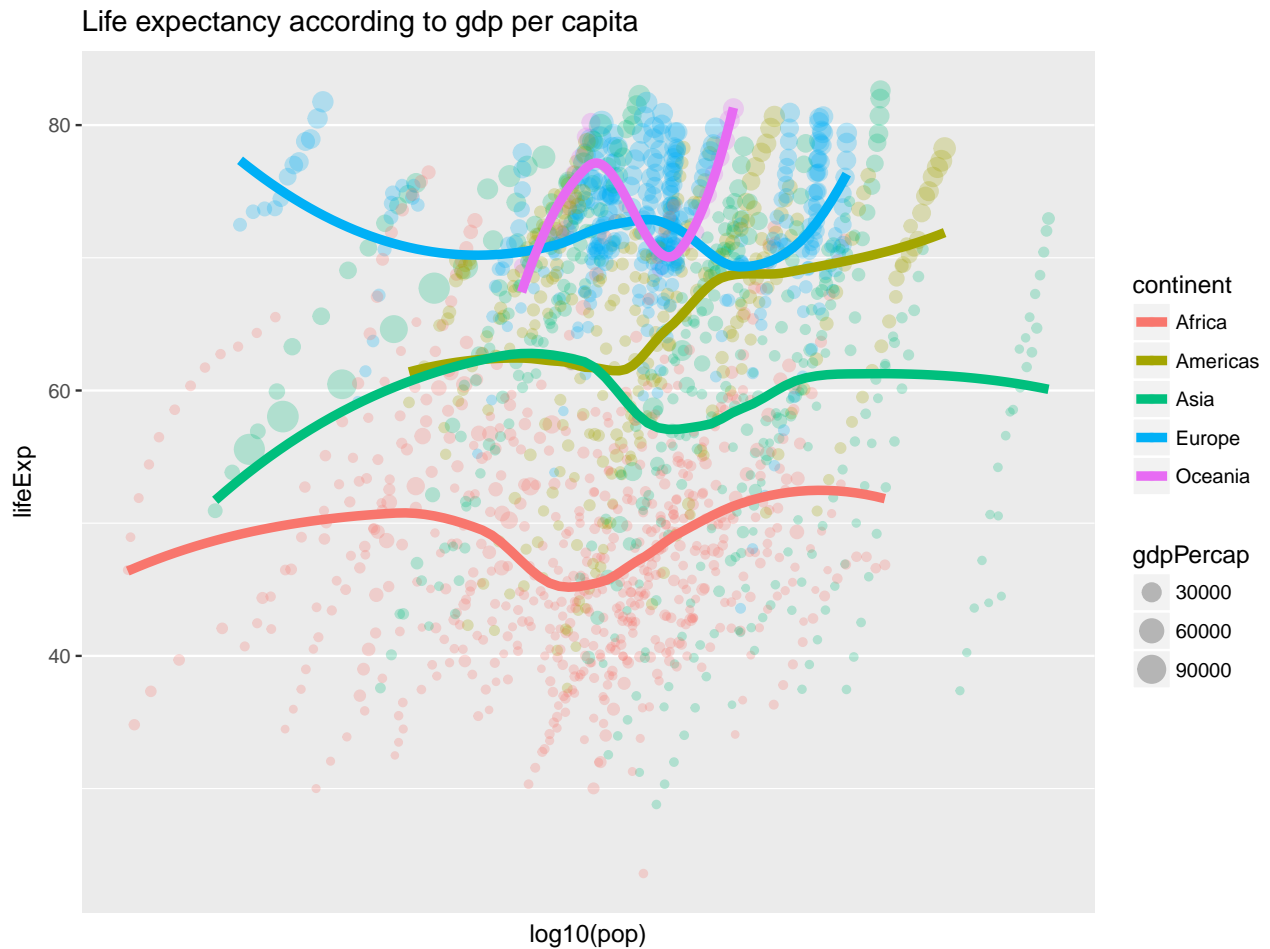
```
p1 + geom_point(aes(size=pop, colour = continent), alpha=0.25)
```



A log scale would be useful to see the lower values better. This resembles the exercise we completed in class last time, so let me switch up the variables a little to population and life expectancy, so that we don't feel like doing the same thing. I will plot the size of the dot as a function of the gdpPerCap this time and add the `geom_smooth()` line with a generalized additive model (gam).

```
p2 <- ggplot(gapminder, aes(x=log10(pop), y=lifeExp)) +
  geom_point(aes(colour=continent, size=gdpPerCap), alpha=0.25) +
  labs(title = "Life expectancy according to gdp per capita") +
  scale_x_log10() +
  geom_smooth(aes(colour = continent), lwd=2, se=FALSE)
p2
```

```
## `geom_smooth()` using method = 'loess'
```



We can immediately take note of obvious discrepancies when looking at this plot. For example, the red dots are at the lower part of the graph, telling us that countries of Africa have lower life expectancy, and the dots are smaller, revealing lower gdpPercap.

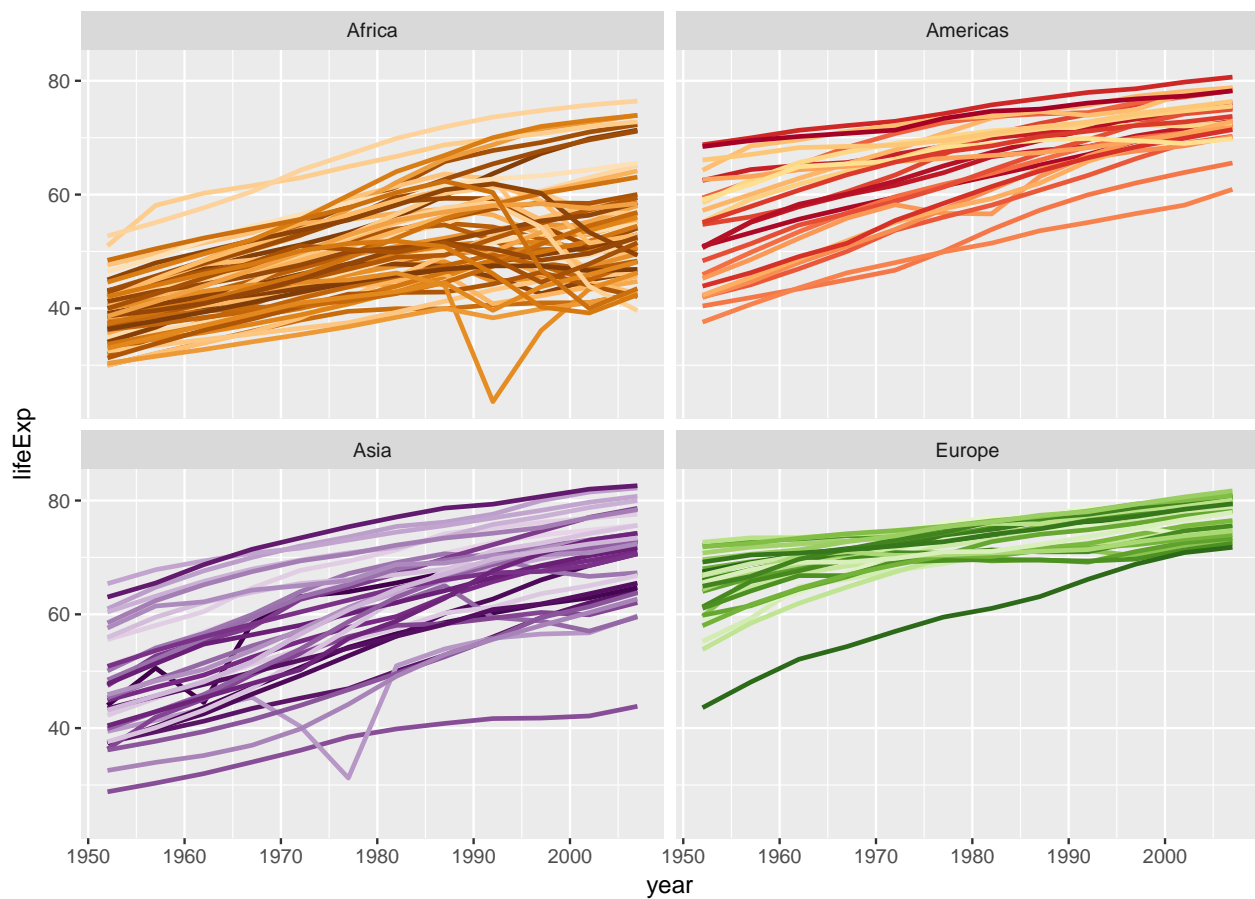
LINE PLOTS

Let's plot some lines tracking the life expectancy per year, but use the `facet_wrap()` function to divide the data by continent into 4 plots (omitting Oceania, given very little data).

```
p3 <- ggplot(subset(gapminder, continent != "Oceania"), aes(x=year, y=lifeExp, group = country, color =
p3 + geom_line(lwd=1, show_guide = FALSE) +
  facet_wrap(~continent) +
  scale_color_manual(values = country_colors) +
  labs(title = "Life expectancy per year and continent")
```

```
## Warning: `show_guide` has been deprecated. Please use `show.legend`
## instead.
```


Life expectancy per year and continent



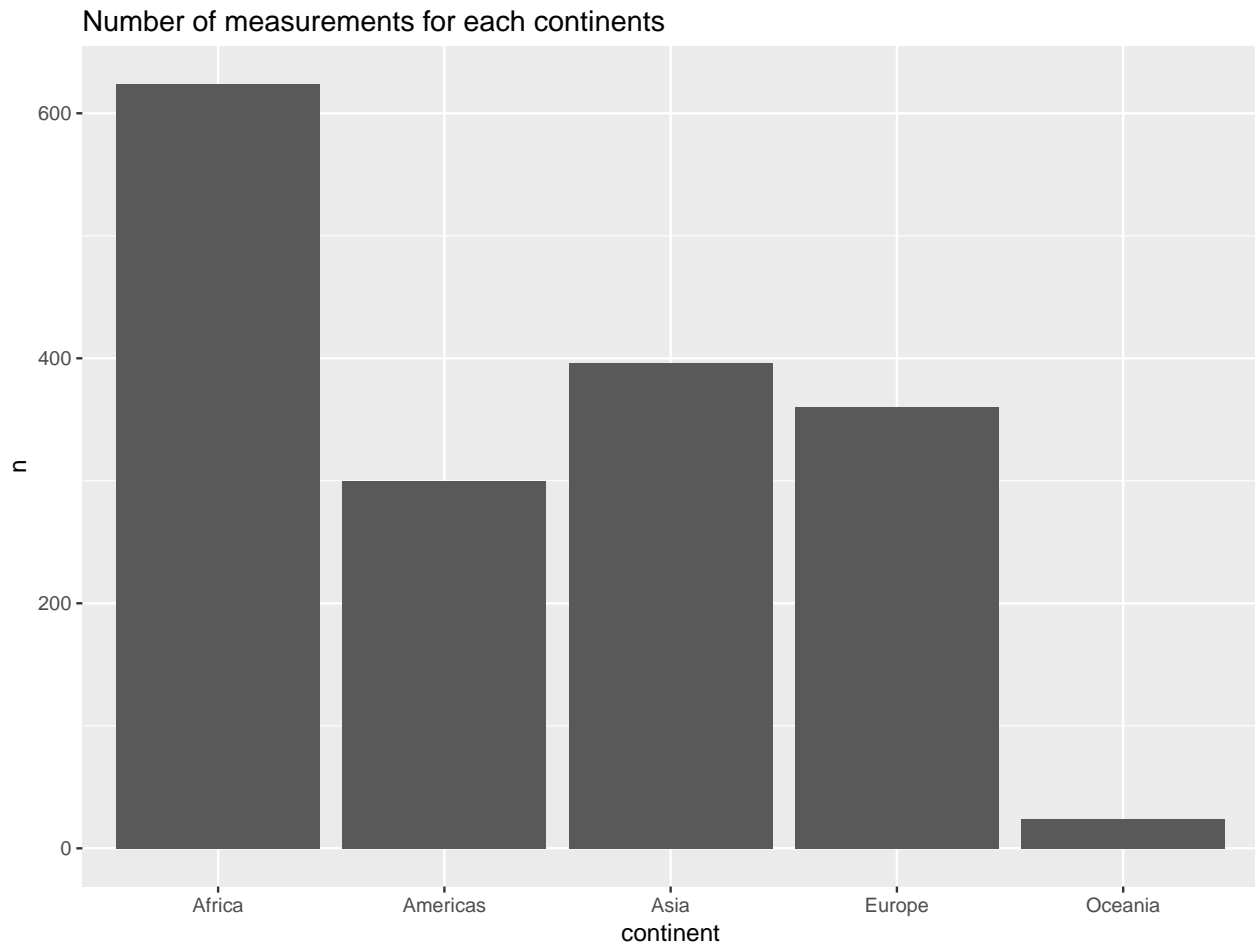
This is very pretty, but I don't know if it is very informative since this plots in a stepwise fashion the data, which collapses the values at each measurement year into a vertical line. I still kept it, for interest.

*I found the reference manual on the tidyverse website very useful.

BAR PLOTS

Let's go back to a plot of the life expectancy and gdp per capita to show a bar plot example.

```
my_gap <- gapminder
continent_freq <- gapminder %>% count(continent)
p4 <- continent_freq %>%
  ggplot(aes(x=continent, y=n))
p4 + geom_bar(stat="identity", na.rm=TRUE) +
  labs(title = "Number of measurements for each continents")
```



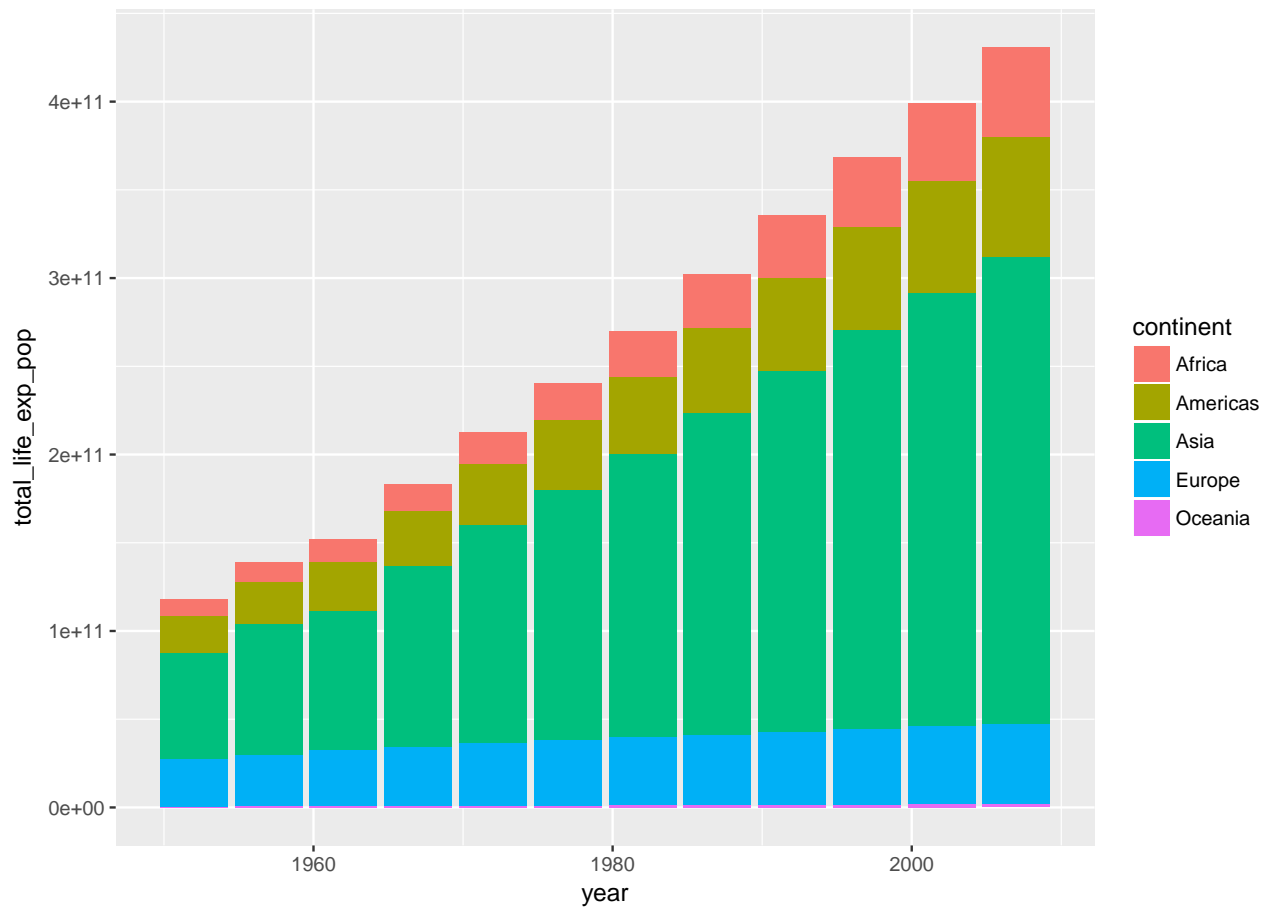
After doing the dataset manipulation exercises (see below), I decided to go back at my plots and change them for presenting more relevant information.

Here is an histogram of the “stacked up” life expectancies of all the countries (population x life expectancy) in each continent, plotted according to the year.

```
my_gap <- gapminder
my_gap_life <- my_gap %>%
  mutate(total_life_exp_pop = pop*lifeExp)
p5 <- my_gap_life %>%
  ggplot(aes(x=year, y=total_life_exp_pop))
p5 + geom_histogram(aes(fill = continent), stat="identity", na.rm=TRUE) +
  labs(title = "Cumulative life expectancy values per year, population and continent")
```

```
## Warning: Ignoring unknown parameters: binwidth, bins, pad
```

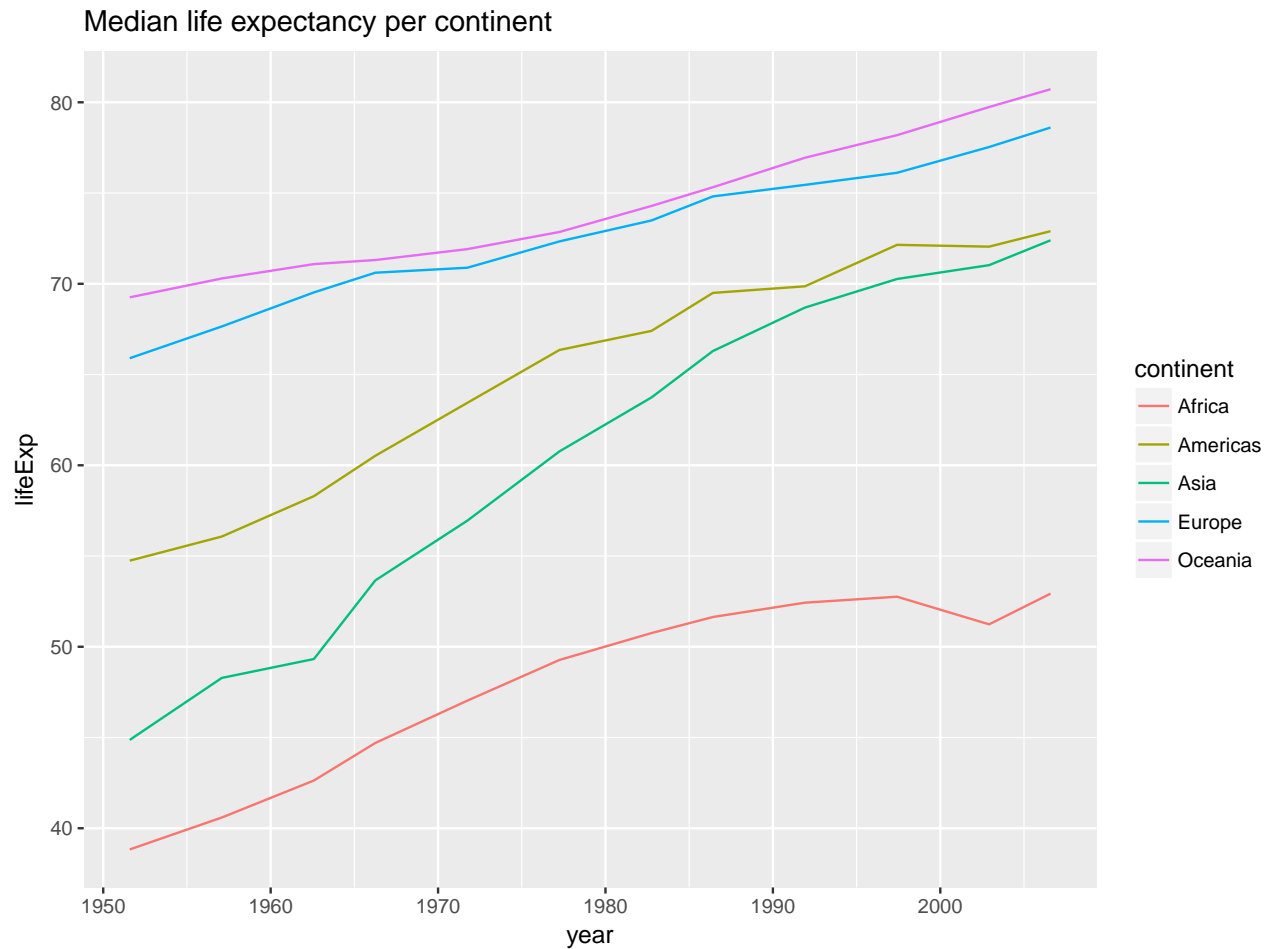
Cumulative life expectancy values per year, population and continent



SUMMARY STATISTICS PLOTS

Here is a way to plot some summary statistics for each continent. Below, you can see the median life expectancy for each continent according to the year.

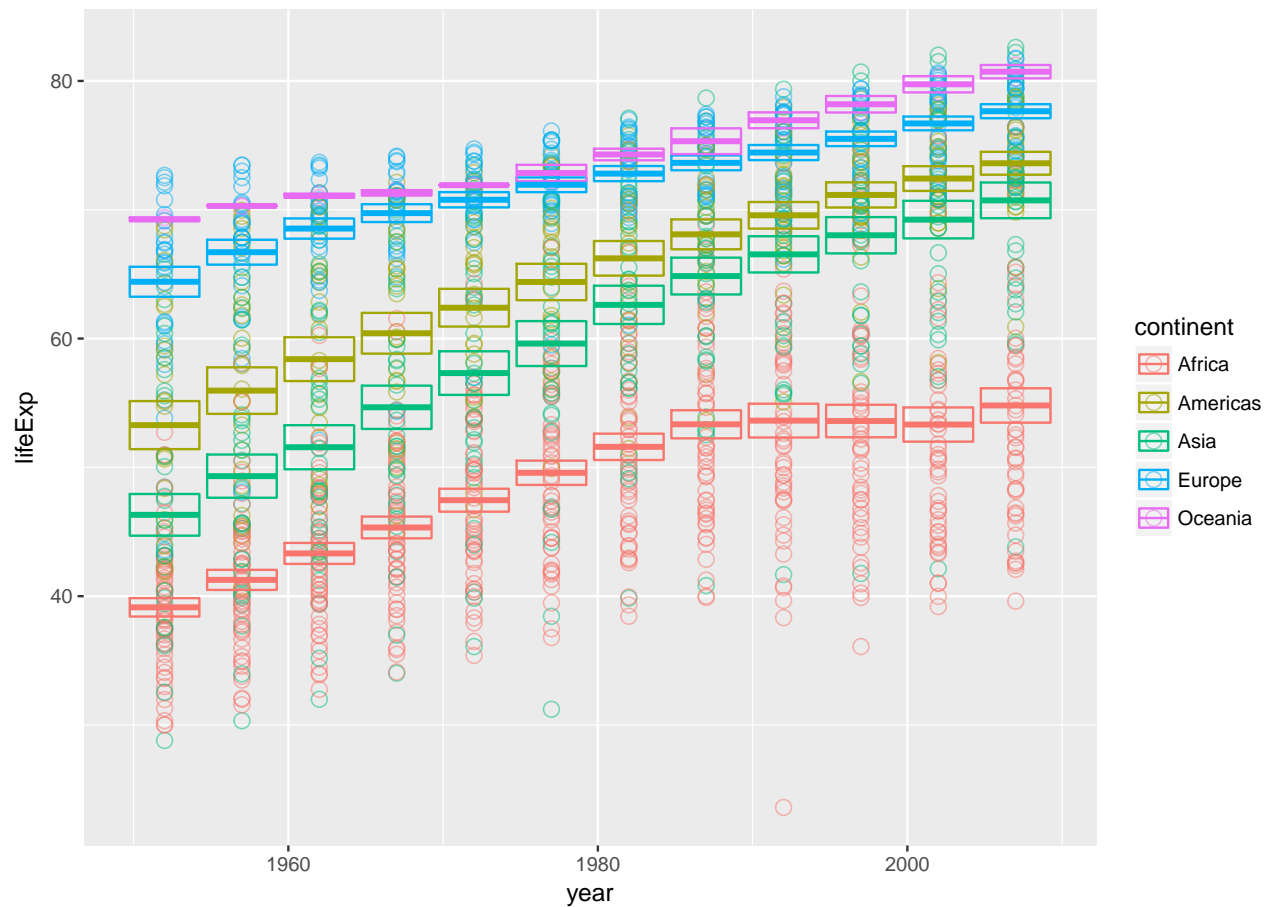
```
p6 <- gapminder %>%
  ggplot(aes(x=year, y=lifeExp))
p6 + stat_summary_bin(mapping = aes(group = continent, colour=continent), fun.y = "median", geom="line",
  labs(title = "Median life expectancy per continent")
```



You can do the same thing with other parameters, like below showing the mean in a b

```
p7 <- gapminder %>%
  ggplot(aes(x=year, y=lifeExp))
p7 +
  geom_point(mapping = aes(colour = continent), alpha=0.5, size=3, shape=21) +
  stat_summary(fun.data = "mean_se", mapping = aes(colour=continent), geom = "crossbar", na.rm=TRUE) +
  labs(title = "Mean life expectancy according to year")
```

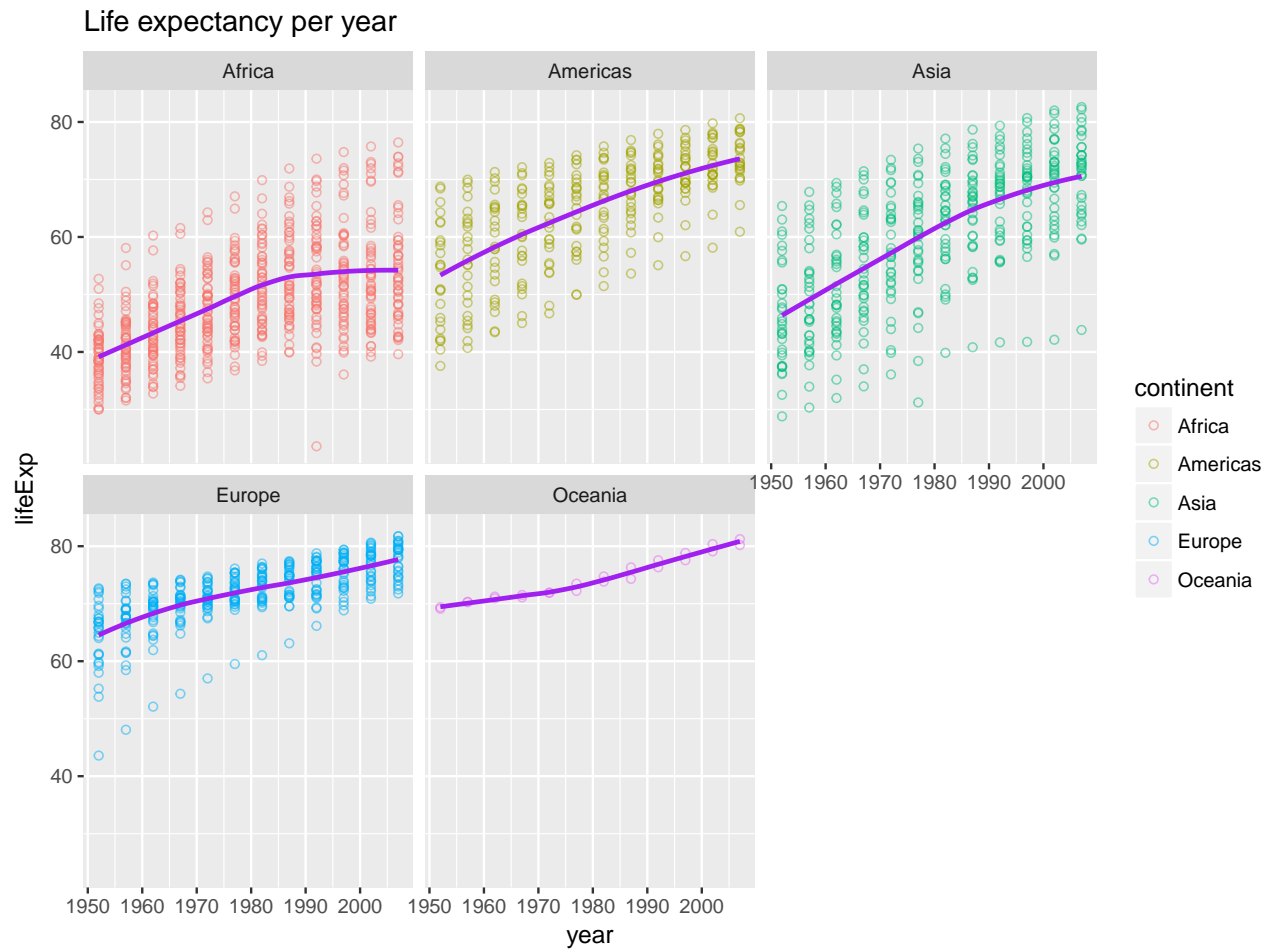
Mean life expectancy according to year



This plot is a bit crowded, we should subdivide it by continent and add a linear regression function to it.

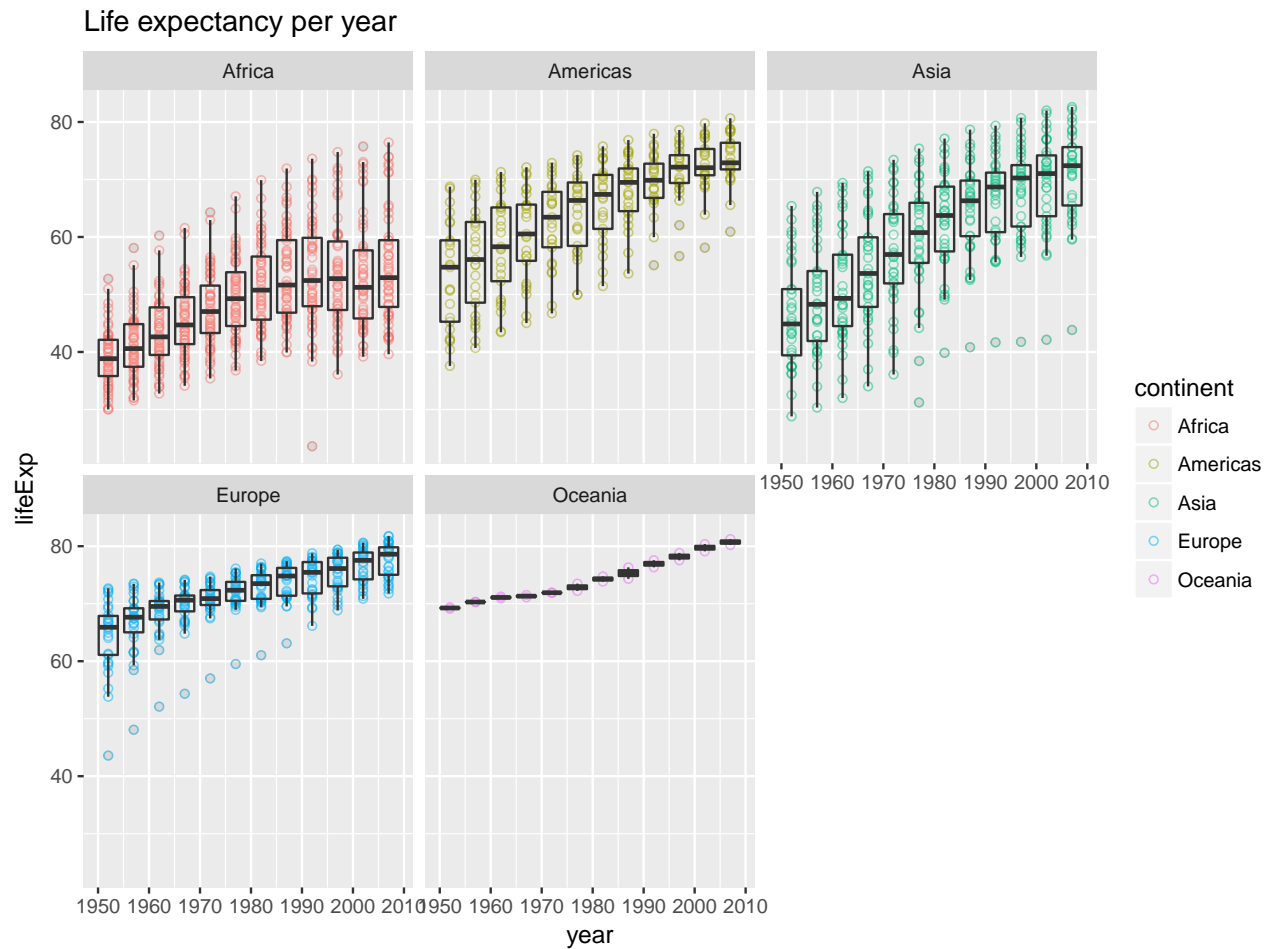
```
p8 <- gapminder %>%
  ggplot(aes(x=year, y=lifeExp)) +
  geom_point(mapping = aes(colour = continent), alpha=0.5, shape=21)
p8 + facet_wrap(~continent) + geom_smooth(se=FALSE, colour = "purple", lwd = 1, alpha = 0.8)+
  labs(title = "Life expectancy per year")

## `geom_smooth()` using method = 'loess'
```



Or we can show the same plot and plot the mean life expectancy (and standard deviation) with `geom_boxplot()` function:

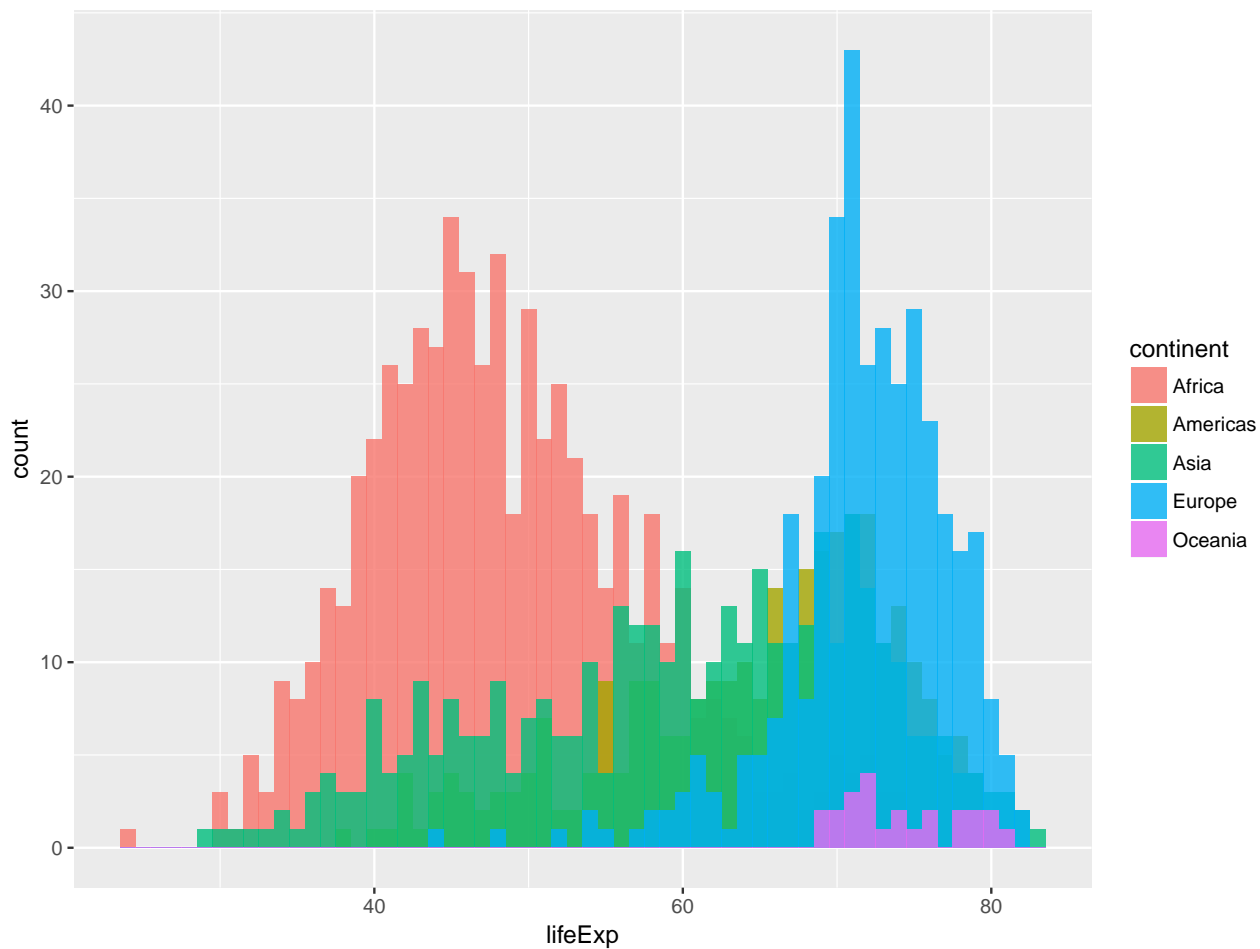
```
p8 + facet_wrap(~continent) + geom_boxplot(aes(group=year), alpha=0.1) +  
  labs(title = "Life expectancy per year")
```



DISTRIBUTION PLOTS

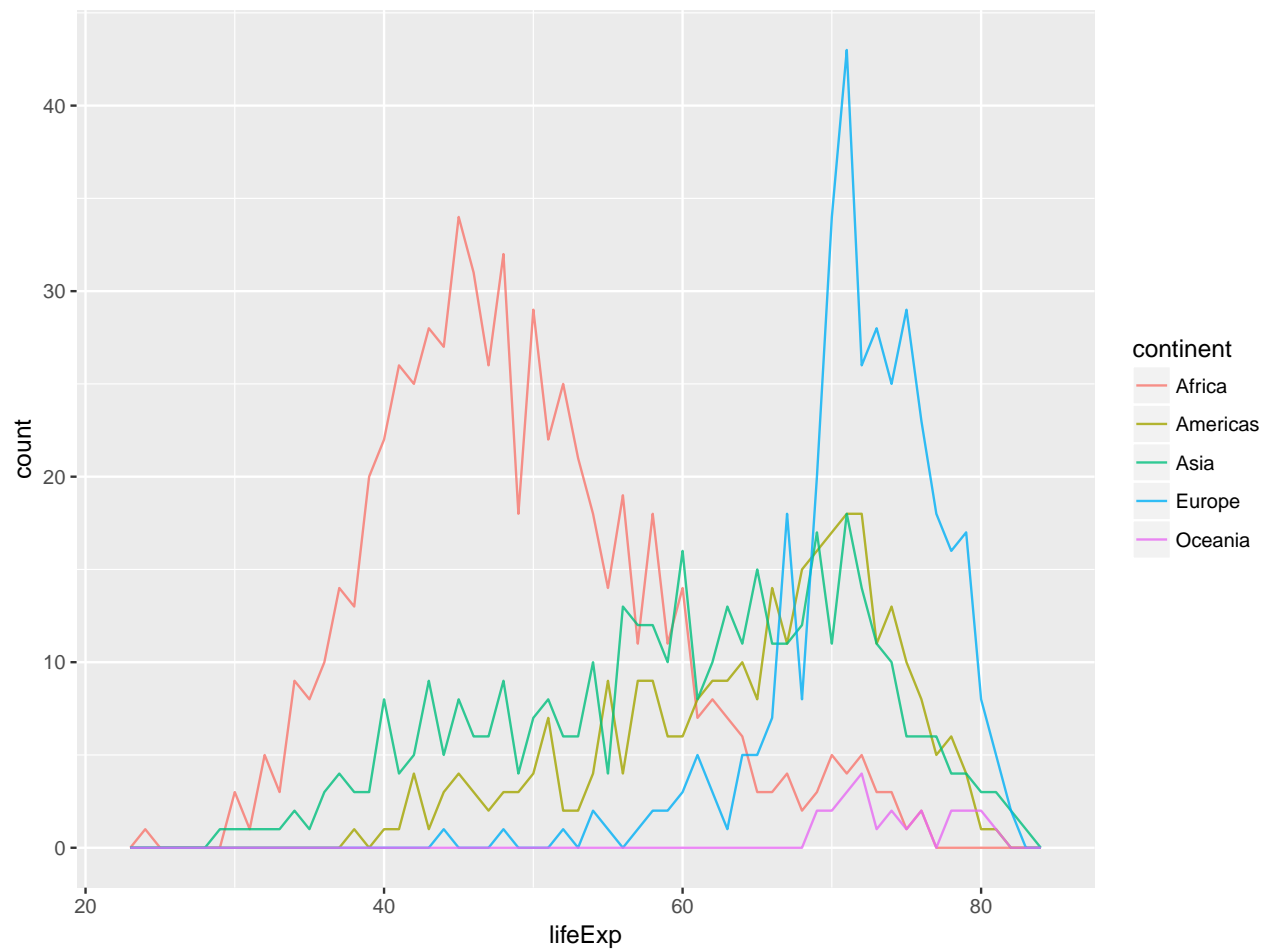
Let us explore a histogram to see the life expectancy measures distribution according to the continent.

```
p9 <- gapminder %>%
  ggplot()
p9 + geom_histogram(binwidth = 1, alpha = 0.8, position = "identity", aes(x=lifeExp, fill = continent))
```



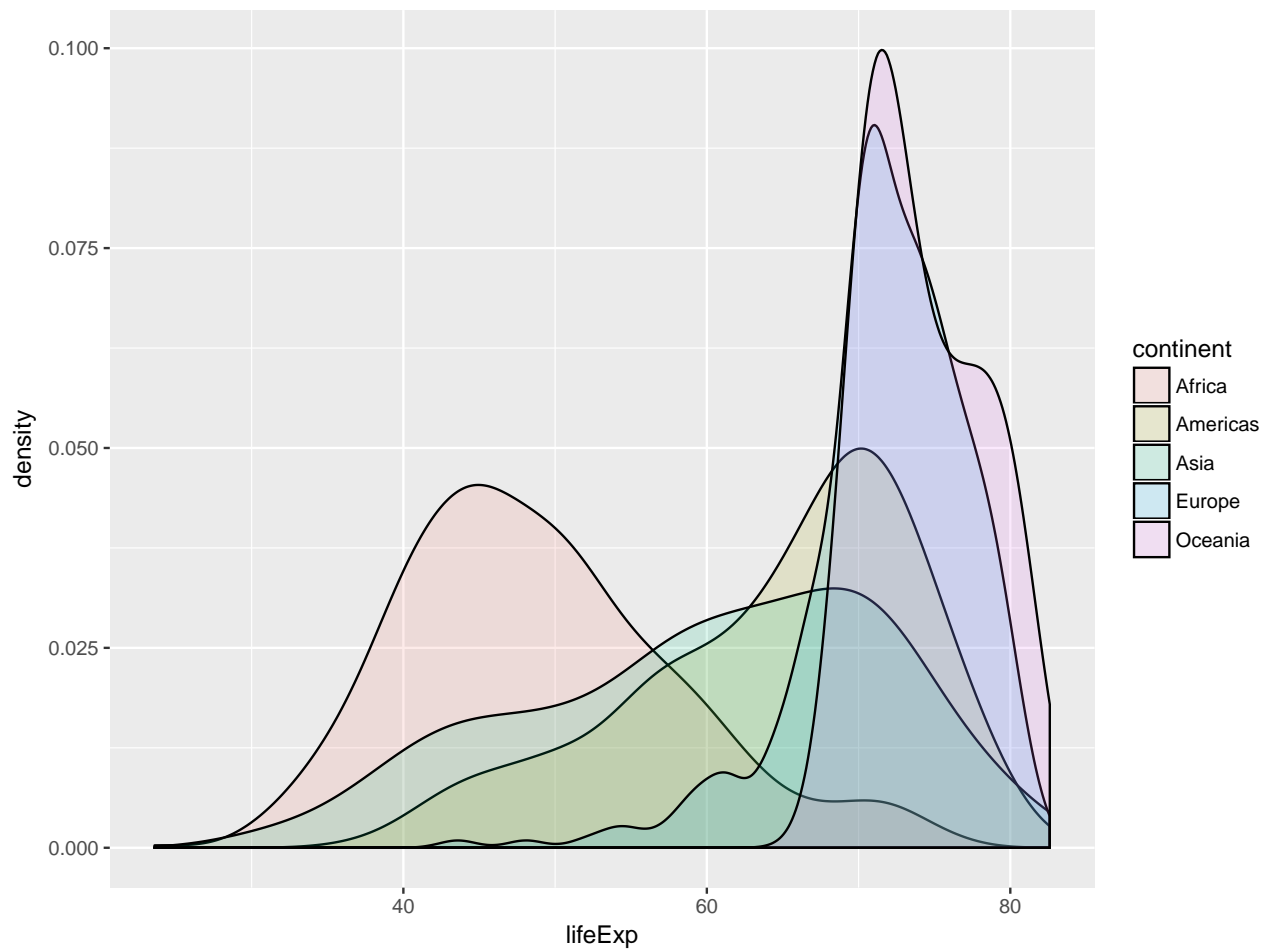
Now, let's use a frequency polygon to show the same information (but let's use a binwidth of 1 year).

```
p10 <- gapminder %>%
  ggplot()
p10 + geom_freqpoly(binwidth = 1, alpha = 0.8, position = "identity", aes(x=lifeExp, color = continent))
```

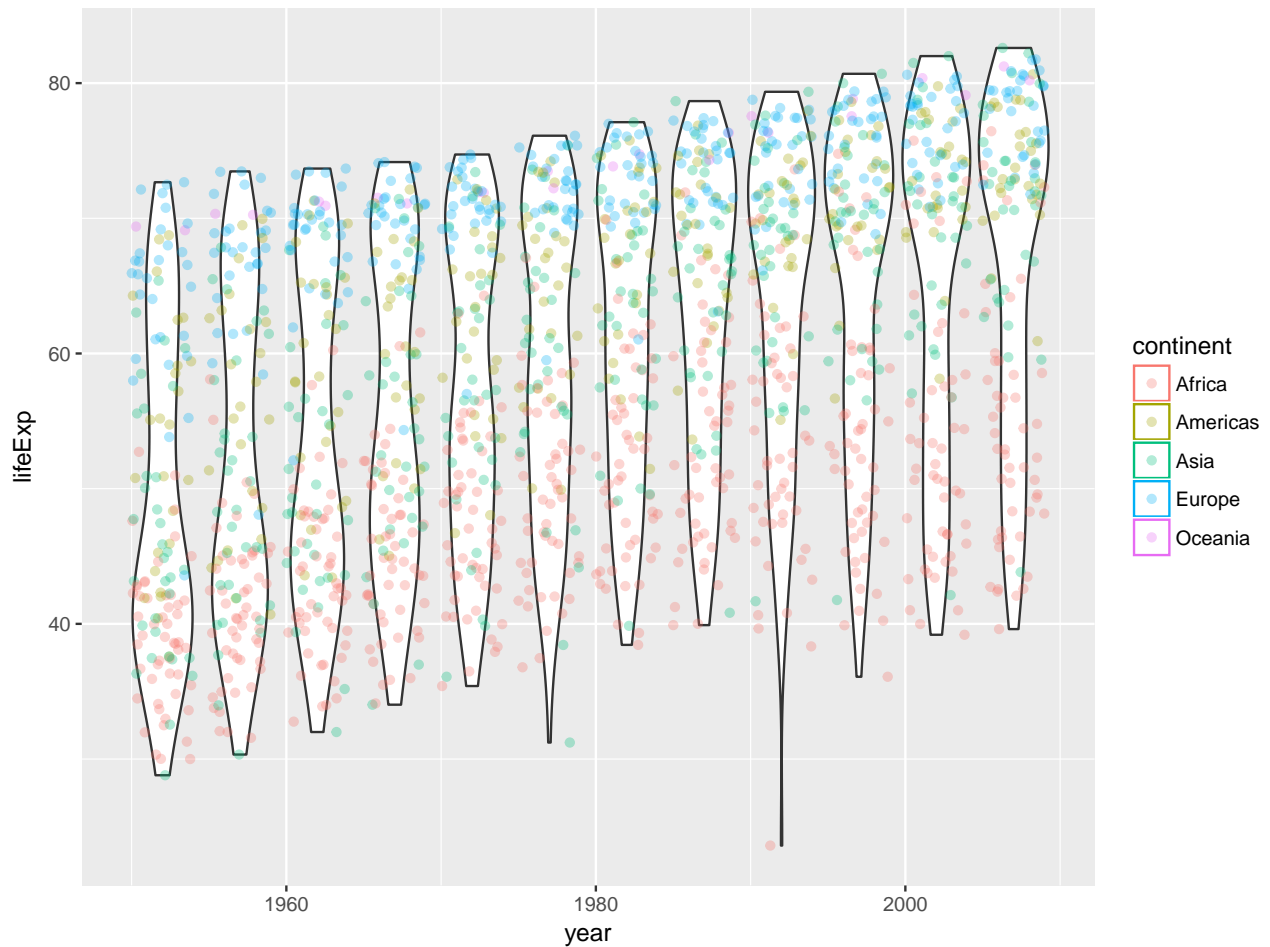
Now, let's look at a smoother plot, which `geom_density()` can help with.

```
p11 <- gapminder %>%
  ggplot(aes(x = lifeExp, fill = continent))
p11 + geom_density(alpha=0.15)
```



Another option is the `geom_violin` and `geom_violin` functions, to present our data in the most readable way.

```
p12 <- gapminder %>%  
  ggplot(aes(x=year, y =lifeExp, colour = continent))  
p12 + geom_violin(aes(group=year)) +  
  geom_jitter(alpha=0.3)
```



Let's explore `filter()`, `select()` and piping (`%>%`), and others

`filter()` exploration

Filter the data to only keep information related to Ireland and France before 2000.

```
my_gap <- gapminder
my_trip <- my_gap %>%
  filter(country %in% c("Ireland", "France"), year <= 2000)
View(my_trip)
```

`select()` and `arrange()` exploration

Then we can retrieve specific columns using `select`, for example, the country, year, `gdpPercap` and `lifeExp` variables. Then we can arrange the data according to the `gdpPercap` for example.

```
my_gap %>%
  filter(country %in% c("Ireland", "France"), year <= 2000) %>%
  select(country, year, gdpPercap, lifeExp) %>%
  arrange(gdpPercap)
```

```
## # A tibble: 20 x 4
```

```
##   country year gdpPercap lifeExp
##   <fctr> <int>      <dbl>   <dbl>
## 1 Ireland  1952   5210.280   66.910
## 2 Ireland  1957   5599.078   68.900
## 3 Ireland  1962   6631.597   70.290
## 4 France   1952   7029.809   67.410
## 5 Ireland  1967   7655.569   71.080
## 6 France   1957   8662.835   68.930
## 7 Ireland  1972   9530.773   71.280
## 8 France   1962  10560.486   70.510
## 9 Ireland  1977  11150.981   72.030
## 10 Ireland 1982  12618.321   73.100
## 11 France  1967  12999.918   71.550
## 12 Ireland 1987  13872.867   74.360
## 13 France  1972  16107.192   72.380
## 14 Ireland 1992  17558.816   75.467
## 15 France  1977  18292.635   73.830
## 16 France  1982  20293.897   74.890
## 17 France  1987  22066.442   76.340
## 18 Ireland 1997  24521.947   76.122
## 19 France  1992  24703.796   77.460
## 20 France  1997  25889.785   78.640
```

mutate() exploration

The cumulative gdp per country can be added to the my_gap dataset using the mutate function.

```
my_gap %>%
  mutate(gdp = pop*gdpPercap)
```

```
## # A tibble: 1,704 x 7
##   country continent year lifeExp      pop gdpPercap      gdp
##   <fctr>      <fctr> <int>   <dbl>   <int>   <dbl>   <dbl>
## 1 Afghanistan Asia   1952   28.801  8425333  779.4453 6567086330
## 2 Afghanistan Asia   1957   30.332  9240934  820.8530 7585448670
## 3 Afghanistan Asia   1962   31.997 10267083  853.1007 8758855797
## 4 Afghanistan Asia   1967   34.020 11537966  836.1971 9648014150
## 5 Afghanistan Asia   1972   36.088 13079460  739.9811 9678553274
## 6 Afghanistan Asia   1977   38.438 14880372  786.1134 11697659231
## 7 Afghanistan Asia   1982   39.854 12881816  978.0114 12598563401
## 8 Afghanistan Asia   1987   40.822 13867957  852.3959 11820990309
## 9 Afghanistan Asia   1992   41.674 16317921  649.3414 10595901589
## 10 Afghanistan Asia   1997   41.763 22227415  635.3414 14121995875
## # ... with 1,694 more rows
```

group_by() and summarize() exploration

Considering only year 2000 and before, if we group by continent, year and country, we can use the summarize_each() function to obtain the mean and median for the lifeExp and gdpPercap variables. Then, we can arrange according to the lifeExp_mean variable to see which continents, years and countries had the lowest lifeExp_mean values.

```
my_gap %>%
  filter(year <=2000) %>%
  group_by(continent, year, country) %>%
```

```
summarize_each(funs(mean, median), lifeExp, gdpPercap) %>%
  arrange(lifeExp_mean)
```

```
## `summarise_each()` is deprecated.
## Use `summarise_all()`, `summarise_at()` or `summarise_if()` instead.
## To map `funs` over a selection of variables, use `summarise_at()`

## # A tibble: 1,420 x 7
## # Groups:   continent, year [50]
##   continent year country lifeExp_mean gdpPercap_mean lifeExp_median
##   <fctr> <int>   <fctr>      <dbl>         <dbl>         <dbl>
## 1 Africa 1992    Rwanda      23.599         737.0686         23.599
## 2 Asia 1952   Afghanistan 28.801         779.4453         28.801
## 3 Africa 1952    Gambia      30.000         485.2307         30.000
## 4 Africa 1952    Angola      30.015        3520.6103         30.015
## 5 Africa 1952 Sierra Leone 30.331         879.7877         30.331
## 6 Asia 1957   Afghanistan 30.332         820.8530         30.332
## 7 Asia 1977    Cambodia    31.220         524.9722         31.220
## 8 Africa 1952   Mozambique  31.286         468.5260         31.286
## 9 Africa 1957 Sierra Leone 31.570        1004.4844         31.570
## 10 Africa 1952 Burkina Faso 31.975         543.2552         31.975
## # ... with 1,410 more rows, and 1 more variables: gdpPercap_median <dbl>
```

Let us explore the outlier values of life expectancy in Europe. I have used this [stat545 tutorial](#) to learn more about these functions.

```
my_gap %>%
  filter(continent == "Europe") %>%
  select(year, country, lifeExp) %>%
  group_by(year) %>%
  filter(min_rank(desc(lifeExp)) <2 | min_rank(lifeExp) <2) %>%
  arrange(year, lifeExp) %>%
  print(n = Inf)
```

```
## # A tibble: 24 x 3
## # Groups:   year [12]
##   year country lifeExp
##   <int>   <fctr>   <dbl>
## 1 1952    Turkey  43.585
## 2 1952    Norway  72.670
## 3 1957    Turkey  48.079
## 4 1957    Iceland 73.470
## 5 1962    Turkey  52.098
## 6 1962    Iceland 73.680
## 7 1967    Turkey  54.336
## 8 1967    Sweden  74.160
## 9 1972    Turkey  57.005
## 10 1972    Sweden  74.720
## 11 1977    Turkey  59.507
## 12 1977    Iceland 76.110
## 13 1982    Turkey  61.036
## 14 1982    Iceland 76.990
## 15 1987    Turkey  63.108
## 16 1987 Switzerland 77.410
## 17 1992    Turkey  66.146
```

```
## 18 1992      Iceland 78.770
## 19 1997      Turkey 68.835
## 20 1997      Sweden 79.390
## 21 2002      Turkey 70.845
## 22 2002 Switzerland 80.620
## 23 2007      Turkey 71.777
## 24 2007      Iceland 81.757
```

Let's re-initialize my European dataset and then perform some further analyses and may be even plotting. Let's start with looking at the worst European life expectancy every year.

```
rm(my_europe)
```

```
## Warning in rm(my_europe): object 'my_europe' not found
```

```
my_europe <- my_gap %>%
  filter(continent == "Europe") %>%
  select(year, country, lifeExp) %>%
  group_by(year) %>%
  top_n(1, wt=desc(lifeExp))
my_europe
```

```
## # A tibble: 12 x 3
## # Groups:   year [12]
##   year country lifeExp
##   <int> <fctr>   <dbl>
## 1 1952 Turkey 43.585
## 2 1957 Turkey 48.079
## 3 1962 Turkey 52.098
## 4 1967 Turkey 54.336
## 5 1972 Turkey 57.005
## 6 1977 Turkey 59.507
## 7 1982 Turkey 61.036
## 8 1987 Turkey 63.108
## 9 1992 Turkey 66.146
## 10 1997 Turkey 68.835
## 11 2002 Turkey 70.845
## 12 2007 Turkey 71.777
```

Wow, Turkey has not been doing well for a long time ...

Using the knitr library to make prettier tables

```
knitr::kable(head(gapminder),
  format = "pandoc",
  digits = 2,
  caption = "Gapminder (head)",
  align = c("l"),
  longtable = TRUE,
  padding = 3)
```

Table 2: Gapminder (head)

country	continent	year	lifeExp	pop	gdpPercap
Afghanistan	Asia	1952	28.80	8425333	779.45

country	continent	year	lifeExp	pop	gdpPercap
Afghanistan	Asia	1957	30.33	9240934	820.85
Afghanistan	Asia	1962	32.00	10267083	853.10
Afghanistan	Asia	1967	34.02	11537966	836.20
Afghanistan	Asia	1972	36.09	13079460	739.98
Afghanistan	Asia	1977	38.44	14880372	786.11

Oh, I just understood that you have to “knit” this document in order to see the nice table made with knitr::kable !

I am planning to use knitr::kable function more in my future homework.

I have used this pandoc website as well to install some complementary packages.

I am looking forward to apply this newly gained knowledge to genomic data analyses such as gene expression, copy number, mutational signatures and pathway analysis.

Now I feel like I have a better grasp on how to use these new tools.