

# Biostats 597 HW1

*Bing Miu*

A look at the data:

```
## # A tibble: 1,704 x 6
##   country    continent year lifeExp      pop gdpPercap
##   <fct>      <fct>    <int>  <dbl>    <int>    <dbl>
## 1 Afghanistan Asia      1952   28.8  8425333    779.
## 2 Afghanistan Asia      1957   30.3  9240934    821.
## 3 Afghanistan Asia      1962   32.0 10267083    853.
## 4 Afghanistan Asia      1967   34.0 11537966    836.
## 5 Afghanistan Asia      1972   36.1 13079460    740.
## 6 Afghanistan Asia      1977   38.4 14880372    786.
## 7 Afghanistan Asia      1982   39.9 12881816    978.
## 8 Afghanistan Asia      1987   40.8 13867957    852.
## 9 Afghanistan Asia      1992   41.7 16317921    649.
## 10 Afghanistan Asia      1997   41.8 22227415    635.
## # ... with 1,694 more rows
```

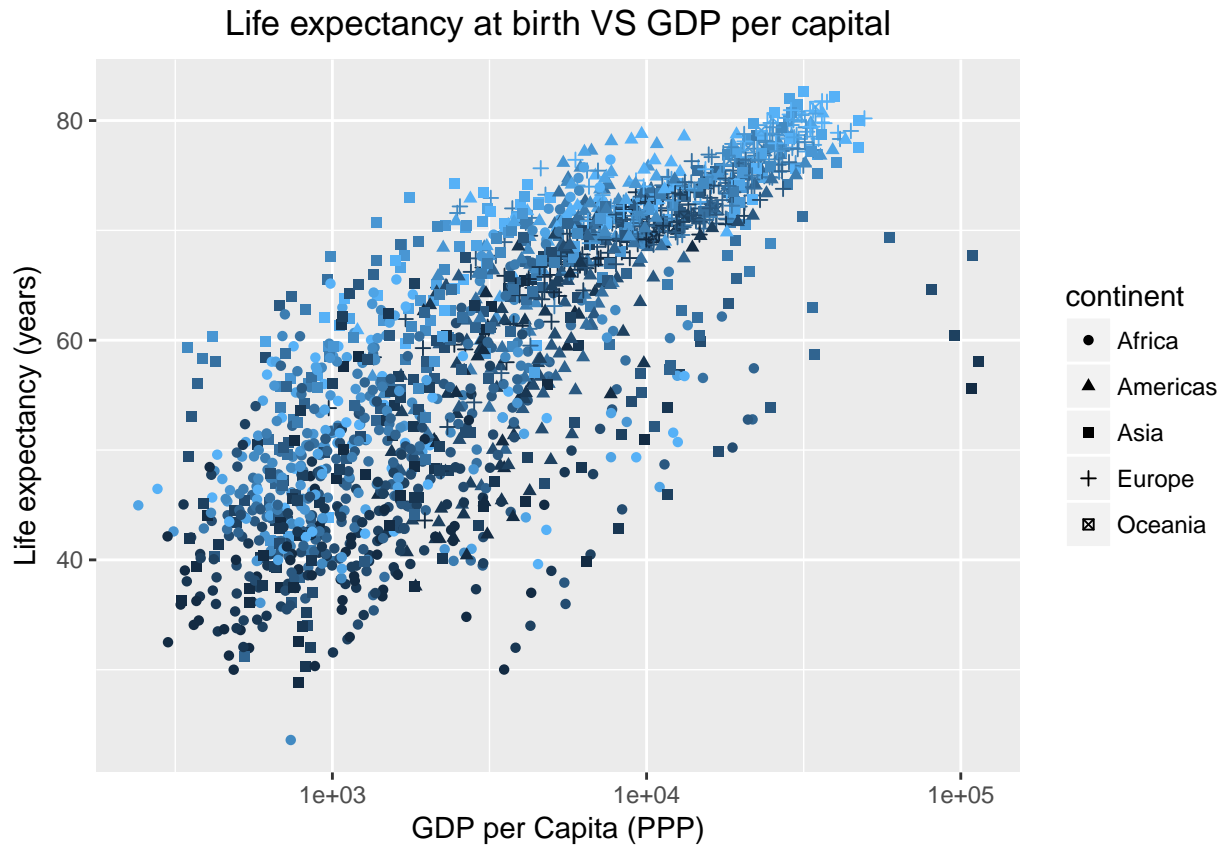
## HW exercises:

use ggplot to make the graphs requested in (1) and (2). Throughout, avoid repetition of code. Goal: gain experience with ggplot and reading help files/using google searches to get help on (arguments of) R functions.

1. make one scatter plot using ggplot with

- Life expectancy on the y-axis, label axis as “Life expectancy”
- GDP per capita on the x-axis, use a log-transform of the x-axis while still displaying the (unlogged) GDP values
- shapes to indicate the continent, add a legend
- color to indicate the year, do NOT add a legend for that

```
ggplot(data = gapminder, mapping = aes(x = gdpPercap, y = lifeExp)) +
  labs(title = "Life expectancy at birth VS GDP per capital",
       x = "GDP per Capita (PPP)", y = "Life expectancy (years)") +
  scale_x_continuous(trans = "log10") + theme(plot.title = element_text(hjust = 0.5)) +
  geom_point(mapping = aes(color = year, shape = continent)) +
  scale_color_continuous(guide = "none")
```

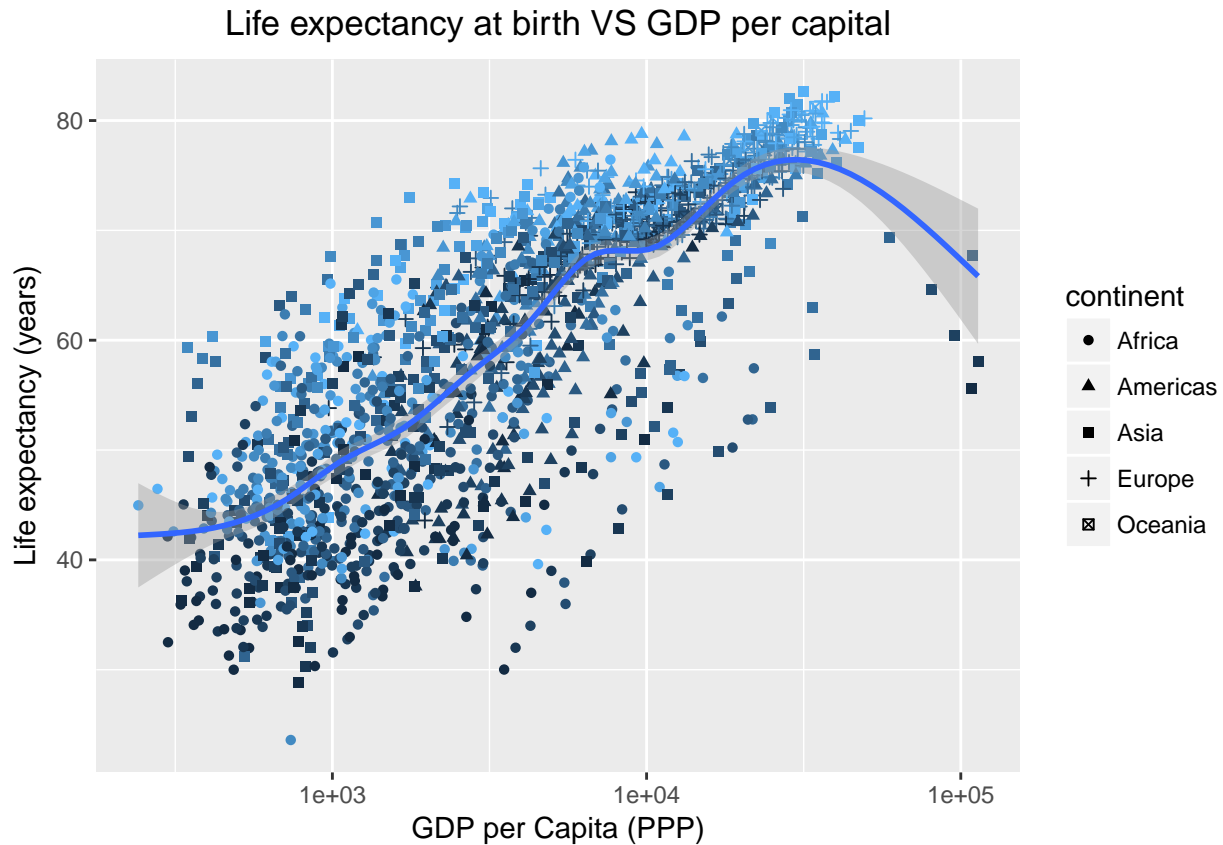


2. add two `geom_smooth` layers (a) and (b) to the plot you made in (1) where

(a) is fitted to the entire dataset and uses default settings for the smoother used and plotting settings

```
ggplot(data = gapminder, mapping = aes(x = gdpPercap, y = lifeExp)) +
  labs(title = "Life expectancy at birth VS GDP per capital",
       x = "GDP per Capita (PPP)", y = "Life expectancy (years)") +
  scale_x_continuous(trans = "log10") + theme(plot.title = element_text(hjust = 0.5)) +
  geom_point(mapping = aes(color = year, shape = continent)) +
  scale_color_continuous(guide = "none") + geom_smooth()
```

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



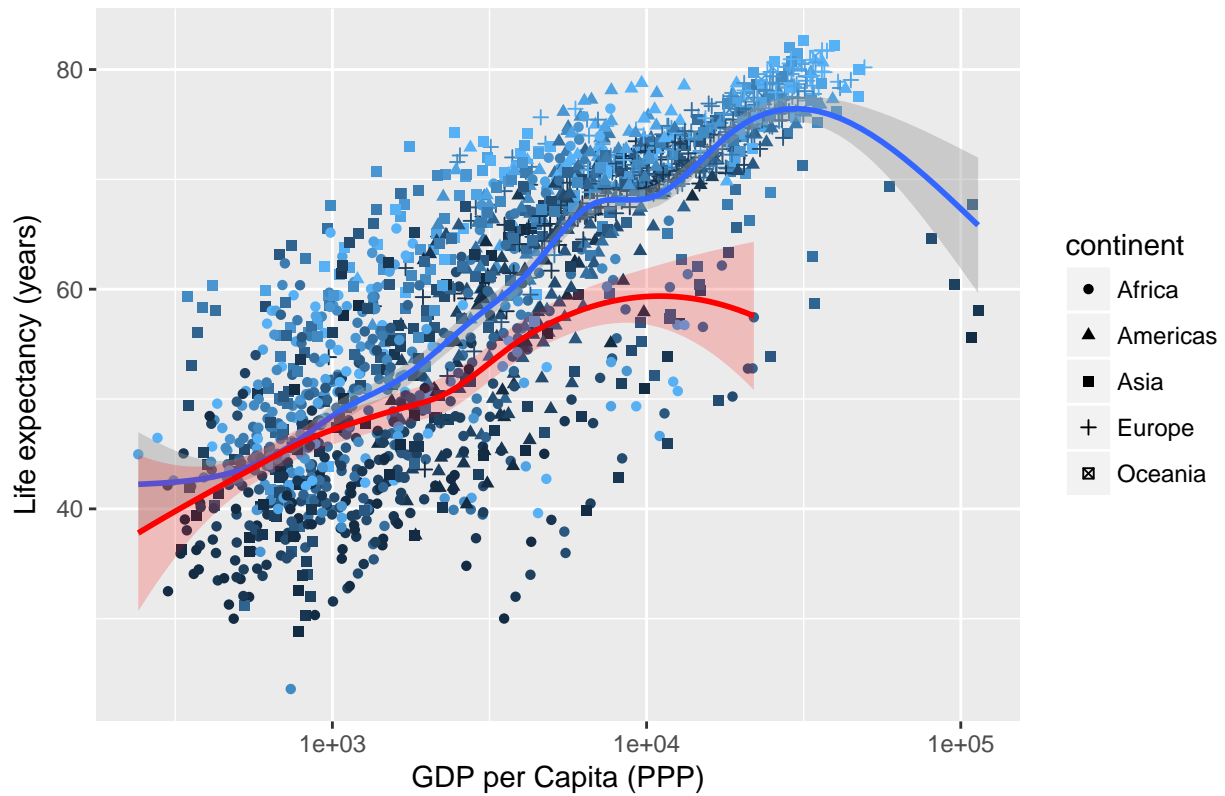
(b) is fitted to data in Africa only, provides a 99% confidence interval, is plotted in red (both point estimates as well as confidence interval), and is transparent (so if it overlaps with (a), (a) is still visible). Hint: to fit to data in Africa only, consider using “data = filter(gapminder, continent == "Africa")”

```
ggplot(data = gapminder, mapping = aes(x = gdpPercap, y = lifeExp)) +
  labs(title = "Life expectancy at birth VS GDP per capital",
        x = "GDP per Capita (PPP)", y = "Life expectancy (years)") +
  scale_x_continuous(trans = "log10") + theme(plot.title = element_text(hjust = 0.5)) +
  scale_color_continuous(guide = "none") + geom_point(mapping = aes(color = year,
    shape = continent)) + geom_smooth() + geom_smooth(data = filter(gapminder,
    continent == "Africa"), color = "red", fill = "red", level = 0.99,
    alpha = 0.2)
```

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

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

Life expectancy at birth VS GDP per capital



Red = fitted to data in Africa only

Blue = fitted to data in all continents