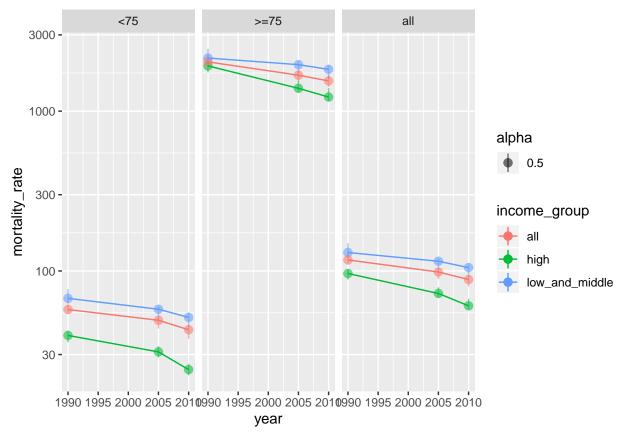
BMI 881 Homework 1

Li Ge 2019-10-03

This is the link to the homework.

Design Choices

- Apparently, the data can be divided into two parts. One part contains the <code>year</code>, <code>age_group</code>, <code>income_group</code>, the other part contains the <code>mortality_rate</code>, <code>interval_low</code>, <code>interval_high</code>. I should find a way to arrange these different groups. And I should always present the mortality rate together with its interval. <code>ggplot2</code>'s facet_grip and <code>geom_pointrange</code> seem to be the perfect fit for it.
- I was originally planning to design a facet_grid with <code>income_group v. age_group</code>, each subplot is the <code>mortality v. year</code>. But then I realized it would be difficult to compare the mortality rate between different income groups. So I decided to use color to indicate different income groups. And it was so much better.
- Another thing worth to mention is that I log-scaled the y-axis. Because the mortality rates are drastically different between >=75 v. rest.



Conclusions

- >=75 age group has the highest mortality rate.
- The higher the income, the lower the mortality rate.
- The mortality rate decrease over the years.

Code

```
knitr::opts_chunk$set(
   echo = FALSE,
   fig.align = "center"
)

library(ggplot2)
# read the data
stroke <- read.csv("https://kbroman.org/BMI881/assets/feigin2014_table1_mortality.csv")

# plot
ggplot(data = stroke, mapping = aes(x = year, y = mortality_rate, color = income_group)) +
   facet_grid(. ~ age_group) +
   scale_y_log10() +
   geom_pointrange(aes(ymin = interval_low, ymax = interval_high, alpha = 0.5)) +
   geom_line()

# this R markdown chunk generates a code appendix</pre>
```