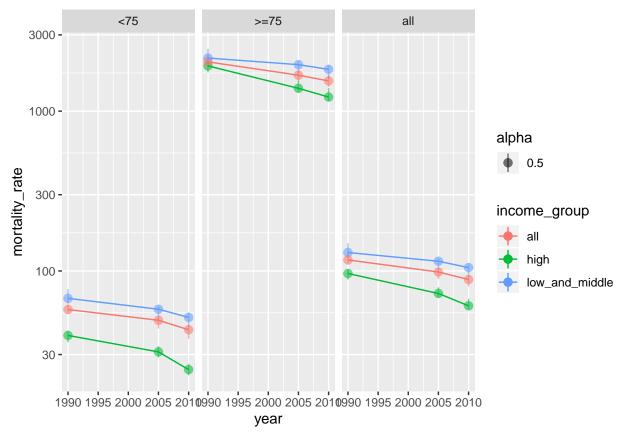
## BMI 881 Homework 1

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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.

## 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>
```