

Analysis on Factors affecting Life Expectancy

```
library(readr)
expect<-read_csv("D:/MITA 2019/Semester 2/Multivariate Analysis/lfe.csv")

## Parsed with column specification:
## cols(
##   .default = col_double(),
##   Country = col_character(),
##   Status = col_character()
## )

## See spec(...) for full column specifications.

head(expect)

## # A tibble: 6 x 22
##   Country Year Status `Life expectanc~` `Adult Mortalit~` `infant deaths`
##   <chr>   <dbl> <chr>         <dbl>         <dbl>         <dbl>
## 1 Afghan~ 2015 Devel~         65           263           62
## 2 Afghan~ 2014 Devel~         59.9         271           64
## 3 Afghan~ 2013 Devel~         59.9         268           66
## 4 Afghan~ 2012 Devel~         59.5         272           69
## 5 Afghan~ 2011 Devel~         59.2         275           71
## 6 Afghan~ 2010 Devel~         58.8         279           74
## # ... with 16 more variables: Alcohol <dbl>, `percentage
## #   expenditure` <dbl>, `Hepatitis B` <dbl>, Measles <dbl>, BMI <dbl>,
## #   `under-five deaths` <dbl>, Polio <dbl>, `Total expenditure` <dbl>,
## #   Diphtheria <dbl>, `HIV/AIDS` <dbl>, GDP <dbl>, Population <dbl>,
## #   `thinness 1-19 years` <dbl>, `thinness 5-9 years` <dbl>, `Income
## #   composition of resources` <dbl>, Schooling <dbl>

library(tidyverse) # metapackage with lots of helpful functions

## -- Attaching packages -----
## ----- tidyverse 1.2.1 -----

## v ggplot2 3.2.1      v purrr 0.3.2
## v tibble 2.1.3       v dplyr 0.8.3
## v tidyr 1.0.0        v stringr 1.4.0
## v ggplot2 3.2.1      v forcats 0.4.0

## -- Conflicts -----
## ---- tidyverse_conflicts() ----
## x dplyr::filter() masks stats::filter()
## x dplyr::lag() masks stats::lag()

library(gridExtra) # grid.arrange to make quick subplots
```

```
##
## Attaching package: 'gridExtra'

## The following object is masked from 'package:dplyr':
##
##      combine

library(reshape2)

##
## Attaching package: 'reshape2'

## The following object is masked from 'package:tidyr':
##
##      smiths

expect <- expect %>%
  drop_na() %>%
  rename('lifexp'='Life expectancy',
        'percexp'='percentage expenditure',
        'totexp'='Total expenditure',
        'admort'='Adult Mortality',
        'infmort'='infant deaths',
        'u5deaths'='under-five deaths',
        'hepb'='Hepatitis B',
        'HIV'='HIV/AIDS') %>%
  filter(!is.na(lifexp), !is.na(admort), !is.na(infmort))
head(expect)

## # A tibble: 6 x 22
##   Country Year Status lifexp admort infmort Alcohol percexp hepb Measles
##   <chr>   <dbl> <chr>   <dbl>  <dbl>  <dbl>   <dbl>  <dbl>  <dbl>
## 1 Afghan~ 2015 Devel~    65    263    62    0.01   71.3    65   1154
## 2 Afghan~ 2014 Devel~   59.9   271    64    0.01   73.5    62    492
## 3 Afghan~ 2013 Devel~   59.9   268    66    0.01   73.2    64    430
## 4 Afghan~ 2012 Devel~   59.5   272    69    0.01   78.2    67   2787
## 5 Afghan~ 2011 Devel~   59.2   275    71    0.01    7.10    68   3013
## 6 Afghan~ 2010 Devel~   58.8   279    74    0.01   79.7    66   1989
## # ... with 12 more variables: BMI <dbl>, u5deaths <dbl>, Polio <dbl>,
## #   totexp <dbl>, Diphtheria <dbl>, HIV <dbl>, GDP <dbl>,
## #   Population <dbl>, `thinness 1-19 years` <dbl>, `thinness 5-9
## #   years` <dbl>, `Income composition of resources` <dbl>, Schooling <dbl>

#Descriptive statistics grouped by status of country
expect %>%
  group_by(Status) %>%
  summarize(count = n(),
            avg_lifexp = mean(lifexp, na.rm=TRUE),
            #avg_infmort = mean(infmort, na.rm=TRUE),
            #avg_admort = mean(admort, na.rm=TRUE),
            avg_GDP = mean(GDP, na.rm=TRUE),
```

```

    avg_Population = mean(Population, na.rm=TRUE),
    avg_totexp = mean(totexp, na.rm=TRUE))

## # A tibble: 2 x 6
##   Status      count avg_lifexp avg_GDP avg_Population avg_totexp
##   <chr>      <int>    <dbl>  <dbl>      <dbl>      <dbl>
## 1 Developed    242      78.7  18977.    8744688.    7.02
## 2 Developing  1407      67.7   3259.   15669946.    5.77

devd <- expect %>%
  filter(Country %in% c('Australia', 'Germany', 'Israel'))
deving <- expect %>%
  filter(Country %in% c('Afghanistan', 'Nigeria', 'Indonesia'))

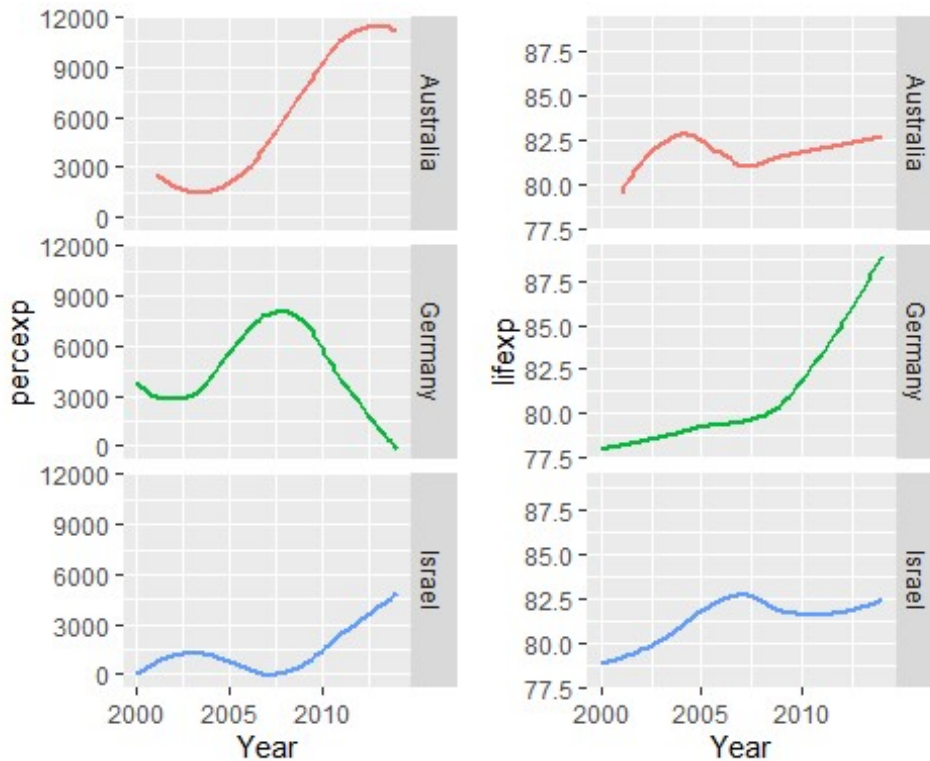
# removing 2015 data, has some NA values
devd <- devd %>%
  filter(Year != 2015)
deving <- deving %>%
  filter(Year != 2015)

#EDA for Developed countries against percentage expenditure and Life
expectancy
devdp1 <- ggplot(devd, aes(Year, percexp))+
  geom_smooth(aes(color=Country), se=FALSE, show.legend=FALSE)+
  facet_grid(Country~.)

devdp2 <- ggplot(devd, aes(Year, lifexp))+
  geom_smooth(aes(color=Country), se=FALSE, show.legend=FALSE)+
  facet_grid(Country~.)
grid.arrange(devdp1, devdp2, nrow=1)

## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'

```

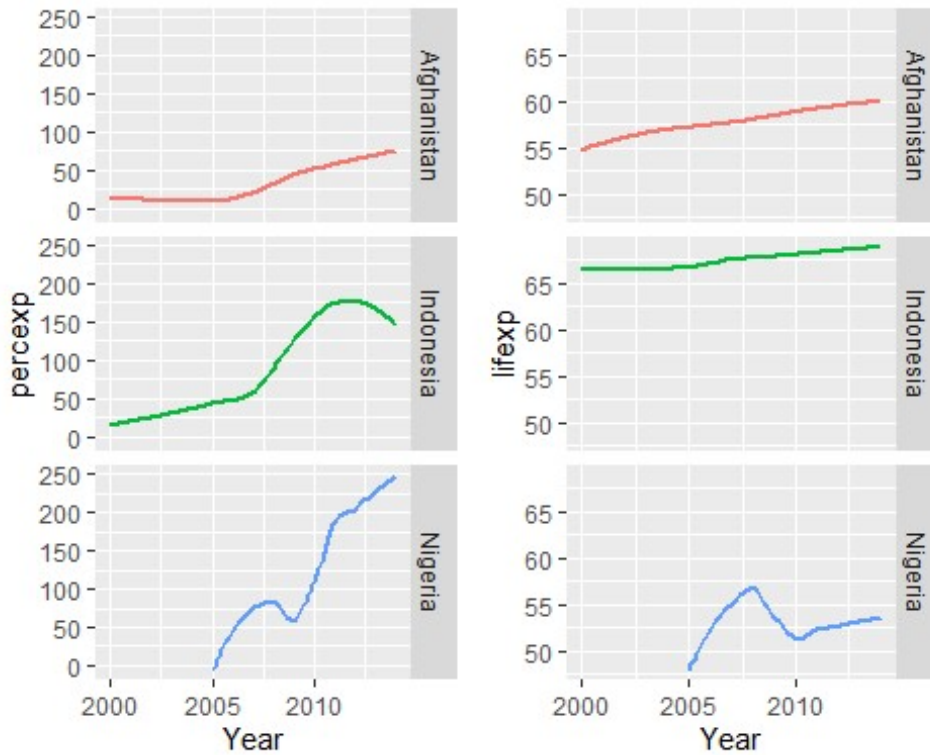


#EDA for developing countries against percentage expenditure and life expectancy

```
devingp1 <- ggplot(deving, aes(Year, percexp))+
  geom_smooth(aes(color=Country), se=FALSE, show.legend=FALSE)+
  facet_grid(Country~.)

devingp2 <- ggplot(deving, aes(Year, lifexp))+
  geom_smooth(aes(color=Country), se=FALSE, show.legend=FALSE)+
  facet_grid(Country~.)
grid.arrange(devingp1, devingp2, nrow=1)

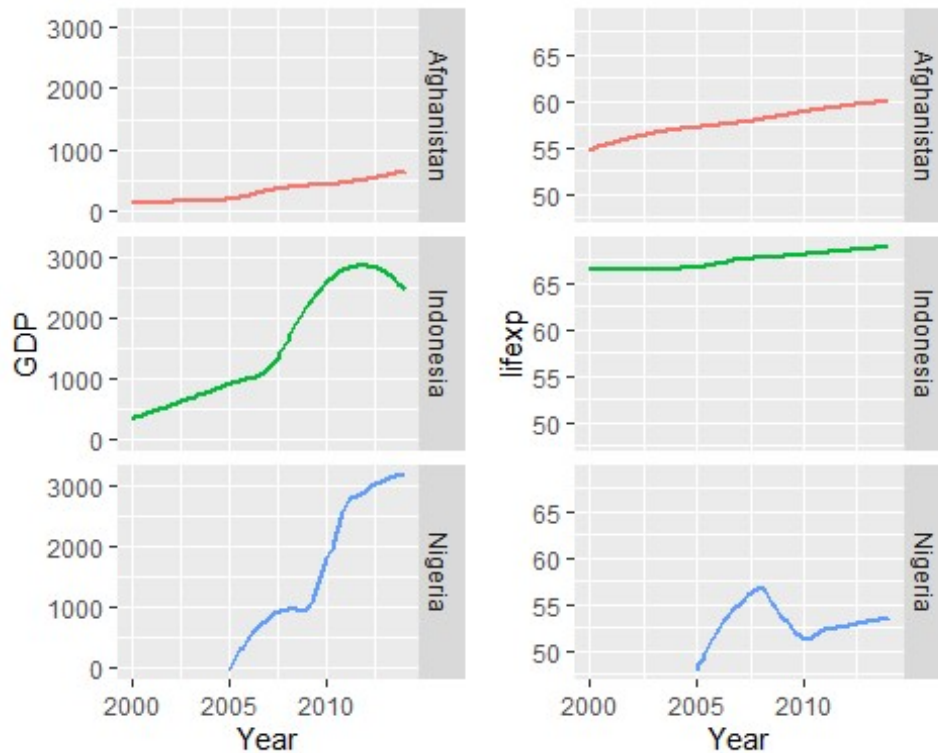
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
```



```
#EDA for developing countries against GDP and Life expectancy
devingp1 <- ggplot(deving, aes(Year, GDP))+
  geom_smooth(aes(color=Country), se=FALSE, show.legend=FALSE)+
  facet_grid(Country~.)

devingp2 <- ggplot(deving, aes(Year, lifexp))+
  geom_smooth(aes(color=Country), se=FALSE, show.legend=FALSE)+
  facet_grid(Country~.)
grid.arrange(devingp1, devingp2, nrow=1)

## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
```



#correlation between life expectancy and percentage expenditure, use cor() to check

```
cor(expect$lifexp, expect$percexp)
```

```
## [1] 0.4096308
```

#correlation between life expectancy and GDP

```
cor(expect$lifexp, expect$GDP)
```

```
## [1] 0.4413218
```

#correlation between life expectancy and Alcohol

```
cor(expect$lifexp, expect$Alcohol)
```

```
## [1] 0.4027183
```

Looking at infant deaths and under-five deaths for developed countries

```
ggplot(devd, show.legend=FALSE)+
```

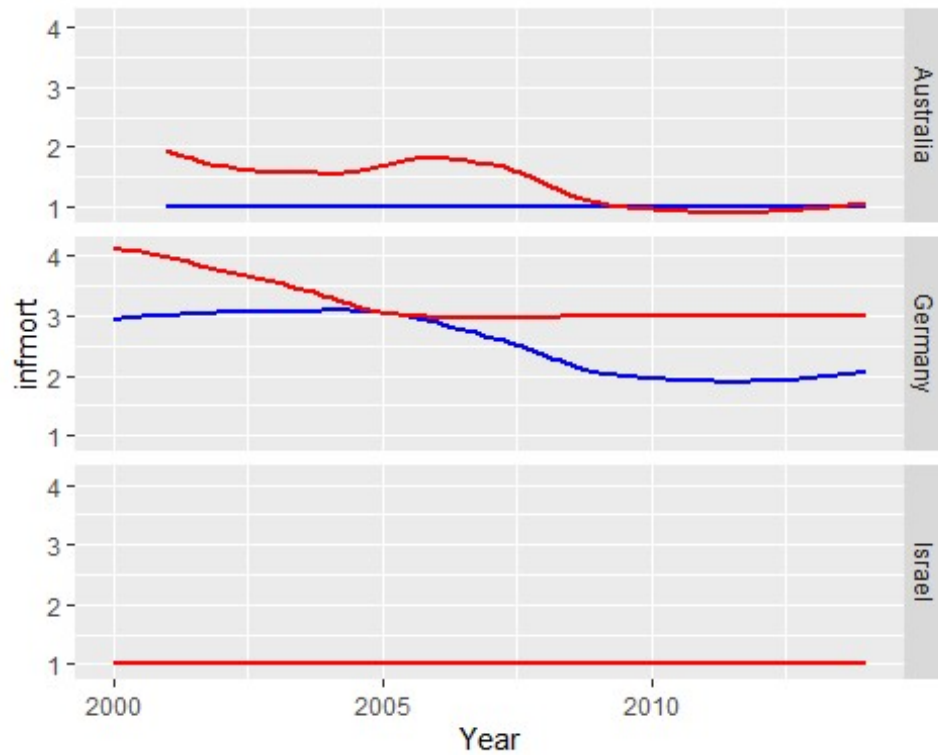
```
  geom_smooth(aes(Year, infmort), color='blue', se=FALSE)+
```

```
  geom_smooth(aes(Year, u5deaths), color='red', se=FALSE)+
```

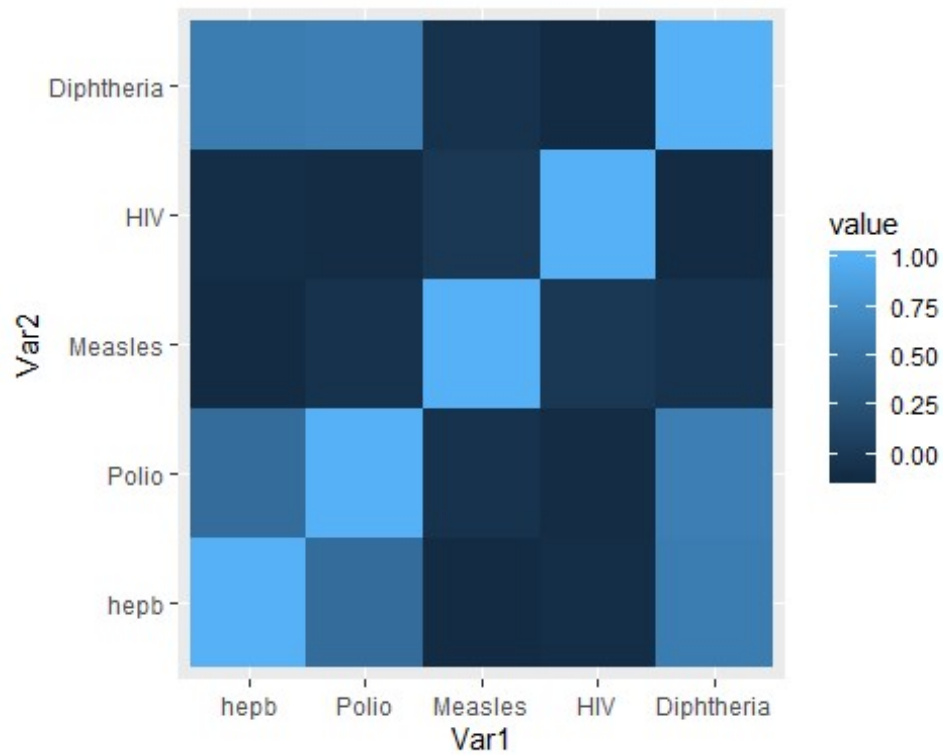
```
  facet_grid(Country~.)
```

```
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
```

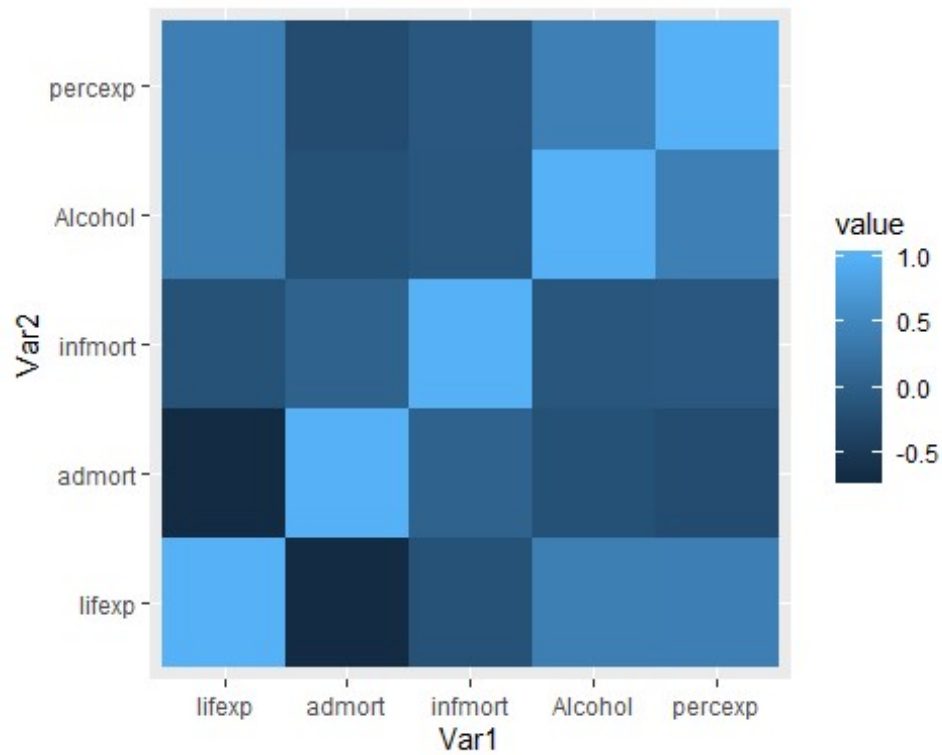
```
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
```



```
# any correlation between vaccinations?
cordf <- expect %>%
  drop_na() %>%
  select(hepb, Polio, Measles, HIV, Diphtheria)
cormat <- cor(cordf)
melted <- melt(cormat)
ggplot(melted)+
  geom_tile(aes(Var1, Var2, fill=value))
```



```
cordf <- expect %>%
  drop_na() %>%
  select(lifexp, admort, infmort, Alcohol, percexp)
cormat <- cor(cordf)
melted <- melt(cormat)
ggplot(melted) +
  geom_tile(aes(Var1, Var2, fill=value))
```

how about adult mortality and infant mortality?

expect %>%

group_by(Year) %>%

summarize(Polio=mean(Polio),
Measles=mean(Measles)) %>%

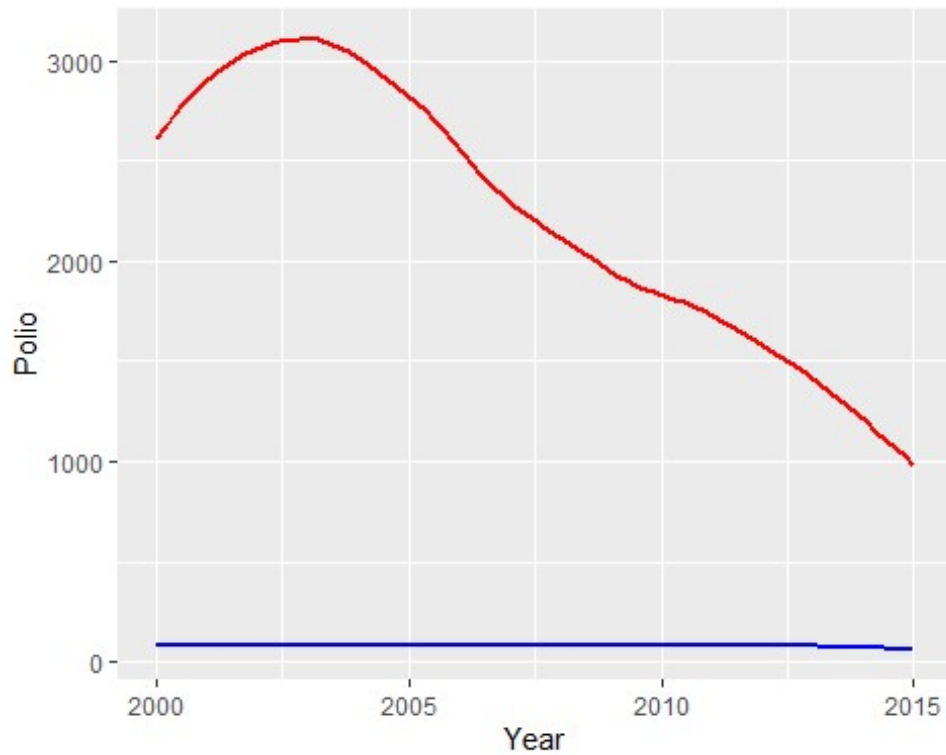
ggplot()+

geom_smooth(aes(Year, Polio), color='blue', se=FALSE)+

geom_smooth(aes(Year, Measles), color='red', se=FALSE)

`geom_smooth()` using method = 'loess' and formula 'y ~ x'

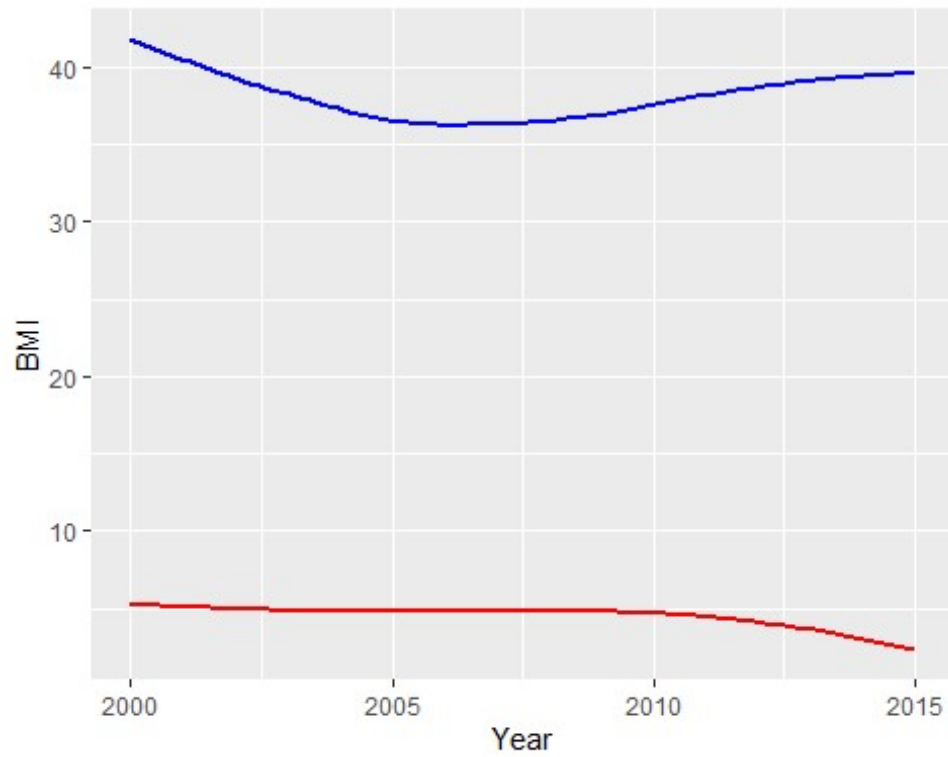
`geom_smooth()` using method = 'loess' and formula 'y ~ x'



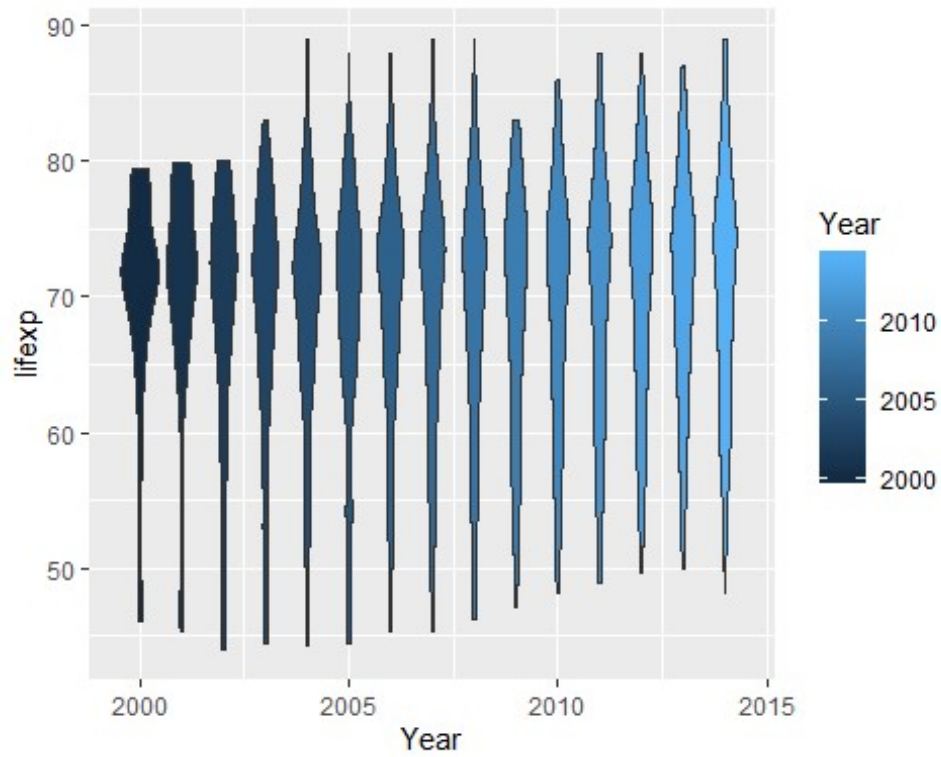
```
#geom_smooth(aes(Year, HIV), color='yellow', se=FALSE)+
#geom_smooth(aes(Year, Diphtheria), color='', se=FALSE)+

expect %>%
  group_by(Year) %>%
  summarize(BMI=mean(BMI),
            Alcohol=mean(Alcohol)) %>%
  ggplot()+
  geom_smooth(aes(Year, BMI), color='blue', se=FALSE)+
  geom_smooth(aes(Year, Alcohol), color='red', se=FALSE)

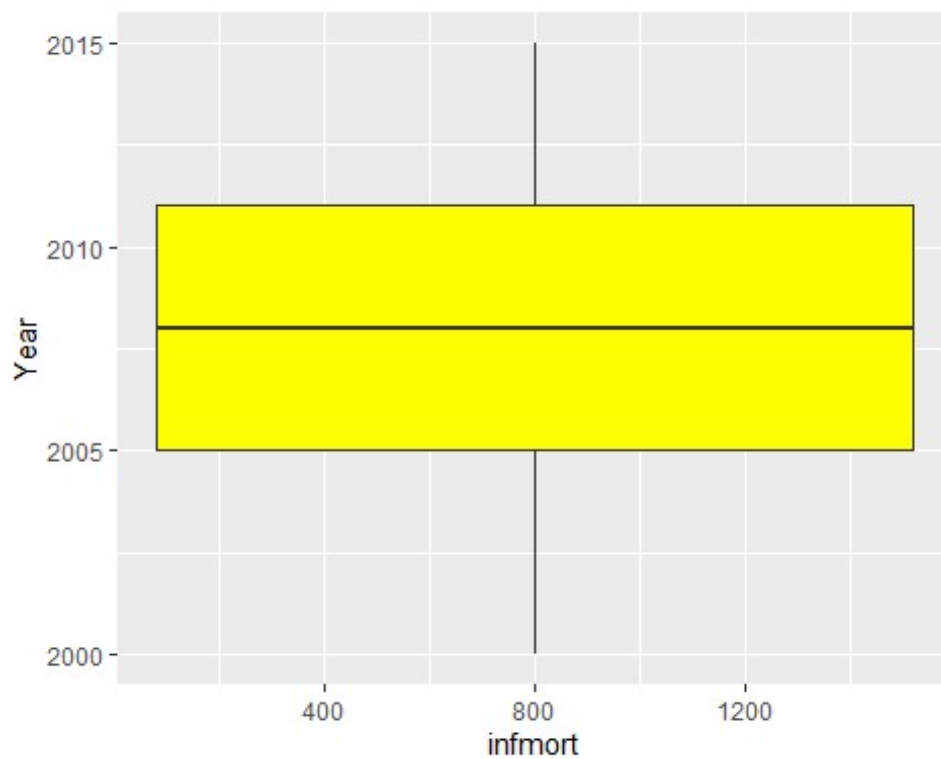
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
```



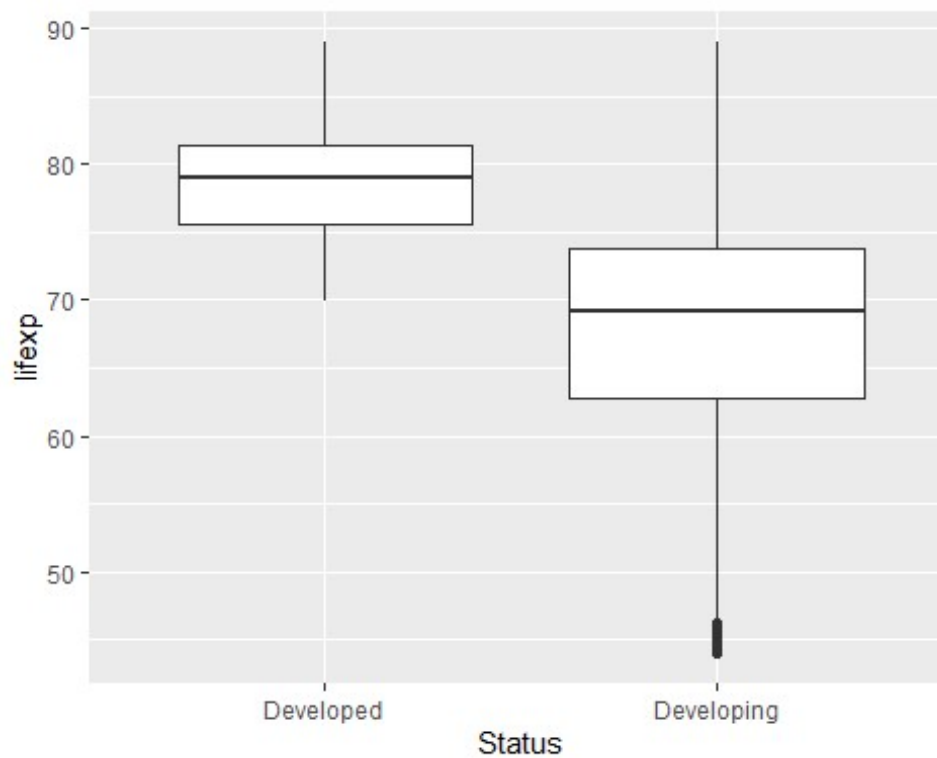
```
#distribution of life expectancy by year  
expect %>%  
  ggplot()+  
  geom_violin(aes(x=Year, y=lifexp, group=Year, fill=Year))
```



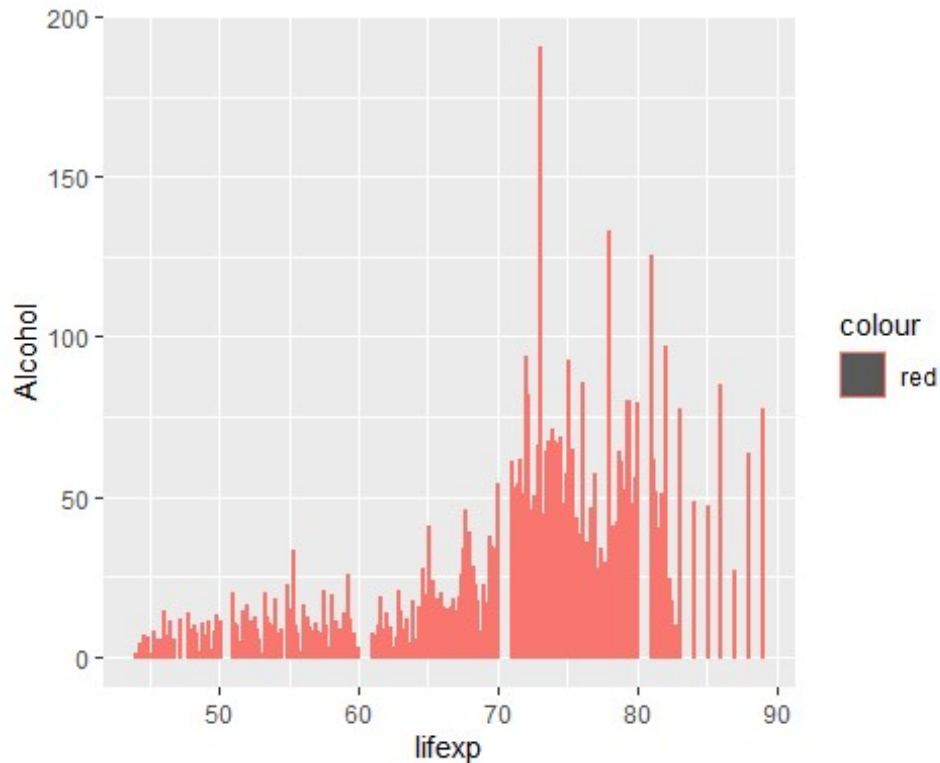
```
ggplot(expect, aes(y=Year, x=infmort)) + geom_boxplot(fill='yellow')
## Warning: Continuous x aesthetic -- did you forget aes(group=...)?
```



```
#Converting status into categorical and plotting boxplot  
expect$factor = as.factor(expect$Status)  
ggplot(expect,aes(y=lifexp,x=Status))+geom_boxplot()
```



```
#Plotting bar graph for Life expectancy and Alcohol  
x<-ggplot(data=expect, mapping=aes(x=lifexp,y=Alcohol))  
x+geom_col(aes(color="red"))
```



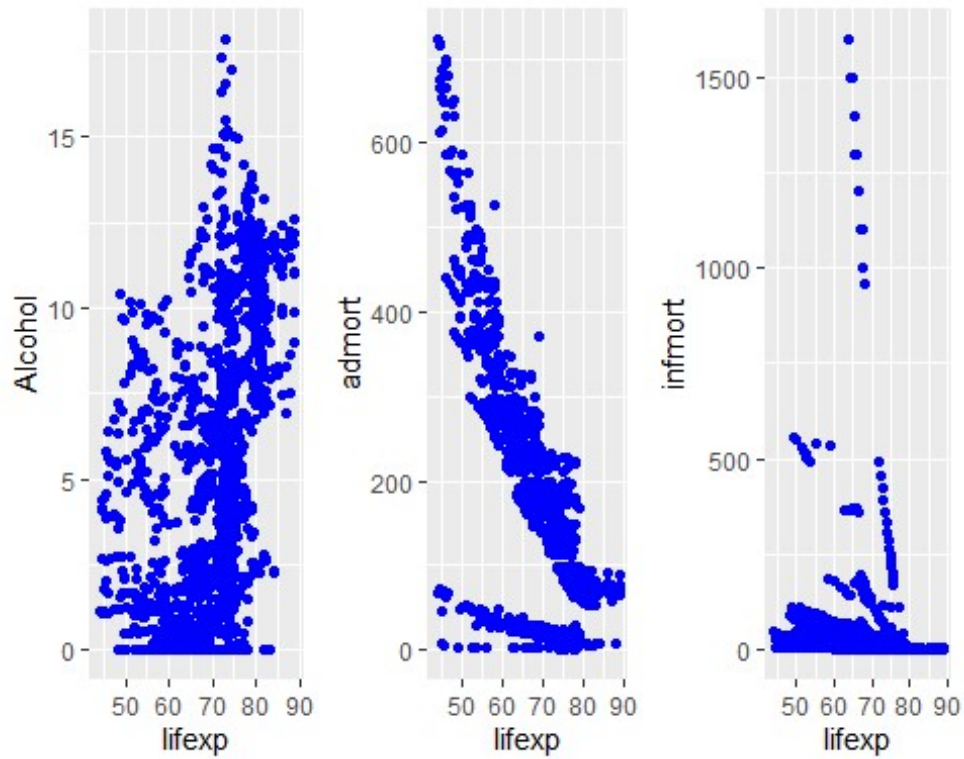
#ScatterPlot- Considering lifexp vs social, immunizations and economic factors.

```
e1<-ggplot(expect,aes(x=lifexp,y=Alcohol))+geom_point(color='blue')
e2<-ggplot(expect,aes(x=lifexp,y=admort))+geom_point(color='blue')
e3<-ggplot(expect,aes(x=lifexp,y=infmort))+geom_point(color='blue')

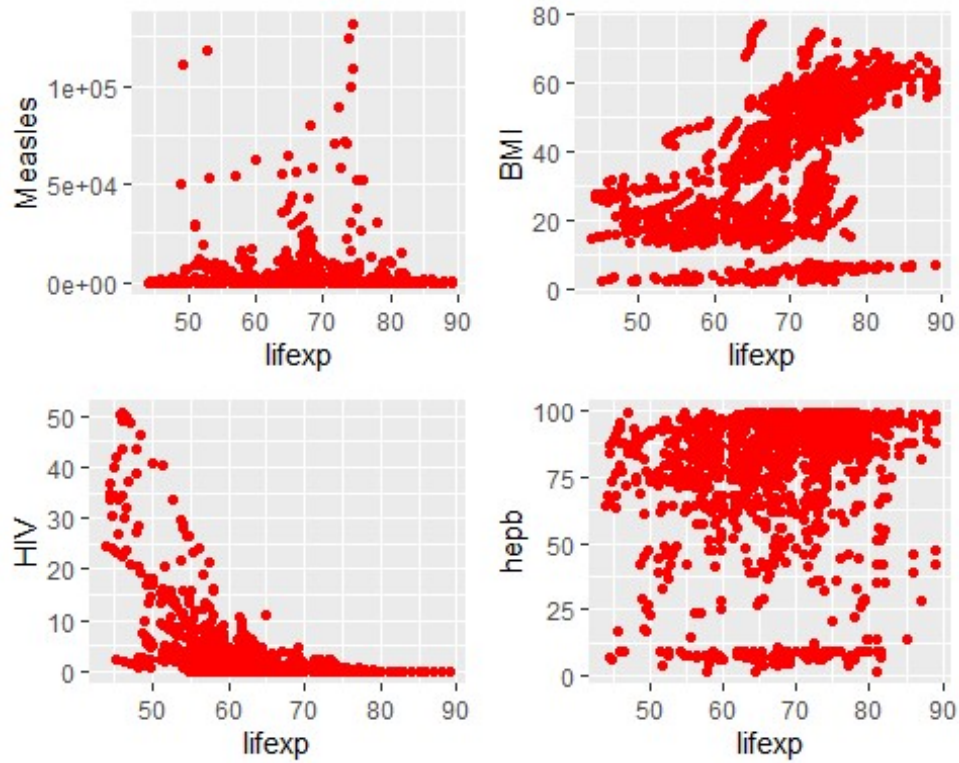
e4<-ggplot(expect,aes(x=lifexp,y=Polio))+geom_point(color='red')
e5<-ggplot(expect,aes(x=lifexp,y=Measles))+geom_point(color='red')
e6<-ggplot(expect,aes(x=lifexp,y=BMI))+geom_point(color='red')
e7<-ggplot(expect,aes(x=lifexp,y=HIV))+geom_point(color='red')
e8<-ggplot(expect,aes(x=lifexp,y=hepb))+geom_point(color='red')

e9<-ggplot(expect,aes(x=lifexp,y=GDP))+geom_point(color='green')
e10<-ggplot(expect,aes(x=lifexp,y=percexp))+geom_point(color='green')
e11<-ggplot(expect,aes(x=lifexp,y=totexp))+geom_point(color='green')

grid.arrange(e1,e2,e3,ncol=3)
```



```
grid.arrange(e5,e6,e7,e8,ncol=2,nrow=2)
```



```
grid.arrange(e9,e10,e11,ncol=3)
```

