End-Semester Project

Bivariate Analysis on GDP per capita, Sanitation and Life Expectancy across Nations in 2010

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1 Introduction

1.1 Overview

This presentation demonstrates the capabilities of *Bivariate Analysis* on datasets, to infer relationship between various features of Nations.

- log of GDP per capita: Logarithm (base e) of Gross Domestic Product (in \$) per citizen. Adjusted for Inflation. [lngdp]
- Sanitation Access %: Percentage of people using at least basic Sanitation facilities, not shared with other households. [snt]
- Life Expectancy: The average number of years a newly born child would live, provided current mortality patterns hold. [lfx]

1.2 Data

```
script.dir <- getSrcDirectory(function(x) {x})</pre>
setwd(script.dir)
numerise = function(x){
 x[grepl("k\$", x)] \leftarrow as.numeric(sub("k\$", "", x[grepl("k\$", x)]))*10^3
 x <- as.numeric(x)</pre>
 return(x)
}
d1_raw = read.csv(file.path(".", "Data", "gdp.csv"), fileEncoding = 'UTF-8-BOM')
d2_raw = read.csv(file.path(".","Data","sanitation.csv"), fileEncoding = 'UTF-8-BOM')
d3_raw = read.csv(file.path(".","Data","life_expectancy.csv"), fileEncoding = 'UTF-8-BOM')
yearname = "X2010"
d1 = d1 raw[!is.na(numerise(d1 raw[, yearname])),][,c("country", yearname)]
colnames(d1)[2] = "lngdp"
d2 = d2_raw[!is.na(numerise(d2_raw[, yearname])),][,c("country", yearname)]
colnames(d2)[2] = "snt"
d3 = d3_raw[!is.na(numerise(d3_raw[, yearname])),][,c("country", yearname)]
colnames(d3)[2] = "lfx"
dtemp = merge(x = d1, y = d2, by = "country")
d = merge(x = dtemp, y = d3, by = "country")
d$lngdp = log(numerise(d$lngdp))
write.csv(d, "./Data/assembled.csv")
kable(head(d, 6L))
```

country	lngdp	snt	lfx
Afghanistan	6.265301	34.9	60.5
Albania	8.183118	95.2	78.1
Algeria	8.273847	87.0	74.5
Andorra	10.454495	100.0	81.8
Angola	8.291547	41.1	60.2
Antigua and Barbuda	9.546813	86.3	75.9

2 Univariate Statistics

2.1 Measures of Central Tendency

Mean or Arithmetic Mean \bar{x} , Geometric Mean $\mathrm{GM}(x)$, Harmonic Mean $\mathrm{HM}(x)$, Median $\mathrm{median}(x)$ and Mode $\mathrm{mode}(x)$ are some measures of central tendency in the sample.

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FREE DATA FROM UN,

$$\begin{split} \bar{x} &= \frac{1}{n} \sum_{i=1}^n (x_i) \quad \text{GM}(x) = \sqrt[n]{\prod_{i=1}^n} a_i \quad \text{HM}(x) = n \sum_{i=1}^n x_i^{-1} \\ \text{median}(x) &= \begin{cases} x_{(n+1)/2} &: n = 1 \mod 2 \\ \frac{x_{(n/2)} + x_{((n/2)+1)}}{2} &: n = 0 \mod 2 \end{cases} \mod (x) = x_{(n)} \end{split}$$

```
getmode <- function(v) {</pre>
uniqv <- unique(v)</pre>
 freq = max(tabulate(match(v, uniqv)))
res = uniqv[which.max(tabulate(match(v, uniqv)))]
 if (freq == 1) res = NULL
 return(res)
}
d_central = data.frame(
 row.names = "Variable",
 Variable = c(
    "*ln(GDP)*",
    "*Sanitation*",
    "*Life Exp.*"
 ),
 Mean = c(
    mean(d$lngdp),
    mean(d$snt),
    mean(d$1fx)
 ),
 GM = c(
```

```
geometric.mean(d$lngdp),
    geometric.mean(d$snt),
    geometric.mean(d$lfx)
  ),
 HM = c(
    harmonic.mean(d$lngdp),
    harmonic.mean(d$snt),
    harmonic.mean(d$lfx)
  ),
  Median = c(
    median(d$lngdp),
    median(d$snt),
    median(d$lfx)
  ),
 Mode = c(
    getmode(d$lngdp),
    getmode(d$snt),
    getmode(d$1fx)
)
kable(
 d_central,
  col.names = c(
    "$\\bar{x}$",
    "$\\operatorname{GM}(x)$",
    "$\\operatorname{HM}(x)$",
    "$\\operatorname{median}(x)$",
    "$\\operatorname{mode}(x)$"
  ),
  digits=5
```

	\bar{x}	GM(x)	HM(x)	median(x)	mode(x)
$\overline{ln(GDP)}$	8.54124	8.42229	8.30248	8.48673	9.23014
Sanitation	72.43857	62.58904	47.61862	85.60000	100.00000
$Life\ Exp.$	70.54603	69.95538	69.28316	72.40000	73.20000

2.2 Measures of Dispersion

Range(x), Semi-int.. SIR(x), Mean Deviation about x' $MD_{(x')}(x)$, Variance s_x^2 , Standard Deviation s_x are some measures of dispersion in the sample.

Note: x_i is the ith observation. $x_{(i)}$ is the ith largest observation.

```
\mathrm{MD}_{(x')}(x) = \frac{\sum_{i=1}^{n} |x_i - x'|}{n} \quad \mathrm{SIR}(x) = \frac{|Q_1 - Q_3|}{2} \quad s_x = \sqrt{\frac{\sum_{i=1}^{n} \left(x_i - \bar{x}\right)^2}{n}} \quad s_x^2 = (s_x)^2
getmd = function(x, center = mean(x)){
  md = mean(
    abs(
       x - rep(center, length(x))
     )
  return(md)
}
d_disp = data.frame(
  row.names = "Variable",
  Variable = c(
     "*ln(GDP)*",
     "*Sanitation*",
    "*Life Exp.*"
  ),
  Range = c(
    max(d$lngdp) - min(d$lngdp),
    max(d$snt) - min(d$snt),
    max(d$lfx) - min(d$lfx)
  ),
  SIR = c(
    IQR(d$lngdp)/2,
     IQR(d\snt)/2,
    IQR(d\$1fx)/2
  ),
  MD = c(
     getmd(d$lngdp),
     getmd(d$snt),
    getmd(d$lfx)
  variance = c(
     (sd(d$lngdp))^2,
     (sd(d$snt))^2,
    (sd(d\$lfx))^2
  ),
  SD = c(
    sd(d$lngdp),
     sd(d$snt),
```

 $\mathrm{Range}(x) = |x_{(n)} - x_{(1)}| \qquad Q_1 = \mathrm{median}(x_{(1)}, \dots, x_{(\lfloor \frac{n+1}{2} \rfloor)}) \qquad Q_3 = \mathrm{median}(x_{(\lfloor \frac{n+2}{2} \rfloor)}, \dots, x_{(n)})$

sd(d\$lfx)

```
kable(
   d_disp,
   col.names = c(
     "$\operatorname{Range}(x)$",
     "$\\operatorname{SIR}(x)$",
     "$\\operatorname{MD}_{(\\bar{x})}(x)$",
     "$\\quad \\quad \\quad \\quad s_x^2$",
     "$\\quad \\quad \\quad \\quad s_x$"
),
   digits=5
)
```

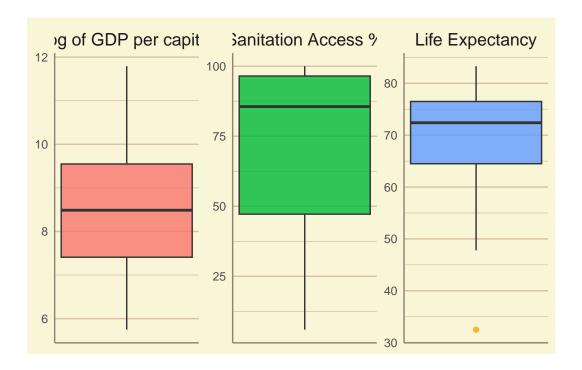
	$\operatorname{Range}(x)$	SIR(x)	$\mathrm{MD}_{(ar{x})}(x)$	s_x^2	s_x
ln(GDP)	6.04435	1.06914	1.17229	2.01791	1.42053
Sanitation	94.03000	24.65000	25.50487	872.29346	29.53461
Life	50.80000	6.00000	6.98712	75.33494	8.67957
Exp.					

2.3 Box Plot

About?

```
labelfunction = function(val1){
 return(list(c(
    "log of GDP per capita",
    "Sanitation Access %",
    "Life Expectancy"
 )))
ggplot(stack(d[2:4]), mapping = aes(y = values))+
geom_boxplot(aes(fill=ind), alpha=0.8, outlier.color = "orange")+
labs(
 x=NULL,
 y=NULL
 )+
mytheme+
mycolor+
facet_wrap(~ind, scales="free", labeller = labelfunction)+
  theme(axis.text.x=element_blank(),
        legend.position="none",
```

```
strip.text.x = element_text(size = rel(1.5)),
panel.grid.minor.x = element_blank(),
panel.grid.major.x = element_blank()
```



2.4 Inferences

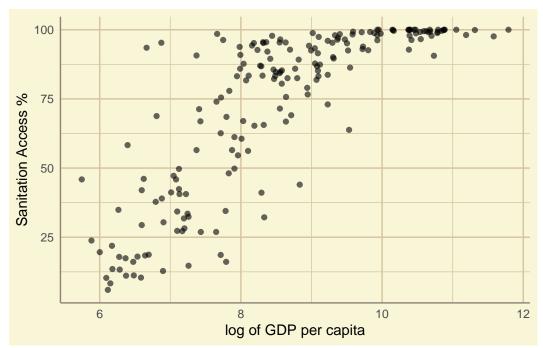
3 Scatter Plot

A *Scatter plot* is a type of Plot using Cartesian coordinate system to display values for two variables for a set of data. The data are displayed as a collection of points, each having one variable determining the *abscissa* and the other variable determining the *ordinate*. It helps us:

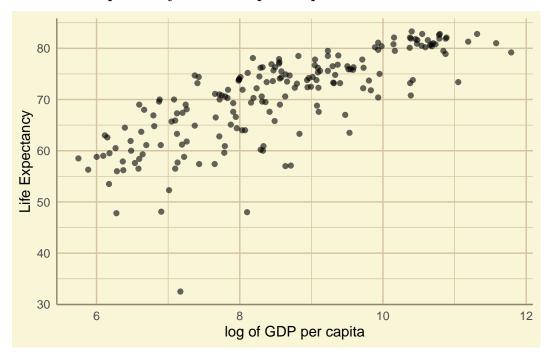
- take a short glance at effect of two variables.
- suggest kinds of correlations between variables.
- estimate the direction of correlation.

```
sctrplot = function(
   d, x_map, y_map,
   x_lab=waiver(), y_lab=waiver(),
   title=waiver()
){
```

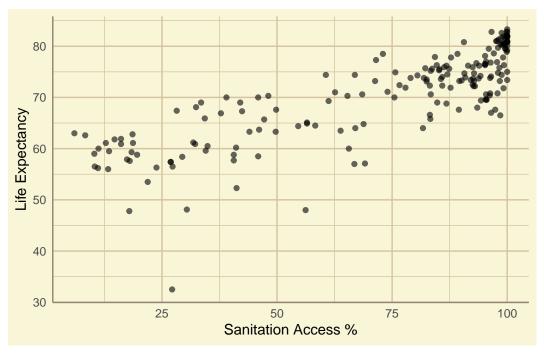
3.1 Sanitation vs. GDP per Capita



3.2 Life Expectancy vs. GDP per Capita



3.3 Life Expectation vs. Sanitation



3.4 Inferences

seems like Linear correlation

4 Bivariate Statistics

4.1 Covariance and Correlation Matrices

Covariance cov(x, y) is a measure of the joint variability of two random variables x, y.

Correlation $r_{x,y}$ is any relationship, causal or spurious, between two random variables x, y. Pearson's r correlation coefficient is considered here.

$$\mathrm{cov}(x,y) = \frac{\sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})}{n} \quad r_{x,y} = \frac{\mathrm{cov}(x,y)}{s_x s_x}$$

```
cov_mat = cov(d[, 2:4])
kable(cov_mat, digits=5)
```

	lngdp	snt	lfx
lngdp snt	2.01791 33.84045 9.52202	33.84045 872.29346 208.54155	9.52202 208.54155 75.33494

$$A_{i,j} = \mathrm{cov}(x_i, x_j)$$

```
cor_mat = cor(d[, 2:4])
kable(cor_mat, digits=5)
```

	lngdp	snt	lfx
lngdp	$ \begin{array}{c} 1.00000 \\ 0.80659 \\ 0.77229 \end{array} $	0.80659	0.77229
snt		1.00000	0.81351
lfx		0.81351	1.00000

$$A_{i,j} = r_{x_i,x_j}$$

4.2 Other Correlation Coefficients

Pearson, Spearman, Kendall #TODO

```
d_cor = data.frame(
 row.names = "Variable",
 Variable = c(
    "*Sanitation vs. ln(GDP)*",
    "*Life Exp. vs. ln(GDP)*",
    "*Life Exp. vs. Sanitation*"
 ),
 Pearson = c(
    cor(d$snt, d$lngdp, method="pearson"),
    cor(d$lfx, d$lngdp, method="pearson"),
    cor(d$lfx, d$snt, method="pearson")
 ),
  Spearman = c(
    cor(d$snt, d$lngdp, method="spearman"),
    cor(d$lfx, d$lngdp, method="spearman"),
    cor(d$lfx, d$snt, method="spearman")
  ),
```

```
Kendall = c(
    cor(d$snt, d$lngdp, method="kendall"),
    cor(d$lfx, d$lngdp, method="kendall"),
    cor(d$lfx, d$snt, method="kendall")
)

kable(
    d_cor,
    digit = 5,
    col.names = c(
        "*Pearson's* $r$",
        "*Spearman's* $r_s$",
        "*Kendall's* $\\tau$"
)
)
```

	Pearson's r	$Spearman's \ r_s$	Kendall's $ au$
Sanitation vs. $ln(GDP)$	0.80659	0.85920	0.67458
Life Exp. vs. $ln(GDP)$	0.77229	0.81639	0.62168
Life Exp. vs. Sanitation	0.81351	0.83513	0.63744

4.3 Partial Correlation

Partial

	Partial Correlation
$\overline{Sanitation \ vs. \ ln(GDP)}$	0.4826925
Life Exp. vs. $ln(GDP)$	0.3377892
Life Exp. vs. Sanitation	0.5075384

4.4 Inferences

Good linear correlation lets try to observe line of best fit.

5 Linear Regression

Simple Univariate Linear Regression is a method for estimating the relationship $y_i = f(x_i)$ of a response variable y with a predictor variable x, as a line that closely fits the y vs. x scatter plot.

$$y_i = \hat{a} + \hat{b}x_i + e_i.$$

Where \hat{a} is the *intercept*, \hat{b} is the *slope*, and e_i is the ith residual *error*. We aim to minimize e_i for better fit.

5.1 Ordinary Least Squares

Ordinary Least squares method reduces e_i by minimizing error sum of squares $\sum e_i^2$.

```
olssmry = function(
   d, x_map, y_map,
    x_lab=waiver(), y_lab=waiver(),
    title=waiver()
){
 model = lm(formula=y_map~x_map)
  smry = summary(model, signif.stars=TRUE)
  smryvec = c(
    as.numeric(model$coefficients["(Intercept)"]),
    as.numeric(model$coefficients["x_map"]),
    smry$r.squared
  return(smryvec)
}
olstab = t(data.frame(
 SvG = olssmry(d, d$lngdp, d$snt),
 LvG = olssmry(d, d$lngdp, d$lfx),
 LvS = olssmry(d, d$snt, d$lfx)
))
row.names(olstab) = c(
  "*Sanitation vs. ln(GDP)*",
 "*Life Exp. vs. ln(GDP)*",
  "*Life Exp. vs. Sanitation*"
kable(
  olstab,
 digit = 5,
 col.names=c(
 "$\\hat{a}$",
  "$\\hat{b}$",
  "$R^2$"
  )
)
```

	\hat{a}	\hat{b}	R^2
Sanitation vs. $ln(GDP)$	-70.79844	16.77006	0.65059
Life Exp. vs. $ln(GDP)$	30.24203	4.71876	0.59643
Life Exp. vs. Sanitation	53.22795	0.23907	0.66180

5.2 Least Absolute Deviation

Least absolute Deviation method reduces e_i by minimizing the sum of absolute deviations $\sum |e_i|$.

```
ladsmry = function(
   d, x_map, y_map,
    x_lab=waiver(), y_lab=waiver(),
    title=waiver()
){
 model = rq(formula=y_map~x_map)
  smry = summary(model)
  smryvec = c(
    as.numeric(model$coefficients[1]),
    as.numeric(model$coefficients[2])
  return(smryvec)
olstab = t(data.frame(
 SvG = ladsmry(d, d$lngdp, d$snt),
 LvG = ladsmry(d, d$lngdp, d$lfx),
 LvS = ladsmry(d, d$snt, d$lfx)
))
row.names(olstab) = c(
  "*Sanitation vs. ln(GDP)*",
  "*Life Exp. vs. ln(GDP)*",
 "*Life Exp. vs. Sanitation*"
kable(
 olstab,
 digit = 5,
  col.names=c(
  "$\\hat{a}$",
```

 R^2 : Coefficient of Determination

```
"$\\hat{b}$"
)
)
```

	\hat{a}	\hat{b}
$\overline{Sanitation \ vs. \ ln(GDP)}$	-71.23153	16.80472
Life Exp. vs. $ln(GDP)$	31.99047	4.61340
Life Exp. vs. Sanitation	53.73041	0.23963

5.3 Line fitting

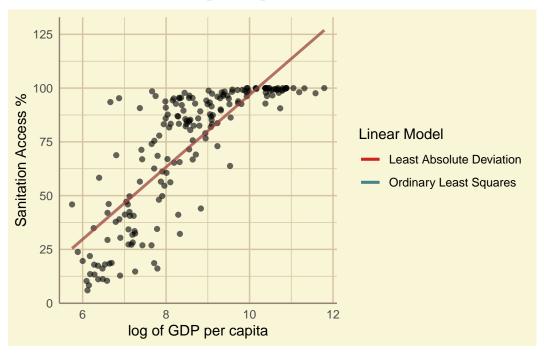
Plotting the estimated *Linear Model* on the Scatter Plot.

```
linearplot = function(
    d, x_map, y_map,
   x_lab=waiver(), y_lab=waiver(),
   title=waiver()
 plot1 = ggplot(d, mapping = aes(x = x_map, y = y_map))+
    geom_point(
      alpha=0.6
   )+
   mytheme+
   labs(
      x=x_lab,
      y=y_lab,
      title=title
    geom_smooth(
      method="lm",
      formula=y~x,
      se=FALSE,
      aes(color = "Ordinary Least Squares")
    )+
    geom_smooth(
      method="rq",
      formula=y~x,
      se=FALSE,
      aes(color = "Least Absolute Deviation")
   )+
    labs(
      color="Linear Model"
    )+
```

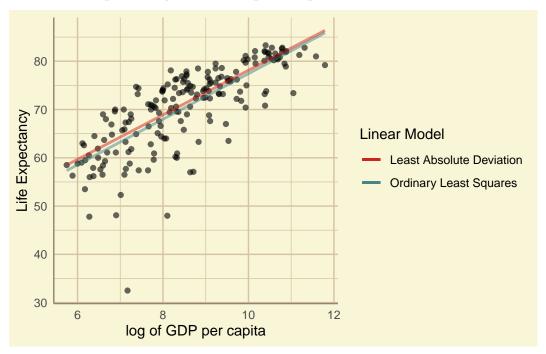
```
mycolor

return(plot1)
}
```

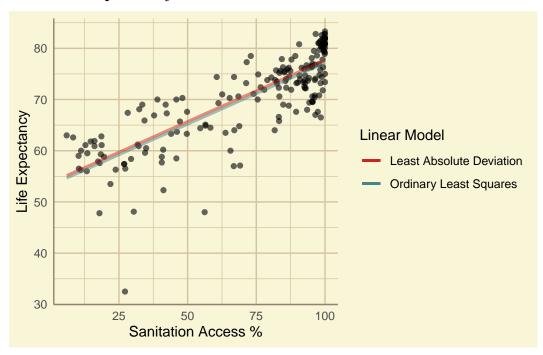
5.4 Sanitation vs. GDP per Capita



5.5 Life Expectancy vs. GDP per Capita



5.6 Life Expectancy vs. Sanitation



5.7 Inferences

6 Conclusion