CA3aVdVeen

Ties van der Veen 12-9-2019

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I. Introduction to the assignment

Packages

```
options(repos="https://cran.rstudio.com")
install.packages("jtools")
## Installing package into 'C:/Users/tiess/OneDrive/Documenten/R/win-library/3.6'
## (as 'lib' is unspecified)
## package 'jtools' successfully unpacked and MD5 sums checked
## The downloaded binary packages are in
## C:\Users\tiess\AppData\Local\Temp\RtmpsvZvvT\downloaded_packages
install.packages("huxtable")
## Installing package into 'C:/Users/tiess/OneDrive/Documenten/R/win-library/3.6'
## (as 'lib' is unspecified)
## package 'huxtable' successfully unpacked and MD5 sums checked
##
## The downloaded binary packages are in
## C:\Users\tiess\AppData\Local\Temp\RtmpsvZvvT\downloaded_packages
install.packages("ggstance")
## Installing package into 'C:/Users/tiess/OneDrive/Documenten/R/win-library/3.6'
## (as 'lib' is unspecified)
## package 'ggstance' successfully unpacked and MD5 sums checked
## The downloaded binary packages are in
## C:\Users\tiess\AppData\Local\Temp\RtmpsvZvvT\downloaded_packages
install.packages("summarytools")
## Installing package into 'C:/Users/tiess/OneDrive/Documenten/R/win-library/3.6'
## (as 'lib' is unspecified)
```

```
## package 'summarytools' successfully unpacked and MD5 sums checked
##
## The downloaded binary packages are in
## C:\Users\tiess\AppData\Local\Temp\RtmpsvZvvT\downloaded_packages
install.packages("pwr")
## Installing package into 'C:/Users/tiess/OneDrive/Documenten/R/win-library/3.6'
## (as 'lib' is unspecified)
## package 'pwr' successfully unpacked and MD5 sums checked
## The downloaded binary packages are in
## C:\Users\tiess\AppData\Local\Temp\RtmpsvZvvT\downloaded_packages
install.packages("knitr")
## Installing package into 'C:/Users/tiess/OneDrive/Documenten/R/win-library/3.6'
## (as 'lib' is unspecified)
## package 'knitr' successfully unpacked and MD5 sums checked
## The downloaded binary packages are in
## C:\Users\tiess\AppData\Local\Temp\RtmpsvZvvT\downloaded_packages
install.packages("lemon")
## Installing package into 'C:/Users/tiess/OneDrive/Documenten/R/win-library/3.6'
## (as 'lib' is unspecified)
## package 'lemon' successfully unpacked and MD5 sums checked
##
## The downloaded binary packages are in
## C:\Users\tiess\AppData\Local\Temp\RtmpsvZvvT\downloaded_packages
install.packages("tinytex")
## Installing package into 'C:/Users/tiess/OneDrive/Documenten/R/win-library/3.6'
## (as 'lib' is unspecified)
## package 'tinytex' successfully unpacked and MD5 sums checked
## The downloaded binary packages are in
## C:\Users\tiess\AppData\Local\Temp\RtmpsvZvvT\downloaded_packages
install.packages("AER")
## Installing package into 'C:/Users/tiess/OneDrive/Documenten/R/win-library/3.6'
## (as 'lib' is unspecified)
```

```
## package 'AER' successfully unpacked and MD5 sums checked
##
## The downloaded binary packages are in
## C:\Users\tiess\AppData\Local\Temp\RtmpsvZvvT\downloaded_packages
library(foreign)
library(tidyverse)
## -- 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 0.8.3 v stringr 1.4.0
## v readr
          1.3.1 v forcats 0.4.0
## -- Conflicts -----
                             ## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                   masks stats::lag()
library(ggdag)
##
## Attaching package: 'ggdag'
## The following object is masked from 'package:ggplot2':
##
##
      expand_scale
## The following object is masked from 'package:stats':
##
##
      filter
library(dplyr)
library(tinytex)
library(jtools)
library(huxtable)
##
## Attaching package: 'huxtable'
## The following objects are masked from 'package:ggdag':
##
      label, label<-
##
## The following object is masked from 'package:dplyr':
##
##
      add_rownames
## The following object is masked from 'package:purrr':
##
##
      every
```

```
## The following object is masked from 'package:ggplot2':
##
##
       theme_grey
library(summarytools)
## Registered S3 method overwritten by 'pryr':
##
     method
                 from
     print.bytes Rcpp
##
##
## Attaching package: 'summarytools'
## The following objects are masked from 'package:huxtable':
##
##
       label, label<-
## The following objects are masked from 'package:ggdag':
##
##
       label, label<-
## The following object is masked from 'package:tibble':
##
##
       view
library(ggstance)
##
## Attaching package: 'ggstance'
## The following objects are masked from 'package:ggplot2':
##
##
       geom_errorbarh, GeomErrorbarh
library(pwr)
library(knitr)
library(lemon)
## Attaching package: 'lemon'
## The following object is masked from 'package:purrr':
##
##
       %11%
library(AER)
## Loading required package: car
```

```
##
## Attaching package: 'car'
  The following object is masked from 'package:dplyr':
##
##
       recode
##
  The following object is masked from 'package:purrr':
##
##
       some
## Loading required package: lmtest
## Loading required package: zoo
##
## Attaching package: 'zoo'
## The following objects are masked from 'package:base':
##
##
       as.Date, as.Date.numeric
## Loading required package: sandwich
## Loading required package: survival
knit_print.data.frame <- lemon_print</pre>
st_options(plain.ascii = FALSE, style = "rmarkdown")
st css()
## <style type="text/css">
           background-color: transparent; border: 0; } .st-table td, .st-table th {
                                                                                            padding: 8px;
Data
theUrl_ca3a <- "https://surfdrive.surf.nl/files/index.php/s/aRyyi4GbwUjF9DF/download"
colonies <- read.dta (file = theUrl_ca3a)</pre>
colonies <- colonies[colonies$excolony==1, ]</pre>
```

II. Potential outcomes

Loading required package: carData

(a) Y(0,i) = low GDP per capita Y(1,i) = high GDP per capita

colonies <- colonies[is.na(colonies\$excolony)==FALSE,]</pre>

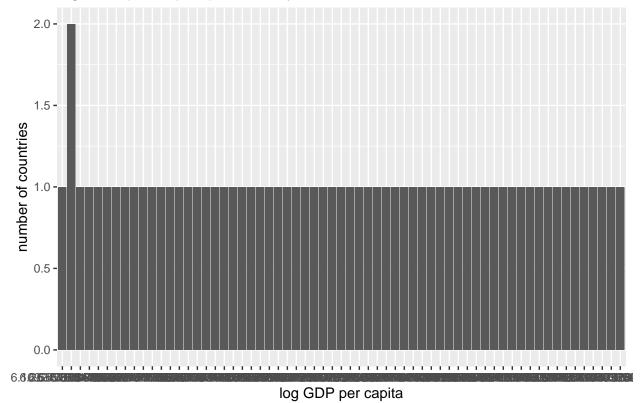
(b) If colonists came to settle, they brought in institutions that helped the country flourish. If they did the opposite and just extracted resources, these institutions were not established and the country had less of an opportunity to develop. These development differences explain current GDP per capita.

III. Descriptive statistics

```
ggplot(colonies, aes(x=as.factor(logpgp95)))+
  geom_histogram(stat='count', binwidth = "0.5")+
  labs(x='log GDP per capita', y='number of countries', title='log GDP per capita per country')
```

Warning: Ignoring unknown parameters: binwidth, bins, pad

log GDP per capita per country



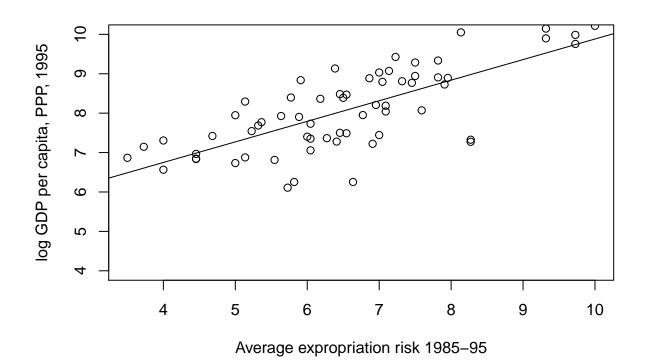
```
summary(is.na(colonies$logpgp95))
```

```
## Mode FALSE
## logical 64
```

No output, so no missing entries.

IV. Estimating the treatment effect using an instrumental variable

(a)



(b)

```
reg3a1 <- lm(logpgp95 ~ avexpr, data=colonies)
summ(reg3a1, confint=TRUE)</pre>
```

```
## MODEL INFO:
## Observations: 64
## Dependent Variable: logpgp95
## Type: OLS linear regression
##
## MODEL FIT:
## F(1,62) = 72.82, p = 0.00
## R<sup>2</sup> = 0.54
## Adj. R<sup>2</sup> = 0.53
##
## Standard errors: OLS
##
## Est. 2.5% 97.5% t val. p
```

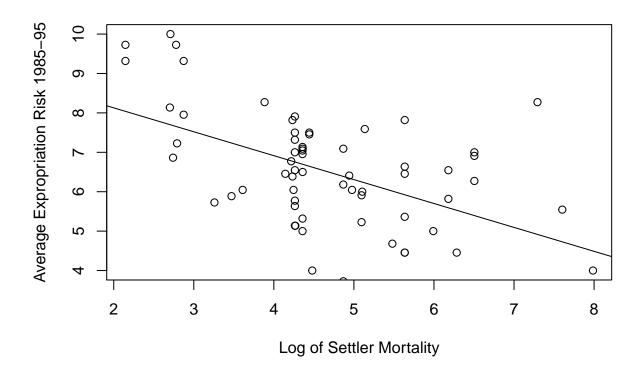
```
## (Intercept) 4.66 3.84 5.48 11.41 0.00
## avexpr 0.52 0.40 0.64 8.53 0.00
```

The regression results suggest a higher GDP per capita for countries with good institutions compared to the countries that don't have good institutions.

(c)

There are multiple confounding factors that influence this result. Seeing this as a causal relationship would be shortsighted, especially since there's such a gap between the assumed variable's origin (strategy of colonization) and "current' (1995) GDP per capita.

(d)



(e)

```
summ(reg3a2, confint=TRUE)
## MODEL INFO:
## Observations: 64
## Dependent Variable: logpgp95
## Type: OLS linear regression
##
## MODEL FIT:
## F(1,62) = 56.60, p = 0.00
## R^2 = 0.48
## Adj. R^2 = 0.47
##
## Standard errors: OLS
                  Est. 2.5% 97.5% t val.
## ----- ---- -----
## (Intercept)
                  10.73 10.00 11.46 29.22 0.00
                  -0.57 -0.73 -0.42
                                       -7.52 0.00
## logem4
```

reg3a2 <- lm(logpgp95 ~ logem4, data=colonies)</pre>

The instrumental variable is statistically significant (p=0.00) as can be seen in the regression output. Economically this variable is not that significant, since there is no causal relationship between mortality in the 1800s and current economic welfare. However, this does not mean that the variable is useless; far from it. It does its job as an instrumental variable.

(f)

```
reg3a3 <- lm(avexpr ~ logem4, data=colonies)
summ(reg3a3, confint=TRUE)</pre>
```

```
## MODEL INFO:
## Observations: 64
## Dependent Variable: avexpr
## Type: OLS linear regression
##
## MODEL FIT:
## F(1,62) = 22.95, p = 0.00
## R^2 = 0.27
## Adj. R^2 = 0.26
##
## Standard errors: OLS
## -----
##
                Est. 2.5% 97.5% t val.
## ----- ---- -----
                9.34 8.12 10.56 15.30 0.00
## (Intercept)
## logem4
                 -0.61 -0.86 -0.35
                                   -4.79 0.00
```

These results show that mortality rate has a sizable effect on settlement strategy, which is what was already predicted. Thus, with the significance reported, these results support the theory.

The F-test reported here gives F=22.95, which is big enough (considering the minimum of 10).

(g)

```
avexpr_hat <- predict(reg3a3)</pre>
colonies$pred_avexpr=avexpr_hat
reg3a4 <- lm(logpgp95 ~ pred_avexpr, data=colonies)
summ(reg3a4, confint=TRUE)
## MODEL INFO:
## Observations: 64
## Dependent Variable: logpgp95
## Type: OLS linear regression
##
## MODEL FIT:
## F(1,62) = 56.60, p = 0.00
## R^2 = 0.48
## Adj. R^2 = 0.47
## Standard errors: OLS
               Est. 2.5% 97.5% t val. p
## ----- ---- -----
## (Intercept) 1.91 0.26
                              3.56
                                    2.32 0.02
## pred_avexpr
                 0.94 0.69 1.20 7.52 0.00
## -----
```

By taking the prediction of the colonization strategy (avexpr), it seems like almost all of the variation in log GDP per capita (logpgp95) is explained. From (b), which showed an estimate of 0.52, this is a big jump.

(h)

```
reg3a5 <- ivreg(logpgp95 ~ avexpr | logem4, data=colonies)
summary(reg3a5, confint=TRUE, diagnostics=TRUE)</pre>
```

```
##
## ivreg(formula = logpgp95 ~ avexpr | logem4, data = colonies)
##
## Residuals:
       Min 1Q Median
                                3Q
                                       Max
## -2.44903 -0.56242 0.07311 0.69564 1.71752
##
## Coefficients:
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 1.9097 1.0267 1.860 0.0676
                        0.1565 6.033 9.8e-08 ***
## avexpr
               0.9443
## Diagnostic tests:
##
                 df1 df2 statistic p-value
## Weak instruments 1 62 22.95 1.08e-05 ***
## Wu-Hausman 1 61
                           24.22 6.85e-06 ***
                  O NA
## Sargan
                              NA
                                       NA
```

```
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.9483 on 62 degrees of freedom
## Multiple R-Squared: 0.187,
                              Adjusted R-squared: 0.1739
## Wald test: 36.39 on 1 and 62 DF, p-value: 9.799e-08
 (i)
reg3a6 <- ivreg(logpgp95 ~ avexpr + lat_abst | lat_abst + logem4, data=colonies)
summary(reg3a6, confint=TRUE, diagnostics=TRUE)
##
## Call:
  ivreg(formula = logpgp95 ~ avexpr + lat_abst | lat_abst + logem4,
##
       data = colonies)
##
## Residuals:
##
      Min
                1Q Median
                                3Q
                                       Max
## -2.5611 -0.6557 0.0732 0.7572
                                   1.8803
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                 1.6918
                            1.2930
                                     1.308
                                              0.196
                 0.9957
                            0.2217
                                     4.492 3.21e-05 ***
## avexpr
## lat abst
                -0.6472
                            1.3351 -0.485
                                              0.630
##
## Diagnostic tests:
##
                    df1 df2 statistic p-value
## Weak instruments
                        61
                                13.09 0.000603 ***
                     1
## Wu-Hausman
                                18.75 5.75e-05 ***
                      1
                         60
                      0
                         NA
## Sargan
                                   NA
                                            NA
##
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.005 on 61 degrees of freedom
## Multiple R-Squared: 0.1025, Adjusted R-squared: 0.07305
## Wald test: 17.01 on 2 and 61 DF, p-value: 1.351e-06
```

Including this covariate allows us to control for environmental situations that would give a certain country an advantage (or disadvantage) over the other, creating differences in GDP growth. The new estimation results show that including this covariate makes the colonization strategy (avexpr) able to pretty much explain the whole effect on GDP per capita.

(j)

The exclusion restriction means that the independent variables do not directly impact the dependent variables. The instrumental variable is used when this exclusion restriction cannot be guaranteed through randomization. In this case, the instrumental variable "mortality rate of settlers in the 1800s" is unlikely to affect other variables in the study, but can be used to check the independence of the colonization strategy variable. Thus it is likely the effect will hold.