```
library(tidycensus)
library(tidyverse)
library(ggspatial)
library(dplyr)
library(ggplot2)
library(ggthemes)
library(fitdistrplus)
library(tibble)
library(skedastic)
library(lmtest)
census_api_key("74060ca8061c55486a9d98bfca2f006fb9772339",install=T)
# 03
variables <- c(</pre>
'DP05_0001E','DP05_0018E','DP03_0062E','DP02_0065PE','DP03_0096PE','DP03_0128PE','DP04_
data <- get_acs(geography = 'tract',</pre>
                 state='IL',
                 county='Cook County',
                output='wide',
                year=2019,
                 geometry=TRUE,
                 variables=variables,
                 show_call=TRUE)
drop.cols <- grep("*M$", colnames(data))</pre>
data <- data[-(drop.cols)]</pre>
colnames(data) <- c(</pre>
'geoid', 'name', 'totpop', 'medage', 'medhhinc', 'propbac', 'propcov', 'proppov', 'proprent', 'g
# 04
ggplot(data) +
  geom_sf(aes(fill=propbac), alpha = 0.8) +
  annotation_scale(bar_cols = c("yellow2", "green")) +
  scale_fill_viridis_c(option='turbo') + theme_economist_white() +
  ggtitle(label = "AES 5-year (2015-2019) Estimates for Cook County, IL",
subtitle = "\nBaccalaureate Attainment Rates")
model <- lm(propbac ~ medhhinc, data)</pre>
summary(model)
# 05
ggplot(data, aes(medhhinc,propbac)) +
  geom_point(color='red') +
  geom_smooth(method = 'lm') +
  theme_economist() +
  labs(x=substitute(paste(bold('Median Household Income ($)'))),
y=substitute(paste(bold('Baccalaureate Attainment Rates (%)'))), title =
'AES 5-year (2015-2019) Estimates for Cook County, IL') +
  theme(
    axis.title.y = element_text(vjust = +4),
    axis.title.x = element_text(vjust = -4)
  )
# 06
mu_resd <- mean(model$residuals)</pre>
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sd_resd <- sd(model$residuals)</pre>
len_resd <- length(model$residuals)</pre>
# ks-test check for normality
test <- ks.test(model$residuals, pnorm, mu_resd, sd_resd)</pre>
format(test$p.value, scientific=FALSE)
# plot to check normality of residuals
frame <- enframe(model$residuals)</pre>
x \leftarrow seq(-25, 25, length=len_resd)
ggplot(frame, aes(x=value)) +
  geom_histogram(fill='white', colour='black', alpha = 0.75, breaks =
seq(-30, 30, by = 2)) +
  stat_function(fun = function(x) dnorm(x, mean = mu_resd, sd = sd resd) *
len_resd * 2, color='red', size=1) +
  theme_economist() +
  theme(
    axis.title.y = element_text(vjust = +4),
    axis.title.x = element_text(vjust = -4),
    plot.title = element_text(hjust = 0.5)
  ) +
  labs(x=substitute(paste(bold('Residual Value'))),
y=substitute(paste(bold('Count')))) +
  ggtitle(label = "Histogram for Residuals of the Model")
# test for serial correlation
dwtest(model)
# plot for serial correlation
par(mfrow = c(1,1))
df.residuals = data.frame(index(model$residuals), model$residuals)
colnames(df.residuals) <- c("Index", "Residuals")</pre>
ggplot(data=df.residuals, aes(x=Index, y=Residuals)) +
  geom_point(col='red', size=2) + ylim(-25, 25) + xlim(0, 1200) +
  theme_economist() + labs(x=substitute(paste(bold("Index"))),
y=substitute(paste(bold('Residuals')))) +
  ggtitle(label = "Plot for Serial Correlation of Residuals") +
  theme(
    axis.title.y = element_text(vjust = +4),
    axis.title.x = element_text(vjust = -4),
    plot.title = element_text(hjust = 0.5)
acf(model$residuals, type = "correlation", lag.max = 1)
# tests for heteroskedasticity
bptest(model)
#white_lm(model)
# plot for heteroskedasticity
ggplot(model, aes(y=model$residuals, x=na.omit(data$medhhinc))) +
  geom_point(col = 'red') + geom_abline(slope = 0) +
  theme_economist() +
  theme(
    axis.title.y = element_text(vjust = +4),
    axis.title.x = element_text(vjust = -4),
    plot.title = element_text(hjust = 0.5)
  labs(x=substitute(paste(bold('Median Household Income'))),
y=substitute(paste(bold('Model Residuals')))) +
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ggtitle(label = "Plot for Heteroskedasticity")
# 07
cor_og <- cor(data$propbac, data$medhhinc, use = "complete.obs")</pre>
samples <- replicate(10000, sample(data$medhhinc, nrow(data),</pre>
replace=TRUE))
cors <- c()
for (i in 1:10000) {
  cors <- c(cors, cor(data$propbac, samples[,i], use = "complete.obs"))</pre>
significant_links <- cors[cors > cor_og]
length(significant_links) / length(cors)
# 08
qplot(seq_along(cors), cors) +
  geom_point(color='red') +
  theme economist() +
  labs(x=substitute(paste(bold(''))), y=substitute(paste(bold('')))) +
  ggtitle(label = "AES 5-year (2015-2019) Estimates for Cook County, IL",
subtitle = "\nCorrelation between 10k samples of Median Household Income
and Baccalaureate Attainment Rates") +
  theme(
    axis.title.y = element_text(vjust = +4),
    axis.title.x = element_text(vjust = -4)
  )
ggplot(enframe(cors), aes(x=value)) +
  geom_histogram(color="black", fill="red") +
  theme_economist() +
  labs(x=substitute(paste(bold('Correlation'))),
y=substitute(paste(bold('Count')))) +
  ggtitle(label = "AES 5-year (2015-2019) Estimates for Cook County, IL",
subtitle = "\nCorrelation between 10k samples of Median Household Income
and Baccalaureate Attainment Rates") +
    axis.title.y = element_text(vjust = +4),
    axis.title.x = element_text(vjust = -4)
  )
# 010
model.intercept <- model$coefficients[1]</pre>
model.slope <- model$coefficients[2]</pre>
slopes \leftarrow seq(-3 * model.slope, 5 * model.slope, 0.001 * model.slope)
sums_of_squares <- c()</pre>
sum_of_squares.get <- function(slope = model.slope, intercept =</pre>
model.intercept) {
  propbac.prediction <- intercept + slope * data$medhhinc</pre>
  sums_of_squares <<- c(sums_of_squares, sum((propbac.prediction -</pre>
data$propbac) ** 2, na.rm = TRUE))
mapply(sum_of_squares.get , slopes, model.intercept)
df.sse <- data.frame(slopes, sums_of_squares)</pre>
colnames(df.sse) <- c('Slope', 'SSE')</pre>
ggplot(df.sse,aes(x=Slope, y=SSE)) +
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geom_point(color='red', size = 1) +
  theme economist() +
  labs(x=substitute(paste(bold('Slope'))),
y=substitute(paste(bold('SSE')))) +
  ggtitle(label = "Sum of Square of Residuals vs Slope", subtitle =
"\nModel: Baccalaureate Attainment Rates vs Median Household Income") +
  theme(
    axis.title.y = element_text(vjust = +4),
    axis.title.x = element_text(vjust = -4)
  ) +
  annotate("point", x=0.0002577586, y=101495.9, color="Black") +
  annotate("text", x=0.0002577586, y=-150000, label="(0.0002577586,
101495.9)")
# 011
intercepts <- seq(-8 * model.intercept, 10 * model.intercept, 0.001 *
model.intercept)
log_likelihood <- c()</pre>
log_likelihood.get <- function(slope = model.slope, intercept =</pre>
model.intercept) {
  propbac.prediction <- intercept + slope * data$medhhinc</pre>
  residuals <- data$propbac - propbac.prediction
  log_likelihood <<- c(log_likelihood, sum(dnorm(residuals, 0,</pre>
summary(model)$sigma, log=TRUE), na.rm = TRUE))
mapply(log likelihood.get , model.slope, intercepts)
df.loglik <- data.frame(intercepts, log_likelihood)</pre>
colnames(df.loglik) <- c('Intercept', 'LogLikelihood')</pre>
ggplot(df.loglik,aes(x=Intercept, y=LogLikelihood)) +
  geom_point(color='red', size = 1) +
  theme_economist() +
  labs(x=substitute(paste(bold('Intercept'))),
y=substitute(paste(bold('Log Likelihood')))) +
  ggtitle(label = "Log Likelihood of Linear Model vs Intercept", subtitle
= "\nModel: Baccalaureate Attainment Rates vs Median Household Income") +
  theme(
    axis.title.y = element_text(vjust = +4),
    axis.title.x = element_text(vjust = -4)
  ) +
  annotate("point", x=4.302643, y=-4720.427, color="Black") +
  annotate("text", x=4.302643, y=-4300, label="(4.302643, -4720.427)")
# 012
medhhinc.robin_hood_policy <- sort(data$medhhinc, index.return=TRUE,</pre>
decreasing=TRUE)
len.medhhin <- length(medhhinc.robin_hood_policy$x)</pre>
medhhinc.robin_hood_policy$x[1:50] <- medhhinc.robin_hood_policy$x[1:50] -</pre>
10000
medhhinc.robin_hood_policy$x[-c(1:(len.medhhin-50))] <-</pre>
medhhinc.robin_hood_policy$x[-c(1:(len.medhhin-50))] + 10000
# Reversing the earlier sort to preserve the relationship between
# data points of propbac and medhhinc
medhhinc.robin_hood_policy <- data.frame(medhhinc.robin_hood_policy)</pre>
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medhhinc.robin_hood_policy <-</pre>
medhhinc.robin_hood_policy[order(medhhinc.robin_hood_policy$ix), ]
propbac.robin_hood_prediction <- predict(model,</pre>
data.frame(medhhinc=medhhinc.robin_hood_policy$x))
# Plot to visualize the effect of Robin Hood Tax Policy on Baccalaureate
Attainment Rates
cols <- c("Before Robin Hood Tax Policy"="Blue", "After Robin Hood Tax
Policy"="Red")
ggplot() +
  geom_point(aes(x=index(na.omit(data$propbac)), y=na.omit(data$propbac),
colour="Before Robin Hood Tax Policy")) +
  geom_point(aes(x=index(propbac.robin_hood_prediction),
y=propbac.robin_hood_prediction, colour="After Robin Hood Tax Policy")) +
  theme_economist() + labs(x=substitute(""),
y=substitute(paste(bold('Baccalaureate Attainment Rates')))) +
  ggtitle(label = "Effect of Robin Hood Policy on Baccalaureate Attainment
Rates") +
  theme(
    axis.title.y = element_text(vjust = +4),
    axis.title.x = element_text(vjust = -4),
    plot.title = element_text(hjust = 0.5),
    legend.text.align = 1
  ) +
  scale_colour_manual(name="", values=cols)
cols <- c("Before Robin Hood Tax Policy"="Blue","After Robin Hood Tax
Policy"="Red")
ggplot() +
  geom histogram(aes(x=na.omit(data$propbac), fill="Before Robin Hood Tax
Policy"), alpha=0.5) +
  geom_histogram(aes(x=propbac.robin_hood_prediction, fill="After Robin
Hood Tax Policy"), , alpha=0.5) +
  theme_economist() + labs(x=substitute(""),
y=substitute(paste(bold('Baccalaureate Attainment Rates')))) +
  ggtitle(label = "Effect of Robin Hood Policy on Baccalaureate Attainment
Rates") +
  theme(
    axis.title.y = element_text(vjust = +4),
    axis.title.x = element_text(vjust = -4),
    plot.title = element_text(hjust = 0.5),
    legend.text.align = 1
  scale_colour_manual(name="", values=cols, aesthetics = "fill")
# Theoretical Comparison of Baccalaureate Attainment Rates before and
after
# Robin Hood Tax Policy
mean(data$propbac, na.rm=TRUE)
mean(propbac.robin_hood_prediction)
var(data$propbac, na.rm=TRUE)
var(propbac.robin_hood_prediction)
```