homework3 report

intersect, setdiff, setequal, union

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##

##

2024-07-02

```
if (!require(ggplot2)) install.packages("ggplot2")

## 穀入需要的程序包: ggplot2

library(ggplot2)

gmp <- read.table("data/gmp.dat")

gmp$pop <- round(gmp$gmp/gmp$pcgmp)

1.

library(dplyr)

## ## 穀入程序包: 'dplyr'

## filter, lag

## The following objects are masked from 'package:stats':

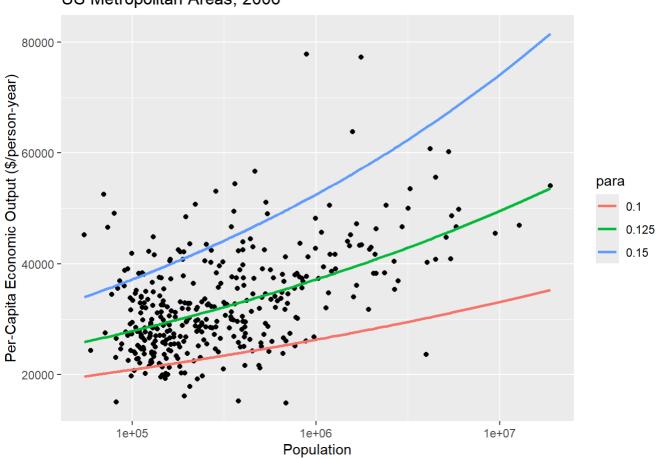
## ## filter, lag

### The following objects are masked from 'package:base':
```

```
library(tidyr)
func \leftarrow function (x, y0, a) {
  return(y0 * x ^ a)
df <- gmp |>
  select(pop, pcgmp) |>
  mutate("0.125" = func(pop, 6611, 1/8),
         "0.1" = func(pop, 6611, 0.1),
         "0.15" = func(pop, 6611, 0.15)) |>
  gather(key = para, value = val, -pop, -pcgmp)
df |>
  ggplot() +
  scale_x_log10() +
  geom_point(aes(x = pop, y = pcgmp)) +
  geom_line(aes(x = pop, y = val, col = para), size = 1) +
  labs(x = "Population", y = "Per-Capita Economic Output ($/person-year)",
  title = "US Metropolitan Areas, 2006")+
rm(df)
```

```
## Warning: Using `size` aesthetic for lines was deprecated in ggplot2 3.4.0.
## i Please use `linewidth` instead.
## This warning is displayed once every 8 hours.
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
## generated.
```

US Metropolitan Areas, 2006



```
mse <- function(para, N = gmp$pop, Y = gmp$pcgmp) {
   return(mean((Y - para[1] * N ^ para[2]) ^ 2))
 mse (c (6611, 0.15))
 ## [1] 207057513
 mse(c(5000, 0.10))
 ## [1] 298459914
3.
 n1m(mse, c(y0=6611, a=1/8))
 ## Warning in nlm(mse, c(y0 = 6611, a = 1/8)): NA/Inf被换成最大的正值
 ## Warning in nlm(mse, c(y0 = 6611, a = 1/8)): NA/Inf被换成最大的正值
 ## Warning in nlm(mse, c(y0 = 6611, a = 1/8)): NA/Inf被换成最大的正值
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 ## Warning in nlm(mse, c(y0 = 6611, a = 1/8)): NA/Inf被换成最大的正值
 ## Warning in nlm(mse, c(y0 = 6611, a = 1/8)): NA/Inf被换成最大的正值
 ## $minimum
 ## [1] 61857060
 ##
 ## $estimate
 ## [1] 6611.0000000
                    0. 1263177
 ##
 ## $gradient
 ## [1] 50.048639 -9.983778
 ##
 ## $code
 ## [1] 2
 ##
 ## $iterations
 ## [1] 3
 n1m(mse, c(y0=6000, a=0.1))
 ## Warning in nlm(mse, c(y0 = 6000, a = 0.1)): NA/Inf被换成最大的正值
 ## Warning in nlm(mse, c(y0 = 6000, a = 0.1)): NA/Inf被换成最大的正值
 ## Warning in nlm(mse, c(y0 = 6000, a = 0.1)): NA/Inf被换成最大的正值
 ## Warning in nlm(mse, c(y0 = 6000, a = 0.1)): NA/Inf被换成最大的正值
```

Warning in nlm(mse, c(y0 = 6000, a = 0.1)): NA/Inf被换成最大的正值 ## Warning in nlm(mse, c(y0 = 6000, a = 0.1)): NA/Inf被换成最大的正值

```
nlm(mse, c(y0=6500, a=0.15))
```

minimum 表示其找到的函数最小值, estimate 表示找到 minimum 值时对应的变量数值。

4.

```
plm(y0 = 6611, a = 0.15)
```

```
## $y0
## [1] 6611
##
## $a
## [1] 0.1263182
##
## $MSE
## [1] 61857060
```

```
p1m(y0 = 5000, a = 0.1)
```

```
## Warning in nlm(mse, c(y0 = y0, a = a), N = N, Y = Y): NA/Inf被换成最大的正值
## Warning in nlm(mse, c(y0 = y0, a = a), N = N, Y = Y): NA/Inf被换成最大的正值
## Warning in nlm(mse, c(y0 = y0, a = a), N = N, Y = Y): NA/Inf被换成最大的正值
## Warning in nlm(mse, c(y0 = y0, a = a), N = N, Y = Y): NA/Inf被换成最大的正值
## Warning in nlm(mse, c(y0 = y0, a = a), N = N, Y = Y): NA/Inf被换成最大的正值
## Warning in nlm(mse, c(y0 = y0, a = a), N = N, Y = Y): NA/Inf被换成最大的正值
```

```
## $y0
## [1] 5000
##
## $a
## [1] 0.1475913
##
## $MSE
## [1] 62521484
```

plm 中调用的 nlm 函数使用的是牛顿型算法来寻找最小值,其结果会与初始值的选取有关。 此处当初始值选为 (y0 = 6611, a = 0.15) 时会得到更小的 MSE。

5.

a.

[1] 481.9195

```
mean (gmp$pcgmp)
## [1] 32922.53
sd(gmp$pcgmp) / (length(gmp$pcgmp) ** 0.5)
## [1] 481.9195
(sum((gmp\$pcgmp - mean(gmp\$pcgmp)) ^ 2) / ((length(gmp\$pcgmp) - 1) * length(gmp\$pcgmp))) ^ 0.5
```

b.

```
mean_pcgmp_except <- function(i) {
  return(mean(gmp$pcgmp[-i]))
}</pre>
```

```
mean_pcgmp_except(i = 2)
```

```
## [1] 32922.62
```

C.

```
jackknifed.means <- c()
for (i in 1:length(gmp$pcgmp))
  jackknifed.means <- c(jackknifed.means, mean_pcgmp_except(i))</pre>
```

d.

```
jackknife.variance <- (length(gmp$pcgmp) - 1) ^ 2 / length(gmp$pcgmp) * var(jackknifed.means)
(jackknife.sd <- jackknife.variance ^ 0.5)</pre>
```

```
## [1] 481.9195
```

和 a 中的结果基本相符。

6

```
plm. jackknife <- function(y0, a, N = gmp$pop, Y = gmp$pcgmp) {
    jackknife.estimates <- list(y0 = c(), a = c())
    for (i in 1:length(N)) {
        estimates <- plm(y0, a, N[-i], Y[-i])
        jackknife.estimates$y0 <- c(jackknife.estimates$y0, estimates$y0)
        jackknife.estimates$a <- c(jackknife.estimates$a, estimates$a)
    }
    n <- length(N)
    y0.variance = (n - 1) ^ 2 / n * var(jackknife.estimates$y0)
    a.variance = (n - 1) ^ 2 / n * var(jackknife.estimates$a)
    return(list(y0.sd = y0.variance ^ 0.5, a.sd = a.variance ^ 0.5))
}</pre>
```

```
options(warn = -1)
plm. jackknife(y0 = 6611, a = 1/8)
```

```
## $y0. sd
## [1] 1.136653e-08
##
## $a. sd
## [1] 0.0009901003
```

```
gmp2013 <- read.table('data/gmp-2013.dat', header = T)
gmp2013$pop <- gmp2013$gmp / gmp2013$pcgmp</pre>
```

```
options(warn = -1)

plm(y0 = 6611, a= 1/8, N = gmp2013$pop, Y = gmp2013$pcgmp)
```

```
## $y0
## [1] 6611
##
## $a
## [1] 0.1433688
##
## $MSE
## [1] 135210520
```

```
plm. jackknife(y0 = 6611, a = 1/8, N = gmp2013$pop, Y = gmp2013$pcgmp)
```

```
## $y0. sd

## [1] 2.692675e-08

##

## $a. sd

## [1] 0.001098548
```