

myHomework2

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```
gmp <- read.table("../data/gmp.dat")
gmp$pop <- round(gmp$gmp/gmp$pcgmp)
```

1

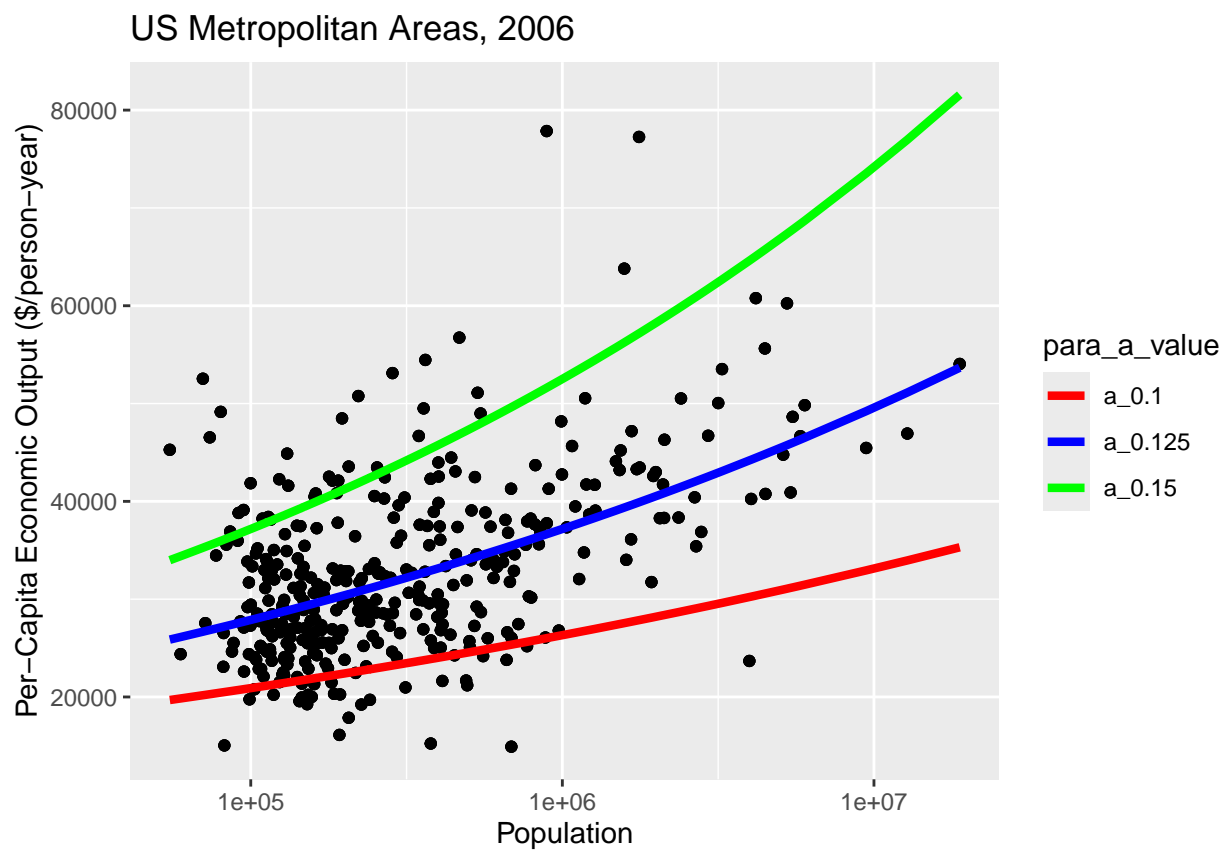
```
gmp <- gmp |>
  mutate(a_0.125 = 6611 * (gmp/pcgmp)^(1/8),
         a_0.1 = 6611 * (gmp/pcgmp)^0.1, a_0.15 = 6611 *
           (gmp/pcgmp)^0.15)
gmp_tidy <- gmp[, 3:7] |>
  gather(para_a_value, nlmfit, -pcgmp,
```

```

    -pop)

gmp_tidy |>
  ggplot() + geom_point(aes(x = pop, y = pcgmp)) +
  labs(x = "Population", y = "Per-Capita Economic Output ($/person-year)",
       title = "US Metropolitan Areas, 2006") +
  geom_line(aes(x = pop, y = nlmfit, color = para_a_value),
           size = 1.5) + scale_x_continuous(trans = "log10") +
  scale_color_manual(values = c("red",
                                "blue", "green"))

```



2.

```

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, 0.1))
```

```
## [1] 298459914
```

3.

```
nlm(mse, c(y0 = 6611, a = 1/8))
```

```
## $minimum
```

```
## [1] 61857060
```

```
##
```

```
## $estimate
```

```
## [1] 6611.0000000 0.1263177
```

```
##
```

```
## $gradient
```

```
## [1] 50.048639 -9.983778
```

```
##
```

```
## $code
```

```
## [1] 2
```

```
##
```

```
## $iterations
```

```
## [1] 3
```

minimum represents the the value of the estimated minimum of f. estimate represents the point at which the minimum value of f is obtained.

4.

```
plm <- function(para, N = gmp$pop, Y = gmp$pcgmp) {  
  t <- nlm(mse, c(para[1], para[2]), N,  
    Y)
```

```

    return(list(parameters = c(t$estimate[1],
                              t$estimate[2]), MSE = t$minimum))
  }
plm(c(6611, 0.15))

```

```

## $parameters
## [1] 6610.9999997    0.1263182
##
## $MSE
## [1] 61857060

```

```
plm(c(5000, 0.1))
```

```

## $parameters
## [1] 5000.0000008    0.1475913
##
## $MSE
## [1] 62521484

```

They differ because their y_0 are different. The starting value, $y_0 = 6611$ and $a = 0.15$, has the lower MSE.

5.

a.

```
mean(gmp$pcgmp)
```

```
## [1] 32922.53
```

```
sem1 <- sd(gmp$pcgmp)/sqrt(nrow(gmp))
sem1
```

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

formula

$$SEM = \sqrt{\frac{\sum_{i=1}^n (x_i - \bar{x})^2}{n(n-1)}}$$

b.

```
mean.jackknife <- function(i, pc = gmp$pcgmp) {  
  return(mean(pc[-i]))  
}
```

c.

```
n = nrow(gmp)  
jackknifed.means <- sapply(1:n, mean.jackknife)
```

d.

```
sem2 <- sqrt((n - 1)^2/n * var(jackknifed.means))  
sem2
```

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

```
abs(sem2 - sem1)/sem1
```

```
## [1] 4.010371e-15
```

非常符合

6.

```
plm.jackknife <- function(para, N = gmp$pop,  
  Y = gmp$pcgmp) {  
  # para = c(6611, 0.125)  
  jackknife <- function(i) {  
    return(plm(para, N[-i], Y[-i])$parameters)  
  }  
  n = length(N)  
  jackknifed.para <- sapply(1:n, jackknife)  
  y.sd <- sqrt((n - 1)^2/n * var(jackknifed.para[1,  
    ]))  
  a.sd <- sqrt((n - 1)^2/n * var(jackknifed.para[2,  
    ]))
```

```

sd <- c(y.sd, a.sd)
names(sd) = c("y.sd", "a.sd")
return(sd)
}

```

```
plm.jackknife(c(6611, 0.125))
```

```
##           y.sd           a.sd
## 1.136653e-08 9.901003e-04
```

7.

```

gmp_2013 <- read.csv("../data/gmp-2013.dat",
  header = T, sep = " ") |>
  mutate(pop = round(gmp/pcgmp))
plm(c(6611, 1/8), N = gmp_2013$pop, Y = gmp_2013$pcgmp)

```

```

## $parameters
## [1] 6611.0000002    0.1433688
##
## $MSE
## [1] 135210524

```

```

plm.jackknife(c(6611, 0.125), N = gmp_2013$pop,
  Y = gmp_2013$pcgmp)

```

```

##           y.sd           a.sd
## 2.692652e-08 1.098548e-03

```

It haven't changed significantly.