Problem 6

```
set.seed(1)
x1 = runif(100)
x2 = (0.5 * x1) + rnorm(100)/10

# Create a linear model in which y is a function of x1 and x2.
y = 2 + (2 * x1) + (0.3 * x2) + rnorm(100)
```

Part A

The form of the linear model is

$$y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \epsilon$$

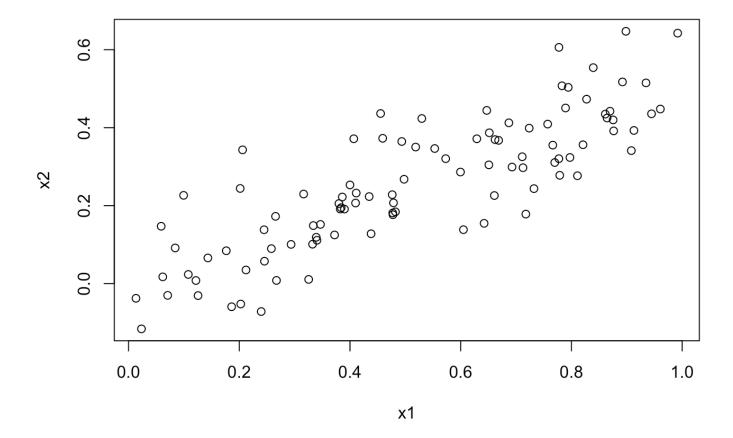
= 2 + 2 \cdot x_1 + 0.3 \cdot x_2 + rnorm(100)
$$\beta_0 = 2, \beta_1 = 2, \beta_2 = 0.3, \epsilon = rnorm(100)$$

Part B

```
cor(x1, x2)
```

```
## [1] 0.8351212
```

```
plot(x1, x2)
```



Part C

```
fit <-lm(y \sim x1 + x2)
summary(fit)
```

```
##
## Call:
## lm(formula = y \sim x1 + x2)
##
## Residuals:
      Min
          1Q Median
                              3Q
                                    Max
## -2.8311 -0.7273 -0.0537 0.6338 2.3359
##
## Coefficients:
##
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.1305 0.2319 9.188 7.61e-15 ***
## x1
               1.4396
                          0.7212 1.996
                                          0.0487 *
               1.0097
                                        0.3754
## x2
                          1.1337 0.891
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.056 on 97 degrees of freedom
## Multiple R-squared: 0.2088, Adjusted R-squared: 0.1925
## F-statistic: 12.8 on 2 and 97 DF, p-value: 1.164e-05
```

The estimated coefficients are $\beta_0=2.1305$, $\beta_1=1.4396$, and $\beta_2=1.0097$, which are at least in the ballpark of the true coefficients (2, 2, 0.3). β_2 is smaller than both β_0 and β_1 in both the true and estimated coefficients.

We can reject the null hypothesis H_0 : $\beta_1=0$, because the p = 0.0487 < 0.05. However, we cannot reject H_0 : $\beta_2=0$, because p = 0.3754 > 0.05.

Part D

```
fit <- lm(y ~ x1)
summary(fit)</pre>
```

```
##
## Call:
## lm(formula = y \sim x1)
##
## Residuals:
       Min
                 1Q Median
                                  3Q
                                          Max
## -2.89495 -0.66874 -0.07785 0.59221 2.45560
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 2.1124 0.2307 9.155 8.27e-15 ***
## x1
                1.9759
                           0.3963 4.986 2.66e-06 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 1.055 on 98 degrees of freedom
## Multiple R-squared: 0.2024, Adjusted R-squared: 0.1942
## F-statistic: 24.86 on 1 and 98 DF, p-value: 2.661e-06
```

We can reject the null hypothesis H_0 : $\beta_1=0$, because the p=2.661e-06<0.05. When we throw out x_2 , we get a much more impressive p value than when we included both x_1 and x_2 in the linear regression.

Part E

```
fit <- lm(y ~ x2)
summary(fit)</pre>
```

```
##
## Call:
## lm(formula = y \sim x2)
##
## Residuals:
##
       Min
                 10 Median
                                  3Q
                                          Max
## -2.62687 -0.75156 -0.03598 0.72383 2.44890
##
## Coefficients:
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.3899 0.1949 12.26 < 2e-16 ***
## x2
                2.8996
                         0.6330 4.58 1.37e-05 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.072 on 98 degrees of freedom
## Multiple R-squared: 0.1763, Adjusted R-squared: 0.1679
## F-statistic: 20.98 on 1 and 98 DF, p-value: 1.366e-05
```

We can reject the null hypothesis H_0 : $\beta_2 = 0$, because the p = 1.366e-06 < 0.05. When we throw out x_1 , we get a much more impressive p value than when we included both x_1 and x_2 in the linear regression.

Part F

In a way, yes, I expected Part E to show a non-significant p -value, but instead it was even lower than in Part D!

Part G

```
x1 = c(x1, 0.1)

x2 = c(x2, 0.8)

y = c(y, 6)
```

```
fit <-lm(y \sim x1 + x2)
summary(fit)
```

```
##
## Call:
## lm(formula = y \sim x1 + x2)
##
## Residuals:
##
       Min
                 1Q Median
                                  3Q
                                          Max
## -2.73348 -0.69318 -0.05263 0.66385 2.30619
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
##
## (Intercept)
                2.2267 0.2314 9.624 7.91e-16 ***
## x1
                0.5394
                          0.5922 0.911 0.36458
               2.5146 0.8977 2.801 0.00614 **
## x2
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.075 on 98 degrees of freedom
## Multiple R-squared: 0.2188, Adjusted R-squared: 0.2029
## F-statistic: 13.72 on 2 and 98 DF, p-value: 5.564e-06
```

```
fit <- lm(y ~ x1)
summary(fit)</pre>
```

```
##
## Call:
## lm(formula = y \sim x1)
##
## Residuals:
      Min
          1Q Median
                              3Q
                                     Max
## -2.8897 -0.6556 -0.0909 0.5682 3.5665
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
                         0.2390 9.445 1.78e-15 ***
## (Intercept) 2.2569
## x1
                1.7657
                           0.4124 4.282 4.29e-05 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.111 on 99 degrees of freedom
## Multiple R-squared: 0.1562, Adjusted R-squared:
## F-statistic: 18.33 on 1 and 99 DF, p-value: 4.295e-05
```

```
fit <- lm(y ~ x2)
summary(fit)</pre>
```

```
##
## Call:
## lm(formula = y \sim x2)
##
## Residuals:
##
       Min
                 10 Median
                                   30
                                          Max
## -2.64729 -0.71021 -0.06899 0.72699 2.38074
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
                         0.1912 12.264 < 2e-16 ***
## (Intercept) 2.3451
                           0.6040 5.164 1.25e-06 ***
## x2
                3.1190
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 1.074 on 99 degrees of freedom
## Multiple R-squared: 0.2122, Adjusted R-squared: 0.2042
## F-statistic: 26.66 on 1 and 99 DF, p-value: 1.253e-06
```

This new observation flips the relationship we saw before. Previously, we saw a more significant p value for x_1 when we fit the lm to both x_1 and x_2 , while now we have a more significant p value for x_2 . However, they still both indicate a low p value in the lms where we fit the two variables independently.