

MTH 4230 R Notes 3

1 Regression Through the Origin

- To fit a regression model *through the origin* (i.e. the model with **no intercept**),

$$Y_i = \beta_1 X_i + \epsilon_i$$

we use the *formula* `y ~ -1 + x` in the call to the `lm()` function.

- For example, consider the data in these vectors `x` and `y`:

```
x <- c(22, 15, 7, 19, 20, 9, 15, 10, 19, 21)
y <- c(12.2, 9.5, 6.7, 5.9, 10.0, 8.9, 11.5, 10.0, 9.9, 10.1)
```

We fit the model with **no intercept** by typing:

```
my.reg <- lm(y ~ -1 + x)
```

Then we look at the results in the usual manner, using `summary()`:

```
summary(my.reg)

##
## Call:
## lm(formula = y ~ -1 + x)
##
## Residuals:
##   Min     1Q   Median     3Q    Max
## -4.709 -1.053  0.520  3.041  4.416
##
## Coefficients:
##   Estimate Std. Error t value Pr(>|t|)
## x    0.5584     0.0571   9.779 4.31e-06
##
## Residual standard error: 2.982 on 9 degrees of freedom
## Multiple R-squared:  0.914, Adjusted R-squared:  0.9044
## F-statistic: 95.62 on 1 and 9 DF,  p-value: 4.31e-06
```

Notice that there's **no intercept term** in the output, just the slope term. A plot of the data with the fitted regression line is shown below. Notice that **the line passes through the origin**.

```
plot(x, y, pch = 19, xlim = c(0,25), ylim = c(0,15),
     main = "Regression Through the Origin")

abline(my.reg, lwd = 2, col = "blue")
```



2 Matrix Approach to Regression

2.1 Obtaining the Design Matrix

- To obtain the $n \times p$ *design matrix* X used in a regression analysis, we use the function:

```
model.matrix() # Returns the design matrix X used in a linear
               # regression analysis carried out by lm()
```

The `model.matrix()` function takes as its main argument either an *lm* object or a *formula* indicating a regression model. It returns the $n \times p$ *design matrix* X .

- For example:

```
x <- c(22, 15, 7, 19, 20, 9, 15, 10, 19, 21)
y <- c(12.2, 9.5, 6.7, 5.9, 10.0, 8.9, 11.5, 10.0, 9.9, 10.1)
```

```
my.reg <- lm(y ~ x)
```

```
X <- model.matrix(my.reg)
```

```
X
##      (Intercept)  x
## 1             1 22
## 2             1 15
## 3             1  7
## 4             1 19
## 5             1 20
## 6             1  9
## 7             1 15
## 8             1 10
## 9             1 19
## 10            1 21
## attr(,"assign")
## [1] 0 1
```

The matrix returned by `model.matrix()` comes equipped with several *attributes*. In R, an *attribute* is bit of extra information (so-called *meta data*) that some *classes* of data objects, including *matrices*, contain.

The "assign" attribute is a *vector* with a 0 representing the intercept and 1 the predictor. For models with more predictors, the "assign" *vector* would have elements 2, 3, ..., p for the additional predictors. To see all of X's *attributes*, type `attributes(X)`.

We can verify that X is *matrix* and look at its dimensions using:

```
is.matrix(X)
## [1] TRUE
dim(X)
## [1] 10 2
```

We see that X indeed a *matrix* and it has $n = 10$ rows and $p = 2$ columns.

- Another way to obtain the *model matrix* is by passing a *formula* to `model.matrix()`:

```
X <- model.matrix(y ~ x)
```

2.2 Performing Computations with the Design Matrix

- All of the usual matrix functions and operators (e.g. `t()`, `%*%`, `solve()`, `det()`, etc.) can be used with the design matrix \mathbf{X} .
- For example, we can use matrix operations to obtain the *vector of estimated coefficients*

$$\mathbf{b} = \begin{bmatrix} b_0 \\ b_1 \end{bmatrix}$$

given by

$$\mathbf{b} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{Y}$$

from the *matrix* \mathbf{X} and *vector* \mathbf{y} created earlier by typing:

```
b <- solve(t(X) %*% X) %*% t(X) %*% y
b

##                [,1]
## (Intercept) 7.3189622
## x           0.1370088
```

We see that the *estimated intercept* is $b_0 = 7.319$ and the *estimated slope* is $b_1 = 0.137$.

2.3 The (Estimated) Variance-Covariance Matrix of b_0 and b_1

- The (estimated) *variance-covariance matrix* $s^2\{\mathbf{b}\}$ of the *estimated coefficient vector*

$$\mathbf{b} = \begin{bmatrix} b_0 \\ b_1 \end{bmatrix}$$

is obtained using the function:

```
vcov()      # Returns the (estimated) variance-covariance matrix of the
            # regression coefficient estimates in a linear regression
            # analysis carried out by lm()
```

The `vcov()` function takes an *lm* object (such as `my.reg`) as its main argument and returns a *matrix* containing the *variances* on the diagonal and *covariances* on the off-diagonals.

- For example, using `my.reg` created earlier:

```
vcov(my.reg)

##                (Intercept)          x
## (Intercept)  3.7119109 -0.2137037
## x           -0.2137037  0.0136117
```

```
vcov.tmp <- vcov(my.reg)
```

We see that:

- The (estimated) *variance* of b_0 is $s^2\{b_0\} = 3.712$, so the (estimated) *standard error* of b_0 is $s\{b_0\} = \sqrt{3.712} = 1.927$.
 - The (estimated) *variance* of b_1 is $s^2\{b_1\} = 0.014$, so the (estimated) *standard error* of b_1 is $s\{b_1\} = \sqrt{0.014} = 0.117$.
 - The (estimated) *covariance* between b_0 and b_1 is $s^2\{b_0, b_1\} = -0.214$.
- Note that the **variance-covariance matrix** of \mathbf{b} is $\text{MSE} \cdot (\mathbf{X}^T \mathbf{X})^{-1}$, so we could also have obtained it using the MSE from `lm()` and the design matrix \mathbf{X} .