This week we'll start out by reviewing some key concepts that we sped over last week. Next, we'll briefly learn how to use loops in R. Finally, we'll work through an example that uses many of the tools we've developed in section to demonstrate why you should be suspicious of any " R^2 maximizers"!

Last section

Using apply, round two: Simply put, apply() is a function that lets you perform operations on different parts of a matrix. I'll show you a few different ways to use apply(), borrowing heavily from Neil Saunders' excellent blog post on the topic¹.

First we use our matrix() command from last week to construct a 2×10 matrix:

Now we'll use the apply function to get the mean of each row. Note that we could also do this manually, using mean(m[1,]) and mean(m[2,]). But that's boring.

```
apply(m, MARGIN = 1, FUN = mean)
```

I've included the parameter names just to be clear about what's happening here. Per the apply() documentation, setting MARGIN = 1 tells R to apply the given function over rows. Setting FUN = mean indicates that the given function is mean. If we actually want the column means, then we just have to set MARGIN = 2:

```
apply(m, 2, mean)
[1] 6 7 8 9 10 11 12 13 14 15
```

Note that including the parameter names is not strictly necessary, so I omitted it above. If you don't pass the parameter names, however, be sure to pass your arguments in the correct order!

Finally, we can apply() a function to every cell in the matrix individually using MARGIN = 1:2:

¹Available at http://nsaunders.wordpress.com/2010/08/20/a-brief-introduction-to-apply-in-r/. He goes into more detail on the entire family of apply() functions, if you're interested.

OLS with matrices: As we discussed last week, the use of canned routines is not permitted for most of this class; you'll have to write the econometric routines from first principles. First, create matrices of the data, since we will be working mainly with matrix operations. Let **y** be the dependent variable, price, and let **X** be a matrix of the other car characteristics, along with a column of ones prepended. The cbind() function binds the columns horizontally and coerces the matrix class.

```
y <- matrix(data$price)</pre>
X <- cbind(1, data$mpg, data$weight)</pre>
head(X)
      [,1] [,2] [,3]
             22 2930
[1,]
[2,]
             17 3350
[3,]
             22 2640
[4,]
             20 3250
[5,]
         1
             15 4080
[6,]
             18 3670
```

Last week I demo'ed how to use the rep() command to create an $n \times 1$ vector of ones. rep(), short for replicate, is an incredibly useful command. However, in this setting cbind() only needs to be passed a single 1 — it's smart enough to do the replication itself in order to ensure that the matrix is filled.

Just to make sure that our matrices will be conformable when we regress y on X, check that the number of observations are the same in both variables.

```
dim(X)[1] == nrow(y)
[1] TRUE
```

Using the matrix operations described in the previous section, we can quickly estimate the ordinary least squared parameter vector.

That's it! And although you're not allowed to use it in your problem sets, lm() is a nice tool for checking our results.

They match! Thank goodness. If you're interested in knowing what lm() does, I highly recommend reading through relevant R documentation².

²Remember, you can do this using ?1m.

Loops in R

Loops are a familiar concept in programming. In general, a loop is a programming statement that allows repeated execution of some piece of code. Let's take a look at one of the most common loop types, a for loop:

```
for (i in 1:5) {
   print(i*i)
}

[1] 1
[1] 4
[1] 9
[1] 16
[1] 25
```

The structure of the for loop always the same. First, we specify the counter variable, i, with for (i in 1:5). This tells R to run the loop five times, first with i=1, second with i=2, and so on. Then, we specify the commands we want to run repeatedly. In this case, we want R to print i^2 for each value of i. And that's it!

I should mention that using loops is somewhat discouraged in R. R is a "vector-based" language, so in theory loops are much less efficient than vectorized functions like %*% or apply(). Of course, the datasets we will work with in this class are small, so any practical difference will be neglible until you get involved with real data. But if you're interested, you can try your hand at writing the equivalent vectorized function using the apply() family.

Centered R^2

We've now laid enough groundwork to start getting into some detailed examples of employing R to learn and do applied econometrics. In other words, I can finally stop torturing you with pointless matrix algebra and start torturing you with matrix algebra with a purpose! The first such example has to do with R^2 values. You will all be calculating this R^2 for your problem sets, but here we'll take it a step farther and look at what happens to R^2 when we add random variables to our regression. Along the way, we'll use many of the tools we've developed in this and the first two sections.

First, we create a random matrix, where each element is drawn from a standard uniform distribution — another context to practice the function() structure. The function randomMat() generates a long vector of length $n \cdot k$ and then reshapes it into an $n \times k$ matrix.

```
randomMat <- function(n, k) {
  v <- runif(n*k)
  matrix(v, nrow=n, ncol=k)
}</pre>
```

You might notice that I didn't include a return statement in this function. That's okay! R automatically returns the output from the last command entered by default. So, the function randomMat() behaves as we would expect:

Another useful function for this section will be to create a square demeaning matrix \mathbf{A} of dimension n. The following function just wraps a few algebraic maneuvers, so that subsequent code is easier to read.

```
demeanMat <- function(n) {
  ones <- rep(1, n)
  diag(n) - (1/n) * ones %*% t(ones)
}</pre>
```

As is described in the notes, pre-multiplying a matrix \mathbf{B} by \mathbf{A} will result in a matrix $= \mathbf{A}\mathbf{B}$ of deviations from the column means of \mathbf{B} . Check that this is true. This may seem like a roundabout way to check the equivalence of the matrices; but it provides the opportunity to practice the apply function.

```
A <- demeanMat(3)
B <- matrix(1:9, nrow=3)
col.means <- apply(B, 2, mean)
C <- apply(B, 1, function(x) {x - col.means})
all.equal(A %*% B, t(C))
[1] TRUE</pre>
```

Alright, we're ready to apply the functions to real data in order to calculate the centered \mathbb{R}^2 . First, read in the data to conform to equation (2.37) on page 14 of the lecture notes, and identify the number of observations n for later use:

```
data <- read.csv("auto.csv", header=TRUE)
names(data) <- c("price", "mpg", "weight")
y <- matrix(data$price)
X2 <- cbind(data$mpg, data$weight)
n <- nrow(X2)</pre>
```

The centered \mathbb{R}^2 is defined according to equation (2.41) as follows:

$$\mathbf{R}^2 = \frac{\mathbf{b}_2' \mathbf{X}_2^{*\prime} \mathbf{X}_2^* \mathbf{b}_2}{\mathbf{y}^{*\prime} \mathbf{y}^*},\tag{1}$$

where $\mathbf{y}^* = \mathbf{A}\mathbf{y}$, $\mathbf{X}_2^* = \mathbf{A}\mathbf{X}_2$, and $\mathbf{b}_2 = (\mathbf{X}_2^{*\prime}\mathbf{X}_2^*)^{-1}\mathbf{X}_2^{*\prime}\mathbf{y}^*$. Noting that \mathbf{A} is both symmetric and idempotent, we can rewrite Equation (1) using \mathbf{A} , \mathbf{X} , and \mathbf{y} as follows. We use that $\mathbf{X}_2^{*\prime}\mathbf{X}_2^* = \mathbf{X}_2^{\prime}\mathbf{A}\mathbf{A}\mathbf{X}_2 = \mathbf{X}_2^{\prime}\mathbf{A}\mathbf{A}\mathbf{X}_2 = \mathbf{X}_2^{\prime}\mathbf{A}\mathbf{X}_2$ and similarly that $\mathbf{y}^{*\prime}\mathbf{y}^* = \mathbf{y}^{\prime}\mathbf{A}\mathbf{y}$ and $\mathbf{X}_2^{*\prime}\mathbf{y}^* = \mathbf{X}_2^{\prime}\mathbf{A}\mathbf{y}$. We'll use this in our code below to construct \mathbf{R}^2 , constructing a general function that we can use with any arbitrary dependent vector and associated cofactor matrix:

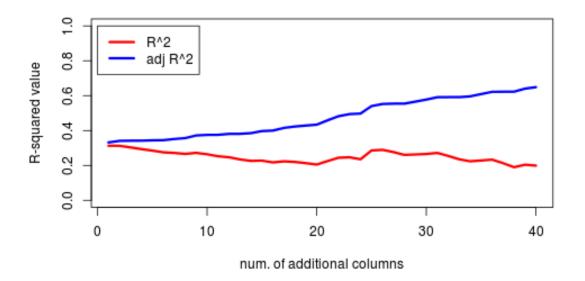
```
R.squared <- function(y, X) {
    n <- nrow(X)
    k <- ncol(X)
    A <- demeanMat(n)
    xtax <- t(X) %*% A %*% X
    ytay <- t(y) %*% A %*% y
    b2 <- solve(xtax) %*% t(X) %*% A %*% y
    R2 <- t(b2) %*% xtax %*% b2 / ytay
    R2.adj <- 1 - ((n-1)/(n-k))*(1-R2)
    return(cbind(R2,R2.adj))
}
R.squared(y, X2)

[,1] [,2]
[1,] 0.2933891 0.2835751</pre>
```

Note that this code also returns $\overline{\mathbf{R}}^2$, the adjusted \mathbf{R}^2 . Now, what happens if we use our randomMat() function to add columns of random covariates to our regression? We'll use a for loop to calculate the \mathbf{R}^2 and $\overline{\mathbf{R}}^2$ as we add additional covariates.

```
n <- nrow(X2); k.max <- 40
X.rnd <- randomMat(n, k.max)
res.R2 <- rep(0, k.max)
res.adjR2 <- rep(0, k.max)

for (i in 1:k.max) {
    X.ext <- cbind(X2, X.rnd[, seq(i)])</pre>
```



As you can see, our \mathbf{R}^2 has gone from a puny 0.3 to a robust 0.7, so our model must be much better, right? Wrong! This is another example of the classic "overfitting" problem: in a regression with 74 observations and 40+ covariates, *something* is always going to explain some portion of the variation. The $\overline{\mathbf{R}}^2$, on the other hand, stays mostly at the same level. But what if we set k.max <-70? k.max <- 100? Why?

That's it for this section. Next week, we'll turn our attention to hypothesis testing.

Additional puzzles

- 1. Write a function wt.coef() that will return the OLS coefficient on weight from the regression of car price on the covariate matrix described above.
- 2. Adjust the function to return a list of coefficients from the same linear regression, appropriately named.
- 3. Find the estimate of the covariance matrix $\sigma^2(\mathbf{X}'\mathbf{X})^{-1}$ and show that the residuals and covariate matrix are orthogonal.