Module 12

R Programming

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R 'programming'

Now we are going to switch gears a little bit, and talk about some of the more traditional programming that you can do in R.

You can do very flexible things, but at a cost of more difficult notation, and having to actually write programming statements. There are slight notation differences as well, including the use of curly { } brackets

We are going to cover for loops and if statements

These allow you to iterate over certain observations or subsets of observations

The syntax is:

```
for(*var* in seq) {
do something
}
```

Typically they look something like:

```
for(i in 1:nrow(dat)) {
   something(dat[i,])
}
```

These are essentially fancier apply statements

For example,

```
> for(i in 1:10) {
+  print(i)
+ }
```

```
[1] 1

[1] 2

[1] 3

[1] 4

[1] 5

[1] 6

[1] 7

[1] 8

[1] 9

[1] 10
```

Here's how they can be more flexible:

```
> Index = c(3,6,7,20,32,100,234,1000,6543)
> for(i in 1:length(Index)) {
+  print(Index[i])
+ }
```

```
[1] 3

[1] 6

[1] 7

[1] 20

[1] 32

[1] 100

[1] 234

[1] 1000

[1] 6543
```

Note that the first time through the body of the loop, i takes the value 1, then evaluates the body. Then, i takes the value 2, and evaluates the body, until i = length(Index), then it stops.

They are essentially more useful than apply statements when you are working with two sets of matching datasets or vectors.

```
> myList = vector("list",length=4)
> mat1=matrix(rnorm(8), nc = 4)
> mat2=matrix(rnorm(8), nc = 4)
> mat1
```

```
[,1] [,2] [,3] [,4]
[1,] -0.2854 -0.01147 -0.3855 -0.3403
[2,] -0.9100 0.52103 1.1067 -1.7502
```

```
> mat2
```

```
[,1] [,2] [,3] [,4]
[1,] 1.429 0.02987 -0.6456 0.5449
[2,] 1.775 0.30341 -0.3780 -0.6708
```

```
> for(i in seq(along=myList)) {
+  myList[[i]] = cbind(mat1[,i],mat2[,i])
+ }
> myList
```

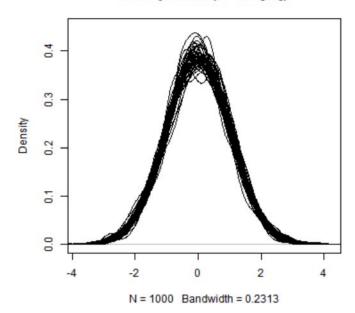
```
[[1]]
        [,1] [,2]
[1,] -0.2854 1.429
[2,] -0.9100 1.775
[[2]]
        [,1] [,2]
[1,] -0.01147 0.02987
[2,] 0.52103 0.30341
[[3]]
        [,1]
               [,2]
[1,] -0.3855 -0.6456
[2,] 1.1067 -0.3780
[[4]]
        [,1] [,2]
[1,] -0.3403 0.5449
[2,] -1.7502 -0.6708
```

```
> i=1
> cbind(mat1[,i],mat2[,i])
        [,1] [,2]
[1,] -0.2854 1.429
[2,] -0.9100 1.775
> i=2
> cbind(mat1[,i],mat2[,i])
         [,1]
               [,2]
[1,] -0.01147 0.02987
[2,] 0.52103 0.30341
> i=3
> cbind(mat1[,i],mat2[,i])
        [,1]
               [,2]
[1,] -0.3855 -0.6456
[2,] 1.1067 -0.3780
```

These are useful for making many columns worth of density plots

```
> mat = matrix(rnorm(1000*50), nc = 50)
> plot(density(mat[,1]), ylim = c(0,0.45))
> for(i in 2:ncol(mat)) {lines(density(mat[,i]))}
```

density.default(x = mat[, 1])



You can also integrate with lists.

```
> outList = vector("list",10)
> start=1:10
> end = sample(1:100, 10)
> for(i in seq(along=outList)) {
+ outList[[i]] = start[i]:end[i]
+ }
> outList
```

```
[[1]]
[1] 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23
[24] 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46
[47] 47 48 49 50 51 52 53 54 55 56 57 58 59
[[2]]
[1] 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24
[24] 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47
[47] 48 49 50 51 52 53 54
[[3]]
[1] 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25
[24] 26 27 28 29 30 31 32 33 34 35 36
[[4]]
[1] 4 5 6 7 8 9 10 11 12 13 14 15 16 17
[[5]]
                                                                                 10/19
[1] 5 6 7 8 9 10
```

'if' statements

You can put 'if' statements inside of 'for' loops

```
for(i in 1:nrow(dat)) {
   if(dat$x > num) {
      dat$y[i] = something
   } else {
      dat$y[i] = something else
   }
}
```

Example

```
> makeIndexes=split(1:nrow(cars), cars$Make)
> lapply(makeIndexes, head, n=4)[1:3]

$ACURA
[1] 10039 13026 13631 14250

$BUICK
[1] 185 233 258 346

$CADILLAC
[1] 3372 4517 8500 9664
```

```
> pval = rep(NA,length(makeIndexes))
> for(i in 1:length(makeIndexes)) {
+    ind = makeIndexes[[i]]
+    if(length(ind)>1) {
+        f = lm(VehBCost~VehOdo, data=cars,subset=ind)
+        pval[i] = summary(f)$coef[2,4]
+    }
+ }
> names(pval)=names(makeIndexes)
> i = 1
> ind = makeIndexes[[i]]
> str(ind)
```

```
int [1:33] 10039 13026 13631 14250 16392 17289 17889 17979 18166 22044 ...
```

```
> f = lm(VehBCost~VehOdo, data=cars, subset=ind)
> summary(f)$coef[2,4]
```

```
[1] 0.4932
```

> pval

CHRYSLER	CHEVROLET	CADILLAC	BUICK	ACURA
1.128e-78	2.834e-06	1.064e-06	1.877e-05	4.932e-01
HUMMER	HONDA	GMC	FORD	DODGE
NA	2.490e-13	1.626e-01	2.58 4e- 27	1.494e-10
KIA	JEEP	ISUZU	INFINITI	HYUNDAI
3.765e-18	2.723e-16	2.697e-04	9.737e-04	1.072e-04
MINI	MERCURY	MAZDA	LINCOLN	LEXUS
8.709e-02	2.953e-04	1.364e-41	7.061e-01	5.014e-09
PONTIAC	PLYMOUTH	OLDSMOBILE	NISSAN	MITSUBISHI
3.305e-154	NaN	6.956e-08	3.594e-10	3.357e-26
TOYOTA	SUZUKI	SUBARU	SCION	SATURN
1.395e-16	4.285e-32	8.062e-01	3.115e-07	4.363e-40
		VOLVO	VOLKSWAGEN	TOYOTA SCION
		1.269e-02	4.738e-06	NA

Note you can also do with with sapply

```
> pval2 = sapply(makeIndexes, function(ind) {
+    if(length(ind)>1) {
+        f = lm(VehBCost~VehOdo, data=cars,subset=ind)
+        summary(f)$coef[2,4]
+    } else NA
+ })
> all.equal(pval,pval2)
```

```
[1] TRUE
```

Example

Now we can read in many files into a list

```
> fn = list.files("Reports/",pattern=".txt",full.names=TRUE)
> name = list.files("Reports/",pattern=".txt",full.names=FALSE)
> head(fn)
```

```
[1] "Reports/April_2009_Report.txt" "Reports/April_2010_Report.txt"
[3] "Reports/April_2011_Report.txt" "Reports/August_2009_Report.txt"
[5] "Reports/August_2010_Report.txt" "Reports/August_2011_Report.txt"
```

```
> fileList = lapply(fn, read.delim, header=TRUE, as.is=TRUE)
> names(fileList) = name
> sapply(fileList,dim)[,1:5]
    April 2009 Report.txt April 2010 Report.txt April 2011 Report.txt
                                                               359
[1,]
                     287
                                          324
                      10
[2,]
                                                               10
                                           10
    August 2009 Report.txt August 2010 Report.txt
[1,]
                      353
                                            369
[2,]
                       10
                                             10
> lapply(fileList[1:5],head,n=2)
$April 2009 Report.txt
                      age bgDrugs height weight block recruitDate
   id
              treat
         sex
1 1072 Female Control 51.00 asprin 63.84 131.3
                                                  d
                                                             21 22.64
2 1073 Female Control 54.81 tylenol 66.10 117.2
                                                  b
                                                             1 18.85
$April 2010 Report.txt
         sex treat age bgDrugs height weight block recruitDate
   id
1 4337 Female Case 46.91 none 64.95 140.6
                                                f
                                                          25 23.43
2 4338 Female Case 47.95 none 66.47 143.3 f
                                                          14 22.81
$April 2011 Report.txt
   id sex treat
                    age bgDrugs height weight block recruitDate
1 7780 Male
             Case 53.93 asprin 70.12 175.0
                                                           29 25.02
2 7781 Male Control 62.77 tylenol 71.02 153.1
                                                           29 21.34
                                                b
$August 2009 Report.txt
   id sex treat
                    age bgDrugs height weight block recruitDate
                                                               bmi
1 2051 Male Control 56.76 tylenol 70.47 168.0
                                                            2 23.78
2 2052 Male Case 50.14 asprin 69.56 172.3 c
                                                            1 25.04
                                                                               17/19
```