Talk 2 - Control Structures in R

Kevin O'Brien, M.S. Vision Sciences Laboratory Fall 2013 University of Georgia Psychology Department

General Notes/Tips

- This presentation is in black and white to make it more easily printable.
- http://www.statmethods.net is a great resource
- This talk assumes you're using RStudio as a GUI for R.
- Feed R a copy of your data, not the original
- Always think before you code, especially in loops, and start small
- "How can I find where my code is failing" is the first step in finding out why your code failed.
- When you're done with your code, clean everything up and run it again. If it works then, you're set and can repeat your analysis.
- While many computer languages have 0 referencing, R has 1 referencing. If this conflicts with your habits, you're probably at a skill level where you'll be able to readily spot those issues.
- Typing ?[function] (e.g. ?mean()) pulls up the help page nicer in RStudio

A Quick Word About Formatting

- #Comments are indicated by an octothorp
- Code in this talk is indicated in blue (should print in gray) and italics
 - print("This demonstrates the print command");
- In RStudio's output console, code is **blue**, output is black, and errors are **red**.
- It may seem silly to put semicolons at the end of individual lines, but make it a habit. Things fail catastrophically in loops without semicolons.
- The quotation marks that you type into one program may not paste correctly in R: "" vs ""

Object Types Are Important

- Vectors are one-dimensional arrays
 - numerals<-c(1,2,3);</pre>
 - names<-c("one","two","three");</pre>
 - capitalNames<-c('ONE','TWO','THREE');</pre>
- Data frames are a collection of vectors:
 - numbers<-data.frame(numerals, names, capitalNames);
 - names(numbers)<-c('Numerals','LowerCase','UpperC ase');

Basic Control Structures

- Boolean comparisons
- If() statement (and if()/else() statement)
- For() loop
- While() loop
- Switch Case
- Sapply / Lapply (R specific)
- Sequences and repeat

Boolean Comparisons

- A statement for a boolean comparison has to have a binary state (TRUE or FALSE)
- == (check if equal)
- <, >, <=, >= (less than, greater than, less than OR equal to, greater than OR equal to)
- |, &, ! (or, and, not)
- isTRUE(1!=1); (checks if contents are true)

Boolean Comparator Examples

- **1==1;** #True
- 1==0; #False
- 1!=0; #True
- 'a'=='a'; #True
- 1<0; #False
- 1>0; #True
- 1<=1; #True
- 1>=1; #True

- (TRUE & TRUE);
- (TRUE & FALSE);
- (TRUE | FALSE);
- (TRUE | TRUE);
- (FALSE | FALSE);
- (!FALSE);
- (!TRUE);

The Humble If() Statement

- "Do this thing in braces if whatever I have in parentheses is true"
- *x*<-10;
- *if(x==10)*{
- print("x is indeed ten");
- }
- If you change the test condition OR the value you're testing so that it evaluates to false, it won't spit out the output from the true condition.
- The if() statement is usually paired with an else().

At long last else

- "if the condition for the if part isn't satisfied, do this thing instead"
- a<-5;
- *if(a==5)*{
- print("Yep. It's a five.")
- }else{
- print("Your contrived demonstration did not satisfy the if() condition");
- }

Well that's pretty boring...

- ...yeah, but it illustrates the mechanics. Let's make a function and cover a new arithmetic operator, because that sounds way less boring (by comparison)!
- %% (modulo / modulus) does integer division and returns the remainder – 10%%3 should return 1, for example
- The code's a bit bulky so it's on the next slide

That next slide I mentioned

- EvenOrOdd<-function(inputValue){
- if(inputValue%%2 == 0){
- print(paste(inputValue,"is even.",sep=""));
- }else{
- print(paste(inputValue,"is odd.", sep=""));
- }
- }
- EvenOrOdd(5);
- EvenOrOdd(6);

Other Nifty Tricks With If/Else

- *p*<-0;
- p<-ifelse(p=="not_potato","potato","not_potato");</pre>
- *z<-0;*
- *if(z==0)*{
- print("Z's zero, so do this thing.");
- }else if(z > 0){
- print("Z's bigger than zero, so do this thing.");
- }else if(z < 0){
- print("Z's smaller than zero, so do this thing.");
- }else{
- print("Wait, if it's not zero, and not larger or smaller...");
- print("WHY DID WE EVEN WRITE THIS PART?");
- }
- It's generally good coding practice to ALWAYS have an ELSE, even if it's just empty or returns an error.

WARNING: LOOPS AHEAD

- Loops make it very easy to do repetitive things a tremendous number of times. DO NOT FORGET THAT THEY ARE POWERFUL.
- You can crash R or crash your computer with an infinite loop or a finite loop that uses too much memory.
- Be absolutely certain you know what you're doing if you do file I/O in a loop you could destroy important stuff outside of R. Seriously.
- Consider yourself warned.

They're actually not that scary

- In most circumstances, you just need to make sure your code runs properly before you put it into a loop. Test the loop with a small amount of data before you let it run on a large amount of data.
- Efficiency increases inside loops are multiplicative, so be mindful of bloated code.
- Be prepared for frustrating errors that will make you feel great to fix.

The For() Loop

- This is the easiest loop to visualize, the hardest loop to break things with, and will cover like 99% of your loop needs.
- For loops require a counter variable and a sequence in R. The next few slides will have several trivial examples before we get into real, useful examples.

For() Loop Baby Steps

- *print(1);*
- print(2);
- print(3);
- print(4);
- *print(5);*
- For something simple like this, a for loop doesn't save us much time, but for something larger, it saves so much time.

- for(i in seq(1:5)){
- print(i);
- }
- for(i in seq(1:100)){
- print(i);
- }
- for(i in seq(from=0, to=1000, by=100)){
- print(i);
- }

Pffft...that still doesn't seem helpful.

- Oh yeah?
- subNum<-seq(1:1000);
- subNum[473]<-4730;
- for(i in 1:length(subNum)){
- if(subNum[i]>1000 | subNum[i]<0){
- subNum[i]=NA;
- print(paste(i, "had an error!", sep=" "));
- }
- }
- Boom! You just changed a value that's impossible to NA so it's flagged properly for your analysis AND had R spit out a message to let you know what value(s) had a problem.

Well, I guess that could be helpful...

•Make this big fake dataset:

- set<-data.frame();
- currentRow = 1;
- for(i in 1:10){
- for(j in 1:10){
- for(k in 1:10){
- set[currentRow,1]<-currentRow;
- set[currentRow,2]<-i;
- set[currentRow,3]<-j;
- set[currentRow,4]<-k;
- set[currentRow,5]<-rnorm(1,mean=100,sd=15);
- set[currentRow,6]<-rnorm(1,mean=100,sd=15);
- set[currentRow,7]<-rnorm(1,mean=100,sd=15);
- set[currentRow,8]<-rnorm(1,mean=100,sd=15);
- set[currentRow,9]<-rnorm(1,mean=100,sd=15);
- currentRow=currentRow+1;
- }
- }
- }

names(set)<-c("SubjectNo", "Cond1", "Cond2", "Cond3", "IQ1", "IQ2", "IQ3", "IQ4", "IQ5");

...why are we doing this?

- for(i in 1:nrow(set)){
- set[i,10]<-sum(set[i,5:9])/5;
- if(set[i,10]>105){
- set[i,11]="HIGH";
- }else if(set[i,10]<95){
- set[i,11]="LOW";
- }else{
- set[i,11]="AVG";
- }
- }
- names(set)[10:11]<-c("Mean","Group");
- We can aggregate, encode, replace, and do a lot of other things in for loops that would otherwise be prone to error and highly time consuming.

Switch Case

- Works like if/else but does not perform boolean assessments
- Improved efficiency under some circumstances (not as good as switch case in other languages)
- demoVariable<-'q';
- switch(demoVariable, a="Got a", b="Got b", c="Got c", "Got something else.");

Sapply / Lapply

- Applies function over specified object or range
- Generally prefer sapply() (neater output)
- someNumbers<-data.frame(rnorm(1000,0,1),rno rm(1000,6,2),rnorm(1000,12,3.6));
- names(someNumbers)<-c("Group1", "Group2", "Group3");
- sapply(someNumbers, summary);
- lapply(someNumbers, summary);

Sequences and Repeat

- seq(from, to, by); rep(thingToRepeat, times);
- seq(from=0, to=1000, by=20);
- rep(1,50);
- *rep(seq(1,5),20);*
- Handy for encoding, generating simulation data, etc.

How do I get my info out?

- (Requires code from slide 21)
- attach(someNumbers);
- output<-t.test(Group1, Group2);
- names(output);
- tVal<-output[[1]];
- tValue<-as.numeric(output[1]);
- The last line grabs just the numeric value, which is handy
- This is essential for making custom functions, running identical tests on massive data collections, etc.

Saving yourself a lot of copy/paste

- source(file=file.choose(new = FALSE));
- corOut<-all.correlations(someNumbers);
- Use the first line to add the function in AllCorrelations.R to your script
- Second line runs it and stores the output
- Note that this script does not correct for multiple comparisons

Handy Bonus Trick

- Need to allow the user to interactively select the working directory?
- library(tcltk);
- setwd(tk_choose.dir(default = "", caption = "Select directory"));