## 18.05 R Tutorial 1B: Random Numbers

In order to run simulations of experiments with random outcomes we make use of R's ability to generate random numbers. More precisely it generates a 'pseudorandom' number, which behaves in many ways like a truly random number.

## The function sample(x, k, replace = TRUE)

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\# sample(x,k) generates a random permutation of k objects from the vector
x. That is, all k choices are different
# start with a vector x with 5 elements.
> x = 1:5
> X
[1] 1 2 3 4 5
# randomly sample 3 of the elements of x.
> sample(x,3)
[1] 2 5 1
> sample(x,3)
[1] 3 1 4
> sample(x,3)
[1] 1 2 5
# Note every element is chosen at most once.
# randomly sampling 5 elements of x is a permutation of all 5 elements.
> sample(x,5)
[1] 1 3 2 5 4
> sample(x,5)
[1] 2 3 1 4 5
# You can't pick more than 5 different elements from a set of 5 things,
so R gives an error.
> sample(x,6)
Error in sample.int(length(x), size, replace, prob) :
  cannot take a sample larger than the population when 'replace = FALSE'
Allowing repeated elements
# Sometimes you want to allow repeats. For example when we roll a die
repeatedly we expect to see numbers repeating. We can think of this as
picking a random element from a set and then putting it back, i.e.
replacing it, so it can be drawn again.
# In R we do this by setting the optional argument replace=TRUE
# Note that we get can get repeated values
```

```
> sample(x,3,replace=TRUE)
[1] 3 1 4
> sample(x,3,replace=TRUE)
[1] 4 5 4
> sample(x,3,replace=TRUE)
[1] 2 1 5
> sample(x,5,replace=TRUE)
[1] 5 1 5 1 4
> sample(x,5,replace=TRUE)
[1] 3 5 5 2 1
> sample(x,5,replace=TRUE)
[1] 4 2 3 2 1
# Now there is no problem asking for more than 5 things from a set of 5
elements.
> sample(x,12,replace=TRUE)
[1] 1 1 2 1 4 2 3 4 1 2 4 5
> sample(x,12,replace=TRUE)
[1] 3 2 5 1 4 2 4 1 4 5 3 1
# To generate an m x n array of random values we can use the function
sample function followed by the matrix function.
# Let's simulate rolling a die.
# We use 1:6 to make vector (1,2,3,4,5,6).
# Generate a 3 x 4 array of random dice rolls.
> y = sample(1:6, 12, replace=TRUE)
> matrix(y,nrow=3,ncol=4)
     [,1] [,2] [,3] [,4]
[1,]
       1
            3
                 6 1
[2,]
        2
             2
                      6
        2
            1
                 4
                      3
[3,]
# Or we could make it a 2 x 6 array.
> matrix(y,nrow=2,ncol=6)
     [,1] [,2] [,3] [,4] [,5] [,6]
[1,]
       1
            2
                 2
                      6
             3
                 1
                      3
                           1
[2,]
       2
                                3
Simulation
# First let's simulate rolling 1 die 3 times and checking if one of the
rolls was a 6.
# First we use sample to generate 3 samples from 1:6
```

> x = sample(1:6,3, replace=TRUE)

> x

```
[1] 6 4 1
# To check if any of the 3 rolls are 6 we use the following command. It
returns a vector of TRUE or FALSE depending on whether that entry of x is
6 or not. Note the use of the double equal sign. We can't use a single
equal sign because that would mean 'set the value of x to 6'. Compare the
result with the value of x above.
> x == 6
[1] TRUE FALSE FALSE
# We can also see which elements of x are less than 6
> x < 6
[1] FALSE TRUE TRUE
# Now let's roll the die 5000 times and see what fraction of the rolls
give 6. We expect that about 1/6 of them will be.
# Simulate 1000 rolls
> x = sample(1:6,1000, replace=TRUE)
# x == 6 gives a vector of TRUE or FALSE. R is clever, when we sum the
vector: each TRUE counts as 1 and each FALSE counts as 0. So the sum is
the number of TRUE's. In this case that means the number of 6's, which
happens to be 168.
> sum(x == 6)
[1] 168
# Divide by 1000 to get the fraction of 6's.
> sum(x == 6)/1000
[1] 0.168
# Compare that with the theoretical value of 1/6.
> 1/6
[1] 0.1666667
# Not bad!
# Now let's estimate the probability of getting at least one 6 in 4
rolls.
# Goal: estimate the probability of getting at least one 6 in 4 rolls.
# Experiment: roll 1 die 4 times and check for a 6.
# Repeat the experiment 10 times and see what fraction of times this
happens.
```

# So you can see all the commands at once, we'll show them all and then
explain them later. For commenting, we'll put a command number before
each '>'
1. > x = matrix(sample(1:6, 4\*10, replace=TRUE), nrow=4, ncol=10)
2. > x

```
[,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10]
[1,]
       5
                                    1
                 1
                      1
                           3
       3
                                                    5
[2,]
            6
                 6
                                              6
       2
            3
                 5
                           5
                                3
                                    4
                                         6
                                                    5
[3,]
       2
[4,]
3. > y = (x==6)
4. > y
     [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10]
[1,] FALSE TRUE FALSE FALSE TRUE FALSE FALSE TRUE FALSE
[2,] FALSE TRUE TRUE FALSE FALSE FALSE FALSE TRUE FALSE
[3,] FALSE FALSE FALSE FALSE FALSE FALSE TRUE FALSE FALSE
[4,] FALSE TRUE TRUE FALSE FALSE FALSE FALSE FALSE FALSE
5. > z = colSums(y)
6. > z
[1] 0 3 2 0 0 1 0 1 2 0
7. > z > 0
[1] FALSE TRUE TRUE FALSE FALSE TRUE FALSE TRUE TRUE FALSE
8. > sum(z > 0)
[1] 5
9. > mean(z > 0)
[1] 0.5
# Command 1: Generate a 4 by 10 random array. Each column represents one
experimental trial of 4 rolls. The 10 columns represent the 10 trials.
# Command 2: Display x
# Command 3: See which of the rolls are 6's. Note the double equals to
check which rolls were 6. The result y is an array of TRUE or FALSE. You
can check that it picks out the 6's.
# Command 4: Display y
# Command 5: Sum each of the columns. The result is then number of 6's in
each
trial (column).
# Command 6: Display z
# Command 7: If the column sum is greater than 0 there was at least one
TRUE in the column, that is at least one 6. This command returns a vector
of TRUE or FALSE representing whether or not the experiment yielded a 6.
```

# command 8: sum the vector in command 7 to get the number of trials that

yielded a 6. We see that 5 out of 10 trials did.

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# command 9: The mean function is just the sum divided by the number of
trials. This is just 5/10 = .5. Half the trials yielded a 10.
# Let's repeat this but with 1000 trials instead of 10. Without all the
comments it's pretty short.
> x = matrix(sample(1:6, 4*1000, replace=TRUE), nrow=4, ncol=1000)
> y = (x==6)
> z = colSums(y)
> sum(z > 0)
[1] 525
> mean(z>0)
[1] 0.525
Our estimate of the probability of at least one 6 in 4 rolls is .525.
This is pretty close to the theoretical value of .518.
The dim() function
# We can always check that x has 4 rows and 1000 columns using the dim()
function.
> dim(x)
       4 1000
[1]
One more simulation
# Goal: estimate the probability of getting a sum
of 7 when rolling two dice.
1. > ntrials = 10000
2. > x = matrix(sample(1:6, 2*ntrials, replace=TRUE), nrow=2,
ncol=ntrials)
3. > y = colSums(x)
4. > mean(y == 7)
[1] 0.1658
# Command 1: We assign the number of trials to the variable ntrials.
Writing code like this makes it easier to modify later. If we want to
change the number of trials to 7 we just have to change this one line of
code.
# Command 2: we create 10000 columns with 2 rows. That is, we run 10000
experiments of rolling 2 dice.
# Command 3: we sum each of the columns, i.e., we sum the two dice.
# Command 4: we find the fraction of sums that are 7.
# Note, this is essentially the exact answer of 1/6.
Exercise: Try to estimate the probability of two sixes
when rolling two dice.
```