

STAT 33B Workbook 7

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This workbook is due **Oct 15, 2020** by 11:59pm PT.

The workbook is organized into sections that correspond to the lecture videos for the week. Watch a video, then do the corresponding exercises *before* moving on to the next video.

Workbooks are graded for completeness, so as long as you make a clear effort to solve each problem, you'll get full credit. That said, make sure you understand the concepts here, because they're likely to reappear in homeworks, quizzes, and later lectures.

As you work, write your answers in this notebook. Answer questions with complete sentences, and put code in code chunks. You can make as many new code chunks as you like.

In the notebook, you can run the line of code where the cursor is by pressing **Ctrl + Enter** on Windows or **Cmd + Enter** on Mac OS X. You can run an entire code chunk by clicking on the green arrow in the upper right corner of the code chunk.

Please do not delete the exercises already in this notebook, because it may interfere with our grading tools.

You need to submit your work in two places:

- Submit this Rmd file with your edits on bCourses.
- Knit and submit the generated PDF file on Gradescope.

If you have any last-minute trouble knitting, **DON'T PANIC**. Submit your Rmd file on time and follow up in office hours or on Piazza to sort out the PDF.

For-loops

Watch the “For-loops” lecture video.

No exercises for this section.

Loop Indices

Watch the “Loop Indices” lecture video.

No exercises for this section.

While-loops

Watch the “While-loops” lecture video.

No exercises for this section.

Preallocation

Watch the “Preallocation” lecture video.

Exercise 1

Use the microbenchmark package to benchmark the “BAD” and “GOOD” example from the lecture video. Benchmark with three different values of *n* (testing both the “BAD” and “GOOD” example for each value). About how much faster is the “GOOD” example?

YOUR ANSWER GOES HERE:

```
#install.packages("microbenchmark")
library(microbenchmark)

bad = function(n) {
  x = c()
  for (i in 1:n) {
    x = c(x, i * 2)
  }
  x
}

good = function(n) {
  x = numeric(n)
  for (i in seq_len(n)) {
    x[i] = i * 2
  }
}

#check with 1e1 iterations
microbenchmark(bad(1e1), times = 100L)

## Unit: microseconds
##      expr    min      lq      mean  median      uq      max  neval
## bad(10) 2.907 3.0625 238.4612   3.171 3.3165 23515.72   100

microbenchmark(good(1e1), times = 100L)

## Unit: microseconds
##      expr    min      lq      mean  median      uq      max  neval
## good(10) 1.64 1.674 48.00721   1.734 1.783 4626.989   100

#check with 1e2 iterations
microbenchmark(bad(1e2), times = 1000L)

## Unit: microseconds
##      expr    min      lq      mean  median      uq      max  neval
## bad(100) 39.995 59.3245 84.32693 62.3475 68.972 20237.97 1000
```

```
microbenchmark(good(1e2), times = 1000L)
```

```
## Unit: microseconds
##      expr    min      lq    mean median      uq    max neval
## good(100) 5.967 6.098 6.368998  6.164 6.2755 51.337  1000
```

#check with 1e3 iterations

```
microbenchmark(bad(1e3), times = 1000L)
```

```
## Unit: milliseconds
##      expr      min      lq    mean  median      uq    max neval
## bad(1000) 1.383179 1.601551 2.641108 1.746335 2.439948 20.90286  1000
```

```
microbenchmark(good(1e3), times = 1000L)
```

```
## Unit: microseconds
##      expr    min      lq    mean  median      uq    max neval
## good(1000) 43.513 47.1005 48.58118 48.0655 49.22 88.498  1000
```

*#From the test we can see that the good example is about
#10-100 times faster than the bad example*

Loops Example

Watch the “Loops Example” lecture video.

Exercise 2

Write a function that returns the first $n + 1$ positions of a 3-dimensional discrete random walk. Return the x, y, and z coordinates in a data frame with columns x, y, and z. Your function should have a parameter n that controls the number of steps.

Hint: For efficiency, use vectors for x, y, and z. Wait to combine them into a data frame until the very last line of your function.

YOUR ANSWER GOES HERE:

```
ThreeDimRandomWalk = function(n) {
  xyz = sample(c(0, 1, 2), n+1, replace = TRUE)
  move = sample(c(-1, 1), n+1, replace = TRUE)
  x <- numeric(n+1)
  y <- numeric(n+1)
  z <- numeric(n+1)
  for (i in seq_len(n+1)) {
    if (xyz[i] == 0) { # x
      x[i + 1] = x[i] + move[i]
      y[i + 1] = y[i]
      z[i + 1] = z[i]
    } else if (xyz[i] == 1) { # y
```

```

      x[i + 1] = x[i]
      y[i + 1] = y[i] + move[i]
      z[i + 1] = z[i]
    } else {
      x[i + 1] = x[i]
      y[i + 1] = y[i]
      z[i + 1] = z[i] + move[i]
    }
  }
  result <- data.frame("x" = x, "y" = y, "z" = z)
  result
}
ThreeDimRandomWalk(10)

```

```

##      x  y  z
## 1    0  0  0
## 2   -1  0  0
## 3    0  0  0
## 4   -1  0  0
## 5   -1 -1  0
## 6   -2 -1  0
## 7   -2 -1 -1
## 8   -2 -2 -1
## 9   -2 -3 -1
## 10  -3 -3 -1
## 11  -3 -3  0
## 12  -3 -4  0

```

Recursion

Watch the “Recursion” lecture video.

Exercise 3

1. Use the microbenchmark package to benchmark `find_fib()` and `find_fib2()` for `n` equal to 1 through 30.
2. Collect the median timings for each into a data frame with a columns `time`, `n`, and `function`. The data frame should have 60 rows (30 for each function).
3. Use `ggplot2` to make a line plot of `n` versus `time`, with a separate line for each `function`.
4. Comment on the the shapes of the lines. Does the computation time grow at the same rate (as `n` increases) for both functions?

YOUR ANSWER GOES HERE:

```

find_fib = function(n) {
  if (n == 1 | n == 2)
    return (1)

```

```

    find_fib(n - 2) + find_fib(n - 1)
  }

find_fib2 = function(n, fib = c(1, 1)) {
  len = length(fib)
  if (n <= len)
    return (fib[n])

  fib = c(fib, fib[len - 1] + fib[len])
  find_fib2(n, fib)
}

#since 30 is too large for my computer (it crashes when I tried to run with 30 iterations),
#I choose to run with 20 iterations for each functions
second = numeric(40)
name = character(40)
n = numeric(40)
for (i in 1:20) {
  second[i] = summary(microbenchmark(find_fib(i), times = 1000L, unit = 'ms'))[1, "median"]
  name[i] = 'find_fib()'
  n[i] = i
}
for (i in 1:20) {
  second[i+20] = summary(microbenchmark(find_fib2(i), times = 1000L, unit = 'ms'))[1, "median"]
  name[i+20] = 'find_fib2()'
  n[i+20] = i
}
data_fib <- data.frame("name" = name, "n" = n, "time" = second)
data_fib

```

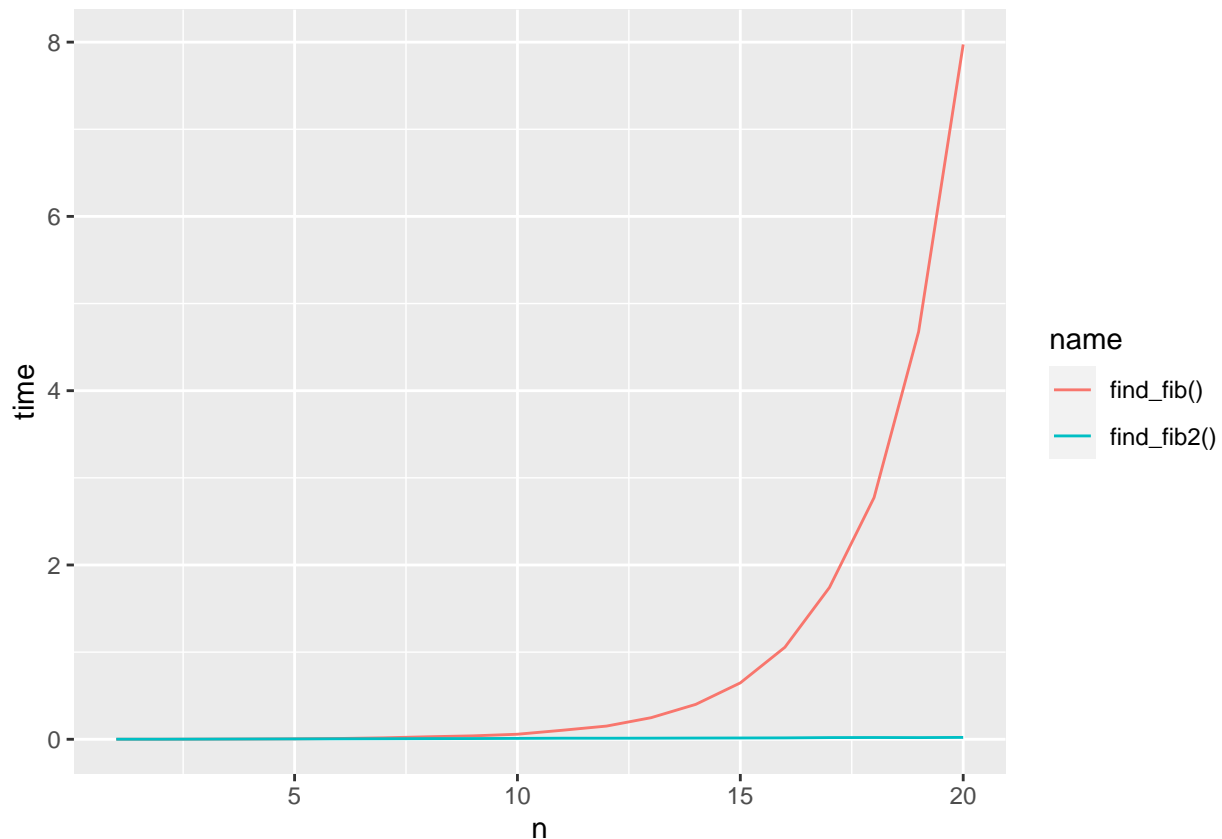
```

##           name  n      time
## 1  find_fib()  1 0.0005260
## 2  find_fib()  2 0.0005130
## 3  find_fib()  3 0.0017940
## 4  find_fib()  4 0.0030835
## 5  find_fib()  5 0.0051810
## 6  find_fib()  6 0.0087135
## 7  find_fib()  7 0.0163770
## 8  find_fib()  8 0.0287085
## 9  find_fib()  9 0.0390705
## 10 find_fib() 10 0.0577030
## 11 find_fib() 11 0.1031210
## 12 find_fib() 12 0.1515715
## 13 find_fib() 13 0.2478040
## 14 find_fib() 14 0.4005430
## 15 find_fib() 15 0.6471620
## 16 find_fib() 16 1.0558620
## 17 find_fib() 17 1.7421690
## 18 find_fib() 18 2.7730785
## 19 find_fib() 19 4.6769565
## 20 find_fib() 20 7.9736650
## 21 find_fib2()  1 0.0008230
## 22 find_fib2()  2 0.0008240
## 23 find_fib2()  3 0.0019130

```

```
## 24 find_fib2() 4 0.0028860
## 25 find_fib2() 5 0.0040580
## 26 find_fib2() 6 0.0051190
## 27 find_fib2() 7 0.0060935
## 28 find_fib2() 8 0.0072470
## 29 find_fib2() 9 0.0082680
## 30 find_fib2() 10 0.0101505
## 31 find_fib2() 11 0.0120700
## 32 find_fib2() 12 0.0118985
## 33 find_fib2() 13 0.0125525
## 34 find_fib2() 14 0.0135715
## 35 find_fib2() 15 0.0145320
## 36 find_fib2() 16 0.0158275
## 37 find_fib2() 17 0.0191220
## 38 find_fib2() 18 0.0200480
## 39 find_fib2() 19 0.0192180
## 40 find_fib2() 20 0.0212565
```

```
#install.packages("ggplot2")
library(ggplot2)
ggplot(data_fib, aes(x = n, y = time, color = name)) + geom_line()
```



```
#from the plot we an see that the function does not grow in the same rate.
# find_fib() grows exponentially large as n gets large, where find_fib2()
#takes about constant time
```

Developing Iterative Code

Watch the “Developing Iterative Code” lecture video.

No exercises for this section. All done!