## Titles and citations — Letchford et al. 2015

Letchford et al. (2015) found an interesting pattern: papers that have shorter titles tend to fare better in terms of citations. They took top-cited papers from a variety of journals, and ranked them by title length (in number of characters), and number citations received (as of November 2014). Then they performed a Kendall's tau test to see whether these rankings are correlated. A negative correlation would mean that the articles with longer titles tend to be ranked low for citations.

The file Letchford2015\_data.csv contains the data needed to replicate their results.

1. Write a program that performs the test described above using all the papers published in 2010. The program should do the following: 1) read the data; 2) extract all the papers published in 2010; 3) rank the articles by citations, and by title length; 4) compute the Kendall's tau expressing the correlation between the two rankings. For this dataset, the Authors got a tau of about -0.07 with a significant p-value.

First, we need to read the data:

```
# Read the data
12015 <- read.csv("../data/Letchford2015_data.csv", stringsAsFactors = FALSE)
# Check dimensions
dim(12015)</pre>
```

```
## [1] 140000
```

```
# Print first few lines
head(12015)
```

```
##
                                      journal title_length cites
     year
             Molecular Biology and Evolution
## 1 2007
                                                         75 17958
## 2 2007
                            Nature Materials
                                                         20 11633
## 3 2007
                               Bioinformatics
                                                         35
                                                           8437
## 4 2007
                                         Cell
                                                         83
                                                            5846
## 5 2007 Ca-A Cancer Journal for Clinicians
                                                         23
                                                            5744
## 6 2007 American Journal of Human Genetics
                                                         84
                                                           5651
```

Now extract only the papers published in 2010:

```
p2010 <- 12015[12015$year == 2010,]
```

To rank the manuscripts according to title length, we can use the function rank (check ?rank), which ranks the entries, resolving possible ties:

```
# Example of use of rank rank(c(1,2,3,2,1,5,3,4))
```

```
## [1] 1.5 3.5 5.5 3.5 1.5 8.0 5.5 7.0
```

Let's store the ranking of title lengths and citations separately:

```
rank_titlelength <- rank(p2010$title_length)
rank_citations <- rank(p2010$cites)</pre>
```

Finally, let's calculate Kendall's tau. A simple way is to invoke cor:

```
cor(rank_citations, rank_titlelength, method = "kendall", use = "pairwise")
```

```
## [1] -0.06551962
```

Which is similar to that reported by the Authors (they performed some filtering of the articles before analyzing them, so we don't expect a perfect match). To get also a p-value, you can use cor.test:

```
cor.test(rank_citations, rank_titlelength, method = "kendall", use = "pairwise")
```

```
##
## Kendall's rank correlation tau
##
## data: rank_citations and rank_titlelength
## z = -13.753, p-value < 2.2e-16
## alternative hypothesis: true tau is not equal to 0
## sample estimates:
## tau
## -0.06551962</pre>
```

2. Write a function that repeats the analysis for a particular journal-year combination. Try to run the function for the top scientific publications Nature and Science, and for the top medical journals The Lancet and New Eng J Med, for all years in the data (2007-2013). Do you always find a negative, significant correlation (i.e., negative tau with low p-value)?

For this point, we need to write a function, that takes as input the data, a journal and a year. The function will then extract the relevant data, and run the test.

We can start building the function from this skeleton:

```
compute_tau_journal_year <- function(my_data, my_journal, my_year){
    # First, filter the data
    my_subset <- my_data[my_data$journal == my_journal & my_data$year == my_year,]
    print(c(my_journal, my_year, "Articles:", dim(my_subset)[1]))
}</pre>
```

and try running it to make sure everything is good:

```
compute_tau_journal_year(12015, "Nature", 2010)
```

```
## [1] "Nature" "2010" "Articles:" "575"
```

Next, we write the analysis proper:

```
## Journal Year tau
## 1 Nature 2010 -0.03833785
```

We can write a fancier version that stores also the p-values, and checks that there are enough articles:

```
compute_tau_journal_year <- function(my_data, my_journal, my_year){</pre>
  # First, filter the data
  my_subset <- my_data[my_data$journal == my_journal & my_data$year == my_year,]</pre>
  if (dim(my_subset)[1] < 2) {</pre>
    tau <- NA
    p.value <- NA
  } else {
    # Rank by title length and citations
    rank_titlelength <- rank(my_subset$title_length)</pre>
    rank_citations <- rank(my_subset$cites)</pre>
    # Run the test
    my_test <- cor.test(rank_citations, rank_titlelength,</pre>
                         method = "kendall", use = "pairwise")
    tau <- as.numeric(my_test$estimate)</pre>
    p.value <- as.numeric(my_test$p.value)</pre>
  return(data.frame(Journal = my_journal,
                     Year = my_year,
                     tau = tau,
                     p.value = p.value))
}
compute_tau_journal_year(12015, "Nature", 2010)
```

```
## Journal Year tau p.value
## 1 Nature 2010 -0.03833785 0.1746622
```

Now let's run it for all years and a few journals:

```
results <- data.frame()
for (year in 2007:2013){</pre>
```

We can see that we can get both positive or negative tau(s):

## results

```
##
            Journal Year
                                            p.value
                                    tau
## 1
             Nature 2007
                           0.0115327643 0.695512738
## 2
            Science 2007 -0.0232164706 0.443764950
## 3
         The Lancet 2007
                                     NA
## 4
      New Eng J Med 2007
                          0.0865486187 0.066693254
## 5
             Nature 2008
                         -0.0804201182 0.004800352
## 6
                          0.0067644073 0.824789544
            Science 2008
## 7
         The Lancet 2008
                          0.1499256823 0.007689070
## 8
      New Eng J Med 2008
                          0.1170901670 0.012675009
## 9
             Nature 2009 -0.0104426422 0.723168401
## 10
            Science 2009 -0.0218930503 0.462036960
## 11
         The Lancet 2009
                          0.0592239256 0.305138499
## 12
     New Eng J Med 2009
                          0.1276904509 0.006284098
             Nature 2010 -0.0383378489 0.174662196
## 13
## 14
            Science 2010 -0.0311213024 0.314116809
## 15
         The Lancet 2010
                          0.1210024231 0.029256634
## 16
     New Eng J Med 2010
                          0.0849086854 0.069678724
             Nature 2011 -0.0279099578 0.323149648
## 17
## 18
            Science 2011
                          0.0171414424 0.579261893
## 19
         The Lancet 2011
                          0.0648182972 0.209265286
      New Eng J Med 2011 -0.0217783333 0.646159372
## 20
## 21
             Nature 2012
                          0.0159125585 0.571748517
## 22
            Science 2012
                          0.0211241243 0.508981567
## 23
         The Lancet 2012
                          0.1024408564 0.066156323
## 24
     New Eng J Med 2012
                          0.1454023269 0.002005165
## 25
             Nature 2013 -0.0344594481 0.242130179
## 26
            Science 2013
                          0.0257910388 0.419193129
## 27
         The Lancet 2013
                          0.1396061773 0.016167487
## 28 New Eng J Med 2013
                          0.0002704937 0.995615193
```

However, to be sure we have a meaningful result, we should correct for multiple testing (when trying very many tests, we can obtain a number of significant *p*-values just by chance), either applying Bonferroni's correction, or using more sophisticated false-discovery-rate approaches.