CS532 Web Science: Assignment 3

Finished on February 18, 2016

Dr. Michael L. Nelson

Naina Sai Tipparti ntippart@cs.odu.edu

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Question

Download the 1000 URIs from assignment #2. "curl", "wget", or "lynx" are all good candidate programs to use. We want just the raw HTML, not the images, stylesheets, etc.

from the command line:

```
% curl http://www.cnn.com/ > www.cnn.com
```

% wget -O www.cnn.com http://www.cnn.com/

% lynx -source http://www.cnn.com/ > www.cnn.com

"www.cnn.com" is just an example output file name, keep in mind that the shell will not like some of the characters that can occur in URIs (e.g., "?", "&"). You might want to hash the URIs, like:

```
\%echo -n "http://www.cs.odu.edu/show_features.shtml?72" |md541d5f125d13b4bb554e6e31b6b591eeb
```

("md5sum" on some machines; note the "-n" in echo - this removes the trailing newline.)

Now use a tool to remove (most) of the HTML markup. "lynx" will do a fair job:

% lynx -dump -force_html www.cnn.com > www.cnn.com.processed Use another (better) tool if you know of one. Keep both files for each URI (i.e., raw HTML and processed).

Answer

Using the python script in Listing 1, 1000 unique URIs were dereferenced and their raw contents were stored in the html/raw/ folder as a file with the filename as the md5 [4]-hashed URI. These were then stripped of all html elements and their processed contents were stored in the html/processed/ folder as the same md5-hashed filename. For reference, the URIs were written as the first line of each of their content files.[8][2][1]

```
1 #! /usr/bin/python
import requests
import concurrent.futures
import md5
from bs4 import BeautifulSoup
import pickle
def convert(uri):
```

```
return md5.new(uri).hexdigest()
11
   def get html(uri):
       print('Getting {}'.format(uri))
       response = requests.get(uri)
       return response.url, response.status code, response.content
16
       name__ == '_main__':
with open('links') as infile:
            uris = [uri.rstrip('\n') for uri in infile]
       with \verb| concurrent.futures.ThreadPoolExecutor(max\_workers=8)| as executor:
21
            uri_futures = [executor.submit(get_html, uri) for uri in uris] for future in concurrent.futures.as_completed(uri_futures):
                 t r y:
                     uri , status code , content = future.result()
                 except Exception as exc:
26
                      print('{} generated an exception: {}'.format(uri, exc))
                     continue
                 if status code == 200:
                     hashed uri = convert (uri)
31
                      print( \( \frac{7}{V}\) Writing \( \{ \} \) as \( \{ \} \) . format(uri, hashed_uri) )
                          with open('html/raw/' + hashed uri, 'w') as outfile:
                               outfile.write(uri + '\n')
                               outfile.write(content)
36
                          with open('html/processed/' + hashed uri + '.processed.txt', 'w') as
                               outfile:
                               outfile.write(uri + '\n')
                               outfile.write(BeautifulSoup(content).get text().encode('utf8'))
                     except Exception as e:
                          print '**** ERROR **** --- ' + uri
41
                          print e
                 else:
                     print('Not writing {}, bad status code: {}'.format(uri, status code))
```

Listing 1: get_html.py

Question

Choose a query term (e.g., "shadow") that is not a stop word (see week 5 slides) and not HTML markup from step 1 (e.g., "http") that matches at least 10 documents (hint: use "grep" on the processed files). If the term is present in more than 10 documents, choose any 10 from your list. (If you do not end up with a list of 10 URIs, you've done something wrong).

As per the example in the week 5 slides, compute TFIDF values for the term in each of the 10 documents and create a table with the TF, IDF, and TFIDF values, as well as the corresponding URIs. The URIs will be ranked in decreasing order by TFIDF values. For example:

Table 1. 10 Hits for the term "shadow", ranked by TFIDF.

```
TFIDF TF IDF URI
0.150 0.014 10.680 http://foo.com/
0.044 0.008 5.510 http://bar.com/
```

You can use Google or Bing for the DF estimation. To count the number of words in the processed document (i.e., the deonminator for TF), you can use "wc":

```
\% wc -w www.cnn.com.processed 2370 www.cnn.com.processed
```

It won't be completely accurate, but it will be probably be consistently inaccurate across all files. You can use more accurate methods if you'd like.

Don't forget the log base 2 for IDF, and mind your significant digits!

Answer

First, the function count_terms was used to count[9] the term frequency for the given term "shadow" in all documents.

```
def count_terms(term, file_list=os.listdir('html/processed')):
    for filename in file_list:
        with open('html/processed/' + filename + '.processed.txt') as infile:
            uri = infile.readline().strip()
            text = infile.read()
            count = text.count(term)
            if count > 0:
                print('{{}} {{}}'.format(count, uri))
                return count, uri
```

Listing 2: count terms function

Ten of the results were chosen at random and stored in the uri_counts file. In order to easily identify which file corresponds to which URI, since the filename is the non-reversible md5-hashed URI string, a mapping from URI to filename was created using the functions in Listing 3 and serialized in the uri_map file using the pickle[7] library.

```
def get_uri(uri):
25
       for filename in os.listdir('html/processed/'):
           with open ('html/processed/' + filename + '.processed.txt') as infile:
               if uri in infile readline():
                    return uri, filename
       return None, None
30
   def get_uris():
       uri file = \{\}
       for uri in open('links').read().split('\n'):
           uri, filename = get_uri(uri)
35
           if not uri:
               continue
           uri file [uri] = filename
```

Listing 3: get uris functions

Reading from the file was done with the line in Listing 4. This loaded the serialized URI to filename map for future use.

```
10 map_file = open('uri_map', 'rb')
```

Listing 4: Loading the uri map

To proceed with processing each of the files to find Term Frequency (TF), Inverse Document Frequency (IDF) and the product of the two (TFIDF), each URI's corresponding file was found using the get_filename function found in Listing 5.

```
def get_filename(uri):
    if uri_map.has_key(uri):
        return uri_map[uri]
```

Listing 5: Getting filename from URI

Then, they were stripped of HTML tags using the strip_html function in Listing 6.

```
45
   def strip html(filename):
       if not filename:
           print
                 'invalid filename'
           return
       with open('html/processed/' + filename + '.processed.txt') as infile:
           # To remove URI in first line
50
           infile.readline()
           # Removing all punctuation
           strs = infile.read()
           r = re.compile(r'[\{\}]'.format(punctuation))
           content = r.sub(
55
                               , strs)
```

Listing 6: Stripping HTML tags from content

And finally the frequencies were calculated for each URI using the functions in Listing 7.

```
def get tf(content, term):
       return float (content.count(term)) / float(len(content.split()))
60
   def get idf(term):
       present = set()
       absent = set()
       for uri, filename in uri map. iteritems():
65
           content = strip_html(filename)
            if not content:
                continue
            if term in content:
                present.add(uri)
70
                absent.add(uri)
       return math.log(float(len(absent)) / float(len(present)), 2)
   def process uri(uri, term):
75
       tf = get tf(strip html(get filename(uri)), term)
       t f i d f = \overline{t} f * i d f
       return tf, tfidf
```

Listing 7: Calculating TF, IDF & TFIDF

These frequencies were then written to the uri_frequencies file using the code in 8.

```
term = sys.argv[2]
with open('uri_counts') as infile:
    uris = uris = [line.split()[1] for line in infile.read().split('\n')]
with open('uri_frequencies', 'w') as outfile:
    outfile.write('{:<7} {:<7} {:<7} \n'.format('TFIDF', 'TF', 'IDF', 'URI')
    ))
    for uri in uris:
        tf, tfidf = process_uri(uri, term)
```

Listing 8: Writing results to uri frequencies file

And the results can be seen in Listing 9:

```
TFIDF
                                    \operatorname{TF}
                                                                         {\rm IDF}
                                                                                                               URI
0.0220
                                    0.0065
                                                                         3.3825
                                                                                                              http://news.google.com/
                                                                                                               http://www.easkme.com/2014/07/mail-merge-in-gmail.html#.VB8GnkIk6rA.
0.0159
                                    0.0047
                                                                          3.3825
                  facebook
0.0113 0.0033
                                                                       3.3825
                                                                                                            http://btc-news-bot.tumblr.com/post/98066358831/we-need-to-do-a-
                  better-job-explaining-bitcoin-in-ways#_=_
0.0109 0.0032 3.3825 http://musicisthedrug-revolution.tumblr.com/post/98067184737/
                  description\_you\_know\_what\_time\_it\_is\_world\_cup\#\_=
0.0099 \quad 0.0029 \quad 3.3825 \quad \text{http://www.ebay.com/itm/4-5-Android-Smartphone-Dual-Sim-Dual-Core-property} \\ \text{Sim-Dual-Core-property} \\ \text{Sim
                  Unlocked-WIFI-3G-GSM-Cell-phone-AT-T-/271610837947? pt=Cell\_Phones\&hash=item3f3d447bbb
 0.0092 0.0027 3.3825 http://www.blogdeizquierda.com/2014/09/mundo-bitcoin-banqueros-de-eu
                  -ven.html?utm_source=twitterfeed&utm_medium=twitter
0.0064 0.0019 3.3825 http://www.valuewalk.com/2014/09/best-apps-apple-iphone-6/
0.0064 0.0019
                                                                      3.3825 http://www.datelinemovies.com/2014/07/bloopers-for-season-4-game-of-
                  t\,h\,ro\,n\,es\,.\,h\,t\,m\,l\#st\,h\,a\,s\,h\,\,.\,y\,YuV0eDx\,.\,\,u\,x\,fs
0.0027 \quad 0.0008 \quad 3.3825 \quad \text{http://abusidiqu.com/its-all-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a-biological-scripted-ebola-virus-is-a
                  weapon-from-the-us-read-this-shocking-report/
0.0019 0.0006 3.3825 http://rss-now.blogspot.com/2014/09/nasa-boeing-space-x-iss.html?
                  \verb"utm_source=dlvr.it&utm_medium=twitter""
```

Listing 9: uri frequencies file

Question

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Answer

Using the Page Rank[6] Checker website to input each of the URIs found in the ten selected URIs from question 2 the results in Listing 10 was determined.

```
PageRank
            URI
0.8
            http://www.ebay.com/
0.8
            http://news.google.com/
0.5
            http://www.valuewalk.com/
            http://www.blogdeizquierda.com/
0.2
0.2
            http://abusidiqu.com/
            http://www.easkme.com/
0.0
            http://www.datelinemovies.com/
0.0
            http://rss-now.blogspot.com/
0.0
            http://musicisthedrug-revolution.tumblr.com/
0.0
0.0
            http://btc-news-bot.tumblr.com/
```

Listing 10: page ranks file

In looking at the similarities and differences in the results of question 2 and question 3 it seems that page rank is unrelated to term frequency measurements. This is logical because the search term isn't taken as an input when calculating page rank. Also, finding page rank has a different goal than measuring search term relevance. It is used to objectively find which pages have a higher probability of a user randomly navigating to the page, which is unrelated to the content of the pages in the given set and is a function of the graph created by links contained in the pages of the set.

Question

Compute the Kendall Tau_b score for both lists (use "b" because there will likely be tie values in the rankings). Report both the Tau value and the "p" value.

See:

http://stackoverflow.com/questions/2557863/measures-of-association-in-r-kendalls-tau-b-and-tau-c

 $http://en.wikipedia.org/wiki/Kendall_tau_rank_correlation_coefficient \# Tau-bhttp://en.wikipedia.org/wiki/Correlation_and_dependence$

Answer

Using R, I input the values of TFIDF and Page rank into two separate vectors. Using the package[5] Kendall[3], it calculated a tau of 0.866 and p of 0.0018457. With a tau number so close to +1, there is a high association between the two measured quantities. In the case of ties, both packages "Kendall and cor produce the same result but cor.test produces a p-value which is not as accurate".

```
#! /usr/bin/Rscript

library(Kendall)
data <- read.table("D:/cs532/a3/q4/kendall", header=TRUE)
tfidf <- data[1]
pagerank <- data[2]
cor(tfidf, pagerank, method="kendall")
Kendall(tfidf, pagerank)</pre>
```

Listing 11: kendall.r

Figure 1: R script that computes the Tau and p value of the Kendall Tau correlation

```
R Console

> Kendall (tfidf, pagerank)
tau = 0.866, 2-sided pvalue =0.0018457
> |
```

Figure 2: Tau and p value of the Kendall

References

- [1] Beautiful soup documentation. http://www.crummy.com/software/BeautifulSoup/bs4/doc/.
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- [7] 11.1. pickle python object serialization. 11.1. pickle Python object serialization.
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