Assignment 8

Fall 2014
CS595 Web Science
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Contents

Listings

1 Question 1

1.1 Question

```
Create a blog-term matrix.  Start by grabbing 100 blogs; include:

http://f-measure.blogspot.com/
http://ws-dl.blogspot.com/

and grab 98 more as per the method shown in class.

Use the blog title as the identifier for each blog (and row of the
matrix).  Use the terms from every item/title (RSS) or entry/title
(Atom) for the columns of the matrix.  The values are the frequency
of occurrence.  Essentially you are replicating the format of the
"blogdata.txt" file included with the PCI book code.  Limit the
number of terms to the most "popular" (i.e., frequent) 500 terms,
this is *after* the criteria on p. 32 (slide 7) has been satisfied.

Create a histogram of how many pages each blog has (e.g., 30
blogs with just one page, 27 with two pages, 29 with 3 pages and
so on).
```

1.2 Answer

To complete this assignment, a blog word count dataset was required. To start off, a list of blog URIs was obtained using the method described in class, implemented as the `get_uris.py` script. Two default blogs, F-Measure and the Old Dominion Web Science and Digital Libraries blogs, were added to the list and then, using the seed URI provided (Listing 2), the remaining 98 URIs from random blogs within the blogger.com family were added. Then, using the `matrix.py` script, the page counts for each blog were extracted and saved to a file called `pagecounts`. The `matrix.py` script is a modified version of `generatefeedvectors.py` from the book *Programming Collective Intelligence* [1].

```
27 if __name__ == '__main__':
28     uris = set()
29     with open('blog_uris', 'a') as outfile:
30         if len(sys.argv) > 1 and sys.argv[1] == 'new':
31             for must_have in must_haves:
32                 uri = get_atom(must_have)
33                 add_uri(uri, uris, outfile)
34         else:
35             with open('blog_uris') as infile:
36                 [uris.add(line.strip()) for line in infile]
37         while len(uris) < 100:
38             uri = get_atom(default)
39             add_uri(uri, uris, outfile)
```

Listing 1: main for get_uris.py

```
7 default = 'http://www.blogger.com/next-blog?navBar=true&blogID=3471633091411211117'
8 must_haves = ['http://f-measure.blogspot.com/', 'http://ws-dl.blogspot.com/']
```

Listing 2: referenced variables in get_uris.py

The `get_uris main` function in Listing 1 was the driver that called the `get_atom` function (shown in Listing 3) to extract the atom [2] URIs from each blog and add them to the set of URIs with the `add_uri` function, shown in Listing 4.

```
def get_atom(uri):
    try:
        r = requests.get(uri)
    except Exception, e:
        return None
    soup = BeautifulSoup(r.text)
    links = soup.find_all('link', {'type':'application/atom+xml'})
    if links:
        return str(links[0]['href'])
    return None
```

Listing 3: get_atom function

```
def add_uri(uri, uris, outfile):
    if uri and uri not in uris:
        uris.add(uri)
        outfile.write(uri + '\n')
        print len(uris), uri
```

Listing 4: add_uri function

The contents of each blog were downloaded and processed by the code shown in Listing 5 and the `get_titles`, `get_words` and `get_next` functions found in Listing 6. This code loops over the URIs that were downloaded with the `get_uris.py` script, parses each entry and extracts all the words in each entry's title. These words were then compiled into a master list for all 100 blogs, with the top 500 words that fit into the range bounded by the code in Listing 7 being used for the final word count.

```
    with open('blog_uris') as infile:
        uris = [line.strip() for line in infile]
    if len(sys.argv) == 2 and sys.argv[1] == 'get':
        with futures.ThreadPoolExecutor(max_workers=8) as executor:
            uri_futures = [executor.submit(get_titles, uri) for uri in uris]
            for future in futures.as_completed(uri_futures):
                uri, title, subtitle, pages, wc = future.result()
                with open('wcs/' + md5.new(uri).hexdigest(), 'w') as out:
                    out.write(title + ': ' + subtitle + '\t' + str(pages) + '\t')
                    json.dump(wc, out)
```

Listing 5: looping over the URIs

```
def get_next(d):
    for item in d.feed.links:
        if item['rel'] == u'next':
            return item['href']
    return None

def getwords(text):
    txt = re.compile(r'<[^>]+>').sub('', text)
    words = re.compile(r'[^A-Z^a-z]+').split(txt)
    return [word.lower() for word in words if word != '']

def get_titles(uri):
    print('processing {}'.format(uri))
    next = uri
    wc = {}
    pages = 0
    while next is not None:
        d = feedparser.parse(next)
        for e in d.entries:
            words = getwords(e.title.encode('utf-8'))
            for word in words:
                wc.setdefault(word, 0)
                wc[word] += 1
        pages += 1
        next = get_next(d)
        print('next {}'.format(next))
    title = d.feed.title.encode('utf-8')
    subtitle = d.feed.subtitle[:50].encode('utf-8')
    print('finished: {}: {}'.format(title, subtitle))
    return uri, title, subtitle, pages, wc
```

Listing 6: processing each blog

```
        for w, bc in sorted(apcount.items(), key=lambda x: x[1], reverse=True):
            frac = float(bc) / len(uris)
            if frac > 0.1 and frac < 0.5:
                wordlist.append(w)
```

Listing 7: bounding the terms

To build a histogram showing the blog page counts, the `pagecounts` file was parsed by the R script in Listing 8 and saved as a pdf, which is shown in Figure 2.

```
#! /usr/bin/Rscript

data <- read.table("pagecounts", sep="\t", header=TRUE, comment.char="")
counts <- table(data$pages)
pdf("hist.pdf")
barplot(counts, ylab="Number of Blogs", xlab="Page Count", main="Page Count per Blog")
dev.off()
```

Listing 8: building the histogram

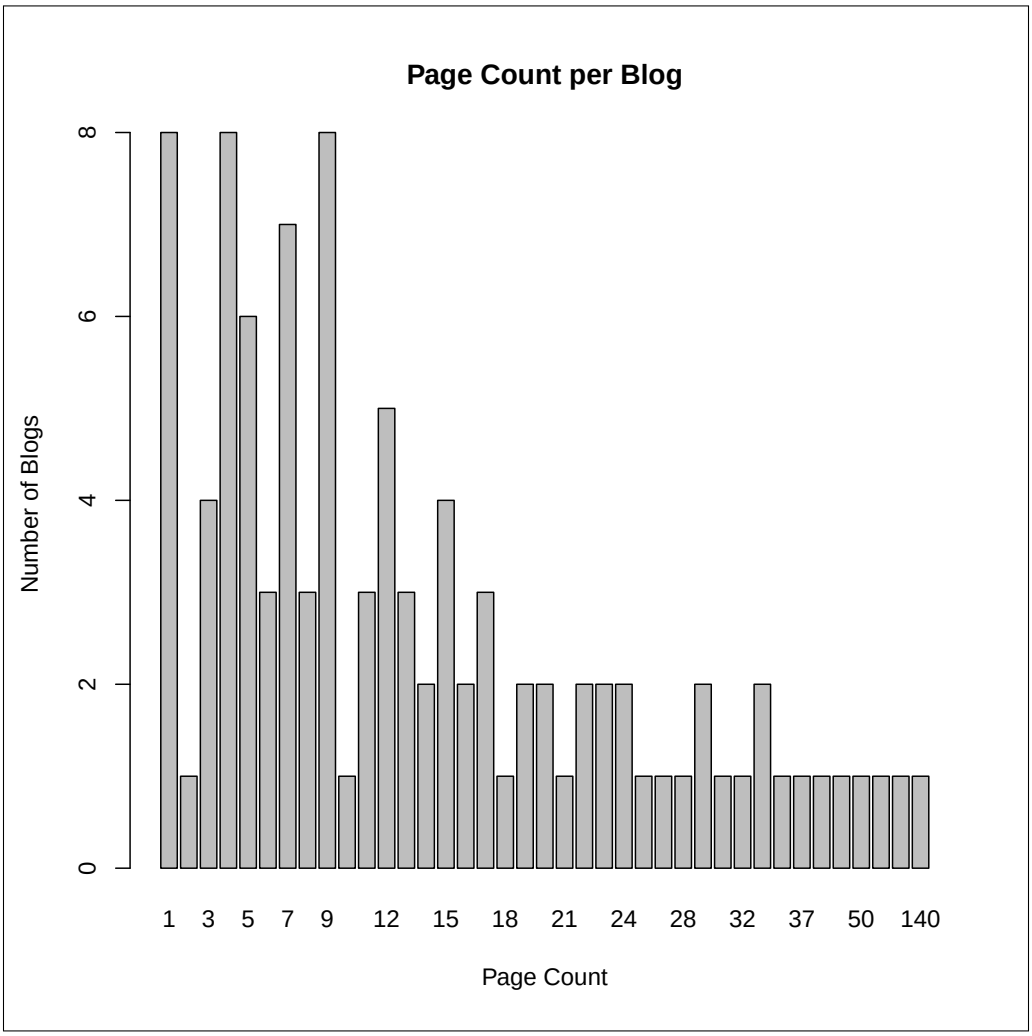

Figure 1: Page Count per Blog

2 Question 2

2.1 Question

Create an ASCII and JPEG dendrogram that clusters (i.e., HAC) the most similar blogs (see slides 12 & 13). Include the JPEG in your report and upload the ascii file to github (it will be too unwieldy for inclusion in the report).

2.2 Answer

The ascii and jpeg dendrograms were created using the method shown in Listing 9, which is modeled after the example from class.

```
286     blognames, words, data = readfile('blogdata1.txt')
287     clust = hcluster(data)
288     with open('dendrogram.txt', 'w') as outfile:
289         stdout = sys.stdout
290         sys.stdout = outfile
291         printclust(clust, labels=blognames)
292         sys.stdout = stdout
293     drawdendrogram(clust, blognames, jpeg='blogclust.jpg')
```

Listing 9: creating the dendrograms

This uses the `readfile` function shown in Listing 10 to read the data that was compiled from Question 1 into the script where it is then processed by the `hcluster` function found in Listing 11 to produce the clustered representation of the blogs.

```
3  def readfile(filename):
4    lines=[line for line in file(filename)]
5
6    # First line is the column titles
7    colnames=lines[0].strip().split('\t')[1:]
8    rownames=[]
9    data=[]
10   for line in lines[1:]:
11     p=line.strip().split('\t')
12     # First column in each row is the rowname
13     rownames.append(p[0])
14     # The data for this row is the remainder of the row
15     data.append([float(x) for x in p[1:]])
16   return rownames,colnames,data
```

Listing 10: creating the dendrograms

```
48 def hcluster(rows,distance=pearson):
49   distances={}
50   currentclustid=-1
51
52   # Clusters are initially just the rows
53   clust=[bicluster(rows[i],id=i) for i in range(len(rows))]
54
55   while len(clust)>1:
56     lowestpair=(0,1)
57     closest=distance(clust[0].vec,clust[1].vec)
58
59     # loop through every pair looking for the smallest distance
60     for i in range(len(clust)):
61       for j in range(i+1,len(clust)):
62         # distances is the cache of distance calculations
63         if (clust[i].id,clust[j].id) not in distances:
64           distances[(clust[i].id,clust[j].id)]=distance(clust[i].vec,clust[j].vec)
65
66         d=distances[(clust[i].id,clust[j].id)]
67
68         if d<closest:
69           closest=d
```

```
70              lowestpair=(i,j)
71
72      # calculate the average of the two clusters
73      mergevec=[
74      (clust[lowestpair[0]].vec[i]+clust[lowestpair[1]].vec[i])/2.0
75      for i in range(len(clust[0].vec))]
76
77      # create the new cluster
78      newcluster=bicluster(mergevec,left=clust[lowestpair[0]],
79                           right=clust[lowestpair[1]],
80                           distance=closest,id=currentclustid)
81
82      # cluster ids that weren't in the original set are negative
83      currentclustid-=1
84      del clust[lowestpair[1]]
85      del clust[lowestpair[0]]
86      clust.append(newcluster)
87
88    return clust[0]
```

Listing 11: hcluster function

The `printclust` function from Listing 12 prints the ascii dendrogram of the cluster object parameter.

```
90  def printclust(clust,labels=None,n=0):
91    # indent to make a hierarchy layout
92    for i in range(n): print ' ',
93    if clust.id<0:
94      # negative id means that this is branch
95      print '-'
96    else:
97      # positive id means that this is an endpoint
98      if labels==None: print clust.id
99      else: print labels[clust.id]
100
101   # now print the right and left branches
102   if clust.left!=None: printclust(clust.left,labels=labels,n=n+1)
103   if clust.right!=None: printclust(clust.right,labels=labels,n=n+1)
```

Listing 12: printclust function

The `drawdendrogram` function from Listing 13 creates a jpeg image of the cluster, which is shown in Figure ??.

```
122 def drawdendrogram(clust,labels,jpeg='clusters.jpg'):
123   # height and width
124   h=getheight(clust)*20
125   w=1200
126   depth=getdepth(clust)
127
128   # width is fixed, so scale distances accordingly
129   scaling=float(w-150)/depth
130
131   # Create a new image with a white background
132   img=Image.new('RGB',(w,h),(255,255,255))
133   draw=ImageDraw.Draw(img)
134
135   draw.line((0,h/2,10,h/2),fill=(255,0,0))
136
137   # Draw the first node
138   drawnode(draw,clust,10,(h/2),scaling,labels)
139   img.save(jpeg,'JPEG')
```

Listing 13: drawdendrogram function

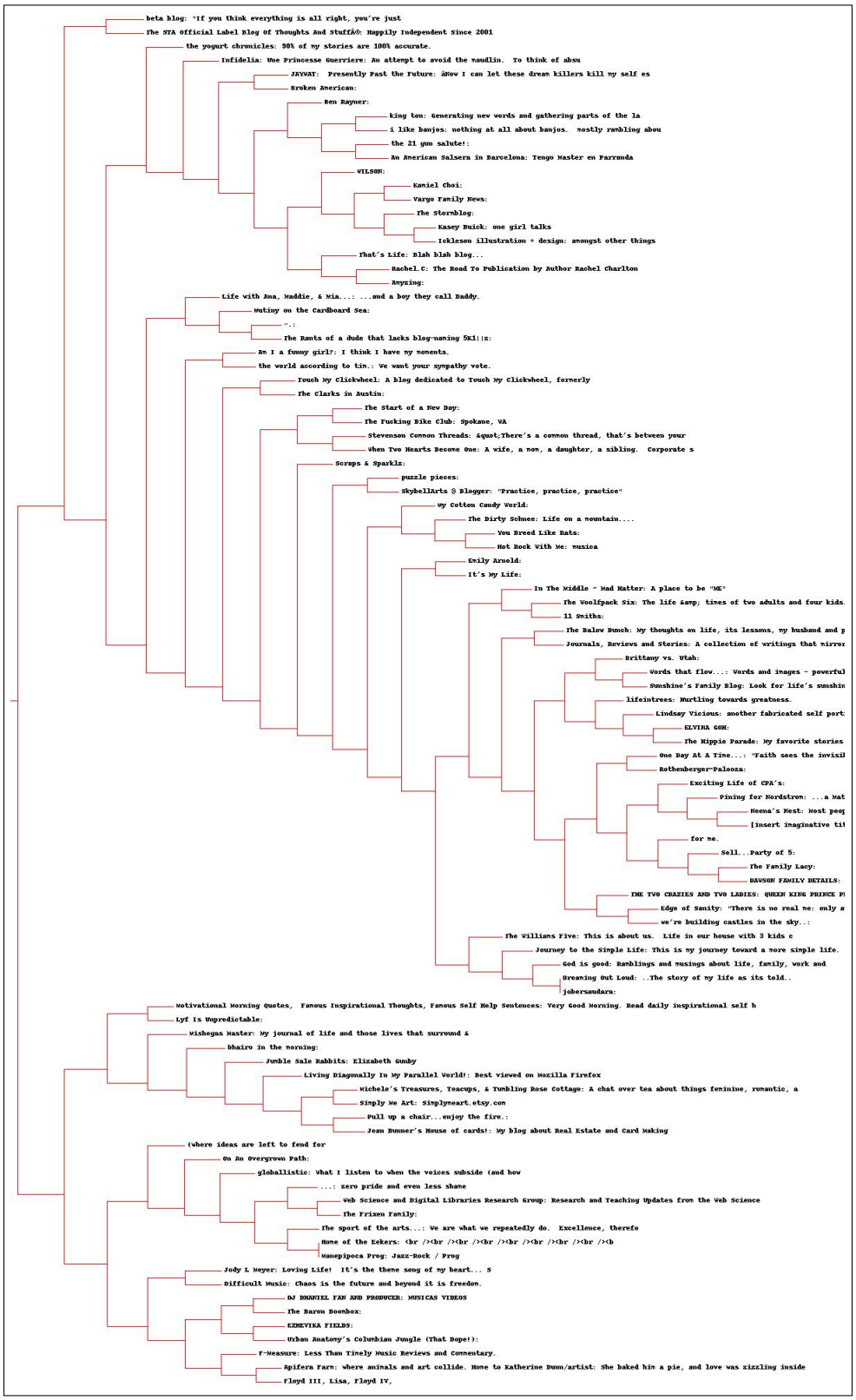

Figure 2: dendrogram

3 Question 3

3.1 Question

Cluster the blogs using K-Means, using k=5,10,20. (see slide 18).
How many interations were required for each value of k?

3.2 Answer

4 Question 4

4.1 Question

```
Use MDS to create a JPEG of the blogs similar to slide 29.
How many iterations were required?
```

4.2 Answer

5 Question 5

5.1 Question

Re-run question 2, but this time with proper TFIDF calculations instead of the hack discussed on slide 7 (p. 32). Use the same 500 words, but this time replace their frequency count with TFIDF scores as computed in assignment #3. Document the code, techniques, methods, etc. used to generate these TFIDF values. Upload the new data file to github.

Compare and contrast the resulting dendrogram with the dendrogram from question #2.

Note: ideally you would not reuse the same 500 terms and instead come up with TFIDF scores for all the terms and then choose the top 500 from that list, but I'm trying to limit the amount of work necessary.

5.2 Answer

6 Appendix A

```
#! /usr/bin/env python

import requests
import sys
from bs4 import BeautifulSoup

default = 'http://www.blogger.com/next-blog?navBar=true&blogID=3471633091411211117'
must_haves = ['http://f-measure.blogspot.com/', 'http://ws-dl.blogspot.com/']

def get_atom(uri):
    try:
        r = requests.get(uri)
    except Exception, e:
        return None
    soup = BeautifulSoup(r.text)
    links = soup.find_all('link', {'type':'application/atom+xml'})
    if links:
        return str(links[0]['href'])
    return None

def add_uri(uri, uris, outfile):
    if uri and uri not in uris:
        uris.add(uri)
        outfile.write(uri + '\n')
        print len(uris), uri

if __name__ == '__main__':
    uris = set()
    with open('blog_uris', 'a') as outfile:
        if len(sys.argv) > 1 and sys.argv[1] == 'new':
            for must_have in must_haves:
                uri = get_atom(must_have)
                add_uri(uri, uris, outfile)
        else:
            with open('blog_uris') as infile:
                [uris.add(line.strip()) for line in infile]
        while len(uris) < 100:
            uri = get_atom(default)
            add_uri(uri, uris, outfile)
```

Listing 14: get_uris.py

```
import feedparser
import futures
import md5
import re
import sys
import json

def get_next(d):
    for item in d.feed.links:
        if item['rel'] == u'next':
            return item['href']
    return None

def getwords(text):
    txt = re.compile(r'<[^>]+>').sub('', text)
    words = re.compile(r'[^A-Z^a-z]+').split(txt)
    return [word.lower() for word in words if word != '']

def get_titles(uri):
    print('processing {}'.format(uri))
    next = uri
    wc = {}
    pages = 0
    while next is not None:
        d = feedparser.parse(next)
        for e in d.entries:
            words = getwords(e.title.encode('utf-8'))
            for word in words:
                wc.setdefault(word, 0)
                wc[word] += 1
        pages += 1
        next = get_next(d)
        print('next {}'.format(next))
    title = d.feed.title.encode('utf-8')
    subtitle = d.feed.subtitle[:50].encode('utf-8')
    print('finished: {}: {}'.format(title, subtitle))
    return uri, title, subtitle, pages, wc

if __name__ == '__main__':
    with open('blog_uris') as infile:
        uris = [line.strip() for line in infile]
    if len(sys.argv) == 2 and sys.argv[1] == 'get':
        with futures.ThreadPoolExecutor(max_workers=8) as executor:
            uri_futures = [executor.submit(get_titles, uri) for uri in uris]
            for future in futures.as_completed(uri_futures):
                uri, title, subtitle, pages, wc = future.result()
                with open('wcs/' + md5.new(uri).hexdigest(), 'w') as out:
                    out.write(title + ': ' + subtitle + '\t' + str(pages) + '\t')
                    json.dump(wc, out)
    else:
        apcount = {}
        wordcounts = {}
        pagecounts = {}
        for uri in uris:
            with open('wcs/' + md5.new(uri).hexdigest()) as infile:
                try:
                    lines = infile.read().split('\t')
                    title = lines[0]
                    pages = int(lines[1])
                    wc = json.loads(lines[2])
                except Exception, e:
                    print('*** {} generated an exception: {}'.format(uri, e))
                    continue
            wordcounts[title] = wc
            pagecounts[title] = pages
            for word, count in wc.items():
                apcount.setdefault(word, 0)
                apcount[word] += count
        wordlist = []
        for w, bc in sorted(apcount.items(), key=lambda x: x[1], reverse=True):
            frac = float(bc) / len(uris)
            if frac > 0.1 and frac < 0.5:
                wordlist.append(w)
        if len(sys.argv) == 2 and sys.argv[1] == 'pages':
            with open('pagecounts', 'w') as outfile:
                outfile.write('blog\tpages\n')
```

```
77                  for blog, pagecount in pagecounts.iteritems():
78                      outfile.write("\"" + blog.replace("\"", "") + "\"" + '\t' + str(
                            pagecount) + '\n')
79          if len(sys.argv) == 2 and sys.argv[1] == 'wc':
80              with open('blogdata1.txt', 'w') as out:
81                  out.write('Blog')
82                  for word in wordlist[:500]:
83                      out.write('\t%s' % word)
84                  out.write('\n')
85                  for blog, wc in wordcounts.items():
86                      print blog
87                      out.write(blog)
88                      for word in wordlist[:500]:
89                          if word in wc:
90                              out.write('\t{}'.format(wc[word]))
91                          else: out.write('\t0')
92                      out.write('\n')
```

Listing 15: matrix.py

7 References

[1] Toby Segaran. *Programming Collective Intelligence*. O'Reilly, first edition, 2007.

[2] Internet Engineering Task Force (IETF). RFC-4287 The Atom Syndication Format. https://tools.ietf.org/html/rfc4287, 2005.