# DATASCI W261: Machine Learning at Scale

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### **Submission Notes:**

- For each problem, we've included a summary of the question as posed in the
  instructions. In many cases, we have not included the full text to keep the final
  submission as uncluttered as possible. For reference, we've included a link to the
  original instructions in the "Useful Reference" below.
- Some aspects of this notebook don't always render nicely into PDF form. In these situations, please reference the complete rendered notebook on Github (<a href="https://github.com/nickhamlin/mids-261-homework/blob/master/HW7/MIDS-W261-2015-HWK-Week07-Hamlin-Thomas-Baek-Danish.ipynb">https://github.com/nickhamlin/mids-261-homework/blob/master/HW7/MIDS-W261-2015-HWK-Week07-Hamlin-Thomas-Baek-Danish.ipynb</a>)

## **Useful References and Notebook Setup:**

Original Assignment Instructions
 (https://www.dropbox.com/s/26ejghkzgdidzwj/HW7-Questions.txt?dl=0)

```
#Use this to make sure we reload the MrJob code when we make changes %load_ext autoreload %autoreload 2
#Render matplotlib charts in notebook
%matplotlib inline

#Import some modules we know we'll use frequently
import numpy as np
import pylab as plt
```

The autoreload extension is already loaded. To reload it, use: %reload ext autoreload

69]: #Use this line of code to kick off a persistent cluster !python -m mrjob.tools.emr.create job flow '--conf-path' 'mrjob.conf'

creating new scratch bucket mrjob-a9a7d2100079455f using s3://mrjob-a9a7d2100079455f/tmp/ as our scratch dir on S3 Creating persistent job flow to run several jobs in... creating tmp directory /var/folders/rz/drh189k95919thyy3gs3tq400000g n/T/no script.nicholashamlin.20160312.024137.341853 writing master bootstrap script to /var/folders/rz/drh189k95919thyy3 gs3tq400000gn/T/no script.nicholashamlin.20160312.024137.341853/b.py creating S3 bucket 'mrjob-a9a7d2100079455f' to use as scratch space Copying non-input files into s3://mrjob-a9a7d2100079455f/tmp/no scri pt.nicholashamlin.20160312.024137.341853/files/ Waiting 5.0s for S3 eventual consistency Creating Elastic MapReduce job flow Can't access IAM API, trying default instance profile: EMR EC2 Defau ltRole Can't access IAM API, trying default service role: EMR DefaultRole Job flow created with ID: j-2VQKH4YTSZQSY j-2VQKH4YTSZQSY

### **HW** 7.0

#### HW 7.0 - Problem Statement

In this part of your assignment you will develop the base of your code for the week.

Write MRJob classes to find shortest path graph distances, as described in the lectures. In addition to finding the distances, your code should also output a distance-minimizing path between the source and target. Work locally for this part of the assignment, and use both of the undirected and directed toy networks.

To proof you code's function, run the following jobs

- shortest path in the undirected network from node 1 to node 4 Solution: 1,5,4
- shortest path in the directed network from node 1 to node 5 Solution: 1,2,4,5

and report your output---make sure it is correct!

### HW 7.0 - MR job definition

For brevity, we've combined the initialization step and the main job into a single unit of code. During the first pass, we check to make sure each node has been properly initialized and, if not, initialize it.

37]:		
•		

```
%%writefile MRbfs.py
from future import division
from mrjob.job import MRJob
from mrjob.job import MRStep
import re
import ast
WORD RE = re.compile(r''[\w']+")
class MRbfs(MRJob):
    def configure options(self):
        super(MRbfs, self).configure options()
        self.add_passthrough_option('--start', default='1', type=str)
        self.add passthrough option('--end', default='4', type=str)
        self.add_passthrough_option('--statuspath', default='', type=str)
        self.add passthrough option('--iteration', default='1', type=str)
        self.add_passthrough_option('--jobtype', default='local', type=str)
    def mapper(self, node, line):
        if self.options.iteration=='1':
            line = line.strip('\n')
            data = line.split("\t")
            status = 'U' #everything is unvisited initially
            # This is the first inialization (from original graph file)
            # there are only 2 elements, the node and adjacency list
            if(len(data) == 2):
                nid = data[0]
                N = ast.literal eval(data[1])
                \# if our node is the start node , initialize the start
                # distance at source = 0.0
                if nid == self.options.start:
                    ds = 0.0
                    status = 'V'
                    path=[nid]
                else:
                    # if this is not the start node, intialize distance to in.
                    ds = float("inf")
                    path=[]
                # yield the root node , (distance, graph structure, status, "
                yield nid, (ds, N, status,path)
                # nor each of the nodes in the adjacency list,
                # we expand frontier with starting distance.
                for m,d in N.iteritems():
                    #new distance is going to be from root node + dist
                    newdist = d+ds
                    new path=path[:]
                    new path.append(m)
```

```
if newdist < float("inf"):</pre>
                    status = 'Q'
                yield m, (newdist, None, status, new path)
    # from Iteration 1 onwards we'll land here...more data items to track
    else:
        if self.options.jobtype=='local':
            inf=float("inf") #THIS IS A TERRIBLE HACKJOB that lets us dea.
            line = line.strip('\n')
            nid,data=eval(line)
        if self.options.jobtype=='emr':
            Infinity=float('inf') #SEE HACKJOB ABOVE
            line = line.strip('\n')
            nid,data=line.split('\t')
            nid=eval(nid)
            data=eval(data)
        dist=data[0]
        N = data[1] # adjacency list - graph to use for next iter
        status = data[2] #status - U, Q, V
        path=data[3] #list of previously visited nodes
        #If a node is in the frontier, expand it
        if status=='Q':
            #Emit original node
            yield nid, (dist, N, 'V',path)
            #Emit linked nodes
            for m,d in N.iteritems():
                newdist=d+dist
                new path=path[:]
                new path.append(m)
                yield m, (newdist, None, 'Q', new path)
        #If it's not in the frontier, pass it through unchanged
        else:
            yield nid, (dist, N, status,path)
def reducer init(self):
    """We originally used these values for playing with
    alternative implementations of the stopping condition, but
    they're not really needed in the final version"""
    self.finished=0
    self.shortest path=[]
    self.min dist=0
def reducer(self, node, distances):
    adjList = {} #adjacency list - to be compiled below
    sdist = float("inf") #this is the shortest distance we've seen to the
```

```
stati = [] #list of all statuses we've encountered for this node
        spath=[] #shortest path we've found to the node so far
        for dist in distances:
            node dist,list of links,temp status,path=dist[:]
            # Extract the original graph structure from the node that has it
            if list of links:
                adjList = list of links
            stati.append(temp status)
            #If we find a record of a visited node, we can move on
            if temp status == 'V':
                sdist=node dist
                spath = path
                break
            # If we find a shorter distance to a node, update our knowledge o.
            if node dist < sdist:</pre>
                sdist = node_dist
                spath=path
                if node not in spath:
                    spath.append(node)
        #Check the current state of the node
        if 'V' in stati:
            status='V'
            #Have we hit our destination node?
            if node==self.options.end :
                self.shortest path=spath
                self.finished=1
                self.min dist=sdist
        elif 'Q' in stati:
            status='0'
        else:
            status='U'
        yield node, (sdist, adjList, status, spath)
    def steps(self):
        return [MRStep(
                        mapper=self.mapper
                        ,reducer init=self.reducer init
                        ,reducer=self.reducer
                      ) ]
if __name__ == '__main__':
    MRbfs.run()
```

Overwriting MRbfs.py

[6]:		

```
## HW7 - Undirected Toy Example, running locally
%reload ext autoreload
%autoreload 2
from MRbfs import MRbfs
from __future__ import division
iterate = 1
stop = False
last total dist = float("inf")
start node = ''
end_node = ''
end sdist = float("inf")
path = \{\}
#### TEST VERSION #####
input dir prefix='undirected toy'
start node='1'
end node='4'
#### RUN THE JOBS #####
#The second condition is insurance to make sure we stop after a finite number
while(not stop and iterate <= 20):</pre>
    output directory=input dir prefix+'Output{0}.txt'.format(str(iterate))
    if iterate==1:
        input directory=input dir prefix+'.txt'
    else:
        input_directory=input_dir_prefix+'Output{0}.txt'.format(str(iterate-1
    #LOCAL VERSION - IN WHICH WE WRITE RESULTS TO A FILE
    mr job = MRbfs(args=[input directory,
                         #'--file=start end.txt',
                         '--no-strict-protocols',
                         '--start', start_node,
                         '--end', end node,
                        #'--statuspath',status_path,
                         '--iteration', str(iterate)])
    with mr job.make runner() as runner:
        runner.run()
        print "Iteration : {0}".format(iterate)
        #Stream output locally
        with open(output directory, 'w+') as f:
            for line in runner.stream output():
                nid,distances = mr job.parse output line(line)
                dist,adjlist,status,path = distances
                f.write(str(mr_job.parse_output_line(line))+'\n')
                if nid==end node and status=='V':
```

```
print "DONE"
    print "Shortest path from {0} to {1} is {2} steps long:"."
    print "The path is: ",path
        stop=True
    if nid==end_node:
        break

#just in case this thing does not stop..
iterate += 1
```

```
Iteration : 1
Iteration : 2
Iteration : 3
DONE
Shortest path from 1 to 4 is 2 steps long:
The path is: ['1', '2', '4']
```

## **HW 7.0 - Directed Toy Test**

[5]:	_

```
## HW7 - Directed Toy Example, running locally
%reload ext autoreload
%autoreload 2
from MRbfs import MRbfs
from future import division
iterate = 1
stop = False
last total dist = float("inf")
start node = ''
end node = ''
end sdist = float("inf")
path = \{\}
#### TEST VERSION #####
input dir prefix='directed toy'
start node='1'
end node='5'
#### RUN THE JOBS #####
#The second condition is insurance to make sure we stop after a finite number
while(not stop and iterate <= 20):</pre>
    output directory=input dir prefix+'Output{0}.txt'.format(str(iterate))
    if iterate==1:
        input directory=input dir prefix+'.txt'
    else:
        input directory=input dir prefix+'Output{0}.txt'.format(str(iterate-1
    #LOCAL VERSION - IN WHICH WE WRITE RESULTS TO A FILE
    mr job = MRbfs(args=[input directory,
                         #'--file=start end.txt',
                         '--no-strict-protocols',
                         '--start', start node,
                         '--end', end node,
                        #'--statuspath',status path,
                        '--iteration', str(iterate)])
    with mr job.make runner() as runner:
        runner.run()
        print "Iteration : {0}".format(iterate)
        #Stream output locally
        with open(output directory, 'w+') as f:
            for line in runner.stream output():
                nid,distances = mr job.parse output line(line)
                dist,adjlist,status,path = distances
                f.write(str(mr_job.parse_output_line(line))+'\n')
                if nid==end node and status=='V':
                    print "DONE"
```

```
print "Shortest path from {0} to {1} is {2} steps long:"."
print "The path is: ",path
stop=True
if nid==end_node:
break

#just in case this thing does not stop..
iterate += 1
```

```
Iteration : 1
Iteration : 2
Iteration : 3
Iteration : 4
DONE
Shortest path from 1 to 5 is 3 steps long:
The path is: ['1', '2', '4', '5']
```

### **HW7.1**

### **HW 7.1 Problem Statement**

Using MRJob, explore the synonyms network data. Consider plotting the degree distribution (does it follow a power law?), and determine some of the key features, like:

number of nodes, number links, or the average degree (i.e., the average number of links per node), etc...

As you develop your code, please be sure to run it locally first (though on the whole dataset). Once you have gotten you code to run locally, deploy it on AWS as a systems test in preparation for our next dataset (which will require AWS).

### **HW 7.1 Calculating Summary Statistics**

Here, we take a simple pairs approach to count nodes and edges:

```
221: | %%writefile mrexplorenltk.py
     from mrjob.job import MRJob
     from mrjob.job import MRStep
     import csv
     from operator import itemgetter
     import re
     import ast
     class mrExploreNLTKTotals(MRJob):
         def mapper(self, _, line):
             """Emit keyless records (since we don't want to group our results)
             Values are (1, node degree)"""
             line = line.strip('\n')
             data = line.split("\t")
             nid = data[0]
             N = eval(data[1])
             node_degree = len(N)
             yield _,(1,node_degree)
         def reducer(self, nid, line):
              """Aggregate node counts and degree counts"""
             nodes=0
             edges=0
             for record in line:
                 nodes+=record[0]
                 edges+=record[1]
             yield None, (nodes,edges)
         def steps(self):
             return [MRStep( mapper=self.mapper
                          ,reducer=self.reducer
                      )
                  ]
     if __name__ == '__main__':
         mrExploreNLTKTotals.run()
```

Overwriting mrexplorenltk.py

```
681: #HW 7.1 - Calculating summary stats on NLTK data locally
     %reload ext autoreload
     %autoreload 2
     from future import division
     from mrexplorenltk import mrExploreNLTKTotals
     mr job = mrExploreNLTKTotals(args=['synNet.txt','--no-strict-protocols'])
     nodes=0
     edges=0
     with mr job.make runner() as runner:
         runner.run()
         for line in runner.stream output():
             #This performs the same function as the reducer, which is needed to co
             #final totals across a larger cluster. It's not really important for
             #but we'll need it for EMR
             ,count = mr job.parse_output_line(line)
             nodes+=count[0]
             edges+=count[1]
     print "Total Nodes = {}".format(nodes)
     print "Total Edges = {}".format(edges)
     print "Average Edges/Node = {}".format(edges/nodes)
     Total Nodes = 8271
```

Total Nodes = 82/1
Total Edges = 61134
Average Edges/Node = 7.39136742836

## HW 7.1 - Plotting the degree distribution

This second (very simple job) returns counts of nodes bucketed by how many degrees they have so that the distribution can be easily visualized.

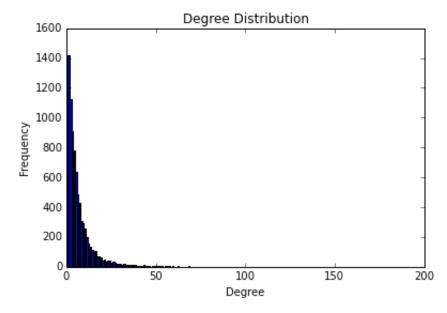
```
76]:
    %%writefile mrexplorenltkdist.py
     from mrjob.job import MRJob
     from mrjob.job import MRStep
     import csv
     from operator import itemgetter
     import re
     import ast
     class mrExploreNLTKDist(MRJob):
         def mapper(self, _, line):
             """Emit one record per node:
             Key: The node's degree (with padded zeros)
             Value: 1
             11 11 11
             line = line.strip('\n')
             data = line.split("\t")
             nid = data[0]
             N = eval(data[1])
             if N:
                 node degree = len(N)
             else:
                 node degree=0
             #Padding the degrees with leading zeros gives us results in order nice
             yield '{0:010d}'.format(node degree), 1
         def reducer(self, degree, count):
             """Aggregate pairs returned by the mapper"""
             total nodes=sum([i for i in count])
             yield degree, total nodes
         def steps(self):
             return [MRStep( mapper=self.mapper
                          ,reducer=self.reducer
                      )
                  ]
     if __name__ == '__main__':
         mrExploreNLTKDist.run()
```

Overwriting mrexplorenltkdist.py

```
77]: #HW 7.1 - Calculating degree distribution on NLTK data locally
     %reload ext autoreload
     %autoreload 2
     from future import division
     from mrexplorenltkdist import mrExploreNLTKDist
     input dir prefix='synNet'
     mr job = mrExploreNLTKDist(args=['{0}.txt'.format(input dir prefix),'--no-str
     nodes=0
     edges=0
     with open ('{0}_degree_distribution.txt'.format(input_dir_prefix),'w') as f:
         with mr_job.make_runner() as runner:
             runner.run()
             for line in runner.stream output():
                 #Write results to a file so we can plot them more easily
                 f.writelines(line)
     print "Done"
```

Done

```
78]:
     # HW 7.1 - Plot the results of the distribution
     x=[]
     y=[]
     with open ('{0} degree distribution.txt'.format(input dir prefix),'r') as f:
         for line in f.readlines():
             data=line.strip('\n').split('\t')
             degree=int(eval(data[0]))
             count=int(data[1])
             x.append(degree)
             y.append(count)
     plt.bar(x,y)
     plt.xlabel('Degree')
     plt.ylabel('Frequency')
     plt.title('Degree Distribution')
     plt.show()
```



As we'd expect, the distribution of degree does follow a power law; there are a handful of pages that are widely linked and lots of pages that have very few links.

## HW 7.1 - Testing on EMR with the same dataset

The only changes here are to the job parameters, but we've rerun everything to ensure that it will work properly in the cloud for 7.3 below.

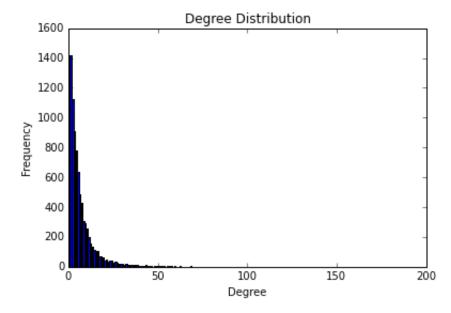
```
82]: #HW 7.1 - Calculating summary stats on NLTK data in EMR
     %reload ext autoreload
     %autoreload 2
     from future import division
     from mrexplorenltk import mrExploreNLTKTotals
     input dir prefix='synNet'
     input directory='s3://hamlin-mids-261/'+input dir prefix+'.txt'
     cluster='j-2VQKH4YTSZQSY'
     #EMR VERSION
     mr job = mrExploreNLTKTotals(args=['-r', 'emr',
                          input directory,
                           '--no-strict-protocols',
                           '--output-dir', 's3://hamlin-mids-261/'+input dir prefix+
                           '--no-output',
                           '--emr-job-flow-id', cluster,
                          ])
     nodes=0
     edges=0
     with mr job.make runner() as runner:
         runner.run()
         for line in runner.stream output():
             _,count = mr_job.parse_output_line(line)
             nodes+=count[0]
             edges+=count[1]
     print "Total Nodes = {}".format(nodes)
     print "Total Edges = {}".format(edges)
     print "Average Edges/Node = {}".format(edges/nodes)
```

```
Total Nodes = 8271
Total Edges = 61134
Average Edges/Node = 7.39136742836
```

```
80]: # HW 7.1 - Calculating degree distribution of NLTK data in EMR
     %reload ext autoreload
     %autoreload 2
     from future import division
     from mrexplorenltkdist import mrExploreNLTKDist
     input dir prefix='synNet'
     input directory='s3://hamlin-mids-261/'+input dir prefix+'.txt'
     cluster='j-2VQKH4YTSZQSY'
     #EMR VERSION
     mr_job = mrExploreNLTKDist(args=['-r', 'emr',
                          input directory,
                           '--no-strict-protocols',
                           '--output-dir', 's3://hamlin-mids-261/'+input dir prefix+
                           '--no-output',
                          '--emr-job-flow-id', cluster,
                         ])
     with open ('{0}_degree_distribution_EMR.txt'.format(input_dir_prefix),'w') as
         with mr_job.make_runner() as runner:
             runner.run()
             for line in runner.stream output():
                 #Write results to a file so we can plot them more easily
                 f.writelines(line)
     print "Done"
```

Done

```
81]:
     # HW 7.1 - Plot the results of the EMR-generated distribution
     x=[]
     y=[]
     with open ('{0} degree distribution.txt'.format(input dir prefix),'r') as f:
         for line in f.readlines():
             data=line.strip('\n').split('\t')
             degree=int(eval(data[0]))
             count=int(data[1])
             x.append(degree)
             y.append(count)
     plt.bar(x,y)
     plt.xlabel('Degree')
     plt.ylabel('Frequency')
     plt.title('Degree Distribution')
     plt.show()
```



## **HW7.2**

### **HW 7.2 Problem Statement**

Write (reuse your code from 7.0) an MRJob class to find shortest path graph distances, and apply it to the NLTK synonyms network dataset.

Proof your code's function by running the job:

• shortest path starting at "walk" (index=7827) and ending at "make" (index=536),

and showing you code's output. Once again, your output should include the path and the distance.

As you develop your code, please be sure to run it locally first (though on the whole dataset). Once you have gotten you code to run locally, deploy it on AWS as a systems test in preparation for our next dataset (which will require AWS).

## **HW 7.2 Implementation**

```
## HW7 - NLTK Example, running locally
%reload ext autoreload
%autoreload 2
from MRbfs import MRbfs
from __future__ import division
iterate = 1
stop = False
last total dist = float("inf")
start node = ''
end_node = ''
end sdist = float("inf")
path = \{\}
#### NLTK VERSION #####
input dir prefix='synNet'
start node='7827'
end node='536'
#### RUN THE JOBS #####
#The second condition is insurance to make sure we stop after a finite number
while(not stop and iterate <= 20):</pre>
    output directory=input dir prefix+'Output{0}.txt'.format(str(iterate))
    if iterate==1:
        input directory=input dir prefix+'.txt'
    else:
        input directory=input dir prefix+'Output{0}.txt'.format(str(iterate-1
    #LOCAL VERSION - IN WHICH WE WRITE RESULTS TO A FILE
    mr job = MRbfs(args=[input directory,
                         '--no-strict-protocols',
                         '--start', start node,
                         '--end', end node,
                         '--iteration',str(iterate)])
    with mr job.make runner() as runner:
        runner.run()
        print "Iteration : {0}".format(iterate)
        #Stream output locally
        with open(output directory, 'w+') as f:
            for line in runner.stream output():
                nid,distances = mr job.parse output line(line)
                dist,adjlist,status,path = distances
                f.write(str(mr job.parse output line(line))+'\n')
                if nid==end node and status=='V':
                    print "DONE"
                    print "Shortest path from {0} to {1} is {2} steps long:".:
                    print "The path is: ",path
                    stop=True
```

# if nid==end\_node: break

#just in case this thing does not stop..
iterate += 1

Iteration : 1
Iteration : 2
Iteration : 3
Iteration : 4

DONE

Shortest path from 7827 to 536 is 3 steps long: The path is: ['7827', '1426', '1668', '536']

44]:			
441.			

```
# NLTK example, running in EMR
%reload ext autoreload
%autoreload 2
from MRbfs import MRbfs
from __future__ import division
iterate = 1
stop = False
#PUT CLUSTER HERE!!!!!
cluster='j-2MM2A2KMPDLBH'
#### NLTK VERSION #####
input dir prefix='synNet'
start node='7827'
end node='536'
#### RUN THE JOBS #####
#The second condition is insurance to make sure we stop after a finite number
while(not stop and iterate <= 10):</pre>
    if iterate==1:
        input directory='s3://hamlin-mids-261/'+input dir prefix+'.txt'
        input_directory='s3://hamlin-mids-261/'+input dir prefix+'Output{0}/'
    #EMR VERSION
    mr job = MRbfs(args=['-r',
                          'emr',
                          input directory,
                          '--start', start_node,
                          '--end', end node,
                          '--no-strict-protocols',
                          '--output-dir', 's3://hamlin-mids-261/'+input dir pre:
                          '--no-output',
                          '--jobtype', 'emr',
                          '--emr-job-flow-id', cluster,
                          '--iteration', str(iterate)
                         ])
    with mr job.make runner() as runner:
        runner.run()
        print "Iteration : {0}".format(iterate)
        for line in runner.stream output():
            nid,distances = mr job.parse output line(line)
            dist,adjlist,status,path = distances
            if nid==end node and status=='V':
                print "DONE"
                print "Shortest path from {0} to {1} is {2} steps long: ".form
                print "The path is: ",path
                stop=True
            if nid==end node:
                break
```

#just in case this thing does not stop..
iterate += 1

```
Iteration : 1
Iteration : 2
Iteration : 3
Iteration : 4
DONE
Shortest path from 7827 to 536 is 3 steps long:
The path is: ['7827', '1426', '1688', '536']
```

-- - - ----

## **HW7.3**

### **HW 7.3 Problem Statement**

Using MRJob, explore the Wikipedia network data on the AWS cloud. Reuse your code from HW 7.1---does is scale well? Be cautioned that Wikipedia is a directed network, where links are not symmetric. So, even though a node may be linked to, it will not appear as a primary record itself if it has no out-links. This means that you may have to ADJUST your code (depending on its design). To be sure of your code's functionality in this context, run a systems test on the directed\_toy.txt network.

## **HW 7.3 Implementation**

The directed toy systems test is shown above in the results for 7.1, so for we have not repeated it here. Running our EDA code on the full dataset just requires updating the input data, so very little needs to change here.

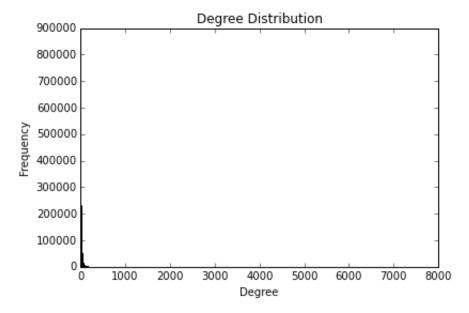
```
83]: | # HW 7.3 - Calculating summary stats on full Wikipedia dataset
     %reload ext autoreload
     %autoreload 2
     from future import division
     from mrexplorenltk import mrExploreNLTKTotals
     input dir prefix='all-pages-indexed-out'
     input directory='s3://hamlin-mids-261/'+input dir prefix+'.txt'
     cluster='j-2VQKH4YTSZQSY'
     #EMR VERSION
     mr_job = mrExploreNLTKTotals(args=['-r', 'emr',
                          input directory,
                           '--no-strict-protocols',
                           '--output-dir', 's3://hamlin-mids-261/WikiEDAOutput',
                           '--no-output',
                           '--emr-job-flow-id', cluster,
                         ])
     nodes=0
     edges=0
     with mr job.make runner() as runner:
         runner.run()
         for line in runner.stream_output():
             _,count = mr_job.parse_output_line(line)
             nodes+=count[0]
             edges+=count[1]
     print "Total Nodes = {}".format(nodes)
     print "Total Edges = {}".format(edges)
     print "Average Edges/Node = {}".format(edges/nodes)
```

```
Total Nodes = 5781290
Total Edges = 142114057
Average Edges/Node = 24.5817208616
```

```
841: # HW 7.3 - Calculating degree distribution on full Wikipedia dataset
     %reload ext autoreload
     %autoreload 2
     from future import division
     from mrexplorenltkdist import mrExploreNLTKDist
     input dir prefix='all-pages-indexed-out'
     input directory='s3://hamlin-mids-261/'+input dir prefix+'.txt'
     cluster='j-2VQKH4YTSZQSY'
     #EMR VERSION
     mr_job = mrExploreNLTKDist(args=['-r', 'emr',
                          input directory,
                           '--no-strict-protocols',
                           '--output-dir', 's3://hamlin-mids-261/WikiDistOutput',
                           '--no-output',
                           '--emr-job-flow-id', cluster,
                         ])
     with open ('{0} degree distribution EMR.txt'.format(input dir prefix), 'w') as
         with mr job.make runner() as runner:
             runner.run()
             for line in runner.stream output():
                 #Write results to a file so we can plot them more easily
                 f.writelines(line)
     print "Done"
```

Done

```
85]:
     # HW 7.3 - Plot the results of the distribution
     x=[]
     y=[]
     with open ('{0} degree distribution EMR.txt'.format(input dir prefix),'r') as
         for line in f.readlines():
             data=line.strip('\n').split('\t')
             degree=int(eval(data[0]))
             count=int(data[1])
             x.append(degree)
             y.append(count)
             #print degree, count
     plt.bar(x,y)
     plt.xlabel('Degree')
     plt.ylabel('Frequency')
     plt.title('Degree Distribution')
     plt.show()
```



Here, we see the general shape of the plot looks similar to what we saw in 7.1, but with an even more extreme power law curve. This is because we're dramatically increasing the size of the data, which means that the most linked pages will have even more links and there will be fewer of them relative to the rest of the population.

Initially our implementation in 7.1 did not scale particularly well and kept crashing our notebook. However, once we refactored the EDA process into two separate jobs we were able to optimize both of them independently. This version scaled much better, generating the results shown above in just a few minutes.

## **HW 7.4**

#### **HW 7.4 - Problem Statement**

Using MRJob, find shortest path graph distances in the Wikipedia network on the AWS cloud. Reuse your code from 7.2, but once again be warned of Wikipedia being a directed network. To be sure of your code's functionality in this context, run a systems test on the directed\_toy.txt network.

When running your code on the Wikipedia network, proof its function by running the job:

 shortest path from "Ireland" (index=6176135) to "University of California, Berkeley" (index=13466359),

and show your code's output.

### HW 7.4 - Running the directed toy example on EMR

26]:	1.	
20].	•	

```
# HW 7.4 - Directed toy example with FULL output, running on EMR
%reload ext autoreload
%autoreload 2
from MRbfs import MRbfs
from __future__ import division
iterate = 1
stop = False
#PUT CLUSTER HERE!!!!!
cluster='j-3AH5RMM6FM1X7'
#### TEST VERSION #####
input dir prefix='testgraph'
start node='1'
end node='5'
#### RUN THE JOBS #####
#The second condition is insurance to make sure we stop after a finite number
while(not stop and iterate <= 10):</pre>
    if iterate==1:
        input directory='s3://hamlin-mids-261/'+input dir prefix+'.txt'
        input directory='s3://hamlin-mids-261/'+input dir prefix+'Output{0}/'
    #EMR VERSION
    mr job = MRbfs(args=['-r',
                          'emr',
                          input directory,
                          '--start', start node,
                          '--end', end node,
                          '--no-strict-protocols',
                          '--output-dir', 's3://hamlin-mids-261/'+input dir pre:
                          '--no-output',
                          '--jobtype', 'emr',
                          '--emr-job-flow-id', cluster,
                          '--iteration', str(iterate)
                        ])
    with mr job.make runner() as runner:
        runner.run()
        print "Iteration : {0}".format(iterate)
        for line in runner.stream output():
            nid,distances = mr job.parse output line(line)
            dist,adjlist,status,path = distances
            print nid, status, path
            if nid==end node and status=='V':
                print "DONE"
                print "Shortest path from {0} to {1} is {2} steps long: ".form
                print "The path is: ",path
                stop=True
        print ""
```

#just in case this thing does not stop..
iterate += 1

```
Iteration: 1
1 V ['1']
2 Q ['1', '2']
3 U []
4 U []
5 U []
6 Q ['1', '6']
Iteration: 2
1 V ['1']
2 V ['1', '2']
3 Q ['1', '2', '3']
4 Q ['1', '2', '4']
5 U []
6 V ['1', '6']
Iteration : 3
1 V ['1']
2 V ['1', '2']
3 V ['1', '2', '3']
4 Q ['1', '2', '4']
5 Q ['1', '2', '4', '5']
6 V ['1', '6']
Iteration : 4
1 V ['1']
2 Q ['1', '2']
3 V ['1', '2', '3']
4 V ['1', '2', '4']
5 Q ['1', '2', '4', '5']
6 V ['1', '6']
Iteration: 5
1 V ['1']
2 Q ['1', '2']
3 Q ['1', '2', '3']
4 Q ['1', '2', '4']
5 V ['1', '2', '4', '5']
DONE
Shortest path from 1 to 5 is 3 steps long:
The path is: ['1', '2', '4', '5']
6 V ['1', '6']
```

Having confirmed that this works in the cloud, now let's run it on the entire dataset

## **HW 7.4 - Full-scale Implementation**

```
93]: # HW 7.4 - Load index into memory so we can look things up
word_dict={}
node_dict={}
with open('./wikipedia/indices.txt') as f:
    for line in f.readlines():
        word,node_id,_,_=line.strip().split('\t')
        node_dict[node_id]=word #Enables us to find words by ID
        word_dict[word]=node_id #Enables us to find IDs by word
```

06]:	
.001.	

```
# HW 7.4 - Full Wikipedia data, running in EMR
%reload ext autoreload
%autoreload 2
from MRbfs import MRbfs
from __future__ import division
iterate = 1
stop = False
#PUT CLUSTER HERE!!!!!
cluster='j-2VQKH4YTSZQSY'
#### WIKIPEDIA VERSION #####
input dir prefix='wiki'
start node='6176135'
end node='13466359'
#### RUN THE JOBS #####
#The second condition is insurance to make sure we stop after a finite number
while(not stop and iterate <= 10):</pre>
    if iterate==1:
        input directory='s3://hamlin-mids-261/all-pages-indexed-out.txt'
    else:
        input_directory='s3://hamlin-mids-261/'+input_dir_prefix+'Output{0}/'
    #EMR VERSION
    mr job = MRbfs(args=['-r',
                          'emr',
                         input directory,
                          '--start', start node,
                          '--end', end node,
                          '--no-strict-protocols',
                          '--output-dir', 's3://hamlin-mids-261/'+input dir pre:
                          '--no-output',
                          '--jobtype', 'emr',
                         #'--pool-emr-job-flows',#
                         #'--max-hours-idle', '1',#
                         #'--num-ec2-instances', '4',#
                         #'--ec2-instance-type', 'm3.xlarge',#
                          '--emr-job-flow-id', cluster,
                         '--iteration', str(iterate)
                         ])
    with mr job.make runner() as runner:
        runner.run()
        print "Iteration : {0}".format(iterate)
        for line in runner.stream output():
            nid,distances = mr_job.parse_output_line(line)
            dist,adjlist,status,path = distances
            #If we've hit our stopping condition, print results and finish up
```

```
if nid==end_node and status=='V':
    word_path=[node_dict[word]+' ('+word+')' for word in path]
    print "DONE"
    print "Shortest path from {0} to {1} is {2} steps long:".formation print "The path is: \n",' ->\n'.join(word_path)
    stop=True
    if nid==end_node:
        break

#Keep track of which iteration we're on
iterate += 1
```

```
Iteration : 1
Iteration : 2
Iteration : 3
DONE
Shortest path from Ireland to University of California, Berkeley is 2 steps long:
The path is:
Ireland (6176135) ->
Seamus Heaney (11607791) ->
University of California, Berkeley (13466359)
```

Sure enough, we find only two steps are required to traverse from "Ireland" (index=6176135) to "University of California, Berkeley" (index=13466359). One of the big performance quirks we noticed is the need to stream data from the S3 bucket on the west coast to our EMR cluster on the east coast. After realizing how long this was taking in the initial transfer, we found it to be more efficient to simply reupload the dataset to a bucket in the east to minimize the delay.

Now let's try running on another path, just to see what happens:

```
# HW 7.4- Lookup ID numbers for some random pages that sound interesting
print word_dict['Teenage Mutant Ninja Turtles']
print word_dict['Pi'] #March 14 is coming up...
```

12621471 10274977

11]:			
111.			

```
# HW 7.4 - Full Wikipedia data, running in EMR
%reload ext autoreload
%autoreload 2
from MRbfs import MRbfs
from __future__ import division
iterate = 1
stop = False
#PUT CLUSTER HERE!!!!!
cluster='j-2VQKH4YTSZQSY'
#### WIKIPEDIA VERSION #####
input dir prefix='wiki'
start node='12621471'
end node='10274977'
#### RUN THE JOBS #####
#The second condition is insurance to make sure we stop after a finite number
while(not stop and iterate <= 10):</pre>
    if iterate==1:
        input directory='s3://hamlin-mids-261/all-pages-indexed-out.txt'
        input_directory='s3://hamlin-mids-261/'+input_dir_prefix+'Output{0}/'
    #EMR VERSION
    mr job = MRbfs(args=['-r',
                         input directory,
                          '--start', start node,
                          '--end', end node,
                          '--no-strict-protocols',
                          '--output-dir', 's3://hamlin-mids-261/'+input dir pre:
                          '--no-output',
                          '--jobtype', 'emr',
                         #'--pool-emr-job-flows',#
                         #'--max-hours-idle', '1',#
                         #'--num-ec2-instances', '4',#
                         #'--ec2-instance-type', 'm3.xlarge',#
                          '--emr-job-flow-id', cluster,
                         '--iteration', str(iterate)
                         ])
    with mr job.make runner() as runner:
        runner.run()
        print "Iteration : {0}".format(iterate)
        for line in runner.stream output():
            nid,distances = mr_job.parse_output_line(line)
            dist,adjlist,status,path = distances
            #print nid, status, path
            if nid==end node and status=='V':
```

```
word_path=[node_dict[word]+' ('+word+')' for word in path]
print "DONE"
print "Shortest path from {0} to {1} is {2} steps long:".formate print "The path is: \n",' ->\n'.join(word_path)
stop=True
if nid==end_node:
    break

#just in case this thing does not stop..
iterate += 1
```

```
Iteration : 1
Iteration : 2
Iteration : 3
Iteration : 4
DONE
Shortest path from Teenage Mutant Ninja Turtles to Pi is 3 steps lon
g:
The path is:
Teenage Mutant Ninja Turtles (12621471) ->
Palladium Books (9937936) ->
Ancient Egypt (1110099) ->
Pi (10274977)
```

### **HW** 7.5

### **HW 7.5 Problem Statement**

Suppose you wanted to find the largest network distance from a single source,i.e., a node that is the furthest (but still reachable) from a single source. How would you implement this task? How is this different from finding the shortest path graph distances? Is this task more difficult to implement than the shortest path distance? As you respond, please comment on program structure, runtimes, iterations, general system requirements, etc...

## HW 7.5 Response

We'd find the largest path using a similar, but not exactly the same, approach that we're using here to find the shortest path. The key difference is that we know in advance that we'll need to thoroughly traverse the entire graph to know that we've found the longest path. In our shortest path implementation, we stop when we've found the shortest path to the node we're interested in. This meant that the task isn't necessarily more "difficult" from a logical perspective, but it may be more computationally intensive as it would likely require more iterations to arrive at a conclusion. Since we know we'll need to run several more iterations (and can't take advantage of the "six-degrees-of-separation" effect that we can when we look for the shortest path), optimizing both our mapreduce implementation and our cluster

infrastructure for rapid turnover is even more important when searching for the longest path. It shouldn't matter whether we use a depth first search or a breadth first search, since we'll still need to visit every node regardless.

### HW 7.5.1:

Can we utilize combiners in the HW 7 to perform the shortest path implementation?

We could probably use combiners to handle some of the logic that's currently addressed in the reducer around deciding what state a node is in. This might entail minimizing network throughput by using the combiner to decide not to emit records to the reducer nodes that have already been visited or to do some precalculation of the shortest path to a given node based only on the records available to that particular mapper. These "subtotals" could then be reconciled in the reducer.

Does order inversion help with the HW 7 shortest path implementation?

We might be able to use order inversion to simplify some of the logic in the reducer by guaranteeing, for example, that the reducer would always receive the original record of a node before records corresponding to expanded links. That said, while this might make the code easier to understand, it may not lead to a significant boost in performance.

### **End of Submission**