## Lecture 12 Seriation/Clustering

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#### Introduction

- We haven't done much with clustering in Data Mining 2.
- In Data Mining 1 we talked about k-means clustering and hierarchical clustering
- Tonight we'll talk about seriation.
- Seriation can be used with clustering to clean up the results or if we're lucky...
  - It can be used by itself for clustering
- Go to <a href="http://www.atgc-montpellier.fr/permutmatrix/manual/SeriationCorps.htm">http://www.atgc-montpellier.fr/permutmatrix/manual/SeriationCorps.htm</a>

### Simple Example Stolen from Previous Website

#### Simple example n°2

- Step 1 The smallest dissimilarity in D is 2,58.

  This value is already near the diagonal.

  No movement is necessary at this step.
- Step 2 The dissimilarity between items 1 and 2 is now the smallest dissimilarity available.

  Dmin = 2,61.

  To put this value against the diagonal,

we swap items 2 and 5.

- Step 3 At this step, the dissimilarity between items 4 and 5 is the currently smallest dissimilarity. Items 4 and 3 were pooled at the first step and we are not allowed to modify this pair. We must put item 5 between items 4 and 1. At the end of this step, items 3, 4 and 5 form a "fragment" of three elements.
- Step 4 For the final step, the dissimilarity between items 2 and 3 is the smallest dissimilarity available. To put this value against the diagonal, we swap fragments {1,2} and {3,4,5}.
- End At the end of this algorithm, small dissimilarities are spread along the main diagonal of D.

  Close elements are close in the seriation.

#### Dissimilaritiy matrix D

D	3	4	1	5	2
3	0,00				
4	2,58	00,00			
1	5,33	7,90	0,00		
5	5,22	2,65	10,55	0,00	
2	2,73	5,29	2,61	7,94	0,00

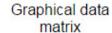
_	_			_	
D	3	4	1	5	2
3	0,00				
4	2,58	0,00			
1	5,33	7,90	00,00		
5	5,22	2,65	10,55	0,00	
2	2.73	5.29	2.61	7.94	0.00

D	3	4	1	2	5
3	00,00				
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1	5,33	7,90	0,00		
2	2,73	5,29	2,61	00,00	
5	5,22	2,65	10,55	7,94	00,00

D	3	4	5	1	2
3	0,00				
4	2,58	0,00			
5	5,22	2,65	0,00		
1	5,33	7,90	10,55	0,00	
2	2,73	5,29	7,94	2,61	00,0

D	1	2	3	4	5
1	00,00				
2	2,61	0,00			
3	5,33	2,73	0,00		
4	7,90	5,29	2,58	0,00	
5	10,55	7,94	5,22	2,65	00,00

#### Graphical representation of D

















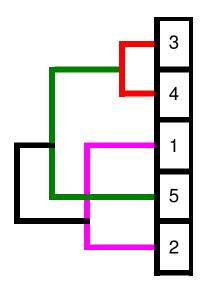


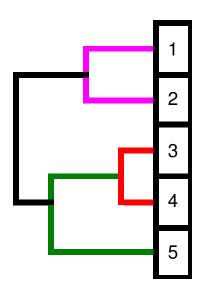




#### Hierarchical Clusters from Before and After Seriation

- Note seriation (from previous page) cleans up the hierarchy
- In the non-seriated hierarchy we have the green cluster crossing the pink cluster
- Note also that things that belong together tend to be next to each other even without the clustering
- That is, 1 is next to 2, 3 is next to 4, (3,4) is next to 5 etc
- If we plotted this as a heatmap, after seriation we could see where we should make our clusters without running a clustering algorithm
- This will become more obvious when we do our real example





#### The Setup...

- Last semester in Text Mining, Chris, John, Matt, Mihir, Parvati, Yousuf and others created a corpus of Linked-in docs, web pages, press releases, google scholar articles etc. on a topic of their choice
- They then removed stopwords, lemmatized, etc. their corpus
- After the semester I put all 1,292 documents into one giant corpus to test whether k-means clustering would be smart enough to separate each set into it's proper topic
- Results are on the next page

#### **Results**

	Cluster #		False Positive (Doesn't Belong)	False Negative (Should be in)	F Score
Chris	9	191	40	0	0.905
Eric	3	153	12	0	0.962
John	0	111	8	0	0.965
Matt	4	78	11	19	0.839
Mihir	7	97	31	2	0.855
Parvati	1	43	2	190	0.309
Stephen	5	98	34	1	0.848
Tim	6	100	12	0	0.943
Tony	2	146	16	0	0.948
Yousuf	8	63	46	0	0.733
all	all	1080	212	212	0.836

	word1	word2	word3	word4	word5	word6	word7	word8	word9	word10
Cluster 0	limb	prosthetic	advanced	arm	artificial	prosthesis	hand	amputee	control	darpa
Cluster 1	biotechnology	university	verified	game	email	edu	video	information	engineering	science
Cluster 2	text	mining	information	document	analysis	literature	gene	tool	biomedical	drug
Cluster 3	trump	donald	clinton	woman	debate	campaign	republican	presidential	president	people
Cluster 4	biophysics	chemistry	journal	biophysical	physical	molecular	et	cell	al	patent
Cluster 5	quantum	compute	computer	physic	qubits	university	institute	science	state	waterloo
Cluster 6	airplane	wing	plane	flight	fly	mode	design	pilot	aircraft	jan
Cluster 7	machine	leaming	learn	proceeding	international	conference	cloud	google	api	workshop
Cluster 8	nanotechnology	nature	nanoscience	ibm	ieee	nanomaterials	tj	nano	center	watson
Cluster 9	game	video	play	player	violent	study	time	world	child	gaming

#### Results (II)

- Note that k-means does the best on cluster 0 (prosthetics) because the topic has nothing to do with the others
- It also does well on the politics, airplane, and gaming clusters
- It inexplicably puts gaming in the biotechnology cluster though
- Overall it does a decent job, and we could make it better by making things like email, et al, jan, edu, etc. into stopwords

#### **Seriation of Documents**

- The question is whether we could do a decent job of clustering the documents with seriation?
- How?
- Take entire corpus; do TF/IDF to get a list of 1000 or 2000 most important words
- (A human edited list would work better, but we don't have time)
- Each document is then a 2000 dimension sparse vector with term frequencies
- Seriation puts closest vectors together which ought to work as a clustering method

### **Step 1: Read in Corpus**

```
In [174]: from sklearn.datasets import load files
     ...: ## folder containing 1292 lemmatized text files (stopwords removed)
     ...: ## about 100 files each under names Chris, Parvati, etc.
     ...: p="E:\Rowan\Classes\TextMining\hw3CaseStudy\LemmatizeWordContainer"
     ...: ## Load all files with one command
     ...: dataset=load files(p, description=None, categories=None,
     ...: load content=True, shuffle=True, encoding=None,
     ...: decode error='strict', random state=0)
     . . . :
     ...: print("%d documents" % len(dataset.data))
     ...: print("%d categories" % len(dataset.target names))
     ...: print(dataset.target names[0:9])
1292 documents
10 categories
['Chris', 'Eric', 'John', 'Matt', 'Mihir', 'Parvati', 'Stephen', 'Tim', 'Tony']
```

 A great thing about sklearn is that if you keep your corpus as a bunch of text files in a hierarchical directory, load\_files will load all of the files and keep the directory structure; all in one line

### Step 2: Make List of all words; sort by freq.

```
In [175]: wordDict = {}
     ...: for i in range(len(dataset.data)):
     ...: words = dataset.data[i].split()
     ...: for word in words:
                 if wordDict.has key(word):
                      wordDict[word]+=1
                 else:
                      wordDict.setdefault(word,1)
     ...: import operator
     ...: wordDist = [sorted(wordDict.iteritems(), key=operator.itemgetter(1),reverse=True)]
     ...: for s in wordDist:
           for (v,k) in s:
                if (k > 1000):
                      print(v+" "+str(k))
     . . . :
game 4623
trump 3727
video 1904
penn 1540
anonymous 1268
time 1190
quantum 1052
airplane 1004
```

- A Python dictionary is useful for keeping track of the occurrences of each word
- Don't know why 'penn' and 'anonymous' appear so frequently.

### Step 3: Make List of Top 2000 Words

```
...: ## kill additional stopwords; we could go crazy, but we'll just kill these 2.
In [176]: wordDict['penn']=0
...: wordDict['anonymous']=0
...:
...:
...: ## put top 2000 words in a list
...: ## we won't bother doing IDF
...: count=0
...: list=[]
...: for s in wordDist:
...: for (v,k) in s:
...: if (count <2000):
...: list.append(v)
...: count+=1</pre>
```

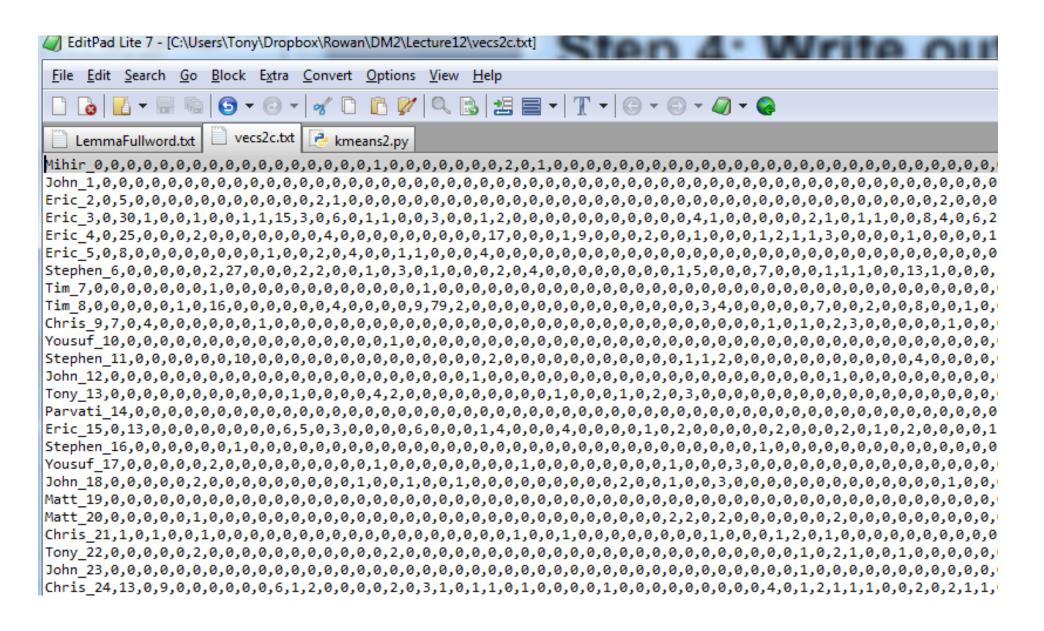
- 2000 is just arbitrary. I don't know if it would work better with 100, 500, 1000, 10000.
- Note also, the first word and 2000<sup>th</sup> word have the same weight here. Don't know if this will be a problem

## Step 4: Write out each document as a sparse vector

```
In [177]: f5=open('vecs2c.txt','w')
     ...: papDict = {}
     ...: for i in range(len(dataset.data)):
            ## build a dictionary for each paper
            words = dataset.data[i].split()
            for word in words:
                if papDict.has key(word):
                      papDict[word]+=1
                else:
                      papDict.setdefault(word,1)
            ## now build a vector containing frequencies
            vecc=[]
            for w in list:
                if papDict.has key(w):
                      vecc.append(papDict[w])
                else:
                      vecc.append(0)
            ## print vector out
            f5.write(dataset.target_names[labels[i]]+"_"+str(i))
            for t in vecc:
                f5.write(","+str(t))
                f5.write("\n")
            papDict={}
     ...: f5.close()
```

 Use a dictionary for each document. Then check to see if any words from the document match the top 2000 words

# Snapshot of output (Total of 1292 vectors; each has 2000 elements)



#### Go to R

Go to Seriation2000.pdf

#### Unrelated Seriation Demo in R

This one has nothing to do with clustering.

 But it might be the coolest thing you've seen in a while

Go to SeriationLouvre.pdf