# Use Clustering to Solve a Mystery in History

### Introduction:

In the late 1780's, to address the U.S Constitution still needing approval from nine of the thirteen states, The Federalist Papers, a series of 85-letters, were written in newspapers urging the ratification of the U.S Constitution. These letters contended that the US Constitution would preserve the Union and allow for the proposed federal government to act in the nations interest. The Federalist Papers have long been regarded as an extremely important source of evidence of the underlying meaning and intentions of the Constitution, as well. In fact, in 1821, in *Cohens v Virginia* Chief Justice John Marshall described the papers as the following:

"complete commentary on our constitution; and its appealed to by all parties in the questions to which that instrument has given birth. Its intrinsic merit entitles it to this high rank, and the part two of its authors performed in framing the constitution, put it very much in their power to explain the views with which it was framed."

The predicament is, for a document with this caliber of importance in creating and ratifying our country's' Constitution, we do not know completely who wrote all the essays that make up the Federalist Papers. In fact, 11 essays are still disputed about regarding whether Madison or Hamilton wrote them. By using the K-Means, EM, and HAC clustering algorithms, we hope to draw a conclusion on who wrote the disputed essays by demonstrating what patterns were learned to predict the disputed papers.

# Analysis:

Data Preparation:

Data Preparation first required loading the .csv file. A general assessment of the file was done using str(data) (Figure 1). We then looked for complete cases by returning a logical vector that indicated which cases are complete. It showed that we have no missing value (Figure 2)

```
str(data)
'data.frame':
                85 obs. of 72 variables:
$ author : Factor w/ 5 levels "dispt","Hamilton",..: 1 1 1 1 1 1 1 1 1 1 ...
$ filename: Factor w/ 85 levels "dispt_fed_49.txt",..: 1 2 3 4 5 6 7 8 9 10 ...
$ a
           : num    0.28    0.177    0.339    0.27    0.303    0.245    0.349    0.414    0.248    0.442    ...
$ all
           : num 0.052 0.063 0.09 0.024 0.054 0.059 0.036 0.083 0.04 0.062 ...
$ also
           : num 0.009 0.013 0.008 0.016 0.027 0.007 0.007 0.009 0.007 0.006 ...
           : num 0.096 0.038 0.03 0.024 0.034 0.067 0.029 0.018 0.04 0.075 ...
$ an
           : num    0.358    0.393    0.301    0.262    0.404    0.282    0.335    0.478    0.356    0.423    ...
$ and
$ any
           : num 0.026 0.063 0.008 0.056 0.04 0.052 0.058 0.046 0.034 0.037 ...
$ are
           : num 0.131 0.051 0.068 0.064 0.128 0.111 0.087 0.11 0.154 0.093 ...
           : num 0.122 0.139 0.203 0.111 0.148 0.252 0.073 0.074 0.161 0.1 ...
$ as
           : num 0.017 0.114 0.023 0.056 0.013 0.015 0.116 0.037 0.047 0.031 ...
$ at
           : num    0.411    0.393    0.474    0.365    0.344    0.297    0.378    0.331    0.289    0.379    ...
$ be
           : num 0.026 0.165 0.015 0.127 0.047 0.03 0.044 0.046 0.027 0.025 ...
$ been
           : num 0.009 0 0.038 0.032 0.061 0.037 0.007 0.055 0.027 0.037 ...
$ but
           : num 0.14 0.139 0.173 0.167 0.209 0.186 0.102 0.092 0.168 0.174 ...
  by
                  0.035 0 0.023 0.056 0.088 0 0.058 0.037 0.047 0.056
```

Figure 1

```
> View(data)
> sum(!complete.cases(data))
[1] 0
```

Figure 2

We then removed the first two columns (author and filename) in order to leave the data frame as just function words and feature values (Figure 3).

```
fp<-data[,c(-1,-2)]
  fp
           all also
       а
                        an
                             and
                                   any
                                                       at
                                                                 been
                                                                        but
                                                                               by
                                          are
   0.280 0.052 0.009 0.096 0.358 0.026 0.131 0.122 0.017 0.411 0.026 0.009 0.140 0
   0.177 0.063 0.013 0.038 0.393 0.063 0.051 0.139 0.114 0.393 0.165 0.000 0.139 0
3
3
   0.339 0.090 0.008 0.030 0.301 0.008 0.068 0.203 0.023 0.474 0.015 0.038 0.173 (
   0.270 0.024 0.016 0.024 0.262 0.056 0.064 0.111 0.056 0.365 0.127 0.032 0.167 0
0
5
   0.303 0.054 0.027 0.034 0.404 0.040 0.128 0.148 0.013 0.344 0.047 0.061 0.209 (
6
   0.245 0.059 0.007 0.067 0.282 0.052 0.111 0.252 0.015 0.297 0.030 0.037 0.186 (
0
   0.349 0.036 0.007 0.029 0.335 0.058 0.087 0.073 0.116 0.378 0.044 0.007 0.102
```

Figure 3

### Processing:

The first step of processing required scaling the data in order to standardize the range of features in the data (Figure 4). By scaling, we can account for any wide range values. We also started preparing for cluster analysis by using the elbow graphing method to figure out the most optimal number of clusters with totalwithinss. We chose to set our seed to 10 and created a function to return the optimal number of clusters. We chose our nstart to be 25, meaning it will generate 25 initial configurations. And then plotted the total within-clusters sum of squares (will be discussed in results).

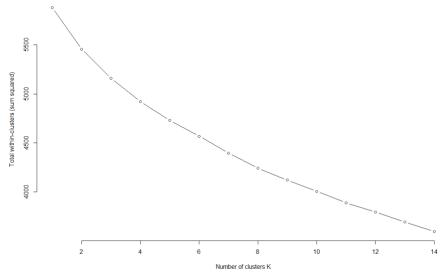


Figure 4

There were two ways of processing the data for K\_means. The first involved computing the K-means clustering in R without specifying the Hartigan-Wong method. The second involves using the Hartingan-Wong method to find the centroid.

In the first method, we run the code shown in figure 5. The cluster is set to four and nstart is at 25. The output is shown in results after it was specified to show centers.

```
#### Option 1: K means_Clustering ####
k_mean<-kmeans(fp2, centers = 4, nstart = 25)
k_mean
k_mean$centers</pre>
```

Figure 5

In the second method, we used the following two references to better understand the Hartigan-Wong method: <a href="https://rstudio-pubs-">https://rstudio-pubs-</a>

static.s3.amazonaws.com/79267\_cf36e5130fa449bb876ee563908c7a27.html and https://en.wikibooks.org/wiki/Data\_Mining\_Algorithms\_In\_R/Clustering/K-Means

The following line was executed to find the centroid. The results will be discussed later. But, we set our cluster to 4, nstart was 25, and max iterations was set to 100. We specified the algorithm was Hartigan-Wong, and then filtered for results with the center value.

```
######Option 2: :K_means using Hartingan-Wong Method###

##HW algorithm to find centroid

km_output1<-kmeans(fp2,centers=4,nstart = 25,iter.max = 100,algorithm = "Hartigan-Wong")

km_output1$centers
```

Next, we used the HAC clustering algorithm noted below and plotted the results. HAC stands for Hierarchical Cluster Analysis and it is an alternative to K-means clustering for identifying groups in the dataset. HAC does not require a pre-specification of clusters, like K\_means did. Additionally, the HAC output is a dendrogram which displays the groups in a tree-shaped plot. The plot can be found in the results section. We set our distance measure to Euclidean and method as complete linkage clustering.

```
# 2nd Algorithm: HAC
hac_output<-hclust(dist(fp2,method = "euclidean"),method = "complete")

#plot HAC
plot(hac_output)

#output desirable number of clusters after modeling
hac_cut<-cutree(hac_output,4)
hac_df1<-data.frame(author=data$author,cluster=hac_cut)
clusplot(fp2,hac_cut,color=TRUE,shade=T,labels = 2,lines =0)</pre>
```

For the third algorithm, we used EM, or Expectation-Maximizing. It is an unsupervised learning algorithm that iteratively finds the maximum likelihood of parameters in a model that depends on unobserved latent variables. We will use the mclust function in R to provide the parameters.

```
#3rd Algorithm: EM
install.packages("mclust")
library(mclust)

#visualize clusters
clPairs(fp2,data$author)
fit<-Mclust(fp2)
summary(fit)</pre>
```

### Results:

**K\_Means** 

Below, the results for both option 1 and 2 are shown (options discussed above). The non-filtered results of both options can be seen in Appendix A. This result was shown to find the clustering sizes. Additionally, it was used to confirm both methods were identical in results. The results shown in Appendix A will confirm this. The four clusters were found to have the following cluster sizes: 4,11,24,46.

The results below show the K\_means for when the results are filter for center. In addition, we plotted the centers to better visually analyze the data (Figure 8 & 9).

Option 1 Results:

Option 1 Results:
> km_output1\$centers
a all also an and any
1 0.28962572 -0.008668348 -0.33688781 0.4564520 -0.4292426 0.31853756 2 -0.57598787 0.176223940 -0.07835166 -0.3045733 0.7401337 -0.88866997
3 -0.06274336
4 -1.37026902 -0.617157145 1.58662115 -1.3968214 2.0034108 0.06092825
are as at be been but 1 -0.1269924 -0.1157504 0.1638521 0.1704703 0.07424644 -0.04491703
1 -0.1269924 -0.1157504 0.1638521 0.1704703 0.07424644 -0.04491703 2 -0.2474613 -0.3541919 0.5272837 -1.3393966 0.27336121 -0.64897176
2 -0.24,4019137 0.3655169 -0.3934412 0.2648322 -0.05293925 0.29481729 4 -0.2705511 0.1120563 -0.9736815 0.1339390 -1.28794190 0.53231438
by can do down even every
1 -0.4488847
3 0.6611626 0.07030163 -0.02037322 -0.0890032 -0.07851415 0.6725072
2 0.4744678 -0.88893879 -0.67604087 -0.4239542 -0.39970839 -0.6189158 3 0.6611626 0.07030163 -0.02037322 -0.0890032 -0.07851415 0.6725072 4 -0.1095873 0.27450777 0.68373426 -0.4239542 -0.36977631 -0.68934921
tor. from had has have her
1 0.02154164 0.071958773 -0.2033678 0.16005265 0.0596352 -0.19775612
2 -0.06056271 0.008195461 1.6808436 0.01947992 0.4388495 0.02101943 3 0.06123478 -0.286643438 -0.2738688 -0.16743504 -0.1427725 -0.00244272
3 0.06123478 -0.286643438 -0.2738688 -0.16743504 -0.1427725 -0.00244272 4 -0.44859010 0.869797219 -0.6403774 -0.88956500 -1.0360062 2.23104823
his if. in. into is it
1 0 1014028 0 09127669 0 4786243 -0 284284711 0 1789476 0 2045855
2 0.2055717 -0.41580531 -0.5515928 0.392435886 -0.8577637 -0.7161222 3 -0.2136100 -0.27657201 -0.4937745 -0.009233076 0.2396741 -0.1122614 4 -0.4497950 1.75321470 -1.0246517 2.245473953 -1.1370926 0.2901717
3 -0.2136100 -0.27657201 -0.4937745 -0.009233076 0.2396741 -0.1122614 4 -0.4497950 1.75321470 -1.0246517 2.245473953 -1.1370926 0.2901717
4 -0.4497950 1.75321470 -1.0246517 2.245473953 -1.1370926 0.2901717 its may more must my no
1 0.2352606 0.06920864 -0.2880071 0.09850013 0.15721793 0.06580667
2 -0.3206233 -1.11663332 -0.1067859 -0.77537462 -0.02090334 -0.77992250
3 -0.2473462 0.31909596 0.1948770 0.25975274 -0.22413956 0.32764604
4 -0.3397059 0.36026651 2.4364815 -0.55898773 -0.40568466 -0.57786607
not now of on one only 1 -0.04176153  0.1049530  0.3506382 -0.5643201 -0.31726578 -0.07651245
2 -0.58955133  0.3780618 -0.2435579  0.2131969  0.09429220 -0.62459918
3 0.29793854 -0.2331719 -0.2382584 1.0056141 0.07239926 0.38610287
4 0.31303232 0.0473373 1.3330044 0.1302333 2.33403737 0.20032370
or our shall should so some 1 0.05350945 0.01390439 0.1675906 0.2178890 -0.05937977 -0.2618477
2 -0.38365788 -0.26303984 -0.3792315 -0.2724530 -0.08521280 0.6392920 3 -0.19677510 -0.08883363 -0.2231519 -0.2394768 0.06804264 0.2661959 4 1.62035105 1.09646083 0.4545054 -0.3196166 0.50894675 -0.3439797
1 1102033103 1103010003 011313031 013130100 0130031013 013133131
such than that the their
1 0.07150055 -0.02059864 0.2084314394 0.1159819 -0.273306977 2 -0.52709523 -0.50106422 -0.5238465684 -0.4025561 0.991849610
2 -0.52709523 -0.50106422 -0.5238465684 -0.4025561 0.991849610 3 -0.02595886 -0.09047437 -0.1592967827 0.3611036 0.008301811 4 0.78300868 2 15765712 -0.0006027944 -2.3933839 0.365633943
then there things this to up
1 -0.07793873
2 -0.04346848 -0.6560934 -0.42002725 -0.3731165 -0.9652637 0.03601269 3 0.12357132 -0.6633081 -0.09329193 -0.1353435 -0.7179521 -0.34049549
3 0.12357132 -0.6633081 -0.09329193 -0.1353435 -0.7179521 -0.34049549 4 0.27440577 0.4137506 -0.60673314 -1.7833116 -0.1442837 0.03935447
upon was were what when which
1 0.6879288 -0.23977418 -0.22843317 0.1146205 0.05561194 0.03705055
2 -0.3728410 1.38567795 1.19163527 -0.3776256 -0.24797150 -0.15054096
3 -1.0156057 -0.08612244 -0.11254063 -0.1307965 -0.13136922 0.15286819 4 -0.7922336 -0.53647667 0.02522827 0.5051132 0.83059962 -0.92930284
who will with would your
1 0.03114336 -0.05828478 -0.11207025 0.2473179 0.02897163
2 0.27465588 -0.84541229 0.17875034 -0.3482766 -0.20876463
3 -0.16193522
4 -0.14184107 0.72202330 1.34607296 1.1361017 0.61638391 > k_mean\$centers
a all also an and any
1 -0.06274336  0.038704552  0.41717596 -0.5024666  0.1495853 -0.21337796
Ti control of the con

Figure 6

Option 2 Results:

```
W algorithm to find centroid
output1<-kmeans(fp2,centers=4,nstart = 25,iter.max = 100,algorithm = "Ha
                       -0.5364/66/ 0.0252282/ 0.5051132

with would

-0.05828478 -0.11207025 0.2473179

-0.84541229 0.17875034 -0.5037495

0.37885591 -0.09147142 -0.5037495

0.72202330 1.34607296 1.1361017
```

Figure 7

# CLUSPLOT(fp2) COMPONENT 1 COMPONENT 1 COMPONENT 1 COMPONENT 1

Figure 8

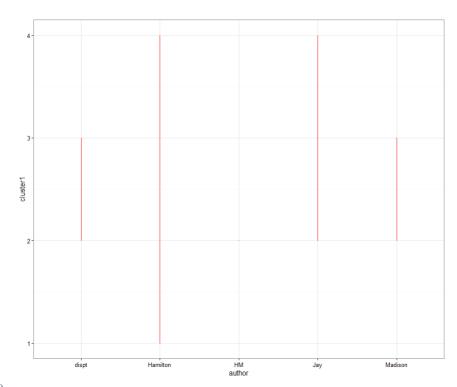


Figure 9

We can use the Figure 9-line plot to see that the disputed authors are divided into clusters 2 and 3. So when compared to the lines for Hamilton and Madison, it is clear that the disputed author line is most similar to Madison. Therefore, it is more likely that the disputed author papers are from Madison.

### HAC Algorithm:

In the second algorithm, HAC, we initially tried two different methods. First, we set the method to average and got the results shown in figure 10 and 11. In these outputs, we saw that most of the data was divided into one cluster. This is why we chose to use the complete agglomeration method.

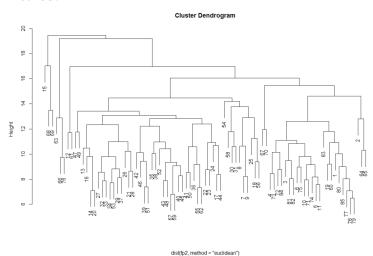
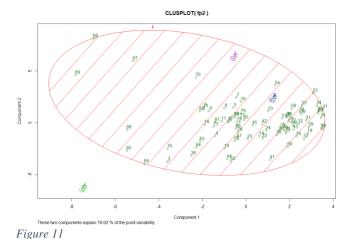


Figure 10



In figure 12 and 13, when we set the agglomeration method to complete, we could see that almost all the disputed papers (1-11) are clustered [located] with Madison's papers (which are 71-85). This confirms the results of the K\_means clustering.

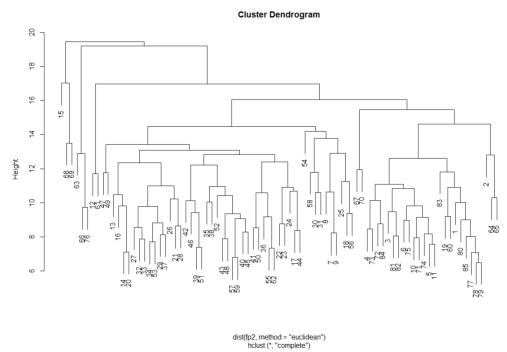


Figure 12

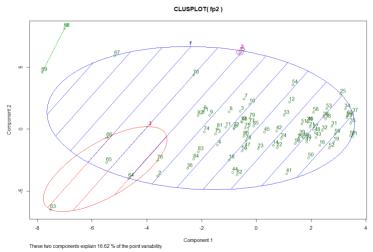


Figure 13

# Expectation Maximum (EM) Algorithm:

We used the following resources to understand EM: <a href="http://rstudio-pubs-static.s3.amazonaws.com/154174\_78c021bc71ab42f8add0b2966938a3b8.html">http://rstudio-pubs-static.s3.amazonaws.com/154174\_78c021bc71ab42f8add0b2966938a3b8.html</a> and <a href="https://rdrr.io/cran/mclust/man/mclustBIC.html">https://rdrr.io/cran/mclust/man/mclustBIC.html</a>

For the third algorithm, we first installed the mclust library in order to first visualize the clusters (Figure 14). We used the Gaussian finite mixture model, which was fitted by the EM algorithm,

```
####3rd Algorithm: EM####
install.packages("mclust")
library(mclust)

#visualize clusters
clPairs(fp2,data$author)
fit<-Mclust(fp2)
summary(fit)</pre>
```

Figure 14

Since we first wanted to visualize the datapoints. We used the clPairs to see the spectrum for the first 30 columns. We mainly could not increase the number of columns due to the size restraints (Figure 15).

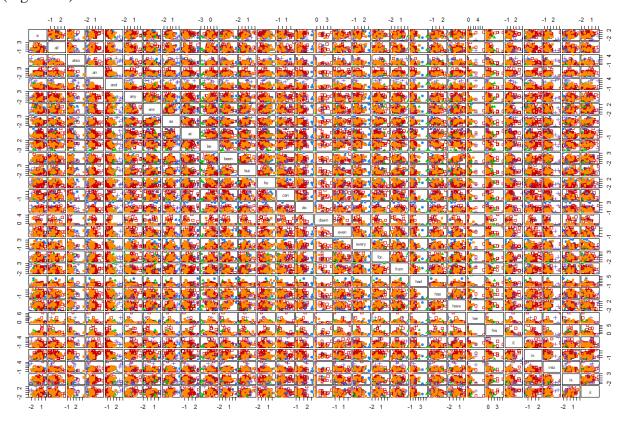
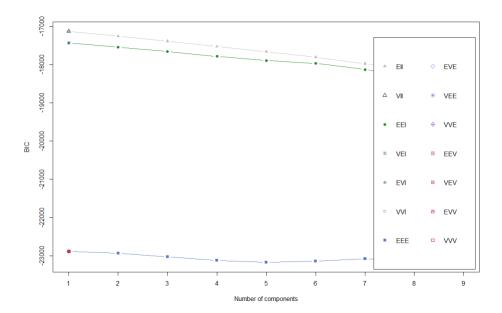


Figure 15

Next, we calculated the fit of the model with one component. The log.likelihood is -8407.477. This is the log likeligood of the BIC Value, which is used for choosing clusters. However, when we plotted our data, the EM algorithm divided all the data points into one cluster, this made it difficult to tell the disputed authors.



### Conclusion:

While the Federalist Papers remain, an important document related to the history of founding and ratification of the US Constitution, understanding who the author of the 11 disputed papers is still considered somewhat of a mystery. While our results were able to, for the most part, show a stronger likelihood that the author was Madison, the results are still not conclusive. One of the main questions raised during this analysis was if the data could be displayed in a hierarchical manner, where records are connected through links. Instead, maybe modeling the papers in a network-like way might be better. Additionally, it is also possible that the papers were written together, by Madison and Hamilton, or even with the assistance of John Jay.

In conclusion, while we did find that the disputed papers are likely the product of Madison, it is also possible that he had help. Further modeling and analysis is required.

# Appendix A: Results

# Option 1 Results:

```
eans clustering with 4 clusters of sizes 24, 4, 11, 46
```

Option 2 Results:

```
> km_output1
K-means clustering with 4 clusters of sizes 46, 11, 24, 4
Cluster means:
```

# Appendix B: R Code

data\_path=file.choose()
data=read.csv(data\_path)
str(data)
sum(!complete.cases(data))
summary(data)
fp<-data[,c(-1,-2)]</pre>

#Scaling fp data

```
fp2<-data.frame(scale(fp,center = T,scale = T))
#Plot (elbow method) to decide optimal number of clusters
set.seed(10)
optim.cluster<-function(k){
 return(kmeans(fp2,k,nstart = 25)\$tot.withinss)
k values<-1:14
oc_values<-purrr::map_dbl(k_values,optim.cluster)
plot(x=k_values, y=oc_values,type="b",frame=F,xlab = "Number of clusters K",ylab="Total
within-clusters (sum squared)")
#### Option 1: K means_Clustering ####
k mean<-kmeans(fp2, centers = 4, nstart = 25)
k mean
k_mean$centers
######Option 2: :K_means using Hartingan-Wong Method####
##HW algorithm to find centroid
km output1<-kmeans(fp2,centers=4,nstart = 25,iter.max = 100,algorithm = "Hartigan-Wong")
km_output1
km output1$centers
#visualize the result
install.packages("cluster")
library(cluster)
install.packages("ggplot2")
library(ggplot2)
clusplot(fp2,km output1$cluster,color=TRUE,shade=T,labels = 2,lines =0)
km_df1<-data.frame(author=data$author,cluster1=km_output1$cluster)
ggplot(km_df1)+geom_polygon(aes(x=author,y=cluster1,group=author,fill=as.factor(cluster1)),c
olor="red")+
 coord_fixed(1.3)+
 guides(fill=F)+
```

```
Parin Patel
```

```
theme bw()
##### 2nd Algorithm: HAC#####
hac_output<-hclust(dist(fp2,method = "euclidean"),method = "average") #average
hac_output2<-hclust(dist(fp2,method = "euclidean"),method = "complete") #complete
#plot HAC
plot(hac_output) ##average
plot(hac output2) ## complete
#output desirable number of clusters after modeling
#average linkage
hac cut<-cutree(hac output,4)
hac_df1<-data.frame(author=data$author,cluster=hac_cut)
avg Clus plot<-clusplot(fp2,hac cut,color=TRUE,shade=T,labels = 2,lines =0)
#compelte linkage
hac_cut<-cutree(hac_output2,4)
hac_df1<-data.frame(author=data$author,cluster=hac_cut)
clusplot(fp2,hac_cut,color=TRUE,shade=T,labels = 2,lines =0 )
####3rd Algorithm: EM####
install.packages("mclust")
library(mclust)
#visualize clusters
clPairs(fp2 [1:30],data$author) ##disputed
fit<-Mclust(fp2)
summary(fit)
#1. BIC (The Bayesian information criterion (BIC)
## is used my mclust with is a test used to assess the fit of a model)
plot(fit,what= "BIC")
#2. classification
plot(fit, what = "classification")
length(fit$classification)
```