PERCEPTION ABOUT GMOS BETWEEN THE USA AND EUROPE THROUGH TWITTER

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Feng Chang

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PERCEPTION ABOUT GMOS BETWEEN THE USA AND EUROPE THROUGH TWITTER

By
Feng Chang

The Supervisory Committee certifies that this disquisition complies with North Dakota State University’s regulations and meets the accepted standards for the degree of

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SUPERVISORY COMMITTEE:

Dr. Kendall E. Nygard
Chair

Dr. Oksana Myronovych

Dr. Supavich Pengnate

Approved:

7/6/2017

Dr. Kendall E. Nygard
Date
Department Chair
ABSTRACT

Twitter is one of the most popular social networking and microblogging services in the internet world. By the end of Jan 24, 2017, there were at least 100 million active users daily around the world. Due to such a huge number of people, it is a great channel to collect information about almost anything existing in the world. Based on this information, we could also analyze the popular topics which are widely discussed. For the work of my paper, the target is to find out the perception about GMOs, but also focused on the difference in perceptions between Europe and the United States. To accomplish this work, a collection of Twitter was collected that all include the text about GMOs, along with their locations. Analytics were performed on the tweets to classify them by sentiment, then statistical tests were carried out to assess differences in perceptions by location.
ACKNOWLEDGEMENTS

I would like to give my sincere gratitude to my advisor, Dr. Nygard, for his valuable suggestions, patient guidance, and unceasing encouragement during my whole research project. I also would like to thank my supervisory committee, Dr. Myronovych and Dr. Pengnate. I really appreciate their time, kindness, and profound comments.

Also, I want to give my wife and family great thanks, for their accompanying, supports, encouragements, and love during my whole life.
DEDICATION

This paper is dedicated to my wife and parents for their endless love, support, and encouragement.
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1. INTRODUCTION

According to the 2015 revision of UN's project, the world population was expected to grow by 2.3 billion between 2009 and 2050. This means that we need to produce at least 70 percent more food production by 2050 for the population growth [1]. Therefore, to provide so much more food is an urgent task on the earth. We could ask people to stop wasting food, or create more land to farm, or come up ways to increase the yield. Genetically modified foods is one of the methods for increasing supply since genetic technology can boost the yield. Many believe that GMOs provide a very promising approach to feed the whole world, especially for those fast growing population countries.

However, we always say everything has two sides. Even though now we could use genetically modified foods technology to increase the yields and feed the whole world, we still cannot ignore its dark side. We wonder whether or not GMO foods are healthy or not. Nowadays, people often to share or express their ideas, opinions, events on those social media networks, especially on Twitter. The purpose of this paper is to collect Twitter feeds and work out analytics on the data to find out people's perceptions about GMOs. Two years ago, people from the same research team at NDSU had published a paper which concluded that regional locations in the United States would influence people's feeling toward GMOs [2]. Also, another researcher from the team wrote another paper on how gender differences [3] would affect people's perception in GMOs. Inspired by those two paper, this paper is to determine the attitude differences about GMOs between Europe and the United States.

To accomplish this goal, a Twitter – based analysis was formulated and applied. The study on Twitter is about collecting the tweets that use the keyword GMO. The tweets are then separated by different locations. The next major step is to classify how many tweets are positive
and how many are negative. After that, statistical calculation or tests were run to find out the
different attitudes between the United States and Europe. Some people may choose to not reveal
their locations in Twitter. That is a distraction for my research, so those tweets were removed.

The rest of the paper is structured as follows. The second chapter includes a literature review. The third chapter describes the way to capture and process the data. The fourth chapter describes the method to classify the data from the processed and cleaned data. In the fifth chapter, the classified data is analyzed, and some statistical results are provided for both United States and Europe. The sixth chapter is the conclusion of the whole paper and suggestions for the future work.
2. LITERATURE REVIEW

In this chapter, it will focus on covering the needed background, which is very important to the paper. Also, some similar research work will be discussed.

2.1. Background

2.1.1. Twitter Streaming API

Twitter provides their APIs to potential developers. Since mobile development is become a hotspot, more and more individual developer started to create their own applications or projects based on the services from Twitter, Facebook or Google. Also, many students and researchers started to use their services for research purposes. Therefore, Twitter allowed a developer to access their global streaming tweets data by using their Streaming API [4]. Twitter provided two APIs for the developer to collect tweets. One is the Search API, the other is the Streaming API. Based on two reasons the Streaming API was chosen. First, it is desirable that the tweets collecting process goes back in time. Second, a higher flow of tweets is available through streaming. The Streaming API delivers a maximum flow of 180,000 tweets per hour, which is two times more than Search API [5]. After Twitter data is collected, some cleaning work was carried out immediately.

2.1.2. Support Vector Machine Classification

Support Vector Machine (SVM) is a machine learning method which was established the 1990s. It partitions the data off into two sides based on a separating classification plane. Normally, a distance between a point or item and a classification plane measures the degree of accuracy. A SVM could maximize the distance, making it as accurate as possible. It enhances generalization ability by its Structural Risk Minimization. To put it simply, even though we might not have enough sample data, we still can achieve a good statistical result. Therefore, it
would be a good classification to use. The tool which this paper uses is scikit – learn’s LinearSVC. It’s because based on Marco’s experiment [6], LinearSVC is much faster than other SVC methods that are provided in scikit – learn. Chapter four will present more detail about this classification method.

2.1.3. Naïve Bayes Classification

Naïve Bayes Classification is based on Bayes theory. We have a formula [7]:

\[ P(A | B) = \frac{P(B | A)P(A)}{P(B)} \]

- \( P(A|B) \): Probability of observing event A given B is true.
- \( P(B|A) \): Probability of observing event B given A is true.
- \( P(A) \): Probabilities of A
- \( P(B) \): Probabilities of B

The Naïve Bayes ignores dependencies among events. This work uses basic Naïve Bayes tool which implements the same process in Python.

To implement this method, we have three steps. First, it requires classified data sample. Next, those classified datasets are provided to train the Naïve Bayes Classifier. Finally, a group of test data is needed to determine the accuracy of the trained Naïve Bayes Classifier. Chapter four will have the detail about the whole process.

2.1.4. TF – IDF

In TF – IDF, there are two terms [8]:
• Term Frequency is the ratio between the count of one word in a document and total word count in a document

• Inverse Document Frequency is the logarithm of the ratio between the total corpus and the corpus which has the specific word plus one.

Then, we multiple TF and IDF to get the result of TF – IDF. The bigger we got, the more the word is. Python NLTK package provides TF – IDF tool. Chapter Five will provide the detail about the information extraction by using TF – IDF.

2.2. Related Work

According to every customer's intention, we have a huge amount of research has been done which is related to GMOs. Also, we had always heard about that Europe was against GMOs the most. One article from The New York Times was written by Mark Lynas [9] attempted to tell us that Europe tries to avoid GMOs even turned against science. We could simply see that how serious Europe was to treated GMOs. They were not only saying that but also some countries like Germany, France, Greece, Italy, started to ban on the cultivation of genetically modified crops. Because Twitter is a good place for people to post their thoughts about something, researchers also start to capture and analyze Twitter feeds for multiple data mining purposes. Hanze, Li [10] examined the opinions and sentiments from his Twitter dataset on GMOs. In the paper, the attitude around the Europe or United States’ on GMOs is the main target. Thus, both United States and Europe are two focused places, and their real difference is and what the real reason might be are the two main purpose needs to be revealed.

2.3. Motivation

Since GMO has been brought out, people start to debate on that. In 2013, two famous people in China had totally opposite ideas about GMOs and they were against each other. The
scientific organization like WHO (National World Health Organization) had stated that GMOs are safe, but still, lots of people don't believe that. They concern the safety, the generic pollution. On the other hand, grocery store provided labels to distinguish GMO and not – GMO food. This also made us consider if GMOs is safe, why they need labeling. Also, we just heard about people dislike GMOs, but do they or we really know what GMOs is? Or do people really dislike it or they just follow other's idea blindly? So many questions make me want to find out the truth behind everything.

A social network is a nice place for people to present their true feeling about something with others. Anytime you post something on Twitter, you will get plenty of replies which include strangers or friends. As time goes on, it became a big data collection which involves almost every topic. Therefore, it became the best dataset for researchers to do data mining things. It is fast to access, less limitation to follow, and more data field to get.

Furthermore, United States is one of the biggest developed countries. Europe is one of the biggest developed union. Therefore, their opinion about GMOs may reveal some real reason about people's attitude. Also, we had a stereotype that compares to the United States, Europe is stricter against GMOs. Based on less evidence or lack of the same type of paper online, the stereotype is doubted, so it is necessary to find out the truth.
3. TWEETS DATA PREPARATION

Even though there are lots of data mining research online, still less paper about GMOs attitude's comparison between the United States and Europe were found. Thus, this paper will make up the lack of the research. This research includes the complete process of collecting Twitter feeds, the whole process of cleaning the collected Twitter feeds, the datasets which are from other universities for classification training, the process of categorizing test data and further analysis.

3.1. Twitter Feeds Aggregation

As I mentioned in the instruction, as an individual developer who wants to do a research or develop their own project, they have to work with Twitter API. However, everyone still needs to configure out the following steps.

3.1.1. Getting Authentication

Whenever you want to start you Twitter project, you will need a Twitter account. Then, you should go to Twitter application management website to create a new application based on filling out the required form which shown in Figure 1.

![Create an application](image)

Figure 1. Create Your New Application.
Once your application is created, you will have a profile page for your project like figure

Figure 2. Application Profile.

Then, you go to Keys and Access Tokens and get consumer key, consumer secret, access token and access token secret. It means that every time when you use Twitter API, you must have those four keys in your application for credentials. If you miss any one of those or you misspell even one letter, your calls to Twitter will fail and you will get nothing.

3.1.2. Use Tweepy with Streaming API

Even though all the credentials are created in the account, the API connection is still not set up yet. The Tweepy package is the connection in Python between developers’ call and API. It is an easy to use the library in Python for accessing Twitter API [11].

3.1.3. Collect and Clean my Tweets

Now, it is the data capturing process. Figure 3 provides the structure of my Twitter feeds collecting and cleaning process.
Figure 3. Collecting and Cleaning Tweets.
Note: For collecting and cleaning Tweets, it has three step totally. All of them are using Python and its tool package.

As the stated information above, this paper will focus on two things in tweets. One is location, the other is the text. However, for Twitter Streaming API, it will return me JSON type data like Figure 4 [12].
Figure 4. A Sample of JSON Response.
Note: The data returns from Twitter Streaming API has its JSON format and includes more than one or two tuple value.

First, Tweepy package in Python will extract username, text, and location for the United States. However, not every tweet has its location. Some people prefer to uncheck the option “show my location”. The tweets which don’t have location will be filtered out. Also, the method to locate a tweet is to set up by its location’s geocode. Second, when those tweets’ locations were identified, most of their text format is a city, state-based. Then, it is convenient to have an array
only includes all the states and plus an extra word, "USA" to match up with locations in tweets to filter out the US locations. This way has two benefits. One is to make sure tweets is from the US, the other is to make sure the location format stays the same. The final step, emojis, URLs, retweets or those special symbols need to get rid of from the tweets still. The tool is re package in Python. It only needs one simple line of code to remove all those things: “re.sub(r"([\w]+)|([\^0-9A-Za-z\s\t])|(https+)|(RT @)"",",item.text)”. Now the final processed US data is in Table 1.
Table 1

20 Lines Processed Tweets for the USA

<table>
<thead>
<tr>
<th>Username</th>
<th>Text</th>
<th>Location</th>
</tr>
</thead>
<tbody>
<tr>
<td>yeah_im_gmo</td>
<td>Chad Reed is Elís last hope</td>
<td>USA</td>
</tr>
<tr>
<td>Samiam01x</td>
<td>Easy Ways to Go GMOFree and Why You Should gruccifer2 pintet america</td>
<td>USA</td>
</tr>
<tr>
<td>NeoProgressive1</td>
<td>VirginiaInCal Boycott all Pepsi products like gruccifer2 pismet america StacysPitaChips amp Sabra for blocking GMO labeling lawsTry Non GMO Verified brands in</td>
<td>KY</td>
</tr>
<tr>
<td>NeoProgressive1</td>
<td>VirginiaInCal Pepsi has dumped 12M into blocking GMO labeling BOYCOTT all Pepsi products like Cheetos for blocking GMO labeling la</td>
<td>KY</td>
</tr>
<tr>
<td>NeoProgressive1</td>
<td>VirginiaInCal Dont be tricked by Pepsi into buying beverages wtoxic GMOs like OceanSpray Boycott all Pepsi products for blocking GMO labeling la</td>
<td>KY</td>
</tr>
<tr>
<td>city_market_CY</td>
<td>Mid night snack I ate the whole tub of guacamole with these Super clean Non GMO</td>
<td>TN</td>
</tr>
<tr>
<td>NeoProgressive1</td>
<td>VirginiaInCal PRODUCT WARNING Smuckers is selling unlabeled GMOs BOYCOTT all Smuckers products like Knotts 4 blocking GMO labeling la</td>
<td>KY</td>
</tr>
<tr>
<td>jayizzo50</td>
<td>trutherbotred GMO Foods YOUR KILLER</td>
<td>CA</td>
</tr>
<tr>
<td>jlacy64</td>
<td>fuck gmos endGMOs</td>
<td>USA</td>
</tr>
<tr>
<td>randybear29</td>
<td>ImBeginningToSuspect depopulation started with GMOs</td>
<td>OH</td>
</tr>
<tr>
<td>tai_writes</td>
<td>xalimoso I cant wait to be a mom id pack my dogs lunch with organic no GMOs kibbles</td>
<td>USA</td>
</tr>
<tr>
<td>egojab</td>
<td>Jollett For most who oppose GSM its not really about safe but about natural</td>
<td>WA</td>
</tr>
<tr>
<td>Porscheey</td>
<td>Stop eating meat its bad for you GMO free</td>
<td>TX</td>
</tr>
<tr>
<td>DefendingBeef</td>
<td>Grass fed finished cattle dont eat GMO corn or fillers Grasses dont require pesticides So just be</td>
<td>CA</td>
</tr>
<tr>
<td>L_Gale517</td>
<td>ACOSorg jdaniel</td>
<td>USA</td>
</tr>
<tr>
<td>Taylor96Taylor</td>
<td>MeosoFunny Doesnt Eat GMO Because of Dangers Diet Consists of Drugs Alcohol and Trail Mix</td>
<td>USA</td>
</tr>
<tr>
<td>DTPORGE</td>
<td>NoGMOsVerified Select Committee report on GM is an insult to science and a danger to the public GMOs RightToKnow GMO gw</td>
<td>NY</td>
</tr>
<tr>
<td>DTPORGE</td>
<td>FarmFairyCrafts Boycott for Advertising in her Magazine GMO So Disappointed</td>
<td>NY</td>
</tr>
<tr>
<td>DTPORGE</td>
<td>FarmFairyCrafts via Second Silent Spring Bird Declines Linked to</td>
<td>NY</td>
</tr>
</tbody>
</table>

Table 1 shows 20 lines of processed Tweets which only have username, text, and location.

Next, a different process needs to be implemented for Europe Twitter feeds. To locate Europe, it is not so efficient to just use its geocode, so to pick up the main cities for Europe
countries is the best way. The reason is easy to find the matched geocode. Those cities are Paris, Berlin, London, Rome. People can see those cities are all main cities for their country, so they should be able to represent their country somehow. The rest of the steps for Europe tweets are the same. Therefore, Europe tweets data is in Table 2.

    Totally, two months were spent to collect tweets. Next, one final step is to put them into one corpus for the next experimental stage. In this step, tweets were separated into the United States and Europe.
Table 2

<table>
<thead>
<tr>
<th>Username</th>
<th>Text</th>
<th>Location</th>
</tr>
</thead>
<tbody>
<tr>
<td>adonisfoods</td>
<td>Erythritol is 100% natural the one we use is organic and GMO free and come from fermented fruits</td>
<td>London, England</td>
</tr>
<tr>
<td>ashmakkar</td>
<td>I support bio</td>
<td>London</td>
</tr>
<tr>
<td>ashmakkar</td>
<td>India India Hindi</td>
<td>London</td>
</tr>
<tr>
<td>TheTapBlog</td>
<td>Dark History of Bayer Crop Science from manufacturing poison gas to hiding side effects of its min</td>
<td>London, England</td>
</tr>
<tr>
<td>Col_Connaughton</td>
<td>Lies Lies and More Lies GMOs Poisoned Agriculture and Toxic Scientific Rants GMO india monsanto</td>
<td>UK</td>
</tr>
<tr>
<td>ActivistFangirl</td>
<td>HELP NEEDED Azure Organic Farm in Oregon will be forcibly mass poisoned with glyphosate by the county government</td>
<td>London, United Kingdom</td>
</tr>
<tr>
<td>weedseeds_UK</td>
<td></td>
<td>London, England</td>
</tr>
<tr>
<td>Col_Connaughton</td>
<td>July2315 False Flag Weekly News falseflag vaccination iran MH17 TTIP israel GMO fraud</td>
<td>London, UK</td>
</tr>
<tr>
<td>buildeven</td>
<td>Modern agritech GMO needs a more lightly regulated US State to thrive but its so not Iowa</td>
<td>London, England</td>
</tr>
<tr>
<td>jasminglynne</td>
<td>gmo is so fucked not from a health standpoint but from a corporate standpoint like wtaf</td>
<td>London</td>
</tr>
<tr>
<td>gmo_crops</td>
<td>GMO news Brazils Mato Grosso Leads Push for GMFree Soy The largest soy-producing state in Brazil Mato Gross</td>
<td>London</td>
</tr>
<tr>
<td>gmo_crops</td>
<td>GMO news Mustard Set to Be Indias First GM Food Gets Regulator Nod</td>
<td>London</td>
</tr>
<tr>
<td>gmo_crops</td>
<td>GMO research TLCUV hyphenated with MALDITOFMS for the screening of invertase substrates in plant extracts</td>
<td>London</td>
</tr>
<tr>
<td>gmo_crops</td>
<td>GMO research The use of Stationary Phase Optimized Selectivity Liquid Chromatography for the development of h</td>
<td>London</td>
</tr>
<tr>
<td>gmo_crops</td>
<td>GMO research Immunoaffinity chromatography combined with tandem mass spectrometry A new tool for the selectiv</td>
<td>London</td>
</tr>
<tr>
<td>vincentdignan</td>
<td>Hey GMO welcome Want to get your posts seen on Facebook Read this</td>
<td>London, England</td>
</tr>
<tr>
<td>pecasyrizos</td>
<td>Nothing less of organic gluten free gmo free raw vegan buckwheat brown bread with sprouted hemp seeds of course</td>
<td>London, England</td>
</tr>
<tr>
<td>bigpicturetv</td>
<td>The latest The Big Picture Daily Thanks to gmo mothersday</td>
<td>London</td>
</tr>
<tr>
<td>bigpicbiz</td>
<td>The latest The Peter Eyres Daily Thanks to tobias gmo sustainability</td>
<td>London</td>
</tr>
</tbody>
</table>
4. TWEETS DATA CLASSIFICATION

The main purpose is to compare tweets between USA and Europe to see the real attitude and to try to find out the real reason that Europe is against GMOs. Obviously, either way, to classify the sentimental results is the first duty. The data only has two polarities, positive and negative. It’s because my training datasets are also having only two polarities. Support Vector Machine classifier will be trained first, then with Naïve Bayes Classification. Also, two datasets online were downloaded for training purpose. The complete structure of classification process is in figure 5.
Figure 5. The Structure of Classification Process.
Note: The man – labeled Tweets dataset will be split into two part, training, and testing. Training dataset trains those classifiers for classifying collect data later. Testing dataset finds out the classifier's accuracy.
4.1. Support Vector Machine Classification

Based on chapter two, even though we might not have enough sample data, we still can result a good statistical regularity with SVM. Therefore, it would be a good classification to deal with the size of my data

4.1.1. Set up Tools

In Python, we have plenty of good tools to deal with sentimental analysis. The main two are NLTK [13] and Sklearn.svm. NLTK is a famous tool go work with human language data. It builds – in some easy to use methods for us to play with. First, every line in data needs to be separated into word format, so nltk.tokenize package help me with that. Next, if the classifier is well – trained, it will be better to test the data, so nltk.classify is chosen. Both of them are very handy. The rest leaves to Sklearn.svm. It provides LinearSVC for me to create my training model.

4.1.2. Train Dataset

The first dataset is from Bo Pang and Lilian Lee’s [14] movie review data [15]. It included 5331 positive and 5331 negative snippets. The size of data separation is like figure 6.

```
training_data = pos_data[:4000] + neg_data[:4000]
testing_data = pos_data[4000:] + neg_data[4000:]
```

*Figure 6. The Size of Training and Testing Data.*

Note: Training data has the first 4000 of 5331 from both positive and negative data. Testing data has the rest from both data.

The first 4000 data from both positive and negative data are used to organize a training data. Then, the rest of the data became the testing data. The ratio is about 3 approximately.

Next, it is a dataset from Alec Go, Richa Bhayani, and Lei Huang [16]. It is pure Twitter feeds data. And according to its size, only five thousand sentences are selected for each positive
and negative tweets. To match the previous ratio, the 3750 are for training and 1250 are for testing like figure 7.

\[
\text{training\_data} = \text{pos\_data}[:3750] + \text{neg\_data}[:3750] \\
\text{testing\_data} = \text{pos\_data}[3750:] + \text{neg\_data}[3750:] 
\]

Figure 7. The Size of Training and Testing Data

4.1.3. Test Accuracy

Finally, figure 8 shows us two groups of quite high results.

Figure 8. Accuracy in SVM for Different Data.
Note: In the SVM model, the accuracy is about 0.755 for movie data and is 0.702 for Tweets data. Even from the chart, the distance from pink top to the orange top is significant, it is 0.05 difference.
4.1.4. Classify Collected Data

Since the accuracy is so high on both training datasets, the research will use them to classify my own processed dataset. Mine is also pure tweets data includes 1097 positive and 1097 negative. This time a surprising result is shown in figure 9.

![Figure 9. Classified Tweets Data Percentage in SVM.](image)

Note: Support Vector Machine has about 0.687 for the USA negative and 0.688 for Europe negative in movie data, but it has about 0.364 for the USA negative and 0.264 for Europe negative in Tweets data.

From this chart, we saw a pair of opposite attitudes based on different data types. Chapter six will dig deep on that. For now, it will generate a txt file which stored those negative tweets from SVM model by using both datasets.
4.2. Naïve Bayes Classification Process

From book Data Mining Algorithms: Explained Using R [17], it stated that the naïve Bayes classifier is one of the simplest methods to achieve classification problem with a reasonable accuracy. Therefore, another way to test how well my dataset works is to use Naïve Bayes Classifier. And the main steps and datasets are the same.

4.2.1. Set up Tools

This time, the only tool is NLTK. It has everything built – in for Naïve Bayes. Those tweets still need to be separated into words format. Then, they will be trained and tested by Naïve Bayes Classifier. Next, the accuracy of trained model will be presented. At the end, the model will be used to classify the collected tweets.

4.2.2. Train Dataset

Due to the same standard, SVM used the same datasets. Also, the separation ratio is the same as figure 6 and 7.

4.2.3. Test Accuracy

For Naïve Bayes, the accuracy for movie data is still higher than tweets data. It’s in figure 10. The value is still quite high.
Figure 10. Accuracy in Naïve Bayes for Different Data.
Note: In the Naïve Bayes model, the accuracy is about 0.777 for Movie data and 0.697 for Tweets data. Even from the chart, the distance from pink top to the orange top is significant, it is 0.08 difference.

4.2.4. Classify Collected Data

Again, the trained Naïve Bayes model could be used to classify the collected tweets data.

The result is showing in figure 11.
Figure 11. Classified Tweets Data Percentage in Naïve Bayes.
Note: Naïve Bayes has about 0.565 for the USA negative and 0.502 for Europe negative in movie data, but it has about 0.488 for the USA negative and 0.382 for Europe negative in Tweets data.

Overall, the Naïve Bayes results don't have any significant difference compared to SVM, but positive attitude increases in Naïve Bayes. Again, the paper will leave a discussion of that in chapter six and generate a txt file which stored those negative tweets from Naïve Bayes model by using both datasets.
5. PROCESSED DATA ANALYSIS

In the previous step, the SVM and Naïve Bayes models generate some txt files only include negative tweets. Those txt files will be analyzed to get a close answer of my question. Next, TF – IDF will be used here to extract the important information [18]. People often use it to retrieve information or mine text. It fits here because the reasons for denying GMOs may come from some keywords in those tweets or Twitter's text, so TF – IDF will help me analyze how important the specific word is a document in a corpus. Also, if the value of TF – IDF is bigger, the word is more important.

5.1. Count Duplicate Words

Before TF – IDF is implemented, the first method comes up in my mind is the simplest one, the most important word should appear the most in the corpus. Therefore, all the duplicate words are counted for each txt file. Based on all the results, table 3 to show a top 20 words which appears more than once from all negative tweets from Standford Twitter Data Naïve Bayes USA (TNBUSA), Standford Twitter Data Naïve Bayes Europe (TNBEURO), Movie Review Data Naïve Bayes EURO (MNBEURO), and Movie Review Data Naïve Bayes USA (MNBEUSA).
Table 3

*Top 20 Words from txt files*

<table>
<thead>
<tr>
<th>TNBUS</th>
<th>TNBEU</th>
<th>MNBEU</th>
<th>MNBUS</th>
</tr>
</thead>
<tbody>
<tr>
<td>gmo:406</td>
<td>gmo:342</td>
<td>gmo:434</td>
<td>gmo:474</td>
</tr>
<tr>
<td>the:214</td>
<td>to:147</td>
<td>the:244</td>
<td>the:250</td>
</tr>
<tr>
<td>to:187</td>
<td>the:126</td>
<td>to:213</td>
<td>to:227</td>
</tr>
<tr>
<td>and:125</td>
<td>of:90</td>
<td>of:115</td>
<td>of:131</td>
</tr>
<tr>
<td>a:119</td>
<td>in:90</td>
<td>in:110</td>
<td>a:121</td>
</tr>
<tr>
<td>in:117</td>
<td>news:87</td>
<td>a:84</td>
<td>in:116</td>
</tr>
<tr>
<td>of:110</td>
<td>and:78</td>
<td>and:81</td>
<td>and:103</td>
</tr>
<tr>
<td>is:109</td>
<td>a:64</td>
<td>on:80</td>
<td>is:100</td>
</tr>
<tr>
<td>i:86</td>
<td>is:63</td>
<td>for:73</td>
<td>i:83</td>
</tr>
<tr>
<td>for:69</td>
<td>for:51</td>
<td>news:70</td>
<td>gmos:83</td>
</tr>
<tr>
<td>gmos:68</td>
<td>i:41</td>
<td>is:68</td>
<td>you:81</td>
</tr>
<tr>
<td>that:62</td>
<td>on:36</td>
<td>i:56</td>
<td>for:78</td>
</tr>
<tr>
<td>on:58</td>
<td>plant:35</td>
<td>latest:43</td>
<td>on:76</td>
</tr>
<tr>
<td>food:56</td>
<td>with:35</td>
<td>via:43</td>
<td>are:73</td>
</tr>
<tr>
<td>are:54</td>
<td>from:34</td>
<td>plant:42</td>
<td>that:68</td>
</tr>
<tr>
<td>non:50</td>
<td>food:34</td>
<td>gmos:41</td>
<td>amp:57</td>
</tr>
<tr>
<td>it:47</td>
<td>no:32</td>
<td>research:39</td>
<td>it:55</td>
</tr>
<tr>
<td>monsanto:47</td>
<td>are:30</td>
<td>that:38</td>
<td>organic:54</td>
</tr>
<tr>
<td>not:46</td>
<td>not:30</td>
<td>daily:38</td>
<td>monsanto:52</td>
</tr>
<tr>
<td>with:45</td>
<td>that:29</td>
<td>you:38</td>
<td>corn:52</td>
</tr>
</tbody>
</table>

Table 3 show the top 20 duplicated words in all txt files.

### 5.2. Count Negative Word

Look at the table from the previous step, it is really hard to get any information from those most counted words. Then, to count only those negative words might be a better idea. To do so, I need to work with some English dictionary. I got a Harvard-IV_NegativeWordList [19].

This time a disappoint result shows in figure 12.

![none of your words are listed in Harvard-IV_NegativeWordList.txt](image)

*Figure 12. Result of Negative Words Count*

No negative words were found in negative tweets which could match with the dictionary.

It is easy to understand because users in Twitter doesn’t type like we speak in daily life.
5.3. TF – IDF Process

In the end, TF – IDF might the best tool. Compare to other two, it is more reliable because it could be used to get important words from each single sentence. For instance, the sentence, “Fashion tips Bowling Green is a very feminine textileTacrobes are not robes but GMO”, the TF – IDF tool will calculate a score or weight for each word in this sentence like table 4. Column Word is one tweet’s sentence word split. Column Document is the sentence position in the txt file. Column Score is the TF – IDF weight for each word in the same sentence.

Table 4
Example of TF – IDF result

<table>
<thead>
<tr>
<th>Word</th>
<th>Document</th>
<th>Score</th>
</tr>
</thead>
<tbody>
<tr>
<td>Fashion</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>tips</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Bowling</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Green</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>is</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>a</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>very</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>feminine</td>
<td>0</td>
<td>0.4</td>
</tr>
<tr>
<td>textileTacrobes</td>
<td>0</td>
<td>0.5</td>
</tr>
<tr>
<td>are</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>not</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>robes</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>but</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>GMO</td>
<td>0</td>
<td>0.2</td>
</tr>
</tbody>
</table>

Table 4 shows the possible example of TF – IDF result.

And, TF – IDF is also built – in Sklearn Python package.

5.4. TF – IDF Results

Combine all those txt files, two final results for the USA and Europe are showing. Figure 13 is for the USA, and figure 14 is for Europe. These are part of the TF – IDF results. Left is the word, right is the score.
Figure 13. TF – IDF Result for the USA.

Note: After implementing TF – IDF, the result is showing the important word which has the value higher than 0.45 for the USA.
Figure 14. TF – IDF Result for Europe.
Note: After implementing TF – IDF, the result is showing the important word which has the value higher than 0.45 for Europe.

Also, after sorting the values in the results, the top 10 word for bother results are showing in Table 5.

<table>
<thead>
<tr>
<th>Top 10 USA</th>
<th>Top 10 Europe</th>
</tr>
</thead>
<tbody>
<tr>
<td>labeling</td>
<td>facts</td>
</tr>
<tr>
<td>la</td>
<td>fears</td>
</tr>
<tr>
<td>twizzler</td>
<td>safety</td>
</tr>
<tr>
<td>hershey</td>
<td>foods</td>
</tr>
<tr>
<td>for</td>
<td>and</td>
</tr>
<tr>
<td>boycotting</td>
<td>of</td>
</tr>
<tr>
<td>reeses</td>
<td>the</td>
</tr>
<tr>
<td>stand</td>
<td>gmo</td>
</tr>
<tr>
<td>virginaincal</td>
<td>explain</td>
</tr>
<tr>
<td>laws</td>
<td>explained</td>
</tr>
</tbody>
</table>

Table 5 shows the top 10 words in TF – IDF results for both USA and Europe.
6. CONCLUSION AND FUTURE WORK

In this chapter, it is an overall review of this paper and also discusses how we can improve this study in the future.

6.1. Conclusion

Overall, for this paper, the purpose is to find what the real attitudes on GMOs are in the USA and Europe and the truth behind people dislike GMOs. All the necessary steps are implemented based on the designed structure.

First, it created a Twitter account for authentication purpose, so that people can get all Twitter API keys include API key, API secret, Access token secret and Access token. Next, those four credentials must be added into Python code to connect Twitter Streaming API with the Python code to collect new dataset. After that, all those easy to use, built-in Python tool helped to extract information from tweets and clean those tweets. Also, SVM and Naïve Bayes are trained by using some good online datasets to classify the new dataset. Once the collected tweets are separated into positive and negative. Then, based on TF – IDF, it presented some clue which might provide some useful information in the future research. And, it got lots of interesting results finally.

In the end, the experimental results gave some answers for most of the questions. Starts now, people would not say they are 100% sure Europe is stricter to GMOs. One of those figures in Chapter four shows that the negativity is higher in the USA. Also, a thought might be true, sometimes people may just follow other’s ideas, like against GMOs. If people really know why they hate it, when count negative word is counted in the dataset, it definitely should have some match results. However, the last doubt still hard to find out, it is the truth behind against GMOs which also brings out the future work for next part.
6.2. Future Work

I want to add a new thing for future work first. This paper is using Streaming API with string “gmo”, but the return the Tweets are never checked. It might not be related to GMOs. Therefore, it is important to check if the Tweets is really about GMOs. Like previous part mentioned above, there is no answer for the truth behind people against GMOs yet. To look at those important words from TF – IDF, it is really hard to get any idea from that, but still, people could see some words like safety, poisonous, healthy, etc. Also, every time when N – gram is added into TF – IDF to assist in getting a better result, the laptop just slows down and not responds for a long time. Therefore, in the future, researchers could use a more powerful computer and combine with more precise language skills to dig deep into those words. Then, they could categorize them, so that they might have some ideas about the real reason that people hate GMOs.
REFERENCES


