MAT 167: Applied Linear Algebra Lecture 22: Text Mining

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Outline

- Introduction
- Preprocessing the Documents and Queries
- The Vector Space Model
- 4 Latent Semantic Indexing

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- 2 Preprocessing the Documents and Queries
- 3 The Vector Space Model
- 4 Latent Semantic Indexing

- Text mining = Methods for extracting useful information from large and often unstructured collections of texts.
- It is also closely related to "information retrieval."
- In this context, keywords that carry information about the contents of a document are called terms.
- A list of all the terms in a document is called an index.
- For each term, a list of all the documents that contain that particular term is called an *inverted index*.
- A typical application is to search databases of scientific papers for given query terms.

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- Because of Lecture 2 and HW #1, you should already be familiar with the concept of term-document matrix.
- Each column represents a document while each row represents a term
- The ijth entry of such a matrix normally represents the frequency of occurrence of term i in document j.
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- In this lecture, we will use the following 'Bags of Words' dataset available from the UCI Machine Learning Repository: http://archive.ics.uci.edu/ml/datasets/Bag+of+Words.
- I his is a collection of 1500 (= n) articles (mostly in the field of machine learning and computational neuroscience) published in the proceedings of *Conference on Neural Information Processing Systems* (NIPS) over certain periods.
- The total number of terms (words) examined for these articles is 12419 (= m).
- More precisely, after tokenization (i.e., breaking a stream of text up into words, phrases, symbols, or other meaningful elements called tokens), and removal of stop words (i.e., common words that do not give useful info; more about these in the next section), the vocabulary of unique words was truncated by only keeping words that occurred more than ten times.

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- First 10 words sorted in the alphabetical order: 'a2i', 'aaa', 'aaai', 'aapo', 'aat', 'aazhang', 'abandonment', 'abbott', 'abbreviated', 'abcde'.
- 10 most frequently used words: 'network', 'model', 'learning' 'function', 'input', 'neural', 'set', 'algorithm', 'system', 'data'.

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- Before the index (a list of terms contained in a given document) is made, we need to do the following two preprocessing steps:
 - Elimination of stop words
 - Stemming
- Stop words are words that can be found in virtually any document (i.e., most likely useless words to characterize the documents), e.g., 'a', 'able', 'about', 'above', 'according', 'accordingly', 'across', 'actually', 'after', . . .
- Stemming is the process of reducing each word that is conjugated or
 has a suffix to its stem. For example, 'fishing', 'fished', 'fish', 'fisher'

 stemming

 if ish' (the root word).
- There are some public domain stemming software systems; see 'Stemming' page in Wikipedia.
- Note that stemming was not performed in the NIPS dataset, e.g., the terms include 'model', 'modeled', 'modeling', 'modelling'.

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- The main idea of this model is to create a term-document matrix, say, $A = (a_{ij}) \in \mathbb{R}^{m \times n}$, where each document is represented by a column vector \mathbf{a}_j that has nonzero entries in the position that correspond to terms found in that document.
- Consequently, each row represents a term and has nonzero entries in those
 positions that correspond to the documents where that term can be found,
 i.e., the inverted index.
- In practice, a text parser (a program) is used to create term-document matrices, and does stemming and stop words removal too.
- The entry a_{ij} is normally set to the term frequency f_{ij} , i.e., the number of times term i appears in document j.
- Can have weights, e.g., $a_{ij} = f_{ij} \log(n/n_i)$ where n_i is the number of documents that contain term i. If term i occurs frequently in only a few documents, then the log factor becomes significant. On the other hand, if term i occurs many documents, the log factor makes $a_{ij} \approx 0$, i.e., term i is not useful. Stop words removal mitigates this to some extent.

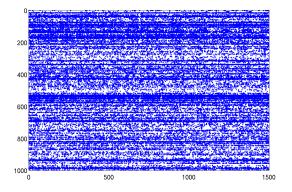
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Usually, the term-document matrix is *sparse*. For example, in the NIPS dataset, the number of nonzero entries in the term-document matrix of size 12419×1500 is 746,316, which is only 4% of the whole matrix entries.



The first 1000 rows of the NIPS term-document matrix. Each dot represents a nonzero entry.

Query Matching

- Query matching = a process of finding the relevant documents for a given query vector $\mathbf{q} \in \mathbb{R}^m$.
- We must define a distance or similarity between q and each document $a_j \in \mathbb{R}^m$, j = 1 : n.
- Often the following cosine distance (in fact, it would be better to say *similarity* rather than distance) is used:

$$\cos(\theta(\mathbf{q}, \mathbf{a}_j)) = \frac{\mathbf{q}^{\mathsf{T}} \mathbf{a}_j}{\|\mathbf{q}\|_2 \|\mathbf{a}_j\|_2}$$

- If $\theta(q, a_i)$ is small enough, then a_i is deemed relevant.
- More precisely, we set some predefined tolerance and if $\cos(\theta(\mathbf{q}, \mathbf{a}_i)) > \cot$, then \mathbf{a}_i is deemed relevant.
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A Query Matching Example

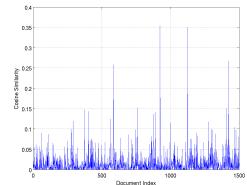
- Let's consider the NIPS dataset and set up $\mathbf{q} \in \mathbb{R}^{12419} = \mathbf{e}_{3528} + \mathbf{e}_{6700} + \mathbf{e}_{6932}$, i.e., only three nonzero entries that correspond to the three terms, 'entropy', 'minimum', 'maximum'.
- Compute $cos(\theta(q, a_i)), j = 1:1500$

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tol=0.2, 0.1, 0.05 correspond to 4, 15, 89 returned documents.

Let us define the following quantities:

Precision:
$$P := \frac{D_r}{D_t}$$
;
Recall: $R := \frac{D_r}{N_r}$,

- If we set tol large in the cosine similarity measure, then we expect to have high P but low R.
- On the other hand, if we set tol small, the situation is the other way around.
- Unfortunately, in the NIPS dataset, there is no information on the documents except those terms used in them. Hence, we cannot really compute "the Recall vs Precision plot" like those in the textbook.

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- Let $A \in \mathbb{R}^{m \times n}$ be a term-document matrix, and let $A_k := U_k \Sigma_k V_k^{\mathsf{T}}$ be the rank k approximation of A using the first k singular values and singular vectors. Let $H_k := \Sigma_k V_k^{\mathsf{T}}$, i.e., $A_k = U_k H_k$.
- For an appropriate value of k, $A \approx A_k$. Hence, we have $a_j \approx U_k h_j$ where a_i and h_i are the jth column vectors of A and H_k , respectively.
- This means that h_j are the expansion coefficients of the best k-term approximation to a_i w.r.t. the ONB vectors $\{u_1, ..., u_k\}$.
- Previously, for a given query vector \mathbf{q} , in order to compute the cosine similarities between \mathbf{q} and \mathbf{a}_j , j=1:n, we had to compute \mathbf{q}^TA followed by the normalization by $\|\mathbf{q}\|_2$ and $\|\mathbf{a}_j\|$.
- Now, let's replace A by its best k-term approximation A_k , i.e., we compute: $\mathbf{q}^{\mathsf{T}}A_k = \mathbf{q}^{\mathsf{T}}U_kH_k = (U_k^{\mathsf{T}}\mathbf{q})^{\mathsf{T}}H_k$.
- Hence, we can simplify the cosine similarity computation as follows

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- The reason why we formed H_k and q_k is that there is no need to explicitly compute and store A_k once we have H_k and \mathbf{q}_k . Directly dealing with A_k by computing and storing it is wasteful and time-consuming particularly for a large A.

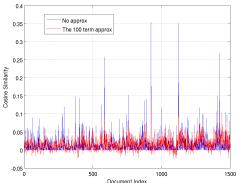
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- Then, the relative error of A_{100} and A in terms of the Frobenius norm, i.e., $||A A_{100}||_F / ||A||_F$ was 0.6074, which is still large.
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With the best 100 term approximation, tol=0.2, 0.1, 0.05 correspond to 0, 4, 72 returned documents; Compare with the no approximation case: 4, 15, 89.

- We know that the vector u_1 is the most dominant basis vector representing the range of the term space (i.e., the column space of A).
- Hence it is of our interest to check what terms u_1 represents (note that the entries of u_1 are nonnegative for this matrix). The 10 terms corresponding to the largest entries of u_1 : 'network', 'model', 'learning', 'input', 'function', 'neural', 'set', 'training', 'data', 'unit'.
- Compare these with the top 10 frequently used terms: 'network', 'model', 'learning', 'function', 'input', 'neural', 'set', 'algorithm', 'system', 'data'. As you can see, they are very close.
- Let's check the entries of u_2 , which contains both positive and negative values. The top 5 positive entries of u_2 : 'network', 'unit', 'input', 'neural', 'output', while the top 5 negative entries of u_2 : 'model', 'data', 'algorithm', 'learning', 'parameter'.
- My interpretation: u_2 tries to differentiate articles related to neuroscience from those related to machine learning algorithms.

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