EECS 767

INFORMATION RETRIEVAL

InProgress Report

By: FiniteLoop Squad

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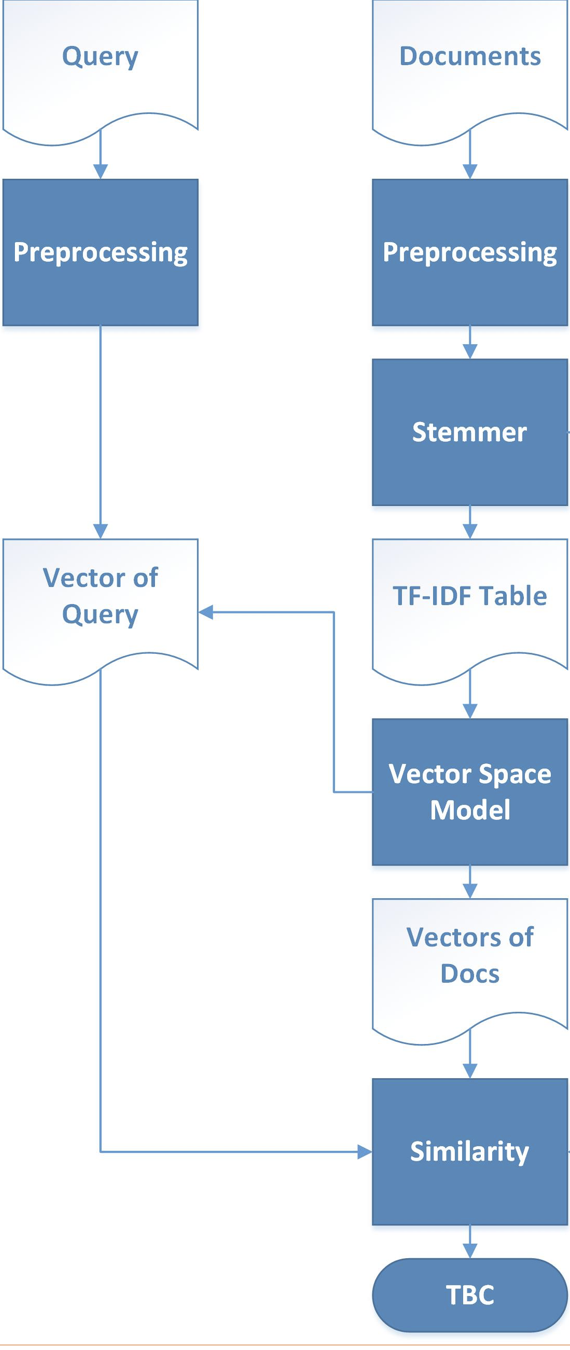
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# Introduction

The FiniteLoop Search Engine is a simple Information Retrieval System for a relatively static web page (document) repository, or corpus, using the Vector Space Model via an inverted index. In order to optimize the results and user experience, we have added techniques such as term proximity and Relevance Feedback for ranking of the results and the ability for a user to refine their search query based on the search results. The FiniteLoop Search Engine utilizes a multi-threaded niche web crawler to collect data from a specific domain and caches the documents locally for ingest and processing. In developing the search engine, we created a control group of documents, based on a quiz provided in class[[1]](#footnote-1), in order to test the ingest, processing, and query capability of the system. This report provides an explanation of the code structure, the data structures employed by the various search engine modules, and our results. The figure to the right, provides a conceptual functional (or process) flow of the search engine, post-retrieval of the documents via the niche web crawler.

# Programming Platform and Version Control Selection

In review of the various programming options available, we focused on those languages which were most capable, natively for this project. Specifically, looking at those languages which supported complex functions such as cosine similarity, web compatibility (Common Gateway Interface, CGI, or apache server module based), and of course, familiarity. After consideration of various options, such as *R*, *Perl*, *C++*, and *Python*, we selected *Python* as our language of choice.

With the current versions available for *Python*, we initially selected to go with version 3.6, being the latest available. As we worked through the various modules of our search engine (pre-processing, processing, query, and human machine interface (HMI), we ran into a few challenges. The Natural Language Toolkit (NLTK) that we selected to facilitate the stop list and lemmer was compatible with *Python* 3.5, not 3.6. Additionally, the Electrical Engineering and Computer Science (EECS) student web server currently provides access to *Python* 2.7 and 3.5. Our conclusion was to go forward with Python 2.7 as it was common to our individual environments as well as the web server. Additionally, we elected to use the EECS web server CGI capability for hosting our search engine.

For our collaboration environment, we set up a GitHub repository specifically for our **FiniteLoop** **Squad** to work and share. In the environment, we are able to coordinate our code development efforts as well as documentation.

# Data Structures

In order to pass the data structures between our modules, we are leveraging a *Python* module called *shelve*. This native module enables us to pass the raw data structures by way of a binary file stored on the file server. The following sub-sections provide the data structures passed between the modules. As a test we utilized a control group of data files, based on the class quiz covering VSM. The text data files are as follows:

test1.txt: Shipment of gold damaged in a fire

test2.txt: Delivery of silver arrived in a silver truck.

test3.txt: Shipment of gold arrived in a truck.

test4.txt: Truck arrive damaged.

# Niche Web Crawler Data Structure

The niche web crawler creates an index of the documents as it crawls and caches the web sites from the frontier. The crawler provides the following data structures to the Ingest module:

1. Download Manifest

The Download Manisfest database provides ingest a dictionary of the filenames that were crawled with a value of the URL that the file was retrieved from.

The generalized data structure look like the following, in *Python* terms:

‘’ = {

|  |  |
| --- | --- |
| DocName1: | DocURL1], |
| DocName2: | DocURL2], |
| …, |  |
| DocNamen: | DocURLn] |

}

The control group does not provide a download\_manifest, as these files are made locally available and not run through the crawler.

# Pre-Processing to Processing Interface Data Structures

Pre-Processing provides the following data structures to the Processing module:

1. Document Key Matrix
2. Term Incidence Matrix, with Frequency
3. Term Proximity Matrix
4. Title mapping
5. Number of documents

The Document Key provides the details for each document as a *Python* dictionary where the document name is the key and the details is a list of values. The details for each document include the document identifier, current location on the local filesystem (cache), and the URL [[2]](#footnote-2)of the document. The document identifier provides the sorting order of the documents and is also the index in the Term Incidence Matrix. This data structure is an *mx3* matrix, where *m* is the number of documents in the corpus.

The generalized data structure looks like the following, in *Python* terms:

doc\_key = {

|  |  |  |  |
| --- | --- | --- | --- |
| DocName1: | [DocID1, | DocLocation1, | DocURL1], |
| DocName2: | [DocID2, | DocLocation2, | DocURL2], |
| …, |  |  |  |
| DocNamen: | [DocIDn, | DocLocationn, | DocURLn] |

}

As a practical example, the contents of the **doc\_key** using the control group is:

doc\_key = {

|  |  |  |  |
| --- | --- | --- | --- |
| 'test1.txt': | [3, | '/EECS767/FiniteLoopSE/test1.txt', | 'no\_url'], |
| 'test3.txt': | [2, | '/EECS767/FiniteLoopSE/test3.txt', | 'no\_url'], |
| 'test4.txt': | [0, | '/EECS767/FiniteLoopSE/test4.txt', | 'no\_url'], |
| 'test2.txt': | [1, | '/EECS767/FiniteLoopSE/test2.txt', | 'no\_url'] |

}

The Term Incidence Matrix provides each term and its occurrence in the corpus as a *Python* dictionary[[3]](#footnote-3) where the term is the key and the document incidence with frequency is a list. Each document incidence list is aligned in order with the list provided in the doc\_key data structure. This data structure is an *nxm* matrix, where *n* is the number of terms and *m* is the number of documents in the corpus.

The generalized data structure looks like the following, in *Python* terms:

index = [

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| { Term1: | [tf1, | tf2, | …, | tfn] }, |
| { Term2: | [tf1, | tf2, | …, | tfn] }, |
| …, |  |  |  |  |
| { Termm: | [tf1, | tf2, | …, | tfn] } |

]

As a practical example, the contents of the **index** using the control group is:

index = {

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| 'truck': | [1, | 1, | 1, | 0], |
| 'arriv': | [1, | 1, | 1, | 0], |
| 'damag': | [1, | 0, | 0, | 1], |
| 'fire': | [0, | 0, | 0, | 1], |
| 'silver': | [0, | 2, | 0, | 0], |
| 'gold': | [0, | 0, | 1, | 1], |
| 'deliveri': | [0, | 1, | 0, | 0], |
| 'shipment': | [0, | 0, | 1, | 1] |

}

The Term Proximity Matrix provides a dictionary of each term, as the key, and a list of tuples as the value. The tuples identify the document and offset from the beginning of the document. Offsets are based on word distance from the beginning of the document after the tokenization and stop word parsing is complete. This data structure is an *nxp* matrix, where *n* is the number of terms and *p* is the number of non-unique terms in the corpus.

The generalized data structure looks like the following, in *Python* terms:

proximity = {

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Term1: | [ [DocID, Prox], | [DocID, Prox], | … | [DocID, Prox] ], |
| Term2: | [ [DocID, Prox], | [DocID, Prox], | … | [DocID, Prox] ], |
| …, |  |  |  |  |
| Termm: | [ [DocID, Prox], | [DocID, Prox], | … | [DocID, Prox] ] |

}

As a practical example, the contents of the **proximity** using the control group is:

proximity = {

|  |  |  |  |
| --- | --- | --- | --- |
| 'truck': | [ (0, 0), | (1, 4), | (2, 3) ], |
| 'arriv': | [ (0, 1), | (1, 2), | (2, 2) ], |
| 'damag': | [ (0, 2), | (3, 2) ], |  |
| 'fire': | [ (3, 3) ], |  |  |
| 'silver': | [ (1, 1), | (1, 3) ], |  |
| 'gold': | [ (2, 1), | (3, 1) ], |  |
| 'deliveri': | [ (1, 0) ], |  |  |
| 'shipment': | [ (2, 0), | (3, 0) ] |  |

}

The title map provides a dictionary of document names and document titles, specifically for HTML pages by leveraging the contents of the title[[4]](#footnote-4). This information is metadata used to provide ‘summary’ text displayed in the results of the search.cgi script. This data structure is an *mx2* matrix, where *m* is the number of documents in the corpus.

The generalized data structure looks like the following, in *Python* terms:

title\_map = {

|  |  |
| --- | --- |
| DocName1: | DocTitle1, |
| DocName2: | DocTitle2, |
| …, |  |
| DocNamen: | DocTitlen, |

}

As the control group does not contain any html pages, a relative example is not provided and left to the reader to imagine.

The number of docs is provided to ensure that there is a check between ingest and processing that the number of documents expected is there number of documents processed. This variable is simply an integer stored in the database file:

num\_docs = x

As in the case of the control group, the database reflects:

num\_docs = 4

# Processing to Query Processing Interfaces Data Structures

Processing provides the following data structures to the Query Processing module:

1. Document Key Matrix
2. Normalized Vector Space Model (VSM) – This file is stored in an artifacts database (processingArtifacts.db) for reference and troubleshooting)
3. Term Dictionary (based on VSM)
4. Term Proximity Dictionary
5. Term IDF Dictionary

The Document Key Matrix is forwarded, unaltered from what was received from the Pre-Processing module, see Section 3.1, Pre-Processing to Processing Interface Data Structures.

The Normalized VSM provides an alphabetically sorted list of vectors (lists). Each vector is in order as identified by the Term Index Look-Up Dictionary and each vector is in order of the Document Key Matrix. The vectors provide the normalized Term Frequency – Inverted Data Frequency (TF-IDF) weight of the term for each document.

The generalized data structure looks like the following, in *Python* terms:

docVector = [

|  |  |  |  |
| --- | --- | --- | --- |
| [WT1,D1, | WT1,D2, | …, | WT1,Dn], |
| [WT2,D1, | WT2,D2, | …, | WT2,Dn], |
| … |  |  |  |
| [WTm,D1, | WTm,D2, | …, | WTm,Dn] |

]

As a practical example, the contents of the **docVector** using the control group is:

docVector = [

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| [0.356, | 0.858, | 0.0, | 0.0, | 0.0, | 0.0, | 0.0, | 0.356], |
| [0.118, | 0.0, | 0.57, | 0.0, | 0.0, | 0.0, | 0.805, | 0.118], |
| [0.27, | 0.0, | 0.0, | 0.0, | 0.65, | 0.65, | 0.0, | 0.27], |
| [0.0, | 0.378, | 0.0, | 0.755, | 0.378, | 0.378, | 0.0, | 0.0] |

]

The docVector, is used to create a look up table for the query module (evoked by search.cgi) to quickly locate terms from the query and perform the cosine similarity process. This resulting data structure is a dictionary of terms such that each term contains the document arrays of normalized weights for that term. This is a reduced set, as there are many 0 weights in the VSM (very sparse) – in order to reduce overhead and look up times, we created a term Dictionary to house tuples of weights and document IDs.

The generalized data structure looks like the following, in *Python* terms:

termDict = {

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Term1: [ | [DocID, Weight], | [DocID, Weight], | … | [DocID, Weight] ], |
| Term2: [ | [DocID, Weight], | [DocID, Weight], | … | [DocID, Weight] ], |
| …, |  |  |  |  |
| Termm: [ | [DocID, Weight], | [DocID, Weight], | … | [DocID, Weight] ] |

}

As a practical example, the contents of the **termDict** using the control group is:

termDict = {

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| 'silver': [ | [1, 0.805] ], |  |  |  |
| 'truck': [ | [0, 0.356], | [1, 0.118], | [2, 0.27] ], |  |
| 'arriv': [ | [0, 0.356], | [1, 0.118], | [2, 0.27] ], |  |
| 'shipment': [ | [2, 0.65], | [3, 0.378] ], |  |  |
| 'damag': [ | [0, 0.858], | [3, 0.378] ], |  |  |
| 'deliveri': [ | [1, 0.57] ], |  |  |  |
| 'gold': [ | [2, 0.65], | [3, 0.378] ], |  |  |
| 'fire': [ | [3, 0.755] ] |  |  |  |

}

The Term Proximity Dictionary provides a listing similar to the Term Dictionary where instead of the weights, it provides a list of the proximities. Specifically, this data structure is constructed for ‘easy’ look up by using the term to find the document ID, to get the proximities.

The generalized data structure looks like the following, in *Python* terms:

proxDict = {

|  |  |  |  |
| --- | --- | --- | --- |
| Term1: { | { DocID: [Prox’s] }, | … | { DocID: [Prox’s] } }, |
| Term2: [ | { DocID: [Prox’s] }, | … | { DocID: [Prox’s] } }, |
| …, |  |  |  |
| Termm: [ | { DocID: [Prox’s] }, | … | { DocID: [Prox’s] } } |

}

As a practical example, the contents of the **proxDict** using the control group is:

proxDict = {

|  |  |  |  |
| --- | --- | --- | --- |
| 'silver': { | 1: [1, 3] }, |  |  |
| 'truck': { | 0: [0], | 1: [4], | 2: [3] }, |
| 'arriv': { | 0: [1], | 1: [2], | 2: [2] }, |
| 'shipment': { | 2: [0], | 3: [0] }, |  |
| 'damag': { | 0: [2], | 3: [2] }, |  |
| 'deliveri': { | 1: [0] }, |  |  |
| 'gold': { | 2: [1], | 3: [1] }, |  |
| 'fire': { | 3: [3] } |  |  |

}

The Term IDF Dictionary provides a dictionary of each term, as the key, and the term IDF weight as well as the index of the term in the VSM (stored in the processingArtifacts). This data structure is used by query to quickly assign IDF values to the query terms.

The generalized data structure looks like the following, in *Python* terms:

termIDF = {

|  |  |
| --- | --- |
| Term1: | [ IDF, Index ], |
| Term2: | [ IDF, Index ], |
| …, |  |
| Termm: | [ IDF, Index ], |

}

As a practical example, the contents of the **termIDF** using the control group is:

termIDF = {

|  |  |
| --- | --- |
| 'silver': | [0.602, 6], |
| 'truck': | [0.125, 7], |
| 'arriv': | [0.125, 0], |
| 'shipment': | [0.301, 5], |
| 'damag': | [0.301, 1], |
| 'deliveri': | [0.602, 2], |
| 'gold': | [0.301, 4], |
| 'fire': | [0.602, 3] |

}

# HTML to Query Processing Interfaces Data Structures

The HTML module provides the following data structure to the Query Processing module:

1. HTML Text

The HTML Text dictionary provides a dictionary of each document, as the key to dictionaries of terms and summary text. This was created due to the large processing time for accessing the information. Each summarized file in the corpus results in a separate summary file. The summary file is the name of the document (from doc\_key) with a ‘.db’ suffix.

The generalized data structure looks like the following, in *Python* terms

htmlText = {

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| DocName1: | { Term1: Text, | Term2: Text, | …, | Termn: Text } |
| DocName2: | { Term1: Text, | Term2: Text, | …, | Termn: Text } |
| …, |  |  |  |  |
| DocNamen: | { Term1: Text, | Term2: Text, | …, | Termn: Text } |

}

As a practical example, the contents of the **htmlText** using the control group (for test1.txt) is:

htmlText = {

|  |  |  |
| --- | --- | --- |
| ‘test1.txt’: | { ‘fire’: gold damaged in a fire’, | ‘damaged’: ‘Shipment of gold damaged in a fire’, |
|  | 'gold': 'Shipment of gold damaged in a fire', | 'shipment': 'Shipment of gold damaged in' } |

}

# Query Processing to HMI Data Structures

The data passed to the HMI is done so directly with the CGI script importing the Query module directly for dynamic processing and results. The Query module provides the following data and data structures to the HMI for display to the user:

* List of results, ordered by Rank
  + Rank by default is cosine similarity based on TF-IDF weights
  + Rank by proximity
  + Rank (and results) based on user Relevance Feedback

The list of results is provided to the HMI as a list of document entries. Each entry contains a list of parameters to display to the user. Additionally, the rank of the results is also provided.

The generalized data structure looks like the following, in *Python* terms:

rankedOutput = [

|  |  |  |  |
| --- | --- | --- | --- |
| [ { DocName1: | [ DocLocation1, | Order1, | Summary1 ] }, |
| { DocName2: | [ DocLocation2, | Order2, | Summary2 ] }, |
| …, |  |  |  |
| { DocNamen: | [ DocLocationn, | Ordern, | Summaryn ] } ], |
| [ Rank1, | Rank2, | …, | Rankn ] |

]

As a practical example, the contents of the **rankedOutput** using the control group with a random query may result in:

rankedOutput = [

|  |  |  |  |
| --- | --- | --- | --- |
| [ { 'test1.txt': | [0, | '/EECS767/FiniteLoopSE/test1.txt', | 'no\_url'] }, |
| { 'test2.txt': | [3, | '/EECS767/FiniteLoopSE/test2.txt', | 'no\_url'] }, |
| {'test3.txt': | [1, | '/EECS767/FiniteLoopSE/test3.txt', | 'no\_url'] }, |
| {'test4.txt': | [2, | '/EECS767/FiniteLoopSE/test4.txt', | 'no\_url'] } ] |
| [0.5968, | 0.5475, | 0.26, | 0.0972] |

]

The proximity scores are calculated concurrently and provided in an identical data structure as the rankedOutput data structure. With the rank and subsequent order based on the proximity relationships.

As an optimization, the query module also provides the queryVector generated – this is passed to the website and back for relevance feedback in order to alleviate the processing overhead of calculating the vector again.

The generalized data structure looks like the following, in *Python* terms:

queryVector = [

|  |  |  |  |
| --- | --- | --- | --- |
| WeightT1, | WeightT2, | …, | WeightTn, |

]

As a practical example, the contents of the **rankedOutput** using the control group with a random query may result in:

queryVector = [

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| 0.707107, | 0.0, | 0.0, | 0.0, | 0.0, | 0.0, | 0.0, | 0.707107 |

]

# Niche Crawler

<BLAKE>

# Ingest

Ingestion is performed by functions contained within the ingest.py file which operate on files stored within a local directory. The path to the directory is currently configured with a static path hard coded within the ingestion function. A function was created to allow for user entry of a different path location, however this functionality is currently deemed unnecessary and has been disabled via inline comments.

The native python library “urllib2” is used to read files and provide compatibility for processing various document formats including html. Each document within the specified directory is loaded into memory as a stream of raw characters and stored in an array, called “data,” with each document representing a single index within the array.

A dictionary called “doc\_key” is created to store the filename, document ID and file path for each document ingested. The document filename is used as the key for the dictionary while the value for each key is represented as an array containing the document ID and file path.

# Preprocessing

# Tokenization

Each document, stored as an index within the data array created during ingestion, is processed by the function “func\_tokenize” within the ingest.py file. The Python Natural Language Tool Kit (NLTK) (<https://www.nltk.org/>) is used to create stop word lists and a stemming function within func\_tokenize. This requires the installation of NLTK as well as downloading the stop word list prior to running the ingestion program. NLTK version 3.2.5 (the latest version as of this report) requires either Python version 2.7 or 3.5. Python 3.5 is not the most current release of Python 3 and may require some systems to install an earlier version of Python.

Preprocessing within func\_tokenize consists of 4 steps executed on each index of the data array:

1. HTML tags are removed from the data stream via regular expression pattern matching.
2. The data stream is converted to lower case, punctuation is removed, and the stream is split into tokens via the Python string.split() method.
3. Stop words contained within the NLTK stop word list are removed from the list of tokens.
4. Each token within the data stream is processed by the Porter stemmer provided by the NLTK library. There is a known issue when processing

Unicode characters with the NLTK Porter stemmer. Because of this, words containing Unicode characters are currently dropped in the stemming phase.

# Indexing

The number of documents processed during ingestion is used to determine the dimensionality of arrays within the terms dictionary. Each data stream, corresponding with a separate document, is parsed for unique terms. If a new term is discovered in a document, the term is added as a key to the dictionary called “terms” with an initial value of an array with multiple indices corresponding with the number of documents processed. The index corresponding to the document, wherein the term was observed, is then incremented by 1. Subsequent observations of the term within the same document, or future documents, results in incrementing the value stored in the term array at the index corresponding with the id of the document being parsed. This results in an array reflecting the term frequency for each term observed within each document.

Concurrently, an additional dictionary called “proximity” is created to record positional data pertaining to each occurrence of a term within the documents parsed. Similar to the “terms” dictionary, unique terms are used as key values in the dictionary. Values within the “proximity” dictionary are represented as an array containing tuples consisting of the document ID and the position the term appeared within the document. A new tuple representing the document id and term position is appended to the term array each time the term is observed within the data stream. This data structure is described in further detail within the “Data Structures” section of this report.

Finally, data is exported to a file called “ingestOutput.db” using the Python shelve library. The “terms” dictionary is exported as “index”, the “doc\_key” dictionary is exported as “doc\_key” and the “proximity” dictionary is exported as “proximity” within the output file.

All of these functions, including the ingest function are run currently out of the ingest.py module.

**Performance**

* Completes ingest and pre-processing of control group in 0.1131 seconds
* Completes ingest and pre-processing of 169 documents (10min web crawl) in XX seconds

# Processing

The processing module reads in the output file from the ingest function in order to acquire the pre-processing data structures. It then generates the TF-IDF, normalizes the vectors and stores the VSM for the query module. Additionally, processing generates a proximity matrix, similar to the VSM data structure and also stores it, along with the doc\_key structure into the output shelve file for use by the query function.

To accomplish this, the index data structure (the term incidence matrix) is sorted and then walked through to evaluate for the Document Frequencies of each term in order to generate the VSM. It does so by first calculating the non-zero indices in the document arrays along with calculating the idf (log n/df), where n is the length of the array (# of documents). With this data, the non-normalized weights are calculated, |Wi| = sqrt(sum(idf2)), and then added to the VSM (docVector). Finally, the module goes through the VSM and normalizes the document vectors and stores the data structure in the shelve output file for the query function.

The proximity file provided by ingest contains a dictionary of tuples where each tuple indicates the term and offset.

The performance for the Processing module is fairly straightforward as the core of the module to massage the *nxm[[5]](#footnote-5)* matrix, into a TF-IDF inverted index – lending itself to O(*nm*) complexity. The performance of this algorithm is dwarfed by the restructuring of the proximity matrix as it is dominated by the number of non-unique terms tracked in the corpus, where the complexity is O(*mp*)[[6]](#footnote-6).

**Performance**

* Completes processing of control group in 0.0736 seconds
* Completes processing of 169 documents (10min web crawl) in XX seconds

# HTML Summaries

Processing the corpus to gain summary text for all of the unique words that appear in a document is a computationally non-trivial feat, though a very simple one to author. Our HTML summary generator goes through every document in the corpus and:

* Performs some basic pruning of the HTML and elimination of the stop words
* Collects all of the unique terms in the document
* Searches through the text for each term individually
* Grabs the words before and after the term for every occurrence in the document (resulting in a list of phrases)
* Stores a randomly selected phrase in a database file dictionary, with the term as the key

The results for each document is stored in a separate file in the cache folder to be accessed by the search.cgi module while parsing results from the query module. This enables the search.cgi module to grab the data structure unique to the result and find a single phrase for each term in the query.

Due to the nature of the brutish script, it is very slow and methodical, running at O(*mp2*), where *m* is the number of documents in the corpus and *p* is the number of terms in the document. Unfortunately, not having a distributed environment available to run the beast – where each node would work on a select set in the corpus, we utilized a manual process of executing multiprocessing on a single server. This was accomplished by using ‘screen’ and spawning multiple instances of the python process to execute multiple sessions on select groupings of the corpus to accomplish the processing. Since this is working with a static repository, the only added reduction in the processing was to not re-process an already summarized document.

**Performance**

* The amount of time to consume, parse, and generate a summary file varies based on the number of terms (size) of a given document. Timing for this has been, on average, between 5 and 10 minutes

# Query

The dictionaries “doc\_vector” and “doc\_key” are passed to query module using a file processingOutput.db which is Python shelve library. “doc\_vector” dictionary provides the alphabetically sorted list of vectors (lists). Each vector is in order as identified by the Term Index Look-Up Dictionary and each vector is in order of the Document Key Matrix. The vectors provide the normalized Term Frequency – Inverted Data Frequency (TF-IDF) weight of the term for each document. Weights in this dictionary are compared with another vector “query\_vector”.

Search query inputted by user is first preprocessed by removing the stop words and processing by the Porter Stemmer provided by NLTK library. Once processed, query tokens are stored in the query vector, whose weights are compared with the document vectors to calculate the cosine similarity. The cosine similarities are calculated using the method similarity() which sorts the similarity, and ranks the results in descending order on the basis of doc\_key associated with each document vector. The ranks are stored in queryOutput.db, which would be passed to CGI to display the results.

Current work which is going on includes getting the proxVector from processing module, and re-rank the top 10 documents obtained from cosine similarity, on the basis of proximities.

**Performance**

* Completes a simple (one term) search of control group in 0.0736 seconds
* Completes a simple (one term) search of 169 documents (10min web crawl) in XX seconds

# Search

The search Common Gateway Interface (CGI) pulls together the user interaction with the results of crawling, ingesting, processing, and active query of the corpus. The execution of the script is quite straightforward:

* Take input from the user
* Pass it to the query module
* Parse the results – searching the summary databases for each result to compose the summary line
* Display to the user

The interface includes an option for the user to incorporate proximity in the search. For each result, the user has an option to pass a specific result back to the engine to perform relevance feedback.

1. EECS 767, Information Retrieval, Spring 2018 [↑](#footnote-ref-1)
2. For instances where a URL isn’t available, such as for local test files, a default value of ‘*no url*’ is used [↑](#footnote-ref-2)
3. Note, *python* dictionaries are not sorted, the order is not guaranteed [↑](#footnote-ref-3)
4. Extracts text between <title>some text</title> [↑](#footnote-ref-4)
5. Where *n* is the number of unique terms and *m* is the number of documents [↑](#footnote-ref-5)
6. Where p is count of total (non-unique) terms tracked in corpus [↑](#footnote-ref-6)