# Faculty of Computer Science, Dalhousie University *23-Sep-2024* CSCI 4152/6509 — Natural Language Processing

# Lecture 6: Counting N-grams

Location: Carleton Tupper Building Theatre C Instructor: Vlado Keselj Time: 16:05 – 17:25

## Previous Lecture

- Regular expressions in Perl
	- Use of special variables
	- Backreferences, shortest match
- Text processing examples
	- tokenization
	- counting letters
- Elements of Morphology

Lemmatization is a word processing method in which a *surface word form*, i.e., the word form as it appears in text, is mapped to its *lemma*, i.e., the canonical form as it appears in a dictionary. For example, the word *working* would be mapped into the verb *work*, or the word *semantically* would be mapped to the lemma *semantics.*

## 6.1 Morphological Processes

A *morphological process* is a word transformation that happens as a regular language transformation. There are tree main morphological processes in English:

- 1. inflection,
- 2. derivation, and
- 3. compounding.

1. Inflection: is a transformation that transforms a word from one lexical class into another related word in the same class. The transformation is performed by adding or changing a suffix or prefix. It is highly regular transformation. Some inflection examples are:  $\text{dog} \rightarrow \text{dog}$ , work  $\rightarrow$  works, work  $\rightarrow$  working, and work  $\rightarrow$  worked.

We will discuss more the concept of *lexical class* or *part of speech* class later, but for now you are probably familiar with the following lexical classes (or types of words): nouns, verbs, adjectives, adverbs, and maybe some other.

Inflection is so regular transformation that usually we do not find inflected variations of a word in a dictionary. It is assumed that a reader of the dictionary will be able to derive these variations by herself. Similarly, we can frequently program inflection in a computer application rather than storing different variations of the word.

2. Derivation: is a transformation that transforms a word from one lexical class into a related word in a different class. Similarly to inflection, it is performed by adding or changing a suffix or prefix. There is also some regularity, but it is less regular than inflection. For example, a derivation is *wide (adjective)* → *widely (adverb)*, but a similar transformation  $\frac{old \rightarrow \text{oldly}}{s}$  not valid. Some other examples are: accept (verb)  $\rightarrow$  acceptable (adjective), acceptable (adjective)  $\rightarrow$  acceptably (adverb), and teach (verb)  $\rightarrow$  teacher (noun).

There are exceptions where a derivation is used to transform a word in a lexical class to another word in the same class but it is a significantly a different word. For example, the transformation of the adjective *red* to *redish* is considered a derivation, rather than an inflection.

Since derivation is not as regular transformation as inflection, derived variations of a word are usually stored in a dictionary, and in a computer application we may want to store them in a lexicon, i.e., a word database, in many cases.



Below you can find a table with some more derivation examples:

3. Compounding: is a transformation where two or more words are combined, usually by concatenation, to create a new word. Some examples are: news + group  $\rightarrow$  newsgroup, down + market  $\rightarrow$  downmarket, over + take  $\rightarrow$ overtake, play + ground  $\rightarrow$  playground, and lady + bug  $\rightarrow$  ladybug.

# 7 Characters, Words, and N-grams

#### *Slide notes:*

## Characters, Words, N-grams

- We saw some experiments with counting characters
- Let us look at Counting Words
- N-grams and Counting N-grams

## 7.1 Counting Words and Zipf's Law



. . .

. . .

One of the basic tasks that we can do using stream-oriented processing of language is to collect statistical values on letters, words, sentences, or similar tokens. We saw previously the code for finding frequency of different letters, and these data can be useful for example for computer identification of a natural language. We can do similar counting but this time of word frequencies. The table above shows the frequencies of words in the novel "Tom Sawyer" by Mark Twain.

Zipf's law is an observation that the product of rank and frequency of the words in a text is "quite constant," if we can use that term. For example, we can test this "law" on the words in the "Tom Sawyer" novel using the following code:

## Counting Words

```
#!/usr/bin/perl
# word-frequency.pl
while (\langle \rangle) {
  while (/'?[a-zA-Z]+/g) { $f{$&}++; $tot++; }
}
print "rank f f(norm) word r*f\n".
      (' – ' x35). "\n";
for (sort { $f{$b} <=> $f{$a} } keys $f) {
  print sprintf("%3d. %4d %lf %-8s %5d\n",
             ++$rank, $f{$_}, $f{$_}/$tot, $_,
             $rank*$f({$_!}\);
}
```
## Program Output (Zipf's Law)



We can present this data in a graphical form and compare it with the function  $f = 10000/r$  to demonstrate the



## Zipf's law:

If we apply a logarithm on both sides of the Zipf's formula we get the formula  $\log r + \log f \approx \text{const.}$ , which means that the Zipf's law implies that the rank-frequency graph using log scales of  $x$  and  $y$  axis should be close to a straight line, descending under an angle of 45 degrees. The following graph illustrates this:



## 7.2 Counting N-grams

Given a sequence of tokens  $T = (t_1, t_2, t_3, \ldots, t_k)$  an n-gram is an arbitrary subsequence of n such tokens, such as  $(t_1, t_2, \ldots, t_n)$ , or  $(t_2, t_3, \ldots, t_{n+1})$ , and so on. Given a textual document, it could be broken into a list of character or list of words, if we assume that a character is a token, or that a word is a token. In those cases, we look at character n-grams or word n-grams, respectively. We typically want to collect all n-grams from a text, and a way to visualize this is to imagine a sliding window over the text.

#### Character N-grams

## – Consider the text:

- The Adventures of Tom Sawyer
- Character n-grams = substring of length  $n$
- $n = 1 \Rightarrow *unigrams*: T, h, e, (space), A, d, v, ...$
- $n = 2 \Rightarrow bigrams:$  Th, he, e<sub>,</sub> \_A, Ad, dv, ve, ...
- $n = 3 \Rightarrow trigrams:$  The, he\_, e\_A, \_Ad, Adv, dve, ...
- and so on; Similarly, we can have word n-grams, such as  $(n = 3)$ : The Adventures of, Adventures of Tom, of Tom Sawyer...
- or normalized into lowercase

For example, if we take another look at the Tom Sawyer novel:

The Adventures of Tom Sawyer

by

Mark Twain (Samuel Langhorne Clemens)

#### Preface

MOST of the adventures recorded in this book really occurred; one or two were experiences of my own, the rest those of boys who were schoolmates of mine. Huck Finn is drawn from life; Tom Sawyer also, but not from an individual -- he is a combination of the characteristics of three boys whom I knew, and therefore belongs to the composite order of architecture.

The odd superstitions touched upon were all prevalent among children and slaves in the West at the period of this story -- that is to say, thirty or forty years ago.

Although my book is intended mainly for the entertainment of boys and girls, I hope it will not be shunned by men and women on that account, for part of my plan has been to try to pleasantly remind adults of what they once were themselves, and of how they felt and thought and talked, and what queer enterprises they sometimes engaged in. ...

## Word and Character N-grams  $(n = 3)$





#### A Program to Extract Word N-grams

The following Perl program word-ngrams.pl lists all word ngrams extracted from the standard input. We set variable \$n to 3 as we want word 3-grams to be extracted. The first while-loop reads input line by line, and in the second while-loop we match string that we assume would be words. We choose any sequence of letters to be a word, and possibly starting with an apostrophe ('). As we will see later, this will recognize usual words, but it will also break complex words like  $I'm$ , you're, or man's into words  $I$  and 'm, you and 're, and man and 's.

```
#!/usr/bin/perl
# word-ngrams.pl
$n = 3;while (\langle \rangle) {
  while \frac{1}{2} (/'?[a-zA-Z]+/q) {
     push @nq, lc(§&;; shift @nq if scalar(@nq) > $n;print "@ng\n" if scalar(@ng) == \varsigman;
  }
}
# Output of: ./word-ngrams.pl TomSawyer.txt
# the adventures of
# adventures of tom
# \dots
```
#### Some Perl List Operators

- push  $\mathfrak{g}_a$ , 1, 2, 3; adding elements at the end
- $-$  pop  $@a$ ; removing elements from the end
- $-$  shift  $@a;$  removing elements from the start
- unshift  $\mathfrak{g}_a$ , 1, 2, 3; adding elements at the start
- $-$  scalar( $@a$ ) number of elements in the array
- $-$  \$#a last index of an array, by default \$#a = scalar (@a) 1
- To be more precise, this is always true:  $\text{scalar}(\mathcal{C}a) = \text{A}^*a \text{A}^*a + 1$
- $-$  \$ [ (by default 0) is the index of first element of an array
- Arrays are dynamic: examples:  $\frac{2}{3}$  [5] = 1,  $\frac{2}{3}$  #a = 5,  $\frac{2}{3}$  #a = -1

Since the first element of an array  $\mathcal{Q}$  a is  $\mathcal{S}$  a  $[0]$  and the last element is  $\mathcal{S}$  a  $[\mathcal{S}$ #a] the number of elements is obviously  $\frac{1}{2}$   $\frac{1}{2}$ . Another way to obtain the number of elements of an array is scalar ( $\frac{2}{3}$ ). The Perl function scalar enforces a scalar context on an expression, and in a scalar context an array is interpreted just a number representing its length.

Perl arrays are dynamic, they expand and also can shrink easily. For example, after the command ' $\beta a$ [5] = 1' the array  $@a$  will be expanded if needed to at least six elements. The command ' $$4a = 5$ ' sets array  $@a$  to exactly six elements. Similarly, ' $\frac{2}{4}a = -1$ ' erases an array by reducing it to zero elements, so it is equivalent to the command ' $@a = ()$ ;'.

#### Extracting Character N-grams (attempt 1)

#!/usr/bin/perl

```
# char-ngrams1.pl - first attempt
sn = 3;while (\langle \rangle) {
 while (\frac{\sqrt{S}}{g}) {
     push @ng, $&; shift @ng if scalar(@ng) > $n;
     print "@ng\n" if scalar(@ng) == $n;
  }
}
# Output of: ./char-ngrams1.pl TomSawyer.txt
# T h e A d v e n t
# h e A d v e n t u
# e A d v e n ...
```
## Extracting Character N-grams (attempt 2)

```
#!/usr/bin/perl
# char-ngrams2.pl - second attempt
$n = 3;while (\langle \rangle) {
 while (\frac{\sqrt{s}}{s+\sqrt{g}}) {
   my $token = $&;
   if ($token = \sqrt{\frac{s+1}{1}} { $token = \sqrt{\frac{s+1}{1}}push @ng, $token;
   shift @nq if scalar(@nq) > \$n;
   print "@ng\n" if scalar(@ng) == $n;
 }
}
# Output of: ./char-ngrams2.pl TomSawyer.txt
# _ T h f _ T _ _ _
# T h e \_ T o \_ \_ M
# h e Tom _M a\# e A om \_ ...
# _ A d m _ S This may be what we want, but
# A d v _ S a probably not.
# d v e S a w
# v e n a w y
# e n t w y e
# n t u y e r
# t u r e r _
# u r e 1 r _ _
# r e s - -# e s _ - _ b
# s \_ o \_ b \ y# _ o f b y _
# o f _ y _
```
This output may be what we want, but probably not. Since we already reduced repeated whitespace characters to one underscore ('\_'), we probably want to treat the new line in the same way.

An easy way to solve the problem is to treat the whole file as one line:

#### Extracting Character N-grams (attempt 3)

```
#!/usr/bin/perl
# char-ngrams3.pl - third attempt
Sn = 3:\zeta = join('', <>); # notice how <> behaves differently
              # in an array context, vs. scalar context
while (\frac{\sqrt{s}}{s+\sqrt{q}}) {
  my $token = $&;if ($token = \sqrt{\frac{s+5}{1}} { $token = \frac{s-1}{1} }
  push @ng, $token;
  shift @ng if scalar(@ng) > $n;
  print "@ng\n\in 'f scalar(@ng) == $n;}
# Output of: ./char-ngrams3.pl TomSawyer.txt
# _ T h f _ T a r k
# T h e _ T o r k _
# h e _ T o m k _ T
\begin{array}{cccccccccccccc} \# & \texttt{e} & \_\texttt{A} & \texttt{o} & \texttt{m} & \_\texttt{m} & \_\texttt{T} & \texttt{w} \end{array}# _ A d m _ S T w a
# A d v _ S a w a i
# d v e S a w a i n
# v e n a w y i n _
# e n t w y e n _ (
# n t u y e r _ ( S
# t u r e r _ ( S a
# u r e r _ b S a m
# r e s _ b y a m u
\# e s by mue
\# s _ o y _ M u e l
# _ o f _ M a e l _
# o f _ M a r ...
```
These days computers have very large working memories (RAM, or Random Access Memories), so reading a whole file in memory as in the above example is normally not a problem. If we want to avoid reading the whole file into memory and still recognize multi-line whitespace as one space character, it can be done but we will leave it for reader as an exercise. One approach would be to write a function next\_char that keeps the current line and on each call reads the next character and returns it. When encoutering a whitespace character, it would read as many lines as needed until a non-whitespace character is found, and it would return a space.

#### Extracting Character N-grams by Line

- We need to handle whitespace spanning multiple line
- Generally, any token may span multiple lines
- Could be done but leads to a bit more complex code

If the files are very large, we may not want to read the whole file in memory and still want to handle multi-line whitespace. This problem may happen in general if any tokens that are part of n-grams span multiple lines. Again, the issue can handled in a brief way if we allow reading the whole file. We want to read the file line by line, without accumulating too many lines at a time, we would need to identify when a token may be spanning multiple lines and only then read lines ahead.

## Word N-gram Frequencies

```
#!/usr/bin/perl
# word-ngrams-f.pl
$n = 3;while (\langle \rangle) {
  while \frac{1}{2} [a - zA - z] + q {
     push \ellng, lc(\frac{5}{6}); shift \ellng if scalar(\ellng) > \frac{5}{6}n;
     \&collect(\Theta ng) if scalar(\Theta ng) == \sin;
  }
}
sub collect {
 my $ng = "@";
  $f{$ng}++; ++$tot;
}
print "Total $n-grams: $tot\n";
for (sort { $f{$b} <=> $f{$a} } keys $f) {
 print sprintf("%5d %lf %s\n",
                $f{$_}, $f{$_}/$tot, $_);
}
# Output of: ./word-ngrams-f.pl TomSawyer.txt
# Total 3-grams: 73522
# 70 0.000952 i don 't
# 44 0.000598 there was a
# 35 0.000476 don 't you
# 32 0.000435 by and by
# 25 0.000340 there was no
# 25 0.000340 don 't know
# 24 0.000326 it ain 't
# 22 0.000299 out of the
# 22 0.000299 i won 't
# 21 0.000286 it 's a
# 21 0.000286 i didn 't
# 21 0.000286 i can 't
# 20 0.000272 it was a
# 19 0.000258 and i 'll
# 18 0.000245 injun joe 's
# 18 0.000245 you don 't
# 17 0.000231 i ain 't
# 17 0.000231 he did not
# 16 0.000218 he had been
```

```
# 15 0.000204 out of his
# 15 0.000204 all the time
# 15 0.000204 it 's all
# 15 0.000204 to be a
# 15 0.000204 what 's the
# 14 0.000190 that 's so
\# \ldots
```
## Character N-gram Frequencies

```
#!/usr/bin/perl
# char-ngrams-f.pl
$n = 3;\zeta = join('', <>); # notice how <> behaves differently
             # in an array context, vs. scalar context
while (\frac{\sqrt{s}}{s+\sqrt{g}}) {
 my $token = $&;
  if ($token = \sqrt{\frac{s+1}{1}}) { $token = \sqrt{\frac{s+1}{1}}push @ng, $token;
  shift @nq if scalar(@nq) > \$n;
  \&collect(\Theta ng) if scalar(\Theta ng) == \sin;
}
sub collect {
 my $ng = "@";
 $f{$ng} ++; ++$tot;
}
print "Total $n-grams: $tot\n";
for (sort { $f{$b} <=> $f{$a} } keys %f) {
print sprintf("%5d %lf %s\n",
                 $f{$_}, $f{$_}/$tot, $_);
}
# Output of: ./char-ngrams-f.pl TomSawyer.txt
# Total 3-grams: 389942
# 6556 0.016813 _ t h
# 5110 0.013105 t h e
# 4942 0.012674 h e _
# 3619 0.009281 n d _
# 3495 0.008963 _ a n
# 3309 0.008486 a n d
# 2747 0.007045 e d _
# 2209 0.005665 _ t o
# 2169 0.005562 i n g
# 1823 0.004675 t o _
# 1817 0.004660 n g _
# 1738 0.004457 _ a _
# 1682 0.004313 _ w a
```

```
# 1673 0.004290 _ h e
# 1672 0.004288 e r _
# 1592 0.004083 d _ t
# 1566 0.004016 _ o f
# 1541 0.003952 a s _
# 1526 0.003913 _ ' '
# 1511 0.003875 ' ' _
# 1485 0.003808 a t _
# \dots
```
## 7.3 Using Ngrams Module

This section is covered in the lab, but you can also read here about the basic use of the Ngrams module.

We will now discuss how different kinds of n-grams can be collected using a Perl module named Text::Ngrams. A program associate with this module is named ngrams.pl, and both files, Ngrams.pm and ngrams.pl, can be found in the directory ˜prof6509/public on bluenose. They can also be found on the course website under the tab 'Misc'. If you use the web-site, for technical reasons the file ngrams.pl was renamed to ngrams-pl.txt and if you download it, you will need to rename it back to ngrams.pl.

The module and the program are open-source code, and can be found in the CPAN archive. The newest version is available on bluenose. The modules are typically installed system-wide and the Perl is configured in such way that it can easily find them. Since you do not have administrative permissions on bluenose, we need to use a way to use the module locally. The Perl modules can be installed on a per-user basis, either in a more systematic way or in more ad-hoc way. We will use here a local ad-hoc installation. You will cover the steps of installing the module in more details in the lab, but for now, we will assume that you are in a convenient sub-directory of your your home directory on bluenose. You would first copy the appropriate files using the commands:

```
cp ˜prof6509/public/ngrams.pl .
cp ˜prof6509/public/Ngrams.pm .
```
These files may actually be installed system-wide on bluenose, but to be sure to use the local version, we will do a couple additional operations and checks. First, create a subdirectory Text and copy the module there:

```
mkdir Text
cp Ngrams.pm Text
```
Check Local ngrams.pl

```
• Use command: more ngrams.pl
```
Let us take a look at the version of ngrams.pl that we use here. (This version is slightly different from the version in the CPAN archive.) We can use the command 'more ngrams.pl and the beginning of the file should look as follows:

```
#!/usr/bin/perl -w
use strict;
use vars qw($VERSION);
$VERSION = 2.005;# $Revision: 1.26 $
use lib '.';
```
use Text::Ngrams;

```
use Getopt::Long;
...
```
natural language processing

The line 'use lib'.';' is important, since it directs Perl to give priority in finding the module in the current directory, rather than some other versions that may be available in the system. You can test the program ngrams.pl but typing:

```
./ngrams.pl
```
then typing some input, and pressing 'C-d'; i.e., Control-D combination of keyboard keys. For example, if you type input:

```
you should get the output:
BEGIN OUTPUT BY Text::Ngrams version 2.005
1-GRAMS (total count: 28)
FIRST N-GRAM: N
LAST N-GRAM: _
-------------------------
= 3
A 4
C 1
E 2
G 3
I 1
L 2
N 3
O 1
P 1
R 2
S 2
T 1
U 2
2-GRAMS (total count: 27)
FIRST N-GRAM: N A
LAST N-GRAM: G \_-------------------------
\_ L 1
P_1A G 1
A L 1
A N 1
A T 1
C E 1
E = 1E S 1
G \qquad 1G E 1
G U 1
```
I N 1  $L = 1$ L A 1 N A 1 N G 2 O C 1 P R 1 R A 1 R O 1 S I 1 S S 1 T U 1 U A 1 U R 1 3-GRAMS (total count: 26) FIRST N-GRAM: N A T LAST N-GRAM: N G \_ ------------------------- \_ L A 1 \_ P R 1 A G E 1 A L  $\_ 1$ A N G 1 A T U 1 C E S 1  $E$   $p$  1 E S S 1  $G E$   $1$ G U A 1 I N G 1  $L$   $\_$   $L$   $\,$   $1$ L A N 1 N A T 1  $\,$  N  $\,$  G  $\,$   $\_$   $\,$  1  $\,$ N G U 1 O C E 1 P R O 1 R A L 1 R O C 1 S I N 1 S S I 1 T U R 1 U A G 1 U R A 1

END OUTPUT BY Text::Ngrams

This are the character n-grams of up to the size 3 of the given text, with their counts.

#### Verifying Version of Ngrams.pm

To test that the program is using the correct version of the module Ngrams.pm we can edit the file Text/Ngrams.pm and temporarily insert a 'die' command at the beginning of the module. The beginning of the module should look as follows:

```
# (c) 2003-2014 Vlado Keselj http://web.cs.dal.ca/˜vlado
#
# Text::Ngrams - A Perl module for N-grams processing
die;
package Text::Ngrams;
use strict;
require Exporter;
use Carp;
...
```
- If we run ngrams.pl it should report error
- Delete 'die;' command from the Ngrams.pm file

It is important to note that this is the copy of the module in the subdirectory Text. After this small test, do not forget to remove again the line 'die;'.

## 8 Elements of Information Retrieval and Text Mining

In the previous sections, we looked at some methods for processing text in a stream mode. Many language processing tasks can be solved in this way, by using mainly regular expressions, extracting some pieces of text, and collecting basic statistics. We will now look at some techniques for working with the documents as whole units withing large collections. First we will look at the task of Information Retrieval, and then the area of Text Mining, with a particular emphasis on Text Classification and brief mentioning of Text Clustering.

The term *Text Mining* was coined at about the same time as *Data Mining*, and it consists of methods for a coarse-grained management of text documents, such as classification and clustering; but also some finer-grained mining of information, such as in information extraction.

## 8.1 Elements of Information Retrieval

– Reading: [JM] Sec 23.1, ([MS] Ch.15)

Information Retrieval is an area of Computer Science mainly concerned with the task of finding a set of relevant documents from a document collection given a user query. A search engine, such as Google, is a information retrieval system.

**Basic Information Retrieval problem definition:** The basic definition of the problem or task of Information Retrieval is also called *ad hoc retrieval* and is given as follows: We are given a set of documents called a *document collection*, where each document is a natural language text. A user has an *information need* which she or he will need to express as a *query*, which is a short text, possibly in natural language or some more specialized format. The task of an Information Retrieval system is to return a subset of documents from the document collection that are *relevant* to the user query. The relevant documents should also be sorted by relevancy, starting from the most relevant document; i.e., a *ranked list of documents*.

## Typical IR System Architecture



#### Steps in Document and Query Processing

- a "bag-of-words" model
- stop-word removal
- rare word removal (optional)
- stemming
- optional query expansion
- document indexing
- document and query representation;
	- e.g. sets (Boolean model), vectors

The document semantics is reduced to the set of stems of content-bearing words.

## 8.2 Vector Space Model

#### Vector Space Model in IR

- We choose a global set of terms  $\{t_1, t_2, \ldots, t_m\}$
- Documents and queries are represented as vectors of weights:

$$
\vec{d} = (w_{1,d}, w_{2,d}, \dots, w_{m,d}) \quad \vec{q} = (w_{1,q}, w_{2,q}, \dots, w_{m,q})
$$

where weights correspond to respective terms

- What are weights? Could be binary (1 or 0), term frequency, etc.
- A standard choice is: *tfidf* term frequency inverse document frequency weights

$$
\text{tfidf} = \text{tf} \cdot \log\left(\frac{N}{\text{df}}\right)
$$

- *tf* is frequency (count) of a term in document, which is sometimes log-ed as well
- *df* is document frequency, i.e., number of documents in the collection containing the term

After preprocessing steps, such as stop-word removal, rare words removal, and stemming, we have a global set of terms  $\{t_1, t_2, \ldots, t_m\}$ , which are used to represent documents and queries.

In a vector space model, document and queries are represented by vectors of weights, such as

$$
\vec{d} = (w_{1,d}, w_{2,d}, \dots, w_{m,d}), \text{ and } \vec{q} = (w_{1,q}, w_{2,q}, \dots, w_{m,q})
$$

where the weights  $w_{i,x}$  correspond to the term  $t_i$ , of the document or query x. There are different ways how weights can be determined. One simple way is to use *binary* weights: 1 if the document contains the term, or 0 if it does not. Another option is to use term counts, or frequency within the document or query. The most widely adopted standard choice is to use *term frequency inverse document frequency* weights (*tfidf*), which are calculated using the following formula:

$$
tfidf = tf \cdot \log\left(\frac{N}{df}\right)
$$

where *tf* is frequency (count) of a term in document, which is sometimes log-ed as well; *df* is document frequency, i.e., number of documents in the collection containing the term; and  $N$  is the total number of documents in the collection. The document frequency  $df$  is the number of documents that contain the term t. We could also calculate it as the portion of the document collection that contain the term; i.e., the fraction of documents that contain the term, which would be  $df/N$ . A term should be more important and have a higher weight if it is more rare, so that is the reason why we use the inverse document frequency, or  $N/df$ . For very rare terms this number could be very large, for example the terms with  $df = 1$  and  $N = 1000000$  it would be 1000000, so to "curb" this growth we apply the slow-growing logarithm function and finally obtain  $tf\cdot df = tf \cdot \log(N/df)$ . In some references, the logarithm is applied to  $tf$  as well.

#### 8.3 Cosine Similarity Measure

A natural measure to measure similarity between a document and a query is the *cosine similarity measure.* It is known that the cosine of the angle between two vectors can be easily computed using the following formula:

$$
sim(q, d) = \frac{\sum_{i=1}^{m} w_{i,q} w_{i,d}}{\sqrt{\sum_{i=1}^{m} w_{i,q}^{2}} \cdot \sqrt{\sum_{i=1}^{m} w_{i,d}^{2}}} = \frac{\vec{q} \cdot \vec{d}}{|\vec{q}| \cdot |\vec{d}|}
$$

The formula gives the cosine of the angle between vectors in 2-dimensional and 3-dimensional space, and although we cannot exactly image the angle in spaces in more dimensions it still preserve some nice properties that match our intuition about similarity between documents, or between a document and a query. For example, if a document and query have exactly the same terms and in exactly the same proportion of their frequencies, the angle will be 0, and the cosine will be 1. On the other hand, if and only if the query and the document have no terms in common, the angle will be  $90^\circ$ , and the cosine will be 0.

The angle between a query and a document vector in the 3-dimensional space is shown in the Figure [1.](#page-15-0)



Figure 1: Cosine Similarity in the 3-dimensional Space

<span id="page-15-0"></span>If the vectors representing documents  $(d)$  and queries  $(\vec{q})$  are normalized in advance, i.e., if they are divided with their length, then the cosine similarity computation becomes simpler and more efficient. Namely, if the normalized vectors are precomputed

$$
\vec{d_0} = \frac{\vec{d}}{|\vec{d}|} = \left(\frac{w_{1,d}}{\sqrt{\sum_{i=1}^m w_{i,d}^2}}, \frac{w_{2,d}}{\sqrt{\sum_{i=1}^m w_{i,d}^2}}, \dots, \frac{w_{m,d}}{\sqrt{\sum_{i=1}^m w_{i,d}^2}}\right)
$$

and

$$
\vec{q}_0 = \frac{\vec{q}}{|\vec{q}|} = \left(\frac{w_{1,q}}{\sqrt{\sum_{i=1}^m w_{i,q}^2}}, \frac{w_{2,q}}{\sqrt{\sum_{i=1}^m w_{i,q}^2}}, \dots, \frac{w_{m,q}}{\sqrt{\sum_{i=1}^m w_{i,q}^2}}\right)
$$

then the similarity value is simply computed as

$$
sim(q, d) = \vec{q}_0 \cdot \vec{d}_0 = \sum_{i=1}^{m} w_{iq_0} w_{id_0}
$$