Information retrieval

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1 Vector space model

Text transformation   The first step to process queries on text is to transform it into a model that can be computed by a machine.

Such a model should be suitable for any kind of text, should be able to capture the semantic of words and then should support the notion of *text similarity*.

The main idea of a vector space model is to map documents in a vector, so they appear as points in a certain space.

For instance, we can collect some words and index them, then build a vector for each document

$$d_k = [x_0, \ldots, x_n]$$

where the value x_i stores the occurrences of the word indexed with i in that document.

Similarity   This model somehow implements the notion of text similarity, represented by some sort of distance function between two documents in space, perhaps the *Euclidean distance* or the difference between the angle of the two vectors. There are a lot of distance measures, and they vary in applications. Some of them assume normalization in the document vectors.

Document-Term matrix   If we arrange document vectors in a vector we obtain the *document-term matrix*, or the *term-document matrix* if we take the opposite. The number of dimensions in the space is equal to the number of words in the vocabulary.
That can be a problem without some sort of lemmatisation and stemming, as sparsity and dimension of the matrix increases.

Vector length problem   By considering the length of the vector in a similarity measure, we are introducing some problems, let's just consider a text and its abstract, the Euclidean distance would be huge, as most of the words do not appear that often.
To solve the problem we can consider the *Cosine similarity* or normalize vector lengths.

The main phases in the generation of a vector space models are: tokenization, normalization, weight, and indexing.

1.1 Tokenization

The goal is to split the text in words, for instance we could split the document at the spaces.

In practice this can be obtained with *Regex*, *Corpus* based and *Machine Learning* based methods.

1.2 Normalization

The idea is to normalize each token, that could mean writing verbs in the normal form, or plurals at the singular.

Stemming One form of normalization is stemming, that simply truncate words to obtain singular or infinite forms. The problems here could be creation of meaningless words and also close words in meaning could be completely different. An example is the *Porter* stemming algorithm.

Lemmatisation Another way to normalize is to lemmatise, It's similar to stemming, but the words are replaced with the dictionary root, this also has a problem, indeed sometime mapping of same words to different lemmas can happen. A way to lemmatise is to use a *Dictionary based* lemmatisation, or something more complicated such as *Wordnet*.

Wordnet Wordnet is a large lexical database, verbs, nouns, adjectives and adverbs are grouped into cognitive synonyms, called synsets, each expressing a different concept. Synsets are connected to each other by conceptual relations and they are also connected to a lemma.

1.3 Term Weighting

Weighting the words is a crucial point in text processing, an easy way could be counting the frequency of the terms in the document.

Term frequency An example of weighing is the term frequency, which counts the frequency of a term given a document:

- *Boolean*: so a 0 or 1 if the word is present or not

- *Natural*: $tf_{t,d} = $ count of a specific word

- *Log*: $1 + \log(tf_{t,d})$

- *Augmented*: $0.5 + \frac{0.5 tf_{t,d}}{\max_{t'} tf_{t',d}}$

- *Log average*: $0.5 + \frac{0.5 \log(tf_{t,d})}{1 + \log(avg_{t'} tf_{t',d})}$

- *Max tf norm*: $k + (1-k)\frac{tf_{t,d}}{tf_{\max}(d)}$

Stop words problem   A problem in this model is the excessive presence of stop words and punctuations. To compensate, we can remove them in the normalization phase.

But sometimes the stop words can be important, for instance with phrasal verbs. A solution can be to count the number of documents that contains that given word.

Inverse document frequency   The idea is to count the number of documents that contains a given words, and to prefer less frequent words:

$$idf = \log \frac{N}{df_t}$$

There are many variants:

- *Smooth*: $\log(\frac{N}{1+df_t}) + 1$

- *Max*: $\log(\frac{\max_{t' \in d} df_t}{1+df_t}) + 1$

- *Probabilistic*: $\log \frac{N-df_t}{df_t}$

TfIdf   By multyplying term frequency and inverse term frequency we obtain this metric.

$$TfIdf(t,d) = tf_{t,d} \cdot idf_t$$

Regarding the value of terms:

- the terms with a high *TfIdf* usually appears in a small number of documents

- the metric is low when a term appears a few times in a document or when it appears in many documents

2   Evaluation on retrieval systems

The goal of evaluation is to assess the quality of the results obtained by an IR system. There's the need of knowing a *ground truth*, so an annotated corpus where for each task we know what documents are relevant. The annotations could be created manually or derived from the data, if it contains annotations.

2.1   The notion of quality

Given a corpus C and a query q, the task is to find a set of documents $A_{q,C}$ that match q, but after retrieving such documents, there's the need to estimate the quality of results.

Precision To formalise the quality of the retrieved documents $A_{q,C}$ we could count how many of these documents are relevant to q, this is the notion of precision:

$$Prec = \frac{relevant\ retrieved}{retrieved}$$

Note that in order to know if a document is relevant or not we need the ground truth or a user feedback.

This measure does not suffice, as for instance, if we retrieve only one correct document we would have maximum precision.

Recall The precision measure does not take in consideration how many relevant documents are there, that is crucial to assess quality of a query result. So we can consider another measure, called Recall:

$$Rec = \frac{relevan\ retrieved}{relevant}$$

Information need Given the two quality measures, should we aim at better precision or more recall? Trivially both, but it actually depends on the information need.
In some cases a really high recall is not necessary, while other times a lower precision could be accepted while not missing anything relevant.

F1 Score To take into consideration both precision and recall, we could take a weighted mean, so a tradeoff between the twos

$$F1 = \frac{2 \cdot Prec \cdot Rec}{Prec + Rec}$$

Baseline system To assess the quality of an IR system, we need a baseline system, something trivial such as tossing a coin to decide whereas a document is relevant or not. Given its quality measure, we can infer the quality of our, hopefully, more complex system, in fact, quality measures can't be seen as absolute values, there's always the need to compare.

2.2 Quality measures

To formally define the notions introduced in the previous section, we need to take into consideration a more detail measure for errors.

Confusion Matrix Given a query q, a ground truth E_q of relevant documents with respect to q, and a set of retrieved documents A_q, we define this matrix:

	$d \in E_q$	$d \notin E_q$
$d \in A_q$	True positive	False positive
$d \notin A_q$	False Negative	True Negative

So now we can redefine the *precision*, *recall* and *F1* measures as follows

$$Prec = \frac{TP}{TP + FP} \quad Rec = \frac{TP}{TP + FN} \quad F1 = \frac{TP}{TP + \frac{1}{2}(FP + FN)}$$

The actual confusion matrix is obtained by counting the documents given the retrieved ones and the ground truth.

The matrix can be useful to estimate the system parameters, for instance, if the number of retrieved documents is set to a certain value k and the value of true positives is really high, probably a smaller k would do the job, while if a lot positives are left out, i.e, the matrix has a high number of false negatives, maybe an higher k would make the system better.

Other measures The are a lot of measures to estimate the quality of a system, such as *specificity*, *negative predictive value*, *miss-rate*, *fall-out* and *accuracy*. Finding the right measure is really important, for instance, using accuracy with an unbalanced dataset is not better than precision, recall and F1.

2.3 Evaluation of ranking systems

We talked about boolean retrieval systems and we took into consideration if a document is retrieved or not.

In a raking scenario, we want to give importance not only to precision and recall, but also to the rank assigned by the system.

Setting a threshold One way to go is to set a specific threshold and compute the precision and recall of those top documents. This method does not take into consideration the ranking of the documents.

Precision at K If we take the first K retrieved documents, ordered by rank, we can estimate the quality of the rank by computing the precision for the top K documents. By iterating the threshold we can better estimate the performances, opposed to setting a unique threshold.
To decide what thresholds to use we could use the distribution of the system ranking.

Discounted cumulative gain This approach takes into consideration the ranking and discount the gain given by a document with respect to its position in it.
$$DCG = \sum_{i=1}^{n} \frac{R_j}{\log(i + 1)}$$
where R_j is the rank assigned to a document.

Precision vs Recall curve   Another approach is to compute precision and recall measures at each point in the ranking. If we take the measurements and plot them, we find a correlation between precision and recall and if we compute the integral of that function, we obtain the average precision.

To have a smoother curve, it's possible to interpolate the precision, so for each point in the recall axis, we take the maximum precision of consecutive points.
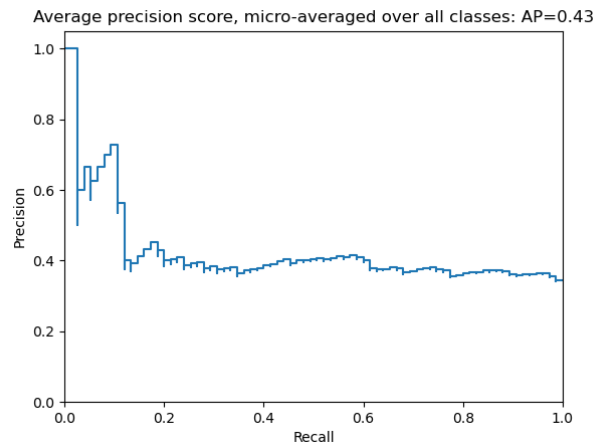

Figure 1: Example of a non-interpolated precision and recall curve

3   Relevance feedback and query expansion

The idea behind this section is the possibility to use some sort of feedbacks to tweak queries before passing them to the information retrieval system.

Relevance feedback   It's any feedback we can collect about correctness of a certain query. For instance a user feedback or a indirect feedback about usefulness of an answer, i.e. the user clicked on a particular website

Query expansion   Transform user queries exploiting the relevance feedback to improve system results over that query.

When expanding a query there are two possible methods, local and global methods, depending on whereas the expansion is applied to a single query or to all queries to the system:

- *Local methods*: related to a particular query, they are based on relevance and indirect feedback of that particular query

- *Global methods*: methods applied to all queries, those can be expansion with a thesaurus, or spelling correction

3.1 Local methods

Vector shifting   Given the sets R and N of relevant and non relevant results respectively, the task is to find a query vector $\vec{q}_o$ that maximizes similarity with R and minimizes the one with N:

$$\vec{q}_o = argmax_{\vec{q}}[sim(\vec{q}, R) - sim(\vec{q}, N)]$$

Given a similarity function, for instance the cosine similarity, the equation becomes

$$\vec{q}_o = \frac{1}{|R|} \sum_{\vec{d_i} \in R} \vec{d_i} - \frac{1}{|N|} \sum_{\vec{d_j} \in N} \vec{d_j}$$

where the two elements are the centroids of R and N.

Rocchio algorithm   This approach is a generalization of the main idea introduced in the section, it proposes a new vector derived by making the query vector closer to the centroid of relevant documents.

$$\vec{q}_m = \alpha \vec{q} + \beta \frac{1}{|R|} \sum_{\vec{d_i} \in R} \vec{d_i} - \gamma \frac{1}{|N|} \sum_{\vec{d_j} \in N} \vec{d_j}$$

where α, β and γ can balance the role of each component.

One of the problems of this approach is the difficulty in estimating a document relevance, also, the sets R and N could be somehow merged.

Pseudo-relevance feedback   Instead of collecting real feedback for a query, we assume that the top k results are correct, we then use them as a feedback for the Rocchio algorithm.

To motivate this choice we can look at the precision and recall curve, indeed a small set of results usually lead to high precision, thus we can move the system in the direction of the top results.

Indirect relevance feedback   In this case the feedback is obtained by looking at user behavior, for instance we could count the number of visits to the results to measure the popularity of documents.

3.2 Global Methods

Query expansion   The idea here is to expand the query by adding new information, this could mean adding terminology or shifting the query vector as before.

3.2.1 How to add information?

To add information to a query we can perform *syntactic analysis* of the original query, using *dictionaries* to add new terms related to the query, or use the *corpus of reference documents* to add terminology.

Dictionaries and knowledge bases The process to enrich a query when having a knowledge base, that could be a dictionary an ontology or whatever, is to:

1. *Lookup* the original query on the external knowledge base

2. Extract *candidate terminology*, so terms that could be added to the query and are somehow related to the one in the original string

3. Use a *word sense disambiguation* tool to split relevant and non relevant terms, and add the relevant ones to the original query

Wordnet As introduced in section 1.2 on page 4, Wordnet can be used to extract lemmas but also to find hypernyms and hyponyms or a given word. For instance, if a query presents the word *play*, with the meaning of a *dramatic composition* we could add the latter to the original query.

Another use case could be searching for something like *President Lincoln.* In this case we check for the definition of the two terms, and find out that there's a matching in some possible definitions, indeed the first one could be related to the president of the US, and also the second, thus we can infer that the query is related to US presidency and enrich the query with this information.

Statistical terminology expansion Given the set of relevant and non relevant documents R and N we can select relevant terminology by using the *pointwise Kullback-Leibler divergence.* We compute the probability of a word to be in R and in N

$$p(w) = \frac{count(w, R)}{\sum_{w_i \in R} count(w_i, R)}, \quad q(w) = \frac{count(w, N)}{\sum_{w_i \in N} count(w_i, N)}$$

and then compare them

$$\delta_w(p \parallel q) = p(w) \log \frac{p(w)}{q(w)}$$

if this quantity is close to zero a term is not useful to separate relevant and non relevant documents. On the other end, words with a positive δ can be used to expand the query.

3.2.2 Spelling correction

There are main phases in spelling correction, the first one is to select the correct versions of a certain query in a set of candidates and then choosing the best one, i.e. the most common or the nearest one.

To detect a *spelling error* we could see few query results, or maybe search for the query tokens in a vocabulary.

String similarity   A measure that computes the distance of two strings, one could be the *Levenshtein distance*, also known as the edit distance. It can be solved with dynamic programming implementing the following recursive definition:

$$DP[i][j] = \begin{cases} j \ if \ i = 0 \\ i \ if \ j = 0 \\ DP[i-1][j-1] \ if \ s[i] = r[i] \\ \min(DP[i-1][j], DP[i][j-1], DP[i-1][j-1]) + 1 \ otherwise \end{cases}$$

Phonetic correction   To minimize errors due to phonetic misspelling the idea is to use a *phonetic hash*, so that similar sounding words have the same value. These algorithm are called *Soundex* and are language-dependent.

K-gram classification and matching   The idea is to consider for each word the subsequences of K characters.

An example, for the word *PLAY*, is to first add a start and end symbol, $\#s$ and $\#e$ respectively and then computing the subsequences, counting how many times they appear in the word.

This creates a vector representation for the words, and to find similar words we can use cosine similarity like we did in section 1 on page 3.

Sequence-to-sequence learning   This technique use deep learning to match a sequence into another one. The idea is to train a model to map misspelled words into they correct counterpart.

4   Probabilistic Information Retrieval

The complexity of text analysis is such that we can't have a notion of correctness of an answer to a query, we therefore switch idea and change *true* to *probable*.

4.1 Review of probability theory

Random variable A variable A represents an events, so a subset of the space of possible outcomes. Another representation of A is through a random variable, that maps outcomes to real numbers.

Joint and conditional probabilities We denote $P(A|B)$ the probability of A occurring given the occurrence of B.

Chain rule The following twos hold

$$P(A, B) = P(A \cap B) = P(A|B)P(B) = P(B|A)P(A)$$

$$P(\neg A, B) = P(\neg A|B)P(\neg A)$$

Partition rule If an event B can be divided into an exhaustive set of disjoint sub-cases, the probability of B is the sum of the sub-cases

$$P(B) = P(A, B) + P(\neg A, B)$$

Bayes rule The following rule holds

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)}$$

This rule is a powerful tool to invert the conditional probability. The first step is to estimate the likelihood of the event A, thus $P(A)$, then we can derive a posterior probability $P(A|B)$ after having seen the event B, based on the likelihood of B occurring when A does or not hold.

The left member of the equation $P(A|B)$ is what we are trying to estimate, this could be for instance *probability of a document to be relevant given that it contains a word X.*

$P(B|A)$ is on the other end the probability of observing the new evidence given the initial hypothesis. This could mean finding the *probability of observing the word X given that a document is relevant.*

$P(A)$ is called the prior. For instance *probability of any document to be relevant.*

$P(B)$ is the marginal likelihood, for instance *probability of observing the word X in any document.*

Odds of an event It is often useful to talk about the odds of an event

$$O(A) = \frac{P(A)}{P(\neg A)} = \frac{P(A)}{1 - P(A)}$$

4.2 Binary indipendence model

In information retrieval we represent the relevant of a document given an information need as a random variable R, Given a document d and a query q we want to estimate the probability

$$P(R = 1|q, d)$$

Given a dictionary of terms $T = \langle t_1, \ldots, t_n \rangle$ we model each document as a binary vector $\vec{x} = (x_1, \ldots, x_n)$ where x_i is 1 if the ith term is in the document.

The goal is to calculate

$$O(R = 1|q, \vec{x}) = \frac{P(R = 1|q, \vec{x})}{P(R = 0|q, \vec{x})}$$

Simplifying the goal function   Applying the Bayes rule leads to

$$O(R = 1|q, \vec{x}) = \frac{P(R = 1|q, \vec{x})}{P(R = 0|q, \vec{x})} = \frac{\frac{P(\vec{x}|R=1,q)P(R=1|q)}{P(q,\vec{x})}}{\frac{P(\vec{x}|R=0,q)P(R=0|q)}{P(q,\vec{x})}}$$

$$= \frac{P(\vec{x}|R = 1, q)}{P(\vec{x}|R = 0, q)} \cdot \frac{P(R = 1|q)}{P(R = 0|q)}$$

The left part of the result represent the odds of having that document given that it is relevant and given that query. On the right part we have the probability of the query producing relevant documents or not.

Assuming that the terms are independent, note that this is a strong assumptions, we can write

$$O(R = 1|q, \vec{x}) = \frac{P(R = 1|q)}{P(R = 0|q)} \prod_{i=1}^{n} \frac{P(x_i|R = 1, q)}{P(x_i|R = 0, q)}$$

This product can be split according to the presence of terms in the document

$$O(R = 1|q, \vec{x}) = \frac{P(R = 1|q)}{P(R = 0|q)} \prod_{i=1}^{n} \frac{P(x_i = 1|R = 1, q)}{P(x_i = 1|R = 0, q)} \prod_{i=1}^{n} \frac{P(x_i = 0|R = 1, q)}{P(x_i = 0|R = 0, q)}$$

To simplify the formula we introduce this values

$$p_i = P(x_i = 1|R = 1, q), \quad q_i = P(x_i = 1|R = 0, q)$$

	Document	$R = 1$	$R = 0$
We observe the term	$x_i = 1$	p_i	q_i
We do not observe the term	$x_i = 0$	$1 - p_i$	$1 - q_i$

Thus the previous equation becomes, where the first product iterates over the words that appear in the document and the query, and the second product goes

for words that do not appear in the document, but do on the queries.

$$O(R = 1|q, \vec{x}) = \frac{P(R = 1|q)}{P(R = 0|q)} \prod_{x=1,q=1} \frac{p_i}{q_i} \prod_{x=0,q=1} \frac{1 - p_i}{1 - q_i}$$

This assumes that $p_i = q_i$ for words that don't appear in a query, the assumption is motivated by the fact that a document is relevant or not given a query, not by itself.

We now extend the right product to cover all the query terms, formally

$$O(R = 1|q, \vec{x}) = \frac{P(R = 1|q)}{P(R = 0|q)} \prod_{x=1,q=1} \frac{\frac{p_i}{q_i}}{\frac{1-p_i}{1-q_i}} \prod_{q=1} \frac{1 - p_i}{1 - q_i}$$

We not observe that the quantity $\frac{P(R=1|q)}{P(R=0|q)}$ represents the odds of the query and does not depend on the document. Also, the quantity $\prod_{q_i=1} \frac{1-p_i}{1-q_i}$ depends on the terms that are in the query, it's so query dependent but it does not change for different documents. This leads to a simplification of the previous formula, with some rewriting

$$O(R = 1|q, \vec{x}) \approx \prod_{x_i=1,q_i=1} \frac{p_i(1 - q_i)}{q_i(1 - p_i)}$$

Retrieval Status Value After the last simplification we can calculate the Retrieval Status Value for a document given a query as the quantity

$$RSV_{q,d} = \log \prod_{x_i=1,q_i=1} \frac{p_i(1 - q_i)}{q_i(1 - p_i)} = \sum_{x_i=1,q_i=1} \log \frac{p_i(1 - q_i)}{q_i(1 - p_i)}$$

Estimating probabilities We now just need to estimate p_i and q_i and we are ready to go. If a dataset has informations about relevant documents given a query, we can easily estimate them, given $R(t)$ and $N(t)$ occurrences of term t in relevant and non relevant documents

$$p_i = \frac{R(t_i) + 0.5}{R + 1.0}, \quad q_i = \frac{N(t_i) + 0.5}{N + 1.0}$$

Of course we don't always have the relevant and non relevant documents for queries, so we need some assumptions:

- $p_i = q_i$ *if* $t_i \notin q$, as discussed above

- $p_i = 0.5$ *if* $t_i \in q$, we have the same probability of observing or not a term in a randomly-picked relevant document

- $q_i \approx \frac{|\{d|t_i \in d\}|}{|D|}$, the set of non-relevant documents is approximated by the whole collection of documents D

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By applying the previous assumptions we obtain

$$RSV_{q,d} = \sum_{x_i=1,q_i=1} \log \frac{p_i(1-q_i)}{q_i(1-p_i)}$$
$$= \sum_{x_i=1,q_i=1} \log \frac{(1-q_i)}{q_i}$$
$$= \sum_{x_i=1,q_i=1} \log \frac{N-N_i+0.5}{N_i+0.5}$$

By having n and N as the number of docs with t and the total number of docs, respectively

$$\log \frac{(1-q_i)}{q_i} \approx \log \frac{N-n}{n} \approx \log \frac{N-n}{n} = IDF$$

This means that in practice, we are intersecting a document and a query terms, summing the *IDF* of the words in the intersection, of course, by considering the initial assumptions.

4.3   Non binary models

The non binary models takes into consideration the frequency of the terms and not simply a binary vector. They remove the binary assumption of the model presented in 4.2 on page 13.

Okapi BM25   In this non binary model, the RSV is estimated as follows

$$RSV_{q,v} = \sum_{t \in q} \log \frac{N}{df_t} \cdot \frac{(k+1)tf_{t,d}}{k((1-b)+b(\frac{L_d}{L_{avg}}))+tf_{t,d}}$$

k regulates the term frequency, while b scales the importance of the document length.

4.4   Probabilistic language modelling

The goal is now to remove the independence assumption made in the model at 4.2 on page 13.

In order to accomplish this, we need a model that can compute the probability of a sentence, i.e. a query, where the order of the words is important:

$$P(S) = P(w_1, \ldots, w_n)$$

Another request is to have a generative model for sentences, so a model that estimates the probability of a word, given the previous terms in the sequence:

$$P(w_k|w_{k-1}, \ldots, w_1)$$

The idea is to build a model on top of each document and calculate the probability $P(q|M_d)$ that a query q has been generated by the model M_d.

Computing a sentence probability   Given a sentence, for instance *Mario plays a lot with Maria*, we need to compute the joint probability:

$$P(\mathit{Mario}, \mathit{plays}, a, \mathit{lot}, \mathit{with}, \mathit{Maria})$$

By applying the chain rule we can calculate the probability of the sequence

$$P(x_1, x_2, \ldots, x_n) = P(x_1)P(x_2|x_1)P(x_3|x_1, x_2) \cdots = \prod_i^n P(w_i|w_1, \ldots, w_{i-1})$$

Calculating the quantity on real world documents could be impossible, as we might not be able to count long sequences of words to calculate the probability. An approximation to the sequence probability is given by restricting the sequence length:

$$P(w_1, \ldots, w_n) \approx \prod_i^n P(w_i|w_{i-k}, \ldots, w_{i-1})$$

k is a parameter, it should be small if the dataset is really small, or higher if there is enough data. For instance, if we take $k = 2$ we are estimating the sequence by bi-grams

$$P(w_1, \ldots, w_n) \approx \prod_i^n P(w_i|w_{i-1})$$

Note that to compute the probability of a bi-gram, we just count occurrences of a given bi-gram over all the possible ones.

Query likelihood model   After building a model M_d for each documents, following the idea presented on the previous paragraph, we can estimate relevance of documents given a query.

So the probability that a query was generated by a document becomes:

$$P(q|M_d) = K_q \prod_i P(w_i|M_d)^{tf_{w_i,d}} \quad K_q = \frac{Len(K_q)!}{tf_{t_1,q}! \ldots tf_{t_N,q}!}$$

The K_q is query dependent, so we can ignore it, it would take into consideration the query permutations.

We can also estimate the probability with this quantity

$$P(q|M_d) \approx \prod_i \frac{tf_{t_i,d}}{L_d}$$

Note that $tf_{t_i,d}$ can be changed to the probability of the k-grams computed at the previous paragraph, it depends on how we choose to estimate the sentence probability.

Smoothing   Note that all zero and non frequent term probabilities are a problem, for instance a long sequence could be interrupted by only one bi-gram not appearing in the document, the solution is to smooth the numbers, for instance with the *Laplace smoothing* that adds one to each bi-gram, the *Bayesian smoothing* or the *Linear interpolation* that takes into consideration the global document model.

4.5   Other models

Kullback-Leibler divergence model   In this model the relevance term is computed as the quantity

$$R(d, q) = \sum_i P(w_i|M_q) \log \frac{P(w_i|M_q)}{P(w_i|M_d)}$$

where M_q is the model computed on the query.

Translation model   The idea here is to use a generative model taking into consideration the possible translation of a query term.

5   Unsupervised text classification

Text classification   The problem of associating texts, so documents, to classes, represented by labels. Classes may be a partition of the document space, or they can overlap. The approaches to solve the problem are basically two, pre-trained models, so supervised, or based only on documents, so unsupervised.

Depending on the number of classes and how the cover the document space, we define four tasks

	2 classes	Many classes
Disjoint classes	Binary classification	Multi-class classification
Overlapping classes	Soft binary classification	Multi-label classification

Note that ranking can be seen as soft binary classification.

5.1   Clustering

Goal   Clustering is the problem of grouping objects in clusters, in particular given a distance metric, we want to minimize the distance within a cluster and maximize the one between different clusters.

Parameters   Choosing the right number of clusters is pretty difficult, a high number leads to consistency inside clusters, but poor aggregation, while a low number increases aggregation but reduce quality of the single clusters.

Types of clustering There are many approaches to clustering, one could for instance consider flat or hierarchical clustering, in the first one there is no structure, where in the second clusters might contain other clusters. There is also a difference in cluster assignment, one could assign in an hard or soft way.

5.1.1 Support to document retrieval

Terminological analysis After computing a clustering of documents we could analyze terminology of documents of the same cluster to extract relevant terms, this information can then be used for query expansion.

Cluster pruning By selecting representative documents for clusters, we can compare a query only to those documents. After finding a match all the documents in the cluster are returned.

5.1.2 Evaluation

Given $\Omega = \{\omega_1, \ldots, \omega_n\}$ the set of clusters and $\mathbb{C} = \{c_1, \ldots, c_n\}$ the set of classes, we can evaluate the clustering by focusing on pairs of documents.

Let's consider the pair (x_i, x_j), we ask ourselves if

$$\exists \omega_k : (x_i, x_j) \in \omega_k \ \ \exists c_z : (x_i, x_j) \in c_z$$

They represent the clustering and the real world scenario. If the two are true, we find a true positive, if they are both false a true negative. In the case the first one is true and the second is not, we have a false positive, a false negative otherwise.

We can observe that choosing a small number of clusters leads to a large quantity of false positives, while a fine grain partition, i.e. a high number of clusters, produces a high number og false negatives.

Measures We can use some measures of quality in clustering, for instance

$$Rand = \frac{TP + TN}{TP + TN + FP + FN}$$

but also *Purity* and *Normalized Mutual Information.*

5.2 K-means

This approach starts with k random centroids and assign each data point to the closets one. We then recompute the centroids with the current clusters and repeat the first point until termination.

The termination criterion could be a fixed number of iterations, no change in documents assignments or *RSS* measure below a threshold:

$$RSS = \sum_{k=1}^{K} \sum_{\vec{x} \in \omega_k} |\vec{x} - \vec{\mu}(\omega_k)|^2$$

where the measure is the sum of the distances between a point and it's centroid squared. Note that at each iteration the *RSS* decrease.

Two issues of this approach is that the initial position of clusters changes the results a lot, and also, the k parameter is crucial to the algorithm. A good value for k is $K = \min_K(\min(RSS_K) + \lambda K)$

5.3 Model-based clustering

A k-means generalization is obtained by interpreting the centroids as a model that generates data. A centroid with some added noise can generate a document.

Idea Instead of generating classes, we start from the points, we assume that they were generated with a generative model and we estimate the latent model parameters.

Latent parameters $\Theta = \{\vec{\mu}_1, \ldots, \vec{\mu}_n\}$ are the centroids to be found by k-means.

$L(D|\Theta)$ is the log-likelihood that the data D wa generated by Θ, this is the objective function, so Θ becomes:

$$\Theta = \max_{\Theta} L(D|\Theta) = \max_{\Theta} \sum_{n=1}^{N} \log P(d_n|\Theta)$$

That means maximizing the sum of probabilities that documents are generated by the model. We are in a soft clustering scenario, as we are basically dealing with probabilities as cluster assignments.

5.3.1 Expectation Maximization algorithm

This is an iterative algorithm that maximizes $L(D|\Theta)$, but can also be used to find latent models in a variety of applications.

Let's consider a multivariate Bernoulli distribution for data, so all documents are binary vectors, we want to estimate the probability that a given document is assigned to a specific cluster given the models parameters.

$$P(d|w_k; \Theta) = \prod_{t_m \in d} q_{mk} \cdot \prod_{t_m \notin d} (1 - q_{mk}) \quad \Theta_k = (\alpha_k, q_{1k}, \ldots, q_{Mk})$$

where $q_{mk} = P(U_m = 1|\omega_k)$ is the probability that a document from the cluster ω_k contains the term t_m.

α_k si the prior of the cluster k, so the probability that of a document to be in that cluster, not having any information about it.

Process We start by generating a document by picking a cluster w_k with probability α_k, generating terms with probability q_{mk}.

We need to estimate the two parameters, and we do that iteratively similarly to k-means. The iteration is composed by two steps, *Maximization* and *Expectation.*

Maximization step The goal here is to estimate the parameters of the model, so the prior and the terms probability given the cluster:

$$q_{mk} = \frac{\sum_{n=1}^{N} r_{nk} I(t_m \in d_m)}{\sum_{n=1}^{N} r_{nk}} \quad \alpha_k = \frac{\sum_{n=1}^{N} r_{nk}}{N}$$

where I can be one or zero wether the term appears or not in the document, and r_{nk} is the soft assignment of the document n to the cluster k.

Expectation step We now estimate the probability of each term to be assigned to a cluster.

$$r_{nk} = \frac{\alpha_k(\prod_{t_m \in d_n} q_{mk})(\prod_{t_m \notin d_n}(1 - q_{mk}))}{\sum_{k=1}^{K} \alpha_k(\prod_{t_m \in d_n} q_{mk})(\prod_{t_m \notin d_n}(1 - q_{mk}))}$$

this is basically, for each cluster, the prior, multiplied by the product of q_{mk} or $1 - q_{mk}$ for each term in the document computed before, normalized by the same quantity over all the clusters.

Why do EM works? The main idea of the algorithm is to push documents that contains the same words in the same cluster. Basically if two words appears in a document they should have a high probability of being assigned to the same cluster, so they have the same q_{mk}. This leads to assigning documents with the same words in the same clusters.

5.4 Affinity propagation clustering

Input The input is a similarly matrix, for instance $s(i, j) = -||\vec{i} - \vec{j}||^2$. There are also some special values for the diagonal of the matrix, where larger values are more likely to be chosen as exemplars for clusters.

Idea Documents exchange messages to decide exemplars and clusters assignments. This approach is similar to *page rank with authorities* [1].

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[1] https://nlp.stanford.edu/IR-book/html/htmledition/hubs-and-authorities-1.html

Responsibility messages   A message $r(i, k)$ that denotes how well k is an exemplar for i. An example can be

$$r(i, k) = s(i, k) - \max_{k' \, s.t. k' \neq k} \{a(i, k') + s(i, k')\}$$

basically we take the similarity of a pair of nodes and subtract the maximum similarity for any other document. This value will be zero for the most similar document, and negative for all other, as the quantity $a(i, k')$ is initially zero.

Availability   This message $a(i, k)$ represent how appropriate is for i to choose k as exemplar.

$$a(i, k) = \min \left\{ 0, r(k, k) + \sum_{i' \, s.t. i' \notin \{i, k\}} \max\{0, r(i', k)\} \right\}$$

Self availability   The availability measure given the same document as argument, is

$$a(k, k) = \sum_{i' \, s.t. i' \neq k} \max\{0, r(i', k)\}$$

Process   At first, documents are connected by similarity, then we recompute the availability with respect to the responsibility and vice versa. This leads to making documents connections stronger and stronger and to the election of an exemplar for clusters. After some iterations convergence is reached.

5.5   Hierarchical clustering

This approach produces a hierarchy of clusters. It does not need the number of clusters but a criterion of optimal cluster selection. Most of this algorithms are deterministic.

Process   Staring from a similarity matrix for the documents, we start taking the pair of documents with maximum similarity and group them in the same cluster. An idea is to have documents as nodes, and to add a cluster node connected to the two most similar documents.

The next step is to delete the rows and columns of the two selected documents, as they are already assigned, but, we add a new row and column for the cluster node. The similarity value between a document and a cluster is the minimum similarity in that cluster for the considered document, but many strategies can be applied.

After rebuilding the table, we continue with the initial step, note that if a cluster and document are selected as most similar items, we are adding a cluster connected to them, ergo we are considering the cluster obtained in the first step, as a sub-cluster for the last one.

The last step is to select the number of clusters. The previous logic created tree, as we grouped documents and clusters by adding nodes connected to them. We can choose a threshold for the intra-cluster similarity, or select a certain number of clusters, or considering the maximum jump in similarity in the obtained cluster tree and cut off the tree there. By cutting off the tree, we take the lower part, so the one with the leaves.
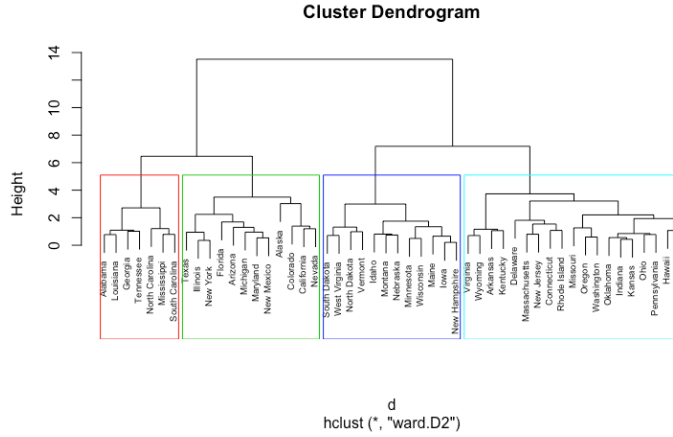

Figure 2: Hierarchical clustering example with 4 clusters

Top-down clustering   This approach starts from a cluster containing all the documents, and recursively splits it keeping track of the parent of each document. The process stops when a threshold of intra-cluster similarity is obtained or when there's a cluster for each document. This approach is the opposite of the bottom up tree creation.

6   Topic modeling

Vector space models problems   The approaches seen so far did not take into consideration semantics of words. We understood that the vector space models suffer from high dimensionality, the terms also could be synonyms, leading to ambiguity, in fact two words that means the same thing are two different dimensions in the model.

6.1   Linear algebra recap

Eigenvalues and eigenvectors   Given a square matrix of size n, if we find a n dimensional vector such that

$$A\vec{x} = \lambda\vec{x}$$

λ is a eigenvalue and $\vec{x}$ is an eigenvector. It also holds that

$$(A - \lambda I)\vec{x} = 0 \implies |A - \lambda I| = 0$$

where I is the identity matrix, and the implication holds as $\vec{x}$ is non-zero. In general the second part of the implication has n solutions, so n eigenvalues associated with a certain eigenvector.

After finding the solutions we can write that

$$AX = A[\vec{x_1}, \ldots, \vec{x_n}] = A\vec{x_1} + \cdots + A\vec{x_n}$$
$$= \lambda_1 \vec{x_1} + \cdots + \lambda_n \vec{x_n} = \Lambda X$$

Where Λ is the diagonal matrix of the eigenvalues, and X the matrix with eigenvectors.

Diagonalization   If the n eigenvector found at the previous step are independent, and the matrix A is invertible, which is not a problem as its square, we can write

$$AX = X\Lambda \rightarrow AXX^{-1} = X\Lambda X^{-1} \rightarrow A = X\Lambda X^{-1}$$

This representation is called diagonalization of A. Also, if A is symmetric, we can write

$$A = X\Lambda X^\top$$

6.2   Latent Semantic Indexing

The goal is to discover topics that motivate data by using matrix factorization techniques, while taking in consideration the possibility of expressing the same topic with different words.

Diagonalization   We learned that if a matrix is symmetric we can write the diagonalization as $A = X\Lambda X^\top$. The question is, what happens if we multiply the document-term matrix by itself?

We find a symmetric matrix that somehow encapsulates a notion of similarity of documents because if two documents share the same words the product will be really high for that two documents.

If we multiply the term-document matrix we obtain a similar result.

Singular Value decomposition   The idea behind the approach is to define the term document matrix A in terms of

$$A = U\Sigma V^\top$$

this means finding something that holds together documents and terms. The terms in the diagonalization represent:

- V is the matrix with the eigenvectors of $AA^\top$

- U is the matrix with the eigenvectors of $A^\top A$

- Σ is the diagonal matrix composed starting from the eigenvalues of $AA^\top$ and $A^\top A$, which are the same

Note that the eigenvalues in Σ are sorted on the diagonal.

Defining topics   After applying singular value decomposition to the document term matrix we have the latent topic matrix Σ. We can consider a new matrix $\Sigma_{k \times k}$, that somehow encapsulate the top k relevant topics for the documents. Note that after this cut of Σ we get a diagonalization that is an approximation of the original matrix.

$$A \approx U \Sigma_{k \times k} V^\top$$

To make topics human understandable we assign each a set of words, and also, we can assign to each document a topic, in a soft clustering fashion.

Basically, given the three matrixes that SVD outputs, after cutting Σ as discussed above, we can discover what topics have the documents by multiplying U and $\Sigma_{k \times k}$. The output will be a matrix with document on rows and topics on columns, where the absolute value of a cell represent how strong a topics is present in a document.

In the same way we can find a relation between topics and words by multiplying $\Sigma_{k \times k}$ and $V^\top$.

6.3   Latent Dirichlet Allocation

This approach relies on probability theory to find topics in documents. In the LSI approach we tried to find a *glue*, formally Σ, between document and terms.

In LDA the goal is to estimate the probabilistic relation between topics and documents, basically the probability that a given document contains a given topic, and the one that a topics contains a certain word.

Elements   The basic elements of the approach are:

- $\phi^{(k)}$ probability distribution over the vocabulary for the kth topic

- θ_d document distribution over topics

- z_i topic index for the word w_i

- α, β hyperparameters that govern the first two distributions

Estimating the generative model To find the topics and document distributions we assume that they were generated by a given distribution and try to estimate them.

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7 Supervised classification

As discussed in section 5 on page 17, text classification is the problem of associating texts to classes, that are identified by labels.

Training data set A set of data examples are provided to the classifier to learn the mapping between data and labels. The goal is then to predict unlabeled data.

Unsupervised vs Supervised In the first case we don't have any information about classification criterions, while in the second we have classes associated to features, represented by the data points associated with labels.

7.1 Models

Decision trees The application of this model to text classification is rather easy. One could compute a vectorization of the text, in one of the many ways seen and apply the tree learning algorithm. This means imposing constraints on the presence and relevance of a certain word in a document.

One of the pros of having a decision tree is quick and easy visualization. This means being able to see how the model computes a classification, to potentially avoid unethical decisions.

Rule-based classifiers These classifiers are based on rules, they are similar to decision trees, basically the antecedent of the rule is a subset of words and the consequent is a label.

Naive Bayes Classifiers The idea is that data has been generated by a mixture of k components, where k is the number of classes. The goal is to estimate the probability

$$P(class \mid features) = \frac{P(class)P(features \mid class)}{P(features)}$$
$$= P(class)P(features \mid class)$$

where the probability of a class is the size of the class over the size of the corpus. Also, the probability of features given the class, is simply the product of the count of a given words over the total count of the words in the class.

K-NN classifiers   The application of the model to text is straight forward after text vectorization.

Linear classifiers   The idea is to learn an hyperplane separating data, again, this is not different to any linear classifier task.

The main idea is to minimize a loss function plus a regularizer term.

Support Vector Machines   A linear classifier is the Support Vector Machine, the goal is to find the optimal hyperplane separating data, i.e. the one with greater margin between the two classes.

To accommodate non linearly separable data one could use the kernel trick. Basically we perform a transformation of the dataset to map it in a high dimensional space where data point could be linearly separable. There are many possible kernels.

7.2   Classification strategies

When the system is efficient in performing binary classification we can apply it to multi-class classification by transforming the problem in a binary task.

One vs Rest   The goal here is to learn a binary classifier between a class and all the others.

One vs One   The idea here is to train a classifier between every pair of classes.

7.3   Multi-label classification

To assign multiple labels to data point we need to exploit something in the previous classifiers.

For instance, we could use the distance of the many one vs one classifiers as confidence measures for labels.

Another way is to take the power set of classes and train many classifiers as before. We are left with a multi-class classification where each class is a combination of labels.

7.4 Hyperparameters tuning and evaluation

Model selection The activity of tuning hyperparameters to find the best model instance to solve a task.

Cross Validation The idea is to split the dataset in k folds and consider $k - 1$ folds as training and the last one as test set. We iterate over the possible training sets and average the error.

Confusion matrix We can arrange the errors in a confusion matrix to see how many elements of class i are classified as class j. Obviously the goal is to have positive values only on the diagonal.

8   Language models

Language model A language model is the probability distribution of a sequence of words, it can be used for all sort of applications.

$$p(w_1, w_2, \ldots, w_n)$$

Examples This probability could be used to predict the next word given a sequence

$$p(w_n | w_1, w_2, \ldots, w_{n-1})$$

Another usage is to predict a feature given the sequence, for instance find the author from the text

$$p(author | w_1, w_2, \ldots, w_n)$$

Bases and evaluation The goal of language modeling is to build a computational model of the language of an entity X, that can be any entity in a domain.

Language models are entirely *data-driven*. In other terms, we need to train models on the right data. One of the disadvantages of this fact is that instead of modeling the language of X, we are modelling the language of the *documents about X*.

8.1   N-grams models

The easier way to estimate a sequence probability is to count frequencies, but that could be unfeasible. We discussed a similar approach in Section 4.4 on page 15 in *Computing a sentence probability.*

Unigram models A first approach is to use word frequencies, the idea is to find the relative frequency of a word, by simply counting it in the corpus and use it to estimate the probability of a sequence.

$$p(w_n, w_{n-1} \ldots, w_1) = p(w_n)p(w_{n-1}) \ldots p(w_1)$$

Note that this approach does not take into consideration the sequence in any way, this is a *Unigram model* .

These models might be useful to compare two documents, where we need to compare the probability of seeing a given word, or to find how much relevant is a given word for a document. We can do that by comparing the word probability in all the dataset and in the single document.

Pointwise Mutual Information

$$pmi(a, b) = \log \frac{p(a, b)}{p(a)p(b)} = \log \frac{p(a|b)}{p(a)} = \log \frac{p(b|a)}{p(b)}$$

We can set the two variables to the event of extracting the document C from the corpus and the probability w of extracting a given word.

$$pmi(w, C) = \log \frac{p(w, C)}{p(w)p(C)}$$

By computing the *pmi* for the same word and two different documents, we can compare the difference of relevance of a word for the two documents.

Chain rule We now want to consider the sequence instead of taking words independently as in the unigram model.

$$p(w_k | w_1, \ldots, w_{k-1}) = \frac{count(w_1, \ldots, w_{k-1}, w_k)}{\sum_w count(w_1, \ldots, w_{k-1}, w)}$$

Basically to compute the probability of finding a given word after a sequence, we count how many times we observe the sequence with that word, over the count of all other words after the sequence.

In practice, we can reduce a long sequence to a shorter one, for instance of two or three words, to potentially have enough data to estimate the probability.

8.2 Evaluation

There are basically two ways of evaluating a language model:

- *Extrinsic evaluation* : embed the system in a real applications and measure improvement

- *intrinsic evaluation* : define a metric and evaluate the model independent from any application, this requires a training set and a testing set, where the second must be different but consistent with the language learned by the model

Perplexity   In the case of intrinsic evaluation, the main idea is to check if the model fits the test data. This measure is called perplexity.

$$pp(T) = p(w_1, \ldots, w_n)^{-\frac{1}{n}}$$

For a two-gram model the measure becomes:

$$pp(T) = \sqrt[n]{\prod_{i=1}^{n} \frac{1}{p(w_i \mid w_{i-1})}}$$

Generalization and zero probabilities   Language models have an issue when a word is present in the test set but not in the training set, this becomes unknown with no null probability.

To fix this we apply smoothing, for instance *Laplace smoothing*, i.e. adding one to each word count, or a parameter k.

Backoff and interpolation   Another way to deal with null probabilities is to apply *backoff and interpolation*. Basically to estimate a n-gram frequency, if we do not have enough evidence, we use the $(n-1)$-gram frequency, until we have just a unigram. An example can be:

$$p(New\ York\ City) = p(City \mid York)\ p(York \mid New)$$

Interpolation means computing a weighted sum over an n-gram levels, this way, even if we only observe the last word of a n-gram, we do not have zero probability.

8.3   Word-Context models

The problem of data sparsity is mitigated by the n-gram models, but not solved completely.

Skip-gram model   The main of the model is to fix the cases where two sentences are really similar, i.e. they give the same information, but they do not have any common n-gram.

We introduce the concept of *word context*, basically we take a n-gram by skipping at most k words, this leads to skip noisy terms. By doing so, we could potentially match two sentences that differs for a few words but that do not share commons grams.

Continuous Bag of Words   The idea of this model is to take the context of a word, for instance a collection of 2-grams, and predict a word.

Word context matrix We can think of a word context matrix, where we store information about appearance of a word w within t other words.

Taking into account the word order depends on the focus of the model, in language the order is important, in semantics it is less relevant.

The context matrix stores in $c_{i,j}$ for each word w_i how many times the word w_j occurs in its context. If we consider the row of that matrix as a vector, two words are close in that space if they have a similar context.

After computing the matrix, we can factorize it with singular value decomposition to obtain a more compact representation. A problem is that such a factorization has many zero terms, to fix it we can *weighted matrix factorization.*