

Email Classification with Naïve Bayes

Mitchell Della Marta, Shu Han Bor

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1 Introduction

1.1 Aim

To implement an accurate Naïve Bayes classifier for email spam detection.

1.2 Importance

As email is one of the main forms of communication, spam detection to remove spam is important. Email spam is an unsolved global issue that negatively affects productivity and uses up valuable resources. Spam emails are usually sent with malicious intent to users for some monetary gain. Email users spend a large amount of time regularly deleting these spam emails, which needlessly occupy storage space and consumes bandwidth. Hence the development of classifiers that are able to separate legitimate from spam emails is required. Classification of emails to separate and remove spam from other emails, aims to save the user from having to spend time and effort sorting through and deleting spam, and protect unaware users from malware.

Traditional non-machine learning based techniques are easily fooled by spammers and require periodic manual updates. Therefore, machine learning based methods are required, which are able to automatically analyse email contents and dynamically update themselves to cope with new spamming techniques.

2 Data Preprocessing

Before classification can be performed, we must represent the files in our sample set *LingSpam-mini600* [2] appropriately. Using the “bag-of-words” model, words are extracted from the file and treated as features. There are two main characteristics in an email; the subject and the body. Thus, we will construct two “bags-of-words”, one for each component. To determine which corpus a feature will belong to, we if the line begins with “Subject:”. If it does the words within the line will be added to the subject corpus, otherwise they will be added to the body corpus.

To each file, we performed the following steps:

1. Replace all punctuation and special symbols with a space.
2. Remove stop words, using a list of stop words “english.stop”.
3. Remove numbers or words containing digits.

The tokenisation schemes used in the example data are different to that used in our stop words list. The data detaches clitics and contraction affixes, whereas the stop words list is tokenised based on whitespace and punctuation characters. Therefore, we chose to replace punctuation with a space (e.g. “n’t” becomes “n”, “t”) instead of simply removing them. This meant that the stop word list would remove clitics as it removes single characters, instead of keeping it as a feature as (“nt” isn’t in the stop words list).

The *document frequency* feature selection method was then used to select the 200 words which occurred in the most documents, and used to build a classifier for the given documents. The following steps were taken to accomplish this:

1. Counter is created for each email, which keeps track of the number of times each word appears in that email.
2. For each word that appeared in the counter for an email, it is added once to the counter for the corresponding subcorpus.
3. In each corpus the words with the top 200 document frequency score are selected as features to represent that corpus.

Note that different runs of our preprocessing script can result in different classifier accuracies. As the counter is stored in a hashmap, words with the same frequency before and after the 200 word cutoff are chosen arbitrarily. Depending on the words chosen, the accuracy of the classifier can be affected.

Each selected feature was then weighted using its *tf-idf* score [1].

$$tfidf(t_k, d_j) = \#(t_k, d_j) \times \log \frac{|Tr|}{\#Tr(t_k)}$$

Then the scores were normalised, using *cosine normalisation* [1].

$$w_{kj} = \frac{tfidf(t_k, d_j)}{\sqrt{\sum_{s=1}^{|T|} (tfidf(t_s, d_j))^2}}$$

In cases where the *tf-idf* weightings for all 200 features for a given document is 0 (i.e. the feature appears in and only in every document in a corpus), the denominator of the cosine normalisation will also be 0. This is impossible, so instead we return $w_{kj} = 0$.

2.1 Data Characteristics

Table I: Characteristics of data set

	Subject	Body
# Features before removing stop words	1074	19886
# Features after removing stop words	915	19386
# $Class_{nonspam}$	600	600
# $Class_{spam}$	200	200

The “bag-of-words” model produced 19886 and 1074 features in the body and subject corpora respectively. After the removal of stop words, there were 19386 unique words remaining in the body corpus, and 915 words in the subject corpus (see Table I).

3 Feature Selection

Feature selection is performed using document frequency. This method involves computing the number of documents a word occurs in, for every word, and selecting the top 200 words with the highest scores to build a classifier.

Shown in Table II and Table III are the top 100 words for the subject and body corpora respectively and their document frequency score. Removing stop words filters out extremely common words that have little value in classification. It is beneficial in text processing, as it removes low quality features, allowing more significant features to have precedence. Most resulting words are sensible, although there are some aren’t (e.g. “qs”, “ll”, “ca”). However, removing these words results in a lower accuracy. Therefore, we have chosen to retain these as some of them may be help the classifier distinguish between spam and non-spam emails. Thus, given our task to process natural language text, the selection of words shown makes sense as it gives a better representation of the contents of the emails, and helps improve the accuracy of the classifier.

A comparison of Table II and Table III shows significant disparities in the document frequency of features and word distribution. The frequencies for the subject is significantly lower than that of the body. Whilst some features are shared between subject and body, most features selected are different.

Table II: Top 100 words in subject corpus and corresponding document frequency (DF) scores

Rank	Word	Score	Rank	Word	Score	Rank	Word	Score
1	sum	30	35	speaker	6	69	intuitions	4
2	summary	26	36	german	6	70	banning	4
3	english	24	37	internet	6	71	school	3
4	language	21	38	business	6	72	resolution	3
5	free	20	39	list	5	73	ary	3
6	disc	19	40	resources	5	74	adjectives	3
7	query	18	41	native	5	75	verbal	3
8	linguistics	15	42	research	5	76	teaching	3
9	comparative	13	43	word	5	77	future	3
10	sex	13	44	spanish	5	78	lists	3
11	opposites	12	45	linguist	5	79	background	3
12	words	12	46	jobs	5	80	synthetic	3
13	book	10	47	needed	5	81	credit	3
14	email	10	48	grammar	5	82	home	3
15	call	9	49	software	5	83	live	3
16	job	9	50	languages	5	84	youthese	3
17	method	9	51	time	5	85	uniformitarianism	3
18	japanese	8	52	fwd	4	86	released	3
19	correction	8	53	summer	4	87	names	3
20	syntax	7	54	address	4	88	opportunity	3
21	program	7	55	books	4	89	decimal	3
22	qs	7	56	information	4	90	world	3
23	chinese	7	57	request	4	91	misc	3
24	announcement	7	58	phonetics	4	92	sites	3
25	million	7	59	pig	4	93	double	3
26	part	6	60	american	4	94	acquisition	3
27	slip	6	61	programs	4	95	site	3
28	workshop	6	62	unlimited	4	96	policy	3
29	armey	6	63	web	4	97	fall	3
30	money	6	64	www	4	98	teach	3
31	lang	6	65	secrets	4	99	hey	3
32	conference	6	66	great	4	100	line	3
33	dick	6	67	read	4			
34	mail	6	68	systems	4			

Table III: Top 100 words in body corpus and corresponding document frequency (DF) scores

Rank	Word	Score	Rank	Word	Score	Rank	Word	Score
1	information	205	35	message	91	69	full	74
2	language	192	36	ll	89	70	system	74
3	mail	183	37	receive	88	71	ac	73
4	university	179	38	check	88	72	today	73
5	time	178	39	phone	88	73	questions	72
6	list	171	40	good	87	74	remove	72
7	address	165	41	day	86	75	interest	72
8	english	159	42	interested	86	76	john	71
9	linguistics	156	43	year	86	77	found	70
10	http	156	44	include	85	78	related	70
11	people	146	45	working	85	79	site	69
12	send	146	46	case	85	80	linguist	69
13	free	144	47	based	84	81	usa	69
14	make	140	48	ve	84	82	text	68
15	email	133	49	note	83	83	point	68
16	number	128	50	home	83	84	read	68
17	work	128	51	made	83	85	ago	67
18	www	122	52	part	83	86	book	67
19	languages	119	53	including	81	87	week	67
20	find	118	54	mailing	81	88	making	66
21	fax	116	55	type	80	89	dear	66
22	order	108	56	give	79	90	cost	66
23	call	103	57	program	79	91	question	65
24	form	101	58	web	79	92	simply	65
25	research	100	59	place	79	93	received	63
26	state	99	60	special	78	94	offer	63
27	linguistic	99	61	line	78	95	general	63
28	subject	98	62	date	78	96	important	62
29	years	98	63	days	77	97	data	62
30	world	98	64	back	76	98	ca	61
31	contact	97	65	internet	76	99	long	61
32	de	96	66	american	75	100	summary	61
33	money	94	67	service	75			
34	word	91	68	business	74			

4 Subject vs Body Analysis

4.1 Results

We ran our preprocessed data against a variety of Weka classifiers; ZeroR, OneR, IBk with $k = 1$ (1-NN) and $k = 3$ (3-NN), NaiveBayes (NB), J48 (DT) and MultilayerPerceptron (MLP). We tested these with Weka's 10 fold cross-validation. We also ran it against our Naïve Bayes classifier, tested against our implementation of the 10 fold cross-validation.

Table IV: Various classifiers tested with 10 fold cross validation for both the subject and body corpora

Accuracy (%)		
Classifier	Subject	Body
ZeroR	66.67	66.67
OneR	70.00	82.00
1-NN	79.00	87.17
3-NN	68.17	85.00
NB	68.50	94.83
DT	66.67	92.50
MLP	76.17	96.67
MyNB	80.83	95.33

Table V: Accuracy of classifiers from lowest (1) to highest (8) for both corpora

Accuracy Ranking								
Corpus	1	2	3	4	5	6	7	8
Subject	ZeroR	DT	3-NN	NB	OneR	MLP	1-NN	MyNB
Body	ZeroR	OneR	3-NN	1-NN	DT	NB	MyNB	MLP

The comparison of the performance (accuracy) of MyNB and NB over the 10 folds was completed in the same execution as the respective results in Table IV. MyNB used our 10 fold cross-validation and NB used Weka's 10-fold cross-validation. The items in each fold (given to MyNB and NB) are expected to be different.

Table VI: Comparison of fold accuracy between our Naïve Bayes classifier (C_1) and Weka's (C_2) for the subject corpus

Fold Number	C1 (%)	C2 (%)	Difference
Fold 1	80.00	60.00	20.00
Fold 2	83.33	73.33	10.00
Fold 3	71.67	63.33	8.34
Fold 4	80.00	76.67	3.33
Fold 5	83.33	71.67	11.66
Fold 6	76.67	70.00	6.67
Fold 7	83.33	73.33	10.00
Fold 8	86.67	63.33	23.34
Fold 9	81.67	73.33	8.34
Fold 10	81.67	60.00	21.67
Mean	80.83	68.50	12.34

Table VII: Comparison of fold accuracy between our Naïve Bayes classifier (C_1) and Weka's (C_2) for the body corpus

Fold Number	C1 (%)	C2 (%)	Difference
Fold 1	98.33	98.33	0.00
Fold 2	98.33	93.33	5.00
Fold 3	96.67	95.00	1.67
Fold 4	90.00	100.00	10.00
Fold 5	91.67	95.00	3.33
Fold 6	96.67	93.33	3.33
Fold 7	93.33	95.00	1.67
Fold 8	95.00	91.67	3.33
Fold 9	96.67	91.67	5.00
Fold 10	96.67	95.00	1.67
Mean	95.33	94.83	3.50

4.2 Discussion

4.2.1 Comparison of Classifiers

The accuracy of a classifier in comparison to the other classifiers, varies slightly against the two corpora. Accuracy from lowest to highest for both corpora are shown in Table V. Although there

are some discrepancies, in general ZeroR, OneR, 3-NN, DT perform more poorly than 1-NN, NB, MyNB, MLP.

The ZeroR classifier performs poorly for both corpora. It returns the majority class given from the training data. As a result ZeroR returns “nonspam” as there are 400 non-spam and 200 spam cases. Thus ZeroR produces a correct answer for and only for test data of the majority class, “nonspam”, which is

$$\begin{aligned}\frac{\#nonspam}{\#documents} &= \frac{360}{540} \\ &= 66.67\%\end{aligned}$$

as obtained.

OneR works similarly to ZeroR, but instead of having no rules, it generates one rule for each item in the data set. After calculating rules for each, it will select the rule that gives the smallest total error. Although its accuracy isn’t high, it is decent for the simplicity and speed of the algorithm.

In contrast to these two classifiers, multilayer perceptron (MLP) is more complex and returns the highest accuracy for body, and the second highest accuracy for subject. MLP is defined as “consists of multiple layers of simple, two-state, sigmoid processing elements (nodes) or neurons that interact using weighted connections.” (Pal 1992, p.684). The Weka MLP assigns the features as output neurons with a random weight. Weka’s default number of hidden neurons[3] was used,

$$\begin{aligned}\frac{(\text{attributes} + \text{classes})}{2} &= \frac{(200 + 2)}{2} \\ &= 101\end{aligned}$$

As the network is then trained using back propagation, which corrects the weights to minimize the error in the entire output, MLP has a high accuracy. However, although it is very accurate, MLP takes significantly more time to train than the other classifiers.

Naïve Bayes (NB) is based on the Bayes rule of conditional probability. It analyses each attribute individually, by assuming they are all of equal importance and independent of one another. Although it is based on a simple concept, our MyNB classifier outperforms MLP which is a comparatively more complex algorithm for the subject corpus, and performs almost as well as MLP for the body corpus.

Whilst NB and MyNB are based on the same principle, MyNB outperforms NB. The difference in performance is due to implementation. The major difference between implementations is that NB uses numeric estimator precision values that are chosen by calculating the differences between adjacent values. This allows them to form a kernel density estimation, that estimates a probability density function (PDF), as opposed to using a normal PDF. On the other hand, we simply assume the data has a normal probability distribution function.

1-NN and 3-NN are both types of the k-nearest neighbour algorithm, where $k = 1$ and $k = 3$ respectively. Although they use similar algorithms, 1-NN performs better than 3-NN. 1-NN takes one neighbour, whereas 3-NN takes three. Therefore, this disparity is probably because in most cases the closest neighbour correctly classifies the new example, but the next two closest

do not. Hence 1-NN gets a greater accuracy than 3-NN. These classifiers work moderately well compared to the other classifiers, as each class is characterised by a large number of features, which the nearest neighbour algorithm takes advantage of. In order to increase the accuracy of this algorithm, it should be given a larger training set. Although the benefits of this have to be weighed against the time it takes to find the nearest neighbour in a larger training set.

Weka uses J48 (implementation of Quinlan's C4.5) to construct the decision tree classifier, which relies on using information gain to select attributes that best classifies the data. It performs averagely on the body corpus, and relatively poorly on the subject corpus.

4.2.2 Comparison of myNB vs Weka's NB

We conduct a paired t-test in our comparison to determine whether the differences in accuracy between our and Weka's Naïve Bayes was statistically significant, with a confidence level of 95%. To carry out the paired t-test:

1. Calculate the difference between the two classifiers in each fold

$$|c_{1_i} - c_{2_i}| \text{ for } i \in [1, \dots, N]$$

where $c_{1_i} \in C_1$ and $c_{2_i} \in C_2$ for all i .

2. Calculate the sample standard deviation of the differences:

$$s_{N-1} = \sqrt{\frac{1}{N-1} \sum_{i=1}^N (x_i - \bar{x})^2}$$

where $X = \{x_1, \dots, x_N\}$ are the observed differences.

3. Calculate the $(1 - \alpha)$ -upper confidence interval (UCL) of the mean:

$$\text{UCL}_{1-\alpha} = \bar{X} \pm t_{(1-\alpha)(k-1)} s_{N-1}$$

for the confidence level $(1 - \alpha) = 0.95$.

4. If the interval $\text{UCL}_{0.95}$ contains 0 then the difference in accuracy is not statistically significant; otherwise, it is.

Using the differences calculated in Table VI for the subject corpus we find that:

$$\begin{aligned} s_{N-1} &= 6.86 \\ \text{UCL}_{0.95} &= \bar{X} \pm t_{0.95,9} \cdot s_{N-1} \\ &= 12.34 \pm 2.262 \times 6.86 \\ &= 12.34 \pm 15.52 \\ &= [-3.18, 27.85] \end{aligned}$$

Similarly, given the differences calculated in Table VII, we obtain for body corpus:

$$\begin{aligned}
 s_{N-1} &= 2.77 \\
 \text{UCL}_{0.95} &= \bar{X} \pm t_{0.95,9} \cdot s_{N-1} \\
 &= 3.50 \pm 2.262 \times 2.77 \\
 &= 3.50 \pm 6.27 \\
 &= [-2.77, 9.77]
 \end{aligned}$$

For the subject corpus $\text{UCL}_{0.95} = [-3.18, 27.85]$ and for body corpus $\text{UCL}_{0.95} = [-2.77, 9.77]$. In both cases the interval $\text{UCL}_{0.95}$ contains 0. Therefore, the difference between the two Naïve Bayes classifiers for both subject and body are not statistically significant.

4.2.3 Comparison Between Subject and Body Corpora

The overall performance of the classifiers regardless of type, perform significantly better on the body than the subject corpus. This suggests that the email body is a better indicator of content.

5 Challenge Analysis

In the extension seven feature selection metrics were implemented. We compared the accuracies of the metrics when used as the features for our Naïve Bayes to empirically determine the best feature selection metric and number of features for classifying [2].

The chosen metrics were Correlation coefficient (CC) [4], Information gain (IG) [5], Chi-square (CHI), Categorical proportional difference (CPD), Mutual information (MI) as defined in [6], Document frequency (DF) and Random feature (RF).

5.1 Feature Selection Metrics

Here we present the seven feature selection metrics used, presented in terms of the following four dependency tuples [5]:

1. (t, c) : presence of t and membership in c
2. (t, \bar{c}) : presence of t and membership in \bar{c}
3. (\bar{t}, c) : absence of t and membership in c
4. (\bar{t}, \bar{c}) : absence of t and membership in \bar{c}

where t represents a term and c and \bar{c} represent the two classes.

Correlation coefficient (CC). Correlation coefficient of a word t with a category $c_i \in \{c, \bar{c}\}$ was used as:

$$CC(t) = \sqrt{N} \frac{|P(t, c)P(\bar{t}, \bar{c}) - P(t, \bar{c})P(\bar{t}, c)|}{\sqrt{P(t)P(\bar{t})P(c)P(\bar{c})}}$$

it is a simplification of the Correlation coefficient given in [5] due to having two classes such that the complement of a class is simply the other class.

Information gain (IG). Information gain measures the decrease in entropy or the number of bits of information obtained when the feature is given vs. absent [7],[5]. Defined in [5] for a term t present in category c_i as:

$$IG(t, c_i) = \sum_{c' \in \{c, \bar{c}\}} \sum_{t' \in \{t, \bar{t}\}} P(t', c') \cdot \log \frac{P(t', c')}{P(t')P(c')}$$

given that we have two classes:

$$IG(t, c_i) = IG(t, c) = IG(t, \bar{c})$$

so the IG value of a term t is independent of the class.

For the remaining metrics we define them in the following terms used in [6]:

1. $A := \#(t, c)$, is the number of times term t and category c occur together.
2. $B := \#(t, \bar{c})$, is the number of times term t occurs with \bar{c} .
3. $C := \#(\bar{t}, c)$, is the number of times category c occurs without term t .
4. $D := \#(\bar{t}, \bar{c})$, is the number of times category \bar{c} occurs without term t .

Chi-square (CHI). Chi-square measures the lack of independence between a term t and a category c' . For a term t' and a category c' it is defined as:

$$\chi^2(t, c') = \frac{N(AD - BC)^2}{(A + C)(B + D)(A + B)(C + D)}$$

we then take the maximum χ^2 value for the term:

$$\chi^2(t) = \max\{\chi^2(t, c), \chi^2(t, \bar{c})\}$$

selecting the highest values as our features.

Categorical Proportional Difference (CPD). CPD is a feature selection method proposed by Simeon, M. and Hilderman, R. in [6]. CPD measures the degree to which a word contributes to differentiating a particular category from other categories in a text corpus. It is defined as:

$$CPD(t, c) = \frac{A - B}{A + B}$$

using a Laplace correction the CPD became:

$$\text{CPD}(t, c) = \frac{A - B}{A + B + 2}$$

we then take the maximum CPD value for each term:

$$\text{CPD}(t) = \max\{\text{CPD}(t, c), \text{CPD}(t, \bar{c})\}$$

selecting the highest CPD values as our features.

Mutual Information (MI). MI measures the mutual dependence between a term t and a category c' . It is defined as [6]:

$$\text{MI}(t, c') = \log_2 \frac{AN}{(A + B)(A + C)}$$

we then take the maximum of the MI values for each term:

$$\text{MI}(t) = \max\{\text{MI}(t, c), \text{MI}(t, \bar{c})\}$$

selecting the highest MI values as our features.

Document frequency (DF). DF measures the number of documents in which term t occurs, independent of category:

$$\text{DF}(t) = A + B$$

selecting the highest DF values as our features.

Random feature (RF). RF randomly selects terms t independently of category:

$$\text{RF}(t) = \text{random}()$$

selecting the highest RF values as our features.

5.2 Results

Our Naïve Bayes classifier was run with feature selection performed with correlation coefficient (CC), information gain (IG), mutual information (MI), chi (CHI), document frequency (DF), categorical proportional difference (CPD) and random feature (RF) metrics.

Table VIII: Accuracy of different feature selection methods using our Naïve Bayes and 10-fold cross-validation for different numbers of features

# Features	Feature Selection Metric						
	CC	IG	MI	CHI	DF	CPD	RF
10	89.17	89.50	67.67	89.17	82.83	83.83	57.33
20	91.17	95.83	36.67	91.50	90.67	88.17	65.67
30	90.67	96.33	70.67	91.33	92.33	90.33	42.83
40	91.00	96.50	59.83	90.83	94.17	91.50	37.17
50	90.33	96.17	76.83	91.50	94.00	92.33	43.17
75	90.83	96.17	79.67	91.83	94.17	94.50	75.17
100	91.67	97.17	79.00	92.17	94.67	96.17	51.33
125	91.50	97.17	81.00	93.00	94.67	96.50	41.67
150	92.50	97.00	82.50	94.00	94.50	96.67	50.17
175	92.33	96.83	83.17	94.83	94.83	97.00	52.50
200	91.67	97.50	84.67	95.83	95.33	97.17	54.83
225	91.33	97.83	84.00	95.67	95.33	97.17	73.83
250	90.83	97.33	86.00	95.83	95.33	97.17	52.83

We further compared DF and IG for large numbers of features as presented in Table IX:

Table IX: Accuracy of information gain (IG) and document frequency (DF) with the Naïve Bayes classifier using different numbers of features

# Features	Accuracy of IG (%)	Accuracy of DF (%)
500	98.17	95.17
1000	98.67	97.00
2500	99.83	96.00
5000	98.83	95.50

With a breakdown of the folds for the most accurate results from Table IX :

Table X: Comparison of fold accuracy at $N = 2500$ between information gain (IG) and document frequency (DF) with our Naïve Bayes Classifier

Fold Number	C1 (%)	C2 (%)
Fold 1	100.0	100.0
Fold 2	100.0	98.33
Fold 3	100.0	98.33
Fold 4	100.0	95.00
Fold 5	100.0	91.67
Fold 6	100.0	98.33
Fold 7	100.0	95.00
Fold 8	98.33	93.33
Fold 9	100.0	95.00
Fold 10	100.0	95.00
Mean	99.83	96.00

5.3 Discussion

Information gain yielded the highest accuracy for most of the feature set sizes (Table VII). IG performed best for 12 of the 13 feature set sizes, always beating our original feature selection method of document frequency. We used a paired t-test as described in 4.2.2.

The mean of the differences, \bar{X} , over the feature selection sizes was determined as $\bar{X} = 2.96$ and the sample standard deviation as $s_{N-1} = 1.44$. Calculating the UCL:

$$\begin{aligned}
 \text{UCL}_{0.95} &= \bar{X} \pm t_{0.95,9} \cdot s_{N-1} \\
 &= 2.96 \pm 2.262 \times 1.44 \\
 &= 2.96 \pm 3.26 \\
 &= [-0.30, 6.22]
 \end{aligned}$$

As $\text{UCL}_{0.95} = [-0.30, 6.22]$ contains 0 information gain and document frequency did not produce significantly different results for different numbers of features. Hence we cannot assume that IG always outperforms DF as a feature selection method for our data.

Next we compare IG and DF for the large feature set sizes in using a paired t-test where IG produced the most accurate results, from Table X this was when we selected 2500 features. The mean of the differences, \bar{X} , over the 10-folds was $\bar{X} = 2.96$ and the sample standard deviation

was $s_{N-1} = 1.44$. Calculating the UCL:

$$\begin{aligned} \text{UCL}_{0.95} &= \bar{X} \pm t_{0.95,9} \cdot s_{N-1} \\ &= 3.83 \pm 2.262 \times 2.49 \\ &= 2.96 \pm 5.63 \\ &= [-1.80, 9.46] \end{aligned}$$

As $\text{UCL}_{0.95} = [-1.80, 9.46]$ contains 0 Information gain and document frequency did not produce statistically significantly different results on each fold.

Document frequency and information gain have a linear computational complexity. We expect that DF performed worse than IG since DF only looks at the dependence between the presence of t and $c' \in \{c, \bar{c}\}$ whilst information gain also takes into account the absence of t and $c' \in \{c, \bar{c}\}$. As a result DF selects features that may offer nothing to category prediction (e.g. when t occurs equally in c and \bar{c}). Meanwhile DF deselects features that may offer significantly to category prediction because the term t does not meet the document frequency threshold (i.e. either (t, c) or (t, \bar{c}) is zero but the other does not occur enough to be selected as a feature). On the other hand, IG takes into account all four dependency tuples as described above to select the features that give the most bits of information for category prediction.

Mutual information (MI) and random frequency (RF) both performed weakly. The low accuracy produced by MI is because the score of each term is strongly influenced by marginal probabilities. As a result it performs poorly on our sparse data set. Random frequency was implemented for our own pleasure and as a ‘minimum’ expectation of performance with our Naïve Bayes.

Chi-square (CHI), categorical proportional difference (CPD) and correlation coefficient (CC) performed comparably to document frequency but were not chosen as our feature selection metric. CHI performed similarly for all feature sizes but was outperformed at every size by IG and so was not used. CC underperformed DF for most feature sizes and so was also rejected. CPD performs very well compared to IG and DF, CPD beat DF for all feature sizes greater than 50, even beating IG with a size of 175. However, CPD was rejected because IG returned a better maximum accuracy.

6 Conclusion and Future Work

We implemented a Naïve Bayes classifier using document frequency as our feature selection method with 200 features selected. Our Naïve Bayes produced an accuracy of 95.33% on body, and 80.83% on subject, these were both more accurate than Weka’s classification accuracy of 94.83% on body, and 68.50%.

From our extension we found the information gain feature selection metric was most accurate. At a feature set size of 2500 it produced the highest accuracy of 99.83% and 97.50% with 200 features. This yielded an improvement in accuracy of 3.83% and 2.16% respectively.

Our future research work will deal with creating more meaningful features. This includes implementing stemming and creating tokens using more than one word. Currently, the tokens used in our classifier are formed from single words. Therefore, it will not analyse common consecutive

words that are found in spam emails. By taking into account permutations of consecutive words, or words that appear within a specified distance of each other, the accuracy of our Bayesian classifier could be increased.

7 Reflection

This assignment has helped us learn more about how to conduct research as a team. We chose to conduct research on different aspects of text classification, before combining our findings to select the best extension to carry out. We also learnt how to conduct experiments that were relevant to our aim to increase the accuracy of our classifier algorithm. We appreciate that this task has allowed us gain insight into an interesting and relevant area in computer science.

8 Instructions

8.1 Prerequisites

- Either python3.3 or python3.4 installed.
- *lingspam-mini600* data set placed in the working directory (i.e. same directory as *process.py* and *naivebayes.py*).

8.2 Running Scripts

1. Use `python process.py` to run preprocessing script to create *body.csv* and *subject.csv*. These files are then accessible in the same directory.
2. Use `python naivebayes.py` to run Naïve Bayes classifier on the csv files. Result of accuracy calculation on both subject and body will be output.

You can change the feature selection metric in the preprocessing by changing the function used to create `body_features` and `sub_features` in *process.py*.

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