

Kernel's Correlation for Improving SVM in Text Documents' Classification¹

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Abstract — Support Vector Machine (SVM) is a powerful method of classification, based on kernels, working with large data sets. In almost all cases there are used two distinct parameters that can be modified for obtaining best results. One of these parameters is easy to infer but the second is usually used as being the number of features that are taken into consideration. This seems to be not a good idea for text classification because it was shown that for a great number of features the classification accuracy is quite poor. In this paper we propose a method to correlate those parameters in order to obtain better results and vary only one parameter. We intend to scale only the degree for the polynomial kernel and only constant C for the Gaussian kernel. We also show that using this method leads in almost all cases to better results. We introduce two formulas to correlate the parameters for polynomial and Gaussian kernels.

Keywords — Learning with Kernels, Support Vector Machine, and Text Classification

I. INTRODUCTION

DOCUMENTS are typically represented as vectors of the features space. Each word in the vocabulary represents a dimension of the feature space. The number of occurrences of a word in a document represents the value of the corresponding component in the document's vector [7]. The native feature space consists of the unique terms that occur into the documents, which can be tens or hundreds of thousands of terms for even a moderate-sized text collection, being a major problem of text categorization.

As a method for text classification we use Support Vector Machine technique, which was proved as being efficient for nonlinear separable input data. This is a relatively recent learning method based on kernels [17], [18]. We use this method both in features selection step and in the classification step. Also we studied the influence of the input data representation on kernel parameters correlation.

The idea of the kernel is to compute the norm of the difference between two vectors in a higher dimensional space without representing those vectors in the new space. Adding a scalar constant to the kernel involves better classifying results. In this paper we tested a new idea to correlate this scalar with the dimension of the space where

the data will be represented. Thus we consider that those two parameters (the degree and the scalar) need to be correlated.

In this paper we present a comparative study of different parameters for two types of kernels, polynomial and Gaussian, and different methods to correlate kernel's parameters. In almost all articles where the SVM method is used, researchers used two parameters to infer the kernel but usually they modified only one of them [5], [6]. For the second parameter it is not explicitly specified the modification rule. Sometimes this parameter is chosen as the number of features. For some kinds of applications, when the number of features is not so large, this can be a good idea; but for text classification, when the number of features can be a great, this can lead to powerless results.

The general process of classifying text data can be considered as having four steps. In the first step is done feature extraction from the text file (text mining). In the second step the features are selected. The third step is the learning step (training). The last is the evaluation step, where the classification process is evaluated [13].

In the first step, we have used text mining like an application of data mining techniques to extract the feature vectors that characterizes a document. Starting with a set of d documents and t terms (words belonging to the documents), we can model each document as a vector v in the t dimensional space \mathbb{R}^t (a feature vector that characterizes the document into the set of documents).

In the second step we used SVM technique as a method of feature selection in order to reduce the features space dimension and to select the best features. In [9], [10], [11] SVM feature selection was proved to be the best one. For the input data we have used three types of representations: *Binary representation*, *Nominal representation*, and *Cornell SMART*.

In the next step (classification), we have also used Support Vector Machine. A great advantage of this technique is that it can use large input sets. We implemented this classification method for two types of kernels: *polynomial kernel* and *Gaussian kernel (Radial Basis Function - RBF)*. We tried to find a simplified form of the kernels, without reducing the performance, actually increasing it, using only one, more intuitive parameter.

For multi-class classification we chose the well-known method "one class versus the rest" [16]. Thus, considering M classes, we repeated two class classification for each topic (the category where the document is classified) obtaining M decision functions [4].

Section II contains prerequisites for the work that we present in this paper. In section III we present the frame and methodology used for our experiments. Section IV

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presents the main results of our experiments. Section V debates and concludes on the most important results obtained and proposes some further work.

II. SUPPORT VECTOR MACHINE

The Support Vector Machine (SVM) is a classification technique based on statistical learning theory [14], [16] that was applied with great success in many challenging non-linear classification problems and was successfully applied to large data sets.

The SVM algorithm finds a hyperplane that optimally splits the training set. The optimal hyperplane can be distinguished by the maximum margin of separation between all training points and the hyperplane (a practical idea about implementation of this algorithm can be found in [2]). Looking at a two-dimensional problem we actually want to find a line that “best” separates points in the positive class from points in the negative class. The hyperplane is characterized by a decision function like:

$$f(x) = \text{sgn}(\langle \mathbf{w}, \Phi(x) \rangle + b), \quad (1)$$

where \mathbf{w} is the weight vector, orthogonal to the hyperplane, “ b ” is a scalar that represents the margin of the hyperplane, “ x ” is the current sample tested, “ $\Phi(x)$ ” is a function that transforms the input data into a higher dimensional feature space and $\langle \cdot, \cdot \rangle$ representing the dot product. Sgn is the signum function that returns 1 if the value is greater or equal to 0 and -1 otherwise. If \mathbf{w} has unit length, then $\langle \mathbf{w}, \Phi(x) \rangle$ is the length of $\Phi(x)$ along the direction of \mathbf{w} . Generally \mathbf{w} will be scaled by $\|\mathbf{w}\|$. The training part the algorithm needs to find the normal vector “ \mathbf{w} ” that leads to the largest “ b ” of the hyperplane.

The problem seems very easy to be solved but we have to keep in mind that the optimal classification line should classify correctly all the elements generated by the same given distribution. There are a lot of hyperplanes that meet the classification requirements but the algorithm tries to determine the optimum one. This learning algorithm can be performed in a dot product space and for data which is linear separable, by constructing f from empirical data. It is based on two facts. First, among all the hyperplanes separating the data, there is a unique optimal hyperplane, distinguished by the maximum margin of separation between any training point and the hyperplane. Second, the capacity of the hyperplane to separate the classes decreases with the increasing of the margin.

For training data which is not separable by a hyperplane in the input space the idea of SVM is to map the training data into a *higher-dimensional feature space* via Φ , and construct a separating hyperplane with the maximum margin there. This yields a non-linear decision boundary into the input space. By the use of a kernel function $\langle \mathbf{w}, \phi(x) \rangle$ it is possible to compute the separating hyperplane without explicitly carrying out the map into the feature space [16].

In order to find the optimal hyperplane, we need to solve the following objective function:

$$\underset{\mathbf{w} \in H, b \in \mathbb{R}}{\text{minimize}} \tau(\mathbf{w}) = \frac{1}{2} \|\mathbf{w}\|^2 \quad (2)$$

subject to $y_i (\langle \mathbf{w}, \mathbf{x}_i \rangle + b) \geq 1$ for all $i=1, \dots, m$

The constraints ensure that $f(x_i)$ will be +1 for $y_i=+1$ and -1 for $y_i=-1$. This problem is computationally attractive because it can be constructed by solving a quadratic programming problem for which efficient algorithms already exist. Function τ is called objective function with inequality constrain. Together, they form a so-called *primal optimization problem*. To solve this type of problems it is more convenient to deal with the dual problem by introducing the Lagrange multipliers $\alpha_i \geq 0$ and the Lagrangian [14], [16] which lead to the so-called *dual optimization problem*.

$$L(\mathbf{w}, b, \alpha) = \frac{1}{2} \|\mathbf{w}\|^2 - \sum_{i=1}^m \alpha_i (y_i (\langle \mathbf{x}_i, \mathbf{w} \rangle + b) - 1) \quad (3)$$

,with Lagrange multipliers $\alpha_i \geq 0$. The Lagrangian L must be maximized with respect to the dual variables α_i , and minimized with respect to the primal variables \mathbf{w} and b . This leads to:

$$\mathbf{w} = \sum_{i=0}^m \alpha_i y_i \mathbf{x}_i, \text{ and } \sum_{i=0}^m \alpha_i y_i = 0 \quad (4)$$

The solution vector thus has an expansion in terms of training examples. Note that although the solution \mathbf{w} is unique (due to the strict convexity of primal optimization problem), the coefficients α_i need not be. According to the *Karush-Kuhn-Tucker (KKT) theorem*, only the Lagrange multipliers α_i that are non-zero at the saddle point, correspond to constraints that are precisely met. Formally, for all $i=1, \dots, m$, it can be written:

$$\alpha_i [y_i (\langle \mathbf{x}_i, \mathbf{w} \rangle + b) - 1] = 0 \text{ for all } i=1, \dots, m \quad (5)$$

The samples x_i for which $\alpha_i > 0$ are called *Support Vectors*. According to the KKT condition they lie exactly on the margin. All remaining training samples are irrelevant. By eliminating the primal variables \mathbf{w} and b in the Lagrangian we arrive to the so-called dual optimization problem, which is the problem that one usually solves in practice.

$$\underset{\alpha \in \mathbb{R}^m}{\text{maximize}} W(\alpha) = \sum_{i=1}^m \alpha_i - \frac{1}{2} \sum_{i,j=1}^m \alpha_i \alpha_j y_i y_j \langle \mathbf{x}_i, \mathbf{x}_j \rangle \quad (6)$$

Thus the hyperplane can be written in the dual optimization problem as:

$$f(x) = \text{sgn} \left(\sum_{i \in \mathbb{R}} y_i \alpha_i \langle \mathbf{x}_i, x \rangle + b \right) \quad (7)$$

Everything was formulated in a dot product space. On the practical level, changes have to be made to perform the algorithm in a higher-dimensional feature space. Thus the new patterns $\Phi(x_i)$ can equally well be the result of mapping the original input patterns x_i into a higher dimensional feature space using function Φ . Maximizing the target function and evaluating the decision function involve the computation of dot products $\langle \phi(x), \phi(x) \rangle$ in a higher dimensional space. These expensive calculations

are reduced significantly by using a positive definite kernel k , such that $k(x, x') = \langle x, x' \rangle$. This substitution, which is referred sometimes as the *kernel trick* [16] is used to extend hyperplane classification to nonlinear Support Vector Machines. The kernel trick can be applied since all feature vectors only occur in dot products. The weight vectors then becomes an expression in the feature space, and therefore Φ will be the function through which we represent the input vector in the new space. Thus we obtain the *decision function* as the following form:

$$f(x) = \text{sgn} \left(\sum_{i \in \mathfrak{R}} y_i \alpha_i k(x, x_i) + b \right) \quad (8)$$

III. EXPERIMENTAL FRAMEWORK

A. The used dataset

Our experiments are performed on the Reuters-2000 collection [15], which has 984Mb of newspapers articles in a compressed format. Collection includes a total of 806,791 documents, with news stories published by Reuters Press covering the period from 20.07.1996 through 19.07.1997. The articles have 9822391 paragraphs and contain 11522874 sentences and 310033 distinct root words. Documents are pre-classified according to 3 categories: by the *Region* (366 regions) the article refers to, by *Industry Codes* (870 industry codes) and by *Topics* proposed by Reuters (126 topics, 23 of them contain no articles). Due to the huge dimensionality of the database we will present here results obtained using a subset of data. From all documents we selected the documents for which the industry code value is equal to “System software”. We obtained 7083 files that are represented using 19038 features and 68 topics. We represent documents as vectors of words, applying a stop-word filter (from a standard set of 510 stop-words) and extracting the word stem. From these 68 topics we have eliminated those topics that are poorly or excessively represented. Thus we eliminated those topics that contain less than 1% documents from all 7083 documents in the entire set. We also eliminated topics that contain more than 99% samples from the entire set, as being excessively represented. After doing so we obtained 24 different topics and 7053 documents that were split randomly in training set (4702 samples) and evaluation set (2351 samples). In the feature extraction part we take into consideration both the article and the title of the article.

B. Used kernel types

The idea of the kernel is to compute the norm of the difference between two vectors in a higher dimensional space without representing those vectors in the new space. In practice we can see that by adding a scalar constant to the kernel we can get better classifying results. In this paper we tested a new idea to correlate this scalar with the dimension of the space where the data will be represented. We consider that those two parameters (the degree and the scalar) need to be correlated.

For the polynomial kernel we change only the polynomial degree and for the Gaussian kernel we change

only the constant C according to the new following formulas (x and x' being the input vectors):

$$\text{Polynomial} \quad k(x, x') = (2 \cdot d + \langle x \cdot x' \rangle)^d \quad (9)$$

d being the only parameter that need to be modified and

$$\text{Gaussian} \quad k(x, x') = \exp \left(- \frac{\|x - x'\|^2}{n \cdot C} \right) \quad (10)$$

where C being the only parameter that need to be modified and n being the number of elements greater than 0 from the input vectors that is automatically computed.

C. Polynomial kernel parameter's correlation

Usually when learning with a polynomial kernel researchers use a kernel that looks like:

$$(\langle \mathbf{x} \cdot \mathbf{x}' \rangle + b)^d \quad (11)$$

where d and b are independent parameters. “ d ” is the degree of the kernel and it is used as a parameter that helps mapping the input data into a higher dimensional features space. This is why this parameter is intuitive. The second parameter “ b ”, is not so easy to infer. In all studied articles, the researchers used it, but they don't present a method for selecting it. We notice that if this parameter was eliminated (i.e., chosen zero) the quality of results can be poor. It is logically that we need to correlate d and b parameters because the offset b needs to be modified as the dimension of the space modifies. Due to this, based on running laborious classification simulations presented in this paper, we suggest using “ $b=2*d$ ” in our application.

D. Gaussian kernel parameter's correlation

Also for the Gaussian kernel we modified the standard kernel used by research community. Usually the kernel looks like:

$$k(x, x') = \exp \left(- \frac{\|x - x'\|}{C} \right) \quad (12)$$

where the parameter C is a number that represents the dimension of the training set (and usually this is a very big number for text classification). We introduce the parameter n that is multiplied by parameter C and is a value that represents the number of distinct features that occur into the current two input vectors (x and x'), having weights greater than 0. We keep the constant C that becomes a small number (usually obtain best results between 1 and 2). As far as we know, we are the first authors proposing a correlation between these two parameters for both polynomial and Gaussian kernels.

E. Representing the input data

Because there are many ways to define the feature-weight, we represent the input data in different formats, and we try to analyze their influence on classification accuracy. We take in consideration three formats for representing data [1].

Binary representation – in the input vector we store “0” if the word doesn't occur in the document and “1” if it occurs without considering the number of occurrences. The weight can only be 0 or 1.

Nominal representation – in the input vector we compute the value of the weight using the formula:

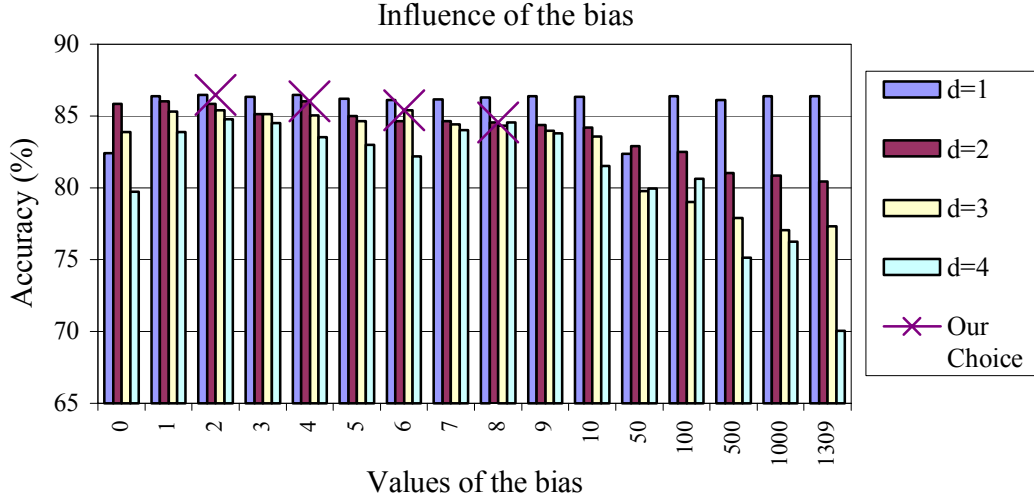


Figure 1 – Influence of bias for Nominal representation of the data

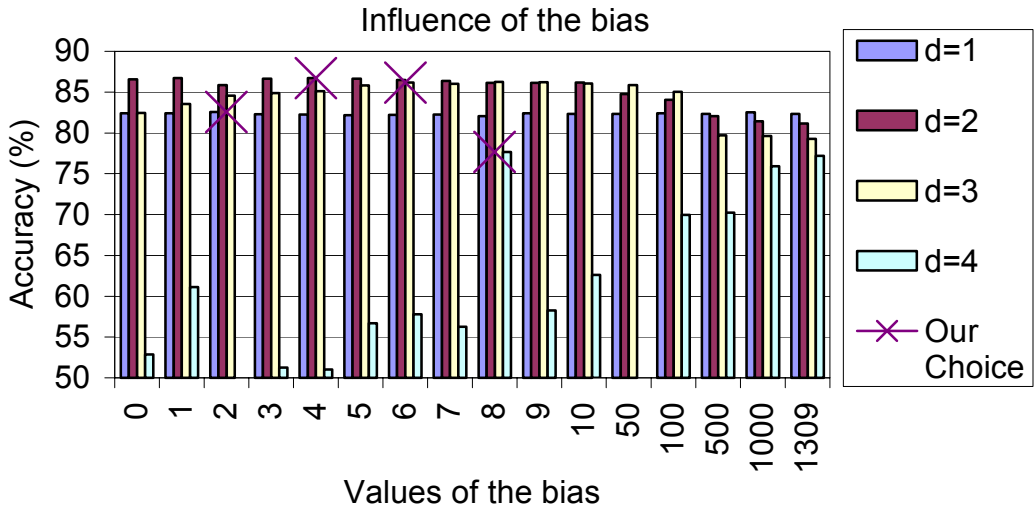


Figure 2 – Influence of bias for Binary data representation

$$TF(d,t) = \frac{n(d,t)}{\max_{\tau} n(d,\tau)}, \quad (13)$$

where $n(d,t)$ is the number of times that term t occurs in document d , and the denominator represents the value of term that occurs the most in document d , and $TF(d,t)$ is the term frequency.

Cornell SMART representation – in the input vector we compute the value of the weight using the formula:

$$TF(d,t) = \begin{cases} 0 & \text{if } n(d,t) = 0 \\ 1 + \log(1 + \log(n(d,t))) & \text{otherwise} \end{cases} \quad (14)$$

where $n(d,t)$ represent number of times term t occurs in document d . In this case the weight can take value 0 if the word does not exist into the document and, greater than 1 if it exists. The scaled value creates a gap between the values for no word and the value of one appearance.

IV. SIMULATION RESULTS

For each test we train the algorithm using the training files. After that we test the decision functions obtained using the test files. We classify the document into the class

for which is obtained the greater absolute value. The result (class) is compared with the results (classes) that were proposed by Reuters. We obtain thus the accuracy of classification for all 24 classes. We'll present all tests using only 1309 features obtained from all 19038 features, using a feature selection method based also to Support Vector Machine [3], [11].

A. Results for polynomial kernel

In order to improve the classification accuracy using polynomial kernel our idea was to correlate the bias of the kernel with the degree of the kernel (Section III.C). In this idea we develop tests for four degrees of the kernel, considering for each of them 16 distinct values of the bias and, respectively, for each input data representation. Thus for each degree of the kernel we vary the value of the bias from 0 to the number of the features. Usually in the literature the bias is selected between 0 and the total number of features.

In Figure 1 we present results obtained with polynomial kernel and Nominal data representation by varying the

Table 1: Influence of bias for CORNELL SMART data representation.

Bias	D=1	D=2	D=3	Choice
0	81.84	86.69	82.35	
1	81.84	86.64	83.37	
2	82.22	86.81	84.01	82.22
3	82.22	86.56	84.77	
4	82.22	86.81	65.12	86.81
5	82.01	86.47	85.54	
6	81.71	86.60	86.39	86.39
7	81.71	86.43	86.18	
8	82.09	86.43	86.47	
9	81.84	86.18	86.47	
10	81.80	85.96	86.26	
50	81.92	84.73	84.90	
100	82.22	83.71	82.82	
500	82.05	81.88	8.34	
1000	82.09	80.94	53.51	
1309	82.09	80.77	50.40	

degree of the kernel and the bias. In “Our choice” entry we put only the values that were obtained using our formula that correlates the polynomial kernel’s parameters. As it can be observed, using our correlation idea (formula 9) assures that in almost all cases there are obtained the best results. In this case, only for degree 4 the best value was obtained for bias equal with 2 and we obtain a value with 0.21% smaller than the best results (84.56% in comparison with the best obtained 84.77%).

Effective values of the accuracy obtained using Cornell Smart data representation, for each kernel degree and for each bias, are presented in Table 1, with bold are represented the best obtained values. For each degree there are multiple bias values involving best results and our proposed formula assures to hit these values in almost all cases. Also an interesting observation is that for kernel degree = 1, we usually obtained the same classification accuracy for all bias values, with only 0.51% smaller than the best value. As it can be observed from Figure 1 and Figure 2, with no bias we obtain the worst results. The same tests were developed also for Binary data representation and we have obtained usually the same results as can be see in Figure 2.

B. Results for Gaussian kernel

For the Gaussian kernel we modified the usually used constant C that represent the number of features, with a product between a small number (noted also by C in our formula 3) and a number that is automatically computed (Section III.D). We made tests with four C distinct values. For each of these values, we vary n from 1 to 1309 (total number of the features used). Because our proposed value for n is automatically computed, this number can not be specified by the command line, so that for each value of constant C we specified a value called “auto” (in Figure 3) that means the value automatically computed using our formula.

We made tests only for Binary and Cornell Smart representations of the input data. Into Gaussian kernel we

fill in a parameter that represents the number of elements greater then zero (parameter “ n ” from equation 12). Nominal representation (equation 13) represents all weight values between 0 and 1. When parameter “ n ” is used, all the weights become very close to zero involving very poor classification accuracies (for example, due to its almost zero weight, a certain word really belonging to the document, might be considered as not belonging to that document). So we don’t present here the results obtained using Nominal representation and Gaussian kernel.

In Figure 3 we present results obtained for Binary data representation. When we use our idea to compute “ n ”, we obtained the best results. Also better results were obtained when the value of n is between 50 and 100. This occurs because usually each document uses a small number of features (on average between 50 and 100) in comparison with total features from entire set of documents. When n is equal with the total number of features (usually used into the literature) the accuracy decrees, in average for all testes, with more than 10% in comparison with using an automatically computed value for n . It can also be observed that when the value of parameter n increases the accuracy decrees substantially. The same tests were also made using Cornell Smart data representation (Figure 4), obtaining the same tendency and usually accuracy with 1% better than in Binary case.

In contrast with polynomial kernel, in Gaussian kernel case we obtained best results only with our proposed formula to compute the parameter “ n ”.

C. Kernel’s influence

In this section we present the influence of correlating kernel’s parameters on classification accuracy using other implementation of SVM. In order to do this we make a short comparison between the results that we obtained with usually used implementation of SVM, called LibSvm [19], and our implemented application called UseSvm [12]. LibSvm uses “one versus the one” method for multiclass classification. Our developed UseSvm program uses “one versus the rest” method, as we already mentioned. Reuter’s database, used in our tests, contains strongly overlapped data and in this case the first method usually obtains powerless results.

We have used also sets with 1309 features, obtained using SVM feature selection method. In order to fairly compare LibSvm with our UseSvm, we eliminated, when possible, Reuters overlapped data (for working only with non-overlapped classes, as far as is possible; formally, $\forall i, j = 1, 13, C_i \cap C_j = \emptyset$ for each $i \neq j$). We choose for each sample only first class that was proposed by Reuters. We also eliminated classes that are poorly or excessively represented. We obtained only 13 classes randomly split in two sets and used for training and testing for both LibSvm and UseSvm. Results obtained by LibSvm are poor in comparison with the results of our application, because, despite our efforts, the data are however slightly overlapped. In Figure 5 and Figure 6 we present results obtained for the polynomial kernel and the Gaussian kernel with nominal data representation for both applications. We are using equivalent parameters for both

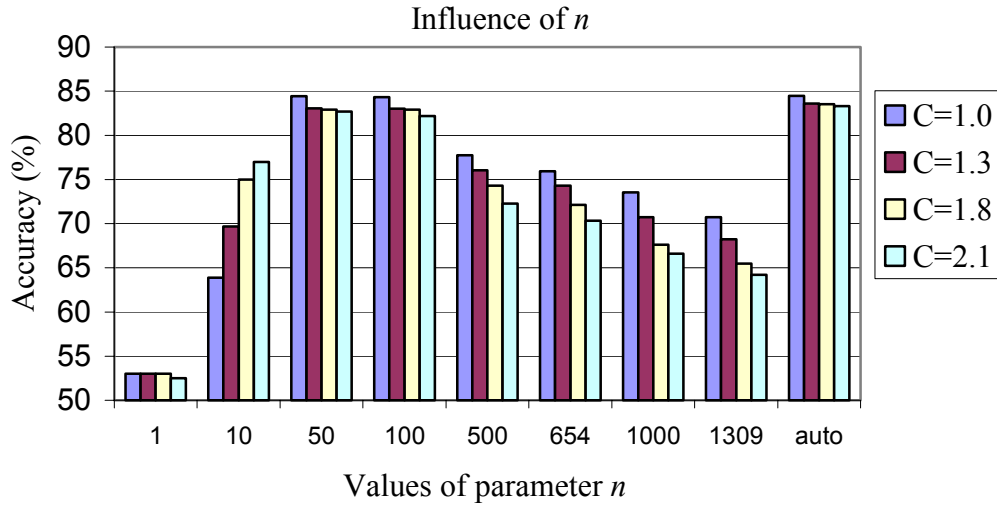


Figure 3 – Influence of value for “ n ” for Gaussian kernel and Binary data representation

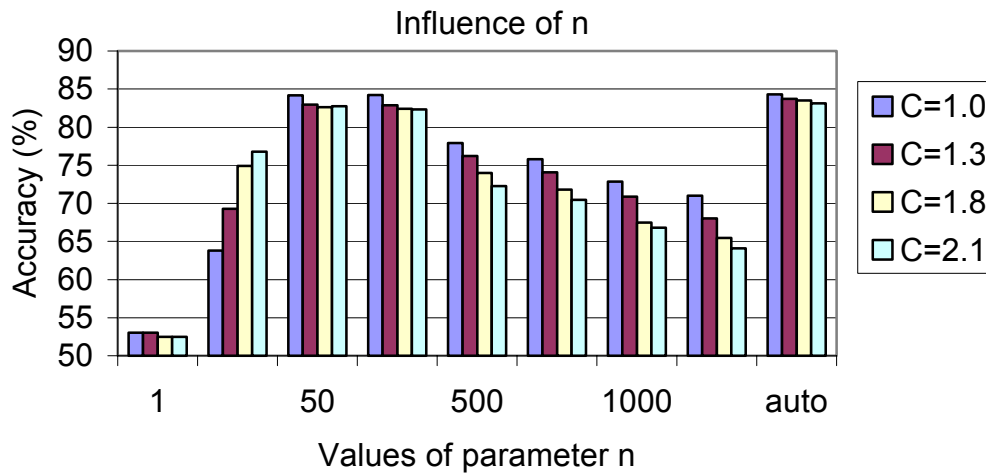


Figure 4 – Influence of parameter n for Gaussian kernel and Cornell Smart data representation

applications. As LibSvm has more parameters than UseSvm, we have left on default value for the parameters that appear only in LibSvm.

As we already specified, for polynomial kernel our suggestion was to make “ $b=2*d$ ” (see Section III.C). We present results using LibSvm with $b=0$ (default value) respectively with $b=2*d$ (specified explicitly by command line) comparing with our UseSvm program.

As it can be observed from Figure 5, our UseSvm program obtains far better results than the well-known LibSvm (with an average gain of 18.82% better). By comparing LibSvm with the default bias with LibSvm with modified bias (according to our formula 9), we noticed that the modified bias leads to better results (with an average gain of 24.26% better). The average gain is computed as average obtained by LibSvm with the default bias divided by the average obtained by LibSvm with modified bias. For degree 1 were obtained similar results because values of default bias and value computed using our formula are quite equal.

For the Gaussian kernel simulations, presented in Figure 6, our suggestion was to multiply the constant C with a

parameter n (like we already explained in Section III.D). It is difficult to give this parameter from the LibSvm’s command line because n is computed dynamically and LibSvm have only one parameter that can be modified, called $gamma$. More precisely, LibSvm uses $gamma=1/n$ only when $gamma$ is default (n means the average of number of attributes in the input data). For LibSvm we have used $gamma$ as $1/C$. For LibSvm+ “ $gamma$ ” we considered “ $gamma$ ” to be equal to $2/nC$, where n is the number of features. The single case of equivalence between these two programs (LibSvm and UseSvm) is obtained for the default value of $gamma$ in LibSvm and respectively for $C=1$ in UseSvm. This case is presented separately as “def” in Figure 6.

As it can be observed, using our idea to modify the Gaussian kernel the results obtained using LibSvm are better in comparison with results obtained using LibSvm with standard kernel (with an average gain of 28.44%). Our UseSvm program obtains far better results than the well-known LibSvm (with an average gain of 25.57% better). For the default parameter of LibSvm our

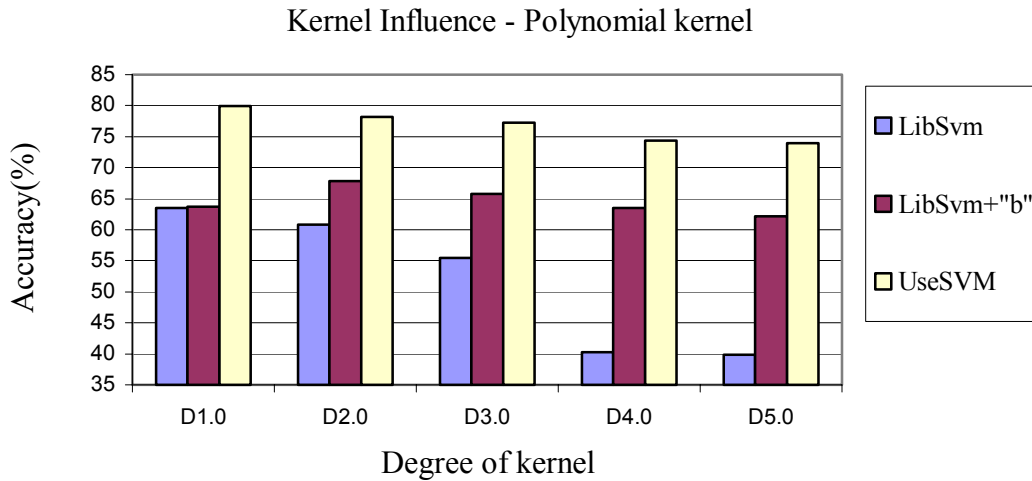


Figure 5 Influence of correlation between parameters from polynomial kernel

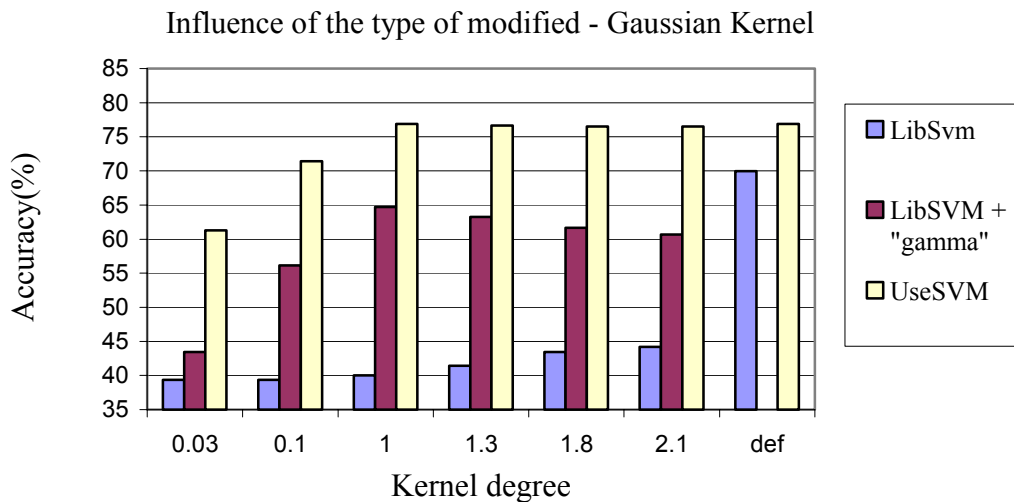


Figure 6 Influence of modified about Gaussian Kernel

application also has obtained better results (76.88% in comparison with 69.97% for LibSvm).

V. CONCLUSIONS AND FURTHER WORK

In this paper, we have proposed a new method to better correlate kernel's parameters. The method correlates the degree of the Polynomial kernel with the bias, respectively correlates the constant from the Gaussian kernel with a value that represents the number of distinct features that occurs into the currently used vectors and having weights greater than 0. In the polynomial kernel case there are more values for which we obtain the best results but our proposed formula assures to hit in almost all cases the best results without having to make more tests to find the good parameter for the bias. For Gaussian kernel our proposed formulas assures to obtain the best results. Also we showed that for text classification problems, using bias or parameter C equal to the number of features, as it is usually used in the literature, is not a good idea. Using our proposed formulas there is obtained in average with 3% better results for polynomial kernel and with 15% better

results for Gaussian kernel.

Also we tested our idea using other SVM implementation, usually used into the literature and called LibSvm, and we obtain the same tendency. We obtained an average accuracy classification gain of 24.26% for polynomial kernel, respectively 28.44% for Gaussian kernel. As far as we know, we are the first authors proposing a correlation between these two parameters for both polynomial and Gaussian kernels.

Work is ongoing to classify larger text data sets (all Reuters database). In this idea we want to develop a pre-classification of all documents, obtaining fewer samples (using simple algorithms like Linear Vector Quantization or Self Organizing Maps). After that we'll use the obtained samples as entry vectors for the already developed features selection and classification methods.

Because almost all available data from a real world are in fact unlabeled data, we will try to combine classifying method with a clustering method, also based on SVM, in order to use labeled and unlabeled data into a hybrid classification algorithm. An interesting natural extension of our algorithm might be an adaptation for a Web mining

application, in order to extract and categorized online news.

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