A Short Introduction to Learning with Kernels^{*}

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Abstract. We briefly describe the main ideas of statistical learning theory, support vector machines, and kernel feature spaces. This includes a derivation of the support vector optimization problem for classification and regression, the ν -trick, various kernels and an overview over applications of kernel methods.

1 An Introductory Example

Suppose we are given empirical data

$$(x_1, y_1), \dots, (x_m, y_m) \in \mathfrak{X} \times \{\pm 1\}.$$
 (1)

Here, the domain \mathfrak{X} is some nonempty set that the patterns x_i are taken from; the y_i are called *labels* or *targets*.

Unless stated otherwise, indices i and j will always be understood to run over the training set, i.e., i, j = 1, ..., m.

Note that we have not made any assumptions on the domain \mathfrak{X} other than it being a set. In order to study the problem of learning, we need additional structure. In learning, we want to be able to *generalize* to unseen data points. In the case of pattern recognition, this means that given some new pattern $x \in \mathfrak{X}$, we want to predict the corresponding $y \in \{\pm 1\}$. By this we mean, loosely speaking, that we choose y such that (x, y) is in some sense similar to the training examples. To this end, we need similarity measures in \mathfrak{X} and in $\{\pm 1\}$. The latter is easy, as two target values can only be identical or different.¹ For the former, we require a similarity measure

$$k: \mathfrak{X} \times \mathfrak{X} \to \mathbb{R},$$

(x, x') $\mapsto k(x, x'),$ (2)

i.e., a function that, given two examples x and x', returns a real number characterizing their similarity. For reasons that will become clear later, the function k is called a *kernel* [13, 1, 8].

A type of similarity measure that is of particular mathematical appeal are dot products. For instance, given two vectors $\mathbf{x}, \mathbf{x}' \in \mathbb{R}^N$, the canonical dot product is defined as

^{*} The present article is based on [23].

¹ If the outputs are not in $\{\pm 1\}$, the situation gets more complex, cf. [34].

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$$(\mathbf{x} \cdot \mathbf{x}') := \sum_{i=1}^{N} (\mathbf{x})_i (\mathbf{x}')_i.$$
(3)

Here, $(\mathbf{x})_i$ denotes the *i*-th entry of \mathbf{x} .

The geometrical interpretation of this dot product is that it computes the cosine of the angle between the vectors \mathbf{x} and \mathbf{x}' , provided they are normalized to length 1. Moreover, it allows computation of the length of a vector \mathbf{x} as $\sqrt{(\mathbf{x} \cdot \mathbf{x})}$, and of the distance between two vectors as the length of the difference vector. Therefore, being able to compute dot products amounts to being able to carry out all geometrical constructions that can be formulated in terms of angles, lengths and distances.

Note, however, that we have not made the assumption that the patterns live in a dot product space. In order to be able to use a dot product as a similarity measure, we therefore first need to embed them into some dot product space \mathcal{H} , which need not be identical to \mathbb{R}^N . To this end, we use a map

$$\begin{split} \Phi &: \mathfrak{X} \to \mathcal{H} \\ & x \mapsto \mathbf{x}. \end{split} \tag{4}$$

The space \mathcal{H} is called a *feature space*. To summarize, embedding the data into \mathcal{H} has three benefits.

1. It lets us define a similarity measure from the dot product in \mathcal{H} ,

$$k(x, x') := (\mathbf{x} \cdot \mathbf{x}') = (\Phi(x) \cdot \Phi(x')).$$
(5)

- 2. It allows us to deal with the patterns geometrically, and thus lets us study learning algorithm using linear algebra and analytic geometry.
- 3. The freedom to choose the mapping Φ will enable us to design a large variety of learning algorithms. For instance, consider a situation where the inputs already live in a dot product space. In that case, we could directly define a similarity measure as the dot product. However, we might still choose to first apply a nonlinear map Φ to change the representation into one that is more suitable for a given problem and learning algorithm.

We are now in the position to describe a pattern recognition learning algorithm that is arguably one of the simplest possible. The basic idea is to compute the means of the two classes in feature space,

$$\mathbf{c}_{+} = \frac{1}{m_{+}} \sum_{\{i:y_i=+1\}} \mathbf{x}_i,\tag{6}$$

$$\mathbf{c}_{-} = \frac{1}{m_{-}} \sum_{\{i:y_{i}=-1\}} \mathbf{x}_{i},\tag{7}$$

where m_+ and m_- are the number of examples with positive and negative labels, respectively (see Figure 1). We then assign a new point **x** to the class whose



Fig. 1. A simple geometric classification algorithm: given two classes of points (depicted by 'o' and '+'), compute their means $\mathbf{c}_+, \mathbf{c}_-$ and assign a test pattern \mathbf{x} to the one whose mean is closer. This can be done by looking at the dot product between $\mathbf{x} - \mathbf{c}$ (where $\mathbf{c} = (\mathbf{c}_+ + \mathbf{c}_-)/2$) and $\mathbf{w} := \mathbf{c}_+ - \mathbf{c}_-$, which changes sign as the enclosed angle passes through $\pi/2$. Note that the corresponding decision boundary is a hyperplane (the dotted line) orthogonal to \mathbf{w} (from [23]).

mean is closer to it. This geometrical construction can be formulated in terms of dot products. Half-way in between \mathbf{c}_+ and \mathbf{c}_- lies the point $\mathbf{c} := (\mathbf{c}_+ + \mathbf{c}_-)/2$. We compute the class of \mathbf{x} by checking whether the vector connecting \mathbf{c} and \mathbf{x} encloses an angle smaller than $\pi/2$ with the vector $\mathbf{w} := \mathbf{c}_+ - \mathbf{c}_-$ connecting the class means, in other words

$$y = \operatorname{sgn} \left((\mathbf{x} - \mathbf{c}) \cdot \mathbf{w} \right)$$

$$y = \operatorname{sgn} \left((\mathbf{x} - (\mathbf{c}_{+} + \mathbf{c}_{-})/2) \cdot (\mathbf{c}_{+} - \mathbf{c}_{-}) \right)$$

$$= \operatorname{sgn} \left((\mathbf{x} \cdot \mathbf{c}_{+}) - (\mathbf{x} \cdot \mathbf{c}_{-}) + b \right).$$
(8)

Here, we have defined the offset

$$b := \frac{1}{2} \left(\|\mathbf{c}_{-}\|^{2} - \|\mathbf{c}_{+}\|^{2} \right).$$
(9)

It will prove instructive to rewrite this expression in terms of the patterns x_i in the input domain \mathcal{X} . To this end, note that we do not have a dot product in \mathcal{X} , all we have is the similarity measure k (cf. (5)). Therefore, we need to rewrite everything in terms of the kernel k evaluated on input patterns. To this end, substitute (6) and (7) into (8) to get the *decision function*

$$y = \operatorname{sgn}\left(\frac{1}{m_{+}}\sum_{\{i:y_{i}=+1\}} (\mathbf{x} \cdot \mathbf{x}_{i}) - \frac{1}{m_{-}}\sum_{\{i:y_{i}=-1\}} (\mathbf{x} \cdot \mathbf{x}_{i}) + b\right)$$
$$= \operatorname{sgn}\left(\frac{1}{m_{+}}\sum_{\{i:y_{i}=+1\}} k(x, x_{i}) - \frac{1}{m_{-}}\sum_{\{i:y_{i}=-1\}} k(x, x_{i}) + b\right).$$
(10)

Similarly, the offset becomes

$$b := \frac{1}{2} \left(\frac{1}{m_{-}^{2}} \sum_{\{(i,j): y_{i} = y_{j} = -1\}} k(x_{i}, x_{j}) - \frac{1}{m_{+}^{2}} \sum_{\{(i,j): y_{i} = y_{j} = +1\}} k(x_{i}, x_{j}) \right).$$
(11)

Let us consider one well-known special case of this type of classifier. Assume that the class means have the same distance to the origin (hence b = 0), and that k can be viewed as a density, i.e., it is positive and has integral 1,

$$\int_{\mathfrak{X}} k(x, x') dx = 1 \quad \text{for all } x' \in \mathfrak{X}.$$
(12)

In order to state this assumption, we have to require that we can define an integral on \mathcal{X} .

If the above holds true, then (10) corresponds to the so-called Bayes decision boundary separating the two classes, subject to the assumption that the two classes were generated from two probability distributions that are correctly estimated by the *Parzen windows* estimators of the two classes,

$$p_1(x) := \frac{1}{m_+} \sum_{\{i:y_i=+1\}} k(x, x_i)$$
(13)

$$p_2(x) := \frac{1}{m_-} \sum_{\{i: y_i = -1\}} k(x, x_i).$$
(14)

Given some point x, the label is then simply computed by checking which of the two, $p_1(x)$ or $p_2(x)$, is larger, which directly leads to (10). Note that this decision is the best we can do if we have no prior information about the probabilities of the two classes. For further details, see [23].

The classifier (10) is quite close to the types of learning machines that we will be interested in. It is linear in the feature space, while in the input domain, it is represented by a kernel expansion in terms of the training points. It is examplebased in the sense that the kernels are centered on the training examples, i.e., one of the two arguments of the kernels is always a training example. The main point where the more sophisticated techniques to be discussed later will deviate from (10) is in the selection of the examples that the kernels are centered on, and in the weight that is put on the individual kernels in the decision function. Namely, it will no longer be the case that *all* training examples appear in the kernel expansion, and the weights of the kernels in the expansion will no longer be uniform. In the feature space representation, this statement corresponds to saying that we will study all normal vectors \mathbf{w} of decision hyperplanes that can be represented as linear combinations of the training examples. For instance, we might want to remove the influence of patterns that are very far away from the decision boundary, either since we expect that they will not improve the generalization error of the decision function, or since we would like to reduce the computational cost of evaluating the decision function (cf. (10)). The hyperplane will then only depend on a subset of training examples, called *support vectors*.

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2 Learning Pattern Recognition from Examples

With the above example in mind, let us now consider the problem of pattern recognition in a more formal setting [28, 29] following the introduction of [19]. In two-class pattern recognition, we seek to estimate a function

$$f: \mathcal{X} \to \{\pm 1\} \tag{15}$$

based on input-output training data (1). We assume that the data were generated independently from some unknown (but fixed) probability distribution P(x, y). Our goal is to learn a function that will correctly classify unseen examples (x, y), i.e., we want f(x) = y for examples (x, y) that were also generated from P(x, y).

If we put no restriction on the class of functions that we choose our estimate f from, however, even a function which does well on the training data, e.g. by satisfying $f(x_i) = y_i$ for all i = 1, ..., m, need not generalize well to unseen examples. To see this, note that for each function f and any test set $(\bar{x}_1, \bar{y}_1), ..., (\bar{x}_{\bar{m}}, \bar{y}_{\bar{m}}) \in \mathbb{R}^N \times \{\pm 1\}$, satisfying $\{\bar{x}_1, ..., \bar{x}_{\bar{m}}\} \cap \{x_1, ..., x_m\} = \{\}$, there exists another function f^* such that $f^*(x_i) = f(x_i)$ for all i = 1, ..., m, yet $f^*(\bar{x}_i) \neq f(\bar{x}_i)$ for all $i = 1, ..., \bar{m}$. As we are only given the training data, we have no means of selecting which of the two functions (and hence which of the completely different sets of test label predictions) is preferable. Hence, only minimizing the training error (or *empirical risk*),

$$R_{emp}[f] = \frac{1}{m} \sum_{i=1}^{m} \frac{1}{2} |f(x_i) - y_i|, \qquad (16)$$

does not imply a small test error (called risk), averaged over test examples drawn from the underlying distribution P(x, y),

$$R[f] = \int \frac{1}{2} |f(x) - y| \, dP(x, y). \tag{17}$$

Statistical learning theory [30, 28, 29], or VC (Vapnik-Chervonenkis) theory, shows that it is imperative to restrict the class of functions that f is chosen from to one which has a *capacity* that is suitable for the amount of available training data. VC theory provides *bounds* on the test error. The minimization of these bounds, which depend on both the empirical risk and the capacity of the function class, leads to the principle of *structural risk minimization* [28]. The best-known capacity concept of VC theory is the VC dimension, defined as the largest number h of points that can be separated in all possible ways using functions of the given class. An example of a VC bound is the following: if h < m is the VC dimension of the class of functions that the learning machine can implement, then for all functions of that class, with a probability of at least $1 - \eta$, the bound

$$R(\alpha) \le R_{emp}(\alpha) + \phi\left(\frac{h}{m}, \frac{\log(\eta)}{m}\right)$$
(18)

holds, where the *confidence term* ϕ is defined as

$$\phi\left(\frac{h}{m}, \frac{\log(\eta)}{m}\right) = \sqrt{\frac{h\left(\log\frac{2m}{h}+1\right) - \log(\eta/4)}{m}}.$$
(19)

Tighter bounds can be formulated in terms of other concepts, such as the *annealed VC entropy* or the *Growth function*. These are usually considered to be harder to evaluate, but they play a fundamental role in the conceptual part of VC theory [29]. Alternative capacity concepts that can be used to formulate bounds include the *fat shattering dimension* [2].

The bound (18) deserves some further explanatory remarks. Suppose we wanted to learn a "dependency" where $P(x,y) = P(x) \cdot P(y)$, i.e., where the pattern x contains no information about the label y, with uniform P(y). Given a training sample of fixed size, we can then surely come up with a learning machine which achieves zero training error (provided we have no examples contradicting each other). However, in order to reproduce the random labelings, this machine will necessarily require a large VC dimension h. Thus, the confidence term (19), increasing monotonically with h, will be large, and the bound (18) will not support possible hopes that due to the small training error, we should expect a small test error. This makes it understandable how (18) can hold independent of assumptions about the underlying distribution P(x, y): it always holds (provided that h < m), but it does not always make a nontrivial prediction — a bound on an error rate becomes void if it is larger than the maximum error rate. In order to get nontrivial predictions from (18), the function space must be restricted such that the capacity (e.g. VC dimension) is small enough (in relation to the available amount of data).

3 Hyperplane Classifiers

In the present section, we shall describe a hyperplane learning algorithm that can be performed in a dot product space (such as the feature space that we introduced previously). As described in the previous section, to design learning algorithms, one needs to come up with a class of functions whose capacity can be computed.

[31] considered the class of hyperplanes

$$(\mathbf{w} \cdot \mathbf{x}) + b = 0 \quad \mathbf{w} \in \mathbb{R}^N, b \in R,$$
(20)

corresponding to decision functions

$$f(\mathbf{x}) = \operatorname{sgn}\left((\mathbf{w} \cdot \mathbf{x}) + b\right),\tag{21}$$

and proposed a learning algorithm for separable problems, termed the *Generalized Portrait*, for constructing f from empirical data. It is based on two facts. First, among all hyperplanes separating the data, there exists a unique one yielding the maximum margin of separation between the classes,

$$\max_{\mathbf{w},b} \min\{\|\mathbf{x} - \mathbf{x}_i\| : \mathbf{x} \in \mathbb{R}^N, (\mathbf{w} \cdot \mathbf{x}) + b = 0, i = 1, \dots, m\}.$$
 (22)

Second, the capacity decreases with increasing margin.



Fig. 2. A binary classification toy problem: separate balls from diamonds. The *optimal* hyperplane is orthogonal to the shortest line connecting the convex hulls of the two classes (dotted), and intersects it half-way between the two classes. The problem being separable, there exists a weight vector \mathbf{w} and a threshold b such that $y_i \cdot ((\mathbf{w} \cdot \mathbf{x}_i) + b) > 0$ (i = 1, ..., m). Rescaling \mathbf{w} and b such that the point(s) closest to the hyperplane satisfy $|(\mathbf{w} \cdot \mathbf{x}_i) + b| = 1$, we obtain a *canonical* form (\mathbf{w}, b) of the hyperplane, satisfying $y_i \cdot ((\mathbf{w} \cdot \mathbf{x}_i) + b) \ge 1$. Note that in this case, the margin, measured perpendicularly to the hyperplane, equals $2/||\mathbf{w}||$. This can be seen by considering two points $\mathbf{x}_1, \mathbf{x}_2$ on opposite sides of the margin, i.e., $(\mathbf{w} \cdot \mathbf{x}_1) + b = 1, (\mathbf{w} \cdot \mathbf{x}_2) + b = -1$, and projecting them onto the hyperplane normal vector $\mathbf{w}/||\mathbf{w}||$ (from [17]).

To construct this *Optimal Hyperplane* (cf. Figure 2), one solves the following optimization problem:

minimize
$$\tau(\mathbf{w}) = \frac{1}{2} \|\mathbf{w}\|^2$$
 (23)

subject to $y_i \cdot ((\mathbf{w} \cdot \mathbf{x}_i) + b) \ge 1, \quad i = 1, \dots, m.$ (24)

This constrained optimization problem is dealt with by introducing Lagrange multipliers $\alpha_i \geq 0$ and a Lagrangian

$$L(\mathbf{w}, b, \boldsymbol{\alpha}) = \frac{1}{2} \|\mathbf{w}\|^2 - \sum_{i=1}^m \alpha_i \left(y_i \cdot \left((\mathbf{x}_i \cdot \mathbf{w}) + b \right) - 1 \right).$$
(25)

The Lagrangian L has to be minimized with respect to the *primal variables* \mathbf{w} and b and maximized with respect to the *dual variables* α_i (i.e., a saddle point has to be found). Let us try to get some intuition for this. If a constraint (24) is violated, then $y_i \cdot ((\mathbf{w} \cdot \mathbf{x}_i) + b) - 1 < 0$, in which case L can be increased by increasing the corresponding α_i . At the same time, \mathbf{w} and b will have to change such that L decreases. To prevent $-\alpha_i (y_i \cdot ((\mathbf{w} \cdot \mathbf{x}_i) + b) - 1)$ from becoming arbitrarily large, the change in \mathbf{w} and b will ensure that, provided the problem is separable, the constraint will eventually be satisfied. Similarly, one can understand that for all constraints which are not precisely met as equalities, i.e., for which $y_i \cdot ((\mathbf{w} \cdot \mathbf{x}_i) + b) - 1 > 0$, the corresponding α_i must be 0: this is the value of α_i that maximizes L. The latter is the statement of the Karush-Kuhn-Tucker complementarity conditions of optimization theory [6].

The condition that at the saddle point, the derivatives of L with respect to the primal variables must vanish,

$$\frac{\partial}{\partial b}L(\mathbf{w}, b, \boldsymbol{\alpha}) = 0, \quad \frac{\partial}{\partial \mathbf{w}}L(\mathbf{w}, b, \boldsymbol{\alpha}) = 0, \tag{26}$$

leads to

$$\sum_{i=1}^{m} \alpha_i y_i = 0 \tag{27}$$

and

$$\mathbf{w} = \sum_{i=1}^{m} \alpha_i y_i \mathbf{x}_i.$$
⁽²⁸⁾

The solution vector thus has an expansion in terms of a subset of the training patterns, namely those patterns whose α_i is non-zero, called *Support Vectors*. By the Karush-Kuhn-Tucker complementarity conditions

$$\alpha_i \cdot [y_i((\mathbf{x}_i \cdot \mathbf{w}) + b) - 1] = 0, \quad i = 1, \dots, m,$$
(29)

the Support Vectors lie on the margin (cf. Figure 2). All remaining examples of the training set are irrelevant: their constraint (24) does not play a role in the optimization, and they do not appear in the expansion (28). This nicely captures our intuition of the problem: as the hyperplane (cf. Figure 2) is completely determined by the patterns closest to it, the solution should not depend on the other examples.

By substituting (27) and (28) into L, one eliminates the primal variables and arrives at the Wolfe dual of the optimization problem (e.g., [6]): find multipliers α_i which

maximize
$$W(\alpha) = \sum_{i=1}^{m} \alpha_i - \frac{1}{2} \sum_{i,j=1}^{m} \alpha_i \alpha_j y_i y_j (\mathbf{x}_i \cdot \mathbf{x}_j)$$
 (30)

subject to
$$\alpha_i \ge 0, \quad i = 1, \dots, m, \text{ and } \sum_{i=1}^m \alpha_i y_i = 0.$$
 (31)

The hyperplane decision function can thus be written as

$$f(\mathbf{x}) = \operatorname{sgn}\left(\sum_{i=1}^{m} y_i \alpha_i \cdot (\mathbf{x} \cdot \mathbf{x}_i) + b\right)$$
(32)

where b is computed using (29).

The structure of the optimization problem closely resembles those that typically arise in Lagrange's formulation of mechanics. Also there, often only a subset of the constraints become active. For instance, if we keep a ball in a box, then it will typically roll into one of the corners. The constraints corresponding to the walls which are not touched by the ball are irrelevant, the walls could just as well be removed.

Seen in this light, it is not too surprising that it is possible to give a mechanical interpretation of optimal margin hyperplanes [9]: If we assume that each support vector \mathbf{x}_i exerts a perpendicular force of size α_i and sign y_i on a solid plane sheet lying along the hyperplane, then the solution satisfies the requirements of mechanical stability. The constraint (27) states that the forces on the sheet sum to zero; and (28) implies that the torques also sum to zero, via $\sum_i \mathbf{x}_i \times y_i \alpha_i \cdot \mathbf{w}/||\mathbf{w}|| = \mathbf{w} \times \mathbf{w}/||\mathbf{w}|| = 0.$

There are theoretical arguments supporting the good generalization performance of the optimal hyperplane [30, 28, 4, 25, 35]. In addition, it is computationally attractive, since it can be constructed by solving a quadratic programming problem.

4 Support Vector Classifiers

We now have all the tools to describe support vector machines [29, 23]. Everything in the last section was formulated in a dot product space. We think of this space as the feature space \mathcal{H} described in Section 1. To express the formulas in terms of the input patterns living in \mathcal{X} , we thus need to employ (5), which expresses the dot product of bold face feature vectors \mathbf{x}, \mathbf{x}' in terms of the kernel k evaluated on input patterns x, x',

$$k(x, x') = (\mathbf{x} \cdot \mathbf{x}'). \tag{33}$$

This can be done since all feature vectors only occured in dot products. The weight vector (cf. (28)) then becomes an expansion in feature space,² and will thus typically no longer correspond to the image of a single vector from input space. We thus obtain decision functions of the more general form (cf. (32))

$$f(x) = \operatorname{sgn}\left(\sum_{i=1}^{m} y_i \alpha_i \cdot (\varPhi(x) \cdot \varPhi(x_i)) + b\right)$$
$$= \operatorname{sgn}\left(\sum_{i=1}^{m} y_i \alpha_i \cdot k(x, x_i) + b\right),$$
(34)

and the following quadratic program (cf. (30)):

² This constitutes a special case of the so-called representer theorem, which states that under fairly general conditions, the minimizers of objective functions which contain a penalizer in terms of a norm in feature space will have kernel expansions [32, 23].



Fig. 3. Example of a Support Vector classifier found by using a radial basis function kernel $k(x, x') = \exp(-||x - x'||^2)$. Both coordinate axes range from -1 to +1. Circles and disks are two classes of training examples; the middle line is the decision surface; the outer lines precisely meet the constraint (24). Note that the Support Vectors found by the algorithm (marked by extra circles) are not centers of clusters, but examples which are critical for the given classification task. Grey values code the modulus of the argument $\sum_{i=1}^{m} y_i \alpha_i \cdot k(x, x_i) + b$ of the decision function (34) (from [17]).)

maximize
$$W(\alpha) = \sum_{i=1}^{m} \alpha_i - \frac{1}{2} \sum_{i,j=1}^{m} \alpha_i \alpha_j y_i y_j k(x_i, x_j)$$
(35)

subject to
$$\alpha_i \ge 0, \quad i = 1, \dots, m, \text{ and } \sum_{i=1}^m \alpha_i y_i = 0.$$
 (36)

In practice, a separating hyperplane may not exist, e.g. if a high noise level causes a large overlap of the classes. To allow for the possibility of examples violating (24), one introduces slack variables [10, 29, 22]

$$\xi_i \ge 0, \quad i = 1, \dots, m \tag{37}$$

in order to relax the constraints to

$$y_i \cdot ((\mathbf{w} \cdot \mathbf{x}_i) + b) \ge 1 - \xi_i, \quad i = 1, \dots, m.$$
(38)

A classifier which generalizes well is then found by controlling both the classifier capacity (via $\|\mathbf{w}\|$) and the sum of the slacks $\sum_i \xi_i$. The latter is done as it can be shown to provide an upper bound on the number of training errors which leads to a convex optimization problem.

One possible realization of a *soft margin* classifier is minimizing the objective function

$$\tau(\mathbf{w}, \boldsymbol{\xi}) = \frac{1}{2} \|\mathbf{w}\|^2 + C \sum_{i=1}^m \xi_i$$
(39)

subject to the constraints (37) and (38), for some value of the constant C > 0 determining the trade-off. Here and below, we use boldface Greek letters as a shorthand for corresponding vectors $\boldsymbol{\xi} = (\xi_1, \ldots, \xi_m)$. Incorporating kernels, and rewriting it in terms of Lagrange multipliers, this again leads to the problem of maximizing (35), subject to the constraints

$$0 \le \alpha_i \le C, \quad i = 1, \dots, m, \text{ and } \sum_{i=1}^m \alpha_i y_i = 0.$$
 (40)

The only difference from the separable case is the upper bound C on the Lagrange multipliers α_i . This way, the influence of the individual patterns (which could be outliers) gets limited. As above, the solution takes the form (34). The threshold b can be computed by exploiting the fact that for all SVs x_i with $\alpha_i < C$, the slack variable ξ_i is zero (this again follows from the Karush-Kuhn-Tucker complementarity conditions), and hence

$$\sum_{j=1}^{m} y_j \alpha_j \cdot k(x_i, x_j) + b = y_i.$$

$$\tag{41}$$

Another possible realization of a soft margin variant of the optimal hyperplane uses the ν -parameterization [22]. In it, the parameter C is replaced by a parameter $\nu \in [0, 1]$ which can be shown to lower and upper bound the number of examples that will be SVs and that will come to lie on the wrong side of the hyperplane, respectively. It uses a primal objective function with the error term $\frac{1}{\nu m} \sum_i \xi_i - \rho$, and separation constraints

$$y_i \cdot ((\mathbf{w} \cdot \mathbf{x}_i) + b) \ge \rho - \xi_i, \quad i = 1, \dots, m.$$

$$(42)$$

The margin parameter ρ is a variable of the optimization problem. The dual can be shown to consist of maximizing the quadratic part of (35), subject to $0 \leq \alpha_i \leq 1/(\nu m)$, $\sum_i \alpha_i y_i = 0$ and the additional constraint $\sum_i \alpha_i = 1$.

5 Support Vector Regression

The concept of the margin is specific to pattern recognition. To generalize the SV algorithm to regression estimation [29], an analog of the margin is constructed in the space of the target values y (note that in regression, we have $y \in \mathbb{R}$) by using Vapnik's ε -insensitive loss function (Figure 4)

$$|y - f(\mathbf{x})|_{\varepsilon} := \max\{0, |y - f(\mathbf{x})| - \varepsilon\}.$$
(43)



Fig. 4. In SV regression, a tube with radius ε is fitted to the data. The trade-off between model complexity and points lying outside of the tube (with positive slack variables ξ) is determined by minimizing (46) (from [17]).

To estimate a linear regression

$$f(\mathbf{x}) = (\mathbf{w} \cdot \mathbf{x}) + b \tag{44}$$

with precision ε , one minimizes

$$\frac{1}{2} \|\mathbf{w}\|^2 + C \sum_{i=1}^m |y_i - f(\mathbf{x}_i)|_{\varepsilon}.$$
(45)

Written as a constrained optimization problem, this reads:

minimize
$$\tau(\mathbf{w}, \boldsymbol{\xi}, \boldsymbol{\xi}^*) = \frac{1}{2} \|\mathbf{w}\|^2 + C \sum_{i=1}^m (\xi_i + \xi_i^*)$$
 (46)

subject to
$$((\mathbf{w} \cdot \mathbf{x}_i) + b) - y_i \le \varepsilon + \xi_i$$
 (47)

$$y_i - ((\mathbf{w} \cdot \mathbf{x}_i) + b) \le \varepsilon + \xi_i^* \tag{48}$$

$$\xi_i, \xi_i^* \ge 0 \tag{49}$$

for all i = 1, ..., m. Note that according to (47) and (48), any error smaller than ε does not require a nonzero ξ_i or ξ_i^* , and hence does not enter the objective function (46).

Generalization to kernel-based regression estimation is carried out in complete analogy to the case of pattern recognition. Introducing Lagrange multipliers, one thus arrives at the following optimization problem: for $C > 0, \varepsilon \ge 0$ chosen a priori,

m

maximize

$$W(\boldsymbol{\alpha}, \boldsymbol{\alpha}^*) = -\varepsilon \sum_{i=1} (\alpha_i^* + \alpha_i) + \sum_{i=1} (\alpha_i^* - \alpha_i) y_i$$
$$-\frac{1}{2} \sum_{i,j=1}^m (\alpha_i^* - \alpha_i) (\alpha_j^* - \alpha_j) k(x_i, x_j)$$
(50)

subject to

$$0 \le \alpha_i, \alpha_i^* \le C, \quad i = 1, \dots, m, \text{ and } \sum_{i=1}^m (\alpha_i - \alpha_i^*) = 0.$$
 (51)

The regression estimate takes the form

$$f(x) = \sum_{i=1}^{m} (\alpha_i^* - \alpha_i) k(x_i, x) + b,$$
(52)

where b is computed using the fact that (47) becomes an equality with $\xi_i = 0$ if $0 < \alpha_i < C$, and (48) becomes an equality with $\xi_i^* = 0$ if $0 < \alpha_i^* < C$.

m

Several extensions of this algorithm are possible. From an abstract point of view, we just need some target function which depends on the vector $(\mathbf{w}, \boldsymbol{\xi})$ (cf. (46)). There are multiple degrees of freedom for constructing it, including some freedom how to penalize, or regularize, different parts of the vector, and some freedom how to use the kernel trick. For instance, more general loss functions can be used for $\boldsymbol{\xi}$, leading to problems that can still be solved efficiently [27]. Moreover, norms other than the 2-norm $\|.\|$ can be used to regularize the solution. Yet another example is that polynomial kernels can be incorporated which consist of multiple layers, such that the first layer only computes products within certain specified subsets of the entries of \mathbf{w} [17].

Finally, the algorithm can be modified such that ε need not be specified a priori. Instead, one specifies an upper bound $0 \le \nu \le 1$ on the fraction of points allowed to lie outside the tube (asymptotically, the number of SVs) and the corresponding ε is computed automatically. This is achieved by using as primal objective function

$$\frac{1}{2} \|\mathbf{w}\|^2 + C\left(\nu m\varepsilon + \sum_{i=1}^m |y_i - f(\mathbf{x}_i)|_\varepsilon\right)$$
(53)

instead of (45), and treating $\varepsilon \geq 0$ as a parameter that we minimize over [22].

We conclude this section by noting that the SV algorithm has not only been generalized to regression, but also, more recently, to one-class problems and novelty detection [20]. Moreover, the kernel method for computing dot products in feature spaces is not restricted to SV machines. Indeed, it has been pointed out that it can be used to develop nonlinear generalizations of any algorithm that can be cast in terms of dot products, such as principal component analysis [21], and a number of developments have followed this example.

6 Polynomial Kernels

We now take a closer look at the issue of the similarity measure, or kernel, k.

In this section, we think of \mathfrak{X} as a subset of the vector space \mathbb{R}^N , $(N \in \mathbb{N})$, endowed with the canonical dot product (3).

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Fig. 5. Architecture of SV machines. The input x and the Support Vectors x_i are nonlinearly mapped (by Φ) into a feature space \mathcal{H} , where dot products are computed. By the use of the kernel k, these two layers are in practice computed in one single step. The results are linearly combined by weights v_i , found by solving a quadratic program (in pattern recognition, $v_i = y_i \alpha_i$; in regression estimation, $v_i = \alpha_i^* - \alpha_i$). The linear combination is fed into the function σ (in pattern recognition, $\sigma(x) = \operatorname{sgn}(x + b)$; in regression estimation, $\sigma(x) = x + b$) (from [17]).

6.1 Product Features

Suppose we are given patterns $x \in \mathbb{R}^N$ where most information is contained in the *d*-th order products (monomials) of entries $[x]_i$ of x,

$$[x]_{j_1}\cdots [x]_{j_d},\tag{54}$$

where $j_1, \ldots, j_d \in \{1, \ldots, N\}$. In that case, we might prefer to *extract* these product features, and work in the feature space \mathcal{H} of all products of d entries. In visual recognition problems, where images are often represented as vectors, this would amount to extracting features which are products of individual pixels.

For instance, in \mathbb{R}^2 , we can collect all monomial feature extractors of degree 2 in the nonlinear map

$$\Phi: \mathbb{R}^2 \to \mathcal{H} = \mathbb{R}^3 \tag{55}$$

$$([x]_1, [x]_2) \mapsto ([x]_1^2, [x]_2^2, [x]_1[x]_2).$$
(56)

This approach works fine for small toy examples, but it fails for realistically sized problems: for N-dimensional input patterns, there exist

$$N_{\mathcal{H}} = \frac{(N+d-1)!}{d!(N-1)!}$$
(57)

different monomials (54), comprising a feature space \mathcal{H} of dimensionality $N_{\mathcal{H}}$. For instance, already 16 × 16 pixel input images and a monomial degree d = 5 yield a dimensionality of 10¹⁰.

In certain cases described below, there exists, however, a way of *computing dot* products in these high-dimensional feature spaces without explicitly mapping into them: by means of kernels nonlinear in the input space \mathbb{R}^N . Thus, if the subsequent processing can be carried out using dot products exclusively, we are able to deal with the high dimensionality.

The following section describes how dot products in polynomial feature spaces can be computed efficiently.

6.2 Polynomial Feature Spaces Induced by Kernels

In order to compute dot products of the form $(\Phi(x) \cdot \Phi(x'))$, we employ kernel representations of the form

$$k(x, x') = (\Phi(x) \cdot \Phi(x')), \tag{58}$$

which allow us to compute the value of the dot product in \mathcal{H} without having to carry out the map Φ . This method was used by [8] to extend the *Generalized Portrait* hyperplane classifier of [30] to nonlinear Support Vector machines. [1] call \mathcal{H} the *linearization space*, and used in the context of the potential function classification method to express the dot product between elements of \mathcal{H} in terms of elements of the input space.

What does k look like for the case of polynomial features? We start by giving an example [29] for N = d = 2. For the map

$$C_2: ([x]_1, [x]_2) \mapsto ([x]_1^2, [x]_2^2, [x]_1[x]_2, [x]_2[x]_1),$$
(59)

dot products in $\mathcal H$ take the form

$$(C_2(x) \cdot C_2(x')) = [x]_1^2 [x']_1^2 + [x]_2^2 [x']_2^2 + 2[x]_1 [x]_2 [x']_1 [x']_2 = (x \cdot x')^2, \quad (60)$$

i.e., the desired kernel k is simply the square of the dot product in input space. The same works for arbitrary $N, d \in \mathbb{N}$ [8]: as a straightforward generalization of a result proved in the context of polynomial approximation (Lemma 2.1, [16]), we have:

Proposition 1. Define C_d to map $x \in \mathbb{R}^N$ to the vector $C_d(x)$ whose entries are all possible d-th degree ordered products of the entries of x. Then the corresponding kernel computing the dot product of vectors mapped by C_d is

$$k(x, x') = (C_d(x) \cdot C_d(x')) = (x \cdot x')^d.$$
(61)

Proof. We directly compute

$$(C_d(x) \cdot C_d(x')) = \sum_{j_1, \dots, j_d=1}^N [x]_{j_1} \cdots [x]_{j_d} \cdot [x']_{j_1} \cdots [x']_{j_d}$$
(62)

$$= \left(\sum_{j=1}^{N} [x]_{j} \cdot [x']_{j}\right)^{a} = (x \cdot x')^{d}.$$
 (63)

Instead of ordered products, we can use unordered ones to obtain a map Φ_d which yields the same value of the dot product. To this end, we have to compensate for the multiple occurrence of certain monomials in C_d by scaling the respective entries of Φ_d with the square roots of their numbers of occurrence. Then, by this definition of Φ_d , and (61),

$$(\Phi_d(x) \cdot \Phi_d(x')) = (C_d(x) \cdot C_d(x')) = (x \cdot x')^d.$$
(64)

For instance, if n of the j_i in (54) are equal, and the remaining ones are different, then the coefficient in the corresponding component of Φ_d is $\sqrt{(d-n+1)!}$ (for the general case, cf. [24]. For Φ_2 , this simply means that [29]

$$\Phi_2(x) = ([x]_1^2, [x]_2^2, \sqrt{2} \ [x]_1[x]_2).$$
(65)

If x represents an image with the entries being pixel values, we can use the kernel $(x \cdot x')^d$ to work in the space spanned by products of any d pixels provided that we are able to do our work solely in terms of dot products, without any explicit usage of a mapped pattern $\Phi_d(x)$. Using kernels of the form (61), we take into account higher-order statistics without the combinatorial explosion (cf. (57)) of time and memory complexity which goes along already with moderately high N and d.

To conclude this section, note that it is possible to modify (61) such that it maps into the space of all monomials up to degree d, defining [29]

$$k(x, x') = ((x \cdot x') + 1)^d.$$
(66)

7 Representing Similarities in Linear Spaces

In what follows, we will look at things the other way round, and start with the kernel. Given some kernel function, can we construct a feature space such that the kernel computes the dot product in that feature space? This question has been brought to the attention of the machine learning community by [1], [8], and [29]. In functional analysis, the same problem has been studied under the heading of *Hilbert space representations* of kernels. A good monograph on the functional analytic theory of kernels is [5]; indeed, a large part of the material in the present section is based on that work.

There is one more aspect in which this section differs from the previous one: the latter dealt with vectorial data. The results in the current section, in contrast, hold for data drawn from domains which need no additional structure other than them being nonempty sets \mathcal{X} . This generalizes kernel learning algorithms to a large number of situations where a vectorial representation is not readily available [17, 12, 33].

We start with some basic definitions and results.

Definition 1 (Gram matrix). Given a kernel k and patterns $x_1, \ldots, x_m \in \mathfrak{X}$, the $m \times m$ matrix

$$K := (k(x_i, x_j))_{ij} \tag{67}$$

is called the Gram matrix (or kernel matrix) of k with respect to x_1, \ldots, x_m .

Definition 2 (Positive definite matrix). An $m \times m$ matrix K_{ij} satisfying

$$\sum_{i,j} c_i \bar{c}_j K_{ij} \ge 0 \tag{68}$$

for all $c_i \in \mathbb{C}$ is called positive definite.

Definition 3 ((Positive definite) kernel). Let \mathfrak{X} be a nonempty set. A function $k : \mathfrak{X} \times \mathfrak{X} \to \mathbb{C}$ which for all $m \in \mathbb{N}, x_i \in \mathfrak{X}$ gives rise to a positive definite Gram matrix is called a positive definite kernel. Often, we shall refer to it simply as a kernel.

The term *kernel* stems from the first use of this type of function in the study of integral operators. A function k which gives rise to an operator T via

$$(Tf)(x) = \int_{\mathcal{X}} k(x, x') f(x') \, dx' \tag{69}$$

is called the kernel of T. One might argue that the term *positive definite kernel* is slightly misleading. In matrix theory, the term *definite* is usually used to denote the case where equality in (68) only occurs if $c_1 = \cdots = c_m = 0$. Simply using the term *positive kernel*, on the other hand, could be confused with a kernel whose values are positive. In the literature, a number of different terms are used for positive definite kernels, such as *reproducing kernel*, *Mercer kernel*, or *support vector kernel*.

The definitions for (positive definite) kernels and positive definite matrices differ only in the fact that in the former case, we are free to choose the points on which the kernel is evaluated.

Positive definiteness implies positivity on the diagonal,

$$k(x_1, x_1) \ge 0 \text{ for all } x_1 \in \mathfrak{X},\tag{70}$$

(use m = 1 in (68)), and symmetry, i.e.,

$$k(x_i, x_j) = \overline{k(x_j, x_i)}.$$
(71)

Note that in the complex-valued case, our definition of symmetry includes complex conjugation, depicted by the bar. The definition of symmetry of matrices is analogous, i.e., $K_{ij} = \bar{K}_{ji}$.

Obviously, real-valued kernels, which are what we will mainly be concerned with, are contained in the above definition as a special case, since we did not require that the kernel take values in $\mathbb{C}\setminus\mathbb{R}$. However, it is not sufficient to require that (68) hold for real coefficients c_i . If we want to get away with real coefficients only, we additionally have to require that the kernel be symmetric,

$$k(x_i, x_j) = k(x_j, x_i).$$
 (72)

Kernels can be regarded as generalized dot products. Indeed, any dot product can be shown to be a kernel; however, linearity does not carry over from dot products to general kernels. Another property of dot products, the Cauchy-Schwarz inequality, does have a natural generalization to kernels:

Proposition 2. If k is a positive definite kernel, and $x_1, x_2 \in \mathcal{X}$, then

$$|k(x_1, x_2)|^2 \le k(x_1, x_1) \cdot k(x_2, x_2).$$
(73)

Proof. For sake of brevity, we give a non-elementary proof using some basic facts of linear algebra. The 2×2 Gram matrix with entries $K_{ij} = k(x_i, x_j)$ is positive definite. Hence both its eigenvalues are nonnegative, and so is their product, K's determinant, i.e.,

$$0 \le K_{11}K_{22} - K_{12}K_{21} = K_{11}K_{22} - K_{12}\bar{K}_{12} = K_{11}K_{22} - |K_{12}|^2.$$
(74)

Substituting $k(x_i, x_j)$ for K_{ij} , we get the desired inequality.

We are now in a position to construct the feature space associated with a kernel k.

We define a map from \mathfrak{X} into the space of functions mapping \mathfrak{X} into \mathbb{C} , denoted as $\mathbb{C}^{\mathfrak{X}}$, via

$$\Phi: \mathfrak{X} \to \mathbb{C}^{\mathfrak{X}}$$
$$x \mapsto k(., x). \tag{75}$$

Here, $\Phi(x) = k(., x)$ denotes the function that assigns the value k(x', x) to $x' \in \mathfrak{X}$.

We have thus turned each pattern into a function on the domain \mathfrak{X} . In a sense, a pattern is now represented by its similarity to *all* other points in the input domain \mathfrak{X} . This seems a very rich representation, but it will turn out that the kernel allows the computation of the dot product in that representation.

We shall now construct a dot product space containing the images of the input patterns under Φ . To this end, we first need to endow it with the linear structure of a vector space. This is done by forming linear combinations of the form

$$f(.) = \sum_{i=1}^{m} \alpha_i k(., x_i).$$
(76)

Here, $m \in \mathbb{N}$, $\alpha_i \in \mathbb{C}$ and $x_i \in \mathcal{X}$ are arbitrary.

Next, we define a dot product between f and another function

$$g(.) = \sum_{j=1}^{m'} \beta_j k(., x'_j)$$
(77)

 $(m' \in \mathbb{N}, \beta_j \in \mathbb{C} \text{ and } x'_j \in \mathfrak{X})$ as

$$\langle f,g\rangle := \sum_{i=1}^{m} \sum_{j=1}^{m'} \bar{\alpha}_i \beta_j k(x_i, x'_j).$$
(78)

To see that this is well-defined, although it explicitly contains the expansion coefficients (which need not be unique), note that

$$\langle f,g\rangle = \sum_{j=1}^{m'} \beta_j \overline{f(x'_j)},\tag{79}$$

using $k(x'_j, x_i) = \overline{k(x_i, x'_j)}$. The latter, however, does not depend on the particular expansion of f. Similarly, for g, note that

$$\langle f,g\rangle = \sum_{i=1}^{m} \bar{\alpha}_i g(x_i).$$
(80)

The last two equations also show that $\langle \cdot, \cdot \rangle$ is bilinear. It is symmetric, as $\langle f, g \rangle = \overline{\langle g, f \rangle}$. Moreover, it is positive definite, since positive definiteness of k implies that for any function f, written as (76), we have

$$\langle f, f \rangle = \sum_{i,j=1}^{m} \alpha_i \alpha_j k(x_i, x_j) \ge 0.$$
(81)

The latter implies that $\langle \cdot, \cdot \rangle$ is actually itself a positive definite kernel, defined on our space of functions. To see this, note that given functions f_1, \ldots, f_n , and coefficients $\gamma_1, \ldots, \gamma_n \in \mathbb{R}$, we have

$$\sum_{i,j=1}^{n} \gamma_i \gamma_j \langle f_i, f_j \rangle = \left\langle \sum_{i=1}^{n} \gamma_i f_i, \sum_{j=1}^{n} \gamma_j f_j \right\rangle \ge 0.$$
(82)

Here, the left hand equality follows from the bilinearity of $\langle \cdot, \cdot \rangle$, and the right hand inequality from (81).

For the last step in proving that it even is a dot product, we will use the following interesting property of Φ , which follows directly from the definition: for all functions (76), we have

$$\langle k(.,x), f \rangle = f(x) \tag{83}$$

-k is the representer of evaluation. In particular,

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$$\langle k(.,x), k(.,x') \rangle = k(x,x').$$
 (84)

By virtue of these properties, positive definite kernels k are also called *reproduc*ing kernels [3, 5, 32, 17].

By (83) and Proposition 2, we have

$$|f(x)|^{2} = |\langle k(.,x), f \rangle|^{2} \le k(x,x) \cdot \langle f, f \rangle.$$

$$(85)$$

Therefore, $\langle f, f \rangle = 0$ directly implies f = 0, which is the last property that was left to prove in order to establish that $\langle ., . \rangle$ is a dot product.

One can complete the space of functions (76) in the norm corresponding to the dot product, i.e., add the limit points of sequences that are convergent in that norm, and thus gets a Hilbert space H, usually called a *reproducing kernel* Hilbert space.³

The case of real-valued kernels is included in the above; in that case, H can be chosen as a real Hilbert space.

8 Examples of Kernels

Besides (61), [8] and [29] suggest the usage of Gaussian radial basis function kernels [1]

$$k(x, x') = \exp\left(-\frac{\|x - x'\|^2}{2\,\sigma^2}\right)$$
(86)

and sigmoid kernels

$$k(x, x') = \tanh(\kappa(x \cdot x') + \Theta).$$
(87)

While one can show that (87) is not a kernel [26], (86) has become one of the most useful kernels in situations where no further knowledge about the problem at hand is given.

Note that all the kernels discussed so far have the convenient property of unitary invariance, i.e., k(x, x') = k(Ux, Ux') if $U^{\top} = U^{-1}$ (if we consider complex numbers, then U^* instead of U^{\top} has to be used).

The radial basis function kernel additionally is translation invariant. Moreover, as it satisfies k(x, x) = 1 for all $x \in \mathcal{X}$, each mapped example has unit length, $\|\Phi(x)\| = 1$. In addition, as k(x, x') > 0 for all $x, x' \in \mathcal{X}$, all points lie inside the same orthant in feature space. To see this, recall that for unit length vectors, the dot product (3) equals the cosine of the enclosed angle. Hence

$$\cos(\angle(\varPhi(x), \varPhi(x'))) = (\varPhi(x) \cdot \varPhi(x')) = k(x, x') > 0,$$
(88)

which amounts to saying that the enclosed angle between any two mapped examples is smaller than $\pi/2$.

The examples given so far apply to the case of vectorial data. Let us at least give one example where \mathfrak{X} is not a vector space.

³ A Hilbert space H is defined as a complete dot product space. Completeness means that all sequences in H which are convergent in the norm corresponding to the dot product will actually have their limits in H, too.

Example 1 (Similarity of probabilistic events). If \mathcal{A} is a σ -algebra, and P a probability measure on \mathcal{A} , then

$$k(A,B) = P(A \cap B) - P(A)P(B)$$
(89)

is a positive definite kernel.

Further examples include kernels for string matching, as proposed by [33] and [12].

There is an analog of the kernel trick for distances rather than dot products, i.e., dissimilarities rather than similarities. This leads to the class of *conditionally positive definite kernels*, which contain the standard SV kernels as special cases. Interestingly, it turns out that SVMs and kernel PCA can be applied also with this larger class of kernels, due to their being translation invariant in feature space [23].

9 Applications

Having described the basics of SV machines, we now summarize some empirical findings.

By the use of kernels, the optimal margin classifier was turned into a classifier which became a serious competitor of high-performance classifiers. Surprisingly, it was noticed that when different kernel functions are used in SV machines, they empirically lead to very similar classification accuracies and SV sets [18]. In this sense, the SV set seems to characterize (or *compress*) the given task in a manner which up to a certain degree is independent of the type of kernel (i.e., the type of classifier) used.

Initial work at AT&T Bell Labs focused on OCR (optical character recognition), a problem where the two main issues are classification accuracy and classification speed. Consequently, some effort went into the improvement of SV machines on these issues, leading to the *Virtual SV* method for incorporating prior knowledge about transformation invariances by transforming SVs, and the *Reduced Set* method for speeding up classification. This way, SV machines became competitive with (or, in some cases, superior to) the best available classifiers on both OCR and object recognition tasks [7, 9, 11].

Another initial weakness of SV machines, less apparent in OCR applications which are characterized by low noise levels, was that the size of the quadratic programming problem scaled with the number of Support Vectors. This was due to the fact that in (35), the quadratic part contained at least all SVs — the common practice was to extract the SVs by going through the training data in chunks while regularly testing for the possibility that some of the patterns that were initially not identified as SVs turn out to become SVs at a later stage (note that without chunking, the size of the matrix would be $m \times m$, where m is the number of all training examples). What happens if we have a high-noise problem? In this case, many of the slack variables ξ_i will become nonzero, and all the corresponding examples will become SVs. For this case, a decomposition algorithm was proposed [14], which is based on the observation that not only can we leave out the non-SV examples (i.e., the x_i with $\alpha_i = 0$) from the current chunk, but also some of the SVs, especially those that hit the upper boundary (i.e., $\alpha_i = C$). In fact, one can use chunks which do not even contain all SVs, and maximize over the corresponding sub-problems. SMO [15] explores an extreme case, where the sub-problems are chosen so small that one can solve them analytically. Several public domain SV packages and optimizers are listed on the web page http://www.kernel-machines.org. For more details on the optimization problem, see [23].

10 Conclusion

One of the most appealing features of kernel algorithms is the solid foundation provided by both statistical learning theory and functional analysis. Kernel methods let us interpret (and design) learning algorithms geometrically in feature spaces nonlinearly related to the input space, and combine statistics and geometry in a promising way. Kernels provide an elegant framework for studying three fundamental issues of machine learning:

- Similarity measures the kernel can be viewed as a (nonlinear) similarity measure, and should ideally incorporate prior knowledge about the problem at hand
- Data representation as described above, kernels induce representations of the data in a linear space
- Function class due to the representer theorem, the kernel implicitly also determines the function class which is used for learning.

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