

Classification of Multivariate Data Using Distribution Mapping Exponent

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Abstract: We introduce distribution-mapping exponent that is something like effective dimensionality of multidimensional space. The method for classification of multivariate data is proposed. It is based on local estimate of distribution mapping exponent q for each point x . Distances of all points of a given class of the training set from a given (unknown) point x are searched and it is shown that the sum of reciprocals of q -th power of these distances can be used as the probability density estimate. The classification quality was tested and compared with other methods using multivariate data from UCI Machine Learning Repository. The method has no tuning parameters.

Keywords multivariate data; classification; distribution-mapping exponent

1 Introduction

Classification of multivariate data is a problem solved by lot of methods from nearest neighbor method to decision trees, neural networks and genetic algorithms. The problem is generally difficult because of several influences, e.g.

- High problem dimensionality where curse of dimensionality causes excessive grow of processing time.
- Presence of noise; true data are rarely “pure”.
- Multicollinearity, i.e. mutual dependence of individual variables. If variables are originally considered independent, i.e. orthogonal to all others, multicollinearity causes distortion of the space; coordinates are not orthogonal already.
- Boundary effect. Due to this effect nearest points seem to be rather far and farther points near so that the distance between the nearest and the farthest

point of finite data set can be smaller than the distance of the nearest neighbor from the given point.

In this paper we deal with distances in multidimensional space and try to simplify a complex picture of probability distribution of points in this space introducing mapping functions of one variable. This variable is the distance from the given point (the query point x [3]) in multidimensional space. From it follows that mapping functions are different for different query points and this is cost we pay for simplification from n variables in n -dimensional space to one variable. We will show that this cost is not too high – at least in application presented here.

The distance is basic notion for all approaches dealing with neighbors, especially nearest neighbors. There is a lot of methods of classification based on the nearest neighbors [1]. They estimate the probability density at point x (a query point [3]) of the data space by ratio i/V_i of number i of points of a given class in a suitable ball of volume V_i with center at point x [5]. These methods need to optimize the best size of the neighborhood, i.e. the number i of points in the neighborhood of the point x or size of volume V_i . The probability density in the feature (data) space is given by training data. The optimal neighborhood size depends on the training data set, i.e. on the character of data as well as on the number of samples of a given class in the training set. Often it is recommended to choose neighborhood size equal to the square root of number of samples of the training set [5].

The method proposed is based on distances of the training set samples x_s , $s = 1, 2, \dots, k$ from point x . It is shown that the sum of reciprocals of q -th power of these distances, where q is a suitable number, is convergent and can be used as a probability density estimate. From the fact of high power of distances in multidimensional Euclidean space, fast convergence, i.e. small influence of distant samples, follows. The speed of convergence is the better the higher dimensionality and the larger q .

The method reminds Parzen window approach [4], [5] but the problem with direct application of this approach is that the step size does not satisfy a necessary convergence condition.

Using distances, i.e. a simple transformation from n -dimensional Euclidean space E_n to one-dimensional Euclidean space E_1 , and no iterations, the curse of dimensionality is straightforwardly eliminated. The method can be also considered as a variant of the kernel method, based on a probability density estimator but using a much simpler metric and does not satisfy some mathematical conditions.

Throughout this paper let us assume that we deal with normalized data, i.e. the individual coordinates of the samples of the learning set are normalized to zero mean and unit variance and the same normalization constants (empirical mean and empirical variance) are applied to all other (testing and of unknown class) data. This transformation does not mean any change in form of the distribution, i.e.

uniform distribution remains uniform, exponential distribution remains exponential (with $\lambda = 1$ and shifted by 1 to the left), etc.

2 Probability Distribution mapping function

Let a query point x be placed without loss of generality in the origin. Let us build balls with their centers in point x and with volumes V_i , $i=1, 2, \dots$

Individual balls are one in another, the $(i-1)$ -st inside the i -th like peels of onion. Then the mean density of points in the i -th ball is $\rho_i = m_i/V_i$. The volume of a ball of radius r in n -dimensional space is $V(r) = \text{const} \cdot r^n$. Thus we have constructed a mapping between the mean density ρ_i in the i -th ball ρ_i and its radius r_i . Then $\rho_i = f(r_i)$. Using tight analogy between density $\rho(z)$ and probability density $p(z)$ one can write $p(r_i) = f(r_i)$ and $p(r_i)$ is the mean probability density in the i -th ball with radius r_i here. This way a complex picture of probability distribution of points in the neighborhood of a query point x is simplified to a function of a scalar variable. We call this function a probability distribution mapping function $D(x, r)$, where x is a query point, and r the distance from it. More exact definitions follow.

Definition

Probability distribution mapping function $D(x, r)$ of the neighborhood of the query point x is function $D(x, r) = \int_{B(x,r)} p(z) dz$, where r is distance from the query point

and $B(x, r)$ is ball with center x and radius r .

Definition

Distribution density mapping function $d(x, r)$ of the neighborhood of the query point x is function $d(x, r) = \frac{\partial}{\partial r} D(x, r)$, where $D(x, r)$ is a probability distribution mapping function of the query point x and radius r .

Note. It is seen that for fixed x the function $D(x, r)$, $r > 0$ is monotonically growing from zero to one. Functions $D(x, r)$ and $d(x, r)$ for x fixed are one-dimensional analogs to the probability distribution function and the probability density functions, respectively. For illustration see Fig. 1.

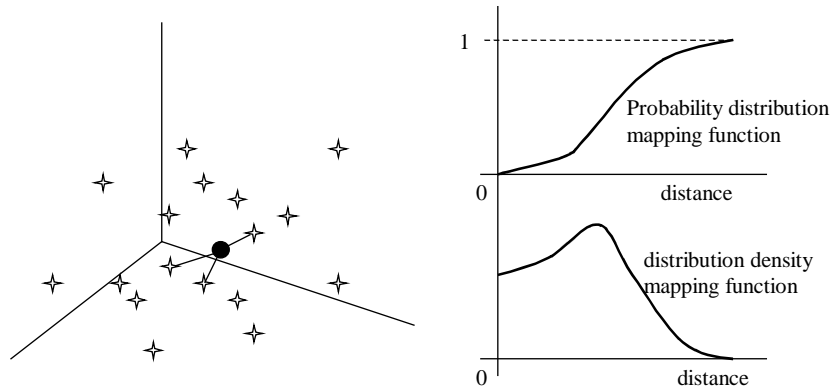


Fig. 1. Data in a multidimensional space and corresponding probability distribution mapping function and distribution density mapping function.

Power approximation of the probability distribution mapping function

Let us approximate the probability distribution mapping function by parabolic function in form $D(x, r^n) = \text{const.}(r^n)^\alpha$. This function is tangent to the vertical axis in point $(0, 0)$ and let it is going through some characteristic points of the distribution.

Definition

Power approximation of the probability distribution mapping function $D(x, r^n)$ is function r^q such that $\frac{D(x, r^n)}{r^q} \rightarrow \text{const}$ for $r \rightarrow 0+$. The exponent q is a distribution-mapping exponent. The variable $\alpha = q/n$ we call distribution mapping ratio.

Note. We often omit a multiplicative constant of the probability distribution mapping function.

Using approximation of the probability distribution mapping function by $D(x, r^n) = \text{const.}(r^n)^\alpha$ the distribution mapping exponent is $q = n\alpha$.

Note that distribution-mapping exponent is influenced by two factors

- True distribution of points of the learning set in E_n .
- Boundary effects, which have the larger influence the larger dimension n and the smaller the learning set size [1], [5].

To overcome the problem of estimation of q using real data, this exponent is estimated by linear regression for each query point as shown in the next Chapter.

3 Distribution mapping exponent estimation

Let the learning set U of total m_T samples be given in the form of a matrix X_T with m_T rows and n columns. Each sample corresponds to one row of X_T and, at the same time, corresponds to a point in n -dimensional Euclidean space E_n , where n is the sample space dimension. The learning set consists of points (rows) of two classes $c \in \{0, 1\}$, i.e. each row (point or sample) corresponds to one class. Then, the learning set $U = U_0 \cup U_1$, $U_0 \cap U_1 = \emptyset$, $U_c = \{x_{cs}\}$, $s = 1, 2, \dots, N_c$, $c = \{0, 1\}$. N_c is the number of samples of class c , $N_0 + N_1 = m_T$, and $x_{cs} = \{x_{cs1}, x_{cs2}, \dots, x_{csn}\}$ is the data sample of class c .

We use normalized data, i.e. each variable x_{csj} (j fixed, $s = 1, 2, \dots, m_T$, $c = 0$ or 1) corresponds to the j -th column of matrix X_T has zero mean and unit variance.

Let point $x \notin U$ be given and let points x_{cs} of one class be sorted so that index $i = 1$ corresponds to the nearest neighbor, index $i = 2$ to the second nearest neighbor, etc. In the Euclidean metrics, $r_i = \|x, x_{ci}\|$ is the distance of the i -th nearest neighbor of class c from point x .

From definition of the distribution mapping exponent it follows that r_i^q should be proportional to index i , i.e.

$$r_i^q = ki, \quad i = 1, 2, \dots, N_c, \quad c = 0 \text{ or } 1, \quad (1)$$

and where k is a suitable constant. Using logarithm we get

$$q \ln(r_i) = k' + \ln(i), \quad i = 1, 2, \dots, N_c. \quad (2)$$

System of these N_c equations with respect to unknown q can be solved using standard linear regression for both classes. Thus we get two values of q , q_0 and q_1 . To get a single value of q we use the weighted arithmetic mean, $q = (q_0 N_0 + q_1 N_1) / (N_0 + N_1)$.

At this point we can say that q is something like effective dimensionality of the data space including true distribution of points of both classes and boundary effect. In the next chapter we use it directly instead of dimension.

4 All learning samples approach

Let us define

$$p_c(x) = \frac{C}{k-1} \sum_{i=2}^k 1/r_i^q \quad (3)$$

where C is a constant. We show below that $p_c(x)$ is probability density estimate. Note that (3) reminds Parzen window approach [4], [5 Chap. 4.3] with weighting function $K(y) = y^{-q}$ for $y > r_1^q$ and $K(y) = 0$ otherwise, $q \in <1, n>$ and with window width $h = 1$. The problem with direct application of this approach here is that h does not satisfy the necessary condition $\lim_{i \rightarrow \infty} h(i) = 0$ [4, eq.(1.8)].

On the one hand due to (1) the series in (3) should be a harmonic series that is divergent. On the other hand we will prove below that it is not true. The series $1/r_i^q$ converges with size of r_i for $q > 1$ and thus we have no reason to limit ourselves to the nearest k points and we can use all points in the learning set using $k = N_c$, $c = 0$ or 1 . At the same time the ordering of individual components is not essential and we need not sort the samples of X_T with respect to their r_i when using the nearest neighbor approach. (But we need to sort them when estimating distribution mapping exponent q .)

In practical procedure for each query point x we first compute the distribution mapping exponent q using (2) by standard linear regression. After it, we simply sum up all components $1/r_i^q$ and, at the same time, we store the largest component which corresponds to the nearest neighbor of point x which has the smallest r_i^q . In the end we subtract it thus excluding the nearest point. This is made for both classes simultaneously getting numbers S_0 and S_1 for both classes. Their ratio gives a value of the discriminant function, here the Bayes ratio. We can get also probability estimation that the point $x \in E_n$ is of class 1:

$$R(x) = \frac{S_1}{S_0} \quad \text{or} \quad p_1(x) = \frac{S_1}{S_1 + S_0} .$$

Then for a threshold (cut) θ chosen, if $R(x) > \theta$ or $p_1(x) > \theta$ then x belongs to class 1 else to class 0.

The method is very close to the nearest neighbor as well as kernel methods. From the point of view of kernel methods, the kernel is or would be $K(x) = \|x - x_i\|^{-q}$ with Euclidean norm $\|\cdot\|$ in E_n . There is no smoothing (bandwidth) parameter. The problem is that this kernel is difficult to consider as a probability distribution function according to the definition of a kernel [1]. Taking $\|x - x_i\| = r$ we have $K(r) = r^{-q}$ and integrals $\int_{-\infty}^{\infty} K(r) dr$ or $\int_0^{\infty} K(r) dr$ are not convergent; they should be equal to 1 or at least finite.

5 Probability Density Estimation

Let us look at the problem what is the relation of part D_i of space E_n which falls on i nearest neighbors of the given point x . We will assume the following:

Assumption 1: Let there be points in the Euclidean space E_n distributed uniformly in the sense that the distribution of each of the n coordinates is uniform. Let i be the order number of the i -th nearest neighbor to the point x . Let r_i be the distance of the i -th nearest neighbor of the given point $x \in E_n$ from point x_i . Let D be a constant, $q \in (1, n)$ be a constant, and \bar{D}_i be the mean value of the variable r_i^q , and let it hold

$$\bar{D}_i = iD .$$

Comment: Under Assumption 1 by “the part D_i of the space E_n ” we do not mean a volume of a ball with the center in the point x and radius r_i but, in fact (except for a multiplicative constant), a ball of the same center and radius but in the space of dimension given by constant q , i.e. in the E_q . The basis for introducing Assumption 1 is finding as follows. By simulation one can find that the relation $\bar{V}_i = iV$ where V is a constant does not hold but for some $q \leq n$ it holds $\bar{D}_i = \bar{r}_i^q = iD$ where i is the number of the i -th nearest neighbor of point $x \in E_n$ and D is a constant. From it follows that the q -th power grows linearly.

Theorem 1

Let Assumption 1 be valid, and let $\bar{\Delta}_i$ be mean of $\Delta_i = r_i^q - r_{i-1}^q$, \bar{D}_i be mean of $D_i = r_i^q$, \bar{V}_i be mean of $V_i = cr_i^n$ where c is a constant. Moreover let exist a constant K such that $p(\bar{\Delta}_i) = K/\bar{\Delta}_i$. Then for the probability density $p(i) = K'i/\bar{V}_i$ of points in the neighborhood of point x it holds $p(\bar{\Delta}_i) = p(\bar{D}_i) = p(i)$, where $p(\bar{D}_i) = \frac{iK}{\bar{D}_i}$.

Proof: The $p(i)$ is probability density and at the same time due to Assumption 1 $1/\bar{D}_i$ is proportional to $p(i)$. Then there is a constant K that $p(\bar{D}_i) = p(i)$. Under Assumption 1 there is $\bar{\Delta}_i = D$ and then $p(\bar{\Delta}_i) = p(\bar{D}_i) = p(i)$. \square

6 The Proof of Convergence

Theorem 1 states that probability density is proportional to $1/r_i^q$ and formula (3) uses the sum of these ratios supposing to get a reasonable number for probability

density estimation. So it is supposed that for a number of samples going to infinity, the sum would be convergent.

Theorem 2

Let exist a mapping of probability density of points of class c in E_n , $E_n \rightarrow E_1$: $p(x_{ci}) = p(r_{ci}^q)$ so that

$$\begin{aligned} K/r_{c1}^q &= p(x_{c1}), \quad K/(r_{c2}^q - r_{c1}^q) = p(x_{c2}), \\ \dots K/(r_{cNc}^q - r_{c(Nc-1)}^q) &= p(x_{cNc}), \end{aligned} \quad (4)$$

where K is a fixed constant that has the same value for both classes. Let exist a constant $\varepsilon > 0$ and index $k > 2$ so that for each $i > k$ it holds

$$p(x_{ci}) \leq \frac{p(x_{c2})}{(1 + (i - k)\varepsilon)^{i-k}}. \quad (5)$$

Then

$$S_c = \sum_{i=2}^{N_c} \frac{1}{r_{ci}^q} = p(x_{c2})K(1 + C_c), \quad (6)$$

where K and C_c are finite constants.

Proof: First we arrange (6) in form

$$S_c = \sum_{i=2}^{N_c} \frac{1}{r_{ci}^q} = \frac{1}{r_{c2}^q} + \sum_{i=3}^{N_c} \frac{1}{r_{c2}^q + \Delta_{c3} + \Delta_{c4} + K + \Delta_{ci}}.$$

Then using mapping (4) introduced we get

$$\begin{aligned} S_c &= Kp_{c2} + K \sum_{i=3}^{N_c} \frac{1}{\frac{1}{p_{c2}} + \frac{1}{p_{c3}} + K + \frac{1}{p_{ci}}} = \\ &= p_{c2}K \left(1 + \sum_{i=3}^{N_c} \frac{1}{1 + \frac{p_{c2}}{p_{c3}} + K + \frac{p_{c2}}{p_{ci}}} \right) \equiv p_{c2}K \left(1 + \sum_{i=3}^{N_c} P_i \right) \end{aligned} \quad (7)$$

For individual elements p_{c2} / p_{ci} in denominators of fractions in the sum it holds

$$\frac{p_{c2}}{p_{ci}} = \frac{p_{c2}(1 + (i - k)\varepsilon)^{i-k}}{p_{c2}} = (1 + (i - k)\varepsilon)^{i-k}.$$

Using condition (5) the summed elements P_k, P_{k+1}, \dots in (7) since the k -th have form

$$\begin{aligned} P_k &= \frac{1}{C}, \quad P_{k+1} = \frac{1}{C + 1 + \varepsilon}, \quad P_{k+2} = \frac{1}{C + 1 + \varepsilon + (1 + \varepsilon)^2}, \\ P_{k+i} &= 1/[C + (1 + \varepsilon) + (1 + 2\varepsilon)^2 + \dots + (1 + i\varepsilon)^i]. \end{aligned}$$

Then according to d'Alembert's criterion

$$\frac{P_{k+i+1}}{P_{k+i}} = \frac{C + (1+\varepsilon) + (1+2\varepsilon)^2 + \dots + (1+i\varepsilon)^i}{C + (1+\varepsilon) + (1+2\varepsilon)^2 + \dots + (1+i\varepsilon)^i + (1+(i+1)\varepsilon)^{i+1}} < 1$$

$\forall i > 0$ and $\forall \varepsilon > 0$. Then the series is convergent.

Notes:

a) In the statement of the theorem the sum need not start just by index $i = 2$. We can start with the nearest neighbor ($i = 1$) or other neighbors ($i > 2$). The value $i = 2$ is given by a compromise between the error caused by the small value and the large variability of $\Delta_{c1} = r_{c1}$, and the inaccuracy caused by the larger distance from point x for $i > 2$, see Chap. 3.

b) The last condition (5) defines the speed of diminishing the tail of the distribution; probably condition that the distribution should have the mean would suffice.

7 Discussion

From formula (7) it is seen that for a "smooth" form of the distribution function around point x and for the large density of points for both classes, the ratios p_{c2}/p_{c1} are very close to 1 for rather large values of i (e.g. 100, but let us take 11 here). For both classes the elements of sum in (7) are $\frac{1}{2}, \frac{1}{3}, \dots, \frac{1}{11}$ and their sum is 2.01987 here, and the other elements have form $\frac{1}{11+(i-1)(1+\delta)}$, where

starting from the index k it is $\delta \geq \varepsilon$. (Index k can be different for both classes.) It is then probable that the values of sums in (7) will be very close for both classes and the ratio of (7) for one and the other class will be close to Bayes ratio $p_1(x_{c2})/p_0(x_{c2}) = S_1/S_0$. In such a case we can also estimate the probability that the sample x belongs among signals:

$$p_1(x) \approx p_1(x_{c2}) \approx \frac{S_1}{S_1 + S_0}.$$

Using neighbor distances for the probability density estimation, the probability density estimation should copy the features of the probability density function based on real data. The idea of most nearest-neighbors-based methods as well as kernel methods [1] does not reflect the boundary effects. That means that for any point x , the statistical distribution of the data points x_i surrounding it is supposed to be independent of the location of the neighbor points and their distances x_i from point x . This assumption is often not met, especially for small data sets and for

higher dimensions. To illustrate this, let us consider uniformly distributed points in a cube $(-0.5, +0.5)^n$. Let there be a ball with its center in the origin and the radius equal to 0.5. This ball occupies $\frac{4}{3}\pi \cdot 0.5^3 = 0.524$, i.e. more than 52 % of that cube in a three-dimensional space, 0.080746, i.e. 8 % of unit cube in 6-dimensional space, 0.0026 in 10-dimensional space, and $3.28e-21$ in 40-dimensional space. It is then seen that starting by some dimension n , say 5 or 6 and some index i , the i -th nearest neighbor does not lie in such a ball around point x but somewhere “in the corner” in that cube but outside this ball (the boundary effect [6]). From it follows that this i -th neighbor lies farther from point x as would follow from the uniformity of distribution. In farther places from the origin the space thus seems to be less dense than near the origin. The function $f(i) = \bar{r}_i^n$, where \bar{r}_i is the mean distance of the i -th neighbor from point x , should grow linearly with index i in the case of uniform distribution without the boundary effect mentioned. In the other case this function grows faster than linearly.

The samples of the learning set are normalized to zero mean and unit variance for each variable as introduced in the beginning. Assume that all thus arising marginal distributions are approximately normal. Our point x has an unknown class and also unknown probabilities $p_1(x)$ and $p_0(x)$ and lies just in the point $(0, 0, \dots, 0)$. For point x we can introduce different neighborhoods, for our estimations let us use three only:

- A. Till the distance of one sigma in all dimensions,
- B. From the distance of one sigma to the distance of two sigmas in all dimensions,
- C. From the distance of two sigmas to infinity in each dimension.

There is assumption of normality of all variables. Then in each dimension, approximately 68 % points of the learning set lie inside A, 95 % points lie inside A and B, i.e. 27 % in B, and 5 % in C. To these three “layers” also some mean distances (0.5, 1.5, and 3) correspond in all dimensions.

The total portion (percentage) of points in layer A are given by 0.68^n , in A and B together by 0.95^n , and in C by what remains to 1.

For computation of sum in (3) all points from layers A, B, and C are used. Each point in these three layers benefit to the total sum (3) by its part. Benefits to the total sum by points in the individual layer are given by the relative number of points in a layer (total points in layer) divided by average distance (which is the average distance in one dimension times the square root of the dimension) to the $(n-1)$ -st power recomputed to 100 %. E.g. for layer A and $n=2$ there is $B_A = 0.4224 \cdot 1/(0.5\sqrt{2})^2 = 0.653932$, $B_B = 0.4401 \cdot 1/(1.5\sqrt{2})^2 = 0.207465$,

and $B_C = 0.0975 \cdot 1/(3\sqrt{2})^2 = 0.022981$. These numbers divided by sum $B_A + B_B + B_C$ give 73.94 %, 23.46 %, 2.60 %, respectively. For $n = 5$ the distribution of points in layers A, B, C are 14.54%, 62.84%, and 22.62%, respectively, and corresponding benefits are in the same ordering 94.83%, 5.06%, and 0.11%. Similarly for $n = 20$ we get numbers 0.044687%, 35.80%, 64.15% for distribution of points, and 99.999931%, 0.000069%, 2.35588E-12 for corresponding benefits of points in layers A, B, and C.

These estimations show that due to the geometry of the multidimensional Euclidean space the share of points corresponding to A with respect to the total number of points lessens essentially with dimension. At the same time, their benefit to the total sum is close to 100 %. This is because parts A, B, C are, in fact, not cubes but n -dimensional balls of radii computed from an average distance in one dimension. It also follows that the share of the layer C to the total sum is negligible for the dimension 5 or 6 and more. With the dimension growing also the convergence of the sum is much faster as the points of the learning set near point x gave practically the whole value of the sum. The larger dimension the lesser percentage of points from the learning set influence the result. On the other hand, for low dimensionality, especially 2 and 3 even the farthest points have strong influence.

"German"			"Heart"		
Algorithm	Error	Note	Algorithm	Error	Note
SFSloc7	0.520	1; 2	SFSloc7	0.357	3
Discrim	0.535		Bayes	0.374	
LogDisc	0.538		Discrim	0.393	
Castle	0.583		LogDisc	0.396	
Alloc80	0.584		Alloc80	0.407	
Dipol92	0.599		QuaDisc	0.422	
Smart	0.601		Castle	0.441	
Cal	0.603		Cal5	0.444	
Cart	0.613		Cart	0.452	
QuaDisc	0.619		Cascade	0.467	
KNN	0.694		KNN	0.478	
Default	0.700		Smart	0.478	
Bayes	0.703		Dipol92	0.507	
IndCart	0.761		Itrule	0.515	
BackProp	0.772		BayTree	0.526	
BayTree	0.778		Default	0.560	
Cn2	0.856		BackProp	0.574	
Ac2	0.878		LVQ	0.600	
Itrule	0.879		IndCart	0.630	
NewId	0.925		Kohonen	0.693	
LVQ	0.963		Ac2	0.744	
Radial	0.971		Cn2	0.767	
C4.5	0.985		Radial	0.781	
Kohonen	1.160		C4.5	0.781	
Cascade	100.0		NewId	0.844	
"Adult"			"Ionosphere"		
Algorithm	Error	Note	Algorithm	Error	Note
FSS Naive Bayes	0.1405		IB3	0.0330	6; 7
NBTree	0.1410		backprop	0.0400	8
C4.5-auto	0.1446		SFSloc7	0.0596	9
IDTM Dec. table	0.1446		Ross Quinlan's C4	0.0600	10
HOODG	0.1482		nearest neighbor	0.0790	
C4.5 rules	0.1494		"non-linear" perceptr.	0.0800	
OC1	0.1504		"linear" perceptron	0.0930	
C4.5	0.1554				
Voted ID3 (0.6)	0.1564				
CN2	0.1600				
Naive-Bayes	0.1612				
Voted ID3 (0.8)	0.1647				
T2	0.1684				
SFSloc7	0.1786	4			
1R	0.1954				
Nearest-neighbor 1	0.2035				
Nearest-neighbor 2	0.2142				
Pebles	Crashed	5			

Table 1. Comparison of classification error of SFSloc7 for different tasks with results for another classifiers as given by [7]. Notes – see the next page.

Notes to Table 1:

- 1 for threshold 0.413
- 2 numeric data
- 3 for threshold 0.24
- 4 for threshold 0.868482
- 5 Unknown why (bounds WERE increased)
- 6 parameter settings: 70% and 80% for acceptance and dropping respectively
- 7 (Aha & Kibler, IJCAI-1989)
- 8 an average of over ..
- 9 for threshold 0.550254
- 10 no windowing

8 Results - testing the classification ability

The classification algorithm was written in c++ as SFSloc7 program and tested using tasks from UCI Machine Learning Repository [7]. Tasks of classification into two classes for which data about previous tests are known were selected: "Adult", "German", "Heart", and "Ionosphere".

The task "Adult" is to determine whether a person makes over 50000 \$ a year.

The task "German" is about whether the client is good or bad to lend him money.

The task "Heart" indicates absence or presence of heart disease for patient.

For the task "Ionosphere" the targets were free electrons in the ionosphere. "Good" radar returns are those showing evidence of some type of structure in the ionosphere. "Bad" returns are those that do not; their signals pass through the ionosphere.

We do not describe these tasks in detail here as all can be found in [7]. For each task the same approach to testing and evaluation was used as described in [7]. In Table 1 results are shown together with results for other methods as given in [7]. For each task methods are sorted according to classification error, the method with the best – lowest error first. It is seen that for some tasks SFSloc7 is good but there are tasks where the method is worse than average – it would be strange to outperform all methods for all tasks. The method is totally parameterless. There is no parameter for tuning to get the best result. The method simply works satisfactorily or not, there is nothing to try more.

Conclusions

In this paper we dealt with simplified representation of probability distribution of points in multidimensional Euclidean space including boundary effects. A new

method for classification was developed. The method is based on notion of distribution mapping exponent and its local estimate q for each query point x . The theorem on convergence was formulated and proved and a convergence estimation was shown. It was found that the higher dimensionality, the better.

The method has no tuning parameters: No neighborhood size, no convergence coefficients etc. need to be set up in advance to assure convergence. There is no true learning phase. In the „learning phase“ only normalization constants are computed and thus this phase is several orders of magnitude faster than the learning phase of neural networks or many other methods [2], [7]. In the recall phase for each sample to be classified the learning set is searched twice, once for finding the local value of the distribution mapping exponent q , and second for all samples of the learning set elements of sum (3) are computed. The amount of computation is thus proportional to the learning set size, i.e. the dimensionality times the number of the learning samples.

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References

- [1] Silverman, B. W.: Density Estimation for Statistics and data Analysis. Chapman and Hall, London, 1986.
- [2] Bock, R. K. et al.: Methods for multidimensional event classification: a case study. To be published as Internal Note in CERN, 2003.
- [3] Hinnenburg, A., Aggarwal, C.C., Keim, D.A.: What is the nearest neighbor in high dimensional spaces? Proc. of the 26th VLDB Conf., Cairo, Egypt, 2000, pp 506-515.
- [4] Parzen, E.: On Estimation of Probability Density Function and Mode. The Annals of Mathematical Statistics, Vol. 33, No. 3 (Sept. 1962), pp. 1065-1076.
- [5] Duda, R., Hart, P., Stork, D.G.: Pattern Classification. John Wiley and Sons, 2000.
- [6] Arya, S., Mount, D.M., Narayan, O., Accounting for Boundary Effects in Nearest Neighbor Searching. Discrete and Computational Geometry Vol.16 (1996), pp. 155-176.
- [7] UCI Machine Learning Repository.
<http://www.ics.uci.edu/~mlern/MLSummary.html>