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NEIGHBORHOOD COUNTING MEASURE METRIC AND MINIMUM RISK METRIC: AN EMPIRICAL COMPARISON

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Neighborhood Counting Measure Metric and Minimum Risk Metric: An empirical comparison

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Abstract—Wang in a PAMI paper proposed Neighborhood Counting Measure (NCM) as a similarity measure for the k-nearest neighbors classification algorithm. In his paper, Wang mentioned Minimum Risk Metric (MRM) an earlier method based on the minimization of the risk of misclassification. However, Wang did not compare NCM with MRM because of its allegedly excessive computational load. In this letter, we empirically compare NCM against MRM on k-NN with k=1, 3, 5, 7 and 11 with decision taken with a voting scheme and k=21 with decision taken with a weighted voting scheme on the same datasets used by Wang. Our results shows that MRM outperforms NCM for most of the k values tested. Moreover, we show that the MRM computation is not so probihibitive as indicated by Wang.

Index Terms—Pattern Recognition, Machine Learning, k-Nearest Neighbors, distance measures, MRM, NCM.

I. Introduction

THE k-nearest neighbors (k-NN) is a well-known **L** algorithm used in machine learning and in pattern recognition for classification tasks [1]. Given a point to classify and a distance (or similarity) function defined on the input space, k-NN finds the k-neighbors of the point and classify it to the majority class of its neighbors. The performance of k-NN depends on the distance/similarity function used to compute the set of neighbors and on the choice of k. The classification decision can weight the vote of each neighbor depending on the values of the used distance or similarity. In literature, there is a large variety of distance functions that are applicable to different data types. Among the most simple distances we can recall the Euclidean distance for numeric attributes and the Hamming distance for categorical attributes. More complex distances, such as HEOM [2] and HVDM [2], were introduced for data with mixed features. Other relevant distances are Minimum Risk Metric (MRM) [3] that minimize the risk of misclassification using conditional probabilities and Neighboring Counting Measure (NCM) proposed by Wang [4] that works counting the neigborhoods in the input space. Paredes and Vidal [5] presented an algorithm to learn weighted metrics for numerical data. The weights are learned by minimizing an approximation of the leave-one-out classification error on a subset of the training set with gradient descent algorithm. MRM is a distance for classification tasks that relies on the estimates of the posterior probabilities to minimize directly the misclassification risk. MRM builds on the approach started by Short and Fukunaga [6] metric that minimize the difference between finite risk and asymptotic risk. MRM uses a Naïve Bayes to estimate the conditional probabilities for this reason the time of execution is high but as showed in [3] MRM outperformed other distance metrics like HEOM, DVDM, IVDM and HVDM. Despite its name, MRM is not a metric because it does not verify the identity of indiscernibles.

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NCM [4], presented by Wang, works on the concept of neighborhood instead of neighbor. Once defined a topological space, the neighborhoods are regions in the data space that include a specific data point in a query. The similarity between two points is given by the number of neighborhoods that cover both points. In order to assess which points are closer to a test point NCM counts the neighborhoods and chooses the points that have more neighborhoods that are common with the test point. There are many ways to define a topological space. Wang defines the topological neighborhoods as a hypertuple and derive a method for counting in an efficient way all the possible neighborhoods. NCM can work with mixed-feature datasets, namely with both numerical and categorical features. NCM is simple to implement and has a polynomial complexity in the number of attributes. In [4] NCM is shown to outperform HEOM, DVDM, IVDM and HVDM, however, Wang did not test NCM against MRM for its high computational cost. In this way the author left unasked the questions on the comparison of the two methods.

In this letter, we compare the performance of NCM and MRM completing the comparison that was missing in [4]. Empirical evaluation shows that MRM outperforms NCM and, although the running time of MRM is higher, but in the same order of the running time of NCM and so it is not prohibitive as Wang suggested. The comparison between MRM, NCM and the technique by Paredes and Vidal [5] is beyond the aim of the present letter.

The paper is organized as follows: Section 2 overviews MRM and the similarity measure NCM. Experimental evaluation procedures and results are presented is Section 3. Finally, conclusions are drawn in Section 4.

II. METHODS

In this section we describe the methods tested in the experimental procedure.

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A. Minimum Risk Metric

MRM proposed by Blanzieri and Ricci [3], is a very simple distance that minimizes the risk of misclassification r(x,y) defined as the "the probability of misclassifying x by the 1-nearest neighbor rule given that the nearest neighbor of x using a particular metric is y". MRM is expressed by:

$$MRM(x,y) = r(x,y) = \sum_{i=1}^{m} p(c_i|x)(1 - p(c_i|y))$$
 (1)

where m is the number of classes $p(c_i|x)$ is the probability that x belong to the class c_i and $(1-p(c_i|y))$ is the probability that the point y does not belong to the class c_i . Given a point x and a point y, respectively belonging one to the test set and one to the training set, the risk to misclassify x when assigning it the same label of y is given by $p(c_i|x)(1-p(c_i|y))$. The total finite risk is the sum of the risks extended to all different classes like in (1). Several estimations of the conditional probabilities in (1) are possible so MRM can be considered a family of distances. A simple choice is to estimate the conditional probabilities $p(c_i|x)$ using the Naïve Bayes estimator. The idea of minimization of the expected risk as a distance has been more recently re-proposed by Mahamud and Herbert [7] who defined r(x, x') as the "conditional risk of assigning input x with the class label corresponding to x'". They also demonstrated the optimality of it in terms of minimization of the expectation of r(x, x') over the sampling of test points and learning points. They pointed out that the (1) holds only in the case the samples are identically and independently distributed. Instead of estimating the risk by means of estimation of $\hat{p}(x|c_i)$ in (1) they estimate r(x,y) directly as a function of a distance d. From their work we can derive that MRM is symmetric, subadditive (the triangular inequality holds), but it does not verify the identity of indiscernibles (namely MRM(x, y) = 0iff x = y does not hold true) so MRM is not a metric but only a distance.

B. Neighborhood Counting Measure

The NCM proposed by Wang [4] is a similarity measure defined as:

$$NCM(x,y) = \prod_{i}^{n} C(x_i, y_i) / C(x_i)$$
 (2)

where

$$C(x_i, y_i) = \begin{cases} (max(a_i) - max(x_i, y_i)) \times \\ (min(x_i, y_i) - min(a_i)), & \text{if } a_i \text{ is numerical} \\ 2^{m_i - 1}, & \text{if } a_i \text{ is categorical and } x_i = y_i \\ 2^{m_i - 2}, & \text{if } a_i \text{ is categorical and } x_i \neq y_i \end{cases}$$

$$(3)$$

$$c(x_i) = \begin{cases} (max(a_i) - x_i) \times (x_i - min(a_i)) & \text{if } a_i \text{ is numerical to outperform this metrics in terms of accuracy.} \\ 2^{m_i - 1}, & \text{if } a_i \text{ is categorical} \end{cases}$$

Where n is the number of attributes of the data, a_i indicates the i-th attribute and m_i is defined as $m_i = |domain(a_i)|$, finally x_i, y_i indicate the value of the i-th attribute in x and y respectively. In (2) each factor in the product is the number of

TABLE I DATA SETS USED IN THE EXPERIMENTS.

DataSet	Instances	N_Att	Clas_Val	Type	Missing
anneal	898	38	5	Mixed	no
auto	205	25	6	Mixed	no
breast-cancer	286	9	2	Mixed	yes
bridges-v1	108	11	6	Mixed	yes
credit-rating	690	15	2	Mixed	yes
german-credit	1000	20	2	Mixed	yes
Z00	101	17	7	Mixed	no
credit	490	15	2	Mixed	yes
hepatitis	155	19	2	Mixed	yes
horse-colic-data	368	22	2	Mixed	yes
bridges-v2	108	11	6	Nominal	yes
vote	435	16	2	Nominal	yes
tic-tac-toe	958	9	2	Nominal	no
soybean	683	35	19	Nominal	yes
audiology	226	69	24	Nominal	no
primary-tumor	339	17	22	Nominal	yes
sonar	208	60	2	Numerical	no
vehicle	846	18	4	Numerical	no
wine	178	13	3	Numerical	no
yeast	1484	8	10	Numerical	no
ecoli	336	7	8	Numerical	no
Glass	214	9	7	Numerical	no
heart-statlog	270	13	2	Numerical	no
pima-diabetes	768	8	2	Numerical	no
ionosphere	351	34	2	Numerical	no
iris	150	4	3	Numerical	no

27 Dataset: 10 Mixed, 6 Nominal, 10 Numerical

neighborhoods for the i-th attribute. The NCM similarity has linear complexity with respect to the number of attributes. The idea underlying (2) is to count the neighborhoods between two points, where neighborhoods are defined basing on the notion of hypertuple. If the *j*-th attribute is categorical the number of neighborhoods is derived by the number of subsets that cover both points, otherwise if the j-th attribute is numerical the number of neighborhoods is represented by the number of intervals that generate a hypertuple that covers both points. Wang first develops the algorithm for counting the neighborhoods for categorical attributes and numerical attribute with finite domain. NCM as expressed in (2) is more general, and it takes into account also numerical attributes with infinite domain and it assumes the attributes to be equally important. The original paper did not mention how to manage the exceptions derived by 0/0 in (2) but following the indication of the author [8] in case of 0/0 we set the ratio equal to 1. In our experiments, we have used the distance 1 - NCM(x, y) as indicated by the author [8].

III. EMPIRICAL EVALUATION

The aim of the experiment is to empirically compare the performance of NCM and MRM in classification tasks using the k-NN algorithm. We do not include HEOM, VDM and the standard Euclidean and Hamming metrics in our experiments because both MRM [3] and NCM [4] has been already shown to outperform this metrics in terms of accuracy.

A. Experimental procedure

The k-NN is applied to NCM and MRM and the decision on the class is taken with a simple voting schema on the k neighbors; in one specific case (with k=21) the decision is

taken with a weighted voting schema. In order to reproduce the results of [4] we considered the same datasets. Table I shows the datasets used. All of them originated from UCI machine learning repository [9]. The datasets have heterogeneous composition (Nominal, Numerical, and Mixed), in order to evaluate the performance in datasets with different kind of features. The datasets used in [4] show some little difference in terms of number of attributes and number of instances. In particular Anneal, Credit and Soybean differs in the numbers of instances, whereas Vote, Zoo and Horse-colic differs in the number of attributes.

We ran 10-fold cross-validation 10 times with random partitions of data for each data set and for each k value. In each test we assess the statistical differences using the two-tail paired t-Test with significance level equal to 0.05. In order to compare directly with the data published by Wang [4] in a first set of experiments we set k = 11, and k = 21 with the weighted voting scheme. MRM is a distance function so we used as weight its inverse 1/distance whereas for NCM we used as weight directly the *similarity* value. We also run tests with the usual values of k=1, 3, 5, 7. We implemented both methods in Java and we used Weka [10] to perform all the tests, the statistical analysis and the measures of the running time. The Naïve Bayes used in MRM is the one provided by Weka with numeric attributes that have been discretized replicating the choice done in [3]. In the training phase we create an hash table containing the estimates of $\hat{p}(c_i|y)$, with y belonging to the training set. In this way, we compute a priori the estimates of conditional probabilities for the training set. All the code and the datasets used in our tests are available on request.

B. Results and Discussion

The results of the experiments are presented as follows. Table II presents the results of k-NN with NCM and MRM and k=11, Table III presents the results with k=21 and the weighted voting scheme. Tables IV-V present the results with k=1, 3, 5, 7. The tables present the statistical significance of the difference of the accuracy of MRM against NCM. In order to show visually the differences between the methods, Fig. 1 report scatter plots of the accuracies of MRM against NCM. Finally, Table VI shows a representative example of the running time of the two methods.

1) Reproducing NCM results: We compared the results for NCM of Tables II-III with the analogous results presented by Wang. The variations in the selection of the folds of the datasets can account for the differences in the accuracies between our results and Wang's, however, the differences are in the same order of the standard deviation so our results are substantially aligned with the results of NCM in its original proposal.

2) NCM vs MRM: MRM demonstrates good performances in all the tests. Considering the number of datasets in which MRM has significative differences with respect to NCM, MRM is better than NCM, especially with k=11 and k=3. For example, with k=1, MRM is significantly better of NCM in half of the datasets. These good results are also evident in the

TABLE II ACCURACY RESULTS OF k-NN using NCM and MRM with k=11. Significative differences are computed with respect to NCM.

Data Set	NCM	MRM
anneal	94.30±1.92	95.93±2.20
audiology	57.76 ± 7.02	72.24±6.03 o
autos	56.66 ± 10.47	65.17±10.930
breast-cancer	73.68 ± 3.85	73.16 ± 6.71
bridges-v1	57.60 ± 8.32	68.50±10.460
bridges-v2	57.95 ± 7.39	65.82±10.220
credit-rating	76.25 ± 5.24	86.22±3.80 o
german-credit	71.70 ± 2.64	75.04±3.56 o
credit	76.31 ± 5.71	87.18±4.37 o
pima-diabetes	72.58 ± 4.22	75.26 ± 4.78
ecoli	80.21 ± 5.45	80.84 ± 4.63
Glass	62.01 ± 8.82	72.60±8.63 o
heart-statlog	70.93 ± 9.51	82.56±6.12 o
hepatitis	84.07 ± 7.69	84.34 ± 10.42
horse-colic-data	78.81 ± 5.97	79.54 ± 5.83
ionosphere	88.70 ± 4.73	89.40 ± 4.81
iris	93.47 ± 5.92	93.33 ± 5.76
primary-tumor	43.30 ± 6.08	48.88±5.16 ∘
sonar	70.81 ± 8.70	76.71 ± 9.61
soybean	47.44 ± 4.13	92.93±2.95 o
tic-tac-toe	98.24 ± 1.38	69.64±4.40 •
vehicle	68.22 ± 3.85	60.96±3.44 ●
vote	93.50 ± 3.52	90.02±3.91 ●
wine	89.26 ± 6.44	98.71±2.64 ∘
yeast	54.50 ± 3.85	57.53±3.79 o
ZOO	84.66 ± 6.29	90.35 ± 7.74

o, • statistically significant improvement or degradation

TABLE III

ACCURACY RESULTS OF k-NN using NCM and MRM with k=21 and weighted voting scheme. Significative differences are computed with respect to NCM.

Data Set	NCM	MRM
anneal	96.27 ± 1.57	95.87 ± 2.22
audiology	68.24 ± 7.65	71.79 ± 6.15
autos	75.91 ± 8.94	65.17±10.93●
breast-cancer	74.64 ± 4.00	73.16 ± 6.71
bridges-v1	63.05 ± 7.94	68.24 ± 10.01
bridges-v2	62.26 ± 9.08	64.29 ± 9.35
credit-rating	77.26 ± 5.22	86.22±3.80 o
german-credit	72.83 ± 2.91	75.04 ± 3.56
credit	77.88 ± 5.57	87.18±4.37 o
pima-diabetes	75.02 ± 4.54	75.26 ± 4.78
ecoli	79.17 ± 4.76	81.05 ± 4.82
Glass	65.80 ± 8.15	71.38 ± 8.62
heart-statlog	74.52 ± 7.83	82.56±6.12 o
hepatitis	81.38 ± 6.01	84.34 ± 10.42
horse-colic-data	83.77 ± 5.69	79.54 ± 5.83
ionosphere	89.52 ± 4.72	89.40 ± 4.81
iris	93.60 ± 5.76	93.33 ± 5.76
primary-tumor	47.08 ± 5.20	49.03 ± 5.26
sonar	66.43 ± 8.10	76.71±9.61 o
soybean	49.69 ± 4.18	92.91±2.95 o
tic-tac-toe	85.31 ± 2.99	69.64±4.40 •
vehicle	68.90 ± 3.73	61.05±3.46 ●
vote	91.72 ± 4.00	90.02 ± 3.91
wine	90.45 ± 5.98	98.71±2.64 ∘
yeast	55.84 ± 3.64	57.37 ± 3.70
ZOO	95.05 ± 6.24	92.81 ± 6.95

 $\circ,\, \bullet$ statistically significant improvement or degradation

scatter plots in Fig. 1. Considering all the results in Tables IV-V we can observe that increasing the value of k the only three datasets in which NCM reaches an accuracy significantly higher than MRM are Vote, Tic-Tac-Toe and Vehicle. This result is not surprising given the optimality of the minimization of the risk as a distance for classification tasks [7].

The running time results in Table VI show that the running time of MRM is in the same order of the running time of NCM. NCM is always faster of MRM except in the datasets

TABLE IV Accuracy results of k-NN of NCM and MRM with k=1 and k=3. Significative differences are computed with respect to NCM.

	k=1		1	2
D + C +	10 -		k=3	
Data Set	NCM	MRM	NCM	MRM
anneal	97.04 ± 1.75	95.95 ± 2.19	96.31 ± 1.74	95.95 ± 2.19
audiology	76.23 ± 7.64	72.64 ± 6.10	67.25 ± 7.60	72.64±6.10 o
autos	77.63 ± 8.90	65.17±10.93●	66.80 ± 10.43	65.17 ± 10.93
breast-cancer	72.62 ± 6.02	73.16 ± 6.71	73.09 ± 4.92	73.16 ± 6.71
bridges-v1	59.43 ± 11.45	68.31±10.900	56.79 ± 10.69	68.10±10.230
bridges-v2	57.87 ± 11.61	67.32±11.160	53.92 ± 11.30	67.33±10.990
credit-rating	78.51 ± 4.76	86.22±3.80 o	78.35 ± 4.73	86.22±3.80 ∘
german-credit	67.85 ± 4.41	75.02±3.58 o	70.89 ± 3.33	75.01±3.59 o
credit	77.53 ± 5.16	87.18±4.37 o	78.76 ± 4.54	87.18±4.37 o
pima-diabetes	64.82 ± 6.28	75.26±4.78 o	67.55 ± 5.65	75.26±4.78 o
ecoli	77.08 ± 6.24	81.84±4.70 o	80.95 ± 5.58	81.93 ± 4.67
Glass	73.05 ± 8.64	69.34 ± 8.22	69.29 ± 8.40	70.45 ± 8.39
heart-statlog	70.22 ± 9.31	82.56±6.12 o	70.96 ± 8.07	82.56±6.12 o
hepatitis	76.54 ± 7.45	84.34 ± 10.42	81.15 ± 8.30	84.34 ± 10.42
horse-colic-data	73.53 ± 8.10	72.22 ± 16.83	72.53 ± 6.71	79.54±5.83 o
ionosphere	86.22 ± 5.27	89.40 ± 4.81	88.69 ± 4.58	89.40 ± 4.81
iris	94.53 ± 5.56	93.33 ± 5.76	94.73 ± 5.22	93.33 ± 5.76
primary-tumor	40.03 ± 6.32	43.48 ± 6.31	42.45 ± 5.42	43.86 ± 5.64
sonar	68.40 ± 8.90	76.71 ± 9.61	69.00 ± 9.01	76.71 ± 9.61
soybean	61.69 ± 4.86	92.94±2.92 o	54.60 ± 4.25	92.94±2.92 o
tic-tac-toe	98.73 ± 1.15	69.64±4.40 •	98.73 ± 1.15	69.64±4.40 •
vehicle	69.53 ± 3.66	59.80±3.97 ●	69.00 ± 4.45	60.74±3.88 ●
vote	81.54 ± 5.79	90.02±3.91 ∘	88.32 ± 4.92	90.02 ± 3.91
wine	92.07 ± 6.03	98.71±2.64 ∘	90.10 ± 6.04	98.71±2.64 ∘
yeast	50.48 ± 3.81	57.14±4.09 o	52.90 ± 4.06	57.45±3.80 ∘
Z00	93.18 ± 6.71	93.21 ± 7.35	94.45 ± 6.37	93.21 ± 7.35

o, • statistically significant improvement or degradation

TABLE V Accuracy results of k-NN of NCM and MRM with k=5 and k=7. Significative differences are computed with respect to NCM.

	k=5		k=7	
Data Set	NCM	MRM	NCM	MRM
anneal	96.15±1.47	95.95±2.19	95.59±1.54	95.95±2.19
audiology	65.71 ± 8.24	72.64±6.10 o	60.97 ± 7.71	72.64±6.10 o
autos	65.15 ± 10.48	65.17 ± 10.93	61.82 ± 10.91	65.17 ± 10.93
breast-cancer	74.07 ± 3.81	73.16 ± 6.71	74.31 ± 3.96	73.16 ± 6.71
bridges-v1	61.50 ± 9.54	68.59 ± 10.69	59.82 ± 8.28	68.42±10.700
bridges-v2	59.70 ± 9.96	66.10±11.260	57.26 ± 9.14	65.90±10.760
credit-rating	78.39 ± 4.77	86.22±3.80 o	77.86 ± 4.83	86.22±3.80 o
german-credit	70.94 ± 3.10	74.93±3.54 o	72.10 ± 2.96	74.93 ± 3.54
credit	77.88 ± 5.21	87.18±4.37 o	77.65 ± 5.51	87.18±4.37 o
pima-diabetes	69.06 ± 5.41	75.26±4.78 o	71.12 ± 4.77	75.26±4.78 o
ecoli	81.90 ± 5.72	81.93 ± 4.61	80.80 ± 5.56	81.63 ± 4.47
Glass	64.98 ± 8.88	72.64±8.64 o	62.75 ± 8.90	72.69±8.57 o
heart-statlog	70.81 ± 8.56	82.56±6.12 o	71.52 ± 8.68	82.56±6.12 o
hepatitis	82.03 ± 8.42	84.34 ± 10.42	84.20 ± 8.48	84.34 ± 10.42
horse-colic-data	78.75 ± 6.06	79.54 ± 5.83	77.39 ± 6.48	79.54 ± 5.83
ionosphere	89.24 ± 4.27	89.40 ± 4.81	89.66 ± 4.33	89.40 ± 4.81
iris	94.00 ± 5.48	93.33 ± 5.76	93.60 ± 5.52	93.33 ± 5.76
primary-tumor	43.33 ± 5.84	48.20±6.20 ∘	43.33 ± 5.93	48.49±5.56 ∘
sonar	66.98 ± 8.88	76.71±9.61 o	66.40 ± 7.92	76.71±9.61 o
soybean	50.79 ± 4.43	92.94±2.92 o	48.99 ± 4.51	92.94±2.92 o
tic-tac-toe	98.73 ± 1.15	69.64±4.40 •	98.73 ± 1.15	69.64±4.40 •
vehicle	67.85 ± 3.79	60.71±3.91 ◆	68.27 ± 3.79	60.86±3.57 •
vote	92.74 ± 3.50	90.02±3.91 ●	92.81 ± 3.94	90.02±3.91 ●
wine	90.15 ± 6.15	98.71±2.64 ∘	89.08 ± 6.49	98.71±2.64 ∘
yeast	54.30 ± 4.12	57.34±3.86 ∘	55.25 ± 4.16	57.45 ± 3.80
Z00	90.03 ± 6.59	93.11 ± 7.32	86.85 ± 6.83	89.85 ± 7.54

 \circ , ullet statistically significant improvement or degradation

Breast-cancer and Vote where MRM has a smaller running time. However, the differences of running time between the two methods are not impressive. We can conclude that the computation of MRM is not prohibitive, far from the results of being ten times slower than NCM reported by [4] for a straightforward implementation. Wang did not report running times so it is not possible a comparison. It is important to note that the only improvement in our implementation with respect to a straightforward implementation is that we compute the estimates $\hat{p}(c_i|y)$ for y in the training set only once.

IV. CONCLUSION

We have presented an empirical comparison of MRM and NCM. The motivation for the empirical comparison arises from the fact that Wang in [4] did not performed the comparison arguing the MRM was computationally heavy. Our comparison shown that MRM outperforms NCM. MRM, with a simple implementation has a computational cost slightly bigger than NCM, does not have a cost prohibitively high as the work by Wang suggested.

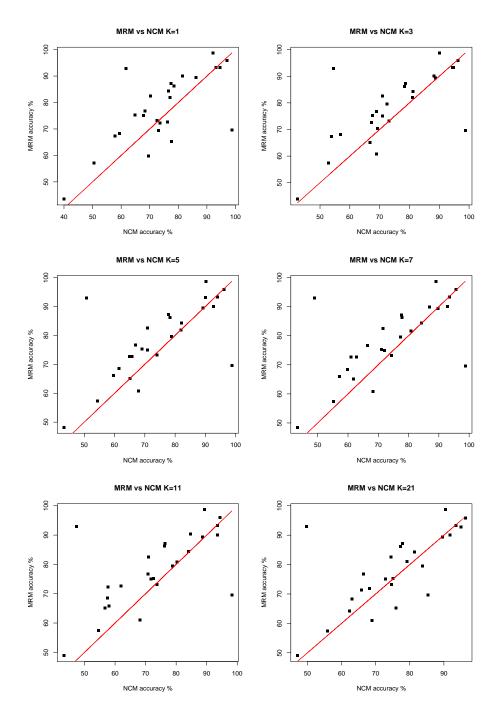


Fig. 1. Scatterplots of the accuracies of Tables II-V. MRM vs NCM for k=1, 3, 5, 7, 11, 21.

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TABLE VI RUNNING TIME IN SECONDS FOR THE TEST PHASE OF k-NN using the Two methods with k=1.

Data Set	NCM	MRM
anneal	1.46±0.65	2.62±0.87
audiology	0.15 ± 0.03	1.02 ± 0.37
autos	0.05 ± 0.03	0.19 ± 0.07
breast-cancer	0.06 ± 0.02	0.03 ± 0.07
bridges-v1	0.00 ± 0.00	0.03 ± 0.02 0.01 ± 0.00
bridges-v2	0.01 ± 0.00	0.01 ± 0.00
credit-rating	0.31 ± 0.05	0.01 ± 0.00 0.29 ± 0.08
german-credit	1.14 ± 0.38	0.25 ± 0.00 0.85 ± 0.31
credit	0.16 ± 0.04	0.03 ± 0.31 0.22 ± 0.13
pima-diabetes	0.08 ± 0.03	0.32 ± 0.19
ecoli	0.03 ± 0.03 0.02 ± 0.00	0.09 ± 0.03
Glass	0.02 ± 0.00 0.01 ± 0.02	0.09 ± 0.03 0.04 ± 0.02
heart-statlog	0.01 ± 0.02 0.02 ± 0.01	0.04 ± 0.02 0.02 ± 0.01
_	0.02 ± 0.01 0.01 ± 0.00	0.02 ± 0.01
hepatitis horse-colic-data	0.01 ± 0.00 0.14 ± 0.04	0.02 ± 0.01 0.13 ± 0.07
	0.14 ± 0.04 0.08 ± 0.05	0.13 ± 0.07 0.25 ± 0.10
ionosphere iris	0.08 ± 0.03 0.00 ± 0.01	0.23 ± 0.10 0.01 ± 0.00
1110		
primary-tumor	0.17 ± 0.06	0.44 ± 0.11
sonar	0.06 ± 0.03	0.24 ± 0.72
soybean	0.86 ± 0.10	4.10 ± 0.16
tic-tac-toe	0.52 ± 0.21	0.41 ± 0.02
vehicle	0.32 ± 0.13	0.96 ± 0.36
vote	0.20 ± 0.08	0.10 ± 0.02
wine	0.01 ± 0.00	0.02 ± 0.03
yeast	0.43 ± 0.13	2.82 ± 0.98
ZOO	0.02 ± 0.00	0.01 ± 0.00



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