

Fig. 3.  $P_{N,sym}^*$  versus N.

observations). Direct calculation of  $P_N^*(\infty)$  was possible for small values of N. For larger values of N it became necessary to upper-bound  $P_N^*(\infty)$ . A Chernoff bound on  $P_N^*(\infty)$  is

$$P_N^*(\infty) \leq (2\sqrt{pq})^N/2.$$

Fig. 3 shows  $P_{N,\text{sym}}^*(m)$  for various numbers of observations N and various sizes of memory m. The Chernoff bound on  $P_N^*(\infty)$  was used for  $N \geq 32$ .

Quite naturally one does better with more memory. The  $P_{N,\text{sym}}^*(m)$ curve for any given value of m follows the  $P_N^*(\infty)$  line for low values of N, diverges from it for larger values of N, and approaches a nonzero limit  $P_{\infty}^*(m)$  as  $N \to \infty$ . This behavior is easily explained. Any given machine can "remember" all of the observations for low values of N. Here infinite memory offers no advantages. For larger values of N, a finite-state machine necessarily loses some information and thus does not do so well as one with infinite memory. As  $N \to \infty$ ,  $P_{N,\text{sym}}^*(m)$ approaches  $P_{\infty}^*(m)$ , the infinite-time lower bound on the probability of error, since from [1] we know that for  $N = \infty$  the optimal machine is symmetric.

### CONCLUSIONS

Experimentally it has been found to be true that the optimal machine has the conjectured form except for the symmetry requirement. Transitions are only between adjacent states. Randomization is found only in transitions to states with a higher probability of error. As  $N \to \infty$ , the probability of a transition out of an end state tends monotonically to 0. It was found that  $P_N^*$  and  $p_{12}(H)$  approach their respective limits  $(P_{\infty}^*)$  and 0) as  $(\ln N)/N$ . A more fundamental argument was given for this behavior.

It was felt that further experiments would only tend to confirm the findings presented. It would be more interesting at this point to attempt mathematical proofs of behavior, specifically of the conjectures. It is hoped that these experimental results will aid in this endeavor. Investigation of deterministic machines for continuousprobability distributions is also of interest and is being pursued [3].

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# The Reduced Nearest Neighbor Rule

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Abstract—A further modification to Cover and Hart's nearest neighbor decision rule, the reduced nearest neighbor rule, is introduced. Experimental results demonstrate its accuracy and efficiency.

The nearest neighbor rule was originally proposed by Cover and Hart [1], [2] and is currently being used by several workers. One reason for the use of this rule is its conceptual simplicity, which leads to straightforward, if not necessarily the most efficient, programming. In a subsequent paper, Hart [5] suggested a means of decreasing memory and computation requirements. This paper introduces a technique, the reduced nearest neighbor rule, that can lead to even further savings. The results of this new rule are demonstrated by applying it to the "Iris" data [6].

The nearest neighbor rule (NN) is described in several places in the literature [1], [2]. For background a simple statement will be included here. First, some notation must be defined. Assume there are Mpattern classes, numbered  $1, 2, \dots, M$ . Let each pattern be defined in an N-dimensional feature space and let there be K training patterns. Each training pattern is a pair  $(x^i, \theta_i)$ ,  $1 \le i \le K$ , where  $\theta_i \in \{1, 2, \dots, M\}$ denotes the correct pattern class and

$$\mathbf{x}^i = (x_1^i, x_2^i, \cdots, x_N^i)$$

is the set of feature values for the pattern. Let  $T_{NN} = \{(x^1, \theta_1),$  $(x^2,\theta_2),\cdots,(x^K,\theta_K)$  be the nearest neighbor training set. Given an unknown pattern x, the decision rule is to decide x is in class  $\theta_I$  if

$$d(x,x^{i}) \leq d(x,x^{i}), \qquad 1 \leq i \leq K,$$

where  $d(\cdot, \cdot)$  is some N-dimensional distance metric.

Actually, the preceding rule is more properly called the 1-NN rule, since it uses only one nearest neighbor. An obvious generalization of this is the k-NN rule, which takes the k nearest patterns  $i_1, i_2, \dots, i_k$ and decides upon the pattern class that appears most frequently in the set  $\theta_{i_1}, \theta_{i_2}, \cdots, \theta_{i_k}$ .

Hart [5] describes a revised rule called the condensed nearest neighbor rule (CNN). Actually, this is not a new decision rule since it still chooses the class of the nearest neighbor. Rather, the word condensed refers to a procedure for choosing a subset of  $T_{NN}$ , which we call  $T_{\text{CNN}}$ , that should perform almost as well as  $T_{\text{NN}}$  in classifying unknown patterns. In this case, a possible drop in performance is being traded for greater efficiency, both in the amount of memory required to store the training set and in the computation time required to reach a decision. Simulation can be used to decide whether the increased efficiency is worth the degradation in performance.

In the formation of  $T_{CNN}$ , the notion of a consistent subset of  $T_{NN}$ is important. It is simply a subset of  $T_{NN}$  that will classify all the patterns in  $T_{NN}$  correctly. Then the minimal consistent subset is the smallest and therefore the most efficient subset of  $T_{NN}$  that will properly classify every pattern in  $T_{\rm NN}$ . The minimal consistent subset of  $T_{\rm NN}$  is important since, in some sense, it has generalized all the important information out of  $T_{NN}$ . It will turn out that  $T_{CNN}$  must be consistent, but cannot be guaranteed to be minimal.

The algorithm for constructing  $T_{\text{CNN}}$  proceeds as follows.

- 1) The first sample pattern is copied from  $T_{NN}$  to  $T_{CNN}$ .
- 2)  $T_{\text{CNN}}$  is used as the training set to classify each pattern of  $T_{\text{NN}}$ , starting with the first. This is done until one of the following two cases

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Experi- ment	Data Set		•	k		IN Error Prob- ability		CNN Error Prob- ability		RNN Error Prob- ability
1	1			1	120	0.0	20	0.0	15	0.0
	1	X		1	120	0.067	31	0.067	28	0.067
2 3 4 5	1	X	X	1	120	0.067	22	0.033	15	0.033
4	1			3	120	0.0	40*	0.0*	30*	0.0*
5	1	$\boldsymbol{X}$		3	120	0.100	62*	0.067*	56*	0.067*
6 -	1	X	X	3	120	0.033	44*	0.033*	30*	0.033*
7	2			1	132	0.667	7	0.611	4	0.611
7 8 9	2	X		1	132	0.500	24	0.556	20	0.389
9	2	$\boldsymbol{X}$	$\boldsymbol{X}$	1	132	0.722	11	0.889	8	0.889
10	2			3	132	0.778	13*	0.444*	8*	0.444*
11	2	X		3	132	0.333	48*	0.556*	40*	0.389*
12	2	X	$\boldsymbol{X}$	3	132	0.833	22*	0.889*	16*	0.889*
13	2 3			1	18	0.833	5	0.811	3	0.833
14	3	$\boldsymbol{X}$		1	18	0.667	6	0.636	4	0.598

TABLE I
RESULTS OF IRIS DATA SIMULATIONS RUN ON THE CDC3600 AT
MICHIGAN STATE UNIVERSITY

- a) every pattern in  $T_{\rm NN}$  is correctly classified, in which case, the process terminates;
- b) one of the patterns in  $T_{\rm NN}$  is classified incorrectly, in which case, go to 3).
- 3) Add the pattern from  $T_{\rm NN}$  that was incorrectly classified to  $T_{\rm CNN}$ . Go to 2).

There are two things that must be noted about this algorithm. First, the number of nearest neighbors to be considered k is not mentioned. Hart [5] mentions  $k \neq 1$  as being a possibility for future research. However, this suggestion contains a fundamental flaw. Take k to be 2n+1. To guarantee a unique solution, at least n+1 of the nearest neighbors must be from the correct pattern class in order for the new pattern to be correctly classified. In forming  $T_{\text{CNN}}$ , an incorrectly classified pattern from  $T_{\text{NN}}$  is added to  $T_{\text{CNN}}$ . In particular, for the worst case, it would have to be added to  $T_{\text{CNN}}$  in particular, for the this does not contradict any assumptions that have been made, it is undesirable in the sense that the  $k \neq 1$  condensed nearest neighbor rule will always make the same decision as the 1-CNN rule. A question that this discussion prompts is, "How many times does this worst case occur?" Experimental evidence will be presented that shows that in 11 out of 12 cases tried, the worst case applied.

An alternative would be to use k=1 to construct  $T_{\text{CNN}}$  and then use k=2n+1 for actual classification of unknown patterns. This method would avoid the problem of ending up with n+1 copies of each training pattern. The results of this procedure are not easy to visualize and would bear further study.

The second item of importance is that the algorithm will terminate for k=1 in all but one case. If  $T_{\rm NN}$  is already a minimal consistent set, then  $T_{\rm CNN}$  will equal  $T_{\rm NN}$ . If this happens, the algorithm will not terminate only if there are two or more patterns  $(x^i,\theta_i)$  and  $(x^j,\theta_j)$  for which  $i \neq j, x^i = x^j$ , but  $\theta_i \neq \theta_j$ . Although this case is not specifically disallowed, it violates the consistency of  $T_{\rm CNN}$ . That is,  $T_{\rm CNN}$  must classify all its member patterns correctly.

One can ask whether this procedure is effective or whether it really reduces the size of the training set or whether  $T_{\rm CNN}$  contains a minimal consistent subset. The answers to these questions come from simulations. In the simulation performed using well-separated data, savings in storage of as great as 90 percent were realized. While this is encouraging, it should be noted that the next rule to be presented, the RNN rule, improved upon CNN in every case tried. This means that  $T_{\rm CNN}$  is not minimal, though how far from minimal it actually is would be difficult to determine.

The last rule to be considered is one that has not appeared in the literature, the reduced nearest neighbor rule RNN. It is an extension of the CNN rule and, like the CNN rule, reduces  $T_{\rm NN}$ . Since it is based on the CNN rule,  $k \neq 1$  will not be considered in this paper. The

algorithm to produce  $T_{\rm RNN}$ , the reduced nearest neighbor training set, is 1) copy  $T_{\rm CNN}$  into  $T_{\rm RNN}$ ; 2) remove the first pattern from  $T_{\rm RNN}$ ; 3) use  $T_{\rm RNN}$  to classify all the patterns in  $T_{\rm NN}$ :

- a) if all patterns are classified correctly, go to 4),
- b) if a pattern is classified incorrectly, return the pattern that was removed and go to 4);

4) if every pattern in  $T_{RNN}$  has been removed once (and possibly replaced) then halt. Otherwise, remove the next pattern and go to 3).

One question of some interest is whether or not  $T_{\rm RNN}$  is a minimal consistent subset of  $T_{\rm NN}$ . Obviously, since  $T_{\rm RNN}$  is a subset of  $T_{\rm CNN}$ , then if  $T_{\rm CNN}$  does not contain the minimal consistent subset of  $T_{\rm NN}$ , neither does  $T_{\rm RNN}$ . An example of a case where  $T_{\rm CNN}$  does not contain a minimal subset for M=2 pattern classes, N=1 features, and K=13 training samples is

$$T_{\text{NN}} = \{(-13,1), (-16,1), (-19,1), (-6,2), (-4,2), (-2,2), (0,2),$$

$$(2,2), (4,2), (6,2), (13,1), (16,1), (19,1)\}$$

$$T_{\text{CNN}} = \{(-13,1), (-6,2), (13,1), (4,2)\}$$

$$T_{\text{RNN}} = T_{\text{CNN}}$$

$$T_{\min}$$
 = minimal consistent subset  
= {(-13,1), (0,2), (13,1)}.

But what happens in the case where  $T_{\text{CNN}}$  does contain a minimal consistent subset? The following example demonstrates just such a case:

$$T_{NN} = \{(0,2), (-3,1), (-2,2), (3,1), (2,2)\}$$

$$T_{CNN} = T_{NN}$$

$$T_{RNN} = \{(-3,1), (-2,2), (3,1), (2,2)\}$$

$$T_{min} = T_{RNN}.$$

If this were always true when  $T_{\min} \subseteq T_{\text{CNN}}$ , the RNN rule would be a very useful tool. It can be shown that this is indeed the case, as follows. Let  $(x^i, \theta_t)$  be any pattern not in  $T_{\min}$ , and let  $(x^m, \theta_m)$  be that pattern in  $T_{\min}$  that causes  $(x^i, \theta_t)$  to be correctly classified. The case that must be considered is when both  $(x^i, \theta_t)$  and  $(x^m, \theta_m)$  are in  $T_{\text{CNN}}$ . By the construction of  $T_{\text{CNN}}$ ,  $(x^t, \theta_t)$  must appear before  $(x^m, \theta_m)$ , since if  $(x^m, \theta_m)$  appears first,  $(x^i, \theta_t)$  would not have been misclassified and therefore would not have been added to  $T_{\text{CNN}}$ . Notice that if there are several elements of  $T_{\min}$  that could properly classify  $(x^t, \theta_t)$ , they must all follow  $(x^t, \theta_t)$  in order for this pattern to be included in  $T_{\text{CNN}}$ .

Now applying RNN to  $T_{\text{CNN}}$ , observe what happens to  $(x^{l}, \theta_{l})$  when it is removed. Since  $T_{\min} \subseteq T_{\text{CNN}} - \{(x^{l}, \theta_{l})\}$  all patterns will be correctly classified and the extraneous pattern will be removed. But what happens when  $(x^{m}, \theta_{m})$  is removed? All patterns that depend on  $(x^{m}, \theta_{m})$  to be correctly classified have already been removed. If they can all be

<sup>\*</sup> Marks the k-CNN and k-RNN rules with  $k \neq 1$ .

correctly classified at this point, then  $(x^m, \theta_m)$  is not required in  $T_{\min}$ which contradicts our initial assumption. Therefore,  $(x^m, \theta_m)$  must be kept and so only those patterns that are not in  $T_{\min}$  are kept.

There were several alternatives possible in the simulations used to study the k-NN, CNN, and RNN rules. Different combinations of these alternatives lead to 14 different experiments. The data used were the so-called "Iris" data [6], which consist of four measurements on each of 150 flowers. There were three pattern classes, Virginica, Setosa, and Versicolor, corresponding to three different types of iris. The 150 samples were divided to form a training set and a test set. This was done in three different ways to give the following three different data

- 1) Random: 120 training patterns were chosen, 40 from each pattern class. The remaining 30 patterns were used to test the training algorithms.
- 2) Pathological: There were 18 of the patterns that were misclassified by the ISODATA clustering algorithm [7] using a Euclidean distance metric. These were used for test patterns. The remaining 132 patterns were used as a training set.
- 3) Inverted Pathological: In an attempt to find out whether all 18 of the patterns mentioned in 2) were misfits, these 18 were used as a training sample to classify the remaining 132 patterns.

A second option that was available was to transform the raw data so that the axes of the data were aligned with the eigenvectors of the data covariance matrix. One can reduce the number of features by selecting only the eigenvectors corresponding to the principal or largest eigenvalues. Finally, the value of k, the number of nearest neighbors, was varied, adding another alternative. The results from these experiments are tabulated in Table I.

Experiment 1 gives some idea of how the 1-NN, CNN, and RNN rules might work in the typical case of well-separated data. Since improvement is impossible, the error probabilities stay the same when going to the three nearest neighbors in experiment 4. In this case the CNN rule offered an 83-percent improvement in memory and time efficiency, and RNN offered an additional 4 percent with no degradation of performance. The average improvement over all the experiments was 84 percent for the CNN rule and an additional 4 percent for the RNN rule. This improvement cost an average increase in the probability of error of 0.0225 for CNN and an average decrease of 0.00037 for the RNN rule.

In only two out of the eight applicable cases did the CNN rule do worse than the NN rule, and in four of the remaining six cases it did better. Similarly, the RNN rule was at least as good as the CNN and NN rules in a majority of the experiments.

As was mentioned above, the k-CNN and k-RNN rules do not appear to be directly useful for  $k \neq 1$ . It was also stated that if k =2n + 1, then the k-CNN and k-RNN training sets could each contain n+1 copies of each training pattern in the corresponding k=1training set. This was considered to be the worst case, and the question was posed of how many times the worst case would come up. Looking at the entries of Table I marked with an asterisk, signifying the k-CNN and k-RNN rules,  $k \neq 1$ , the worst case can be seen to appear in 11 out of 12 possible cases.

To conclude, CNN and RNN are two revised versions of an intuitively appealing decision rule. The question of whether or not they offer any general advantage over a k-NN rule must be answered with more detailed simulations. With the well-separated "Iris" data, the CNN rule appears to be very beneficial, confirming Hart's contention. The RNN rule gives an added advantage, but it is doubtful that the additional time to compute  $T_{RNN}$  required, usually more than twice as much, is worth the extra 3-4-percent gain in efficiency. However, for extremely large training sets, this 3-4 percent might be crucial.

Finally, answering Hart's question about extending the CNN rule for  $k \neq 1$ , which applies to the RNN rule as well, if the k nearest neighbors are used to construct  $T_{CNN}$ , then  $T_{CNN}$  will contain multiple copies of each pattern. On the other hand, if the one nearest neighbor is used to construct  $T_{CNN}$ , there is no assurance that there will be enough patterns from a given class to ever guarantee a majority. Therefore, the conjecture is that k-CNN and k-RNN offer no further improvements.

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# On the Minimum Distortion of Block Codes for a Binary Symmetric Source

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Abstract—It is shown that the mixture of two codes of block length n can have a distortion strictly less than the distortion of the best possible code of block length n having the same rate. The seeming contradiction is explained by offering two interpretations for the mixture of two codes and then explaining why the bound for the performance of the best code cannot be applied directly to the mixture.

Consider a source that emits a sequence of statistically independent equiprobable binary digits. We are concerned with encoding blocks of n source digits, using as codewords a set of M binary n-tuples. The distortion measure is taken as the normalized Hamming distance between a block of n source digits and its associated codeword.

Let  $W = (W_1, W_2, \dots, W_n)$  be a binary *n*-tuple representing a source output sequence and let  $Q(W) = Z = (Z_1, Z_2, \dots, Z_n)$  be the binary n-tuple representing the codeword associated with that sequence. The distortion between W and Z = Q(W) is

$$D(W,Q(W)) = D(W,Z) = \frac{1}{n} \sum_{i=1}^{n} d_H(W_i,Z_i),$$

where

$$d_{H}(W_{i},Z_{i}) = \begin{cases} 1, & W_{i} \neq Z_{i} \\ 0, & W_{i} = Z_{i}. \end{cases}$$

A code is then described by two parameters, its rate R and its average distortion  $\bar{D}$ , defined as

$$R = (1/n)\log_2 M$$

$$\bar{D} = E\{D(W,Z)\},\,$$

where the average is taken over all 2<sup>n</sup> possible source sequences. The rate of the code depends only on the number of codewords, while the average distortion depends upon the choice of the codewords and the choice of a deterministic mapping Q(W), which maps the source outputs into codewords.

The following is a short derivation of a well-known lower bound to D as a function of R and n. Let  $\alpha_j$ ,  $j = 0,1,2,\dots,n$ , be the number of source sequences W for which D(W,Q(W)) = j/n. Then since all source sequences are equally likely

$$\bar{D} = \frac{1}{n2^n} \sum_{j=0}^n j\alpha_j. \tag{1}$$

For any code

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