Nearest neighbor based classifiers

Nearest Neighbor (NN) rule classifies a sample based on the category of it's nearest neighbor. The NN based classifiers use some or all of patterns available intraining set to classify a test pattern. The logic of classification involves finding the similarity between the test pattern and neighboring patterns in the training set.

Nearest neighbor algorithm — The NN algorithm assigns the class label of it's closest neighbor to a test pattern.

If Pisthe test pattern, then if $d(P, X_K) = min \{ d(P, X_i) \}$, where i = 1, 2, ..., n

Pattern P is assigned to the class Ox associated with Xx

Let the training set consist of the following patterns - $X_1 = (0.8, 0.8, 1)$; $X_2 = (1.0, 1.0, 1)$; $X_3 = (1.2, 0.8, 1)$ $X_4 = (0.8, 1.2, 1)$; $X_5 = (1.2, 1.2, 1)$; $X_6 = (4.0, 3.0, 2)$ $X_7 = (3.8, 2.8, 2)$; $X_8 = (4.8, 2.8, 2)$; $X_9 = (3.8, 3.2, 2)$ $X_{10} = (4.2, 3.2, 2)$; $X_{11} = (4.4, 2.8, 2)$; $X_{12} = (4.4, 3.2, 2)$ $X_{13} = (3.2, 0.4, 3)$; $X_{14} = (3.2, 0.7, 3)$; $X_{15} = (3.8, 0.5, 3)$ $X_{16} = (3.5, 1.0, 3)$; $X_{17} = (4.0, 1.0, 3)$; $X_{18} = (4.0, 0.7, 3)$ Test pattern is P = (3.0, 2.0, 2).

> 5 4 3 ×××× 2 P 1 +++ * *

Computing Euclidean distance, e.g. $d(X_1, P) = \sqrt{(0.8-3.0)^2 + (0.8-2.0)^2} = 2.51$ and so on.

It can be verified that $d(X_{16}, P) = 1.12$, which is the smallest distance. Since $X_{16} \in class 3$, point P will be classified to belong to class 3.

K-nearest neighbor algorithm — Here instead of finding just I nearest neighbor, k'neighbors are found. The majority class of these K nearest neighbors is the class label assigned to the new pattern. Typically k' is chosen to be anodd number to break the ties, and arrive at a majority.

Consider the previous example. Based on the distance computed, the 5 nearest neighbors are –

XIG E Class 3

X7 E Class 2

Hence point P will be classified to belong to class 3 based on majority.

XI4 E Class 3

x₆ ∈ Class 2

XIT E Class 3

This method reduces the error in classification when training patterns are noisy, or the chosen training data is challenging and contains outliers. For large data sets, k can be larger to reduce the error. The value of k can be determined to meet the requirement through experimentation.

Modified K-nn (MKNN) algorithm — This algorithm also considers k-neighbors. However, these k-neighbors are weighted according to their distance from the test point. Each of the k' neighbors is associated with the weight w defined as

$$W_j = \frac{dk - dj}{dk - dj}$$
 if $dk \neq d$,

1 if dk = d1 , where j = 1,2,, k.

The value of wij varies from a maximum of I for the nearest nearest neighbor, upto 0 for the most distant nearest

The test pattern P is assigned to that class, for which the weights of the representatives among the k-nearest neighbors sums to the largest value.

Since MKNN applies a weighted majority role instead of simple majority rule, it means that outlier patterns have lesser effect on classification.

Consider again, P = (3.0, 2.0, 9). For the five nearest points, the distances from P are, $d(P, X_{16}) = 1.12$; $d(P, X_7) = 1.13$; $d(P, X_{14}) = 1.32$; $d(P, X_6) = 1.41$; $d(P, X_{17}) = 1.41$.

The values of w will be,

$$W_7 = \frac{1.41 - 1.13}{1.41 - 1.12} = 0.97$$
 $X_7 \in class 2$

$$\omega_{14} = \frac{1.41 - 1.32}{1.41 - 1.12} = 0.31$$
 $\times_{14} \in \text{class 3}$

$$W_6 = 0$$
 $X_6 \in class 2$

$$\omega_{17} = 0$$
 $x_{17} \in class 3$

r-near algorithm - If the nearest neighbor of the point of interest is very far away, then there is no relevance of such point in decision making. Hence, instead of looking at nearest neighbors, only the neighbors within some distance r'of the point of interest can be considered. The algorithm steps are -

i) Given the point P, determine the subdata in hypersphere of radius 'r' centered at P

ii) If B, (p) is empty then output the majority class of the entire data set.

iii) If B, (p) is not empty, output the majority class of the

data points in it

This algorithm can be used to identify outliers. If a pattern does not have any similarity with the patterns within the chosen radius, it can be identified as an outlier.

The nearest neighbor classification method is very time consuming, especially when the number of training patterns is large. To overcome this, many efficient algorithms have been proposed. Some of those use a pre-processing step, which requires additional computing time, but it needs to be done only once.

Branch and bound algorithm — If the data are stored in an ordered fashion, all the points need not be required to be considered for finding the nearest neighbor. If the data is stored in a tree like data structure, the search for the nearest neighbor can be performed efficiently by not having to search down the branches where one can obtain a lower bound for the distance that is higher than the current best value found.

The data is clustered into representative groups Si where each group has the smallest possible radius. Each of

such clusters has center U; and radius v;.

After searching one of the clusters, point X^* is found, which is the nearest neighbor from the cluster to the point of interest P. This distance is say $d = d(P, X_j^*)$. Then lower bounds can be computed for the distances in the other clusters.

For a point $X_k \in S_j$, by triangular inequality $d(P_j N_j) \leq d(P_j X_k) + d(X_k, N_j) \leq d(P_j X_j) + Y_j$.

Hence, a lower bound b; can be computed for the distance between any $X_k \in S_j$ and P as

min d(P,XK) > bj = d(P,Uj)-Vj

Clusters which satisfy the inequality b; ≥d need not be

searched. The following steps can be followed -

i) Cluster the data points into L sets S_i , i=1,2,... L. These are the first level clusters. For each cluster, maintain the centre U_i and the radius Y_i . Within each cluster S_i , cluster the data into L sub-clusters $S_{i,j}$, j=1,2,..., L. Maintain the centre $U_{i,j}$ and the radius $Y_{i,j}$. These are second level clusters. Continue this type of recursive clustering until there are clusters of one point.

ii) To find the nearest neighbor to a new point P,

a. Branch step - First compute bj. Then find the nearest neighbor in the cluster with the smallest bj. If the clusters have subclusters, this step is done recursively.

b. Bound step-If that cluster does not satisfy the bound, look into the cluster with the next highest bj. Otherwise stop and

output the result.

It can be shown that on an average, there is considerable improvement over the standard NN algorithm.

Consider the data for the 18 points mentioned earlier. Let the points of class I be taken as a cluster & similarly points of the other 2 classes as the other two clusters. Further cluster I is subclustered as cluster Ia and Ib & similarly clusters 2a, 2b and 3a, 3b are made. At next level, each point is taken to be a sub-cluster.

_	Cluster 1			(luster 2								
$(X_1 X_2 X_3 X_4 X_5)$				(X ₆ X ₇ X ₈ X ₉ X ₁₀ X ₁₁ X ₁₂)				(X	8)			
(luster la (X, X2 X3) laa lab lac				Cluster 2a (X6 X7 X8 X9) 2aa 2ab 2ac 2ad				Cluster 3a CX13 X14 X15) 3aa 3ab 3ac				
11	X ₁ X ₂	X ₃	20		X ₇		_		XI3	X14	X ₁₅	
\perp	19.6 8.8		74									
1	Clust	er lb	+		luste	v 2h	- 1	1		Cluster	(3b	1
+	(X4 X5)			(X10 X11 X12)				CX16 X17 X18)				+1-
\perp	Iba	lbb			266		c	Ш	3 ba	366	36c	
	X4	Xs		X _t	×II	X		Ш	X16	X17	XIB	2000
$\perp \downarrow$												

It can be seen that centre of cluster 1 is (1.0,1.0). Now consider $X_1 = (0.8,0.8)$. Calculating the radius as $\sqrt{(1-0.8)^2 + (1-0.8)^2} = 0.283$. Similarly calculating the value of the radius for other elements of cluster 1, and choosing maximum amongst them, we get the radius $x_1 = 0.283$. Similar computations are made for the other two clusters.

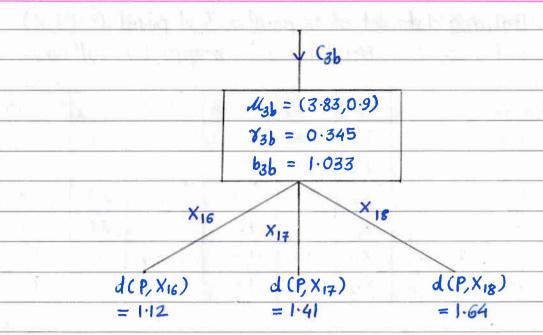
$$\mathcal{U}_{1} = (1 \cdot 0, 1 \cdot 0)$$
 $\mathcal{U}_{2} = (4 \cdot 11, 3 \cdot 0)$ $\mathcal{U}_{3} = (3 \cdot 62, 0 \cdot 72)$
 $\mathcal{V}_{1} = 0 \cdot 283$ $\mathcal{V}_{2} = 0 \cdot 369$ $\mathcal{V}_{3} = 0 \cdot 528$

The test point is P (3.0,2.0).

At the first level, by for each cluster i can be found

Since bz is smallest, the sub-clusters of cluster 3 are searched.

Hence Cab is chosen as bab < bae



The second sub-cluster of cluster 3 is searched and X16 is the point found to be closest to point P. The bound 'd' is calculated as the distance from P to X16, which is 1.12.

Since b_1 and b_2 are greater than 'd', clusters I and 2 need not be searched and X_{16} is considered as the nearest neighbor of P.

Nearest Meighbor search by projection

For a I dimensional space, if the point set is ordered, the nearest neighbor of point P can be found by locating position of P'in the order. Finding the successor and predecessor, and taking

the closest, gives the nearest neighbor.

In the 2-D case, the points are sorted in increasing order using the first co-ordinate value. For finding nearest neighbor of P, locate position of P' in the sorted list of the first co-ordinate value. Then nearest neighbors of P are searched. We examine the point that is closer to P in the x-distance first and remember the closest point to P 60 far observed, say point Q. The search will be pruned as soon as it is known that any other point is further away from P than Q is.

This process is modified to project the points onto both the x and y-axes. The first step is on the x axis, the second on the y-axis, third on x-axis & so on. The search stops, when the

search in either direction stops

Example-Training data set of 18 points. Test point P= (3,2). The entire set of x-coordinates can be mapped as follows-

ē.	Point	Co-ord.	Difference	Point	X Co-ord-	Difference	Point	X co-ord	Difference	
	X	0.8	2.2	X ₇	3.8	0.8	X ₁₃	3.2	0.2	
2.	X ₂	1	2	X8	4.2	1.2	X14	3.2	0.2	
	X3	1.2	1.8	Xg	3.8	0.8	X15	3.8	0.8	
	X4	0.8	2.2	X,	4.2	1.2	X16	3.5	0.5	
	X5	1.2	1.8	XII	4.4	1.4	X 17	4	I	
	X ₆	4	7.71-6	X12	4.4	1.4	XIS	4	1	

These points are rank ordered, and their actual distance from P is found. The point closest to P so far is kept in memory. Hence the rank ordering is - X13, X14, X16, X7, X9, X15, X6, X17,

X18, X8, X10, X11, X12, X3, X5, X2, X1, X4.

Starting from X_{13} , $d(P, X_{13}) = 1.612$ is calculated. X_{13} is kept in memory. After checking for X_{14} , $d(P, X_{14}) = 1.315$ is found and X_{14} is stored in memory. For X_{16} , $d(P, X_{16}) = 1.118$ and hence X_{16} is stored in memory. Continuing further, checking each time that difference $< d(P, X_{16})$, various distances are calculated, which continue to be $> d(P, X_{16})$ and hence X_{16} continues to be stored in memory. The next point X_{8} has a distance in x-direction of 1.2, which is larger than the actual distance of the closest point X_{16} . Then it is not necessary to check any more points.

The entire set of y-coordinates can be mapped as follows-

96	Point	co-ord.	Difference	Point	Co-ord	Difference	2,11(Point	Co-ord.	Difference	
	Xı	0.8	1.2	X ₇	2.8	0.8		X ₁₃	0.4	1.6	
	X ₂	1	Haliz	X ₈	2.8	0.8		X14	0.7	1.3	
	X ₃	0.8	1.2	X ₉	3.2	1.2	108	X15	0.5	1.5	
	X4	1.2	0.8	X _{Io}	3.2	1.2	H	X 16	Los	1.75	
H.	X ₅	1.2	0.8	XII	2.8	0.8	10	X17	7/1	1	
Т	X ₆	3	t Lore	X ₁₂	3.2	1.2) D	X18	0.7	1.3	

For the y-dimension, the rank ordering is - X4, X5, X7, X8, X11, X2, X6, X16, X17, X1, X3, X9, X10, X12, X14, X18, X15, X13. Similar activity like along the x-dimension is carried out. d(P, X16) = 1.118 is the minimum so far and hence X16 is stored. d (P, X17) = 1.414, hence X16 stays stored in memory. The next point X, has a distance in y-direction of 1.2, which is larger than the actual distance of the closest point X16. Hence it is not required to check any more points. Since X16 emerges as the data point stored along x-direction as well as y-direction, after a single pass only X16 is identified as the nearest neighbor.