

Machine Learning CSCE 5215

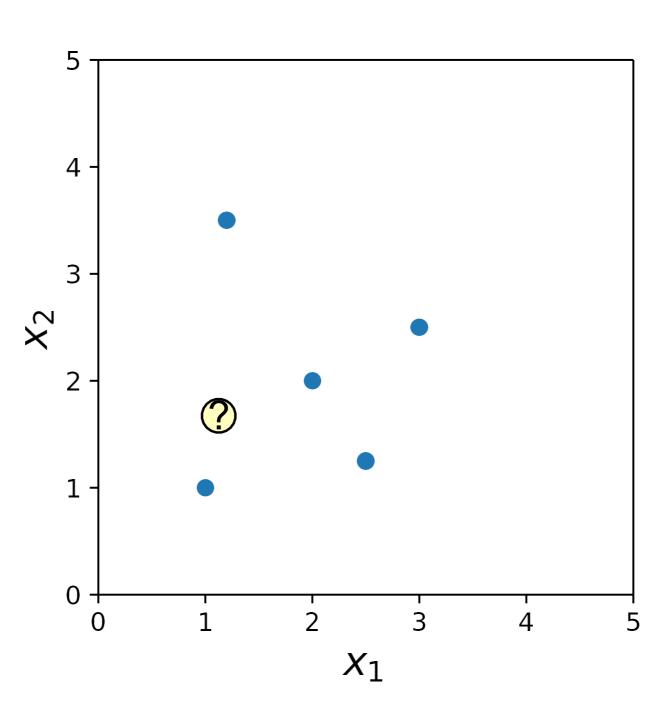
Nearest Neighbor Methods kNN

Instructor: Zeenat Tariq

1-Nearest Neighbor

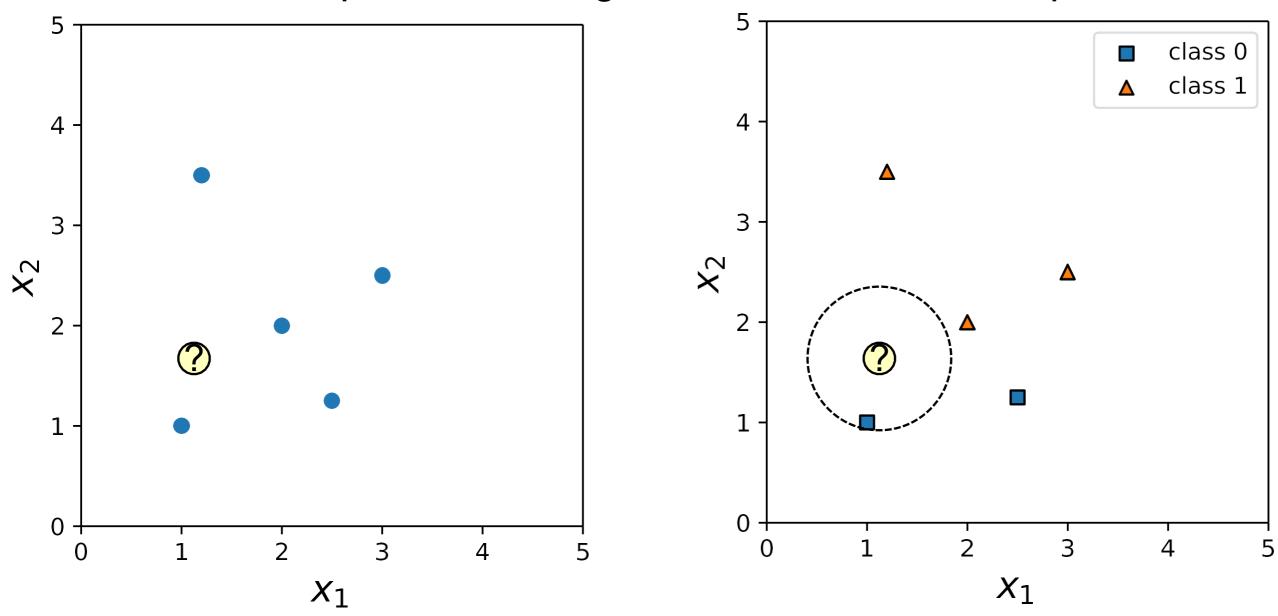
1-Nearest Neighbor

Task: predict the target / label of a new data point



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How? Look at most "similar" data point in training set

1-Nearest Neighbor Training Step

$$\langle \mathbf{x}^{[i]}, y^{[i]} \rangle \in \mathcal{D} \quad (|\mathcal{D}| = n)$$

How do we "train" the 1-NN model?

1-Nearest Neighbor Training Step

$$\langle \mathbf{x}^{[i]}, y^{[i]} \rangle \in \mathcal{D} \quad (|\mathcal{D}| = n)$$

To train the 1-NN model, we simply "remember" the training dataset

1-Nearest Neighbor Prediction Step

Given:
$$\langle \mathbf{x}^{[i]}, y^{[i]} \rangle \in \mathcal{D} \quad (|\mathcal{D}| = n)$$

 $\langle \mathbf{x}^{[q]}, ??? \rangle$

Predict: $f(\mathbf{x}^{[q]})$

Algorithm:

closest_point := None

closest_distance := ∞

query point

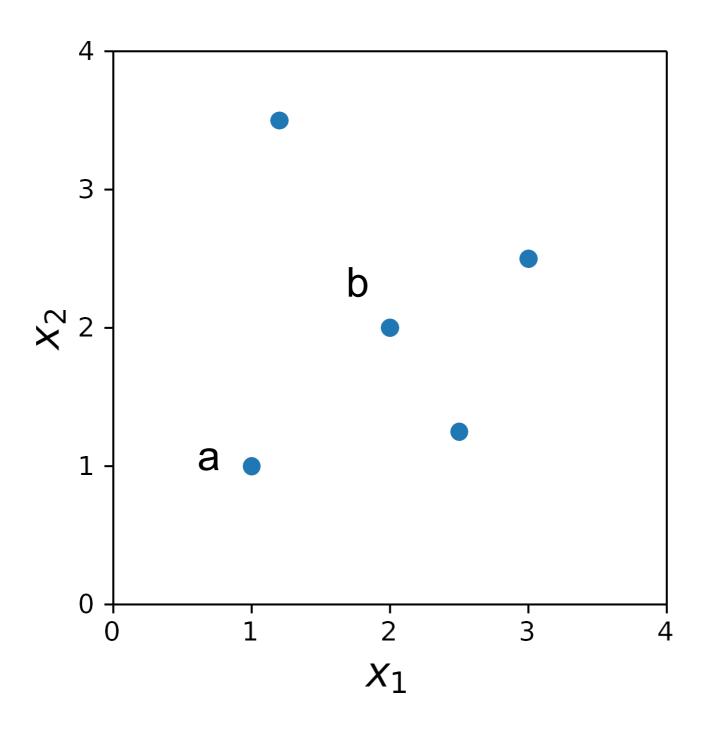
- for i = 1, ..., n:
 - \circ current_distance := $d(\mathbf{x}^{[i]}, \mathbf{x}^{[q]})$
 - if current_distance < closest_distance:</p>
 - closest_distance := current_distance
 - closest_point := $\mathbf{x}^{[i]}$
- return f(closest_point)

Commonly used: Euclidean Distance (L2)

$$d(\mathbf{x}^{[a]}, \mathbf{x}^{[b]}) = \sqrt{\sum_{j=1}^{m} (x_j^{[a]} - x_j^{[b]})^2}$$

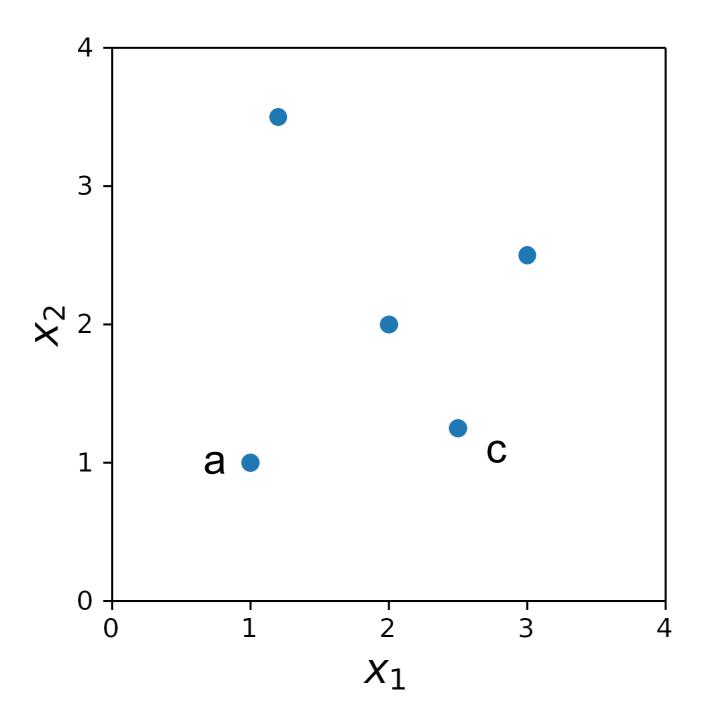
Nearest Neighbor Decision Boundary

Decision Boundary Between (a) and (b)



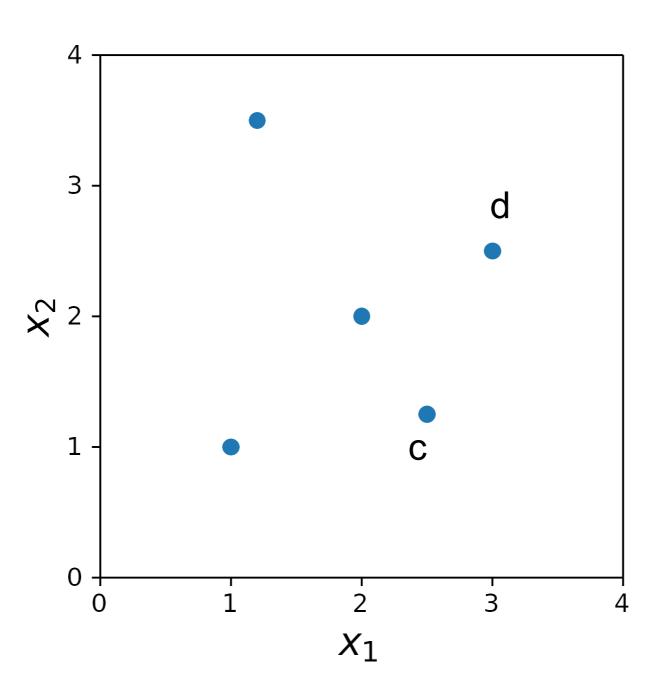
How does it look like?

Decision Boundary Between (a) and (c)

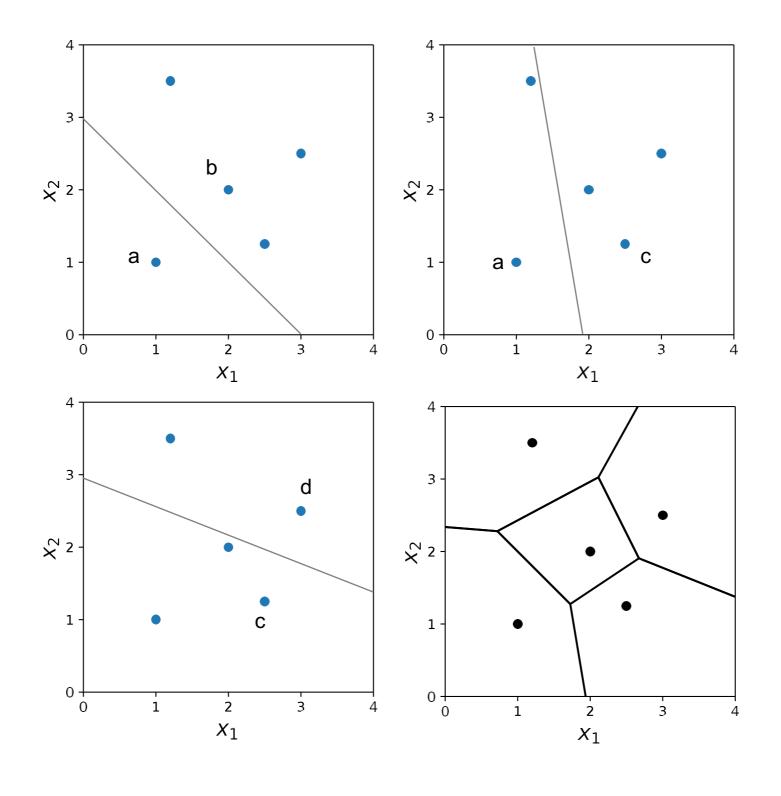


How does it look like?

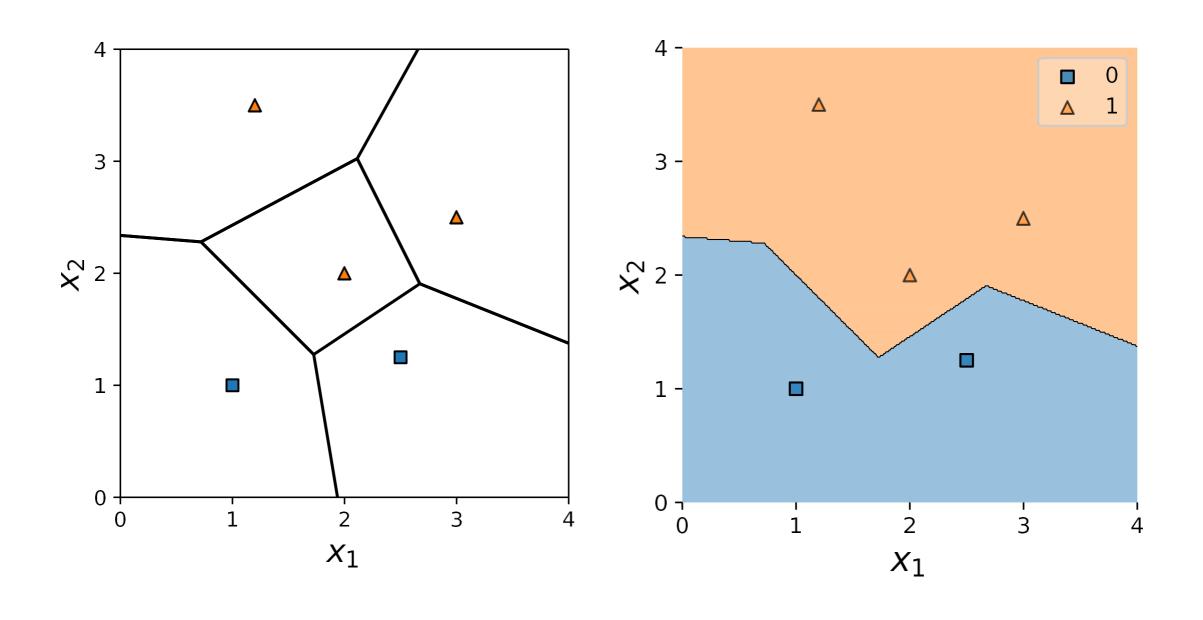
Decision Boundary Between (a) and (c)



Decision Boundary of 1-NN



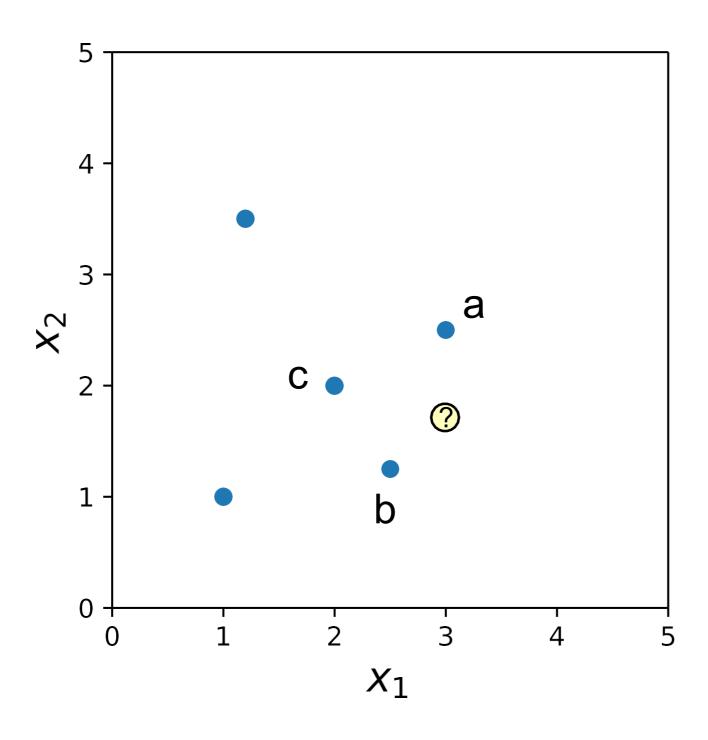
Decision Boundary of 1-NN



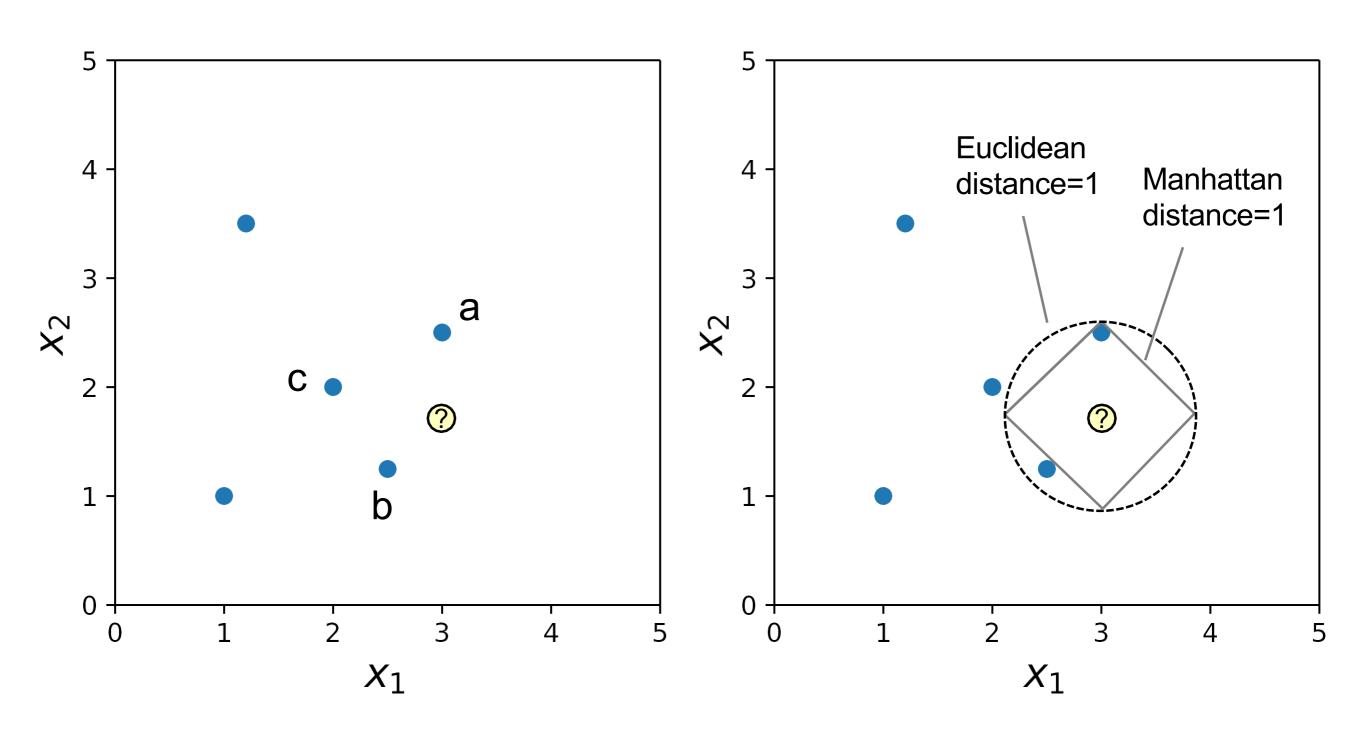
Which Point is Closest to ?







Depends on the Distance Measure!



Some Common Continuous Distance Measures

Euclidean

Manhattan

Minkowski:
$$d(\mathbf{x}^{[a]}, \mathbf{x}^{[b]}) = \left[\sum_{j=1}^{m} \left(\left|x_{j}^{[a]} - x_{j}^{[b]}\right|\right)^{p}\right]^{\frac{1}{p}}$$

Mahalanobis

Cosine similarity

. . .

Some Discrete Distance Measures

Hamming distance:
$$d(\mathbf{x}^{[a]}, \mathbf{x}^{[b]}) = \sum_{j=1}^{m} \left| x_j^{[a]} - x_j^{[b]} \right|$$
 where $x_j \in \{0, 1\}$

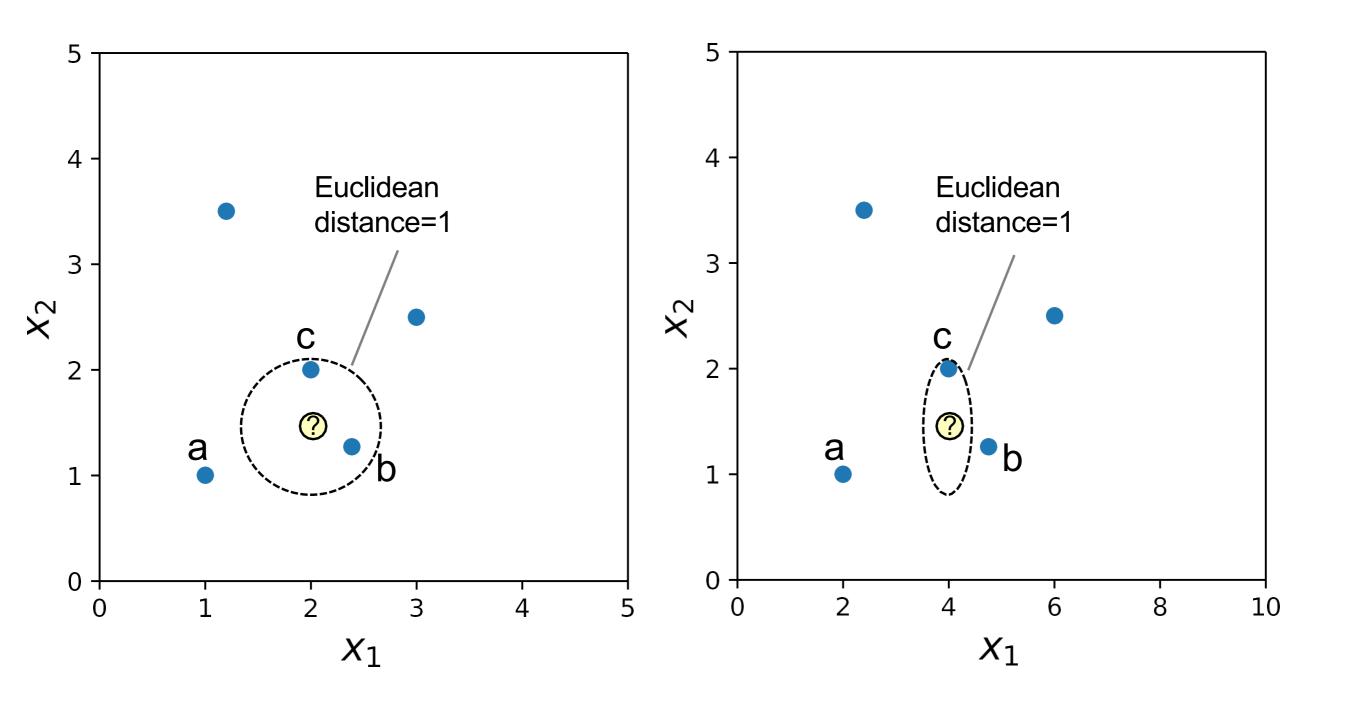
Jaccard/Tanimoto similarity:

$$J(A,B) = \frac{|A \cap B|}{|A \cup B|} = \frac{|A \cap B|}{|A| + |B| - |A \cap B|}$$

Dice:
$$D(A,B) = \frac{2|A \cap B|}{|A| + |B|}$$

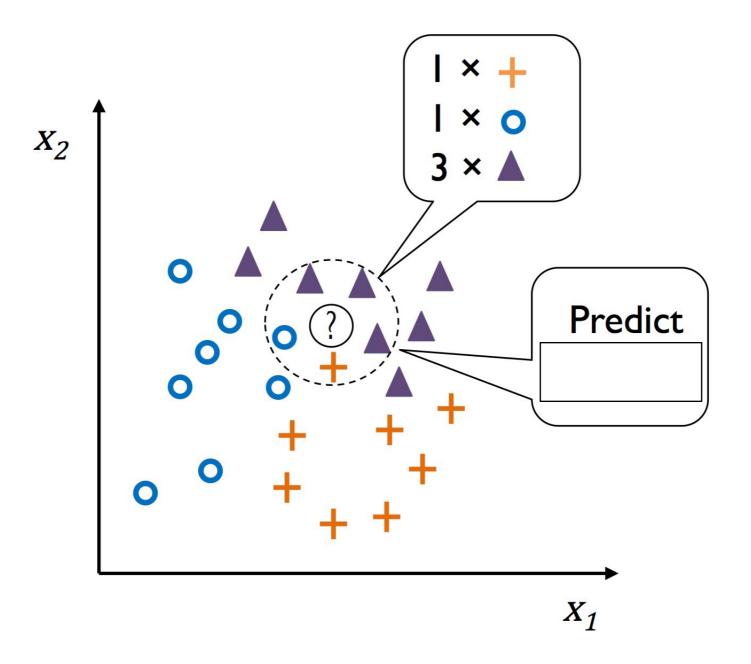
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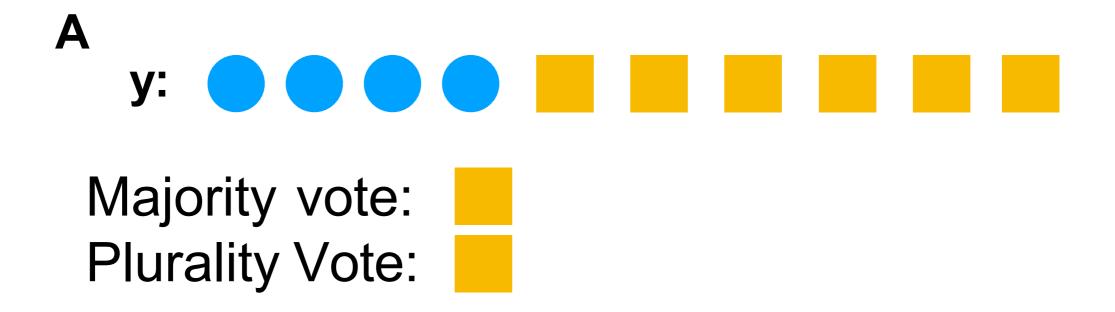
Feature Scaling

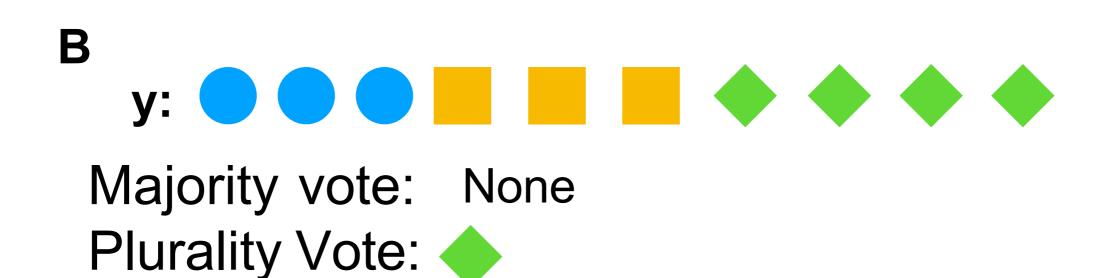


K-nearest neighbors

k-Nearest Neighbors







kNN for Classification

$$\mathcal{D}_k = \{\langle \mathbf{x}^{[1]}, f(\mathbf{x}^{[1]}) \rangle, \dots, \langle \mathbf{x}^{[k]}, f(\mathbf{x}^{[k]}) \rangle\} \qquad \mathcal{D}_k \subseteq \mathcal{D}$$

$$h(\mathbf{x}^{[q]}) = arg \max_{y \in \{1, \dots, t\}} \sum_{i=1}^{k} \delta(y, f(\mathbf{x}^{[i]}))$$
$$\delta(a, b) = \begin{cases} 1, & \text{if } a = b, \\ 0, & \text{if } a \neq b \end{cases}$$

$$h(\mathbf{x}^{[t]}) = \mathsf{mode}(\{f(\mathbf{x}^{[1]}), ..., f(\mathbf{x}^{[k]})\})$$

kNN for Regression

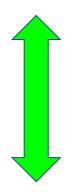
$$\mathcal{D}_k = \{\langle \mathbf{x}^{[1]}, f(\mathbf{x}^{[1]}) \rangle, \dots, \langle \mathbf{x}^{[k]}, f(\mathbf{x}^{[k]}) \rangle\} \qquad \mathcal{D}_k \subseteq \mathcal{D}$$

$$h(\mathbf{x}^{[t]}) = \frac{1}{k} \sum_{i=1}^{k} f(\mathbf{x}^{[i]})$$

Finding the right k value

k too low

"Overfit"



k too high

"Overgeneralize"

The "k" of k Nearest Neighbors is a "hyperparameter".

Almost all machine learning algorithms have hyperparameters.

- k too low \rightarrow "Overfitting" \rightarrow Decisions based on noise
- k too high \rightarrow "Underfitting" \rightarrow Decisions not sensitive to important subgroups in the data set

We'll cover exactly how to pick the right k at the end, but let's understand how this concept applies across many machine learning algorithms.

Applications of Nearest Neighbor Methods



Saudi Computer Society, King Saud University

Applied Computing and Informatics

(http://computer.org.sa) www.ksu.edu.sa www.sciencedirect.com



ORIGINAL ARTICLE

Automated web usage data mining and recommendation system using K-Nearest Neighbor (KNN) classification method



D.A. Adeniyi, Z. Wei, Y. Yongquan *

The major problem of many on-line web sites is the presentation of many choices to the client at a time; this usually results to strenuous and time consuming task in finding the right product or information on the site. In this work, we present a study of automatic web usage data mining and recommendation system based on current user behavior through his/her click stream data on the newly developed Really Simple Syndication (RSS) reader website, in order to provide relevant information to the individual without explicitly asking for it. The K-Nearest-Neighbor (KNN) classification method has been trained to be used on-line and in Real-Time to identify clients/visitors click stream data, matching it to a particular user group and recommend a tailored browsing option that meet the need of the specific user at a particular time. [...]

Distance Metric Learning for Large Margin Nearest Neighbor Classification

Weinberger, Kilian Q., John Blitzer, and Lawrence K. Saul. "Distance metric learning for large margin nearest neighbor classification." *Advances in Neural Information Processing Systems*. 2006.

Kilian Q. Weinberger, John Blitzer and Lawrence K. Saul

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Among 3 nearest neighbors after but not before training:

Among 3 nearest neighbors before but not after training:

We show how to learn a Mahanalobis distance metric for knearest neighbor (kNN) classification by semidefinite programming. The metric is trained with the goal that the knearest neighbors always belong to the same class while examples from different classes are separated by a large margin. On seven data sets of varying size and difficulty, we find that metrics trained in this way lead to significant improvements in kNN classification—for example, achieving a test error rate of 1.3% on the MNIST handwritten digits. As in support vector machines (SVMs), the learning problem reduces to a convex optimization based on the hinge loss. Unlike learning in SVMs, however, our framework requires no modification or extension for problems in multiway (as opposed to bi- nary) classification.

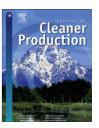
Test Image: 0 1 1 2 2 3 3 4 4 5 5 6 6 7 1 8 8 9 9 9 1 Nearest neighbor after training: 0 2 2 1 0 8 3 7 1 6 6 0 7 9 1 3 5 9 1



Contents lists available at ScienceDirect

Journal of Cleaner Production

journal homepage: www.elsevier.com/locate/jclepro

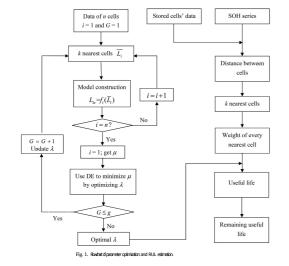


Remaining useful life estimation of lithium-ion α s based on k- nearest neighbor regression with differential evolution optimization



Yapeng Zhou^a, Miaohua Huang^{a,*}, Michael Pecht^b

- ^a Hubei Key Laboratory of Advanced Technology for Automotive Components, Wuhan University of Technology, Wuhan, 430070, PR China
- ^b Center for Advanced Life Cycle Engineering, University of Maryland, College Park, MD, 20742, USA





elatifule material contains LIPFs, EC, and DEC, and the rated voltage is 3.7 V. The cathode and anode lavers of groups A and B are wrapped around orthogonal rotation

syles of groups A and is are wrappen airound orangems rotation orestant. Voltage product and discharged with constant current and constant. Voltage product and discharged with constant current to 2.7 V under 2.4 °C. The detailed specifications and draughfischarge method free set are from in Tuble 1. As shown in Tuble 1. As

- 1. Program the charge/discharge with Bits Pro software on compute
- Cornect the cast to the circuit, and put them into the thermal chamber.

 Turn on the thermal chamber and set the temperature at 2.4 °C. and set 1. b.
- Start the charge/discharge cycling with the Bits Pro software

calculated by integrating the discharge

Remaining useful life estimation is of great importance to customers who use battery-powered products. This paper develops a remaining useful life estimation model based on **k-nearest neighbor regression** by incorporating data from all the cells in a battery pack. A differential evolution technique is employed to optimize the parameters in the estimation model. In this approach, remaining useful life is estimated from a weighted average of the useful life of several nearest cells that share a similar degradation trend to the cell whose remaining useful life needs to be estimated. The developed method obtains a remaining useful life estimation result with average error of 9 cycles, and the best estimation only has an error of 2 cycles. [...]