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- 1. Introduction to k-nearest neighbors
- 2. Measuring distance
- 3. Making predictions
- 4. Standardization, scaling, and normalization

# Example Revisited

Last week we introduced the following toy example:



Our goal was to learn rules involving  $x_1$  and  $x_2$  that can accurately classify a *new observation* as healthy or unhealthy.

A simple rule is to classify each new data-point using its *nearest neighbor*, or the observation closest to it's  $x_2$  and  $x_1$  coordinates:



A new observation at  $\{x_1 = 8, x_2 = 4\}$  (purple) is classified as "healthy", while another new observation at  $\{x_1 = 8, x_2 = 8\}$  (blue) is classified as "unhealthy":



To implement this approach, we need to define how to determine the nearest neighbor:

$$d_{a,b} = \left(\sum_{j=1}^{K} |x_{a,j} - x_{b,j}|^p\right)^{1/p}$$

- Minkowski distance, d<sub>a,b</sub>, measures the distance between data-points a and b
  - The formula sums pairwise coordinate differences across K dimensions
  - The parameter p is chosen by the analyst

Two of the most popular choices are p = 2 and p = 1:

$$egin{aligned} d_{ ext{euclidean}} &= \sqrt{\sum\limits_{j=1}^{K} (x_{a,j} - x_{b,j})^2} \ d_{ ext{manhattan}} &= \sum\limits_{j=1}^{K} |x_{a,j} - x_{b,j}| \end{aligned}$$

When might these measures lead to different neighbors?

Differences in x<sub>a,j</sub> - x<sub>b,j</sub> increase Euclidean distance quadratically, but increase Manhattan distance linearly
This means large differences are further magnified by Euclidean distance

## Distance and the curse of dimensionality

The *curse of dimensionality* describes a multitude of problems that arise when working with many variables.

It's relatively easy to trap a rat inside a pipe (which can move in 1-dimension). It's harder to trap a dog running in a field (which can move in 2-dimensions). It's even harder to trap a bird flying in a field (which moves in 3-dimensions). It's impossible to trap a ghost...

What problems might arise when using Euclidean distance to find neighbors in a high-dimensional dataset?

 $\label{eq:analogy} {\tt Credit: https://stats.stackexchange.com/questions/169156/explain-curse-of-dimensionality-to-a-child and the state of the sta$ 

- In our earlier examples, we classified new observations using the single nearest neighbor - a high variance procedure
  - We can decrease variance (at the expense of introducing additional bias) by using multiple neighbors
  - The k-nearest neighbors algorithm aggregates the outcomes of the k nearest data-points to generate a prediction for a new observation

For a classification task, each of the k-nearest data-points contributes to the predicted class:



x1

There are two schemes by which neighbors can contribute to the predicted class:

- 1. Uniform weighting the predicted probability of each class equals the proportion of neighbors belonging to that class.
- 2. *Distance weighting* neighbors that are weighted by the inverse of their distance, allowing closer data-points to contribute more to the predicted class.

Under uniform weighting, we estimate P(blue = healthy) = 4/5, but under distance weighting we'd estimate P(blue = healthy) > 4/5since the unhealthy neighbor is also the furthest



- In our simple example, x<sub>1</sub> and x<sub>2</sub> have similar scales (ie: standard deviations of similar magnitude)
  - In practice, we'll frequently encounter features with different scales
  - Unless rescaled, features on larger scales will have more influence on distance calculations than features on smaller scales

### Scaling and standardization

A few popular ways to re-scale data are:

1. Standardization:

$$\widetilde{x}_i = rac{x_i - \operatorname{mean}(x)}{\operatorname{sd}(x)}$$

2. Robust scaling:

$$\tilde{x}_i = rac{x_i - \text{median}(x)}{\text{IQR}(x)}$$

3. Min-Max scaling:

$$\tilde{x}_i = \frac{x_i - \min(x)}{\max(x) - \min(x)}$$

4. Max-Absolute scaling:

$$\tilde{x}_i = \frac{x_i}{\max(|x|)}$$

- Standardization forces features to have a mean of zero and a standard deviation of one
  - Robust scaling forces features to have a median of zero, and it can be beneficial for data with large outliers
- Min-Max scaling maps each feature onto a [0,1] interval, which can have computational advantages
  - Max-Absolute scaling is similar to Min-Max scaling, but the output range is [-1,1] and it will *preserve exact zeros* (important for sparse data)

# Scaling vs. Normalization

Scaling changes the range of your data, it does not change the distributional shape:



However, the choice of scaler does matter for k-nearest neighbors, since dimensions are rescaled individually

## Normalization

If you'd like to change the distributional shape of your data to reduce the effects of skew/outliers, two strategies are:

- 1. Log-transformation simply taking the logarithm of each of the variable's values
- 2. Box-Cox transformation  $\tilde{X} = \frac{X^{\lambda}-1}{\lambda}$  for  $\lambda \neq 0$  and X > 0



So far, this presentation has focused exclusively on *classification tasks*, but *k*-nearest neighbors is easily applied to *regression tasks* with the following modifications:

- Outcomes are predicted by the arithmetic average (or distance weighted average) of the numeric outcomes for the identified neighbors
- Model performance is assessed using measures such as root mean squared error (RMSE) or mean absolute error (MAE)

To apply the k-nearest neighbors machine learning algorithm the following decisions must be made:

- 1. Whether/how the data should be standardized/scaled and/or normalized
- 2. How to measure distance
- 3. The number of neighbors to be used
- 4. Whether to use uniform or distance weighting

Today's lab will cover Python implementations of each of these; our next lab will cover data-driven methods for making these decisions.