# Introduction to supervised learning, training vs. validation, and *k*-nearest neighbors

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#### Overview

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  - Methods like scree plots of silhouette scores provide some objective guidance, but it's difficult to judge one analysis to be quantitatively better than another



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- Applications of *unsupervised learning* to tend to be open-ended and prone to all sorts of subjective choices
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▶ In contrast, *supervised learning* tends to be highly objective

- We have a predetermined outcome that we're aiming to predict, which allows us to quantify how accurate our predictions are
- We'll focus on two types of supervised learning
  - Classification predicting the class or category of a data-point
  - Regression predicting a numerical characteristic of a data-point



#### Supervised learning framework

Consider data consisting of an *n*-dimensional vector of outcomes, y, an *n* by *p* matrix of features, X

Further, suppose the true relationship between y and X is given by the following equation:

$$\mathbf{y} = f(\mathbf{X}) + \epsilon$$

- The function f() determines how the features in X influence y
  - $\epsilon$  is an *n*-dimensional vector of errors
- ln this setting, we aim to accurately approximate f()
  - If  $\epsilon = \mathbf{0}$ , we may be able to perfectly approximately f()
  - However,  $\epsilon \neq \mathbf{0}$  introduces *irreducible error*



#### Example

Consider two predictors,  $X_1$  and  $Y_2$ , and a binary outcome Y of "healthy" or "unhealthy". Can these predictors be used to accurately *classify* an observation?



x1



## Example (cont)

In this example, f() is shown below (blue ellipse):



We can see that it's possible to use the data to learn a good approximation of f().



## Example (cont.)

As a human, you might observe that healthy data-points tend to fall between 2 and 8 in both  $x_1$  and  $x_2$ , so you might propose the following *classification model*:



This simple model correctly classifies 178 of 200 data-points.



## Example (irreducible error)

Let's revisit the true relationship between  $X_1$ ,  $X_2$ , and Y. Notice how some "healthy" datapoints are outside the ellipse, and some "unhealthy" ones are inside it:



- The misclassification of these data-points reflects this scenario's irreducible error (sometimes called "Bayes error")
  Even the best passible approximation of f() cannot perfectly.
  - Even the best possible approximation of f() cannot perfectly classify eveny data-point



#### Irreducible error in other contexts

Is a digit a "5" or something else?



How might the concept of irreducible error manifest in this application?



#### Irreducible error in other contexts

We could know the exact "rules" used to make a "5", but it's possible we encounter examples of "5" that look more like a "6".



Even state-of-the-art classifiers (which approach the irreducible error rate) incorrectly classify  $\sim 0.5\%$  of handwritten digits (source)



#### Reducible Error

Achieving the best approximation of f() amounts to minimizing *reducible error*. Consider the following classifier:



ced the error rate to zero?



- We aren't really interested in the error rate for observed examples
  - Instead, we'd like to minimize reducible error on new examples that our model hasn't seen yet



- We aren't really interested in the error rate for observed examples
  - Instead, we'd like to minimize reducible error on new examples that our model hasn't seen yet
- Standard protocol is to split the available data into training and testing sets
  - The training set is used to *learn* a collection of rules
  - The testing set is used to validate how well these rules perform on unseen data



#### Training, Testing, and Error

- Consider a hypothetical example with an irreducible error of "20 units"
  - Training error can be reduced by increasing model complexity (ie: learning more rules)
  - Test error is bounded in probability by the irreducible error



Model Complexity



Reducible error can arise in one of two ways: bias or variance

- Bias is when a learner lacks the structural flexibility to detect aspects of the true relationship between the predictors and the outcome
- Variance is when a learner is overly sensitive to chance artifacts present in the data (ie: the manifestations of irreducible error)

Poor performance due to high bias is called *underfitting*, while poor performance due to high variance is called *overfitting* 



How would you compare the bias and variance of the following models (a rectangle vs. an n-dimensional polygon)?





## Quantifying error

- Our toy example used a binary categorical outcome, a scenario where *classification accuracy* provides a natural way to understand error
  - Later this week we'll consider other ways to measure the error of a classification model
- For a numeric outcome, it's most natural to measure error by summarizing the distances between predicted and observed outcomes:
  - Root Mean Squared Error:  $RMSE = \sqrt{\frac{1}{n}\sum_{i=1}^{n}(y_i \hat{y}_i)^2}$
  - Mean Absolute Error:  $MAE = \frac{1}{n} \sum_{i=1}^{n} |y_i \hat{y}_i|$
- Where  $\hat{y}_i$  our model's prediction for the  $i^{th}$  data-point



#### k-nearest neighbors

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:



Statistics

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

$$d(\mathbf{x}_a, \mathbf{x}_b) = \left(\sum_{j=1}^p |x_{a,j} - x_{b,j}|^q\right)^{1/q}$$

- 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:

$$d_{\text{euclidean}} = \sqrt{\sum_{j=1}^{p} (x_{a,j} - x_{b,j})^2}$$
$$d_{\text{manhattan}} = \sum_{j=1}^{p} |x_{a,j} - x_{b,j}|$$

When might these measures lead to different neighbors?



There are two schemes by which neighbors can contribute to a prediction:

- 1. Uniform weighting each neighbor contributes equally
- 2. *Distance weighting* neighbors are weighted by the inverse of their distance to the new data-point, allowing closer neighbors to contribute more strongly

For regression tasks, the prediction is either a simple or weighted average. For classification tasks, the predicted probabilities are given by either simple or weighted voting.



## Other hyperparameters (number of neighbors)

Notice the gap in performance on the test vs. training data:





### Scaling and standardization

Because KNN relies upon distance calculations, rescaling the predictors is important:

1. Standardization:

$$x_i^* = \frac{x_i - \mathrm{mean}(x)}{\mathrm{sd}(x)}$$

2. Robust scaling:

$$x_i^* = \frac{x_i - \text{median}(x)}{\text{IQR}(x)}$$

3. Min-Max scaling:

$$x_i^* = \frac{x_i - \min(x)}{\max(x) - \min(x)}$$

4. Max-Absolute scaling:

$$x_i^* = \frac{x_i}{\max(|x|)}$$



- 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  $x_i^* = \frac{x_i^{\lambda} 1}{\lambda}$  for  $\lambda \neq 0$  and X > 0





*k*-nearest neighbors is a simple model that is useful in illustrating the basic workflow of machine learning:

- 1. Split the data into training and validation sets
- 2. Set up a data preparation pipeline (ie: rescaling, normalization, dimension reduction, etc.)
- 3. Determine hyperparamters (ie: weighting, number of neighbors, etc.)
- 4. Evaluate the final model on the validation set

This week's lab will cover each of these steps in greater detail

