Data Science

Lecture 2-2: Data Science Fundamentals (Modeling)



Lecturer: Yen-Chia Hsu

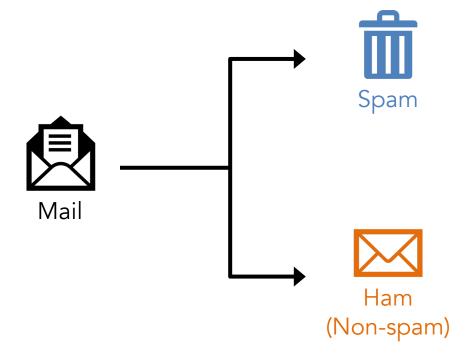
Date: Feb 2025

This lecture recaps classification and regression techniques for modeling data.

For this lecture, let us now use the following text classification task as

an example: identifying whether a text message is spam or ham (non-spam).

Hi Yen-Chia, may we have our meeting on 5/15 by just email update to buy some time? if not, zero worries if you need to talk.



To classify spam messages, we need examples: a dataset with

observations (messages) and labels (spam or non-spam).



Hi Yen-Chia, may we have our meeting on 5/15 by just email update to buy some time? if not, zero worries if you need to talk.



Ham

Would you be willing to meet with me on 3/26 Thursday when I was in TU Delft after (or before) giving the guest lecture (10:35am-11:50am)?



Ham



: Observations

We can extract features (information) using human knowledge, which

can help distinguish spam and ham messages.





Number of special characters = 34 Number of digits = 22

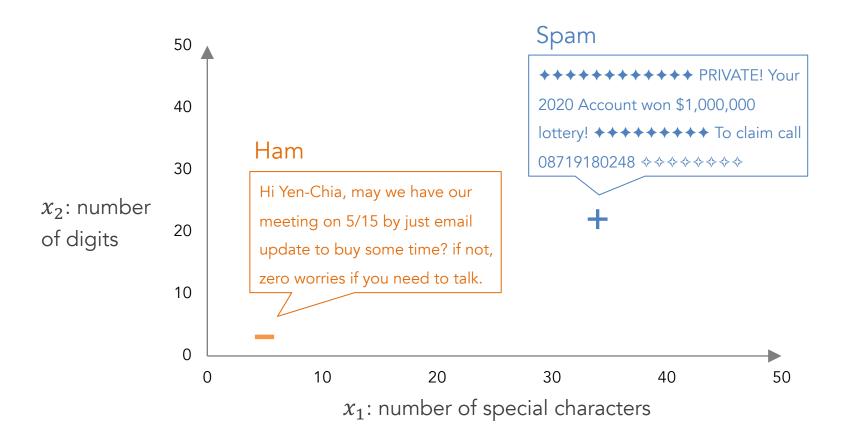
Hi Yen-Chia, may we have our meeting on 5/15 by just email update to buy some time? if not, zero worries if you need to talk.



Number of special characters = 5 Number of digits = 3

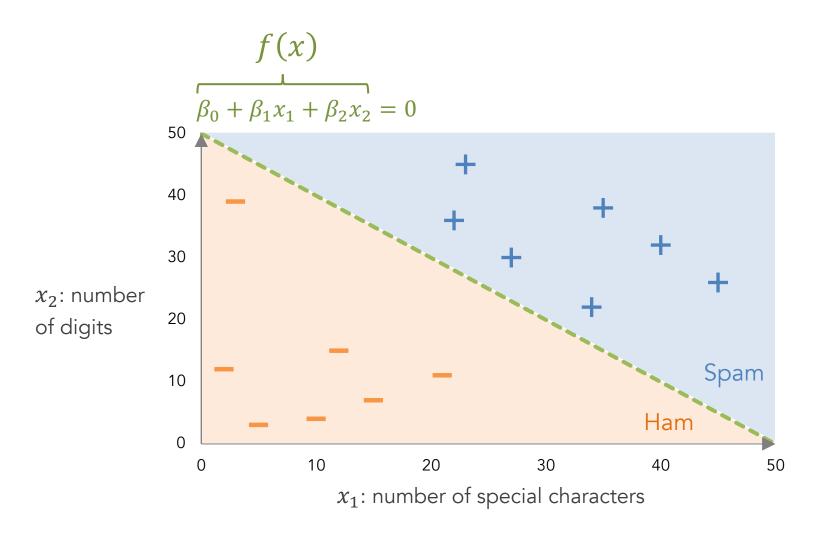
Using features x (which contains x_1 and x_2), we can represent each

message as one data point on an p-dimensional space (p = 2 in this case).

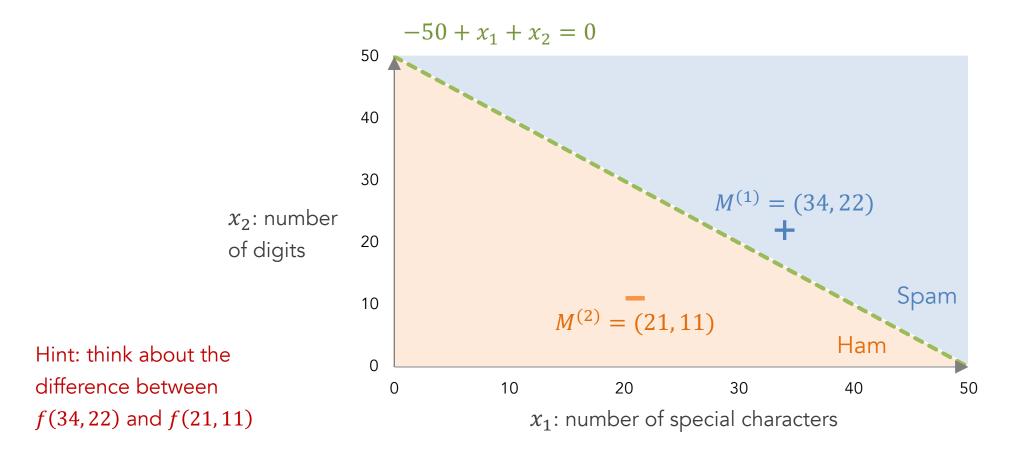


We can think of the model as a function f that can separate the

observations into groups (i.e., class labels y) according to their features $x = \{x_1, x_2\}$.

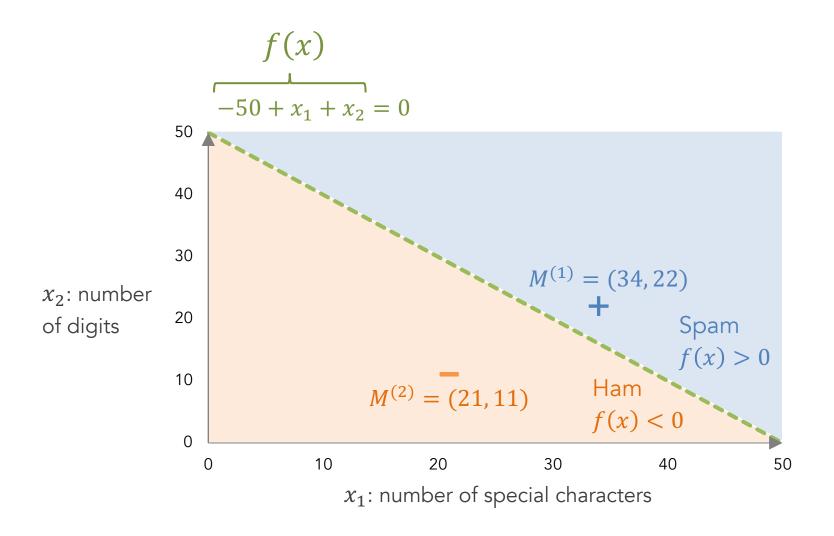


Exercise 2.1: Given a classifier $f(x) = -50 + x_1 + x_2$ and two messages $M^{(1)}$ and $M^{(2)}$, explain how the model classifies the message as spam or ham mathematically. $M^{(1)}$ has 34 special characters and 22 digits. $M^{(2)}$ has 21 special characters and 11 digits.



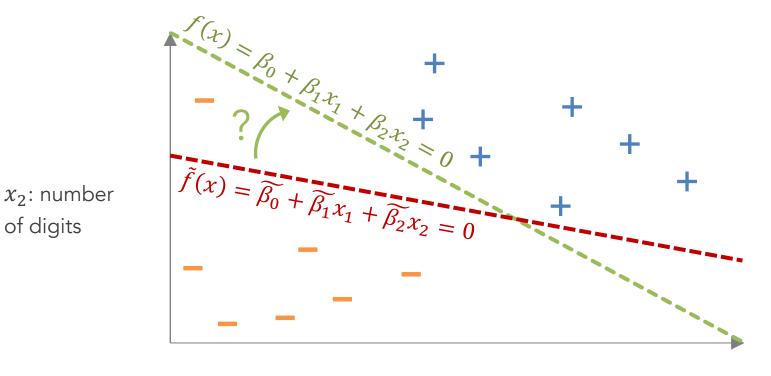
We can plug the features $x = \{x_1, x_2\}$ into the classifier equation f(x)

to determine if it is spam or ham by checking if f(x) is larger or smaller than zero.



of digits

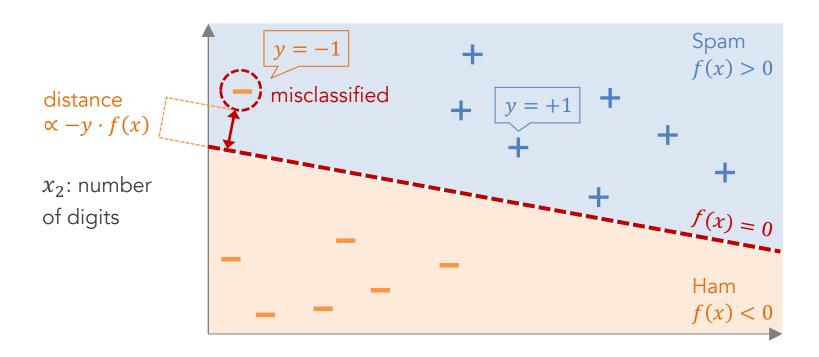
To find a good function f, we start from some f and train it until satisfied. We need something to tell us which direction and magnitude to update.



 x_1 : number of special characters

First, we need an error metric (i.e., cost or objective function). For example, we can use the sum of distances between the misclassified points and line f.

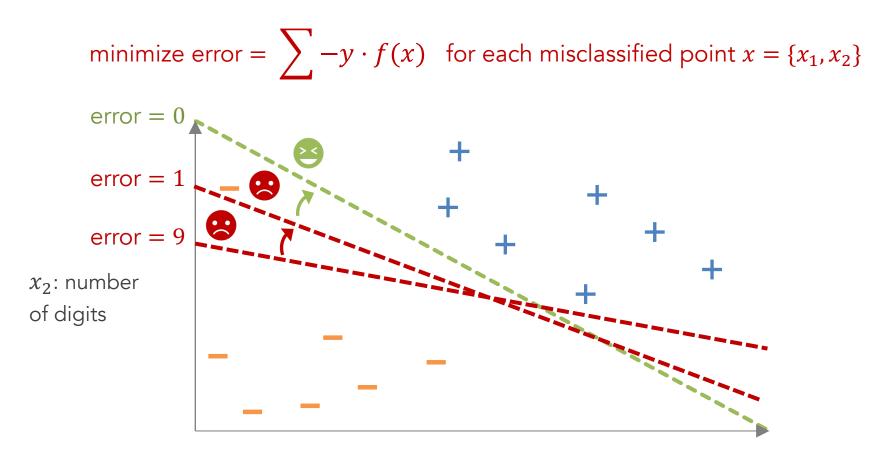
error =
$$\sum -y \cdot f(x)$$
 for each misclassified point $x = \{x_1, x_2\}$



 x_1 : number of special characters

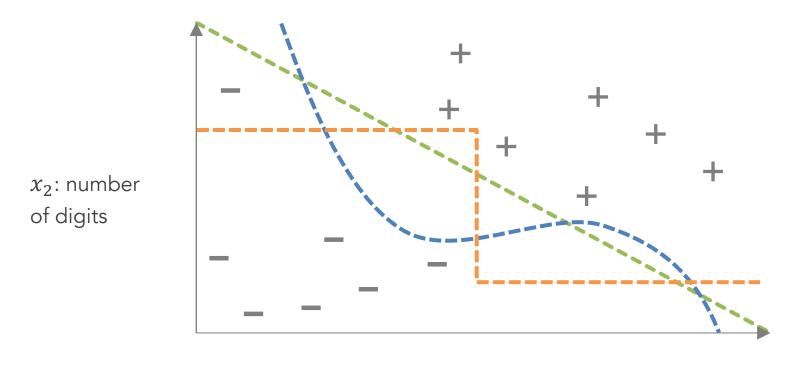
We can use gradient descent (an optimization algorithm) to minimize

the error to train the model f iteratively. This example is the Perceptron algorithm.



 x_1 : number of special characters

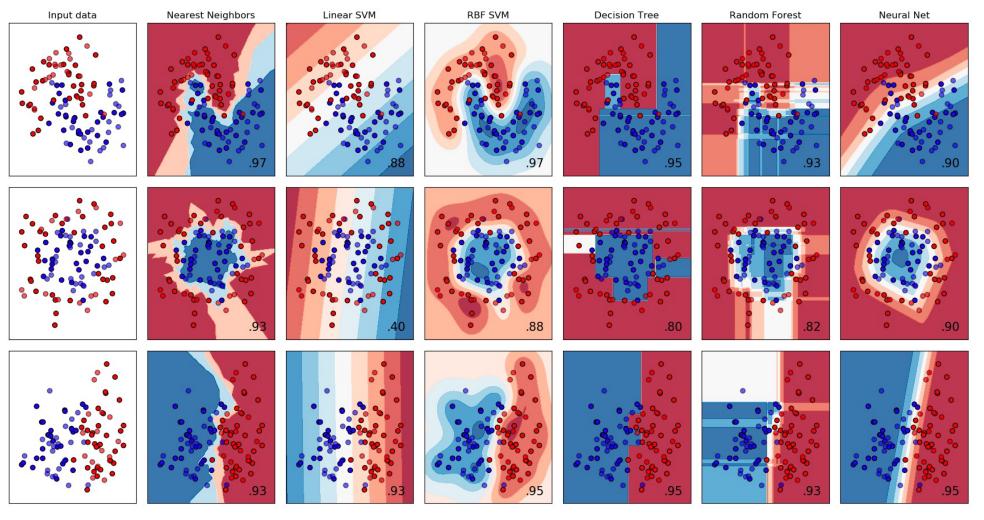
Depending on the needs, we can train different models (using different loss functions) with various shapes of decision boundaries.



 x_1 : number of special characters

Depending on the needs, we can train different models (using

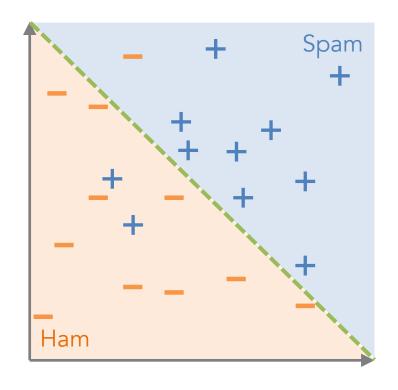
different loss functions) with various shapes of decision boundaries.



Retrieved from https://scikit-learn.org/stable/auto-examples/#classification

To evaluate our classification model, we need to compute evaluation

metrics to measure and quantify model performance, such as the accuracy of all data.



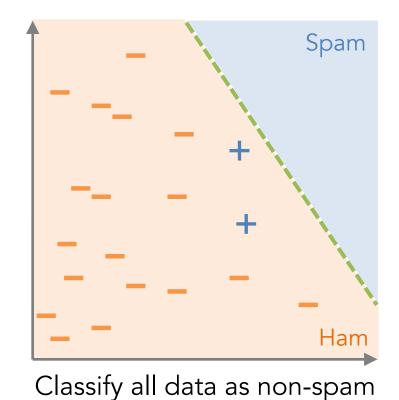
Accuracy for all data

$$= \frac{\text{# of correctly classified points}}{\text{# of all points}}$$

$$=\frac{19}{22}=0.86$$

But what if the dataset is imbalanced (i.e., some classes have far less

data)? In this case, the accuracy of all data is a bad evaluation metric.



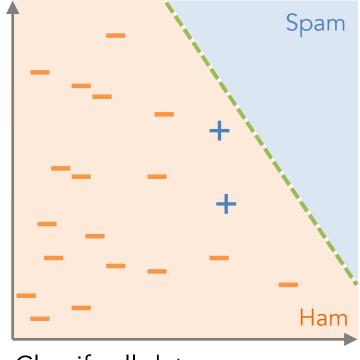
Accuracy for all data

$$= \frac{\text{# of correctly classified points}}{\text{# of all points}}$$

$$=\frac{18}{20}=0.9$$

Instead of computing the accuracy for all the data, we can compute

accuracy for each class, which allows us to see the performance of different labels.



Classify all data as non-spam

Accuracy for spam
$$=\frac{0}{2}=0$$

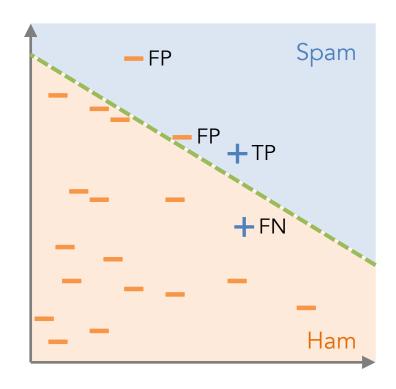
(true positive rate, recall, sensitivity)

Accuracy for ham
$$=\frac{18}{18}=1$$

(true negative rate, specificity)

If we care more about the positive class (e.g., spam), we can use

precision and recall, with its best value at 1 and the worst value at 0.



$$TP = 1$$
 (True Positive)

$$FP = 2$$
 (False Positive)

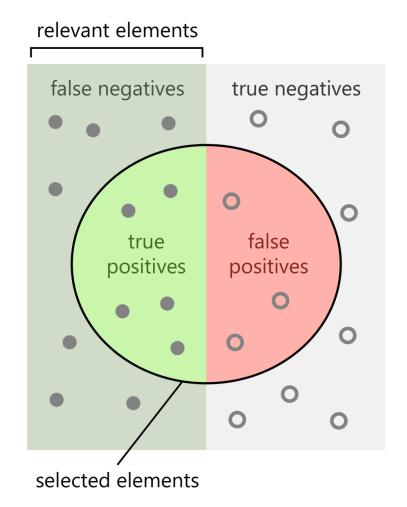
$$FN = 1$$
 (False Negative)

$$Precision = \frac{TP}{TP + FP} = 0.33$$

$$Recall = \frac{TP}{TP + FN} = 0.5$$

Precision and recall can be aggregated into F-score as a general

model performance, with its best value at 1 and worst value at 0.



How many selected items are relevant?

How many relevant items are selected?

$$F-score = 2 \cdot \frac{Precision \cdot Recall}{Precision + Recall}$$

Exercise 2.2: Suppose that we fit a binary classification model in identifying spam and ham (i.e., non-spam). Spam is the positive label, and ham is the negative label.

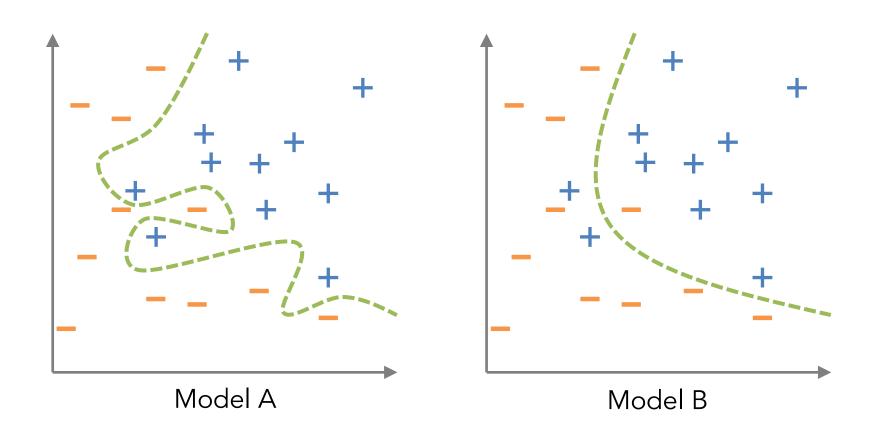
- 40 samples are predicted as spam, and they are indeed spam in reality
- 20 samples are predicted as spam, but it turns out that they are not spam in reality
- 60 samples are predicted as ham, but it turns out that they are spam in reality
- 80 samples are predicted as ham, and they are indeed ham in reality

What are the precision, recall, and f-score (F) of the model?

$$Precision = \frac{TP}{TP + FP} \qquad \qquad Recall = \frac{TP}{TP + FN} \qquad \qquad F = 2 \cdot \frac{Precision \cdot Recall}{Precisio + Recall}$$

We can train different types of models. But how do we know which

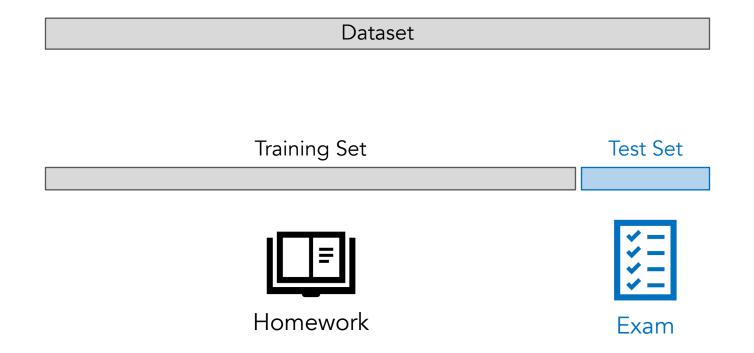
one is better? Can we just pick an evaluation metric to determine which model is good?

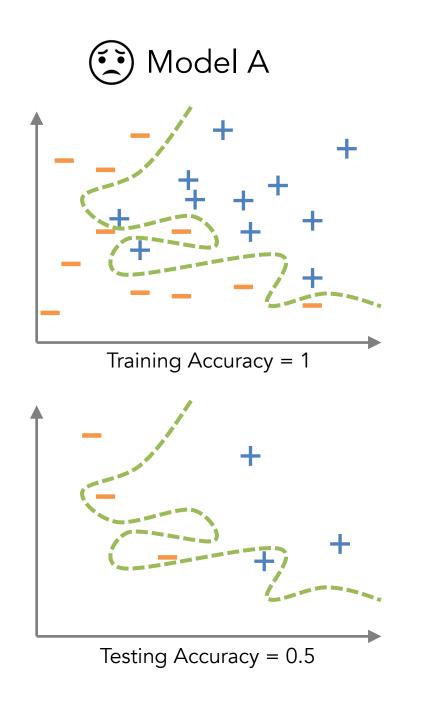


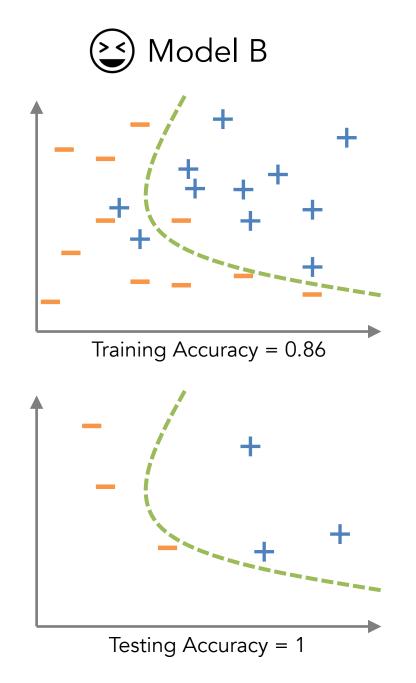
Model Training

To choose models, we need a test set, which contains data that the

models have not yet seen before during the training phase.



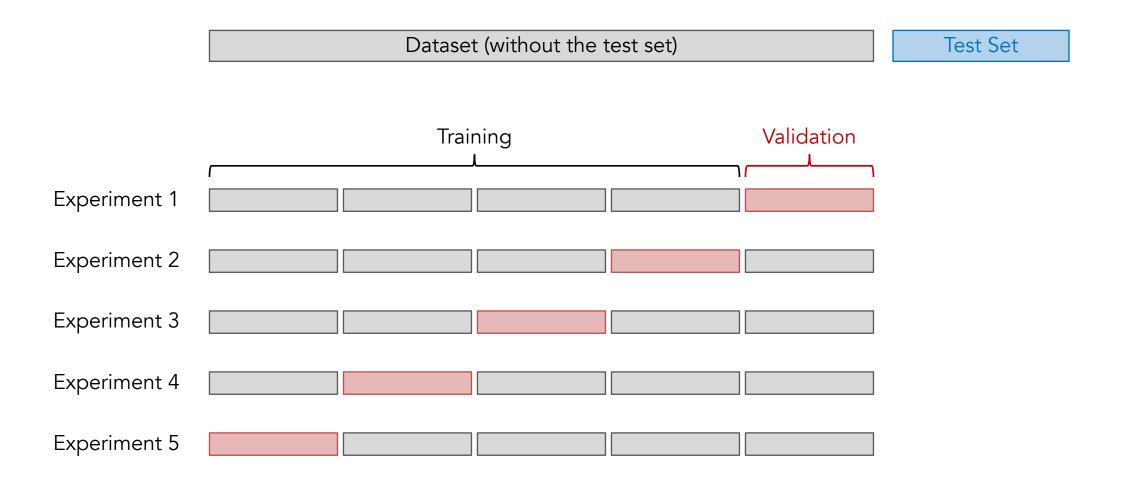




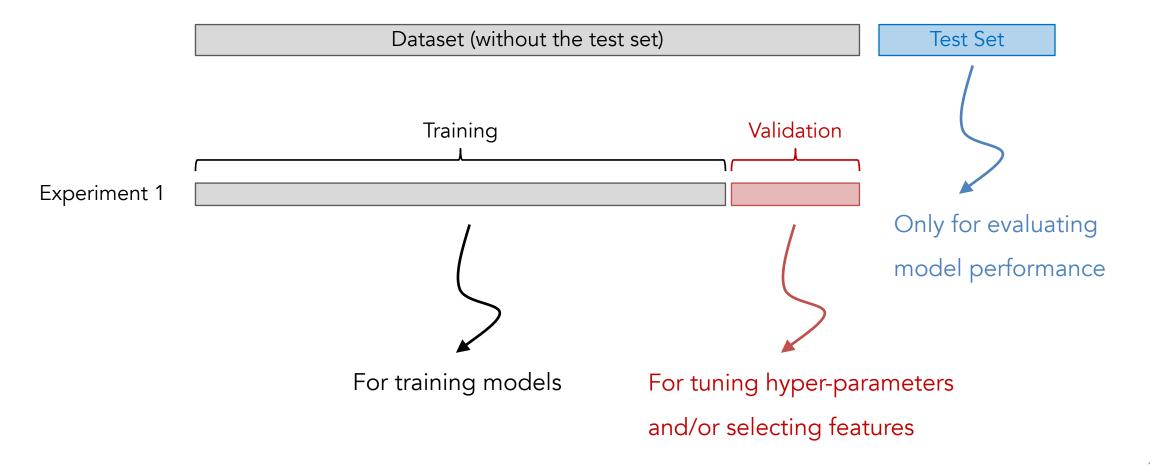
Model Training

To tune hyper-parameters or select features for a model, we use

cross-validation to divide the dataset into folds and use each fold for validation.



Model Training You should not use the test set to tune hyper-parameters or select features, which will lead to information leakage. The test set is used to do an unbiased check of generalization performance after all modeling decisions are made.



Model Training

One way to select features is to recursively eliminate the less

important ones by using metrics like permutation importance (which means permuting a feature several times and measuring the decrease in model performance).

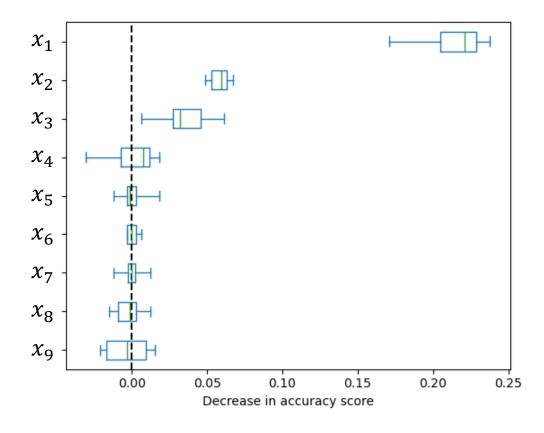


Figure source -- https://scikit-learn.org/1.5/modules/permutation_importance.html

If two highly correlated features exist, the model can access the **Model Training** information from the non-permuted feature. Thus, it may appear that both features are not important (which can be false). A better way is to cluster the correlated features first.

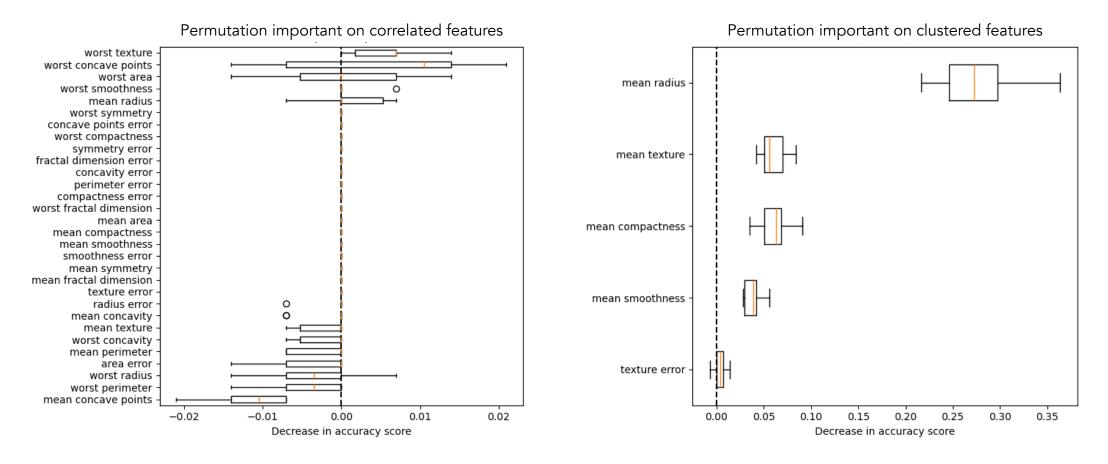
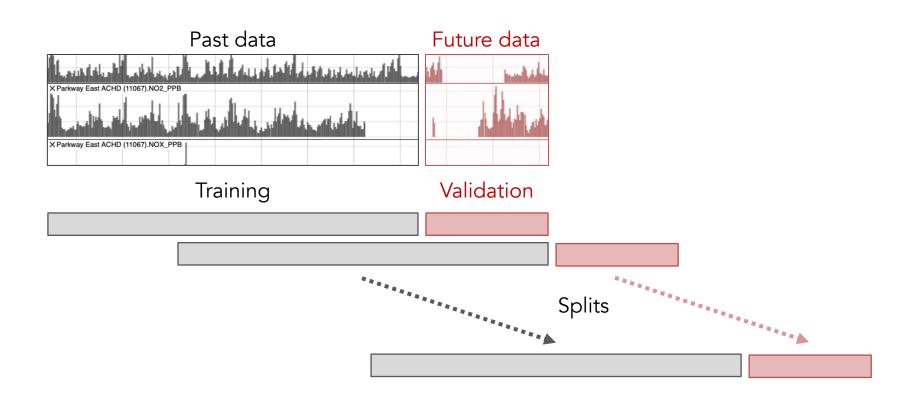


Figure source -- https://scikit-learn.org/1.5/modules/permutation_importance.html

Model Training

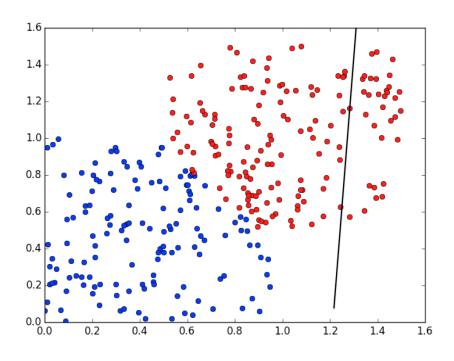
For time-series data, it is better to do the split for cross-validation

based on the order of time intervals, which means we only use data in the past to predict the future, but not the other way around.

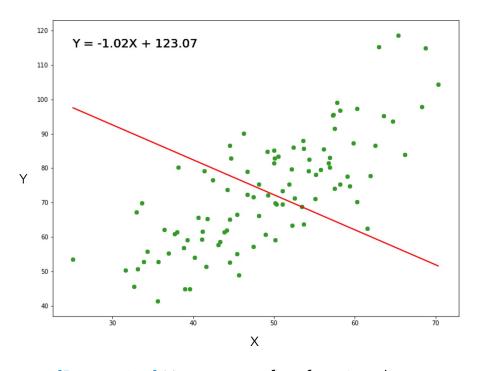


Unlike classification (which separates data into categories), regression

fits a function that maps features x to a continuous variable y (i.e., the response).

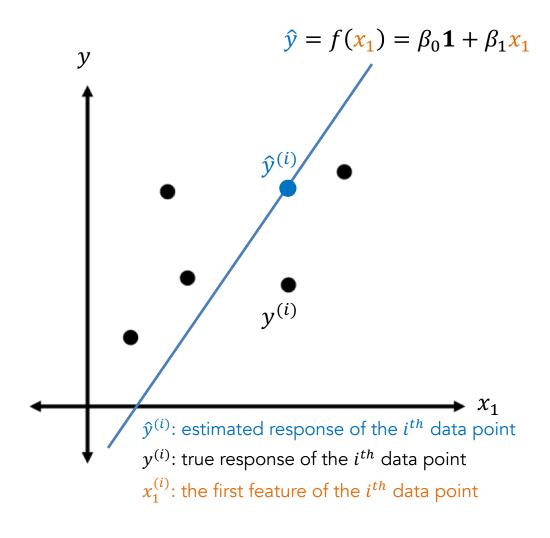


• [Classification] How can we fit a function that separates data points into different groups?



 [Regression] How can we fit a function that maps features (input) to a continuous variable (output)? Linear regression fits a linear function f that maps x_1 (e.g., the first

feature vector of something) to y, which can best describe their linear relationship.



y: true response

 \hat{y} : estimated response

 x_1 : predictor/feature

 β_0 and β_1 : intercept and slope

$$\begin{bmatrix} \hat{y}^{(1)} = f\left(x_1^{(1)}\right) = \beta_0 \cdot 1 + \beta_1 \cdot x_1^{(1)} \\ \vdots \\ \hat{y}^{(n)} = f\left(x_1^{(n)}\right) = \beta_0 \cdot 1 + \beta_1 \cdot x_1^{(n)} \end{bmatrix}$$

$$\begin{bmatrix} \hat{y}^{(1)} \\ \vdots \\ \hat{y}^{(n)} \end{bmatrix} = f \begin{pmatrix} \begin{bmatrix} x_1^{(1)} \\ \vdots \\ x_1^{(n)} \end{bmatrix} \end{pmatrix} = \beta_0 \begin{bmatrix} 1 \\ \vdots \\ 1 \end{bmatrix} + \beta_1 \begin{bmatrix} x_1^{(1)} \\ \vdots \\ x_1^{(n)} \end{bmatrix}$$

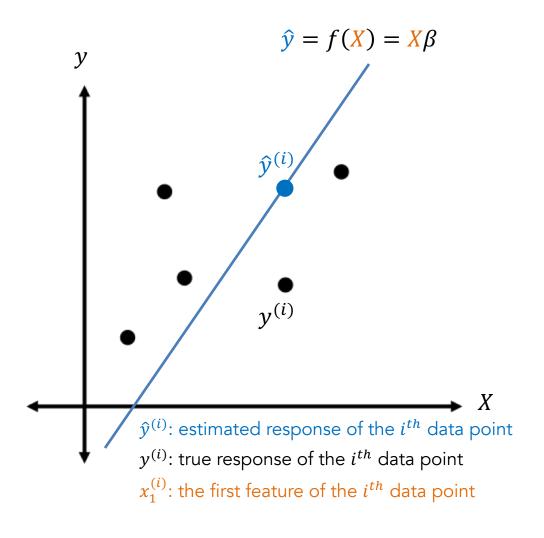
$$\hat{y}$$

$$\mathbf{1}$$

$$x_1$$

We can now create a feature matrix X that includes the intercept term

 β_0 , which gives us a compact form of equation.



y: true response (vector)

 \hat{y} : estimated response (vector)

X: predictor/feature (matrix)

 β : coefficients (vector)

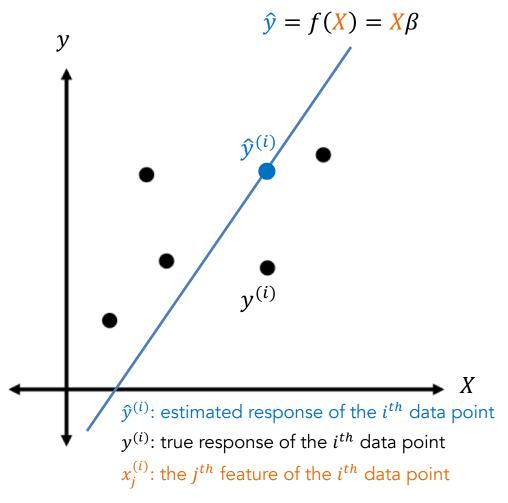
$$\begin{bmatrix} \hat{y}^{(1)} \\ \vdots \\ \hat{y}^{(n)} \end{bmatrix} = \beta_0 \begin{bmatrix} 1 \\ \vdots \\ 1 \end{bmatrix} + \beta_1 \begin{bmatrix} x_1^{(1)} \\ \vdots \\ x_1^{(n)} \end{bmatrix}$$

$$\hat{\mathbf{y}} = \beta_0 \mathbf{1} + \beta_1 x_1 = \begin{bmatrix} 1 & x_1^{(1)} \\ \vdots & \vdots \\ 1 & x_1^{(n)} \end{bmatrix} \times \begin{bmatrix} \beta_0 \\ \beta_1 \end{bmatrix}$$

$$X \qquad \beta$$

We can now generalize linear regression to have multiple predictors

(i.e., multiple linear regression) and keep the compact mathematical representation.



y: true response (vector)

 \hat{y} : estimated response (vector)

X: predictor/feature (matrix)

 β : coefficients (vector)

$$\begin{bmatrix} \hat{y}^{(1)} \\ \vdots \\ \hat{y}^{(n)} \end{bmatrix} = \beta_0 \begin{bmatrix} 1 \\ \vdots \\ 1 \end{bmatrix} + \beta_1 \begin{bmatrix} x_1^{(1)} \\ \vdots \\ x_1^{(n)} \end{bmatrix} + \dots + \beta_p \begin{bmatrix} x_p^{(1)} \\ \vdots \\ x_p^{(n)} \end{bmatrix}$$

$$\hat{y} = f(X) = \begin{bmatrix} 1 & x_1^{(1)} & \dots & x_p^{(1)} \\ \vdots & \vdots & \dots & \vdots \\ 1 & x_1^{(n)} & \dots & x_p^{(n)} \end{bmatrix} \times \begin{bmatrix} \beta_0 \\ \beta_1 \\ \vdots \\ \beta_p \end{bmatrix}$$

$$X \qquad \beta$$

We use the vector and matrix forms to simplify equations.

Vector Matrix Scalar
$$X = \begin{bmatrix} \mathbf{1} & x_1 & \cdots & x_p \end{bmatrix} = \begin{bmatrix} 1 & x_1^{(1)} & \cdots & x_p^{(1)} \\ \vdots & \vdots & \vdots & \vdots \\ 1 & x_1^{(n)} & \cdots & x_p^{(n)} \end{bmatrix}$$

$$\beta = \begin{bmatrix} \beta_0 \\ \beta_1 \\ \vdots \\ \beta_p \end{bmatrix}$$

$$\hat{y} = f(x) = X\beta = \begin{bmatrix} \mathbf{1} & x_1 & \cdots & x_p \end{bmatrix} \times \begin{bmatrix} \beta_0 \\ \beta_1 \\ \vdots \\ \beta_p \end{bmatrix} = \beta_0 + \beta_1 x_1 + \cdots + \beta_p x_p = \begin{bmatrix} \hat{y}^{(1)} \\ \vdots \\ \hat{y}^{(n)} \end{bmatrix}$$

Vector

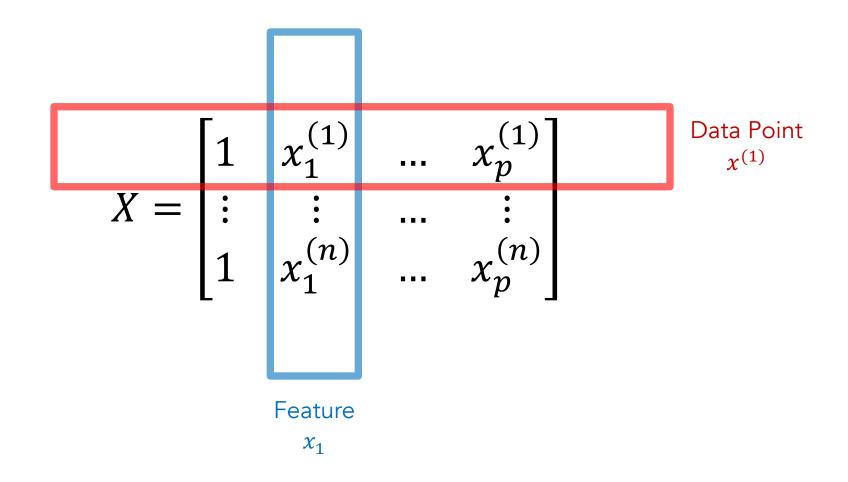
We can map vector and matrix forms to data directly.

Sum of smell ratings		H2S in PPM	SO2 in PPM	•••	Wind direction in DEG	Wind speed in MPH
	25	0.019	0.020		215.0	3,2
	40	0.130	0.033	•••	199.0	3.4
	45	0.095	0.044		184.0	2.9
Vector $\begin{bmatrix} \hat{y}^{(1)} \end{bmatrix}$		Vector x	1			
	$\left[egin{array}{c} dots \ \hat{y}^{(n)} \end{array} ight]$	Matrix $\begin{bmatrix} x_1^{(1)} & \cdots & x_p^{(1)} \\ \vdots & \vdots & \vdots \end{bmatrix}$				
				$x_1^{(n)}$	$x_p^{(n)}$	

Regression

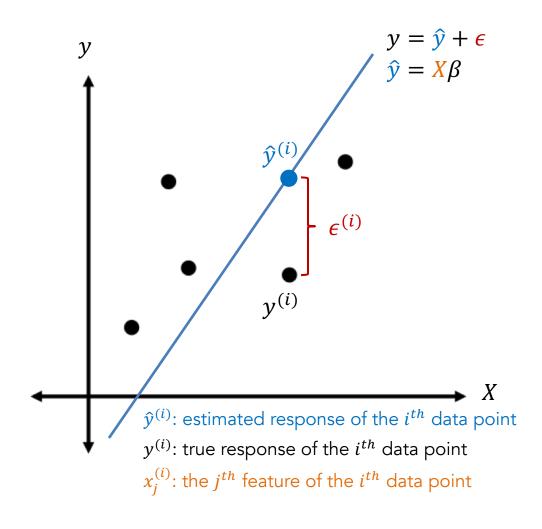
We can look at the feature matrix *X* from two different directions: one

represents features, and the other one represents data points.



Regression

Finally, we need an error metric between the estimated response \hat{y} and the true response y to know if the model fits the data well.



y: true response (vector)

 \hat{y} : estimated response (vector)

X: predictor/feature (matrix)

 β : coefficients (vector)

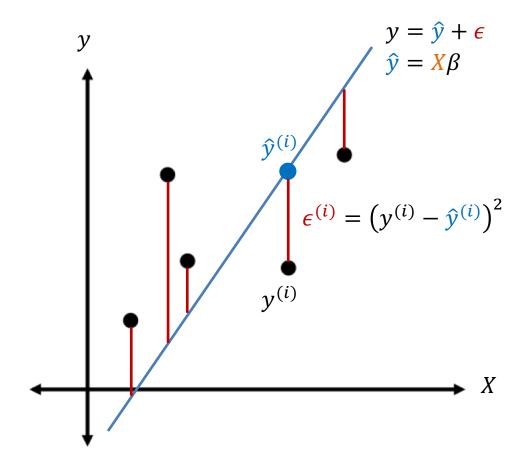
 ϵ : error/noise/residual (vector)

$$\hat{y} = f(X) = \begin{bmatrix} 1 & x_1^{(1)} & \dots & x_p^{(1)} \\ \vdots & \vdots & \dots & \vdots \\ 1 & x_1^{(n)} & \dots & x_p^{(n)} \end{bmatrix} \times \begin{bmatrix} \beta_0 \\ \beta_1 \\ \vdots \\ \beta_p \end{bmatrix}$$

$$X \qquad \beta$$

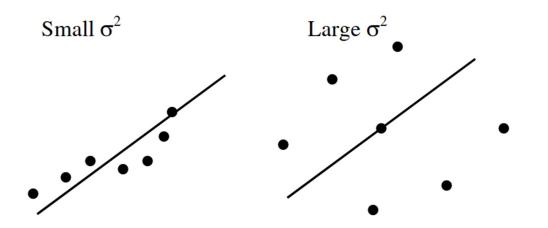
Regression

Usually, we assume that the error ϵ is IID (independent and identically distributed) and follows a normal distribution with zero mean and some variance σ^2 .



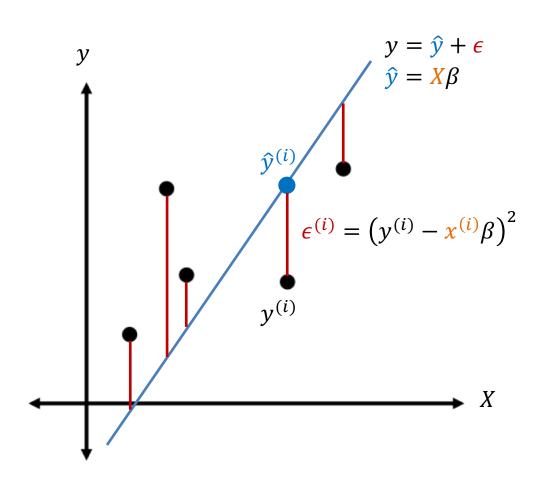
$$\epsilon \sim^{iid} N(0, \sigma^2)$$

total errors =
$$\sum_{i=1}^{n} \epsilon = \sum_{i=1}^{n} (y^{(i)} - \hat{y}^{(i)})^2$$

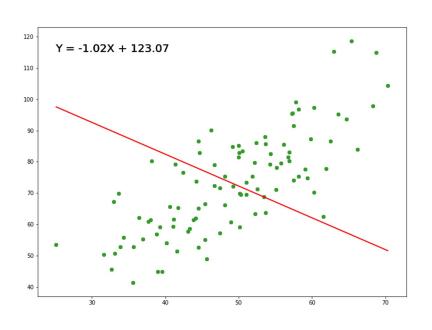


To find the optimal coefficient β , we need to minimize the error (the

sum of squared errors) using gradient descent or taking the derivative of its matrix form.

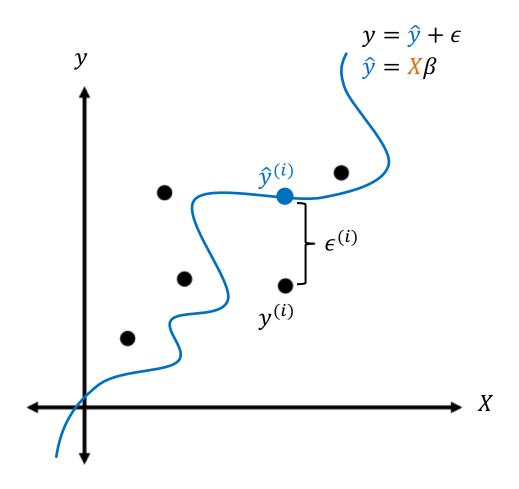


$$\min_{\beta} \sum_{i=1}^{n} \epsilon^{(i)} = \min_{\beta} \sum_{i=1}^{n} (y^{(i)} - x^{(i)}\beta)^{2}$$
$$= \min_{\beta} (y - X\beta)^{T} (y - X\beta)$$



We can model a non-linear relationship using polynomial functions

with degree k. The example below uses one predictor x_1 .



y: true response (vector)

 \hat{y} : estimated response (vector)

X: predictor/feature (matrix)

 β : coefficients (vector)

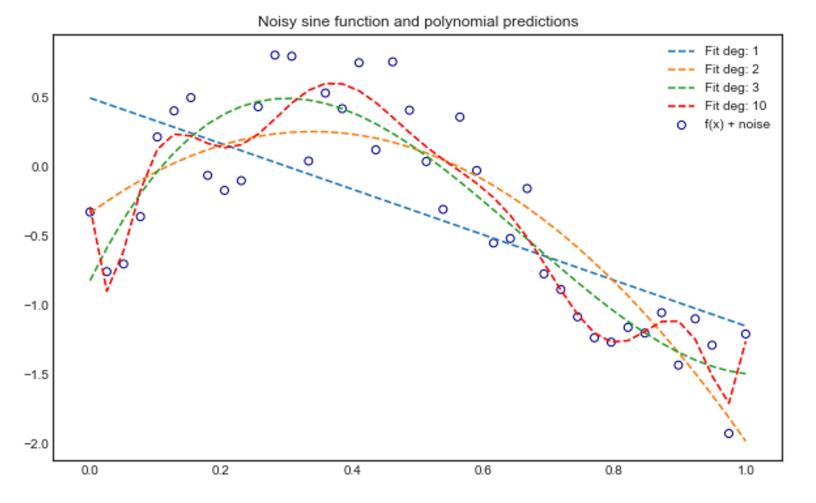
$$X = [\mathbf{1} \quad x_1 \quad (x_1)^2 \quad \cdots \quad (x_1)^k]$$

$$\beta = \begin{bmatrix} \beta_0 \\ \beta_1 \\ \vdots \\ \beta_k \end{bmatrix}$$

$$\hat{y} = X\beta = \beta_0 + \beta_1 x_1 + \beta_2 (x_1)^2 + \dots + \beta_k (x_1)^k$$

Regression

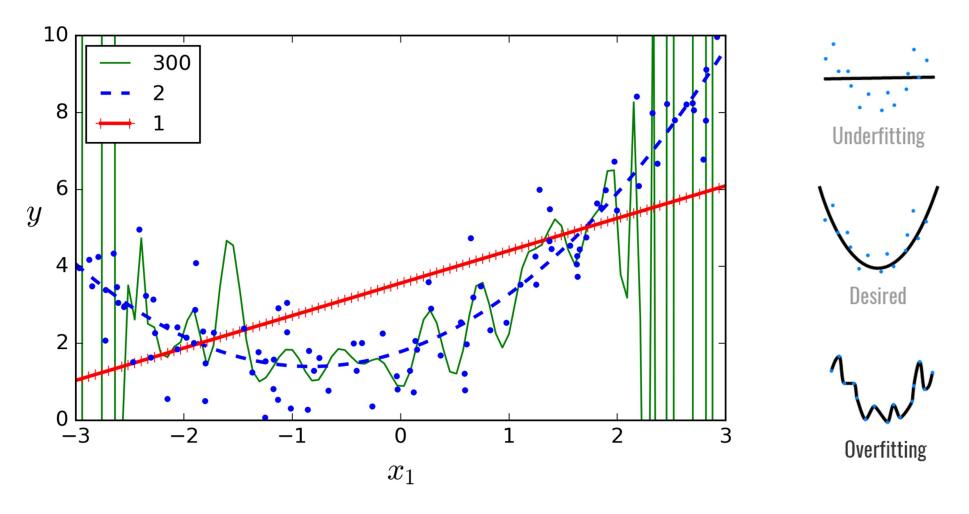
Here is an example of applying linear and polynomial regression to the data that is created using a sine function with some random noise.



Source -- https://stats.stackexchange.com/questions/350130/why-is-gradient-descent-so-bad-at-optimizing-polynomial-regression

Using too complex/simple models can lead to overfitting/underfitting,

which means the model fits the training set well but generalizes poorly on the test set.



Regression

To evaluate regression models, one common metric is the coefficient of determination (R-squared, R^2). There exist other metrics such as AIC (Akaike's Information Criterion) that is based on likelihood, which is not covered in this lecture.

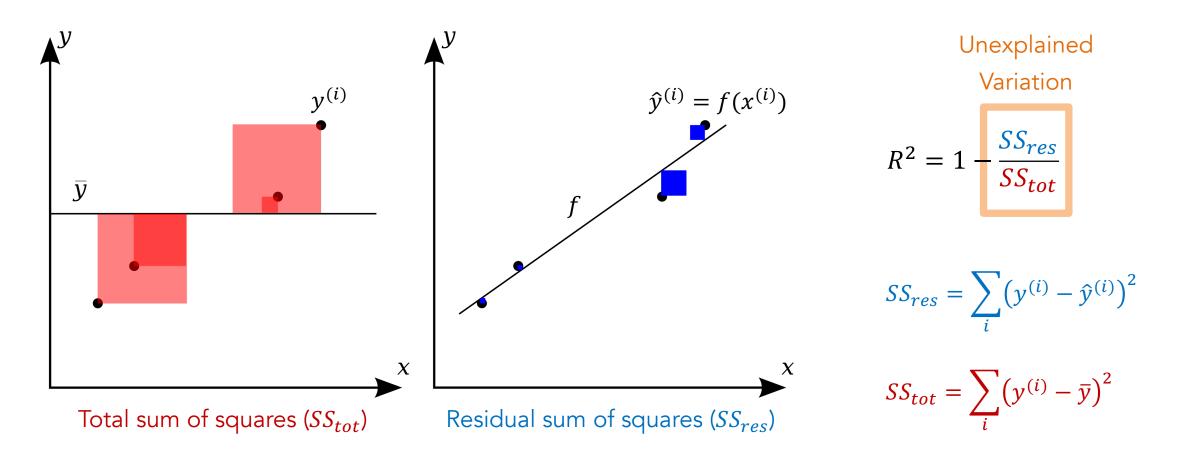
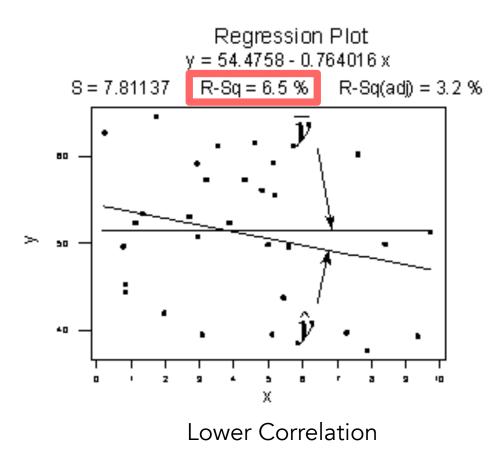
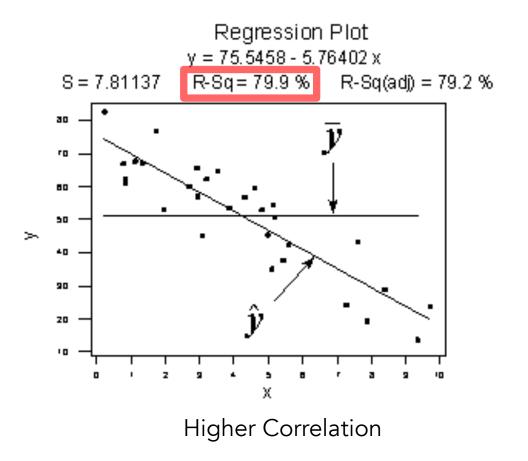


Figure source -- https://en.wikipedia.org/wiki/Coefficient_of_determination

For simple/multiple linear regression, R^2 equals the square of

Pearson correlation coefficient r between the true y and the estimated $\hat{y} = f(X)$.





 R^2 increases as we add more predictors (because the optimization

always want to decrease the residual sum of squares) and thus is not a good metric for model selection. We need the adjusted R^2 , which considers the number of predictors.

$$R_{adj}^2 = 1 - \frac{SS_{res}/df_{res}}{SS_{tot}/df_{tot}}$$

$$df_{res} = n - p - 1$$

$$df_{tot} = n - 1$$

$$SS_{res} = \sum_{i} (y^{(i)} - \hat{y}^{(i)})^2$$

$$SS_{tot} = \sum_{i} (y^{(i)} - \bar{y})^2$$

p: number of features/predictors

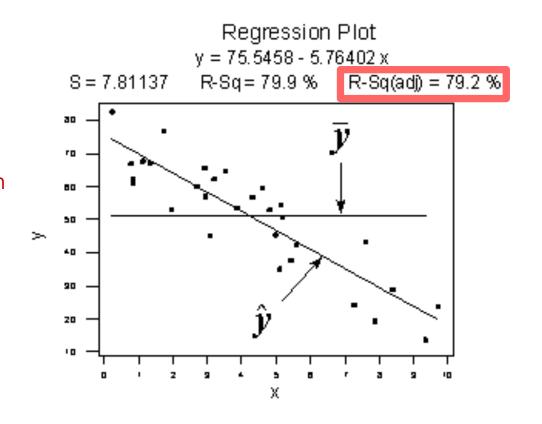
 R_{adj}^2 : adjusted value of R^2

 df_{res} : residual degree of freedom

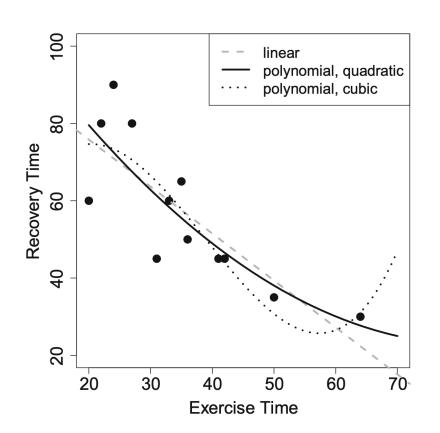
 df_{tot} : total degree of freedom

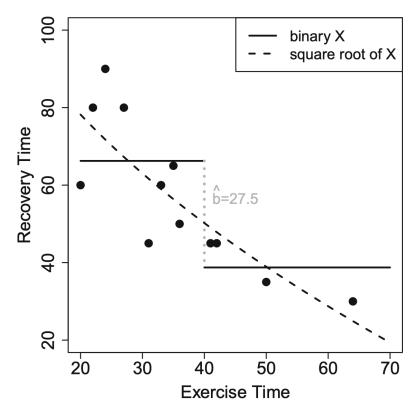
 SS_{res} : residual sum of squares

 SS_{tot} : total sum of squares



In the example below, R^2 is larger for the model with more predictors Regression (i.e., the cubic model that has three predictors). The adjusted R^2 , which considers the number of predictors (model complexity), favors the the square-root model.

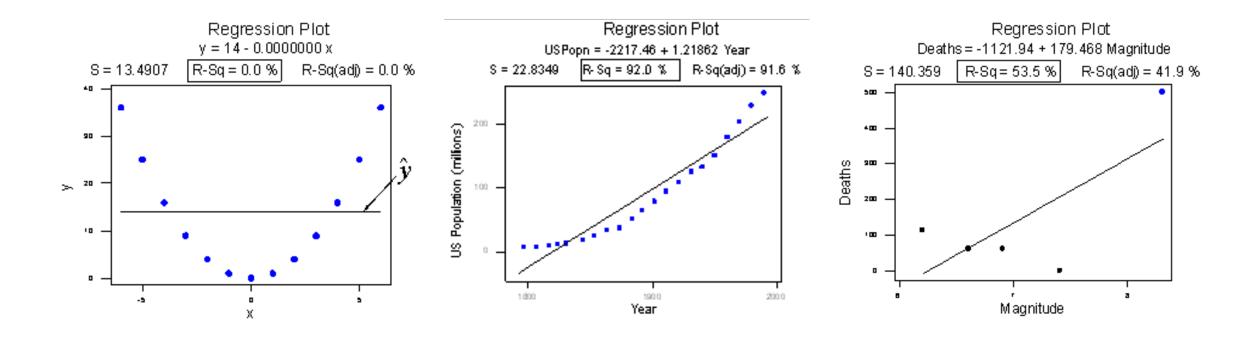




	R^2	R_{adj}^2
Linear	0.6584	0.6243
Quadratic	0.6787	0.6074
Cubic	0.7151	0.6083
Square root	0.6694	0.6363

Be careful when using and explaining R^2 in your findings. A bad

 R^2 does not always mean no pattern in the data. A good R^2 does not always mean that the function fits the data well. And R^2 can be greatly affected by outliers.



Take-Away Messages

- Classification outputs discrete labels, while regression outputs continuous values.
- Precision, recall, and F-score are common metrics for evaluating classification models.
- R-squared is a common evaluation metric for regression models.
- Feature engineering is an important step for models that do not use deep learning techniques.
- To train and update a model iteratively, you need a loss function to measure errors.
- Generally, it is a good practice to divide datasets into different parts for model training and testing.
- A model can perform extremely well on the training set but badly on the test set (i.e., overfitting).
- Cross-validation is a good technique to prevent overfitting.



Questions?