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# Machine Learning Basics

Prof. Kuan-Ting Lai 2021/3/27 THE CLASS IS DECREMENT FOR FOLLOWING

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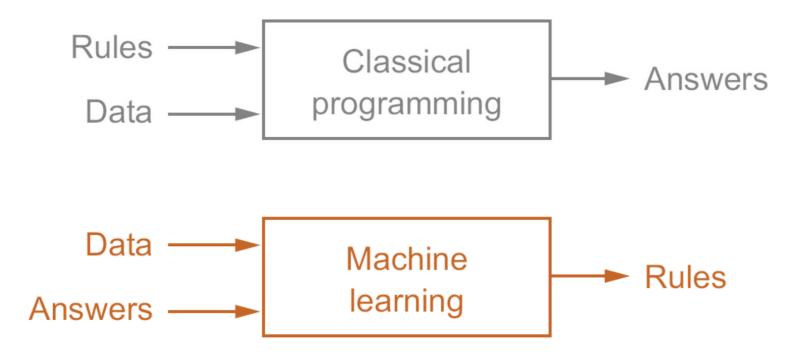
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## Machine Learning



Francois Chollet, "Deep Learning with Python," Manning, 2017

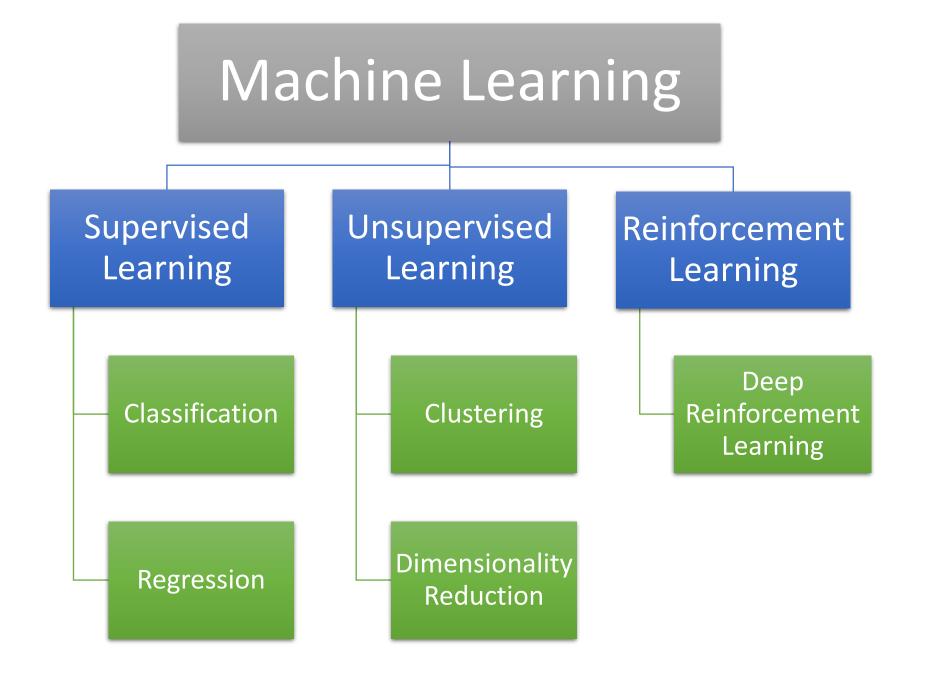
#### Machine Learning Flow

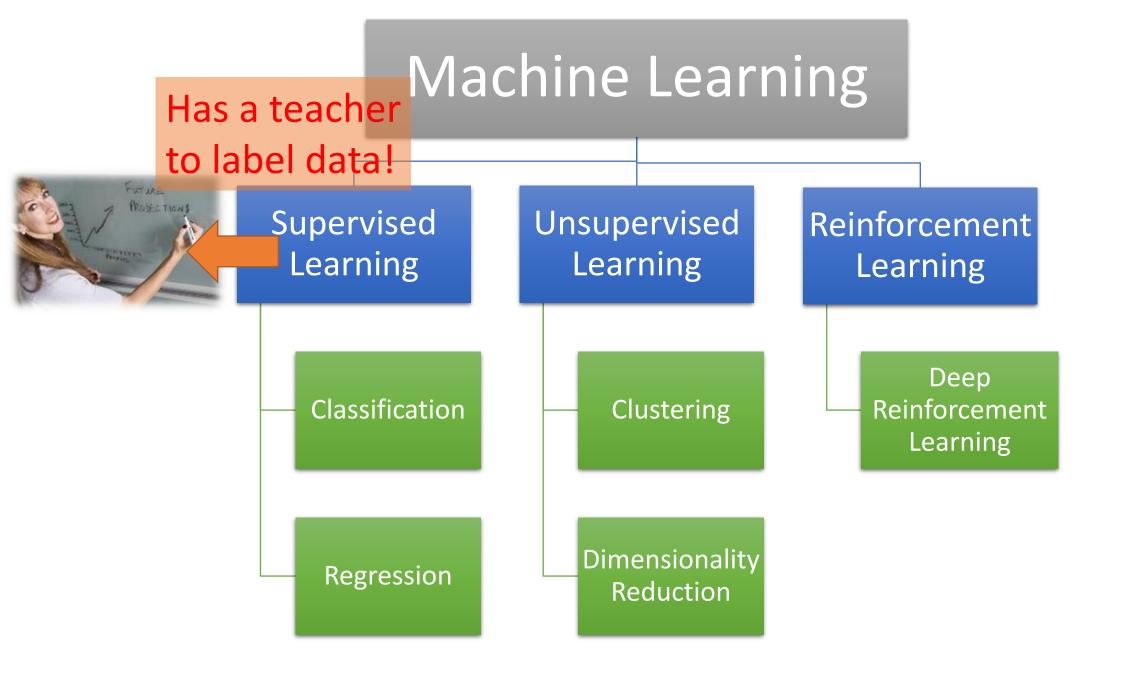


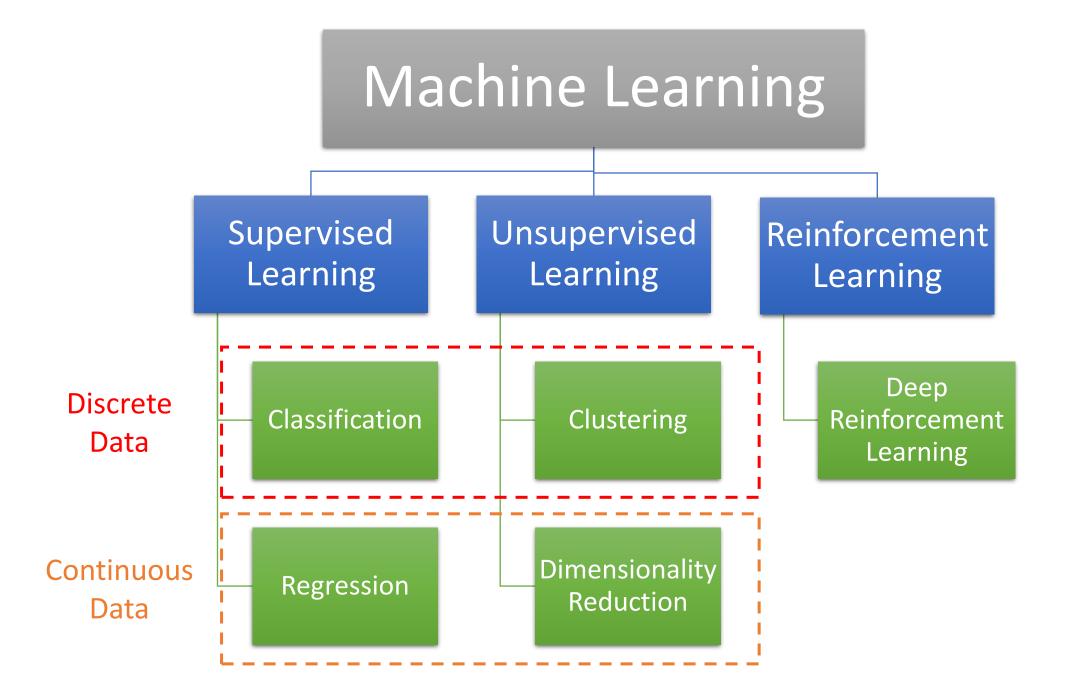
Evaluation

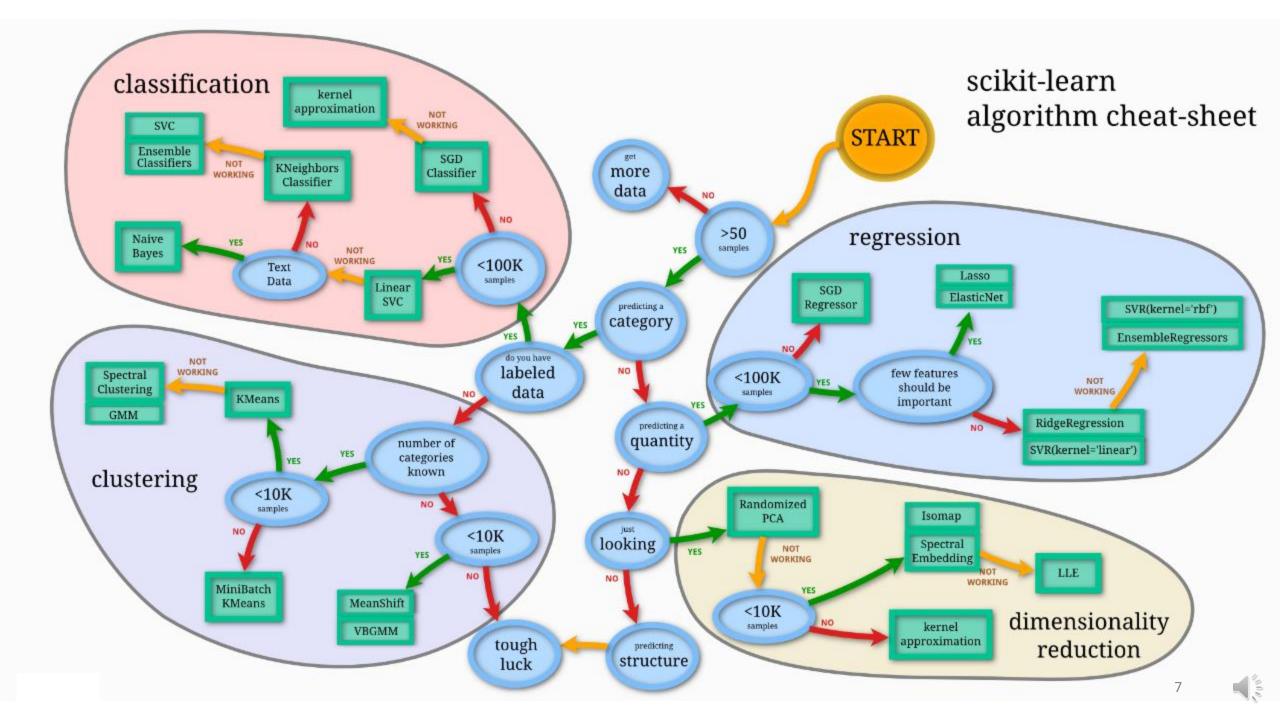
(Loss Function)

Training (Optimization)





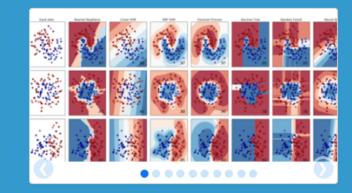




#### scikit-learn.org



Home Installation Documentation - Examples



scikit-learn

Machine Learning in Python

- Simple and efficient tools for data mining and data analysis
- · Accessible to everybody, and reusable in various contexts
- Built on NumPy, SciPy, and matplotlib
- · Open source, commercially usable BSD license

#### Classification

Identifying to which category an object belongs to.

Applications: Spam detection, Image recognition. Algorithms: SVM, nearest neighbors, random forest, ... — Examples

#### Dimensionality reduction

Reducing the number of random variables to consider.

Applications: Visualization, Increased efficiency Algorithms: PCA, feature selection, nonnegative matrix factorization. — Examples

#### Regression

Predicting a continuous-valued attribute associated with an object.

Applications: Drug response, Stock prices. Algorithms: SVR, ridge regression, Lasso, — Examples

#### Model selection

Comparing, validating and choosing parameters and models.

Goal: Improved accuracy via parameter tuning Modules: grid search, cross validation, metrics. — Examples

#### Clustering

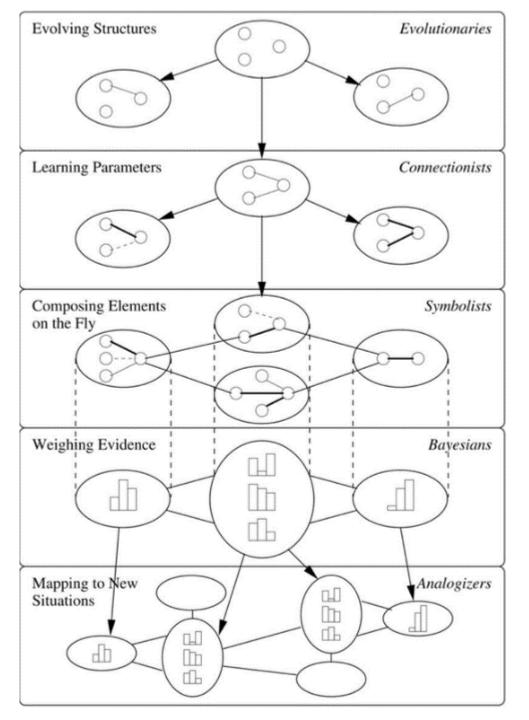
Automatic grouping of similar objects into sets.

Applications: Customer segmentation, Grouping experiment outcomes Algorithms: k-Means, spectral clustering, mean-shift, ... — Examples

#### Preprocessing

Feature extraction and normalization.

Application: Transforming input data such as text for use with machine learning algorithms. Modules: preprocessing, feature extraction.



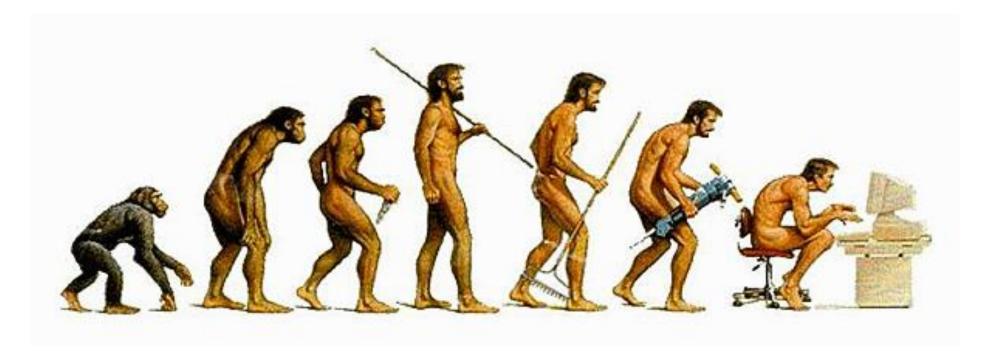
# Five Tribes of Machine Learning

- Evolutionaries
- Connectionists
- Symbolists
- Bayesians
- Analogizers

Pedro Domingos, "The Master Algorithm," 2015

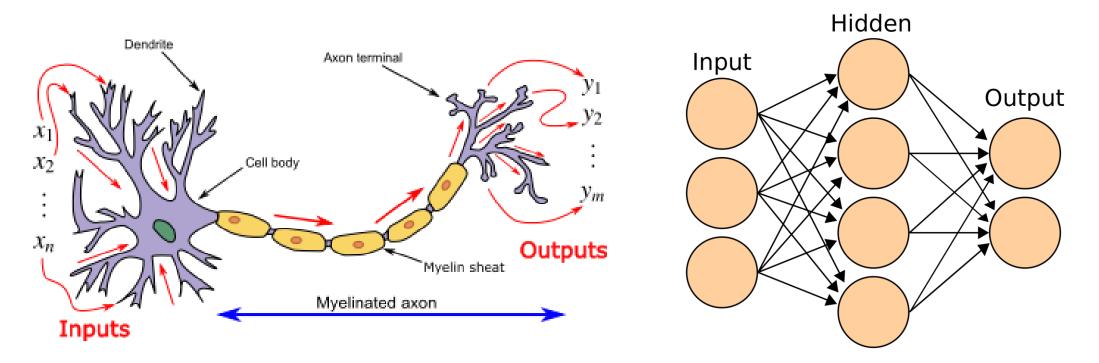
#### Evolutionaries: Survival of the Fittest

- Genetic algorithms, evolutionary algorithms
  - Crossover, mutation



#### Connectionists: Neural Networks

• Neural Networks, deep learning

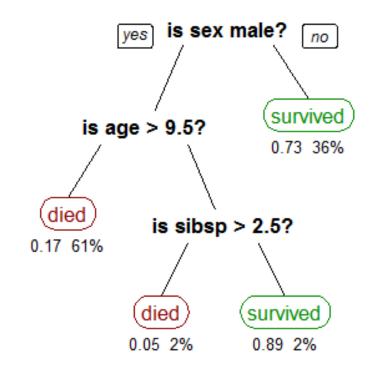


https://en.wikipedia.org/wiki/Artificial neural network

### Symbolists: Finding the Rules

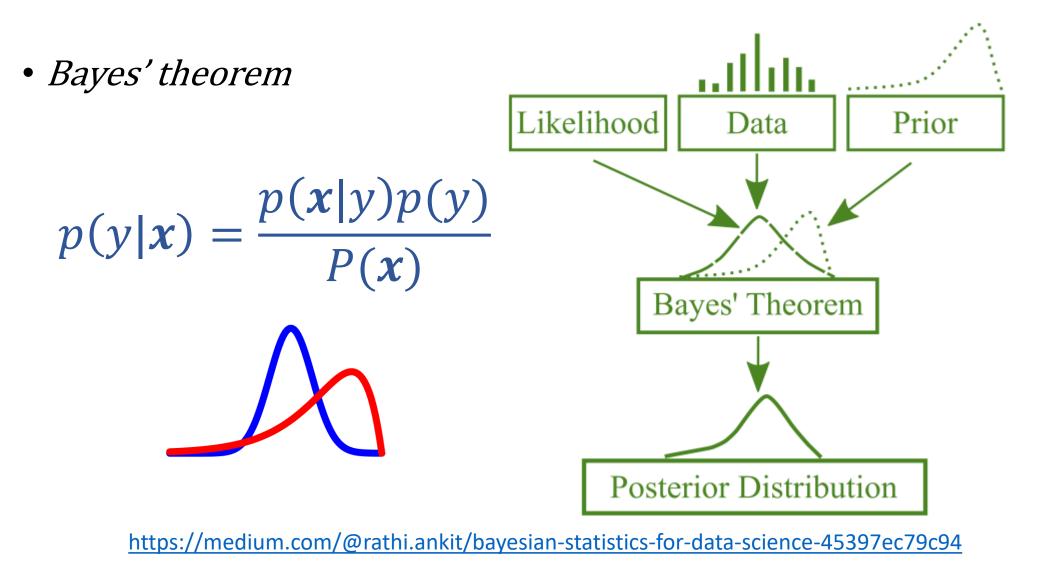
• Decision Tree, expert system, rule-based system

Survival rate of Passengers on Titanic



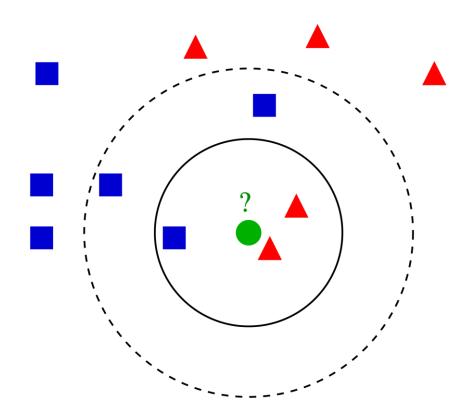


### Bayesians: Anything happens, just the Probability



#### Analogizers: Like Father Like Son

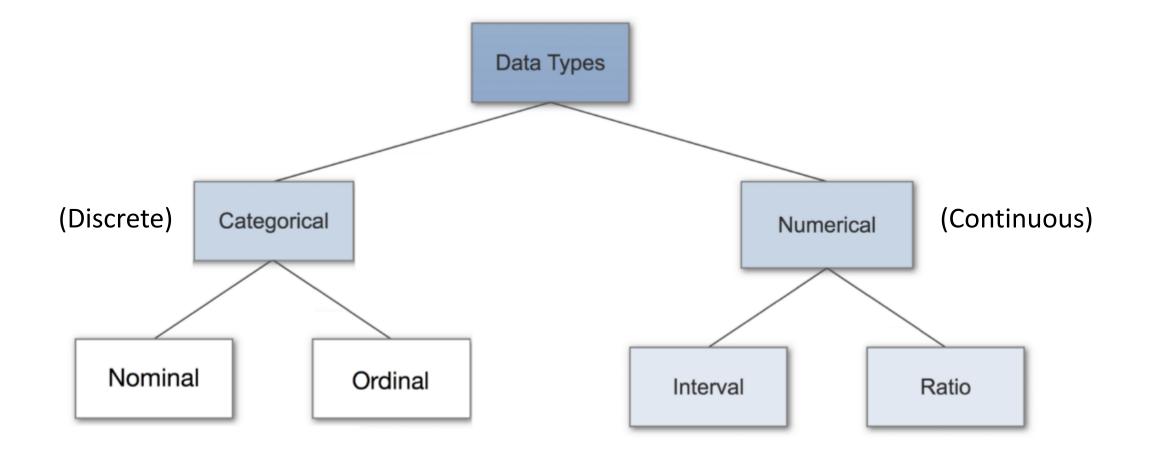
• k Nearest-Neighbors (kNN), SVM



https://en.wikipedia.org/wiki/K-nearest\_neighbors\_algorithm

# Types of Data

#### Data Types (Measurement Scales)



https://towardsdatascience.com/data-types-in-statistics-347e152e8bee

16

### Nominal Data (Labels)

- Nominal data are labeling variables without any quantitative value
- Encoded by one-hot encoding for machine learning
- Examples:

What is your Gender?	What languages do you speak?
O Female	O Englisch
O Male	O French
	O German
	O Spanish

#### Ordinal Data

- Ordinal values represent discrete and ordered units
- The order is meaningful and important

What Is Your Educational Background?

- 1 Elementary
- 🔘 2 High School
- 3 Undegraduate
- 🔵 4 Graduate

#### Interval Data

- Interval values represent ordered units that have the same difference
- Problem of Interval: Don't have a true zero
- Example: Temperature Celsius (°C) vs. Fahrenheit (°F)

Temperature? - 10 -5 0 0 + 5 + 10 + 15



- Same as interval data but have absolute zero
- Can be applied to both descriptive and inferential statistics
- Example: weight & height



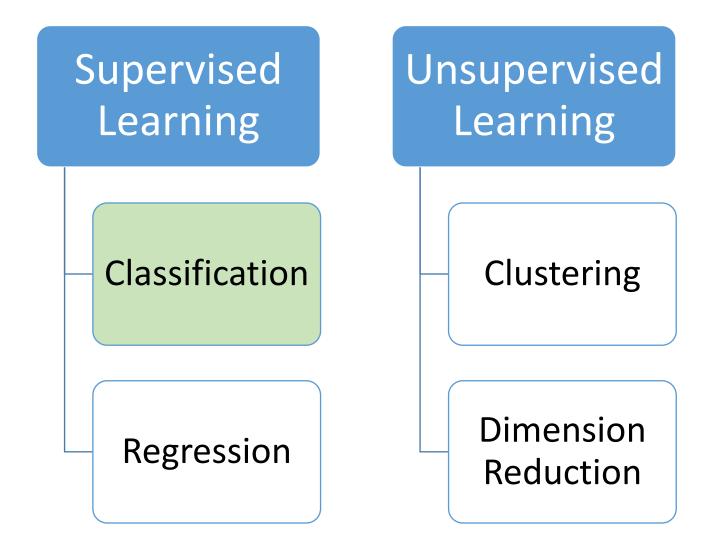


#### Machine Learning vs. Statistics

<u>https://www.r-bloggers.com/whats-the-difference-between-machine-learning-statistics-and-data-mining/</u>

Machine learning	Statistics
network, graphs	model
weights	parameters
learning	fitting
generalization	test set performance
supervised learning	regression/classification
unsupervised learning	density estimation, clustering
large grant = \$1,000,000	large grant = \$50,000
nice place to have a meeting: Snowbird, Utah, French Alps	nice place to have a meeting: Las Vegas in August

#### Supervised and Unsupervised Learning



#### Iris Flower Classification



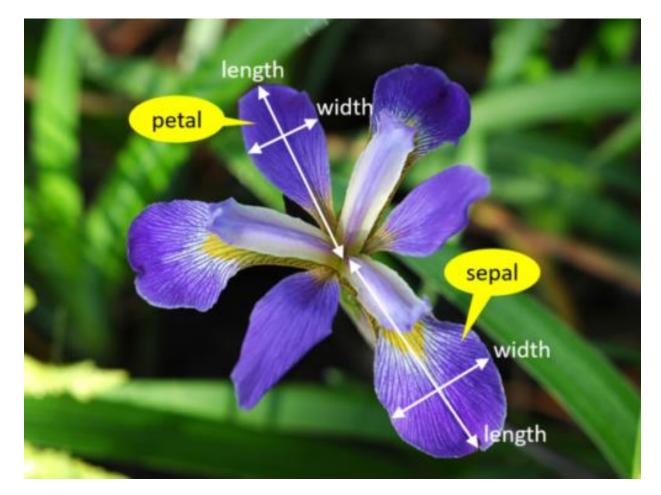
**Iris Versicolor** 

Iris Setosa

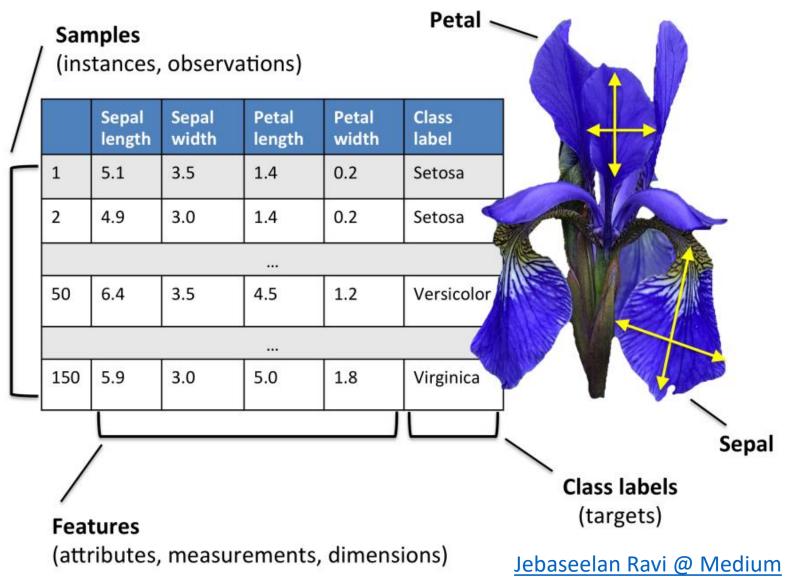
#### Iris Virginica

#### Extracting Features of Iris

• Width and Length of Petal and Sepal

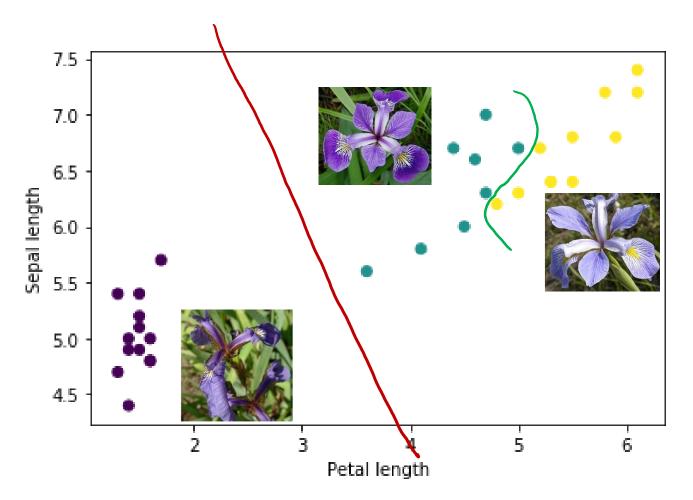


#### Iris Flower Dataset

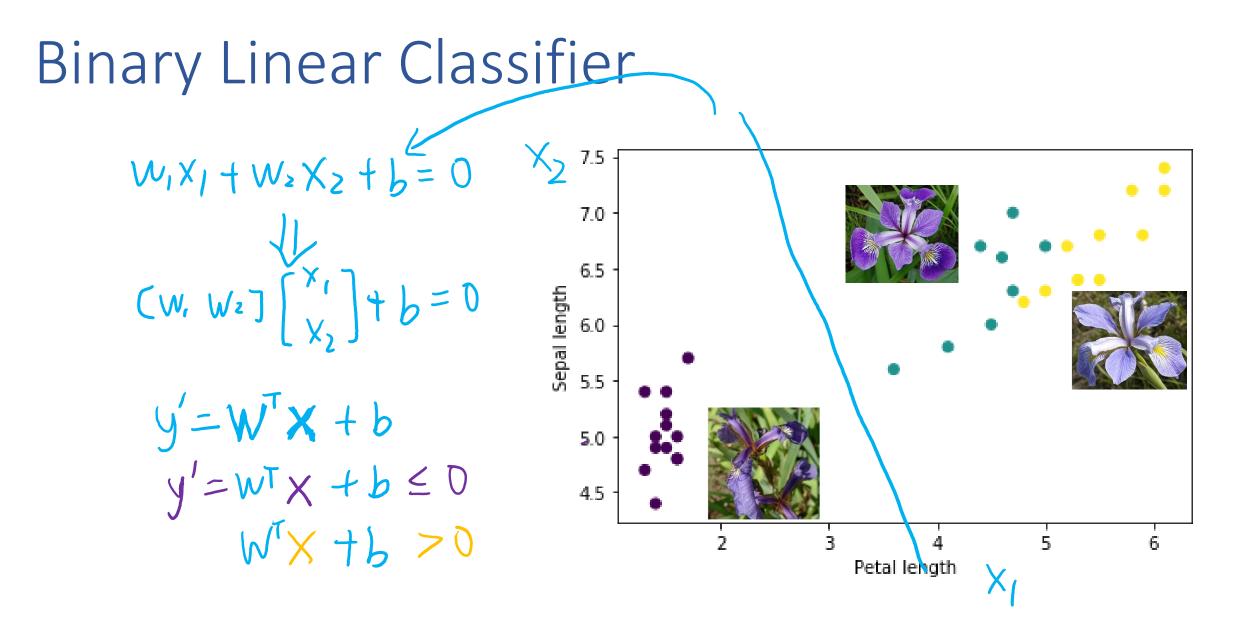


### Classify Iris Species via Petals and Sepals

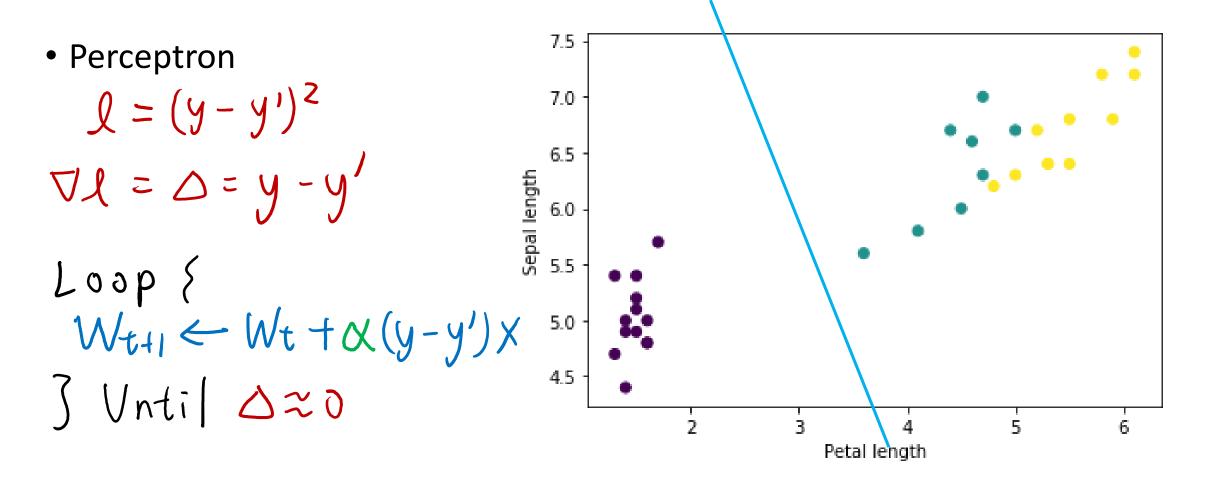
 Iris versicolor and virginica are not linearly separable



https://www.tensorflow.org/tutorials/customization/custom\_training\_walkthrough



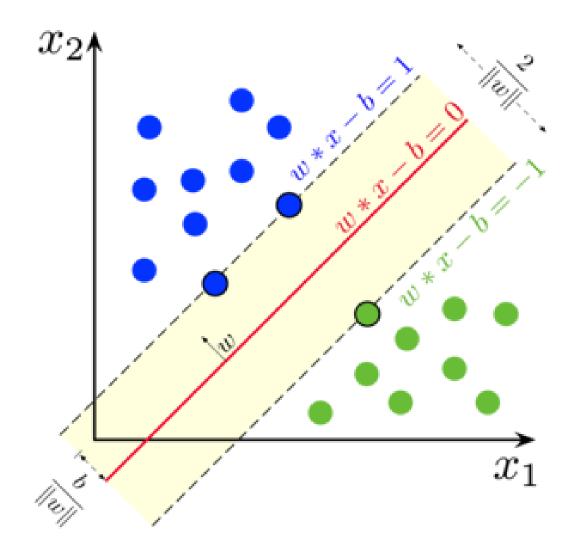
#### Training Linear Classifier



 $y' = \boldsymbol{w}^T \boldsymbol{x} + b$ 

#### Support Vector Machine (SVM)

• Choose the hyperplanes that have the largest separation (margin)



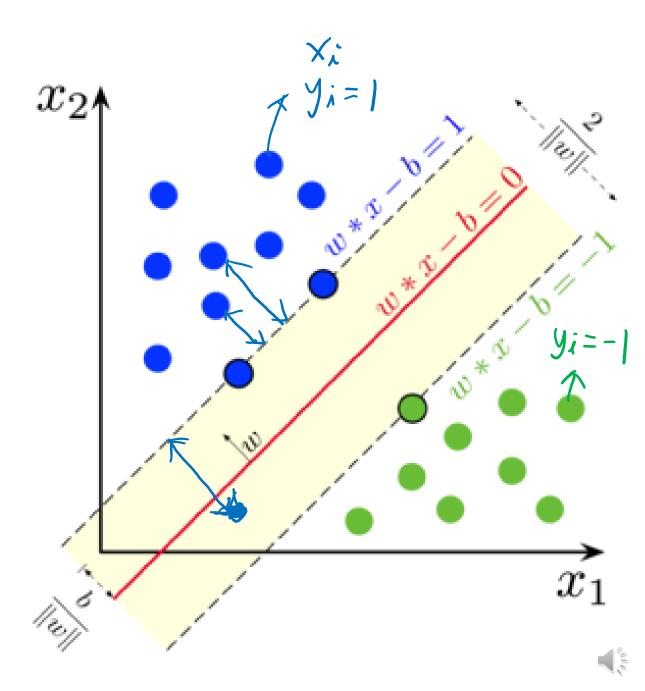
### Loss Function of SVM

• Hinge Loss

$$y_{i} = w^{T} x_{i} - b \ge 1$$
  

$$y_{i} = w^{T} x_{i} - b \le -1$$
  

$$y_{i} (w^{T} x_{i} - b) \ge 1$$



#### SVM Optimization

• Maximize the margin while reducing hinge losses

$$\min_{x \to 1} \frac{1}{||w||^2} x_2$$

$$s. b. t. \frac{1}{N} \sum_{x \to 0}^{N} \max(0, 1 - y_i(w_{x_i - b}))$$

$$J(w) = \frac{1}{2} ||w||^2 + C\left[\frac{1}{N} \sum_{x \to 0}^{N} \max(0, 1 - y_i(w_{x_i}))\right]$$

$$\frac{\partial J(w)}{\partial w} = \frac{1}{N} \sum_{x \to 0}^{N} \left( W \quad \text{if } \max(0, 1 - y_i(w_{x_i}))\right) = 0$$

$$W = \frac{1}{N} \sum_{x \to 0}^{N} \left( W \quad \text{if } \max(0, 1 - y_i(w_{x_i}))\right) = 0$$

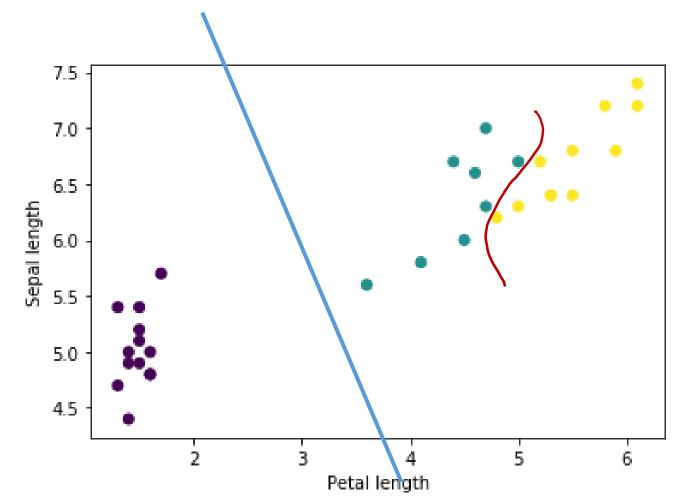
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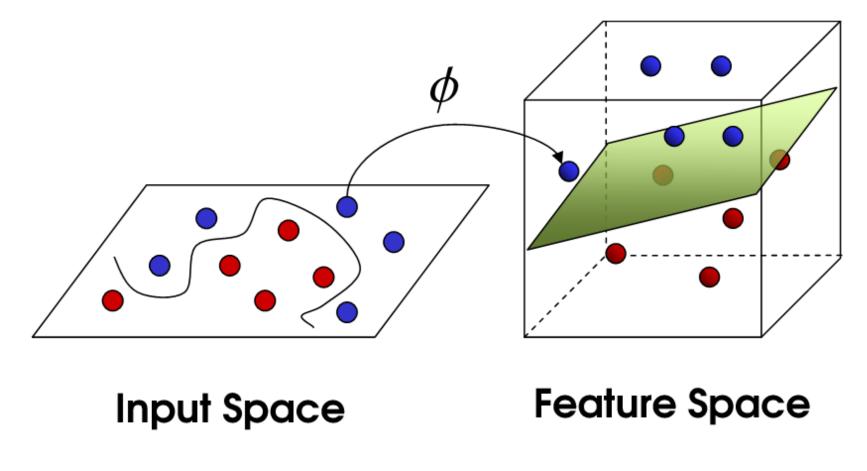
#### Nonlinear Problem for SVM?

• How to separate Versicolor and Virginica?



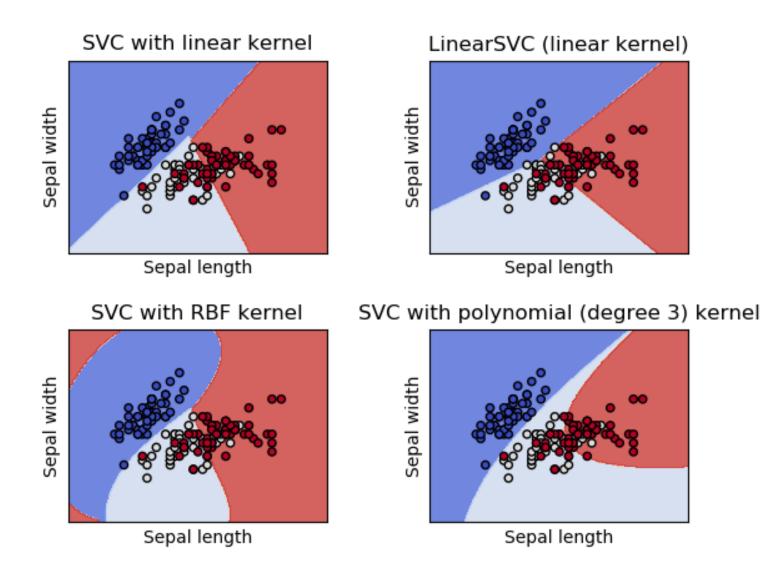
#### SVM Kernel Trick

• Project data into higher dimension and calculate the inner products



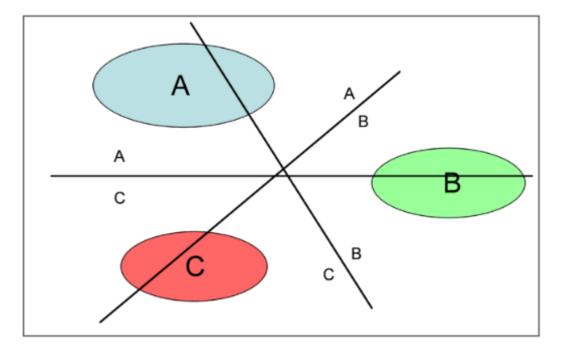
https://datascience.stackexchange.com/questions/17536/kernel-trick-explanation

### Nonlinear SVM for Iris Classification

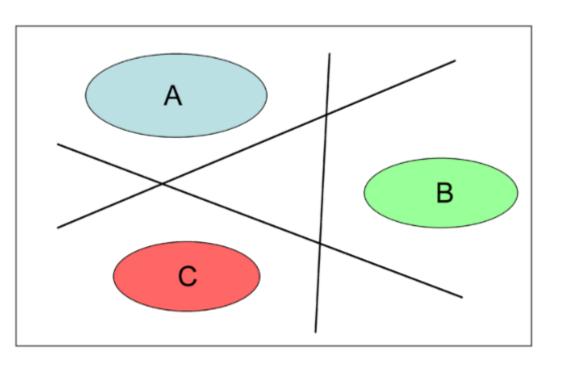


#### Multi-class SVM

### • One-against-One $\begin{pmatrix} n \\ 2 \end{pmatrix}$

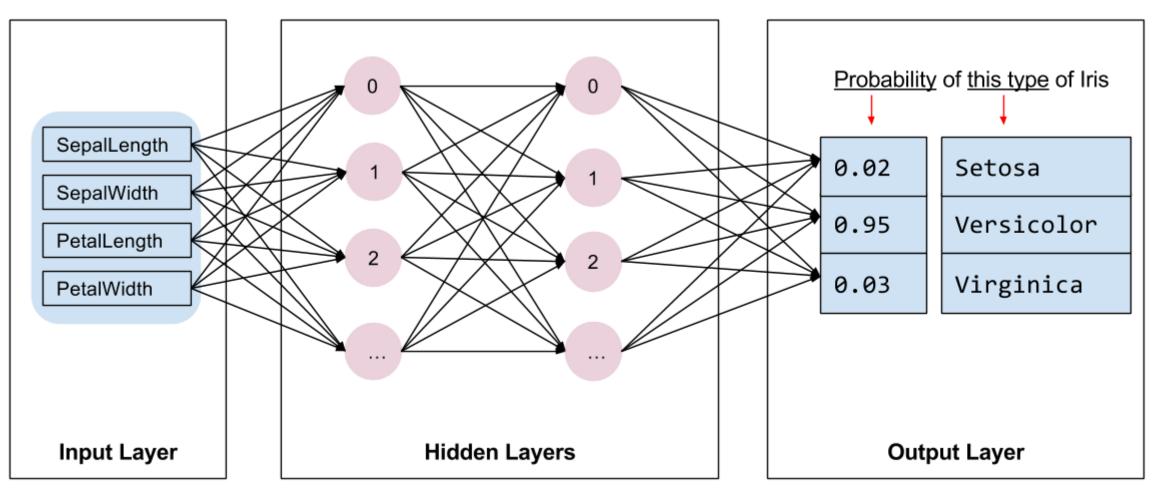


#### • One-against-All



https://courses.media.mit.edu/2006fall/mas622j/Projects/aisen-project/

### Using Neural Network



https://www.tensorflow.org/tutorials/customization/custom\_training\_walkthrough\_

## Feature Engineering & Feature Learning

#### • Data preprocessing

- Vectorization
- Normalization
- handling missing value
- Feature engineering
  - Good features still make learning easier

Raw data: pixel grid		
Better features: clock hands' coordinates	{x1: 0.7, y1: 0.7} {x2: 0.5, y2: 0.0}	{x1: 0.0, y2: 1.0} {x2: -0.38, 2: 0.32}
Even better features: angles of clock hands	theta1: 45 theta2: 0	theta1: 90 theta2: 140

#### Naïve Bayes

• Probabilistic classifier based on Bayes' theorem

$$P(y|\mathbf{x}) = P(y|x_1, x_2, ..., x_n)$$

$$P(y|\mathbf{x}) = \frac{P(y)P(\mathbf{x}|y)}{P(\mathbf{x})}$$

$$Posterior = \frac{Prior \times Liklihood}{Evidence}$$

#### Naïve Bayes

• Probabilistic classifier based on Bayes' theorem

$$P(y|\mathbf{x}) = P(y|x_1, x_2, ..., x_n)$$

$$P(y|\mathbf{x}) = \frac{P(y)P(\mathbf{x}|y)}{P(\mathbf{x})} \qquad Posterior = \frac{Prior \times Liklihood}{Evidence}$$

• Bayes assumes features  $x_i$  are conditional independent

$$P(x|y) = P(x_1|y) P(x_2|y) \cdots P(x_n|y) = \prod_{i=1}^{n} P(x_i|y)$$

$$\Rightarrow P(y|\mathbf{x}) = \frac{P(y)\prod_{i=1}^{n} P(x_i|y)}{P(\mathbf{x})} \propto P(y) \prod_{i=1}^{n} P(x_i|y)$$
$$\Rightarrow \hat{y} = \arg\max_{y} P(y) \prod_{i=1}^{n} P(x_i|y)$$

#### Gaussian Naive Bayes in Scikit

• P(
$$x_i | y_k$$
) =  $\frac{1}{\sqrt{2\pi}\sigma_k} e^{\left(-\frac{(x-\sigma\mu_k)^2}{2\sigma_k^2}\right)}$ 

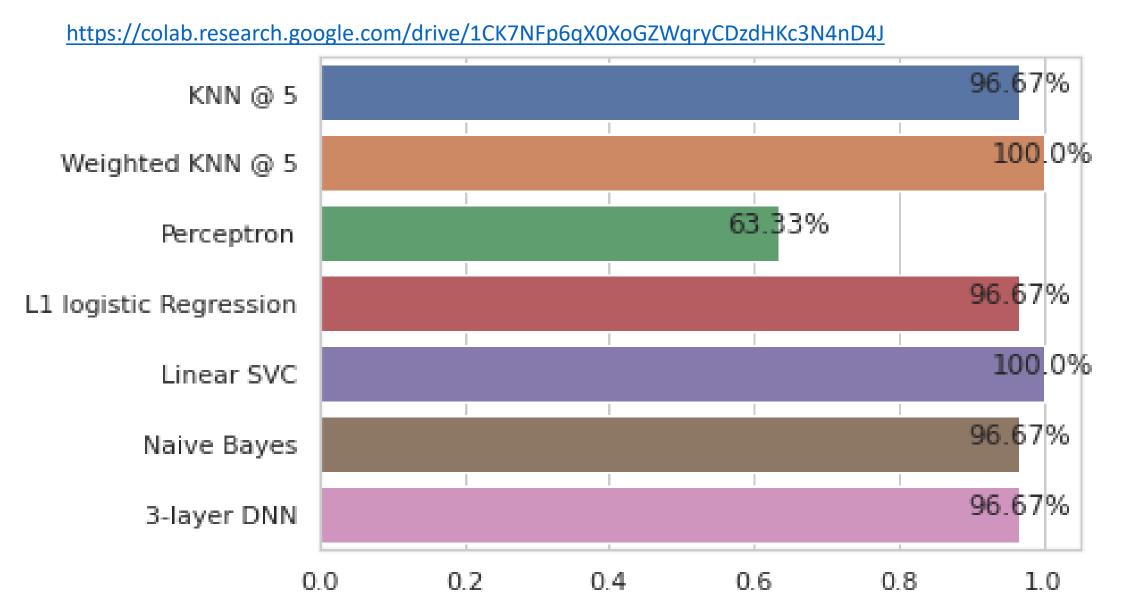
#### >>> from sklearn import datasets

- >>> iris = datasets.load\_iris()
- >>> from sklearn.naive\_bayes import GaussianNB
- >>> gnb = GaussianNB()
- >>> y\_pred = gnb.fit(iris.data, iris.target).predict(iris.data)
- >>> print("Number of mislabeled points out of a total %d points : %d"
- ... % (iris.data.shape[0],(iris.target != y\_pred).sum()))
- Number of mislabeled points out of a total 150 points : 6

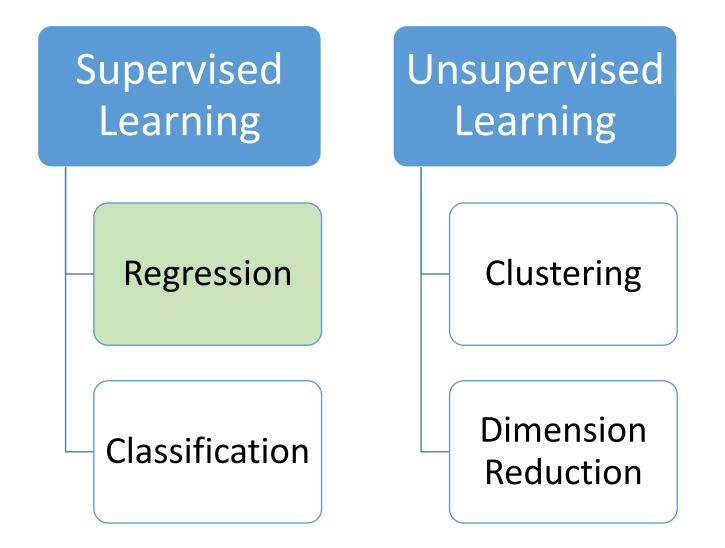
#### Evaluating All Classifiers on Iris Flower Dataset

```
from sklearn.metrics import accuracy score
                                                                                           Notebook Link
    from sklearn.linear model import *
    from sklearn.svm import SVC
    from sklearn.neighbors import KNeighborsClassifier
    from sklearn import datasets
    from sklearn.naive_bayes import GaussianNB
[2] iris = datasets.load iris()
    X = iris.data
    v = iris.target
    print('There are {} training samples in the Iris dataset'.format(len(X)))
    There are 150 training samples in the Iris dataset
[3] # Create different classifiers.
    knn = 5
    Cost = 10 # For regularization
    classifiers = {
           'KNN @ ' + str(knn): KNeighborsClassifier(knn, 'uniform'),
           'Weighted KNN @ ' + str(knn): KNeighborsClassifier(knn, 'distance'),
           'Perceptron': Perceptron(),
           'L1 logistic Regression': LogisticRegression(C=Cost, penalty='11', solver='saga', multi_class='multinomial', max_iter=10000),
           'Linear SVC': SVC(kernel='linear', C=Cost, probability=False, random state=0),
           'Naive Bayes': GaussianNB()
[4] # Training
    for index, (name, classifier) in enumerate(classifiers.items()):
           classifier.fit(X, y)
           y_pred = classifier.predict(X)
            accuracy = accuracy score(y, y pred)
           print ("Accuracy (train) for %s: %0.1f%% " % (name, accuracy * 100))
```

### Classifier Evaluation on Iris dataset

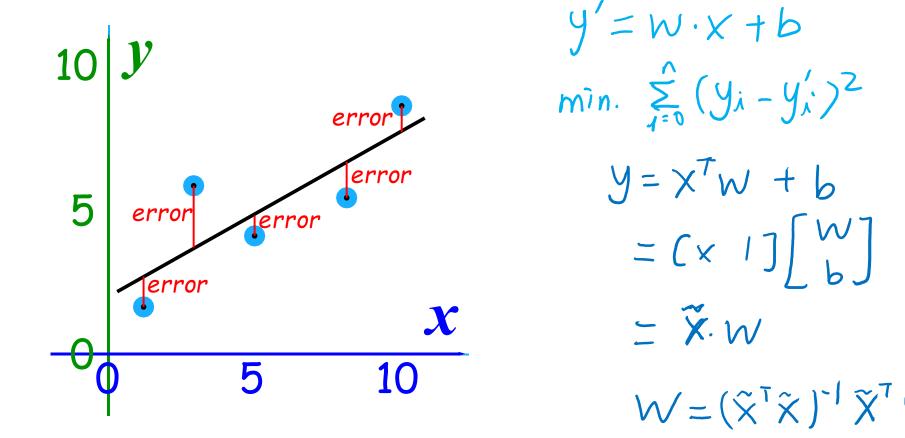


#### Supervised and Unsupervised Learning



#### Linear Regression (Least squares)

• Find a "line of best fit" that minimizes the total of the square of the errors



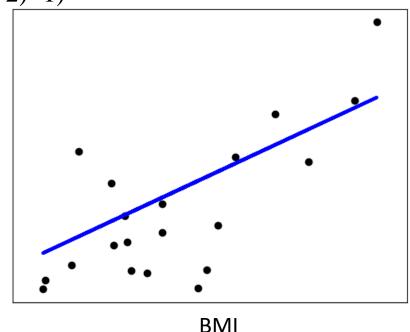
#### Least Squares Solution

- Pseudo Inverse:  $w = (x^T x)^{-1} x^T y$
- xw = y has a unique solution if and only if -The columns of x are linearly independent.  $-x^T x$  is invertible.

#### Scikit-Learn Diabetes Dataset

- 10 features: age, sex, body mass index, average blood pressure, and six blood serum measurements
- Obtained from n = 442 diabetes patients
- Normalized to have mean 0 and squared length =  $1 (sum(x^2)=1)$

Total sample	442
Dimensionality	10
Features	real,2 < x < .2
Targets	integer 25 - 346

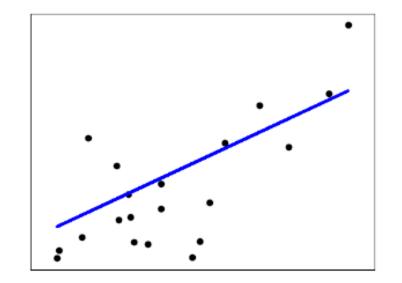


https://www4.stat.ncsu.edu/~boos/var.select/diabetes.html

#### Linear Regression

• Least Squares

$$\min_{w} \left\| y - (w^T x + b) \right\|_2^2$$



```
import matplotlib.pyplot as plt
import numpy as np
from sklearn import datasets, linear_model
from sklearn.metrics import mean_squared_error, r2_score
# Load the diabetes dataset
diabetes = datasets.load_diabetes()
# Use only one feature
diabetes_X = diabetes.data[:, np.newaxis, 2]
diabetes_X_train = diabetes_X[:-20]
diabetes_y_train = diabetes.target[:-20]
diabetes_X_test = diabetes_X[-20:]
diabetes_y_test = diabetes.target[-20:]
```

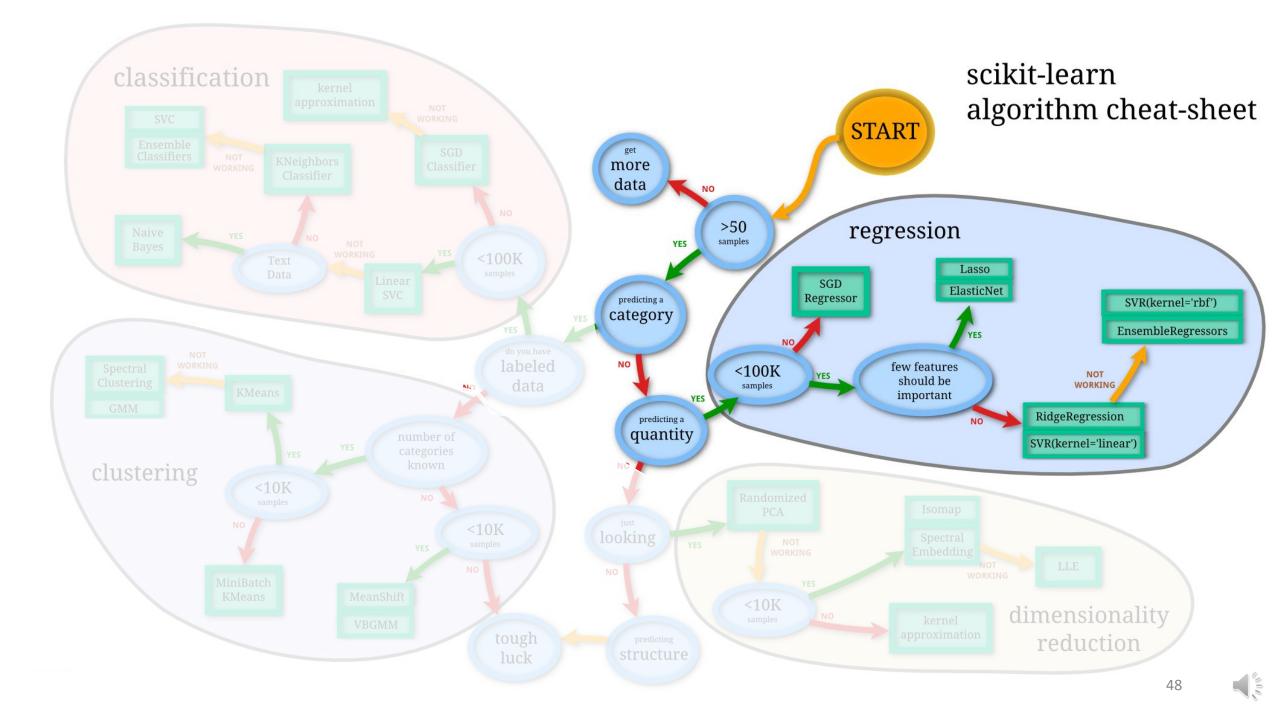
```
# Create linear regression object
regr = linear_model.LinearRegression()
```

```
# Train the model using the training sets
regr.fit(diabetes_X_train, diabetes_y_train)
```

```
# Make predictions using the testing set
diabetes_y_pred = regr.predict(diabetes_X_test)
```

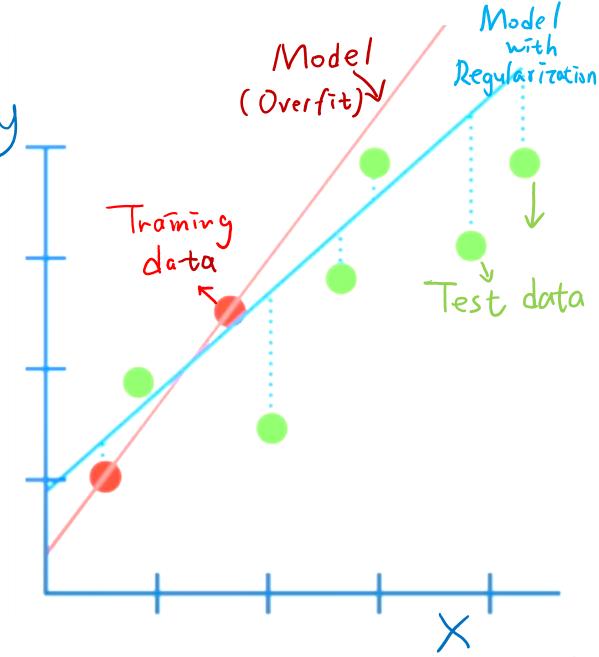
```
# Plot outputs
```

plt.scatter(diabetes\_X\_test, diabetes\_y\_test, color='black')
plt.plot(diabetes\_X\_test, diabetes\_y\_pred, color='blue',
linewidth=3)
plt.xticks(())
plt.yticks(())
plt.show()

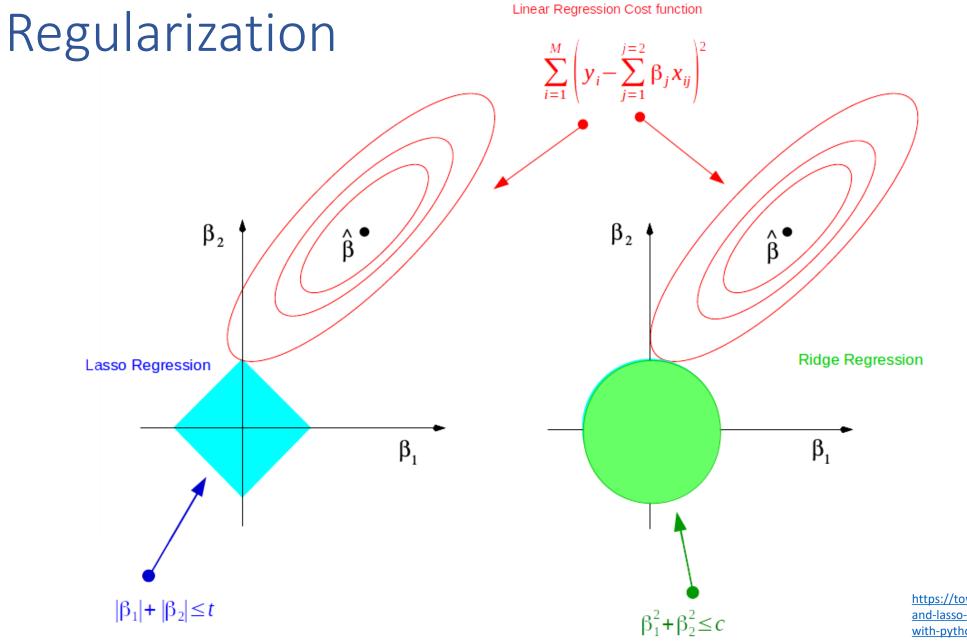


### Regularization

- Linear regression is sensitive to outliers and easy to overfit training data
- Regularization add small amount of bias during training to reduce variance on test data



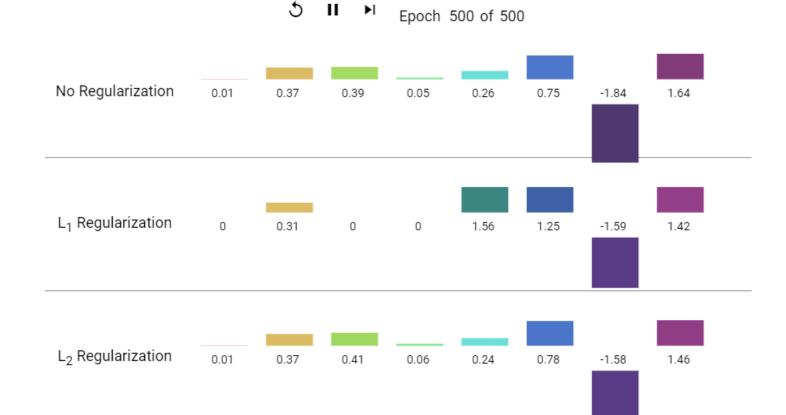
**Dimension Reduction of Feature Space with LASSO** 



https://towardsdatascience.com/ridgeand-lasso-regression-a-complete-guidewith-python-scikit-learn-e20e34bcbf0b

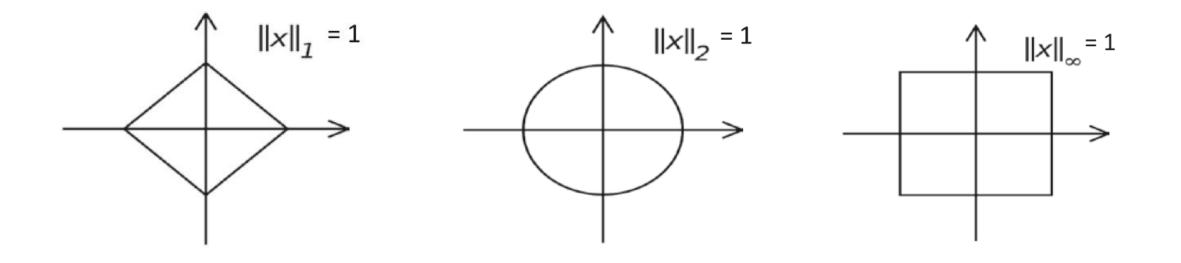
### Regularization

<u>https://developers.google.com/machine-learning/crash-course/regularization-for-sparsity/l1-regularization</u>



#### Unit norms in 2D Vectors

• The set of all vectors of norm 1 in different 2D norms



#### Ridge, Lasso and ElasticNet

• Ridge regression:  $\min_{w} ||Xw - y||_2^2 + \alpha ||w||_2^2$ 

• Lasso regression: 
$$\min_{w} \frac{1}{2n_{\text{samples}}} ||Xw - y||_{2}^{2} + \alpha ||w||_{1}$$

• Elastic Net: 
$$\min_{w} \frac{1}{2n_{\text{samples}}} ||Xw - y||_{2}^{2} + \alpha \rho ||w||_{1} + \frac{\alpha(1 - \rho)}{2} ||w||_{2}^{2}$$



#### Predicting Boston House Prices

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#### Boston House Price Dataset

- Objective: predict the median price of homes
- Small dataset with 506 samples and 13 features
  - <u>https://www.kaggle.com/c/boston-housing</u>

1	crime	per capita crime rate by town.	8	dis	weighted mean of distances to five Boston employment centres.
2	zn	proportion of residential land zoned for lots over 25,000 sq.ft.	9	rad	index of accessibility to radial highways.
3	indus	proportion of non-retail business acres per town.	10	tax	full-value property-tax rate per \$10,000.
4	chas	Charles River dummy variable (= 1 if tract bounds river; 0 otherwise).	11	ptratio	pupil-teacher ratio by town.
5	nox	nitrogen oxides concentration	12	black	1000(Bk - 0.63) <sup>2</sup> where Bk is the proportion of blacks by town.
6	rm	average number of rooms per dwelling.	13	lstat	lower status of the population (percent).
7	age	proportion of owner-occupied units built prior to 1940.			55

#### Normalize the Data

- Make features center around 0 and have unit standard deviation
- Note that the quantities (mean, std) used for normalizing the test data are computed using the training data!

```
# Nomalize the data
mean = train_data.mean(axis=0)
train_data -= mean
std = train_data.std(axis=0)
train_data /= std
test_data -= mean
test_data /= std
```

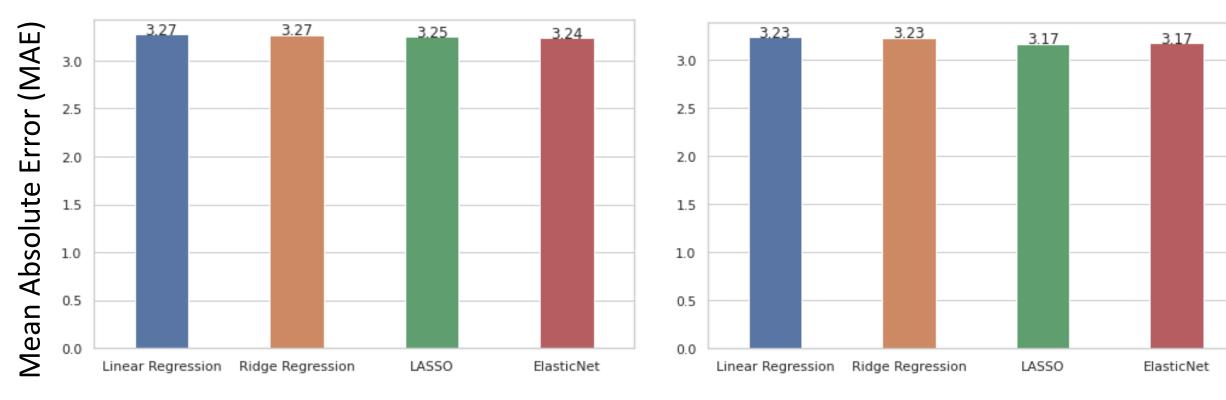
```
+ 程式碼 + 文字
                             Notebook link...
\equiv
            from sklearn.metrics import mean_squared_error, mean_absolute_error
       [1]
            from sklearn.linear model import *
Q
            from sklearn import datasets
<>
            data = datasets.load_boston()
        X = data.data
y = data.target
            y_mean = y.mean()
            print('There are {} training samples in the Boston House dataset. The average price is {}'.format(len(X), y_mean))
            There are 506 training samples in the Boston House dataset. The average price is 22.532806324110677
       [3] mean = X.mean(axis=0)
            X -= mean
            std = X.std(axis=0)
            X /= std
       [4] regressors = {
                    'Linear Regression': LinearRegression(),
                   'Ridge Regression': Ridge(alpha=.5),
                   'LASSO': Lasso(alpha=0.1),
                   'ElasticNet': ElasticNet(alpha=0.1)
       [5] # Training
            mae_list = []
            for index, (name, regressor) in enumerate(regressors.items()):
                   regressor.fit(X, y)
                   y_pred = regressor.predict(X)
                   mse = mean_absolute_error(y, y_pred)
                   print("The Mean Absolute Error for %s: %0.4f " % (name, mse))
                   mae_list.append(mse)
            The Mean Absolute Error for Linear Regression: 3.2709
            The Mean Absolute Error for Ridge Regression: 3.2691
            The Mean Absolute Error for LASSO: 3.2464
=:
            The Mean Absolute Error for ElasticNet: 3.2387
```

### Regressor Comparison

#### **Comparison of Regularization Methods**

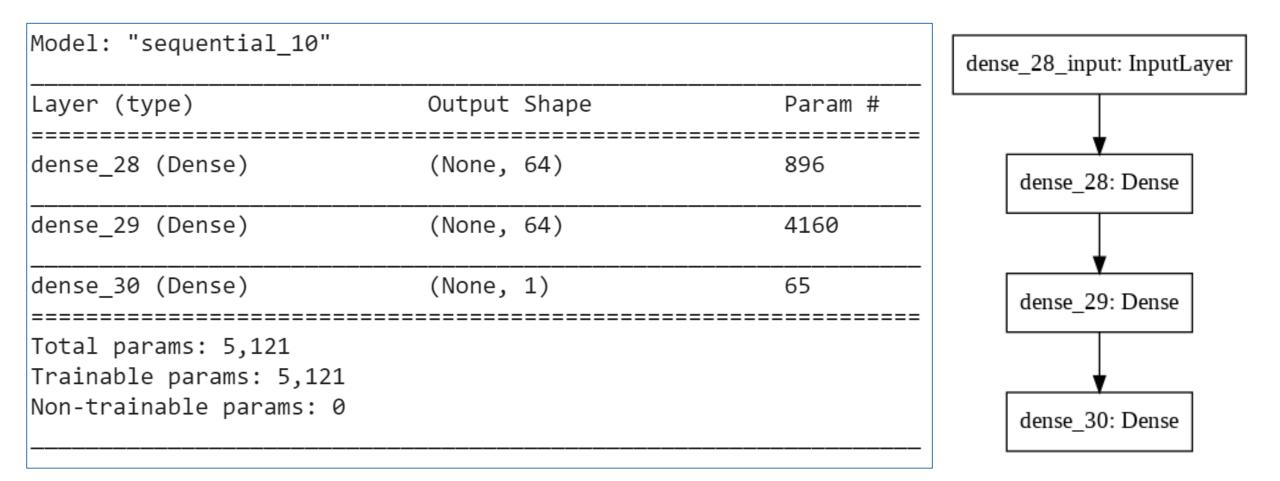
**Training Data (506 samples)** 

#### Test Data (102 samples)



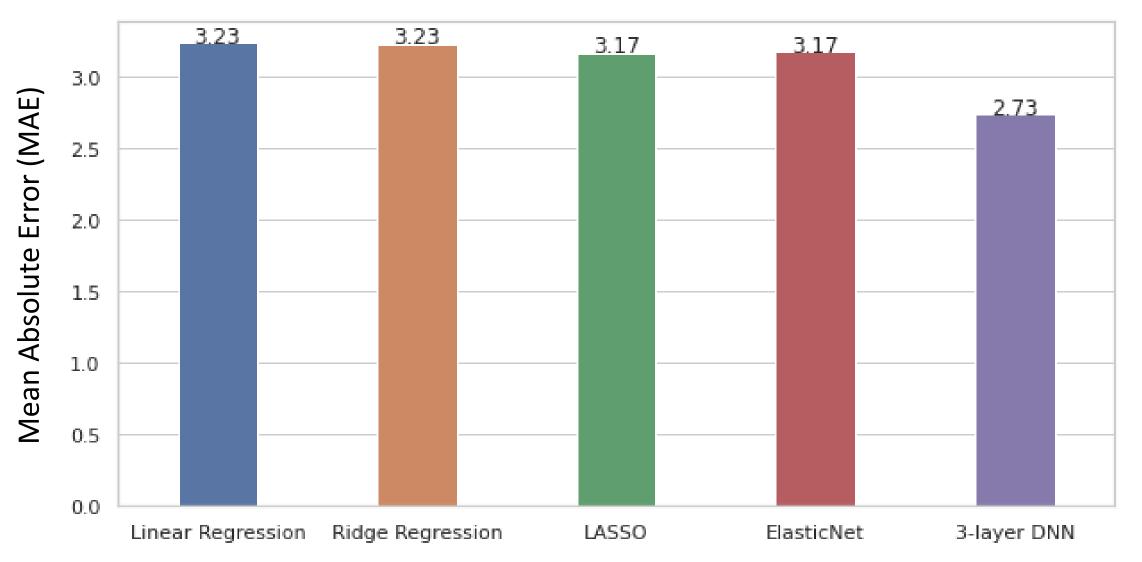
https://colab.research.google.com/drive/1lgITg2vEmKfgqp7yDtrOCbWmtYuzRwIm

#### Predicting House Price using DNN



https://colab.research.google.com/drive/1tJztaaOIxbk\_VuPKm8NpN7Cp\_XABqyPQ

#### **Final Results**



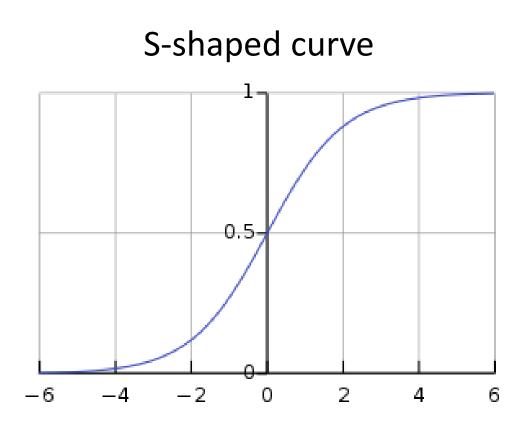
#### Logistic Regression

Sigmoid function

$$S(x) = \frac{e^x}{e^x + 1} = \frac{1}{1 + e^{-x}}$$

• Derivative of Sigmoid

S(x) = S(x)(1 - S(x))



https://en.wikipedia.org/wiki/Sigmoid\_function

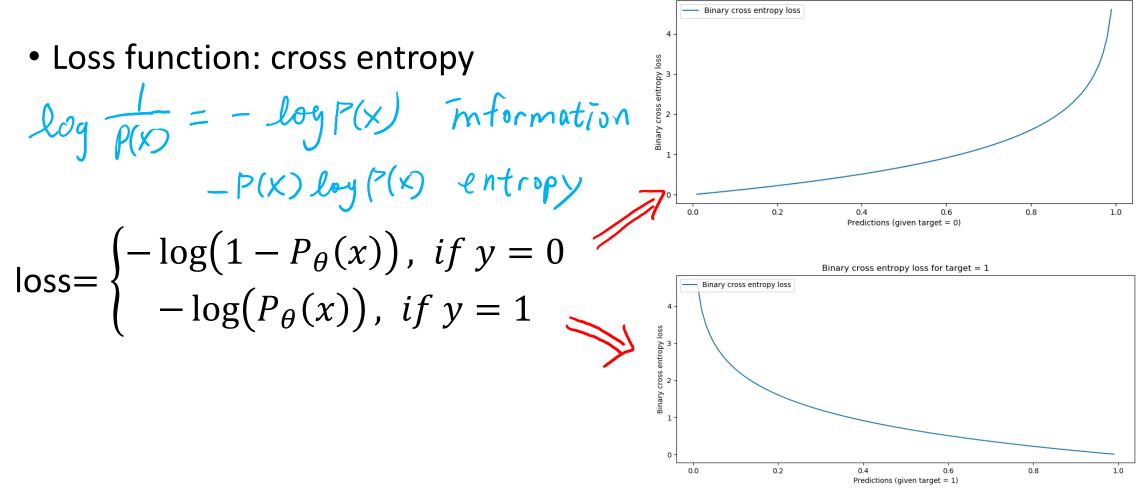
#### **Decision Boundary**

• Binary classification with decision boundary t

$$P(x,w) = P_{\theta}(x) = \frac{1}{1 + e^{-(w^T x + b)}}$$

$$y' = \begin{cases} 0, & P_{\theta}(x) < t \\ 1, & P_{\theta}(x) \ge t \end{cases}$$

#### Cross Entropy Loss



Binary cross entropy loss for target = 0

#### Cross Entropy Loss

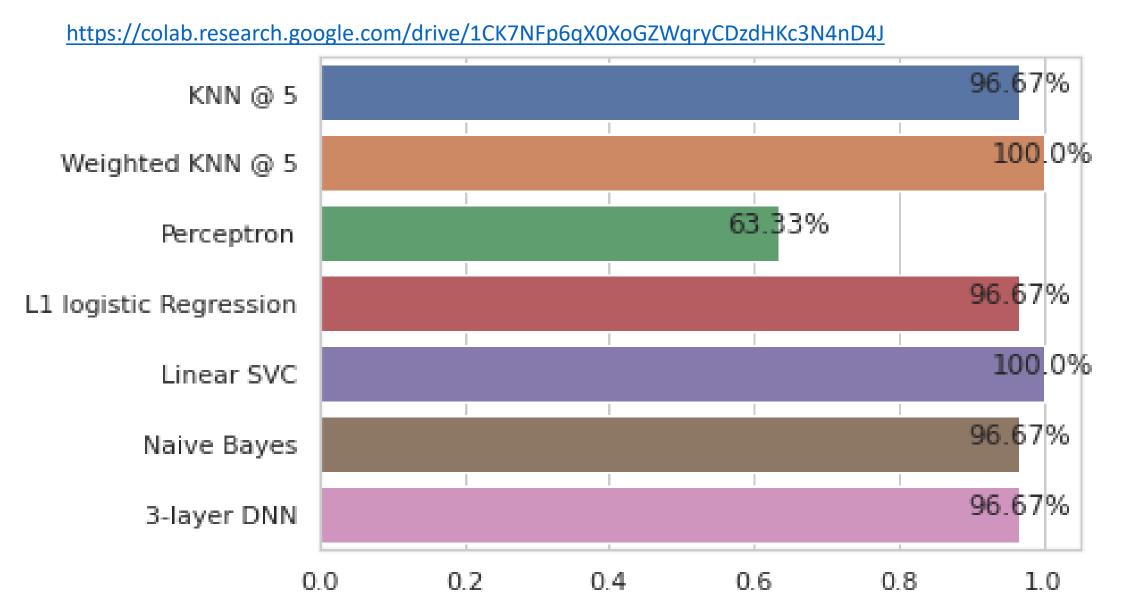
Loss function: cross entropy

$$\operatorname{loss} = \begin{cases} -\log(1 - P_{\theta}(x)), & \text{if } y = 0\\ -\log(P_{\theta}(x)), & \text{if } y = 1 \end{cases}$$

$$\Rightarrow L_{\theta}(\mathbf{x}) = -y \log (P_{\theta}(\mathbf{x})) + -(1-y) \log (1-P_{\theta}(\mathbf{x}))$$

$$\nabla L_W(\mathbf{x}) = -(y - P_\theta(\mathbf{x}))\mathbf{x}$$

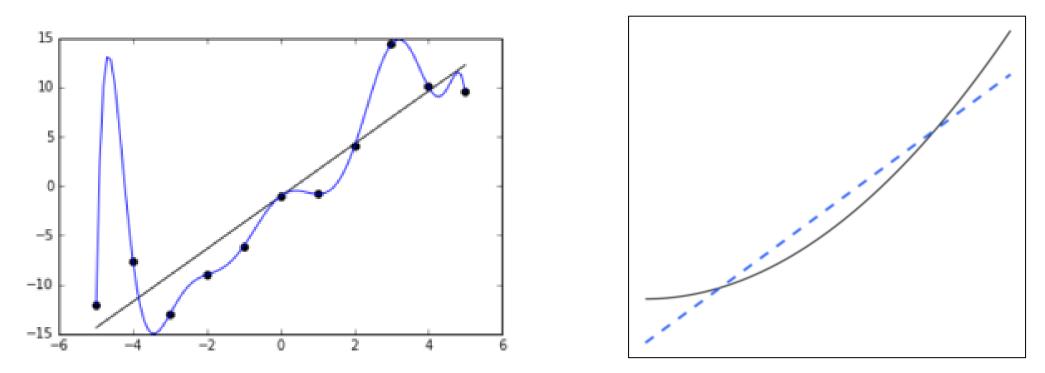
### Classifier Evaluation on Iris dataset



# **Overfitting & Underfitting**

Overfitting

#### Underfitting



https://en.wikipedia.org/wiki/Overfitting

#### Neural Network Urban Legend: Detecting Tanks

• Detector learned the illumination of photos



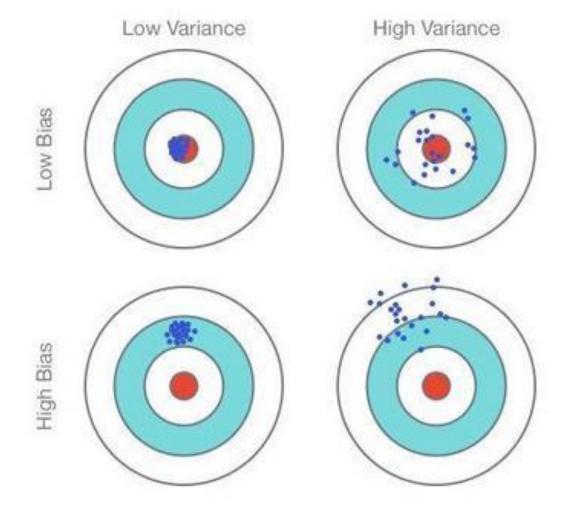


#### Prediction Error: Bias and Variance Trade-off

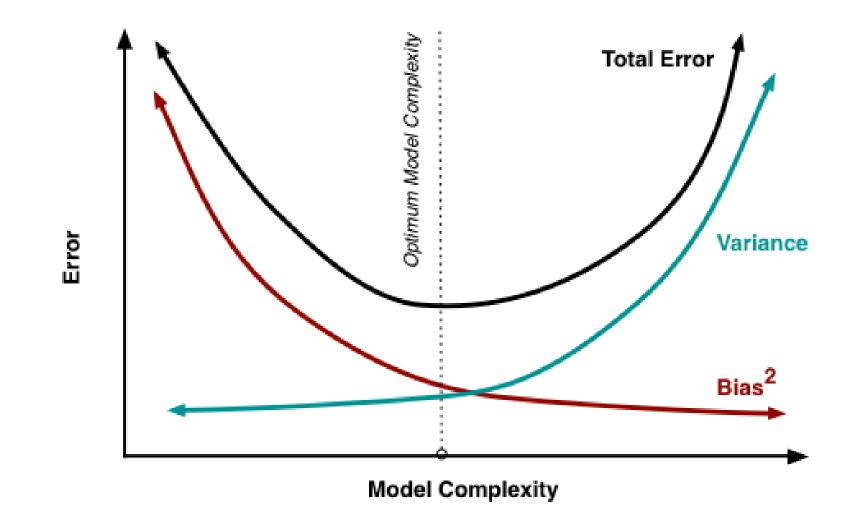
• Model with high variance overfits to training data and does not generalize on unseen test data

$$Err(x) = E\left[(Y - \hat{f}(x))^2
ight]$$
 $Err(x) = \left(E[\hat{f}(x)] - f(x)
ight)^2 + E\left[\left(\hat{f}(x) - E[\hat{f}(x)]
ight)^2
ight] + \sigma_e^2$ 
 $Err(x) = ext{Bias}^2 + ext{Variance} + ext{Irreducible Error}$ 

https://towardsdatascience.com/understanding-the-bias-variance-tradeoff-165e6942b229

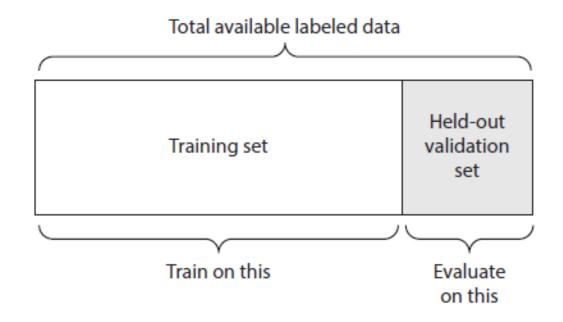


#### Model Selection



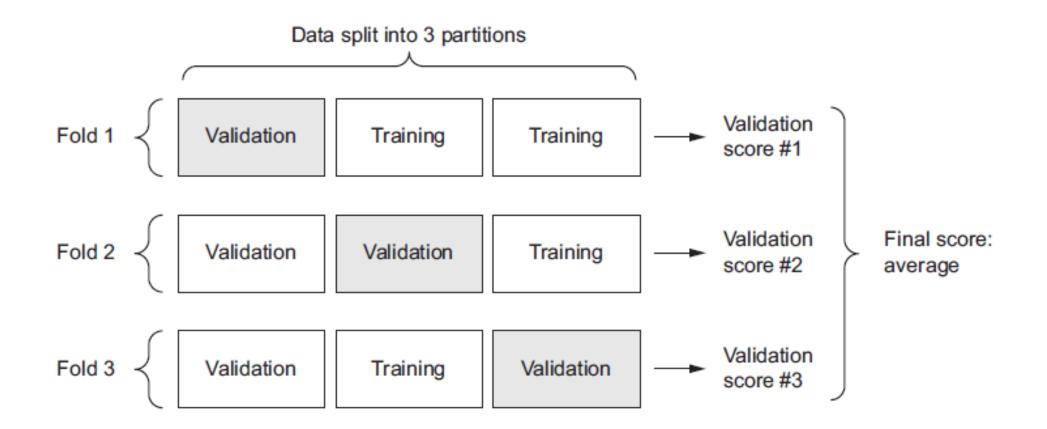
#### Training, Validation, Testing

- Never leak test data information into our model
- Tuning the *hyperparameters* of our model on validation dataset
- Better generalize the model to future unseen data



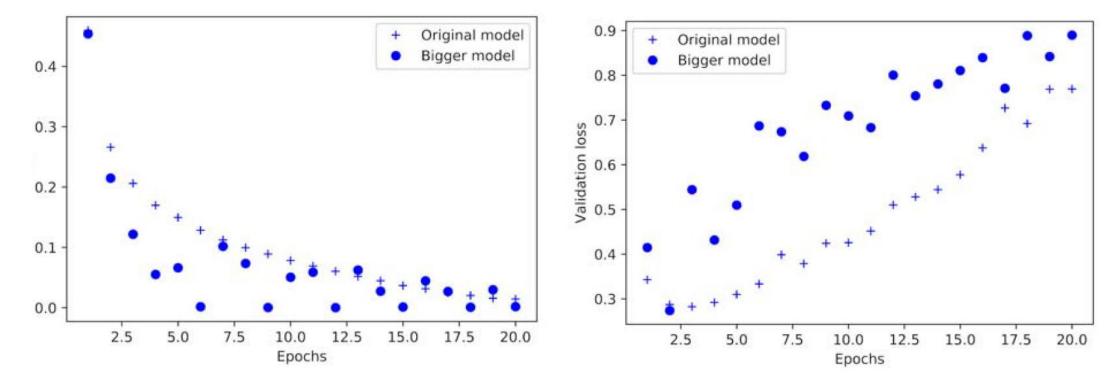
#### **K-Fold Cross Validation**

• Lower the variance of validation set



### Overfitting with Bigger Model

• The more capacity the network has, the more quickly it can model the training data, but the more susceptible it is to overfitting

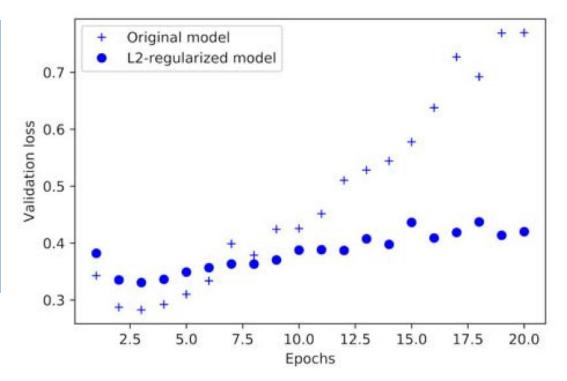


#### Training Loss

Validation Loss

### Regularization

• Weight regularization – L1 and L2 norm



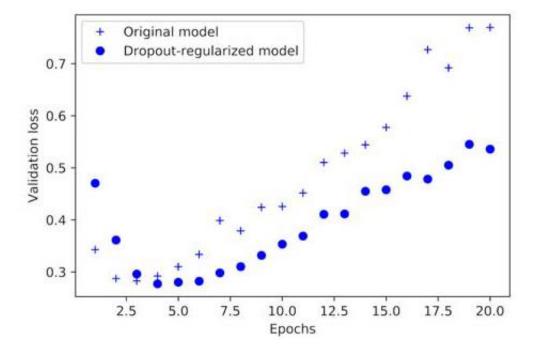
#### Dropout

- randomly dropping out (setting to zero) a number of output features of the layer during training
- Dropout applied to an activation matrix at training time
- At test time, the activation matrix is unchanged

0.3	0.2	1.5	0.0		0.0	0.2	1.5	0.0
0.6	0.1	0.0	0.3	50% dropout	0.6	0.1	0.0	0.3
0.2	1.9	0.3	1.2		0.0	1.9	0.3	0.0
0.7	0.5	1.0	0.0		0.7	0.0	0.0	0.0

#### Adding Dropout to the IMDB Network

```
model = models.Sequential()
model.add(layers.Dense(16, activation='relu', input_shape=(10000,)))
model.add(layers.Dropout(0.5))
model.add(layers.Dense(16, activation='relu'))
model.add(layers.Dropout(0.5))
model.add(layers.Dense(1, activation='sigmoid'))
```



### Machine Learning Workflow

- 1. Defining the problem and assembling a dataset
- 2. Choosing a measure of success (accuracy, MAE,...)
- 3. Deciding on an evaluation protocol (squared loss, cross entropy,...)
- 4. Preparing your data (normalization)
- 5. Developing a model that does better than a baseline
- 6. Scaling up: developing a model that overfits
- 7. Regularizing your model and tuning hyperparameters

# Deep Learning for Classification & Regression

• Choosing the right last-layer activation and loss function

Problem type	Last-layer activation	Loss function
Binary classification	sigmoid	binary_crossentropy
Multiclass, single-label classification	softmax	categorical_crossentropy
Multiclass, multilabel classification	sigmoid	binary_crossentropy
Regression to arbitrary values	None	mse
Regression to values between 0 and 1	sigmoid	mse <b>or</b> binary_crossentropy

# Metrics: Accuracy vs. Precision in Binary Classification

# **Confusion Matrix**

• Binary classification (positive vs. negative)

		True condition Y (Labels)		
	Total population	Condition positive	Condition negative	
Predicted	Predicted condition positive	True positive	<b>False positive,</b> Type I error	
condition	Predicted condition negative	<b>False negative,</b> Type II error	True negative	

https://en.wikipedia.org/wiki/Confusion\_matrix

### **Confusion Matrix**

#### https://en.wikipedia.org/wiki/Confusion\_matrix

		True con					
	Total population	Condition positive	Condition negative	$\frac{\text{Prevalence}}{\sum \text{ Condition positive}} = \frac{\Sigma \text{ Condition positive}}{\Sigma \text{ Total population}}$	<u>Σ True positiv</u>	racy (ACC) = ve + Σ True negative al population	
Predicted	Predicted condition positive	True positive	<b>False positive</b> , Type I error	Positive predictive value (PPV), Precision = $\Sigma$ True positive $\overline{\Sigma}$ Predicted condition positive	Σ Fa	False discovery rate (FDR) = $\Sigma$ False positive $\Sigma$ Predicted condition positive	
condition	Predicted condition negative	<b>False negative</b> , Type II error	True negative	False omission rate (FOR) = $\Sigma$ False negative $\Sigma$ Predicted condition negative	Negative predictive value (NPV) = $\frac{\Sigma \text{ True negative}}{\Sigma \text{ Predicted condition negative}}$		
		True positive rate (TPR), Recall, Sensitivity, probability of detection, Power = $\frac{\Sigma \text{ True positive}}{\Sigma \text{ Condition positive}}$	False positive rate (FPR), Fall-out, probability of false alarm = $\frac{\Sigma \text{ False positive}}{\Sigma \text{ Condition negative}}$	Positive likelihood ratio (LR+) = $\frac{\text{TPR}}{\text{FPR}}$	Diagnostic odds ratio	F <sub>1</sub> score =	
		False negative rate (FNR), Miss rate = $\frac{\Sigma \text{ False negative}}{\Sigma \text{ Condition positive}}$	Specificity (SPC), Selectivity, True negative rate (TNR) = $\frac{\Sigma \text{ True negative}}{\Sigma \text{ Condition negative}}$	Negative likelihood ratio (LR-) = <u>FNR</u> TNR		2 · <u>Precision · Recall</u> Precision + Recall	

#### Coronavirus Confusion

• Error rate = 80%

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- Did the medical staff misuse the test kit?
- Can we just swap positive class and negative class and achieve 80% accuracy?

植試劑

#### 80% of Rapid COVID-19 Tests the Czech Republic Bought From China are Wrong

🎴 by prague morning 🕔 march 26, 2020 🕔 % 1K shares 🖓 2 minute read



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CZECHIA

Up to 80 percent of the 300,000 rapid coronavirus test kits ordered that the Czech Republic ordered from China are not working properly, according to regional hygienists who have tried the tests.

The test kits, worth 54 million korunas ( $\leq$ 1.83 million), show false positive as well as negative results. This is because rapid tests cannot reliably detect infection in its initial phase.

https://praguemorning.cz/80-pf-rapid-covid-19-tests-the-czech-republic-bought-from-china-are-wrong/

### Popular Metrics

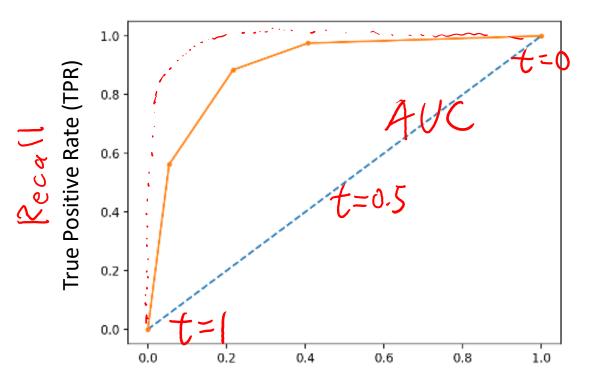
#### Notations

-P: positive samples, N: negative samples, P': predicted positive samples, TP: true positives, TN: true negatives

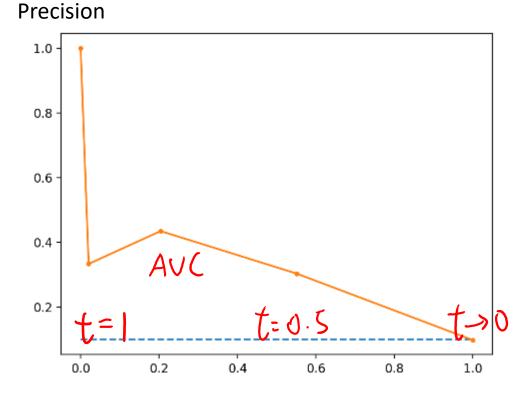
- Recall =  $\frac{TP}{P}$ • Precision =  $\frac{TP}{P'}$ • Accuracy =  $\frac{\hat{TP} + TN}{-}$
- F1 score =  $-\frac{1}{1}$  $\frac{1}{recall} + \frac{1}{precision}$
- Miss rate = false negative rate = 1 recall

#### Evaluate Decision Boundary t

 ROC (Receiver Operating Characteristic) Curve



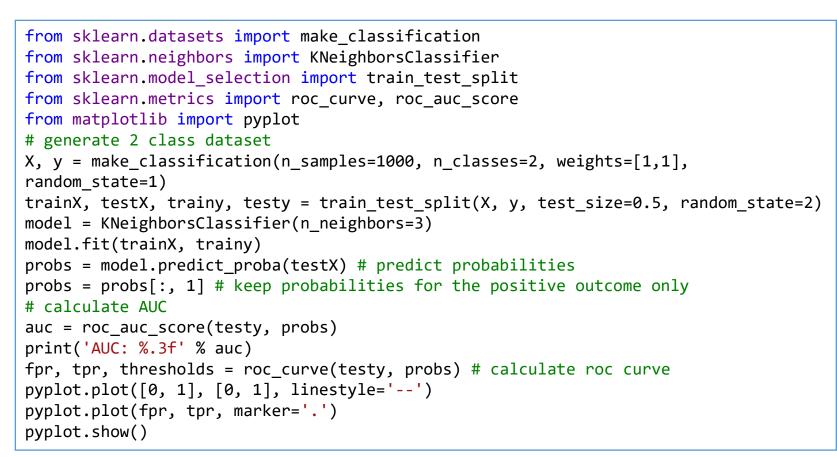
• Precision-Recall (PR) Curve

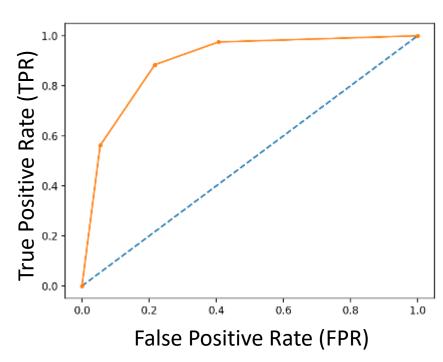


False Positive Rate (FPR)

# ROC (Receiver Operating Characteristic)

- Evaluate binary classifier's ability
- Plot the true positive rate (TPR) against the false positive rate (FPR) at various thresholds (decision boundaries)
- Use area under curve (AUC) to evaluate performance

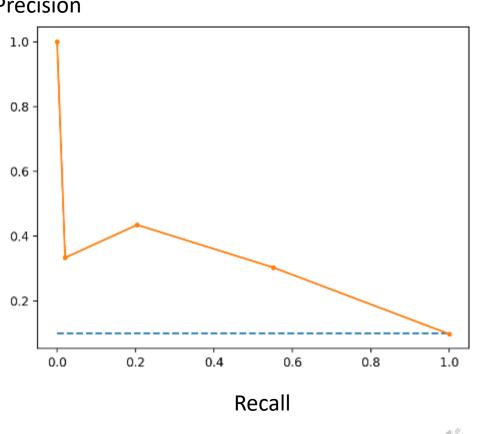




# Precision-Recall (PR) Curve

- Plot Precision vs. Recall
- Popular in information retrieval

```
Precision
from sklearn.datasets import make classification
from sklearn.neighbors import KNeighborsClassifier
                                                                           1.0 -
from sklearn.model_selection import train_test_split
from sklearn.metrics import precision_recall_curve
from matplotlib import pyplot
                                                                           0.8
# generate 2 class dataset
X, y = make classification(n samples=1000, n classes=2,
weights=[0.9,0.09], random state=1)
                                                                           0.6
trainX, testX, trainy, testy = train_test_split(X, y, test_size=0.5,
random state=2)
# fit a model
                                                                           0.4
model = KNeighborsClassifier(n neighbors=3)
model.fit(trainX, trainy)
probs = model.predict proba(testX)[:, 1]
                                                                           0.2 -
# predict class values
yhat = model.predict(testX)
                                                                                      0.2
                                                                               0.0
                                                                                              0.4
# Calculate precision recall curve
precision, recall, thresholds = precision recall curve(testy, probs)
pyplot.plot(recall, precision, marker='.')
pyplot.show()
```



#### ROC Curve & PR Curve

- ROC curves should be used when there are roughly equal numbers of observations for each class
- Precision-Recall curves should be used when the data are imbalance and we only care about the positive class
- ROC may be harmful
  - ROC curves can present an overly optimistic view of an algorithm's performance if there is a large skew in the class distribution

#### Accuracy

• Example: Classifying tumors as malignant or benign (Google Machine Learning Crash Course )

True Positive (TP):	False Positive (FP):
Reality: Malignant	Reality: Benign
ML model predicted: Malignant	ML model predicted: Malignant
Number of TP results: 1	Number of FP results: 1
False Negative (FN):	True Negative (TN):
Reality: Malignant	Reality: Benign
ML model predicted: Benign	ML model predicted: Benign
Number of FN results: 8	Number of TN results: 90

 $\label{eq:accuracy} {\rm Accuracy} = \frac{TP+TN}{TP+TN+FP+FN} = \frac{1+90}{1+90+1+8} = 0.91$ 

#### https://developers.google.com/machine-learning/crash-course/classification/accuracy

#### Accuracy

• Example: Classifying tumors as malignant or benign (Google Machine Learning Crash Course )

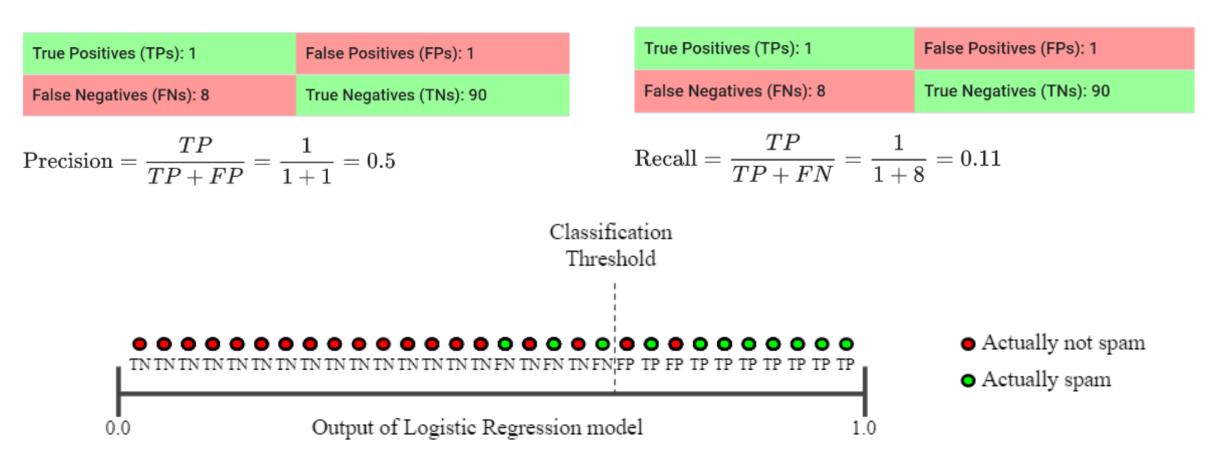
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Number of TP results: 1	Number of FP results: 1
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#### https://developers.google.com/machine-learning/crash-course/classification/accuracy

#### Precision and Recall

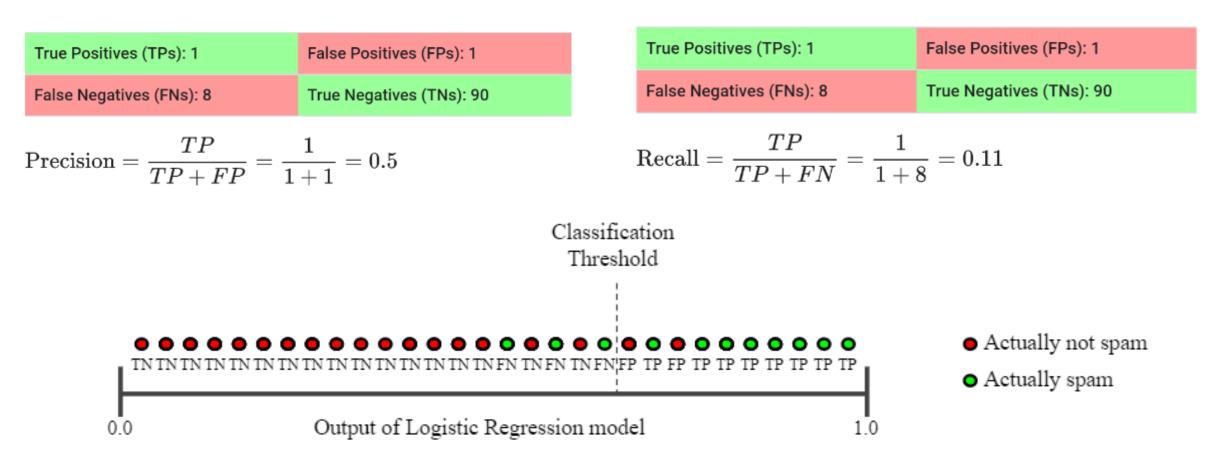
• Classifying tumors as malignant or benign



https://developers.google.com/machine-learning/crash-course/classification/precision-and-recall

#### Precision and Recall

• Classifying tumors as malignant or benign



https://developers.google.com/machine-learning/crash-course/classification/precision-and-recall

# Useful Things to Know about Machine Learning

- 1. It's generalization that counts
- 2. Data alone is not enough
- 3. Overfitting has many faces
- 4. Intuition fails in high dimensions
- 5. Theoretical guarantees are not what they seem
- 6. More data beats a cleverer algorithm
- 7. Learn many models, not just one



Pedro Domingos, "A Few Useful Things to Know about Machine Learning," Commun. ACM, 2012

# Key Takeaways

- Classification
  - KNN, Linear Classification, SVM, Naïve Bayes, Logistic regression
- Regression
  - Linear regression, Ridge Regression, Lasso Regression, Elastic Net

#### • Avoid Overfitting

Regularization, k-Fold Cross validation

#### Confusion Matrix

- Accuracy vs. Precision, Recall
- ROC curve, PR curve and AUC

### References

- Francois Chollet, "Deep Learning with Python", Chapter 4
- Pedro Domingos, "A Few Useful Things to Know about Machine Learning," Commun. ACM, 2012
- <u>https://ml-cheatsheet.readthedocs.io/en/latest/index.html</u>
- <u>https://machinelearningmastery.com/roc-curves-and-precision-recall-curves-for-classification-in-python/</u>
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- <a href="https://en.wikipedia.org/wiki/Naive\_Bayes\_classifier">https://en.wikipedia.org/wiki/Naive\_Bayes\_classifier</a>
- <a href="https://en.wikipedia.org/wiki/Confusion\_matrix">https://en.wikipedia.org/wiki/Confusion\_matrix</a>
- <a href="https://developers.google.com/machine-learning/crash-course/classification/accuracy">https://developers.google.com/machine-learning/crash-course/classification/accuracy</a>