

Scikit-learn/sklearn

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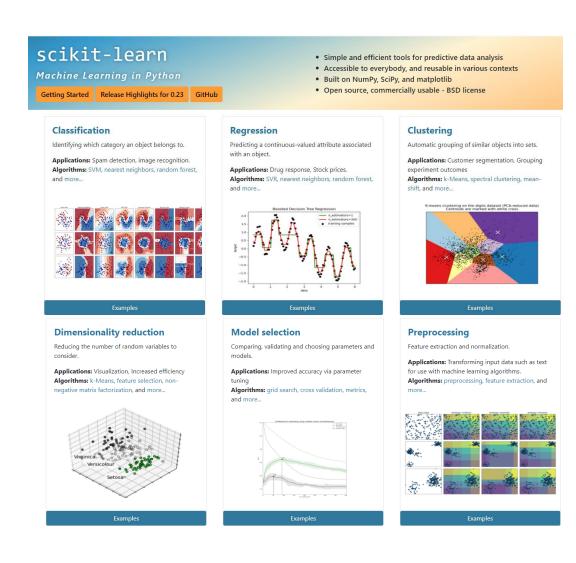
Topics

- Sklearn
- Iris classification in sklearn (binary classification)
- Plot of decision boundary
- Split data into training and testing
- A perceptron for multiclass classification/one-vs.-rest (OVR) approach
- Logistic regression classifier
- Logistic regression for IRIS dataset
- Separable and inseparable dataset
- Confusion matrix
- Multilayer neural network (MLP)
- MLP classifier in sklearn
- MLP classifier in Keras/model.summary()



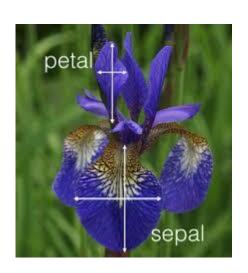
Scikit-learn

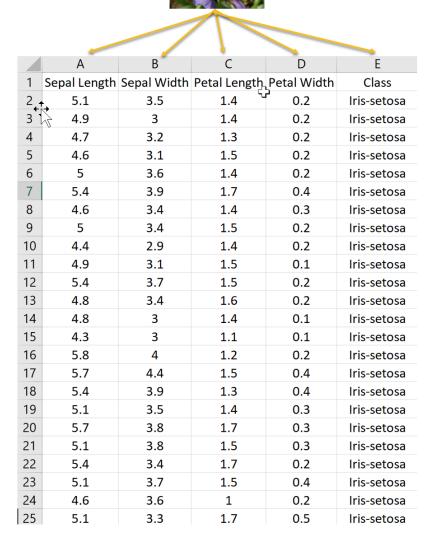
- Is a Python
 machine learning
 library offering a
 vast collection of
 machine learning
 algorithms from
 data processing,
 model selection,
 model training, fine
 tune, and
 evaluation.
- sciPy Toolkit developed by Google.
- https://scikitlearn.org/stable/





Iris dataset classification







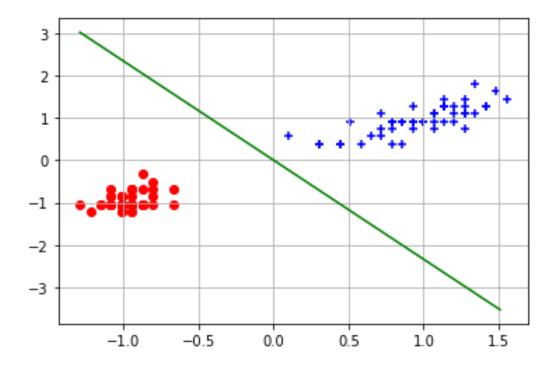
Implementation

```
# implement a perceptron to classifiy iris using Scikit-Learn
import numpy as np
from sklearn import datasets
import matplotlib.pyplot as plt
from sklearn.linear model import Perceptron
from sklearn.preprocessing import StandardScaler
iris = datasets.load_iris()
x = iris.data[0:100,[2,3]] # petal length and petal width
y = iris.target[0:100]
print(x[0:5,:])
print('Class labels:', np.unique(y))
labels = ['setosa', 'versicolor', 'virginica']
# standarization
sc = StandardScaler()
sc.fit(x) # to estimate mean and standard deviation
xstd = sc.transform(x)
print(xstd[0:5,:])
# import a model
#!pip install ....
model = Perceptron(max_iter=100, eta0=0.01, random_state=1)
model.fit(xstd, y)
y pred=model.predict(x)
print(model.coef , model.intercept )
error=np.mean((y-y pred)**2)
print(error)
```



```
#plot decision line
plt.scatter(xstd[y==0,0],xstd[y==0,1], color='red', marker='o',label=labels[0])
plt.scatter(xstd[y==1,0],xstd[y==1,1], color='blue', marker='+',label=labels[1])

xx = np.arange(xstd[:,0].min(),xstd[:,0].max(),0.1)
yy = -model.coef_[0,0]/model.coef_[0,1] * xx - model.intercept_/model.coef_[0,1]
plt.plot(xx,yy,'g-')
plt.grid()
plt.show()
```





Plot decision region

```
# Another way for plotting the decision line/region
def plot decision region(x, y):
 x1 = np.arange(x[:,0].min()-1, x[:,0].max()+1,0.1)
 x2 = np.arange(x[:,1].min()-1,x[:,1].max()+1, 0.1)
 xg1, xg2 = np.meshgrid(x1, x2)
  #print(xg2.flatten().T) # ravel()/flatten() returns a flattened one-dimensional array
  z = model.predict(np.array([xg1.ravel(), xg2.ravel()]).T)
  plt.contourf(xg1, xg2, z.reshape(xg1.shape))
  #plot the scatter plot of the dataset
  plt.scatter(x[y==0,0],x[y==0,1],color='yellow', marker='o')
  plt.scatter(x[y==1,0],x[y==1,1],color='black', marker='d')
  plt.xlabel('x1')
  plt.ylabel('x2')
  plt.show()
                                                 2
plot_decision_region(x,y)
                                                 0
                                                -1
                                                                              x1
```



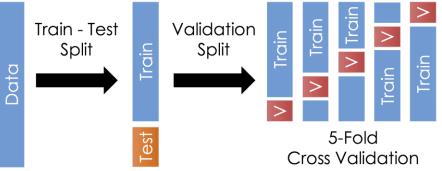
Splitting the dataset into training and testing sets

 It is important to evaluate the trained model performance on unseen data

```
# split the dataset into training and testing sets using (train_test_split) function
# this function shuffels the training sets internally before splitting
from sklearn.model_selection import train_test_split

xtarin, xtest, ytrain, ytest = train_test_split(x, y, test_size=0.3, random_state=1)
print(ytest)
print('proportion of class labels in train and test sets:\n',np.bincount(ytrain), np.bincount(ytest))
#stratify: to have the same proportional of class labels in both training and testing sets (balanced sets)
xtarin, xtest, ytrain, ytest = train_test_split(x, y, test_size=0.3, random_state=1, stratify=y)
print('proportion of class labels in train and test sets:\n',np.bincount(ytrain), np.bincount(ytest))
```

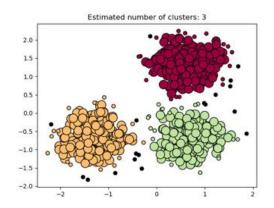
```
[1 1 0 1 1 0 0 1 1 1 1 0 1 1 1 0 0 0 1 0 0 1 1 1 0 0 1 0 0 0] proportion of class labels in train and test sets:
[36 34] [14 16] proportion of class labels in train and test sets:
[35 35] [15 15]
```

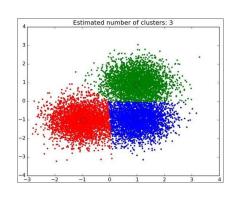


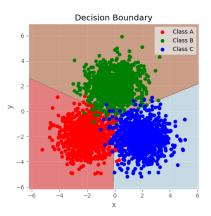


Linear models for multiclass classification

- Many linear classification models are for binary classification only, and do not extend naturally to the multiclass case (except for logistic regression)
- A common technique to extend a binary classification algorithm to a multiclass classification algorithm is the one-vs.-rest (OVR) approach.
- In OVR approach, a binary model is learned for each class that tries to separate that class from all of the other classes, resulting in as many binary models as there are classes.
- Having one binary classifier per class results in having one vector of coefficients (w) and one intercept (b) for each class.
- Sklearn provides a sample dataset generator to help you to create fast and easy to use a custom dataset.
- make_blobs generates isotropic Gaussian blobs for clustering.

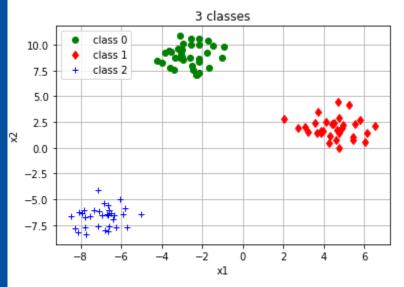


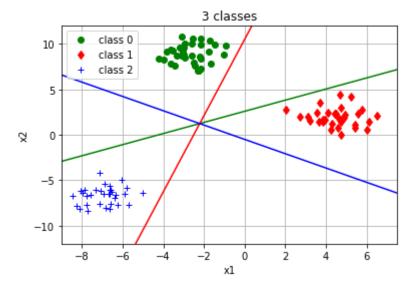






Make_blobs dataset: logistic regression





one-vs.-rest (OVR) approach

	w1	w2	b
Class 0	-0.38	0.63	-1,63
Class 1	0.74	-0.18	1.86
Class 2	-0.36	-0.45	-0.23



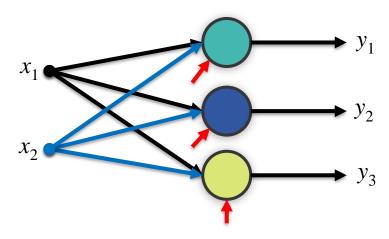
```
#make blobs dataset for multiclass classification
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import make blobs
from sklearn.linear model import LogisticRegression
                                                        # build a model
                                                        model = LogisticRegression()
                                                        model.fit(x,y)
x, y = make blobs(random state=42)
                                                        y pred = model.predict(x)
print(y.shape)
                                                        print(model.coef )
print(np.bincount(y))
                                                        print(model.intercept )
plt.plot(x[y==0,0], x[y==0,1], 'go', label='class 0')
plt.plot(x[y==1,0], x[y==1,1], 'rd', label='class 1')
plt.plot(x[y==2,0], x[y==2,1], 'b+', label='class 2')
plt.xlabel('x1')
                                   # plot decision lines
plt.ylabel('x2')
                                   plt.plot(x[y==0,0], x[y==0,1], 'go', label='class 0')
plt.title('3 classes')
                                   plt.plot(x[y==1,0], x[y==1,1], 'rd', label='class 1')
plt.legend(loc='upper left')
                                   plt.plot(x[y==2,0], x[y==2,1], 'b+', label='class 2')
plt.grid()
plt.show()
                                   x1 = np.linspace(-10, 10)
                                   color = ['g-','r-','b-']
                                   for coef, intercept, color in zip(model.coef_, model.intercept_, color):
                                     x2 = -(coef[0]*x1+intercept)/coef[1]
                                     plt.plot(x1, x2, color)
                                   plt.xlabel('x1')
                                   plt.ylabel('x2')
                                   plt.title('3 classes')
                                   plt.legend(loc='upper left')
                                   plt.ylim([-12.0, 12.0])
                                   plt.xlim([-9.0, 7.5])
                                   plt.grid()
                                   plt.show()
```



Extending a perceptron for multiclass classification

Iris dataset:

- 2 input feature: [sepal length (cm), sepal width (cm), petal length (cm), and petal width (cm)]
- 3 classes (setosa, versicolor, and virginica)



Weights		Bias
-0.00220757	-0.00237113	-0.002
0.00540865	-0.00210787	-0.002
0.00628849	0.00683785	-0.008



Implementation

```
# In this example we will use a perceptron from sklearn to classify three iris classes (setosa, versicolor, and virginica)
# upload python liberaries
import numpy as np
from sklearn import datasets
import matplotlib.pyplot as plt
# upload your iris datset
iris = datasets.load iris()
# select two features
x = iris.data[:,[2,3]] # petal length and petal width
y = iris.target
print(np.unique(y))
print(iris.target names)
print(iris.feature names)
[0 1 2]
['setosa' 'versicolor' 'virginica']
['sepal length (cm)', 'sepal width (cm)', 'petal length (cm)', 'petal width (cm)']
# standardize your feature vector
from sklearn.preprocessing import StandardScaler
sc=StandardScaler()
sc.fit(x)
xstd=sc.transform(x)
print(x[1:2,:], xstd[1:2,:])
[[1.4 0.2]] [[-1.34022653 -1.3154443 ]]
#split dataset into training and testing sets
from sklearn.model selection import train test split
x = xstd
xtrain, xtest, ytrain, ytest = train_test_split(x,y,test_size=0.3, random_state=1,stratify=y)
print(np.bincount(ytrain), np.bincount(ytest))
[35 35 35] [15 15 15]
```

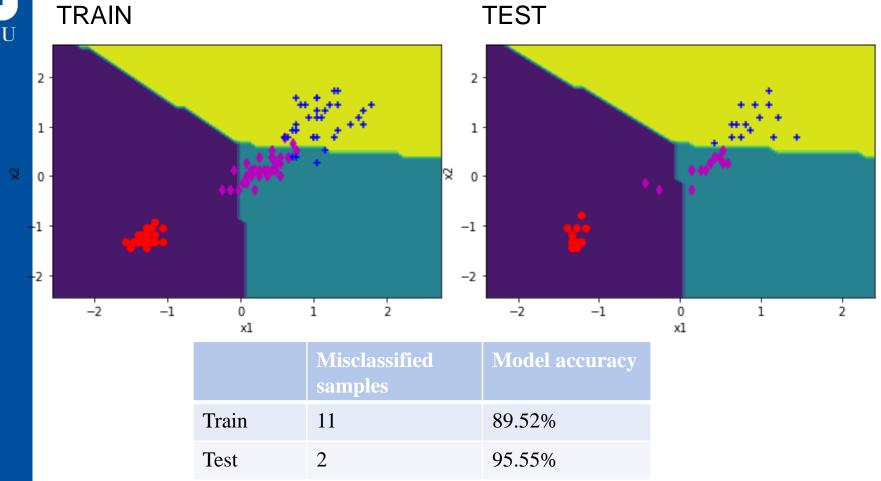


```
# select a model for training
from sklearn.linear model import Perceptron
model = Perceptron(max iter=20, eta0=0.002, random state=1)
model.fit(xtrain, ytrain)
vpred = model.predict(xtest)
missclassified = (ytest != ypred).sum()
print('missclassified samples = ', missclassified)
print('Model accuracy = ', (len(ytest)-missclassified)/len(ytest)*100)
print('Model accuracy = ', model.score(xtest, ytest)*100, '%')
print('weights: \n', model.coef )
print('bias : \n', model.intercept )
missclassified samples = 2
Model accuracy = 95.555555555556
Model accuracy = 95.555555555556 %
weights:
[[-0.00220757 -0.00237113]
[ 0.00540865 -0.00210787]
[ 0.00628849  0.00683785]]
bias :
[-0.002 -0.002 -0.008]
```

```
# plot decision regions in 2D space
x1 = np.arange(x[:,0].min()-1, x[:,0].max()+1, 0.1)
x2 = np.arange(x[:,1].min()-1, x[:,1].max()+1, 0.1)
x1g, x2g = np.meshgrid(x1, x2)
z = model.predict(np.array([x1g.flatten(), x2g.flatten()]).T)
plt.contourf(x1g, x2g, z.reshape(x1g.shape))
plt.scatter(x[y==0,0],x[y==0,1],color='red', marker='o')
plt.scatter(x[y==1,0],x[y==1,1],color='m', marker='d')
plt.scatter(x[y==2,0],x[y==2,1],color='blue', marker='+')
plt.xlabel('x1')
plt.ylabel('x2')
plt.show()
```



Decision regions

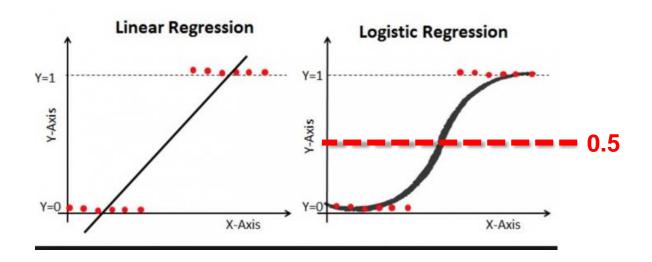


- Notes that a perceptron will never converge if the classes are not perfectly linearly separable.
- In iris dataset, there will be always at least one misclassified sample present in each epoch.



Modeling class probabilities via logistic regression

- Logistic regression is one of the most popular machine learning algorithms in industry.
- It is a classification model that performs very well for linearly separable binary classes (YES/NO, TRUE/FALSE) instead of predicting something continues.
- Instead of fitting a line to the data, logistic regression fits an 'S' shaped logistic function
- Although it gives the probability as a number between 0 and 1, it is usually used for classification using a threshold value.
- Examples: ham/spam email, fraudulent (yes/no) online transactions, etc.





Logistic regression

$$y \in \begin{cases} 1 & Positive Class (Yes / True) \\ 0 & Negative Class (No / False) \end{cases}$$

It can also be extended into multiclass classification

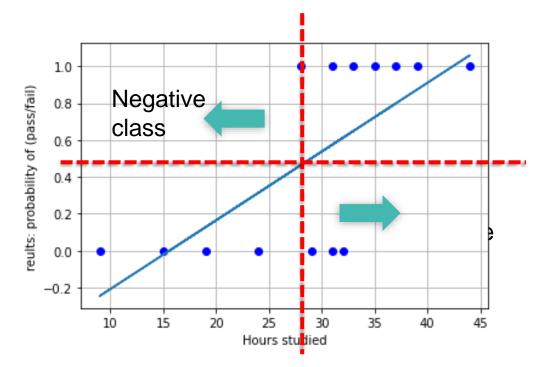
$$y \in \{0,1,2,3\}$$



It is all about likelihood of an event: hours studied to pass example

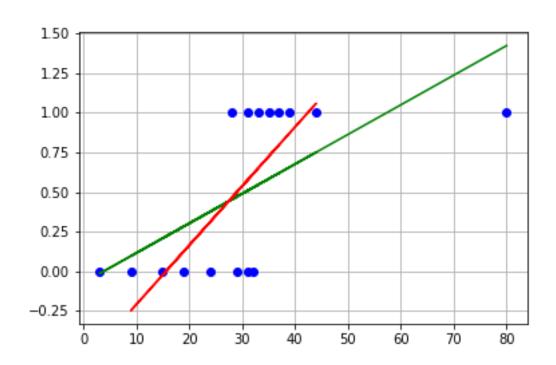
Hours studied	Exam result (1=pass, 0=fail)
29	0
15	0
33	1
28	1
39	1
44	1
31	1
19	0
9	0
24	0
32	0
31	0
37	1
35	1

Can we fit a straight line?





Adding outliers





Fitting a logistic regression model: fit a sigmoid/logistic function

$$net = w^T x = wx + b = 0.4x - 12.05$$

$$g(net) = \frac{1}{1+e^{-net}} = \frac{e^{net}}{1+e^{net}}$$

from sklearn.linear_model import LogisticRegression

```
# fit a logistic regression model
logreg = LogisticRegression()
```

fit the model with data
logreg.fit(x.reshape(-1,1),y.reshape(-1,1))

print(logreg.coef_)
print(logreg.intercept)

xx = np.arange(5, 50,0.1).reshape(-1,1)
y pred = logreg.predict(xx)

def sig(x):
 return(np.exp(x)/(1+np.exp(x)))

net = logreg.intercept_ + logreg.coef_ * xx

plt.plot(x,y,'bd')
plt.plot(xx, sig(net), 'g-')
plt.plot(xx,y_pred,'r-',LineWidth=3.0)

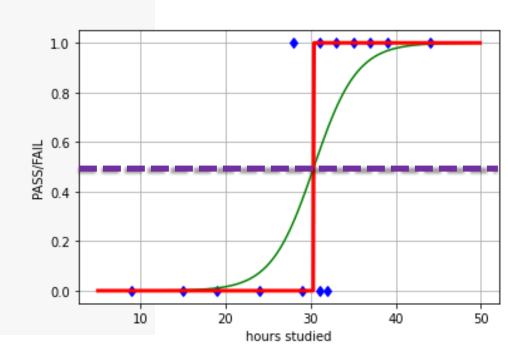
plt.show()

plt.grid()

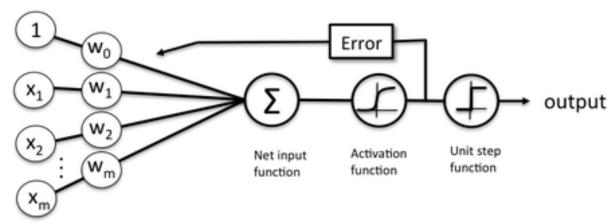
Hypothesis:

$$y = 1$$
 when $g(w^T x) \ge 0.5$

$$y = 0$$
 when $g(w^{T}x) < 0.5$



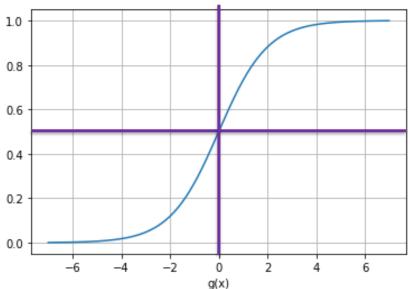




Schematic of a logistic regression classifier.



Decision boundary



Y = 1

$$Y = 0$$

$$p(y=1) + p(y=0) = 1$$

$$p(y=1) = 1 - p(y=0)$$

$$y = g(x) \to 1 \text{ as } x \to \infty (1/(1+0) = 1)$$

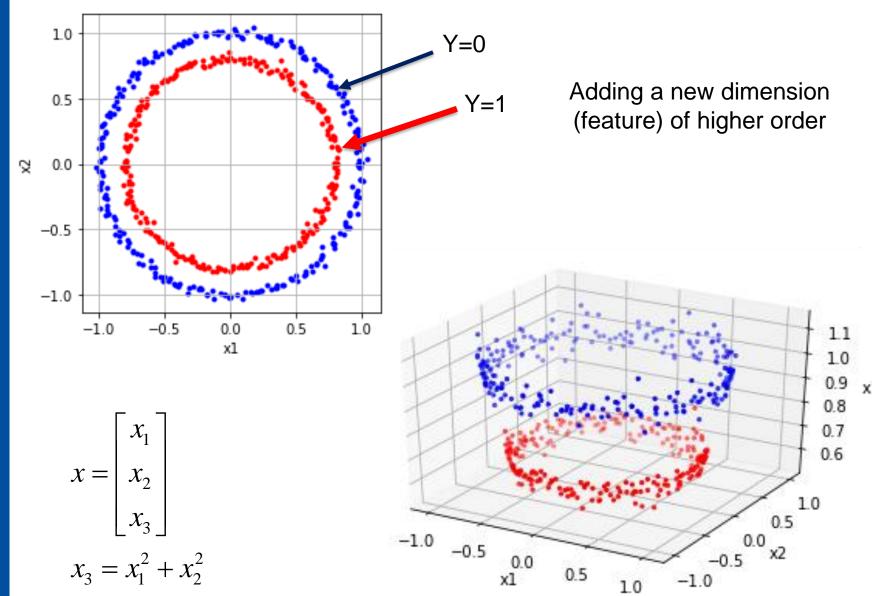
$$y = g(x) \to 0 \text{ as } x \to -\infty (1/(1+\infty) = 0)$$

Hypothesis:

$$y = 1$$
 when $g(w^T x) \ge 0.5 \rightarrow w^T x \ge 0$
 $y = 0$ when $g(w^T x) < 0.5 \rightarrow w^T x < 0$

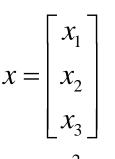


Linearly inseparable dataset



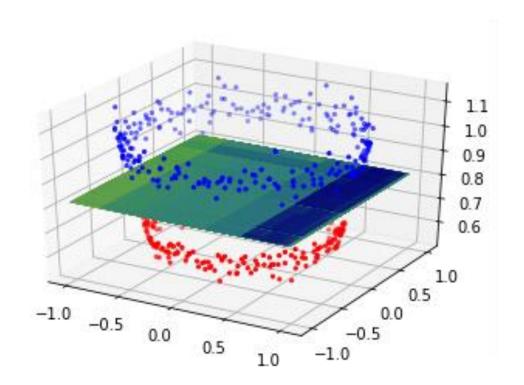


Fit a logistic regression model



$$x_3 = x_1^2 + x_2^2$$

$$net = w_1 x_1 + w_2 x_2 + w_3 x_3 + b$$





Logistic regression: Iris example

```
# implement a perceptron to classifiy iris using Scikit-Learn
import numpy as np
from sklearn import datasets
import matplotlib.pyplot as plt
from sklearn.linear_model import LogisticRegression
from sklearn.preprocessing import StandardScaler
```

```
iris = datasets.load iris()
x = iris.data
y = iris.target
print(x[0:5,:])
print('Class labels:', np.unique(y))
labels = ['setosa', 'versicolor', 'virginica']
# standarization
sc = StandardScaler()
sc.fit(x) # to estimate mean and standard deviation
x = sc.transform(x)
[[5.1 3.5 1.4 0.2]
[4.9 3. 1.4 0.2]
 [4.7 3.2 1.3 0.2]
 [4.6 3.1 1.5 0.2]
 [5. 3.6 1.4 0.2]]
Class labels: [0 1 2]
# import a model
#!pip install ....
model = LogisticRegression(C=1, random_state=1)
model.fit(x, y)
y_pred=model.predict(x)
print(model.coef_, model.intercept_)
error=np.mean((y-y pred)**2)
print(error)
print(np.sum(y!=y pred))
[[-1.07404149 1.16006342 -1.93062866 -1.81168873]
  0.58780051 -0.36182377 -0.36346274 -0.82619289]
 0.02666666666666667
```



Assignment

- Implement logistic regression
- Convert Adaline into a logistic regression
- Training set (m-examples)

$$\{(x^{(1)}, y^{(1)}), (x^{(2)}, y^{(2)}), \dots, (x^{(m)}, y^{(m)})\}, y \in \{0, 1\}$$

· Hypothesis:

$$\phi_w(x) = \frac{1}{1 + e^{-net}} = \frac{1}{1 + e^{-w^T x}}$$
 (notes: net = $w^T x + b = w^T x$)

Cost function:

$$J(w) = \frac{1}{m} \sum_{i=1}^{m} \frac{1}{2} \left(\phi_{w}(x^{(i)}) - y^{(i)} \right)^{2} = \frac{1}{m} \sum_{i=1}^{m} \frac{1}{2} \left(\phi(net) - y^{(i)} \right)^{2}$$

$$\hat{y} = \begin{cases} 1 & \text{if } \phi(net) \ge 0.5 \\ 0 & \text{otherwise} \end{cases} = \begin{cases} 1 & \text{if } net = w^{T} x \ge 0 \\ 0 & \text{otherwise} \end{cases}$$

$$J(\phi(net), y) = \begin{cases} -\log(\phi(net)) & \text{if } y = 1 \\ -\log(1 - \phi(net)) & \text{if } y = 0 \end{cases}$$

$$J(\phi(net), y) = -y \log(\phi(net)) - (1 - y) \log(1 - \phi(net))$$

Update the weights:

$$w_j = w_j + \Delta w_j$$

$$\Delta w_j = -\eta \frac{\partial J}{\partial w_i} = \frac{1}{m} \sum_{i=1}^m (y(i) - \phi(w^T x_j^{(i)})) x_j^{(i)}$$



For Iris classification

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import load_iris
iris = load iris()
x = iris.data[1:100,[1,3]]
y = iris.target[1:100]
list(iris.target names)
print(x.shape)
plt.plot(x[y==0,0],x[y==0,1],'g+',label='1')
plt.plot(x[y==1,0],x[y==1,1],'bo',label='0')
                                                 1.75
plt.legend(loc='upper right')
                                                 1.50
plt.xlabel('Feature1')
plt.ylabel('Feature2')
                                                 1.25
plt.grid()
                                              Feature 7.75
plt.show()
                                                 0.50
                                                 0.25
                                                      2.0
                                                                2.5
                                                                         3.0
                                                                                   3.5
                                                                                            4.0
                                                                                                     4.5
                                                                           Feature1
```



A perceptron (/Adaline) with Gaussian activation function

```
# implement logistic regression classified
class logistic_regression(object):
 def init (self, epochs = 100, eta = 0.1, random state=1):
   self.epochs = epochs
   self.eta = eta
    self.random_state = random_state
 def fit(self, x,y):
   np.random.RandomState(self.random_state)
    self.w = np.random.normal(loc = 0.0, scale = 1.0, size = x.shape[1]+1)
    print(self.w_)
    self.cost = []
   for i in range(self.epochs):
     net = np.dot(x,self.w_[1:])+self.w_[0]
     #print(net)
     #print(net.shape)
     output = 1./(1.+np.exp(-net))
     #print(output)
     error = (y-output)
     #print(np.dot(x.T,error))
     #print(x.T.dot(error))
     self.w [1:]+=self.eta*np.dot(x.T,error)
     self.w_[0]+=self.eta*error.sum()
     #print(y.shape)
     #print(output.shape)
     #print(np.log(output))
     cost = error.mean()#
     cost=-np.dot(y, np.log(output))-np.dot((1-y), np.log(1-output))
      self.cost_.append(cost)
    return(self)
  def predict(self, x):
   net = np.dot(x,self.w_[1:])+self.w_[0]
   output = 1./(1.+np.exp(-net))
   return(np.where(output>=0.5,1,0))
```



```
model = logistic_regression(epochs=20, eta = 0.05)
model.fit(x,y)
y_pred = model.predict(x)
error=np.sum(np.abs(y-y_pred))
print(error)
# decision line
xx1 = np.arange(x[:,0].min()-1,x[:,0].max()+1,0.1)
xx2 = -(model.w_[1]*xx1+model.w_[0])/model.w_[2]
plt.plot(x[y_pred==0,0],x[y_pred==0,1],'g+',label='1')
plt.plot(x[y_pred==1,0],x[y_pred==1,1],'bo',label='0')
plt.legend(loc='upper left')
plt.plot(xx1, xx2, 'm-')
                                                      1.75
plt.xlabel('Feature 1')
plt.ylabel('Feature 2')
                                                      1.50
plt.grid()
plt.show()
                                                      1.25
                                                   Feature 2
plt.plot(range(len(model.cost_)), model.cost_)
                                                      1.00
plt.xlabel('Epochs')
                                                      0.75
plt.ylabel('Error')
plt.grid()
                                                      0.50
plt.show()
                                                                                            +++
                                                                                                    +
                                                      0.25
    1000
     800
                                                                                  Feature 1
     600
```

200

7.5

5.0

2.5

0.0

10.0

Epochs

12.5

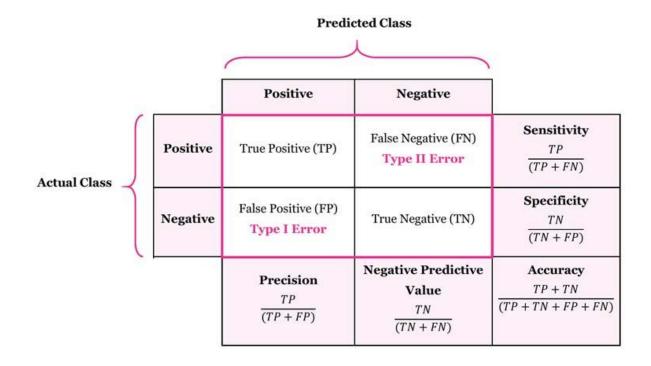
15.0

17.5



Confusion matrix

 A confusion matrix is a table that is often used to describe the performance of a classification model (or "classifier") on a set of test data for which the true values are known.





Basic terms of a confusion matrix (CM)

- True positives (TP): These are cases in which we predicted yes (they have the disease), and they do have the disease.
- True negatives (TN): We predicted no, and they don't have the disease.
- False positives (FP): We predicted yes, but they don't have the disease. (Also known as a "Type I error")
- False negatives (FN): We predicted no, but they do have the disease.
 (Also known as a "Type II error")
- Prediction error (ERR): the sum of all false predictions divided by the total number of predictions.
- Accuracy (ACC): the sum of correct (true) predictions divided by the total number of predictions.
- True positive rate (TPR)
- False positive rate (FPR)
- Precision (PRE):
- Recall (REC):

$$ERR = \frac{FP + FN}{FP + FN + TP + TN}$$

$$ACC = \frac{TP + TN}{FP + FN + TP + TN} = 1 - ERR$$

$$FPR = \frac{FP}{N} = \frac{FP}{FP + TN}, \quad TPR = \frac{TP}{P} = \frac{FP}{FN + TP}$$

$$PRE = \frac{TP}{TP + FP}, \quad REC = \frac{TP}{FN + TP}$$



Confusion matrix example

 A binary classifier for medical testing to determine if a patient has certain disease or not (e.g. tumor diagnosis).

	Predicted:	Predicted:	
n=165	NO	YES	
Actual:			60
NO	50	10	
Actual:			105
YES	5	100	103
	55	110	•

- 2) There are two possible predicted classes: "yes" and
- 3) If We were predicting the presence of a disease, "yes" would mean a patient has the disease, and "no" would mean he/she doesn't have the disease.
- 4) The classifier made a total of **165** predictions (e.g., 165 patients were being tested for the presence of that disease).
- 5) Out of those 165 cases, the classifier predicted "yes" 110 times, and "no" 55 times.
- 6) In reality,105 patients in the sample have the disease, and 60 patients do not.



n=165	Predicted: NO	Predicted: YES	
Actual: NO	50	10	60
Actual: YES	5	100	10

110

- Accuracy: Overall, how often is the classifier correct?
 - (TP+TN)/total = (100+50)/165 = 0.91
- Misclassification Rate: Overall, how often is it wrong?
 - (FP+FN)/total = (10+5)/165 = 0.09
 - equivalent to 1 minus Accuracy
 - also known as "Error Rate"
- True Positive Rate: When it's actually yes, how often does it predict yes?
 - TP/actual yes = 100/105 = 0.95
 - also known as "Sensitivity" or "Recall"
- False Positive Rate: When it's actually no, how often does it predict yes?
 - FP/actual no = 10/60 = 0.17
- True Negative Rate: When it's actually no, how often does it predict no?
 - TN/actual no = 50/60 = 0.83
 - equivalent to 1 minus False Positive Rate
 - also known as "Specificity"
- Precision: When it predicts yes, how often is it correct?
 - TP/predicted yes = 100/110 = 0.91
- Prevalence: How often does the yes condition actually occur in our sample?
 - actual yes/total = 105/165 = 0.64



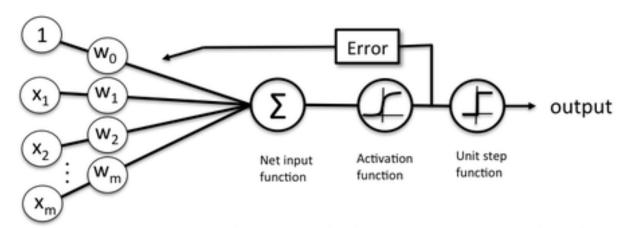
CM for iris

```
from matplotlib.colors import ListedColormap
cmap = ListedColormap(['W', 'Y', 'M'])
# Compute confusion matrix
y true = y
class_names = iris.target_names
print(class names[0:-1])
#cnf matrix = confusion matrix(y true, y pred, labels=[0, 1]) # 0 = 'setosa', 1 = 'versicolor'
cm = confusion matrix(y, y pred) # y true and y pred - 0 = 'setosa', 1 = 'versicolor'
print(cm)
plt.figure()
                                                                             0
                                                                                                1
plt.matshow(cm, cmap=cmap)
for i in range(cm.shape[0]):
  for j in range(cm.shape[1]):
    plt.text(x=j, y=i, s=cm[i,j], va='center', ha='center')
                                                                             49
                                                                                                0
plt.xlabel('Predicted label')
                                                                  0 -
plt.ylabel('True label')
plt.show()
                                                               Frue label
                                                                                               50
                                                                                 Predicted label
```

34



Recap: single-layer neural network



Schematic of a logistic regression classifier.

$$J(w) = \frac{1}{2} \sum_{i=1}^{m} (y(i) - \phi(w^{T} x_{j}^{(i)}))^{2}$$

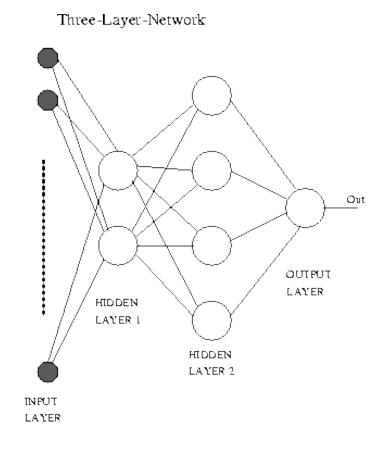
$$w_{j} = w_{j} + \Delta w_{j}$$

$$\Delta w_{j} = -\eta \frac{\partial J(w)}{\partial w_{j}} = \frac{1}{m} \sum_{i=1}^{m} (y(i) - \phi(w^{T} x_{j}^{(i)})) x_{j}^{(i)}$$



Multilayer neural network

Two-Layer-Network Out L Outo Out 3 Ουτρυτ LAYER HIDDEN LAYER

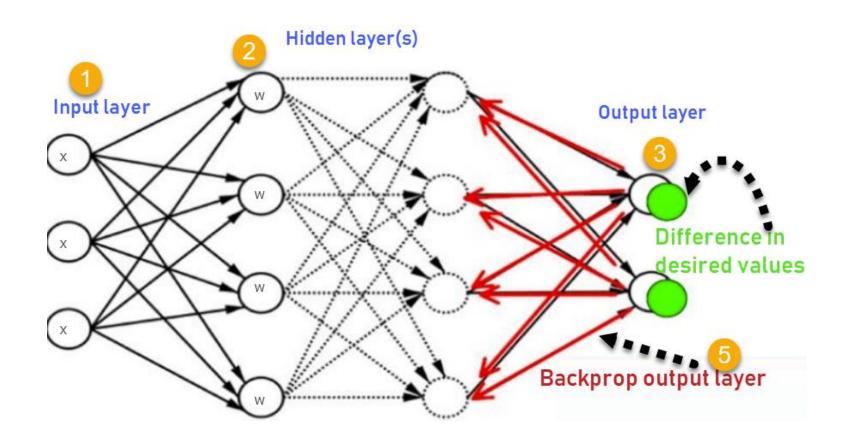


INPUT

LAYER



How backpropagation works?





MLP in sklearn

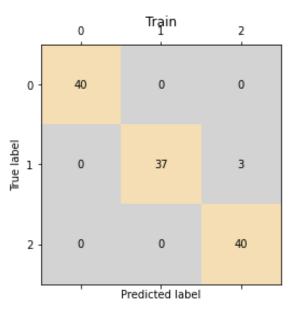
```
# MLP
import numpy as np
import matplotlib.pyplot as plt
from sklearn.datasets import load_iris
from sklearn.neural_network import MLPClassifier
from sklearn.model_selection import train_test_split
from sklearn.metrics import confusion_matrix
from matplotlib.colors import ListedColormap

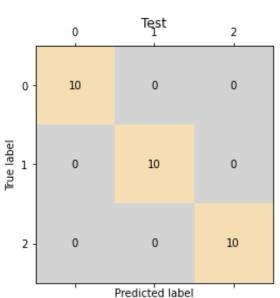
iris = load_iris()
x = iris.data
y = iris.target
print(iris.target_names)

x_train, x_test, y_train, y_test = train_test_split(x, y, stratify=y,random_state=1,test_size=0.2)
np.bincount(y), np.bincount(y_train), np.bincount(y_test)
```



```
mlp =MLPClassifier(solver='sgd', random state=0, hidden layer sizes=[5], alpha = 0.5,max iter=1000)
# optimizer = 'lbfgs': quasi-Newton methods, 'sgd': stochastic gradient descent, and 'adam': stochastic gradient-based optimizer
mlp.fit(x_train, y_train)
y pred = mlp.predict(x test)
print(np.sum(y_test!=y_pred)/len(y_test)*100)
#confusion matrix
def cm(y_test, y_pred):
  cm = confusion matrix(y test, y pred)
 cmap = ListedColormap(['lightgrey', 'silver', 'ghostwhite', 'lavender', 'wheat'])
 cm = confusion matrix(y test, y pred)
 plt.figure()
 plt.matshow(cm, cmap=cmap)
 for i in range(cm.shape[0]):
   for j in range(cm.shape[1]):
      plt.text(x=j, y=i, s=cm[i,j], va='center', ha='center')
  plt.xlabel('Predicted label')
 plt.ylabel('True label')
 plt.show()
cm(y_test, mlp.predict(x_test))
cm(y_train, mlp.predict(x_train))
```







MLP for MNIST dataset

- The MNIST database of handwritten digits has a training set of 60,000 examples, and a test set of 10,000 examples.
- The digits have been size-normalized and centered in a fixed-size image.
- It is a good database for people who want to try learning techniques and pattern recognition methods on real-world data while spending minimal efforts on preprocessing and formatting.
- Each digit is 28x28 pixels gray image.





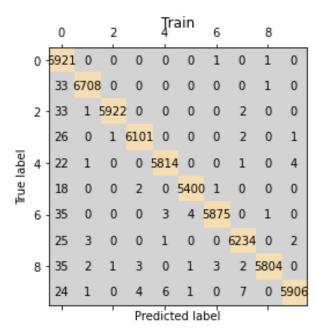
MLP Classifier

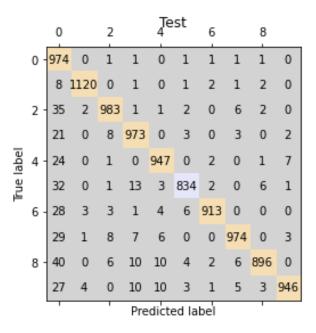
```
# MLP for MNIST
import numpy as np
import matplotlib.pyplot as plt
from keras.datasets import mnist
from keras.utils import to categorical
from sklearn.neural network import MLPClassifier
from sklearn.metrics import confusion matrix
from sklearn.metrics import plot confusion matrix
from matplotlib.colors import ListedColormap
(x_train, y_train), (x_test, y_test)=mnist.load_data() #60000 samples for training, 10000 samples for testing
print('train:', x train.shape, y train.shape)
print('Test set:', x_test.shape, y_test.shape)
# visualize some
fig, ax = plt.subplots(2,5,sharex=True,sharey=True)
                                                             50419
ax = ax.flatten()
for i in range(10):
  ax[i].imshow(x_train[i,:])
plt.show()
# prepare the data for training
x train = x train.reshape(60000,28*28)
x train = x_train.astype('float32')/255
x_{\text{test}} = x_{\text{test.reshape}}(10000,28*28)
x test = x test.astype('float32')/255
y_train = to_categorical(y_train)
y test = to_categorical(y_test)
```



```
mlp = MLPClassifier(solver='sgd', activation='logistic',alpha=1e-4,hidden layer sizes=(100,30),
                    random state=1, max iter=100, verbose=True, learning rate init=.1, tol=1e-4)
mlp.fit(x train, y train)
print("Training set score: %f" % mlp.score(x train, y train))
print("Test set score: %f" % mlp.score(x test, y test))
y pred = mlp.predict(x test)
cmap = ListedColormap(['lightgrey', 'silver', 'ghostwhite', 'lavender', 'wheat'])
#confusion matrix
def cm(y test, y pred, title):
 cm = confusion matrix(y test, y pred)
 plt.figure()
  plt.matshow(cm, cmap=cmap)
 for i in range(cm.shape[0]):
   for j in range(cm.shape[1]):
      plt.text(x=j, y=i, s=cm[i,j], va='center', ha='center')
  plt.title(title)
 plt.xlabel('Predicted label')
 plt.ylabel('True label')
  plt.show()
cm(y_train.argmax(1), mlp.predict(x_train).argmax(1), title='Train')
cm(y test.argmax(1), mlp.predict(x test).argmax(1), title='Test')
```







	Training	Testing	
Accuracy	99.31	94.84	



MNIST using Keras

import matplotlib.pyplot as plt

from keras.datasets import mnist

from keras.layers import Dense

from keras.models import Sequential

from keras.utils import to categorical

MLP for MNTST

import keras

import numpy as np

```
Model: "sequential"
```

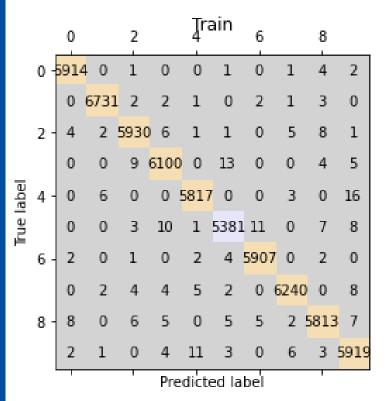
```
Layer (type)
                              Output Shape
                                                         Param #
dense (Dense)
                              (None, 350)
                                                         274750
dense 1 (Dense)
                              (None, 50)
                                                          17550
dense 2 (Dense)
                              (None, 10)
                                                          510
```

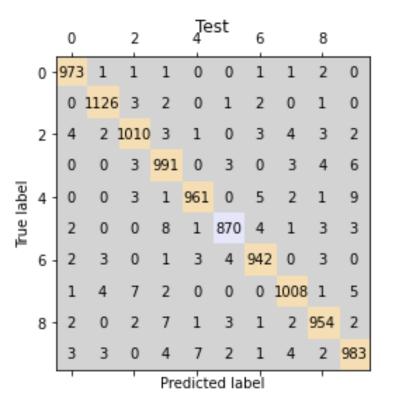
Total params: 292,810 Trainable params: 292,810 Non-trainable params: 0

from sklearn.metrics import confusion_matrix

```
# classification model
input shape=x train.shape[1]
num classes = y train.shape[1]
model = Sequential()
model.add(Dense(350, activation='relu', input shape=(28*28,)))
model.add(Dense(50, activation='relu'))
model.add(Dense(num classes, activation='softmax'))
model.compile(loss='categorical crossentropy', optimizer='adam', metrics=['accuracy'])
model.summary()
model.fit(x train, y train, epochs=100, batch size=512, verbose=1, validation split=0.2)
# Test the model after training
test_results = model.evaluate(x_test, y_test, verbose=1)
print(f'Test results \n - Loss : {test_results[0]: 0.3f} \n - Accuracy: {test_results[1]: 0.3f}')
```







	Loss	Accura cy
Categorical cross entropy	0.122	0.982



Saving and loading models

- Models take a long time to train.
- It is required to save trained models, so we do not have to train them every time.