

Introduction to TensorFlow

Fully Connected Deep Networks with TensorFlow

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大綱

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關於 Fully Connected Deep Networks

Fully Connected Deep Networks 多層感知器

單一個感知器（Perceptron）的公式：

$$y = \sigma(WX + b)$$

σ 即所謂的激活函數（activation function）

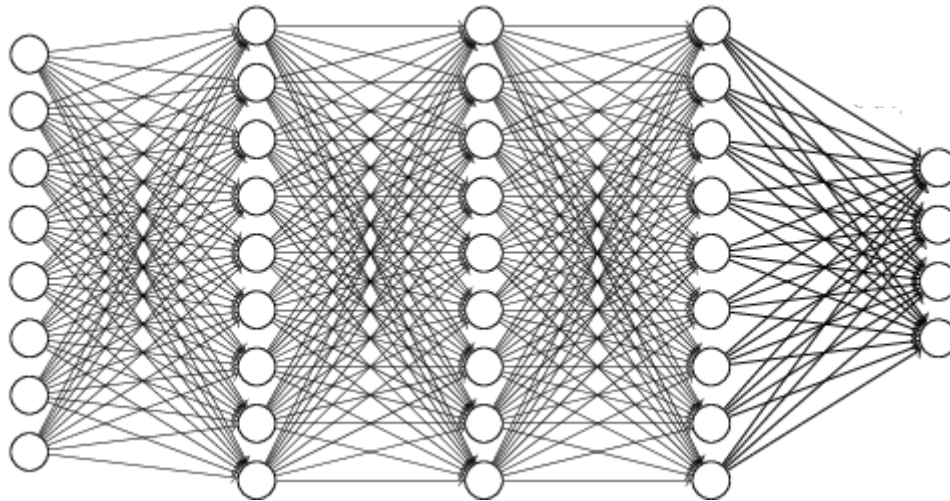
激活函數用來將線性關係映射至非線性關係

常用的激活函數：

- Sigmoid
- TanH
- ReLU

連結多個感知器建構出多層感知器，透過 backpropagation 求出係數

- 輸入層
- 隱藏層
- 輸出層



Source: <https://math.stackexchange.com/questions/2048722/a-name-for-layered-directed-graph-as-in-a-fully-connected-neural-network>
(<https://math.stackexchange.com/questions/2048722/a-name-for-layered-directed-graph-as-in-a-fully-connected-neural-network>).

取得資料

簡單、作為測試目的即可

Scikit-Learn Breast Cancer 資料集

```
In [1]: from sklearn.datasets import load_breast_cancer
```

```
breast_cancer = load_breast_cancer()
print(breast_cancer.feature_names)
print(breast_cancer.DESCR)
```

```
['mean radius' 'mean texture' 'mean perimeter' 'mean area'
 'mean smoothness' 'mean compactness' 'mean concavity'
 'mean concave points' 'mean symmetry' 'mean fractal dimension'
 'radius error' 'texture error' 'perimeter error' 'area error'
 'smoothness error' 'compactness error' 'concavity error'
 'concave points error' 'symmetry error' 'fractal dimension error'
 'worst radius' 'worst texture' 'worst perimeter' 'worst area'
 'worst smoothness' 'worst compactness' 'worst concavity'
 'worst concave points' 'worst symmetry' 'worst fractal dimension']
.. _breast_cancer_dataset:
```

Breast cancer wisconsin (diagnostic) dataset

****Data Set Characteristics:****

:Number of Instances: 569

:Number of Attributes: 30 numeric, predictive attributes and the class

:Attribute Information:

- radius (mean of distances from center to points on the perimeter)
- texture (standard deviation of gray-scale values)
- perimeter
- area
- smoothness (local variation in radius lengths)
- compactness (perimeter² / area - 1.0)
- concavity (severity of concave portions of the contour)
- concave points (number of concave portions of the contour)
- symmetry

- fractal dimension ("coastline approximation" - 1)

The mean, standard error, and "worst" or largest (mean of the three largest values) of these features were computed for each image, resulting in 30 features. For instance, field 3 is Mean Radius, field 13 is Radius SE, field 23 is Worst Radius.

- class:
 - WDBC-Malignant
 - WDBC-Benign

:Summary Statistics:

=====	=====	=====
	Min	Max
=====	=====	=====
radius (mean):	6.981	28.11
texture (mean):	9.71	39.28
perimeter (mean):	43.79	188.5
area (mean):	143.5	2501.0
smoothness (mean):	0.053	0.163
compactness (mean):	0.019	0.345
concavity (mean):	0.0	0.427
concave points (mean):	0.0	0.201
symmetry (mean):	0.106	0.304
fractal dimension (mean):	0.05	0.097
radius (standard error):	0.112	2.873
texture (standard error):	0.36	4.885
perimeter (standard error):	0.757	21.98
area (standard error):	6.802	542.2
smoothness (standard error):	0.002	0.031
compactness (standard error):	0.002	0.135
concavity (standard error):	0.0	0.396
concave points (standard error):	0.0	0.053
symmetry (standard error):	0.008	0.079
fractal dimension (standard error):	0.001	0.03
radius (worst):	7.93	36.04
texture (worst):	12.02	49.54

perimeter (worst):	50.41	251.2
area (worst):	185.2	4254.0
smoothness (worst):	0.071	0.223
compactness (worst):	0.027	1.058
concavity (worst):	0.0	1.252
concave points (worst):	0.0	0.291
symmetry (worst):	0.156	0.664
fractal dimension (worst):	0.055	0.208
=====	=====	=====

:Missing Attribute Values: None

:Class Distribution: 212 - Malignant, 357 - Benign

:Creator: Dr. William H. Wolberg, W. Nick Street, Olvi L. Mangasarian

:Donor: Nick Street

:Date: November, 1995

This is a copy of UCI ML Breast Cancer Wisconsin (Diagnostic) datasets.
<https://goo.gl/U2Uwz2>

Features are computed from a digitized image of a fine needle aspirate (FNA) of a breast mass. They describe characteristics of the cell nuclei present in the image.

Separating plane described above was obtained using Multisurface Method-Tree (MSM-T) [K. P. Bennett, "Decision Tree Construction Via Linear Programming." Proceedings of the 4th Midwest Artificial Intelligence and Cognitive Science Society, pp. 97-101, 1992], a classification method which uses linear programming to construct a decision tree. Relevant features were selected using an exhaustive search in the space of 1-4 features and 1-3 separating planes.

The actual linear program used to obtain the separating plane in the 3-dimensional space is that described in:

[K. P. Bennett and O. L. Mangasarian: "Robust Linear Programming Discrimination of Two Linearly Inseparable Sets", Optimization Methods and Software 1, 1992, 23-34].

This database is also available through the UW CS ftp server:

```
ftp ftp.cs.wisc.edu
cd math-prog/cpo-dataset/machine-learn/WDBC/
```

.. topic:: References

- W.N. Street, W.H. Wolberg and O.L. Mangasarian. Nuclear feature extraction
for breast tumor diagnosis. IS&T/SPIE 1993 International Symposium on
Electronic Imaging: Science and Technology, volume 1905, pages 861-870,
San Jose, CA, 1993.
- O.L. Mangasarian, W.N. Street and W.H. Wolberg. Breast cancer diagnosis and
prognosis via linear programming. Operations Research, 43(4), pages 570-5
77,
July-August 1995.
- W.H. Wolberg, W.N. Street, and O.L. Mangasarian. Machine learning techniques
to diagnose breast cancer from fine-needle aspirates. Cancer Letters 77
(1994)
163-171.

```
In [2]: X_arr = breast_cancer.data  
y_arr = breast_cancer.target  
print(X_arr.shape)  
print(y_arr.shape)
```

```
(569, 30)
```

```
(569,)
```

```
In [3]: from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test = train_test_split(X_arr, y_arr, test_size=0.3, r
andom_state=123)
print(X_train.shape)
print(X_test.shape)
print(y_train.shape)
print(y_test.shape)

(398, 30)
(171, 30)
(398,)
(171,)
```

Benchmark

```
In [4]: from sklearn.linear_model import LogisticRegression
        from sklearn.metrics import accuracy_score

        clf = LogisticRegression(solver="liblinear")
        clf.fit(X_train, y_train)
        y_pred = clf.predict(X_test)
        acc = accuracy_score(y_test, y_pred)
```



```
In [5]: print(clf.coef_)  
print(clf.intercept_)  
print(acc)
```

```
[[ 1.93815613e+00  1.95132301e-01 -2.79314889e-03 -9.98660343e-03  
 -1.31191303e-01 -3.94395839e-01 -5.87981725e-01 -3.08507409e-01  
 -2.34633450e-01 -2.07557569e-02 -1.29122502e-03  1.25273443e+00  
  8.82364969e-03 -8.58922482e-02 -1.44638024e-02 -2.16113982e-04  
 -4.51757012e-02 -3.74554453e-02 -3.87686384e-02  6.76008093e-03  
  1.00073745e+00 -3.88446001e-01 -1.28867607e-01 -1.78769588e-02  
 -2.50482381e-01 -1.09777946e+00 -1.45824047e+00 -6.35683325e-01  
 -7.02842419e-01 -9.58336644e-02]]  
[0.41220102]  
0.9824561403508771
```

Benchmark 完整程式碼

```
In [6]: from sklearn.datasets import load_breast_cancer
        from sklearn.model_selection import train_test_split
        from sklearn.linear_model import LogisticRegression
        from sklearn.metrics import accuracy_score

        breast_cancer = load_breast_cancer()
        X_arr = breast_cancer.data
        y_arr = breast_cancer.target
        X_train, X_test, y_train, y_test = train_test_split(X_arr, y_arr, test_size=0.3, r
        andom_state=123)
        clf = LogisticRegression(solver="liblinear")
        clf.fit(X_train, y_train)
        y_pred = clf.predict(X_test)
        acc = accuracy_score(y_test, y_pred)
```

建構 TensorFlow 計算圖形

準備 Placeholders 作為 Input layers

```
In [7]: import tensorflow as tf

n_features = X_train.shape[1]
with tf.name_scope("input-layer"):
    X = tf.placeholder(tf.float32, (None, n_features)) # ?
    y = tf.placeholder(tf.float32, (None,))           # ?
```

Placeholders 的觀測值個數設定為 None 有什麼好處？

- 多層感知器的應用常會使用 Mini-batching 的技巧
- 訓練結束之後可以餵入 `X_test` 獲得 `y_pred_arr`

準備 Variables 作為 Hidden layers

```
In [8]: import numpy as np

n_classes = np.unique(y_train).size
n_neurons = int((n_features + n_classes) / 2)
with tf.name_scope("hidden-layer"):
    W = tf.Variable(tf.random_normal((n_features, n_neurons))) # (30, 16)
    b = tf.Variable(tf.random_normal((n_neurons,))) # (16,)
    X_hidden = tf.nn.relu(tf.matmul(X, W) + b) # (398, 30) x (30, 16) + (16,)
```

如何決定 hidden-layer 的層數與神經元數量？

- 絕大多數情境：使用一層 hidden-layer 就足夠
- 神經元數量：(輸入層神經元個數 + 輸出層神經元個數) / 2

```
n_neurons = int((n_features + n_classes) / 2)
```


準備 Variables 作為 Output layers

```
In [9]: with tf.name_scope("output-layer"):
        W = tf.Variable(tf.random_normal((n_neurons, 1))) # (16, 1)
        b = tf.Variable(tf.random_normal((1,)))           # (1,)
        y_logit = tf.squeeze(tf.matmul(X_hidden, W) + b) # (398, 16) x (16, 1) + (1,)
        y_one_prob = tf.sigmoid(y_logit)
        y_pred = tf.round(y_one_prob)
```

寫下成本函數的公式

```
In [10]: with tf.name_scope("loss"):
          entropy = tf.nn.sigmoid_cross_entropy_with_logits(logits=y_logit, labels=y)
          loss = tf.reduce_sum(entropy)
```

宣告 Optimizer 與學習速率

```
In [11]: learning_rate = 0.01
with tf.name_scope("optimizer"):
    opt = tf.train.AdamOptimizer(learning_rate)
    optimizer = opt.minimize(loss)
```

建構 TensorFlow 計算圖形完整程式碼

```
In [12]: import tensorflow as tf
import numpy as np

tf.reset_default_graph()
n_features = X_train.shape[1]
n_classes = np.unique(y_train).size
n_neurons = int((n_features + n_classes) / 2)
learning_rate = 0.0001

with tf.name_scope("input-layer"):
    X = tf.placeholder(tf.float32, (None, n_features))
    y = tf.placeholder(tf.float32, (None,))
with tf.name_scope("hidden-layer"):
    W = tf.Variable(tf.random_normal((n_features, n_neurons))) # (30, 16)
    b = tf.Variable(tf.random_normal((n_neurons,))) # (16,)
    X_hidden = tf.nn.relu(tf.matmul(X, W) + b) # (398, 30) x (30, 16) + (16,)
with tf.name_scope("output-layer"):
    W = tf.Variable(tf.random_normal((n_neurons, 1))) # (16, 1)
    b = tf.Variable(tf.random_normal((1,))) # (1,)
    y_logit = tf.squeeze(tf.matmul(X_hidden, W) + b) # (398, 16) x (16, 1) + (1,)
    y_one_prob = tf.sigmoid(y_logit)
    y_pred = tf.round(y_one_prob)
with tf.name_scope("loss"):
    entropy = tf.nn.sigmoid_cross_entropy_with_logits(logits=y_logit, labels=y)
    loss = tf.reduce_sum(entropy)
with tf.name_scope("optimizer"):
    opt = tf.train.AdamOptimizer(learning_rate)
    optimizer = opt.minimize(loss)
```

訓練

```

In [13]: n_steps = 50000
         file_writer_path = "./graphs/fully-connected-deep-networks"
         loss_history = []

         with tf.Session() as sess:
             sess.run(tf.global_variables_initializer()) # 初始化所有的變數張量!
             train_writer = tf.summary.FileWriter(file_writer_path, tf.get_default_graph())
             for i in range(n_steps):
                 feed_dict = {
                     X: X_train,
                     y: y_train
                 }
                 _, loss_ = sess.run([optimizer, loss], feed_dict=feed_dict)
                 loss_history.append(loss_)
                 if i % 1000 == 0:
                     print("step {}, loss: {}".format(i, loss_))
             w_final, b_final = sess.run([W, b])
             y_pred_arr = sess.run(y_pred, feed_dict={X: X_test})

```

```

step 0, loss: 414025.875
step 1000, loss: 158248.84375
step 2000, loss: 2681.51953125
step 3000, loss: 2467.9853515625
step 4000, loss: 2411.2861328125

step 5000, loss: 2320.51806640625
step 6000, loss: 2178.854736328125
step 7000, loss: 1970.0159912109375
step 8000, loss: 1686.59521484375
step 9000, loss: 1358.921875
step 10000, loss: 987.6146850585938
step 11000, loss: 609.17138671875
step 12000, loss: 372.76959228515625
step 13000, loss: 255.74600219726562
step 14000, loss: 155.87393188476562
step 15000, loss: 118.18053436279297
step 16000, loss: 94.63573455810547
step 17000, loss: 73.91920471191406

```

```
In [14]: import matplotlib.pyplot as plt

plt.plot(range(n_steps), loss_history)
plt.title("Loss Summary")
plt.xlabel("Epochs")
plt.ylabel("Loss")
plt.show()
```

<Figure size 640x480 with 1 Axes>

```
In [15]: acc = accuracy_score(y_test, y_pred_arr)
```

```
print(w_final)
```

```
print(b_final)
```

```
print(acc)
```

```
[[-0.7065138 ]
```

```
 [-0.66401774]
```

```
 [-2.1829813 ]
```

```
 [ 1.1565275 ]
```

```
 [-0.40028438]
```

```
 [ 0.9624974 ]
```

```
 [-0.2515118 ]
```

```
 [-0.6297551 ]
```

```
 [-1.0843177 ]
```

```
 [ 0.5629655 ]
```

```
 [ 1.0997338 ]
```

```
 [-1.6978433 ]
```

```
 [ 1.5582514 ]
```

```
 [-0.769502  ]
```

```
 [ 1.513353  ]
```

```
 [-2.228517  ]]
```

```
[0.8738572]
```

```
0.9824561403508771
```


隨堂練習

使用 titanic 資料集並利用 TensorFlow 建立一個 Fully Connected Deep Networks Classifier

```
In [16]: import pandas as pd

train_url = "https://s3-ap-northeast-1.amazonaws.com/kaggle-getting-started/titanic/train.csv"
test_url = "https://s3-ap-northeast-1.amazonaws.com/kaggle-getting-started/titanic/test.csv"
train_df = pd.read_csv(train_url)
test_df = pd.read_csv(test_url)
```