### 3장 Getting started with neural networks

"기회와 준비가 만났을 때 ... "

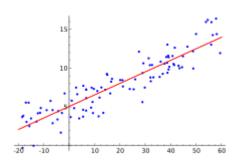


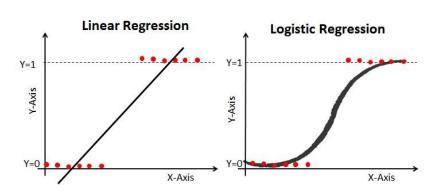


- The two previous examples predict a single discrete label of an input data point.
- Another common type of machine-learning problem is *regression* predicting a continuous value: temperature tomorrow, time, or price

NOTE Don't confuse *regression* and the algorithm *logistic regression*. Confusingly, logistic regression isn't a regression algorithm—it's a classification algorithm.

regression 이라고해서 다 regression은 아니고 Linear나 Logistic이 불으면 Classification이다!







### 3.6.1 The Boston Housing Price dataset

- predict the median price of homes in a given Boston suburb with the crime rate in the mid-1970s
- only 506 samples 404 training samples and 102 test samples.
- And each *feature* in the input data (for example, the crime rate, the local property tax rate) has a different scale. For instance, some values are proportions, which take values between 0 and 1; others take values between 1 and 12, others between 0 and 100, and so on.

### Listing 3.24 Loading the Boston housing

Let's look at the data:

```
>>> train_data.shape
(404, 13) # 13 features
>>> test_data.shape
(102, 13)
```



### 3.6.1 The Boston Housing Price dataset

- As you can see, you have 404 training samples and 102 test samples, each with 13 numerical features capita crime rate, average number of rooms per dwelling, accessibility to highways, and so on.
- ▶ The targets are the median values of owner-occupied homes, in thousands of dollars:

```
>>> train_targets
[15.2, 42.3, 50. ... 19.4, 19.4, 29.1]# 404 in $(*1000)
```

The prices are typically between \$10,000 and \$50,000

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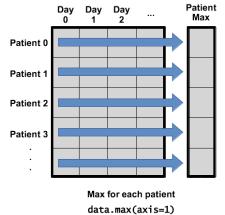
### 6. Predicting house prices: a regression example

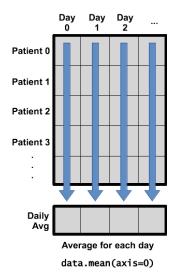


#### 3.6.2 Preparing the data

• feature-wise normalization: feature is centered around 0 and has a unit standard deviation.

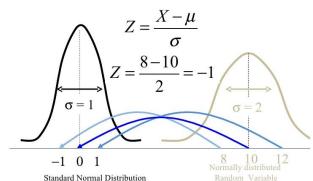
#### Listing 3.25 Normalizing the data





Note that the quantities used for normalizing the test data are computed using the training data.







#### 3.6.3 Building your network

- > small samples small network with two hidden layers, each with 64 units.
- less training data worse overfitting, a small network is one way to mitigate overfitting.

#### **Listing 3.26 Model definition**

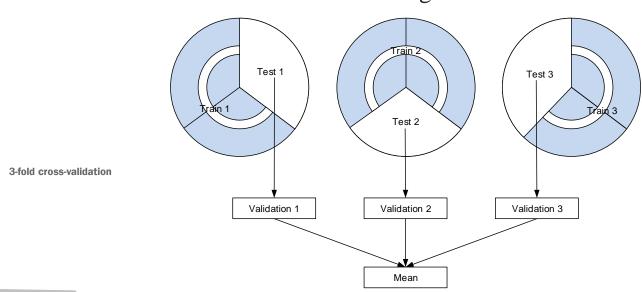
- output a single unit and no activation (it will be a linear layer) for scalar regression to predict a single continuous value, free to learn to predict values in any range
- ▶ sigmoid activation function predict values between 0 and 1
- ▶ mean absolute error (MAE) loss function for regression problems  $0.5 \rightarrow$  off by \$500 on average.



#### 3.6.4 Validating your approach using K-fold validation

매우 중요(Validation하는 방법)

- very small validation set (for instance, about 100 examples) high variance with regard to the validation split
- $\blacktriangleright$  K-fold cross-validation (see figure) splitting the available data into K partitions (typically K = 4 or 5), instantiating K identical models, and training each one on K - 1partitions while evaluating on the remaining partition.
- ▶ The validation score for the model average of the *K* validation scores





#### 3.6.4 Validating your approach using K-fold validation

#### **Listing 3.27 K-fold validation**

```
import numpy as np
k = 4
num val samples = len(train data) // k #나눗셈의몫
num epochs = 100
all scores = []
for i in range(k): \# i = 0,1,2,3
   print('processing fold #', i)
  val data = train data[i * num val samples: (i + 1) * num val samples]
  val targets = train targets[i * num val samples: (i + 1) * num val samples]
  partial train data = np.concatenate ([train data[:i* num val samples],
      train data[(i + 1) * num val samples:]],
      axis=\overline{0})
   partial train targets = np.concatenate( [train targets[:i * num val samples],
      train targets[(i + 1) * num val samples:]],
      axis=0)
   model = build model()
   history = model.fit(partial train data, partial train targets,
        validation data=(val data, val targets), epochs=num epochs,
        batch size=1, verbose=0)
   mae history = history.history['val mean absolute error']
   all mae histories.append (mae history)
```



### 3.6.4 Validating your approach using K-fold validation

Running this with num\_epochs = 100 yields the following results:

```
>>> all_scores
[2.588258957792037, 3.1289568449719116, 3.1856116051248984, 3.0763342615401386]
>>> np.mean(all_scores)
2.9947904173572462
```

- different validation scores, from 2.6 to 3.2.
- ▶ The average (3.0) is a much more reliable metric K-fold cross-validation
- ▶ \$3,000 on average significant considering with \$10,000 to \$50,000.



#### 3.6.4 Validating your approach using K-fold validation

Let's try training the network a bit longer: 500 epochs. To keep a record of how well the model does at each epoch, you'll modify the training loop to save the per-epoch validation score log.

#### Listing 3.28 Saving the validation logs

```
num epochs = 500
all mae histories = []
for i in range(k):
   print('처리중인 폴드 #', i)
   val data = train data[i * num val samples: (i + 1) * num val samples]
   val targets = train targets[i * num val samples: (i + 1) * num val samples]
   partial train data = np.concatenate(
        [train data[:i * num val samples],
        train data[(i + 1) * num val samples:]],
        axis=0)
   partial train targets = np.concatenate(
        [train targets[:i * num val samples],
        train targets[(i + 1) * num val samples:]],
        axis=0)
   model = build model()
   history = model.fit(partial train data, partial train targets,
                        validation data=(val data, val targets),
                        epochs=num epochs, batch size=1, verbose=0)
   mae history = history.history['val mean absolute error']
    all mae histories.append(mae history)
```



### 3.6.4 Validating your approach using K-fold validation

• the average of the per-epoch MAE scores for all folds.

#### **Listing 3.29 Building the history of successive**

```
average_mae_history = [np.mean([x[i] for x in all_mae_histories])
  for i in range(num_epochs)]
```

Let's plot this; see figure 3.12.

### **Listing 3.30 Plotting validation scores**

```
import matplotlib.pyplot as plt
plt.plot(range(1,len(average_mae_history)+1),
    average_mae_history)
plt.xlabel('Epochs')
plt.ylabel('Validation MAE')
plt.show()
```

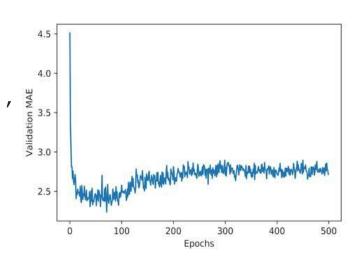


Figure 3.12 Validation MAE by epoch

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# 6. Predicting house prices: a regression example



- ▶ validation MAE stops improving significantly after 80 epochs.
- ▶ adjust the size of the hidden layers, and then look at its performance on the test data.

#### **Listing 3.32 Training the final model**

```
model = build_model()
model.fit(train_data, train_targets,epochs=80, batch_size=16, verbose=0)
test_mse_score, test_mae_score = model.evaluate(test_data, test_targets)
```

Here's the final result:

```
>>> test_mae_score 2.5532484335057877
```

You're still off by about \$2,550



### 3.6.5 Wrapping up

- ▶ Here's what you should take away from this example:
- Mean squared error (MSE) is a loss function commonly used for regression.
- The concept of accuracy doesn't apply for regression. A common regression metric is mean absolute error (MAE). output 노드의 개수는 1개, activation 함수 따로 없음
- When features in the input data have values in different ranges, each feature should be scaled independently as a preprocessing step.
- When there is little data available, using K-fold validation is a great way to reliably evaluate a model.
- When little training data is available, it's preferable to use a small network with few hidden layers (typically only 1 or 2), in order to avoid severe overfitting.



### Chapter summary

- binary classification, multiclass classification, and scalar regression
- preprocess raw data before feeding it into a neural network.
- features with different ranges, scale each feature independently
- As training progresses, neural networks eventually begin to overfit on never-before-seen data.
- If you have small training data, use a small network with only one or two hidden layers, to avoid severe overfitting.
- If your data is divided into many categories, you may cause information bottlenecks if you make the intermediate layers too small.
- Regression uses different loss functions and different evaluation metrics than classification.
- When you're working with little data, K-fold validation can help reliably evaluate your model.