

# VL Deep Learning for Natural Language Processing

06. Text Classification

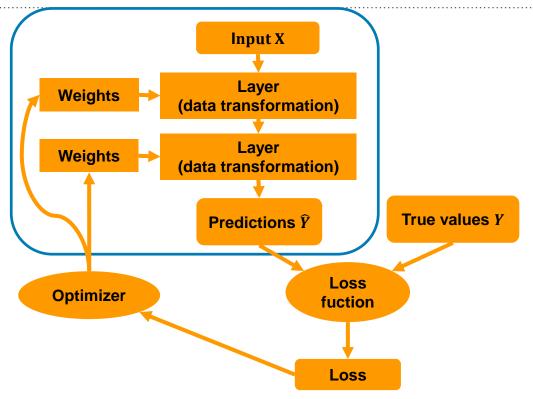
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# Anatomy of a Neural Network









# Learning Goals for this Chapter





- Use neural nets to solve simple problems
  - Binary classification
  - Multi-Class Classification
  - Regression
- Understand ensembles

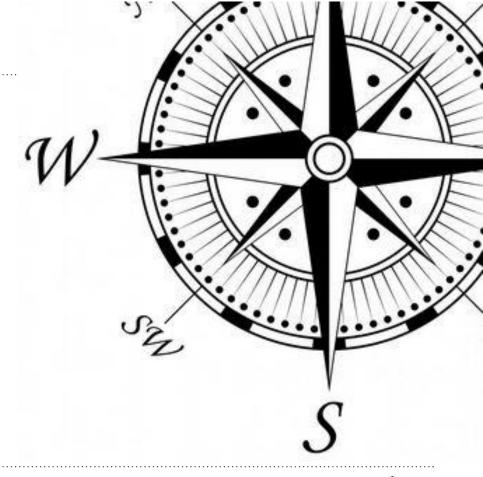
- Relevant chapters:
  - P3, D8





# **Topics Today**

- 1. Classification of Movie Reviews
- 2. Classification of News Articles
- 3. Prediction of Real Estate Prices
- 4. Ensembles







### Loading the Data

from keras.datasets import imdb



```
(train data, train labels), (test data, test labels) = imdb.load data(num words=10000)
                                                           Only the top-10k most
                                           0=negative
                               Word
                                                              frequent words
                              indices
                                           1=postitive
def get review text(sample):
        word index = imdb.get word index()
        reverse word index = dict([(value, key) for (key, value) in word index.items()])
        decoded review = ' '.join([reverse word index.get(i - 3, '?')
                               for i in train data[sample]])
        return decoded review
```



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0=padding; 1=startMarker; 2=unknown

# Preprocessing of Textual Data



Two options to represent text

#### 1. Sequence of words

- Ordering is retained
- Input tensor of shape (samples, wortIndices)
- Input texts need have the same length → padding
- Next layert: embedding layer

#### 2. One-hot encoding

- Ordering is lost
- Frequency of words gets lost
- Input tensor of shape (samples, vocabularySize)

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- o Almost all entries in input vector are 0
- Next layer: dense layer



#### Vectorization



```
import numpy as np
def vectorize sequences(sequences, dimension=10000):
       results = np.zeros((len(sequences), dimension))
       for i, sequence in enumerate(sequences):
               results[i, sequence] = 1.
       return results
x train = vectorize sequences(train data)
x test = vectorize sequences(test data)
y train = np.asarray(train labels).astype('float32')
y test = np.asarray(test labels).astype('float32')
x val = x train[:10000]
partial x train = x train[10000:]
y val = y train[:10000]
partial y train = y train[10000:]
```



### The Neural Network

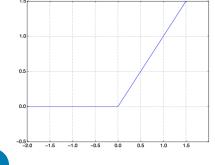


#### 1. Network architecture

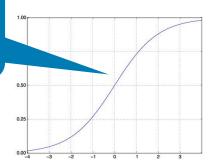
- What kind of layers?
  - Dense (fully connected)
- What kind of activation function?
  - o relu (rectified linear unit) for hidden layers
  - o sigmoid for output layer

#### 2. Configuration

- How many layers?
  - Two hidden layers + one output layer
- How many units per layer?
  - $\circ$  2x16 + 1x1 for output



Output in range [0,1];
Can be interpreted as probabilities!





#### **Activation Function**



- The activation function is key!
- A dense layer only consists of a scalar product and addition:

- Only linear transformation possible
- The hypothesis space of the layer would be all linear transformations from the input data into a space with dimention=number of units of layer.
- Multiple layers in a row would be more powerful.
- What is needed: a non-linearity → activation fuction
   output = relu(dot(W, input) + b)
- relu is the most popular one
  - Others, less popular ones include prelu, elu, etc.



#### The Network



```
from keras import models
from keras import layers
model = models.Sequential()
model.add(layers.Dense(16, activation='relu', input shape=(10000,)))
model.add(layers.Dense(16, activation='relu'))
model.add(layers.Dense(1, activation='sigmoid'))
model.compile(optimizer='rmsprop', loss='binary crossentropy'
metrics=['accuracy'])
                                             model.compile(optimizer='rmsprop',
                                             loss='binary crossentropy',
from keras import optimizers
                                             metrics=['accuracy'])
from keras import losses
from keras import metrics
model.compile(optimizer=optimizers.RMSprop(lr=0.001),
       loss=losses.binary crossentropy,
       metrics=[metrics.binary accuracy])
```



# Training the Model



```
history = model.fit(partial_x_train,
partial_y_train,
epochs=20,
batch_size=512,
validation_data=(x_val, y_val))

>>> history_dict = history.history
>>> history_dict.keys()
[u'acc', u'loss', u'val acc', u'val loss']
```



# Training and Validation Loss



```
import matplotlib.pyplot as plt
                                                                    Training and validation loss
history dict = history.history
loss values = history dict['loss']
val loss values = history dict['val loss']
epochs = range(1, len(loss values) + 1)
plt.plot(epochs, loss values, 'bo', label='Training loss')
plt.plot(epochs, val loss values, 'b', label='Validation loss')
plt.title('Training and validation loss')
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.legend()
plt.show()
```



# Training and Validation Accuracy



```
Training and validation accuracy
plt.clf()
                                                               0.95
acc values = history dict['acc']
                                                              0.90
val acc values = history dict['val acc']
                                                               0.85
                                                                                  15.0
                                                                            Epochs
plt.plot(epochs, acc values, 'bo', label='Training acc')
plt.plot(epochs, val acc values, 'b', label='Validation acc')
plt.title('Training and validation accuracy')
plt.xlabel('Epochs')
plt.ylabel('Accuracy')
plt.legend()
plt.show()
```



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# Overfitting



- Classical example of over optimization
  - Best validation accuracy after 3 epochs
  - Thus, e.g. stop training early (early stopping)

```
Training and validation accuracy

1.00
Training acc
Validation acc

0.95

0.80

2.5 5.0 7.5 10.0 12.5 15.0 17.5 20.0 Epochs
```

```
model = models.Sequential()
model.add(layers.Dense(16, activation='relu', input_shape=(10000,)))
model.add(layers.Dense(16, activation='relu'))
model.add(layers.Dense(1, activation='sigmoid'))
model.compile(optimizer='rmsprop', loss='binary_crossentropy',
metrics=['accuracy'])
model.fit(x_train, y_train, epochs=4, batch_size=512)
results = model.evaluate(x_test, y_test)
>>> results
[0.2929924130630493, 0.8832799999999995]
>>> model.predict(x_test)
array([[ 0.98006207][ 0.99758697]...,[ 0.65371346]], dtype=float32)
```



# Take Away Message



- Preprocessing is a very important step
  - Different representations of the input
  - Lead to different architectures,
  - Leads to different results
- Multiple dense layers with relu are suitable for a variety of task
- For binary classification the output layer should be dense and should contain one unit with a sigmoid activation function
- Scalar sigmoid output + binary classification = binary cross-entropy as loss function
- Rmsprop Optimizer works well in general
- Overfitting needs to avoided by monitoring training progress and appropriate actions



# **Binary Classification**





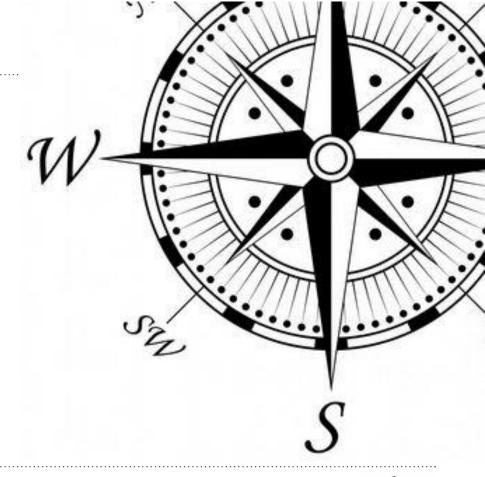
- Reimplement the given example
- Experiment with different settings
  - Network architecture
    - Number of layers (+/- 1), Units (8/32)
  - Loss function (mse)
  - Activation function (tanh)
- State-of-the-art models achieve 95% accuracy





# **Topics Today**

- 1. Classification of Movie Reviews
- 2. Classification of News Articles
- 3. Prediction of Real Estate Prices
- 4. Ensembles





### Preprocessing



```
from keras.datasets import reuters
(train data, train labels), (test data, test labels) =
reuters.load data(num words=10000)
import numpy as np
def vectorize sequences(sequences, dimension=10000):
       results = np.zeros((len(sequences), dimension))
       for i, sequence in enumerate(sequences):
               results[i, sequence] = 1.
       return results
x train = vectorize sequences(train data)
                                                     Vectorization
x test = vectorize sequences(test data)
x val = x train[:1000]
partial x train = x train[1000:]
y val = y train[:1000]
partial y train = y train[1000:]
```



# **Encoding Multi-Class Labels**



```
def to one hot(labels, dimension=46):
       results = np.zeros((len(labels), dimension))
       for i, label in enumerate(labels):
               results[i, label] = 1.
       return results
one hot train labels = to one hot(train labels)
                                                                 Alternative I:
one hot test labels = to one hot(test labels)
from keras.utils.np utils import to categorical
one hot train labels = to categorical(train labels)
                                                               Alternative II:
one hot test labels = to categorical(test labels)
```



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#### The Network



```
from keras import models
                                    units wouldn't be
from keras import layers
                                       enough!
model = models.Sequential()
model.add(layers.Dense(64, activation='relu', input shape=(10000,)))
model.add(layers.Dense(64, activation='relu'))
model.add(layers.Dense(46, activation='softmax'))
model.compile(optimizer='rmsprop',
loss='categorical crossentropy',
metrics=['accuracy'])
```

For 46 classes, 16

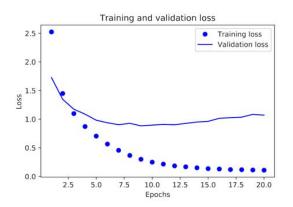
**Probability** distribution over all classes  $\sum_{c} p(c|d) = 1$ 

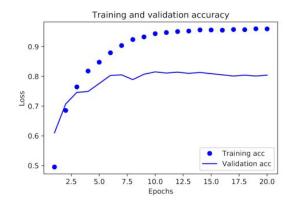
Measures the distance between two probability distributions



# Training the Model









#### Results



```
model.fit(partial x train, partial y train, epochs=9, batch size=512,
validation data=(x val, y val))
results = model.evaluate(x test, one hot test labels)
>>> results
[0.9565213431445807, 0.79697239536954589]
>>> import copy
                                                        Baseline:
>>> test labels copy = copy.copy(test labels)
                                                         random
>>> np.random.shuffle(test labels copy)
>>> hits array = np.array(test labels) == np.array(test labels copy)
>>> float(np.sum(hits array)) / len(test labels)
0.18655387355298308
```



### **Alternative Encoding of Labels**



```
y train = np.array(train labels)
                                                  In case the
y test = np.array(test labels)
                                                category labels
                                                 are integers
model.compile(optimizer='rmsprop',
loss='sparse categorical crossentropy', metrics=['acc'])
predictions = model.predict(x test)
>>> predictions[0].shape
(46,)
>>> np.sum(predictions[0])
1.0
>>> np.argmax(predictions[0])
```



# Take Away Message



- The output layer should have as many units as there are different classes.
- For single-label, multi-class problems, use the softmax activation function
- Categorical cross-entropy minimizes the distance between the true distribution of the labels and the predicted distribution of the network.
- One-hot encoding or using integer
  - Loss function has different name
- Avoid information bottlenecks!



#### Multi-Class Classification





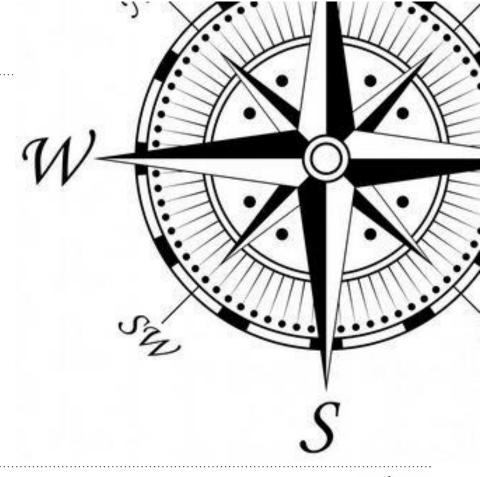
- Reimplement the given example with the 20 news groups dataset.
- Create an information bottleneck by assigning only 4 or 8 units to the second hidden layer.
- Experiment with different network sizes
  - layers (+/- 1)
  - units (32/128)





# **Topics Today**

- 1. Classification of Movie Reviews
- 2. Classification of News Articles
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### Loading the Data



```
from keras.datasets import boston_housing
(train_data, train_targets), (test_data, test_targets) =
boston_housing.load_data()

>>> train_data.shape
(404, 13)
>>> test_data.shape
(102, 13)

Features contain e.g. crime
rate, number of rooms,
highway access, etc.

>>> train_targets
[ 15.2, 42.3, 50. ... 19.4, 19.4, 29.1]
```

Average price in 1000\$



### Normalizing 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
```

NEVER look at the test data! Not even for preprocessing or normalization!





#### The Network



```
from keras import models
                                               No activation
from keras import layers
                                              function: Linear
def build model():
                                               output layer
       model = models.Sequential()
       model.add(layers.Dense(64, activative='relu',
input shape=(train data.shape[1],)))
       model.add(layers.Dense(64, accivation='relu'))
       model.add(layers.Dense(1))
       model.compile(optimizer='rmsprop', loss='mse', metrics=['mae'])
       return model
                                            Mean Squared
                                                                 Mean Absolute
                                                Error
                                                                     Error
```



#### K-Fold Cross-Validation



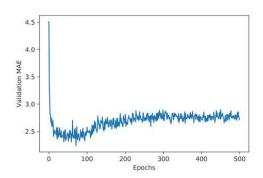
```
import numpy as np; k=4; num val samples=len(train data) // k; num epochs=100;
all scores=[]
for i in range(k):
       print('processing fold #', i)
       val data = train data[i * num val samples: (i + 1) * num val samples]
       val targets = train targets[i * num val samples: (i + 1) * num val samples]
       partial train data = np.concatenate([train data[:i * num val samples],
                      train data[(i + 1) * num val samples:]], axis=0)
       partial train targets = np.concatenate([train targets[:i * num val samples],
                      train targets[(i + 1) * num val samples:]], axis=0)
       model = build model()
                                                                          Silent
       history = model.fit(partial train data, partial train targets,
                                                                          mode
                      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)
```



#### Results



- For num\_epochs=100
  - MAE-values for each fold [2.5882589577920, 3.12895684497191, 3.18561160512489, 3.07633426154013]
  - Average2.9947904173572462
- Monitoring the individual folds for 500 epochs





#### **Better Visualization**



```
Moving
def smooth curve(points, factor=0.9):
                                                              2.80
                                             Average!
                                                              2.75
     smoothed points = []
                                                              2.70
     for point in points:
                                                             ₩ 2.65
        if smoothed points:
                                                              2.60
             previous = smoothed points[-1]
                                                             ® 2.55
                                                              2.50
             smoothed points.append(previous * factor
                                                              2.45
                           + point * (1 - factor))
        else:
             smoothed points.append(point)
                                                                             Ignore first
     return smoothed points
                                                                             10 points!
smooth mae history = smooth curve(average mae history[10:])
plt.plot(range(1, len(smooth mae history) + 1), smooth mae history)
plt.xlabel('Epochs')
plt.ylabel('Validation MAE')
plt.show()
```



### **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)

>>> test_mae_score
2.5532484335057877
```



# Take Away Message



- For regression problems we need a different loss function.
  - MSE (Mean Squared Error) is very popular.
- Evaluation metric for regression problems:
  - MAE (Mean Absolute Error)
- Features with values in different ranges need to be normalized independentley of each other.
- If there are only a few training samples, use k-fold cross-validation.
- If there are only a few training samples, the network needs to be small,
   i.e. it should have only a low number of trainable parameters.
  - One or two hidden layers the most



### Regression





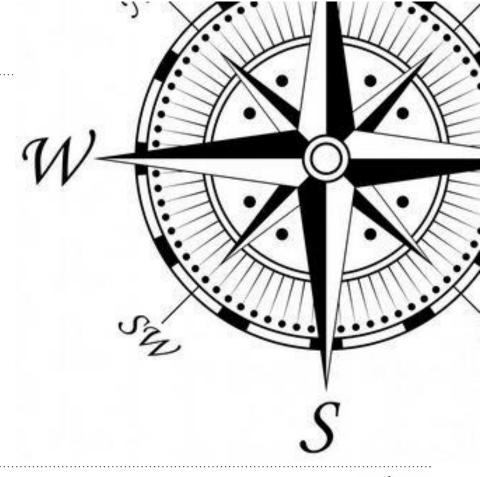
- Reimplement the example with one of the following datasets:
  - https://www.kaggle.com/datasets/eswarchandt/amazon-music-reviews
  - https://www.kaggle.com/datasets/septa97/100k-courseras-course-reviews-dataset
  - https://www.kaggle.com/datasets/andrewmvd/trip-advisor-hotel-reviews
- Experiment with different network sizes
  - layers (+/- 1)
  - units (32/128)





# **Topics Today**

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### Ensembling



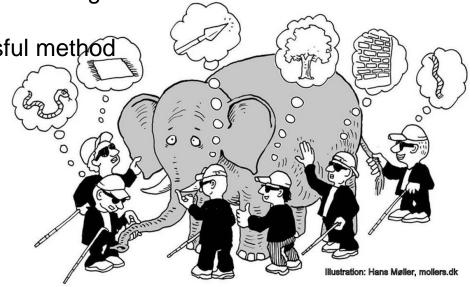
- An Ensemble is the combination of multiple models.
  - The predictions of different, diverse models are combined.

Each model "grasps" one aspect of the training data.

The blind and the elephant

In practice (e.g. Kaggle) very successful method

- General machine learning approach
  - (Election)
  - Bagging
  - Boosting
  - Stacking







#### In the Wild



- Better than just taking the average:
  - Learn weights using validation data!
    - E.g. with the Nelder-Mead-method (similar to steepest descent)

```
preds_a = model_a.predict(x_val)
preds_b = model_b.predict(x_val)
preds_c = model_c.predict(x_val)
preds_d = model_d.predict(x_val)
final_preds = 0.5*preds_a + 0.25*preds_b + 0.1*preds_c + 0.15*preds_d
```

- Goal: Many diverse models
  - The can have a high bias, as long as it differs from model to model.
  - The same network with different initializations does not work.
- Often promising:
  - Ensembles from decision trees
    - random forest or gradient boosting trees
  - and deep neural networks



# Learning Goals for this Chapter





- Use neural nets to solve simple problems
  - Binary classification
  - Multi-Class Classification
  - Regression
- Understand ensembles

- Relevant chapters:
  - P3, D8



