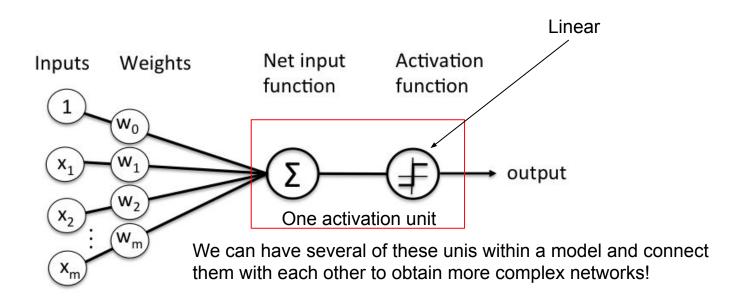
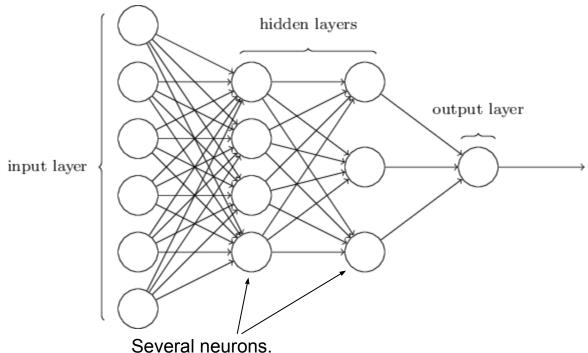
Multi Layer Perceptron

Concept and implementation in TensorFlow/Keras

Single Layer ANN



Multi-Layer ANN



TensorFlow & Keras

Computer Power

- Computer power has increased rapidly allowing us to train very complex and powerful learning systems and so to improve the predictive performance of our machine learning models.
- We can even take advantage of multi-core CPUs and spread the computations over multiple processing units.
- This is even possible with your laptop or desktop computer!
- However, even the CPU with the most number of processing units is overloaded when the task is to train very complex deep learning models where we need to learn > Millions of parameters. Solution?
- GPUs instead of CPUs!

Calling GPUs

- Challenge: Writing code to target GPUs
- Special packages such as CUDA and OpenCL, however writing code in those packages is hard.
- TensorFlow helps us to overcome the challenge

TensorFlow

What is TensorFlow

- TensorFlow is a multiplatform programming interface for implementing and running machine learning algorithms with wrappers for deep learning
- Developed by the researchers and engineers of the Google Brain team -- contributions happen also through open source communities.
- Was developed for Google internal use but was released in 2015 for public under a permissive open source licence.
- TensorFlow runs on both CPU and GPU (it shines however with GPUs)
- Supports CUDA-enabled GPUs however, OpenCL will likely be supported in near future
- There are support for a number of programming languages such as Python.

Companies using TensorFlow

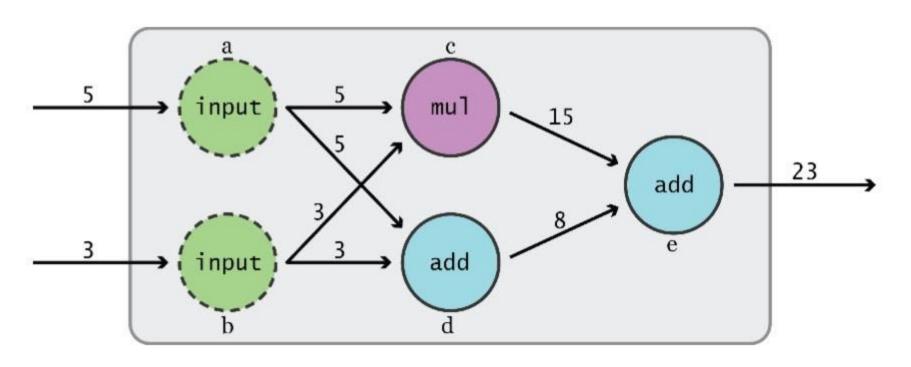
- Google
- OpenAl
- DeepMind
- Snapchat
- Uber
- Airbus
- eBay
- Dropbox
- A bunch of startups
- And we:)

Get Started with TensorFlow

Import TensorFlow

import tensorflow as tf

TensorFlow Graph and Sessions



Edges and Nodes in the Graph

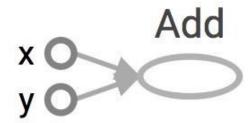
Edges in the graph are the tensors! In other words, tensors are DATA.

An n-dimensional array

- 0-d tensor: scalar (number)
- 1-d tensor: vector
- 2-d tensor: matrix
- and so on

Nodes in the graph are the operators, variables, and constants

import tensorflow as tf a = tf.add(3, 5)



Why x, y?

TF automatically names the nodes when you don't explicitly name them.

$$x = 3$$

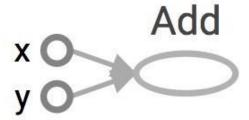
$$y = 5$$

What is the output of a?

import tensorflow as tf

a = tf.add(3, 5)

print a



What is the output of a?

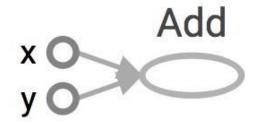
import tensorflow as tf

a = tf.add(3, 5)

print a

>> Tensor("Add:0", shape=(), dtype=int32)

Not 8!



How to get the value of a?

Two steps:

Create a session, assign it to variable sess so we can call it later.

Within the session, evaluate the graph to fetch the value of a.

```
import tensorflow as tf a = tf.add(3, 5)
```

```
sess = tf.Session()
print sess.run(a)
sess.close()
```

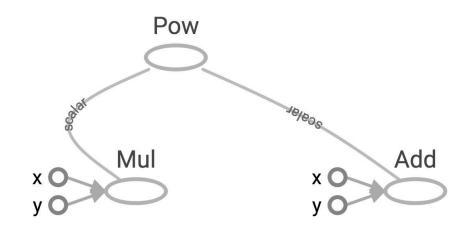


Or

```
with tf.Session() as sess: print sess.run(a)
```

More Graphs

```
x = 2
y = 3
op1 = tf.add(x, y)
op2 = tf.mul(x, y)
op3 = tf.pow(op2, op1)
with tf.Session() as sess:
    op3 = sess.run(op3)
```



Keras

Keras

TensorFlow can sometimes be hard to code.

Keras comes to rescue:

- Built on top of TensorFlow
- Simple to get started, simple to keep going
- Written in python and highly modular; easy to expand
- Deep enough to build serious models
- General idea is to based on layers and their input/output
- The layers are computed in sequences (but there is also graph structures)

Keras

https://keras.io/

We will be using *Conda* to manage our course exercises, and *Python 3* for all of our code

Conda provides an easy way to manage environments, so all of the course dependencies can be installed without any hassle

```
# Get the appropriate package for your Operating System
https://repo.continuum.io/miniconda/Miniconda3-latest-Linux-x86_64.sh
https://repo.continuum.io/miniconda/Miniconda3-latest-MacOSX-x86_64.sh
https://repo.continuum.io/miniconda/Miniconda3-latest-Windows-x86_64.exe
```

- # Create an environment for the course exercises
- > conda create --name dl4nlp python=3.6

```
# Activate the environment
# You will ALWAYS need to do this before running anything
related to the course
```

```
# For linux/macOS
```

> source activate dl4nlp

```
# For windows
```

> activate dl4nlp

```
# Install dependencies
```

- > conda install keras numpy matplotlib jupyter
 scikit-learn pydot graphviz nltk nb_conda
- > conda install -c conda-forge ffmpeg

- # Run python code in this environment
- > python some_code.py

- # We also usually just work in Jupyter/iPython notebooks
- > jupyter notebook

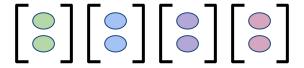
This opens a tab in your browser where you can create notebooks, write and run code

Setup

```
# Deactivate environment
# Once you are done for the day, its usually good
practice to deactivate the environment
```

- # For linux/macOS
- > source deactivate

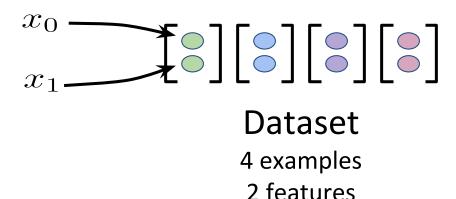
- # For windows
- > deactivate



Dataset

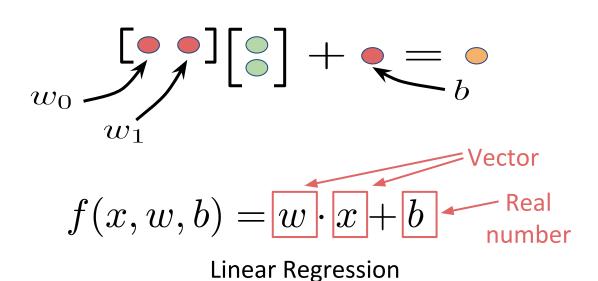
4 examples

2 features



4 examples 2 features





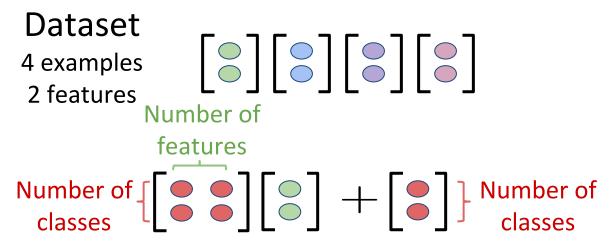
Dataset

4 examples

2 features

$$W \longrightarrow \begin{bmatrix} \vdots \\ \vdots \\ b \end{bmatrix} + \begin{bmatrix} \vdots \\ b \end{bmatrix}$$

$$f(x, W, b) = W \cdot x + b$$



$$f(x,W,b) = W \cdot x + b$$

3 class classification

$$f(x, W, b) = W \cdot x + b$$

Dataset

4 examples

2 features

$$\begin{bmatrix} \bullet & \bullet \end{bmatrix} \begin{bmatrix} \bullet & \bullet \end{bmatrix} + \begin{bmatrix} \bullet & \bullet \end{bmatrix} = \begin{bmatrix} \bullet & \bullet \end{bmatrix}$$

In this case, we are performing the above computation per example

$$f(x, W, b) = W \cdot x + b$$

Dataset

4 examples

2 features

$$\begin{bmatrix} \bullet & \bullet \end{bmatrix} \begin{bmatrix} \bullet & \bullet \end{bmatrix} + \begin{bmatrix} \bullet & \bullet \end{bmatrix} = \begin{bmatrix} \bullet & \bullet \end{bmatrix}$$

In this case, we are performing the above computation per example

$$f(x, W, b) = W \cdot x + b$$

Dataset

4 examples 2 features

$$\begin{bmatrix} \bullet & \bullet \end{bmatrix} \begin{bmatrix} \bullet & \bullet \end{bmatrix} + \begin{bmatrix} \bullet & \bullet \end{bmatrix} = \begin{bmatrix} \bullet & \bullet \end{bmatrix}$$

In this case, we are performing the above computation per example

$$f(x, W, b) = W \cdot x + b$$

Dataset

4 examples 2 features

$$\begin{bmatrix} \bullet & \bullet \end{bmatrix} \begin{bmatrix} \bullet & \bullet \end{bmatrix} + \begin{bmatrix} \bullet & \bullet \end{bmatrix} = \begin{bmatrix} \bullet & \bullet \end{bmatrix}$$

In this case, we are performing the above computation *per example*

$$f(x, W, b) = W \cdot x + b$$

Dataset
4 examples

2 features

What if we can process all the examples in one go?

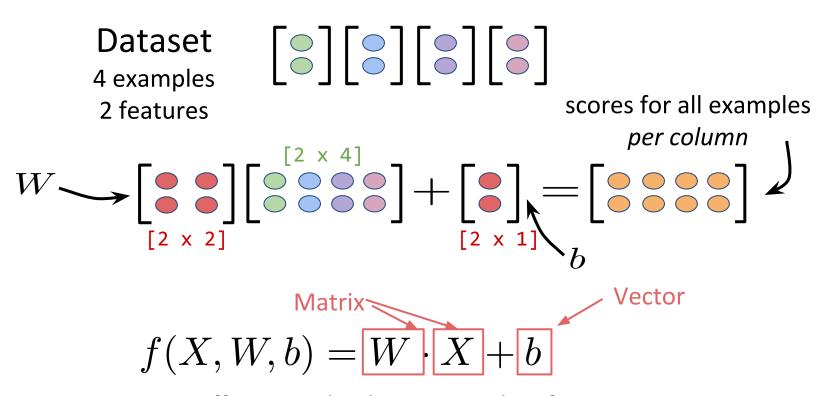
$$f(x, W, b) = W \cdot x + b$$

Dataset
4 examples
2 features

What if we can process all the examples in one go?

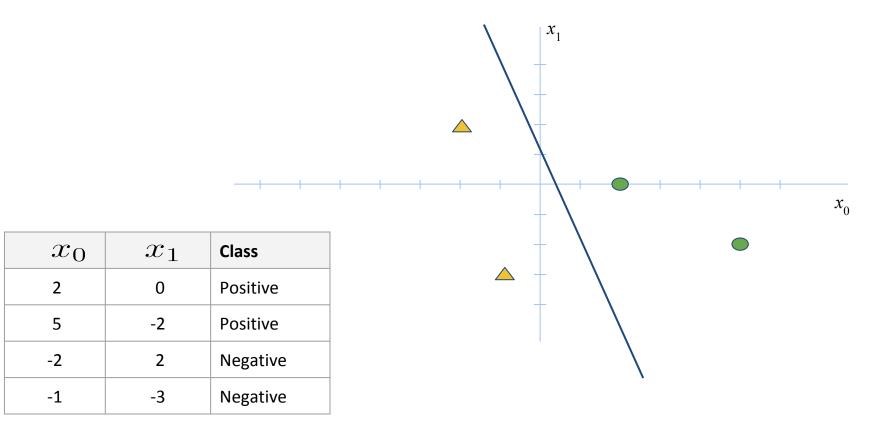
How: Stack all examples into one big matrix!

$$f(x, W, b) = W \cdot x + b$$



Efficient Multi-class Linear Classification

Linear Classifier using Regression



Data setup

```
data = [(2,0),(5,-2),(-2,2),(-1,-3)]
labels = [-1,-1,1,1]
```

- Usually data is loaded from an external source
- Eventually, all data is represented in some structured form like in matrices
- Data for supervised learning is normally composed of the actual data points and the labels for each point

Data setup

```
X = np.array(data)
y = np.array(labels)
```

- Eventually, all data is represented in some structured form like in matrices
- Here, we convert all of our data and labels into Numpy arrays

Model definition

```
model = Sequential()
model.add(Dense(1, input_shape=(2,)))

model.compile(loss="mse", optimizer="sgd", metrics=['acc'])
model.summary()
```

- In this case, Dense is the objective function for a linear classifier
- Dense corresponds to the equation of f which is Wx + b
- loss computes mean squared error

$$f(x, W, b) = w_0 \cdot x_0 + w_1 \cdot x_1 + b$$
$$MSE(x, W, b, y) = (f(x, W, b) - y)^2$$

Model definition

```
model = Sequential()
model.add(Dense(1, input_shape=(2,)))

model.compile(loss="mse", optimizer="sgd", metrics=['acc'])
model.summary()
```

Input shape defines the number of features.

In our case, this is 2

Model definition

```
model = Sequential()
model.add(Dense(1, input_shape=(2,)))

model.compile(loss*"mse", optimizer="sgd", metrics=['acc'])
model.summary()
```

The number of units of the Dense layer - this corresponds to the number of outputs (neuron within a hidden layer).

In our case, we only want to output 1 number (regression)

Model definition

```
model = Sequential()
model.add(Dense(1, input_shape=(2,)))
model.compile(loss="mse", optimizer="sgd", metrics=['acc'])
model.summary()
        Mean squared
                                   Optimization
           error loss
                                     function is
                                 gradient descent
```

```
parameter history = []
for epoch in range(50):
   # Perform one step over the entire dataset
   loss history = model.fit(X, y, epochs=1, verbose=False)
    # Get predictions (value of the objective function, f)
   y pred = model.predict(X, verbose=False)
   # See how well our model is doing
                                                               Optimization loop:
   # Recall our classes are [-1,-1,1,1]
   num correct = 0
                                                               We will run the
    if y pred[0] < 0: num correct += 1</pre>
   if y pred[1] < 0: num correct += 1</pre>
                                                               optimization for 50
    if y pred[2] > 0: num correct += 1
   if y pred[3] > 0: num correct += 1
                                                               epochs
   acc = num correct / 4.0
   loss = loss history.history['loss'][-1]
   print("Epoch %d: %0.2f (acc) %0.2f (loss)"%(epoch+1, acc, loss))
   # Not mandatory: Save parameters for later analysis
   w, b = model.layers[0].get weights()
   parameter history.append((w,b))
```

```
parameter history = []
for epoch in range(50):
   # Perform one step over the entire dataset
                                                                Here we fit over
   loss history = model.fit(X, y, epochs=1, verbose=False)
                                                                our data once.
   # Get predictions (value of the objective function, f)
                                                                In the fit function,
   y pred = model.predict(X, verbose=False)
                                                                Keras automatically
   # See how well our model is doing
   # Recall our classes are [-1,-1,1,1]
                                                                computes the
   num correct = 0
   if y pred[0] < 0: num correct += 1</pre>
                                                                objective function,
   if y pred[1] < 0: num correct += 1</pre>
   if y pred[2] > 0: num correct += 1
                                                                computes the loss
   if y pred[3] > 0: num correct += 1
   acc = num correct / 4.0
                                                                and uses the
   loss = loss history.history['loss'][-1]
   print("Epoch %d: %0.2f (acc) %0.2f (loss)"%(epoch+1, acc, loss
                                                                optimizer to adjust
   # Not mandatory: Save parameters for later analysis
                                                                the parameters of
   w, b = model.layers[0].get weights()
   parameter history.append((w,b))
                                                                the model
```

```
parameter history = []
for epoch in range(50):
    # Perform one step over the entire dataset
                                                              fit also returns the
    loss history = model.fit(X, y, epochs=1, verbose=False)
                                                              history of losses for
    # Get predictions (value of the objective function, f)
   y pred = model.predict(X, verbose=False)
                                                              each epoch, along
   # See how well our model is doing
                                                              with whatever
   # Recall our classes are [-1,-1,1,1]
   num correct = 0
                                                              metrics we
   if y pred[0] < 0: num correct += 1</pre>
    if y pred[1] < 0: num correct += 1</pre>
                                                              requested for when
   if y pred[2] > 0: num correct += 1
                                                              defining the model
    if y pred[3] > 0: num correct += 1
    acc = num correct / 4.0
   loss = loss history.history['loss'][-1]
   print("Epoch %d: %0.2f (acc) %0.2f (loss)"%(epoch+1, acc, loss))
   # Not mandatory: Save parameters for later analysis
   w, b = model.layers[0].get weights()
   parameter history.append((w,b))
```

```
parameter history = []
for epoch in range(50):
   # Perform one step over the entire dataset
    loss history = model.fit(X, y, epochs=1, verbose=False)
    # Get predictions (value of the objective function, f)
                                                               predict takes as
   v pred = model.predict(X, verbose=False)
                                                               input some data
   # See how well our model is doing
                                                               and returns the
    # Recall our classes are [-1,-1,1,1]
   num correct = 0
                                                               value of the
    if y pred[0] < 0: num correct += 1</pre>
   if y pred[1] < 0: num correct += 1</pre>
                                                               objective function
    if y pred[2] > 0: num correct += 1
    if y pred[3] > 0: num correct += 1
                                                               (In this case, one
    acc = num correct / 4.0
   loss = loss history.history['loss'][-1]
                                                              value per data
   print("Epoch %d: %0.2f (acc) %0.2f (loss)"%(epoch+1, acc, 1
                                                               point)
   # Not mandatory: Save parameters for later analysis
   w, b = model.layers[0].get weights()
   parameter history.append((w,b))
```

```
parameter history = []
for epoch in range(50):
    # Perform one step over the entire dataset
    loss history = model.fit(X, y, epochs=1, verbose=False)
    # Get predictions (value of the objective function, f)
   y pred = model.predict(X, verbose=False)
    # See how well our model is doing
    # Recall our classes are [-1,-1,1,1]
    num correct = 0
    if y pred[0] < 0: num correct += 1</pre>
    if y pred[1] < 0: num correct += 1</pre>
    if y pred[2] > 0: num correct += 1
    if y pred[3] > 0: num correct += 1
    acc = num correct / 4.0
    loss = loss_history.history['loss'][-1]
    print("Epoch %d: %0.2f (acc) %0.2f (loss)"%(epoch+1, acc, 1
    # Not mandatory: Save parameters for later analysis
    w, b = model.layers[0].get weights()
    parameter history.append((w,b))
```

Here, we compare the value of the objective function and assign classes. Recall that a value of < 0 is Class 1, and > 0 is Class 2

Usually this is not relevant but helps to see how the model learns.

```
parameter history = []
for epoch in range(50):
   # Perform one step over the entire dataset
   loss history = model.fit(X, y, epochs=1, verbose=False)
   # Get predictions (value of the objective function, f)
   y pred = model.predict(X, verbose=False)
   # See how well our model is doing
                                                               Print the progress.
    # Recall our classes are [-1,-1,1,1]
                                                               The value of the
   num correct = 0
   if y pred[0] < 0: num correct += 1</pre>
                                                               loss should go down
   if y pred[1] < 0: num correct += 1</pre>
   if y pred[2] > 0: num correct += 1
                                                               with each epoch
    if y pred[3] > 0: num correct += 1
   acc = num correct / 4.0
   loss = loss history.history['loss'][-1]
    print("Epoch %d: %0.2f (acc) %0.2f (loss)"%(epoch+1, acc, loss))
   # Not mandatory: Save parameters for later analysis
   w, b = model.layers[0].get weights()
```

parameter history.append((w,b))

```
parameter history = []
for epoch in range(50):
    # Perform one step over the entire dataset
    loss history = model.fit(X, y, epochs=1, verbose=False)
    # Get predictions (value of the objective function, f)
   y pred = model.predict(X, verbose=False)
    # See how well our model is doing
    # Recall our classes are [-1,-1,1,1]
    num correct = 0
    if y pred[0] < 0: num correct += 1</pre>
    if y pred[1] < 0: num correct += 1</pre>
    if y pred[2] > 0: num correct += 1
    if y pred[3] > 0: num correct += 1
    acc = num correct / 4.0
    loss = loss history.history['loss'][-1]
    print("Epoch %d: %0.2f (acc) %0.2f (loss)"%(epoch+1, acc, 1
```

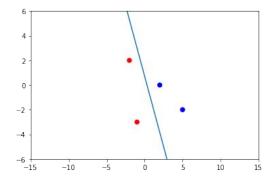
Save parameters after each epoch to visualize later

```
# Not mandatory: Save parameters for later analysis
w, b = model.layers[0].get_weights()
parameter_history.append((w,b))
```

Bonus: Plotting

Bonus: Plotting

Bonus: Plotting



Lets see it in action!



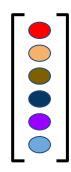
Linear Classification by Regression

Softmax function

In binary classification (as we saw in the example) we can decide what is "1" and what is "-1". When the output was > 0 we took it as "1" otherwise "-1".

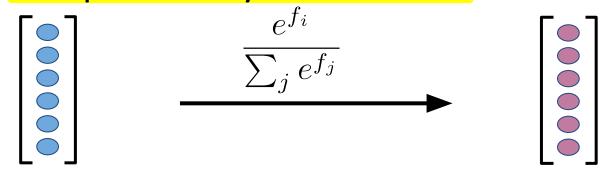
For multi-class classification we can do similar game:

Arg max (
$$\begin{bmatrix} \bullet \\ \bullet \end{bmatrix}$$
) = •, then class 1 is active



However, these scores are not *interpretable*. Their absolute values don't give us any insight, we can only compare them relatively

The softmax function helps us transform these values into probability distributions:



Scores from the classifier

f

Scores as a probability distribution

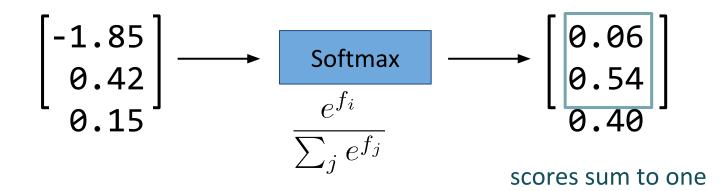
The softmax function helps us transform these values into probability distributions:

$$\begin{bmatrix} -1.85 \\ 0.42 \\ 0.15 \end{bmatrix} \longrightarrow \underbrace{\begin{array}{c} \text{Softmax} \\ e^{f_i} \\ \hline \sum_{j} e^{f_j} \end{array}} \longrightarrow \begin{bmatrix} 0.06 \\ 0.54 \\ 0.40 \end{bmatrix}$$

The softmax function helps us transform these values into probability distributions:

each output can be treated as the

probability of that class



Recall MSE:

Mean Squared Error

$$L = \sum_{i=1}^{n} (f_i - y_i)^2$$

Recall MSE:

Mean Squared Error

In practice, we use *Cross Entropy loss*, which generally performs better for more complex models.

$$H_y(f) = -\sum_i y_i \log(f_i)$$

Here, y represents the true probability distribution (so $y_i = 1$ for the correct class i, and 0 otherwise)

 f_i represents the score of class i from our classifier

$$H_y(f) = -\sum_i y_i \log(f_i)$$
$$= -y_c \log(f_c)$$

Simplifying for our case,

if c is the correct class, then $y_c = 1$, and all other y_i 's are 0Therefore, we only have one element left from the summation

$$H_y(f) = -\sum_i y_i \log(f_i)$$
$$= -y_c \log(f_c)$$
$$= -\log(f_c)$$

Mean Squared Error

Cross Entropy

$$L = \sum_{i=1}^{n} (f_i - y_i)^2$$

$$L = -\log(f_c)$$

Why cross entropy?

Consider three people, Person1 is a *Democrat*, Person2 is a *Republican* and Person3 is *Other*. We have two models to classify these people:

	S _{Other}	S _{Republican}	S _{Democrat}
Person1	0.3	0.3	0.4
Person2	0.3	0.4	0.3
Person3	0.1	0.2	0.7

	S _{Other}	S _{Republican}	S _{Democrat}
Person1	0.1	0.2	0.7
Person2	0.1	0.7	0.2
Person3	0.3	0.4	0.3

Model 1 Model 2

	S _{Other}	S _{Republican}	S _{Democrat}
Person1	0.3	0.3	0.4
Person2	0.3	0.4	0.3
Person3	0.1	0.2	0.7

	S _{Other}	S _{Republican}	S _{Democrat}
Person1	0.1	0.2	0.7
Person2	0.1	0.7	0.2
Person3	0.3	0.4	0.3

Model 1 Model 2

Both models misclassify *Person3*, but is one model better than the other?

	S _{Other}	S _{Republican}	S _{Democrat}
Person1	0.3	0.3	0.4
Person2	0.3	0.4	0.3
Person3	0.1	0.2	0.7

	S _{Other}	S _{Republican}	S _{Democrat}
Person1	0.1	0.2	0.7
Person2	0.1	0.7	0.2
Person3	0.3	0.4	0.3

Model 1 Model 2

Model 2 is better, since it classifies *Person1* and *Person2* with higher scores on the correct class, and mis-classifies *Person3* with a smaller error in the scores

	S _{Other}	S _{Republican}	S _{Democrat}
Person1	0.3	0.3	0.4
Person2	0.3	0.4	0.3
Person3	0.1	0.2	0.7

	S _{Other}	S _{Republican}	S _{Democrat}
Person1	0.1	0.2	0.7
Person2	0.1	0.7	0.2
Person3	0.3	0.4	0.3

Model 1

Mean Squared Error

Model 2

Person1: 0.54

Person2: 0.54

Person3: 1.34

Model 1 Average: 0.81

Person1: 0.14

Person2: 0.14

Person3: 0.74

Model 2 Average: 0.34

	S _{Other}	S _{Republican}	S _{Democrat}
Person1	0.3	0.3	0.4
Person2	0.3	0.4	0.3
Person3	0.1	0.2	0.7

	S _{Other}	S _{Republican}	S _{Democrat}
Person1	0.1	0.2	0.7
Person2	0.1	0.7	0.2
Person3	0.3	0.4	0.3

Model 1

Cross Entropy

Model 2

Person1: $-\log(0.4) = 0.92$

Person1: 0.36

Person2: $-\log(0.4) = 0.92$

Person2: 0.36

Person3: $-\log(0.1) = 2.30$

Person3: 1.20

Model 1 Average: 1.38

Model 2 Average: 0.64

	S _{Other}	S _{Republican}	S _{Democrat}
Person1	0.3	0.3	0.4
Person2	0.3	0.4	0.3
Person3	0.1	0.2	0.7

	S _{Other}	S _{Republican}	S _{Democrat}
Person1	0.1	0.2	0.7
Person2	0.1	0.7	0.2
Person3	0.3	0.4	0.3

Model 1

Model 2

Mean Squared Error

Model 1 Average: 0.81

Model 2 Average: 0.34

Cross Entropy

Model 1 Average: 1.38

Model 2 Average: 0.64

Mean Squared Error

Model 1 Average: 0.81

Model 2 Average: 0.34

Cross Entropy

Model 1 Average: 1.38

Model 2 Average: 0.64

Cross Entropy Loss difference between the two models is greater than the Mean Squared Error!

In general, *Mean Squared Error* penalizes incorrect predictions much more than *Cross Entropy*

A more principled reason arises from the underlying mathematics of MSE and Cross Entropy

MSE causes the gradients to become very small as the network scores become better, so learning slows down!

Cross Entropy is mathematically defined to compare two probability distributions

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Our ground truth is already represented as a probability distribution (with all the probability mass on the correct class)

$$y = \begin{bmatrix} 0.00 \\ 1.00 \\ 0.00 \end{bmatrix}$$

Cross Entropy is mathematically defined to compare two probability distributions

However, the scores directly from a linear classifier do not form any such distribution:

$$f = \begin{bmatrix} -1.85 \\ 0.42 \end{bmatrix}$$

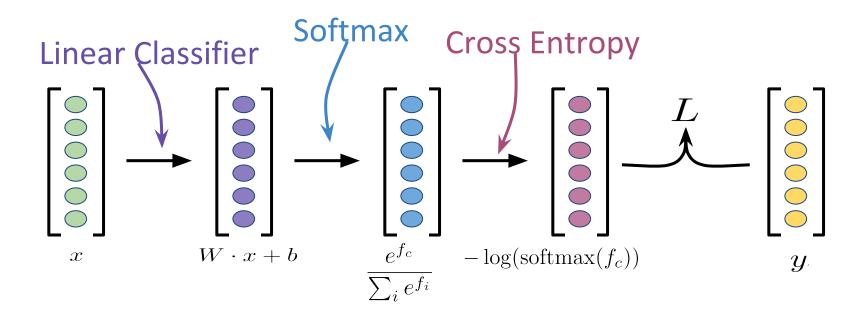
$$0.15$$

Cross Entropy is mathematically defined to compare two probability distributions

Solution: Use softmax!

$$softmax(f) = \begin{bmatrix} 0.06 \\ 0.54 \\ 0.40 \end{bmatrix}$$

Putting it all together



Binary classifier in Keras

Lets see it in action!



Binary Classification

Multi-class classification



Multi-class Classification