Tensorflow and Keras

CSE 4309 – Machine Learning
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Pros and Cons of Using a Library

- Many machine learning methods are already implemented in programming libraries.
- A common question is: which library is the "best".
- The answer depends a lot on the context.
- When studying and trying to understand a method, sometimes the answer is "no library".
 - It is important to do some implementations from scratch.
 - In such an implementation, every single detail must be explicitly addressed.
 - Oftentimes, implementations expose issues that we have not yet fully understood, and that we need to revisit.
- Using a library can boost productivity, minimize coding time.
- Library code can also be optimized to run very efficiently.

Keras, Tensorflow, PyTorch

- Tensorflow is a Python programming library for implementing neural networks.
- Keras is an additional library, that is implemented on top of Tensorflow.
- Keras provides a more simple, high-level interface than Tensorflow.
- Tensorflow allows you to do more things "under the hood", if you do not want to simply implement standard "off-the-shelf" methods.
- PyTorch is an alternative to Tensorflow and Keras for implementing neural networks, that is also commonly used.

Installing Keras on Your Computer

- Install Anaconda
- Open Anaconda Prompt or Anaconda Power Shell
- Type:

pip install --upgrade tensorflow

- You can use the Spyder editor (comes with Anaconda) to write code, test and debug.
- Some people recommend that you create an Anaconda "environment" and install Keras within the environment.
 - That is useful for computers where Keras is installed multiple times, possibly by different users, and Keras is combined in multiple ways with different other packages.
 - For this class, if you use your own computer, this is not needed.

Running Keras Online

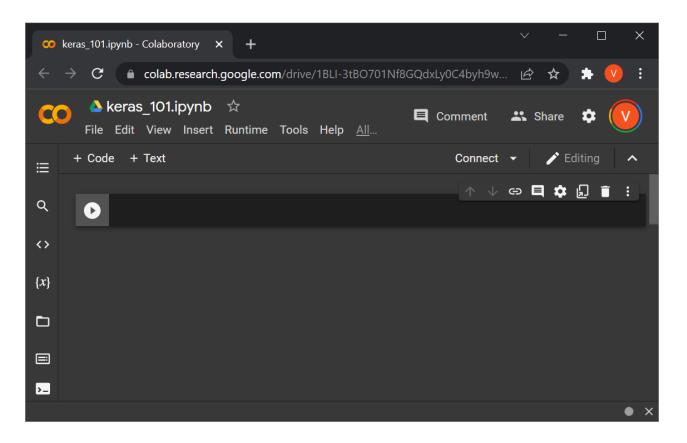
 Google Colab is an online environment where you can write and run Python and Keras code.

https://colab.research.google.com/

- Keras is already installed, so this is very convenient.
- It is a useful job skill to get familiar with Colab.
 - Easy to pick up, if you do everything else you need to do in this class.
- Given that there are many different Python libraries with many different versions, that may not be backward compatible, using Colab ensures that we can run your code.
- Most of the times, the code I write on Anaconda runs on Colab with no problems.

Colab Basics

- Go to Google Colab, sign in with your Google Account.
- Go to File->New notebook
- Give a name to your notebook, like keras_101.ipynb.



If Using Colab, Some Prep

- On Google Drive, I created a top-level folder called cse4392.
- In that folder, I copied:
 - uci_data.py
 - This is posted on the website, under this lecture, and contains some helper code.
 - The uci_datasets directory with all the text files.
- Then, on Google Colab:
 - Create and run a cell with this code:

from google.colab import drive drive.mount('/content/drive')

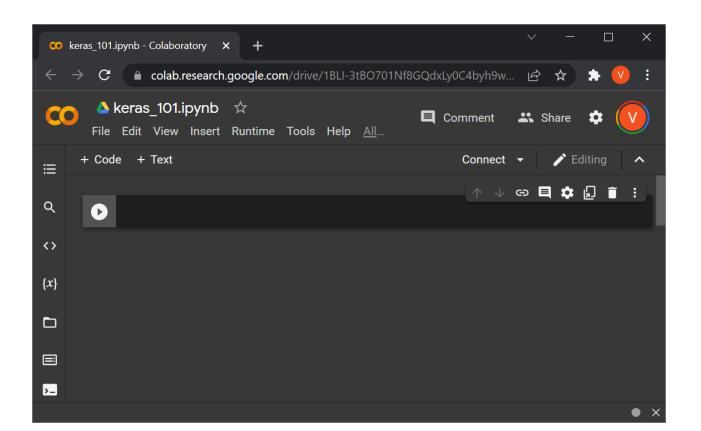
— Create and run another cell with this code:

cd /content/drive/MyDrive/cse4392

The next slides show these steps in more detail.

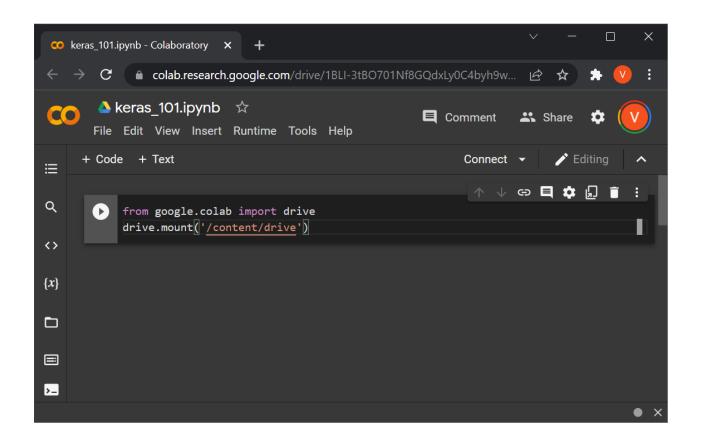
Code Cells in Colab

- To create a new code cell in Colab, click the "+ Code" button.
- To run the code on that cell, click the "play" button to the left of the cell, or press SHIFT+ENTER.

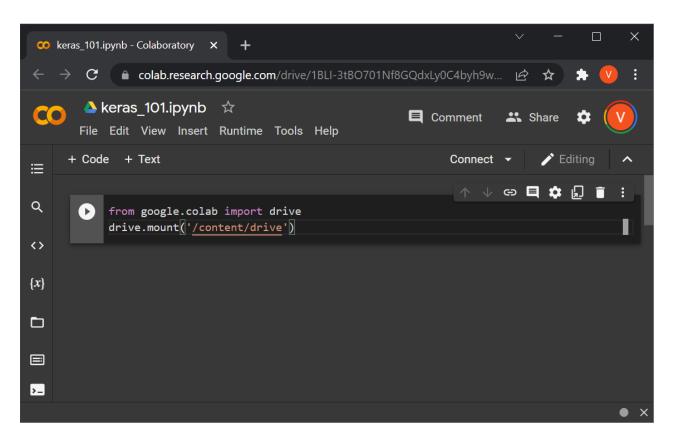


Pasting Code to Colab

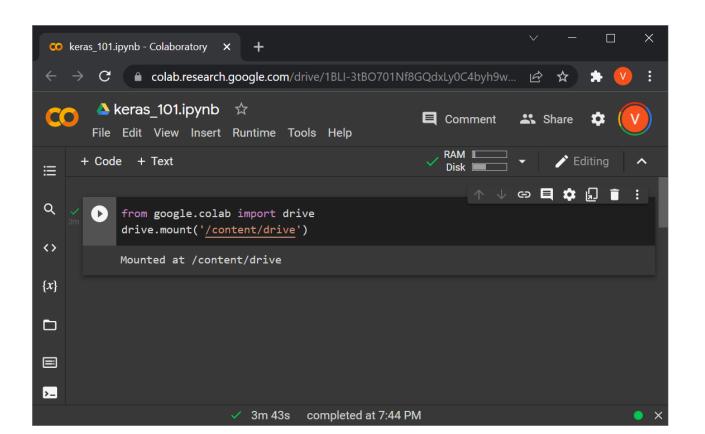
- Pasting code to Colab from Powerpoint does not work for me.
- First I paste from Powerpoint to a Google Drive text file.
- Then I copy-paste from the Google Drive file to the Colab notebook.



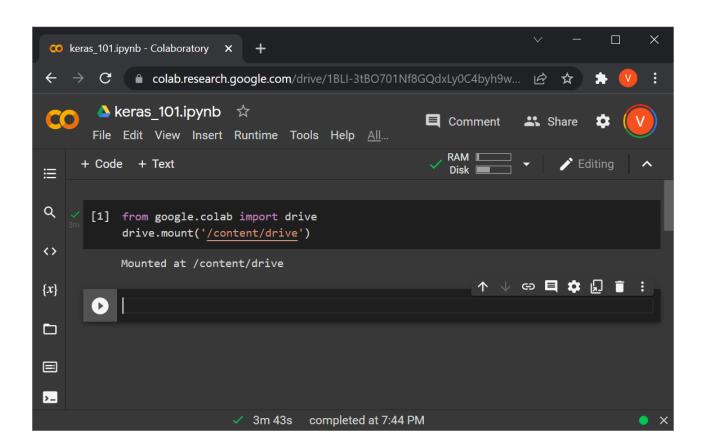
- Put in the code that you see, and run the cell.
- You may get this prompt "Permit this notebook to access your Google Drive files?" If so, click "Connect to Google Drive". You get a couple more prompts to connect to your account and give permissions to Google Drive.



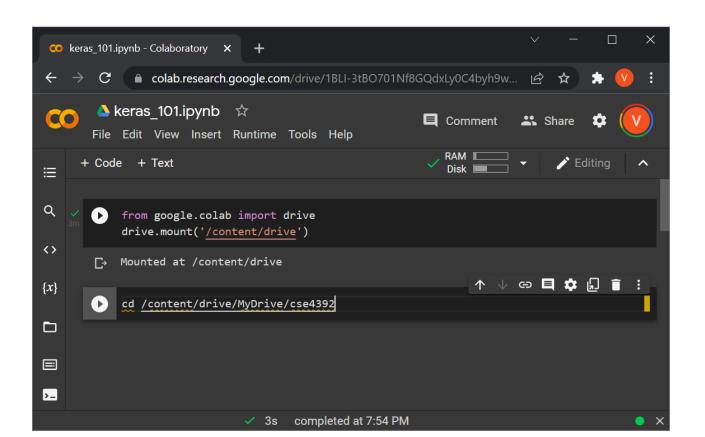
 Once you deal with all the prompts, you get a confirmation that the drive is mounted.



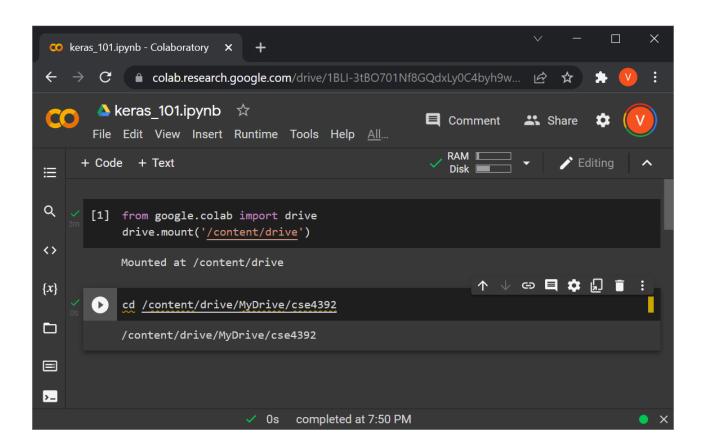
Click on "+ Code" again, to add a new cell.



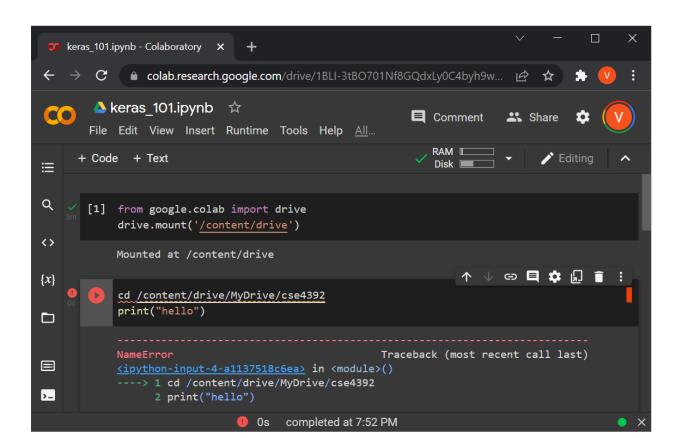
In the new cell, type (or paste) the code that you see.



- Execute the new cell.
- You get a confirmation that the working directory has changed.

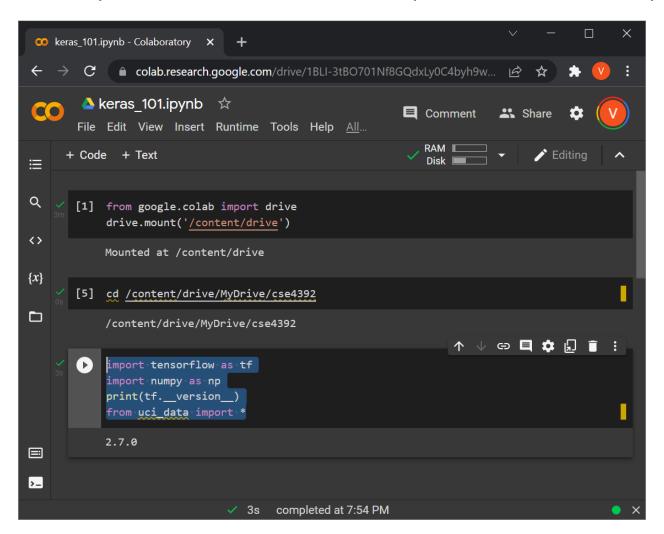


- For some weird reason, the cd command only works (for me) when it is the ONLY line in the cell.
- Here is how it fails otherwise:



Continuing with Rest of Code

Now you can create new cells and put in whatever code you like.



Keras: A First Example

```
import tensorflow as tf
import numpy as np
from uci data import *
(training set, test set) = read uci1("uci datasets", "pendigits")
(training inputs, training labels) = training set
(test inputs, test labels) = test set
input shape = training inputs[0].shape
number of classes = np.max([np.max(training labels), np.max(test labels)]) + 1
model = tf.keras.Sequential([
  tf.keras.Input(shape = input shape),
  tf.keras.layers.Dense(number of classes, activation='sigmoid')
1)
model.compile(optimizer='adam',
        loss=tf.keras.losses.SparseCategoricalCrossentropy(),
        metrics=['accuracy'])
model.fit(training inputs, training labels, epochs=10)
test_loss, test_acc = model.evaluate(test_inputs, test_labels, verbose=0)
print('\nTest accuracy: %.2f%%' % (test acc * 100))
```

- This is a small Keras example.
- It trains and tests a neural network on the pendigits dataset.
- In the next slides we will see what each part does.

Imports

import tensorflow as tf
import numpy as np
from uci_data import *

- uci_data.py is my code, with some auxiliary functions.
 - It is posted on the class website, under this lecture.
 - It has code that reads the UCI dataset files that we have been using.

Loading a Dataset

```
(training_set, test_set) = read_uci1("uci_datasets", "pendigits")
(training_inputs, training_labels) = training_set
(test_inputs, test_labels) = test_set
```

- No tensorflow or Keras code is here yet.
- We are just loading the pendigits dataset, using my read_uci1 helper function, defined at uci_data.py.
- training_inputs is a numpy array
 - training_inputs.shape gives the dimensions of the array, which is (7494, 16)
 - 7494 rows (number of training inputs)
 - 16 columns (number of attributes)

Loading a Dataset

```
(training_set, test_set) = read_uci1("uci_datasets", "pendigits")
(training_inputs, training_labels) = training_set
(test_inputs, test_labels) = test_set
```

- training_labels is a 7494x1 2D array.
 - It could also be a 1D array of 7494 elements, I tested that and it works.
- My code makes sure that class labels have been mapped to consecutive integers starting at 0.
 - The Keras code later assumes that this mapping has been done.

Loading a Dataset

```
(training_set, test_set) = read_uci1("uci_datasets", "pendigits")
(training_inputs, training_labels) = training_set
(test_inputs, test_labels) = test_set
```

- test_inputs is a 3498x16 2D array.
 - 3498 rows, which is the number of test objects.
 - 16 columns, which is the number of attributes.
 - Obviously, the number of attributes in the test inputs should match the number of attributes in the training inputs.
- test_labels is a 3498x1 2D array.

Verifying the Array Sizes

- A (very) common source of bugs in Tensorflow and Keras is wrong array dimensions and sizes. For example:
 - Using an MxN matrix where NxM is expected.
 - Using an Mx1 2D matrix where a 1D matrix of size M is expected.
- It is useful to keep track of what array dimensions, and sizes for those dimensions, are expected.
- When testing and debugging, you can use code like this:

```
print(training_inputs.shape)
print(training_labels.shape)
print(test_inputs.shape)
print(test_labels.shape)
```

Output:

(7494, 16) (7494, 1) (3498, 16)

Parameters for the Network

- Variable input_shape is the dimensionality of the data.
 - It will be the number of units on the input layer.
- Variable number_of_classes is the number of classes.

```
print("input_shape = ", input_shape)
print("number_of_classes = ", number_of_classes)
```

Output:

input_shape = (16,)
number of classes = 10

```
model = tf.keras.Sequential([
    tf.keras.Input(shape = input_shape),
    tf.keras.layers.Dense(number_of_classes, activation='sigmoid')
])
```

- Here is our first piece of Keras code.
- We are creating a two-layer network.
- Function tf.keras.Sequential creates a network that is a sequence of layers.
 - That is the only type of networks we will cover in this class. We will
 discuss several other types of networks in CSE 4392 in the Spring.
- The argument is a list, where every element specifies a layer.

```
model = tf.keras.Sequential([
    tf.keras.Input(shape = input_shape),
    tf.keras.layers.Dense(number_of_classes, activation='sigmoid')
])
```

- tf.keras.Input(shape = input_shape) creates an input layer, and input_shape specifies the number of input units.
- In our case, input_shape will be a list of a single number, because our inputs are simple vectors.
- In cases that we will not cover in this class, input_shape can be a list of 2 or 3 numbers, because each input is a 2D or 3D arrays representing images or other types of data.

```
model = tf.keras.Sequential([
    tf.keras.Input(shape = input_shape),
    tf.keras.layers.Dense(number_of_classes, activation='sigmoid')
])
```

- tf.keras.layers.Dense(number_of_classes, activation='sigmoid') creates a fully connected layer.
 - This is the last layer we specify, so this will be the output layer.
 - We want one output unit for each class, so this is specified by passing number_of_classes as first argument.
 - The activation parameter specifies that we will use the sigmoid activation function
 - There are other options for activation functions, which we will not cover in this class.

- model.compile needs to be called before we train the network.
 - This is one of many Keras-specific "rituals" that we just have to learn and follow.
- The arguments we pass to **model.compile** specify some important hyperparameters of the network.
 - optimizer='adam' specifies one specific variant of gradient descent, that is commonly used.

Loss Function

- loss=tf.keras.losses.SparseCategoricalCrossentropy() specifies that we will use a loss function called "Cross Entropy".
 - The loss parameter specifies the loss function (which we can also call the "error" function).
 - The cross-entropy loss function is an alternative to the sum-of-squared differences that we have seen in the backpropagation slides.

Loss Function

- metrics=['accuracy'] makes no difference in training.
 - It simply specifies that we would like to see the classification accuracy on the training set printed after each training round.
 - In general, it is useful to print out some metrics after each training round, to see how much progress is being made.

Loss Function

model.fit(training_inputs, training_labels, epochs=10)

- model.fit is the function that actually trains the network.
- We specify the training inputs and training labels.
 - Obviously, their dimensions must match the input units and output units that we have already specified.
 - All class labels in training_labels should be non-negative integers that are smaller than the number of output units.
 - Our previous lines of code made sure of that.
- Parameter epochs specifies the number of training rounds.
 - "Epochs" is the term commonly used to specify the number of training rounds in backpropagation.

Output During Training

```
Epoch 1/10 235/235 [==============] - 0s 955us/step - loss: 43.5318 - accuracy: 0.1715

Epoch 2/10 235/235 [=============] - 0s 1ms/step - loss: 15.4048 - accuracy: 0.3979

Epoch 3/10 235/235 [================] - 0s 983us/step - loss: 8.0585 - accuracy: 0.5701
...
```

- For every epoch, we see various statistics. Most important for now:
 - loss shows the value of the loss function after each epoch.
 - accuracy shows the classification accuracy on the training set after each epoch.

Testing the Network

```
test_loss, test_acc = model.evaluate(test_inputs, test_labels, verbose=0) print('\nTest accuracy: %.2f%%' % (test_acc * 100))
```

- model.evaluate applies the network on a test set.
- test_inputs and test_labels specify the test set.
- The verbose parameter specifies how much extra output we want (none in our case).
- Output:

Test accuracy: 96.28%

Some Benchmarks

- I tried different setups for running code that trains (10 epochs) and tests a 2-layer network on the pendigits dataset.
- Running the training and test code (model.fit() and model.evaluate() on Colab took about 5 seconds.
- The same code, running locally on my computer (with Spyder/Anaconda), took less than two seconds.
- My own implementation of backpropagation (which uses numpy, but not Tensorflow or Keras) took about 5 seconds running on my computer.

Some Benchmarks

- Another experiment: a 4-layer network for the pendigits dataset,
 with 50 nodes in each hidden layer.
 - Still training for 10 epochs.
- Running the training and test code (model.fit() and model.evaluate() on Colab took about 11 seconds.
- The same code, running locally on my computer (with Spyder/Anaconda), took about 3 seconds.
- My own implementation of backpropagation (which uses numpy, but not Tensorflow or Keras) took about 8 seconds running on my computer.

GPU vs. CPU Processing

- Note: computers with graphics cards should run the code several times faster than computers without such cards.
 - The computer I used here did NOT have a graphics card.
 - I will try to add some results with a graphics card in the next few days.
- A nice feature of Tensorflow and Keras is that they automatically exploit GPUs, if your system has them.
 - Your code remains the same.

Code for a 4-Layer Network

```
model = tf.keras.Sequential([
    tf.keras.Input(shape = input_shape),
    tf.keras.layers.Dense(50, activation='sigmoid'),
    tf.keras.layers.Dense(50, activation='sigmoid'),
    tf.keras.layers.Dense(number_of_classes, activation='sigmoid')
])
```

- To create the 4-layer network used in the previous timing experiments, just call tf.keras.Sequential() as shown above.
- It is very straightforward.
 - Each hidden layer has 50 units, is fully connected, and uses the sigmoid activation function.

Evaluating the Network on an Input

- Suppose that we have already trained a 2-layer model on the pendigits dataset, as shown in the previous slides.
- Here we see how to (and how not to) apply the model to a new input vector.
- To do that, we use the predict() method.
- Here is an example of how NOT to do it. We get an error.

```
test_index = 10 # nothing special about this value
input_vector = test_inputs[test_index,:]
nn_output = model.predict(input_vector)
```

 We would hope that nn_output contains the output of the model (which is a vector of as many dimensions as the number of classes).

Evaluating the Network on an Input

```
test_index = 10 # nothing special about this value
input_vector = test_inputs[test_index,:]
nn_output = model.predict(input_vector)
```

- The code above (same as in the previous slide) gives an error.
 - This is a good example of often frustrating errors we get because of incompatible array shapes.
- Problem: input_vector is a 1D vector. The predict() method wants its argument to be a 2D array where each row is an input.
- Fix:

```
test_index = 10 # nothing special about this value
input_vector = test_inputs[test_index,:]
input_vector = np.reshape(input_vector, (1, 16))
nn_output = model.predict(input_vector)
```

Evaluating the Network on an Input

• Fix:

```
test_index = 10 # nothing special about this value
input_vector = test_inputs[test_index,:]
input_vector = np.reshape(input_vector, (1, 16))
nn_output = model.predict(input_vector)
```

- The **numpy.reshape()** method is used to convert input vector from a 1D array of 16 elements to a 1x16 2D array.
- Now model.predict() is happy.
- nn_output is a 1x10 2D array.
- If we want to convert nn_output to a 1D array we do:

```
nn_output = nn_output.flatten()
```

From Output to Class Prediction

- As we know, to convert the output to a class prediction, we need to find the argmax.
 - In other words, we find the output unit that gave the highest response.

predicted class label = 0 actual class label = 7

Classification Ties

- It can happen that two or more output units tie for the highest output value.
- Your code should be able to detect that.
- It so happens that our 2-layer network produces lots of ties.
 - In a few slides we will see what causes that and how to fix it.
 - For test_index = 10, six units tie for the highest value.

```
(indices,) = np.nonzero(nn_output == nn_output[predicted_class])
print("indices =", indices)
```

Output:

indices = [0 3 4 5 6 7]

Classification Accuracy and Ties

- In that example, how accurate was the classification?
- We cannot say 100% accurate, because we did not get a single correct answer.
- We cannot say 100% wrong, because the correct class label,
 "7", was one of six class labels that tied for highest value.
- In such cases, it is fair to say that the result was 16.67% accurate.
 - This is 100% divided by the number of classes that tied.
- Obviously, if there are ties but the correct class label is not included in those ties, then the result is 0% accurate.
 - Which means, it is 100% wrong.

Code for Measuring Accuracy

```
predicted_class = np.argmax(nn_output)
actual class = test labels[test index]
print("predicted: %d\nactual: %d\n" % (predicted_class, actual_class))
(indices,) = np.nonzero(nn output == nn output[pred])
print("indices =", indices)
number_of_ties = np.prod(indices.shape)
if (nn_output[actual_class] == nn_output[predicted_class]):
  accuracy = 1.0 / number of ties
else:
                                           Output:
  accuracy = 0
                                           indices = [0 3 4 5 6 7]
                                           accuracy = 0.1667
print("accuracy = %.4f" % (accuracy))
```

Normalizing Input Values

- The reason for that many ties is numerical.
- If we print nn_output for our example, we see that six values are numerically equal to 1.
- This can happen, if the input to the sigmoid function is too high.
- To avoid such numerical issues, it is good to normalize the input values.
- Without such normalization, input values can be arbitrarily high or low.
- The neural networks assignment specifies that you should divide all input vectors by the "MAXIMUM ABSOLUTE value over all attributes over all training objects".
 - Then, all attribute values are between -1 and 1.

Normalizing Input Values

 This code does the required normalization, feel free to use in your assignment:

```
(training_set, test_set) = read_uci1("uci_datasets", "pendigits")
(training_inputs, training_labels) = training_set
(test_inputs, test_labels) = test_set
max_value = np.max(training_inputs)
training_inputs = training_inputs / max_value
test_inputs = test_inputs/ max_value
```

 Normalizing like that before training the model, my code found no classification ties in any of the 3498 test objects of the pendigit dataset.

Recap

- We have seen how to use Keras to train and test multi-layer networks.
- With Keras, we do not have to write code for backpropagation.
 - We just need to specify the network design.
 - Keras figures out on its own how to do the backpropagation in a way that is correct for our specific design.