You must implement a Matlab function or a Python executable file called neural\_network, that uses backpropagation to train a neural network, and then applies that neural network to classify test data. Your function should be invoked as follows:

neural\_network(<training\_file>, <test\_file>, <layers>, <units\_per\_layer>, <rounds>)

If you use Python, just convert the Matlab function arguments shown above to command-line arguments. The arguments provide to the function the following information:

* The first argument, <training\_file>, is the path name of the training file, where the training data is stored. The path name can specify any file stored on the local computer.
* The second argument, <test\_file>, is the path name of the test file, where the test data is stored. The path name can specify any file stored on the local computer.
* The third argument, <layers>, specifies how many layers to use. Note that the input layer is layer 1, so the number of layers cannot be smaller than 2 (any neural network should have at least an input layer and an output layer).
* The fourth argument,<units\_per\_layer>, specifies how many perceptrons to place at each HIDDEN layer. Note that this number is not applicable either to the input layer or to the output layer.
* The fifth argument, <rounds>, is the number of training rounds that you should use. Each training round consists of using the whole training set once (i.e., using each training example once to update the weights).

The training and test files will follow the same format as the text files in the [UCI datasets](https://athitsos.utasites.cloud/courses/cse4309_fall2021/assignments/uci_datasets) directory. A description of the datasets and the file format can be found [on this link](https://athitsos.utasites.cloud/courses/cse4309_fall2021/assignments/uci_datasets/dataset_description.html). For each dataset, a training file and a test file are provided. The name of each file indicates what dataset the file belongs to, and whether the file contains training or test data. Your code should also work with ANY OTHER training and test files using the same format as the files in the [UCI datasets](https://athitsos.utasites.cloud/courses/cse4309_fall2021/assignments/uci_datasets) directory.

As the [description](https://athitsos.utasites.cloud/courses/cse4309_fall2021/assignments/uci_datasets/dataset_description.html) states, **do NOT use data from the last column (i.e., the class labels) as features**. In these files, all columns except for the last one contain example inputs. The last column contains the class label.

**Training**

In your implementation, you should use these guidelines:

* For each dataset, for all training and test objects in that dataset, you should normalize all attribute values, by dividing them with the MAXIMUM ABSOLUTE value over all attributes over all training objects for that dataset. This is a single MAXIMUM value, you should NOT use a different maximum value for each dimension. In other words, for each dataset, you need to find the single highest absolute value across all dimensions and all training objects. Every value in every dimension of every training and test object should be divided by that highest value.
* The weights of each unit in the network should be initialized to random values, between -0.05 and 0.05. The probability of each random value should follow a uniform distribution between -0.05 and 0.05.
* You should initialize your learning rate to 1 for the first training round, and then multiply it by 0.98 for each subsequent training round. So, the learning\_rate used for training round r should be 0.98r - 1.
* Your stopping criterion should simply be the number of training rounds, which is specified as the fifth argument. The number of training rounds specifies how many times you iterate over the entire training set. A single training round consists of a single execution of steps 3, 4, 5, 6 in the backpropagation summary, on slide 45 of the [neural networks slides](https://athitsos.utasites.cloud/courses/cse4309_fall2021/lectures/09c_neural_networks.pdf). If you want your code to correspond step-by-step to that summary, then in step 5 of that summary you should use a threshold equal to -1, so that the error is never used as the stopping criterion.

The number of layers in the neural network is specified by the <layers> argument. If we have L layers:

* Layer 1 is the input layer, that contains no perceptrons, it just specifies the inputs to the neural network. Thus, 2 is the minimum legal value for L.
* Each perceptron at layer 2 has D inputs, where D is the number of attributes. The attributes of the input object provide these D input values to each perceptron at layer 2.
* Layer L is the output layer, containing as many perceptrons as the number of classes.
* If L > 2, then layers 2, ..., L-1 are the hidden layers. Each of these layers has as many perceptrons as specified in <units\_per\_layer>, the third argument. If L = 2, then the fourth argument (<units\_per\_layer>) is ignored.
* If L > 2, each perceptron at layers 3, ..., L has as inputs the outputs of ALL perceptrons at the previous layer.
* Note that each dataset contains more than two classes, so your output layer needs to contain a number of units (perceptrons) equal to the number of classes, as discussed in the slides.

There is no need to output anything for the training phase.

**Classification**

For each test object you should print a line containing the following info:

* Object ID. This is the line number where that object occurs in the test file. Start with 1 in numbering the objects, not with 0.
* Predicted class (the result of the classification). If your classification result is a tie among two or more classes, choose one of them randomly.
* True class (from the last column of the test file).
* Accuracy. This is defined as follows:
  + If there were no ties in your classification result, and the predicted class is correct, the accuracy is 1.
  + If there were no ties in your classification result, and the predicted class is incorrect, the accuracy is 0.
  + If there were ties in your classification result, and the correct class was one of the classes that tied for best, the accuracy is 1 divided by the number of classes that tied for best.
  + If there were ties in your classification result, and the correct class was NOT one of the classes that tied for best, the accuracy is 0.

To produce this output in a uniform manner, use:

fprintf('ID=%5d, predicted=%3d, true=%3d, accuracy=%4.2f\n',

object\_id, predicted\_class, true\_class, accuracy);

After you have printed the results for all test objects, you should print the overall classification accuracy, which is defined as the average of the classification accuracies you printed out for each test object. To print the classification accuracy in a uniform manner, use:

fprintf('classification accuracy=%6.4f\n', classification\_accuracy);