```
In [179]: import numpy as np
          import pandas as pd
          from sklearn.model selection import KFold
          def getData(filePath):
               data = np.genfromtxt(filePath, delimiter=',')
              x, y = np.array(data[:,0:-1], dtype=float), np.array(data[:,-1],dtype=int)
              y = y.reshape(1, len(y)).T
              return x,y
          def splitInputOutput(data):
              x, y = np.array(data[:,0:-1], dtype=float), np.array(data[:,-1],dtype=int)
              y = y.reshape(1, len(y)).T
              return x,y
          def sigmoid(x):
              return 1/(1+np.exp(-x))
          def make sigmoid prime(x):
               return x*(1-x)
          def trainNeuralNet(synapse0, synapse1, epochs, activator, activator prime):
              for j in range(epochs):
                  11 = activator(np.dot(X,synapse0))
                  12 = activator(np.dot(l1,synapse1))
                  12 delta = (y - 12)*activator prime(np.dot(l1,synapse1))
                  11_delta = 12_delta.dot(synapse1.T) * activator_prime(np.dot(X,synapse0))
                  synapse1 += l1.T.dot(12 delta) #adjust our synapses up or down as necessary
                  synapse0 += X.T.dot(l1 delta)
          def transformTestData(x, syn0, syn1, func):
              layer1 transform = func(np.dot(x,syn0))
              return func(np.dot(layer1 transform,syn1))
```

Test our neural network trainer against a simple dataset: X will contain binary tuples and Y will be the XOR result of rows in X.

```
In [185]: # trivial dataset
X, y = getData('data/prepared/trivial.csv')

# X = np.array([[0,0],[0,1],[1,0],[1,1]])
# y = np.array([[0,1,1,0]]).T # XOR(X)

np.random.seed(seed=42)
syn0 = 2*np.random.random((X.shape[1],X.shape[0])) - 1
syn1 = 2*np.random.random((y.shape[0],y.shape[1])) - 1
epochs = 10000

trainNeuralNet(syn0, syn1, epochs, lambda x: sigmoid(x), lambda x: sigmoid(x)*(1-sigmoid(x)))

result = transformTestData(X,syn0,syn1,lambda x: sigmoid(x))
# layer1_transform = sigmoid(np.dot(X,syn0))
# result = sigmoid(np.dot(layer1_transform,syn1))

print("MSE: ",0.5*np.sum((y - result)**2))
print("Output of predicted y (2nd and 3rd rows should be close to 1):")
print(result)
```

```
MSE: 0.000477306752279

Output of predicted y (2nd and 3rd rows should be close to 1):

[[ 0.01869055]
        [ 0.98941286]
        [ 0.98293159]
        [ 0.0142077 ]]
```

Looks good. Now lets load our accute inflamation dataset. Our dataset was randomized. It contained 120 rows and we split 80/20 for train&validation(96) / test(24). We'll use scikit learn's KFold utility class to get our indices for a 5 k-fold cross validation and pick the best model to run our test data against.

```
In [197]: # X,y = qetData('data/prepared/dataWithTemp.csv')
          df = pd.read csv('data/prepared/dataWithTempRandomized.train.csv',sep=',',names=["Temp", "Nausea", "Lumbar", "Pushing","Micturi
          df["Temp"] = df.transform(lambda x: x - 37)
          X,y = splitInputOutput(df.as matrix())
          kf = KFold(n splits=5,random state=None, shuffle=True)
          lowest mse = 1e8 #arbitrary high value
          lowest syn0 = []
          lowest syn1 = []
          epochs = 10000
          print("Performing K-fold cross validation, splits = 5")
          # do our cross validation with training data
          for train index, test index in kf.split(X):
              X train, X test = X[train index], X[test index]
              y train, y test = y[train index], y[test index]
              np.random.seed(seed=42)
              syn0 = 2*np.random.random((X_train.shape[1],X_train.shape[0])) - 1
              syn1 = 2*np.random.random((y_train.shape[0],y_train.shape[1])) - 1
              trainNeuralNet(syn0, syn1, epochs, lambda x: sigmoid(x), lambda x: sigmoid(x)*(1-sigmoid(x)))
              layer1 transform = sigmoid(np.dot(X test,syn0))
              result = sigmoid(np.dot(layer1 transform,syn1))
              mse = 0.5*np.sum((y test - result)**2)
              print("
                         MSE: ",mse)
              if (mse < lowest_mse):</pre>
                  lowest mse = mse
                  lowest_syn0 = syn0
                  lowest syn1 = syn1
          df = pd.read csv('data/prepared/dataWithTempRandomized.test.csv',sep=',',names=["Temp", "Nausea", "Lumbar", "Pushing","Micturit
          df["Temp"] = df.transform(lambda x: x - 37)
          X,y = splitInputOutput(df.as matrix())
          layer1 transform = sigmoid(np.dot(X,lowest syn0))
          result = sigmoid(np.dot(layer1 transform,lowest syn1))
          mse = 0.5*np.sum((y - result)**2)
          result = np.double(result > 0.5)
          print()
          print("MSE against Test data: ",mse)
          print("Accuracy: ",1-np.sum(y-result)/y.shape[0])
```

Performing K-fold cross validation, splits = 5

MSE: 4.49999997067 MSE: 2.99999982413 MSE: 0.99999969645 MSE: 0.99999960494 MSE: 1.50000074508

MSE against Test data: 1.9999999116

Accuracy: 0.833333333333

In []: