In [1]:

```
# Import required packages here (after they are installed)
import numpy as np
import matplotlib.pyplot as mp
from pylab import show

# Load data. csv file should be in the same folder as the notebo ok for this to work, otherwise
# give data path.

data = np.loadtxt("transfusion.csv", dtype = float, delimiter=', ', skiprows=1)
```

In [2]:

```
features = []
donated = []

for row in data:
    features.append(row[:4])
    donated.append(str(row[4]))

# print( features[0])
# print( donated[0])
```

```
In [3]:
```

```
from sklearn.neural network import MLPClassifier
from sklearn.model_selection import cross val score
scores = [[],[],[]]
i = 0
for hidden layers in [1, 2, 5]:
    for num nodes in [2, 5, 10]:
        the tuple = [num_nodes] * hidden_layers
#
          the tuple = []
          for x in range (hidden layers):
#
              the tuple.append( num nodes)
#
        the tuple = tuple(the tuple)
        print(the tuple)
        model = MLPClassifier(hidden layer sizes=(the_tuple),act
ivation='relu',epsilon=0.001,max iter=10000,alpha=0,solver='adam
')
        scores[ i].append( 1 - np.average( cross val score( mode
1, features, donated, cv=10)))
    i += 1
(2,)
(5,)
(10,)
```

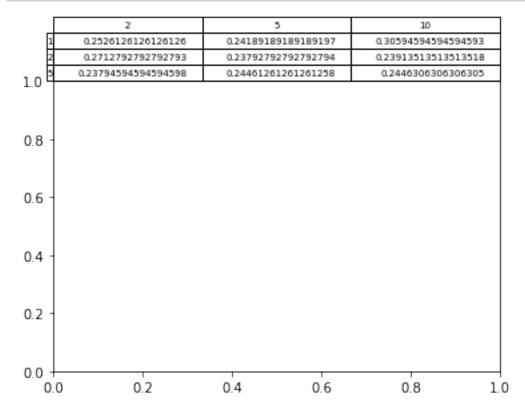
```
(2, 2)
(5, 5)
(10, 10)
(2, 2, 2, 2, 2)
(5, 5, 5, 5, 5)
(10, 10, 10, 10, 10)
```

In [13]:

```
print ( scores)
print("Highest accuracy found with 2 hidden layers and 5 nodes p
er layer")
```

```
[[0.2526126126126126, 0.24189189189189197, 0.3059459 4594594593], [0.2712792792792793, 0.23792792792794, 0.23913513513513518], [0.23794594594594598, 0.244 61261261261258, 0.2446306306306305]] Highest accuracy found with 2 hidden layers and 5 no des per layer
```

In [5]:



In [6]:

```
hwldata = np.loadtxt("data.csv")
#shuffle the data and select training and test data
```

```
np.random.seed(100)
np.random.shuffle(data)
featuresHw1 = []
digits = []
for row in hwldata:
#import the data and select only the 1's and 5's
    if(row[0]==1 \text{ or } row[0]==5):
        featuresHw1.append(row[1:])
        digits.append(str(row[0]))
#Select the proportion of data to use for training. #Notice that
we have set aside 80% of the data for testing
numTrain = int(len(features)*.2)
trainFeatures = features[:numTrain]
testFeatures = features[numTrain:]
trainDigits = digits[:numTrain]
testDigits = digits[numTrain:]
#Convert the 256D data (trainFeatures) to 2D data
#We need X and Y for plotting and simpleTrain for building the m
odel.
#They contain the same points in a different arrangement
X = []
Y = []
simpleTrain = []
#Colors will be passed to the graphing library to color the poin
ts.
#1's are blue: "b" and 5's are red: "r"
colors = []
for index in range(len(trainFeatures)):
    #produce the 2D dataset for graphing/training and scale the
data so it is in the [-1,1] square
    xNew = 2*np.average(trainFeatures[index])+.75
    yNew = 3*np.var(trainFeatures[index])-1.5
    X.append(xNew)
    Y.append(yNew)
    simpleTrain.append([xNew,yNew])
```

In [7]:

```
import time
scoresHw1Features = [[],[],[],[]]
runtimes = [[],[],[],[]]
i = 0
for hidden_layers in [1, 2, 5, 10]:
    for num nodes in [2, 5, 10, 50, 100]:
        the tuple = [num nodes] * hidden layers
        the tuple = tuple(the tuple)
        print(the tuple)
        start time = time.time();
        model = MLPClassifier(hidden layer sizes=(the tuple), ac
tivation='relu', epsilon=0.001, max iter=10000, alpha=0, solver=
"adam")
        scoresHw1Features[ i].append( 1 - np.average( cross val
score( model, trainFeatures, trainDigits, cv=10)))
        end time = time.time();
        total time = end time - start time
        runtimes[ i].append( total time)
    i += 1
for i in range( len( runtimes)):
    for j in range( len( runtimes[0])):
        runtimes[ i][ j] = runtimes[ i][ j] * 1000 # convert sec
onds to milliseconds
```

```
(2,)
(5,)
(10,)
(50,)
(100,)
(2, 2)
(5, 5)
(10, 10)
(50, 50)
(100, 100)
(2, 2, 2, 2, 2)
(5, 5, 5, 5, 5)
(10, 10, 10, 10, 10)
(50, 50, 50, 50, 50)
(100, 100, 100, 100, 100)
(2, 2, 2, 2, 2, 2, 2, 2, 2, 2)
(5, 5, 5, 5, 5, 5, 5, 5, 5, 5)
(10, 10, 10, 10, 10, 10, 10, 10, 10, 10)
(50, 50, 50, 50, 50, 50, 50, 50, 50, 50)
```

```
In [8]:
```

```
print( "CV-Errors")
for i in range( len( scoresHw1Features)):
    print ( scoresHw1Features[i])
print("Runtimes")
for i in range( len( runtimes)):
    print ( runtimes[i])
CV-Errors
```

```
[0.3080952380952382, 0.38142857142857145, 0.34142857
14285714, 0.2880952380952382, 0.30333333333333345
[0.2814285714285715, 0.41285714285714303, 0.31000000
000000005, 0.429999999999994, 0.34809523809523811
[0.26142857142857145, 0.2947619047619048, 0.31666666
66666665, 0.3280952380952382, 0.3480952380952381]
[0.26142857142857145, 0.26142857142857145, 0.2614285]
7142857145, 0.26142857142857145, 0.30190476190476191
Runtimes
[1813.6482238769531, 927.0899295806885, 947.90625572
20459, 175.66609382629395, 312.875986099243161
[2732.5289249420166, 1495.4001903533936, 462.2550010
6811523, 323.5502243041992, 366.77026748657227
[3323.112964630127, 3598.2279777526855, 1844.0201282
50122, 1237.2093200683594, 1158.6182117462158]
[5783.318042755127, 3155.7679176330566, 2534.7840785
980225, 3986.4871501922607, 4590.992689132691
```

In [14]:

```
figure = mp.figure(dpi=100)
ax = figure.add subplot(2, 2, 2)
the table = ax.table(cellText=scoresHw1Features,
                      rowLabels=[1,2,5,10],
                      colLabels=[2, 5, 10, 50, 100],
                    loc='top')
the table.scale(1,4)
the_table.set_fontsize(14)
ax.axis('off')
mp.show()
figure = mp.figure(dpi=100)
ax = figure.add subplot(2, 2, 2)
the table = ax.table(cellText=runtimes,
                      rowLabels=[1,2,5,10],
                      colLabels=[2, 5, 10, 50, 100],
                    loc='top')
the table.scale(1,4)
the table.set fontsize(14)
ax.axis('off')
mp.show()
```

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BILLIMAGOLY	\$61277773666	BRESHERS	SETZONOMINEMA	DIVERSITY
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- a) The runtime seems to increase as the number of layers increase, but decrease as the number of nodes per layer increases. So, they seem to have the opposite effect.
- b) The optimum result seems to be found at 10 layers and 50 nodes per layer.

^{*}Note my runtimes vary drastically with each run

In [10]:

```
the_tuple = tuple([50] * 10)

optimalscores = []
optimalruntime = []

for learnrate in [0.00001, 0.0001, 0.001, 0.01, 0.1]:
    start_time = time.time();
    model = MLPClassifier(learning_rate_init=learnrate, hidden_l
ayer_sizes=(the_tuple), activation='relu', epsilon=0.001, max_it
er=10000, alpha=0, solver="adam")
    optimalscores.append( 1 - np.average( cross_val_score( model
, trainFeatures, trainDigits, cv=10)))
    optimalruntime.append( time.time() - start_time)

print( optimalscores)
print( optimalruntime)
```

```
[0.35476190476190483, 0.2947619047619048, 0.26142857 142857145, 0.25476190476190474, 0.26142857142857145] [11.959640741348267, 4.220972061157227, 5.3577220439 91089, 3.5515692234039307, 1.4335618019104004]
```

- c) Decreasing the learning rate from the default seems to increase the error but increasing it from the default does not seem to affect the accuracy. Runtime seems to decrease as learning rate is increased.
- d) The neural networks is returning different accuracy values with each run, which is likely due to it always starting with random weights. However, the optimum model seems to be the same one almost every time, if not every. It has a slight impact on the expected fit, which might be very slightly more fitted, but almost about the same.

In [11]:

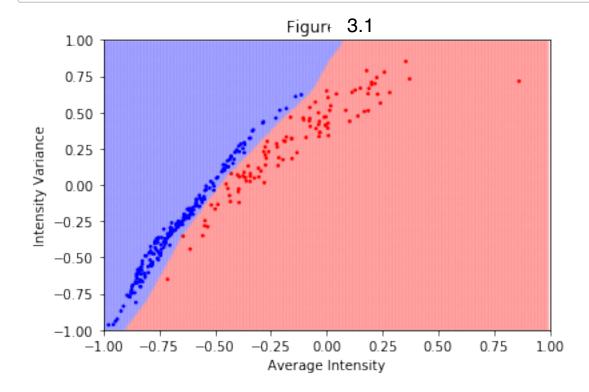
```
data = np.loadtxt("data.csv")

#shuffle the data and select training and test data
np.random.seed(100)
np.random.shuffle(data)
```

```
reatures = []
digits = []
for row in data:
    #import the data and select only the 1's and 5's
    if(row[0]==1 or row[0]==5):
        features.append(row[1:])
        digits.append(str(row[0]))
#Select the proportion of data to use for training.
#Notice that we have set aside 80% of the data for testing
numTrain = int(len(features)*.2)
trainFeatures = features[:numTrain]
testFeatures = features[numTrain:]
trainDigits = digits[:numTrain]
testDigits = digits[numTrain:]
#Convert the 256D data (trainFeatures) to 2D data
#We need X and Y for plotting and simpleTrain for building the m
odel.
#They contain the same points in a different arrangement
X = []
Y = []
simpleTrain = []
#Colors will be passed to the graphing library to color the poin
ts.
#1's are blue: "b" and 5's are red: "r"
colors = []
for index in range(len(trainFeatures)):
    #produce the 2D dataset for graphing/training and scale the
data so it is in the [-1,1] square
    xNew = 2*np.average(trainFeatures[index])+.75
    yNew = 3*np.var(trainFeatures[index])-1.5
    X.append(xNew)
    Y.append(yNew)
    simpleTrain.append([xNew,yNew])
    #trainDigits will still be the value we try to classify. Her
e it is the string "1.0" or "5.0"
    if(trainDigits[index]=="1.0"):
        colors.append("b")
```

```
else:
        colors.append("r")
# create the model
#
# Declare Model
model = MLPClassifier( hidden layer sizes=(the tuple), activatio
n='relu', epsilon=0.001, max iter=10000, alpha=0, solver="adam")
# Fit model to our data
model.fit(simpleTrain,trainDigits)
# Lists to hold inpoints, predictions and assigned colors
xPred = []
yPred = []
cPred = []
# Use input points to get predictions here
for xP in range(-100,100):
    xP = xP/100.0
    for yP in range(-100,100):
        yP = yP/100.0
        xPred.append(xP)
        yPred.append(yP)
        if(model.predict([[xP,yP]])=="1.0"):
            cPred.append("b")
        else:
            cPred.append("r")
## Visualize Results
#plot the points
mp.scatter(X,Y,s=3,c=colors)
#plot the regions
mp.scatter(xPred,yPred,s=3,c=cPred,alpha=.1)
#setup the axes
mp.xlim(-1,1)
mp.xlabel("Average Intensity")
mp.ylim(-1,1)
mp.ylabel("Intensity Variance")
#label the figure
mp.title("Figure 3.1")
```

show()



In []: