```
In [277]: import numpy as np
    import matplotlib.pyplot as plt
    from torch import nn
    import torch
    from torchvision.datasets import MNIST
    import torchvision.transforms as transforms
    from sklearn.metrics import accuracy_score
    %matplotlib inline
```

For this homework you will be using pytorch and torchvision library for neural networks and datasets. You can install them with pip install torch torchvision.

Question 1 Principal Component Analysis

This problem will guide you through the principal component analysis. You will be using a classical dataset, the MNIST hand written digit dataset.

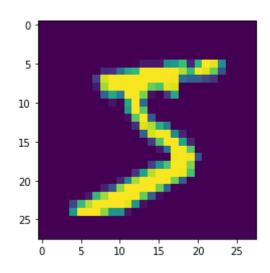
```
In [278]: # Load the MNIST dataset
    mnist = MNIST('.', download=True)
    data = mnist.train_data.numpy()
    labels = mnist.train_labels.numpy()
    print('shapes:', data.shape, labels.shape)
    plt.imshow(data[0])
    print('label:', labels[0])
```

/Users/joshuakang/opt/anaconda3/lib/python3.7/site-packages/torchvisio n/datasets/mnist.py:53: UserWarning: train_data has been renamed data warnings.warn("train_data has been renamed data") /Users/joshuakang/opt/anaconda3/lib/python3.7/site-packages/torchvisio

/Users/joshuakang/opt/anaconda3/lib/python3.7/site-packages/torchvisio n/datasets/mnist.py:43: UserWarning: train_labels has been renamed targ ets

warnings.warn("train_labels has been renamed targets")

```
shapes: (60000, 28, 28) (60000,)
label: 5
```



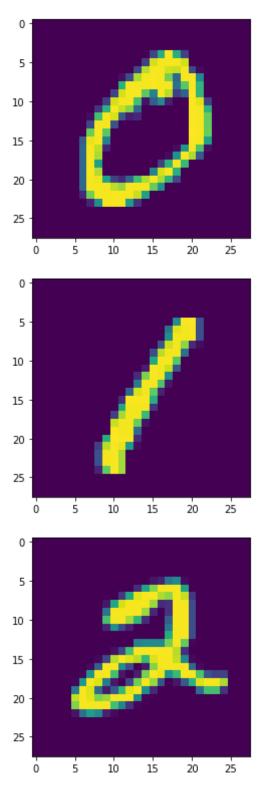
Question 1.1 Familiarize yourself with the data [5pt]

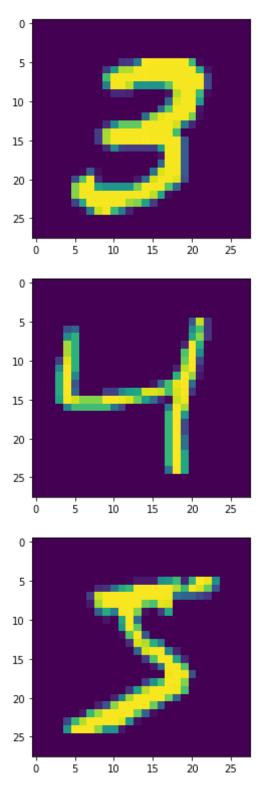
For this task, you will be using the torchvision package that provides the MNIST dataset. For each digit class(0-9), plot 1 image from the class and store those 10 images for each digit class in the array digit images.

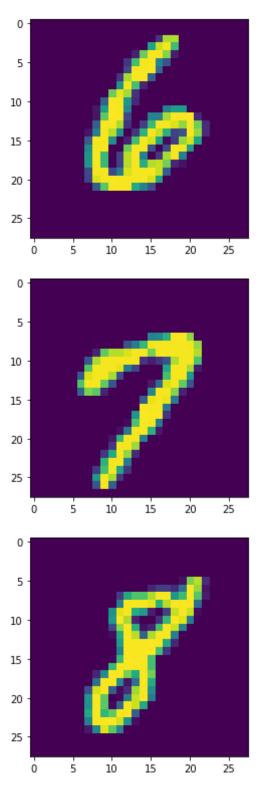
```
In [280]: digit_images = np.zeros([10, 28, 28])
    import random

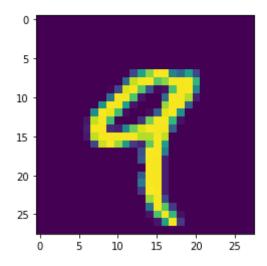
test_images = []

for i in range(10):
    index = np.where(labels == i)[0][0]
    test_images.append(index)
    digit_images[i] = data[index]
    plt.figure()
    plt.imshow(data[index])
### YOUR CODE HERE
### END OF CODE
```









Question 1.2 PCA

The following questions will guide you through the PCA algorithm.

Question 1.2.1 Centering the data [5pt]

For each image, flatten it to a 1-D vector. To perform PCA on the dataset, we first move the data points so they have 0 mean on each dimension. Store the centered data in variable <code>data_centered</code> and the mean of each dimension in variable <code>data_mean</code>.

```
In [295]: data_centered = np.zeros((60000, 784))
    data_mean = None
    flat_data = []
    for datapoint in data:
        flat_data.append(datapoint.flatten())
    flat_data = np.array(flat_data)

    data_mean = np.mean(flat_data,axis = 0)

for i in range(flat_data.shape[0]):
        data_centered[i] = np.array(flat_data[i] - data_mean)

### YOUR CODE HERE
### END OF CODE
```

Question 1.2.2 Compute the covariance matrix of the data [5pt]

You need to store the covariance matrix of the data in variable data_covmat . You may **not** use numpy.cov

```
In [296]: data_covmat = (data_centered.T@data_centered)/data_centered.shape[1]
# data_covmat = np.cov(data_centered)
### YOUR CODE HERE
### END OF CODE
print(data_covmat.shape)
(784, 784)
```

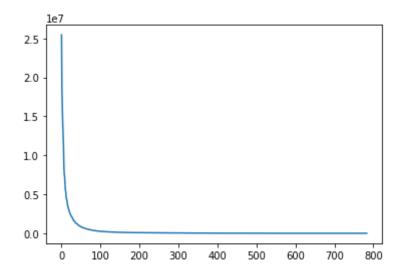
Question 1.2.3 Compute the eigenvalues of the covariance matrix [5pt]

You need to store the eigenvalues of the covariance matrix in variable <code>covmat_eig</code>, sorted in descending order. Then you need to plot the eigenvalues with <code>plt.plot</code>. You can use any numpy function.

```
In [307]: eig_val,covmat_eig = np.linalg.eig(data_covmat)
    eig_loc = np.argsort(eig_val)[::-1]
    new_basis = covmat_eig[:,eig_loc[:2]]
    x = []
    for i in range(784):
        x.append(i)
    plt.plot(x, eig_val)
    print(new_basis.shape)

### YOUR CODE HERE
### END OF CODE
```

(784, 2)



Question 1.2.4 Project data onto the first 2 principal components [5pt]

Now you need to project the centered data on the 2D space formed by the eigenvectors corresponding to the 2 largest eigenvalues. Create a 2D scatter plot where you need to assign a unique color to each digit class.

```
In [308]: ### YOUR CODE HERE
    Z = data_centered @ new_basis

print(Z.shape)

colors = ["red", "orange", "yellow", "green", "blue", "indigo", "violet"
    , "black", "brown", "pink"]

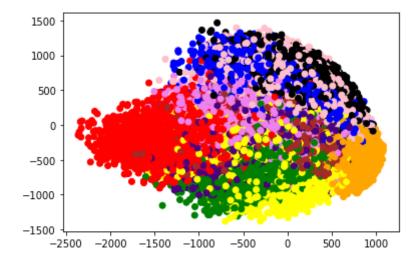
fig = plt.figure()
    ax = fig.add_subplot(111)

projected_x = [x for x in Z[:,0]]
    projected_y = [y for y in Z[:,1]]
    color_map = [colors[label] for label in labels]
    plt.scatter(projected_x, projected_y, c=color_map)

### END OF CODE

(60000, 2)
```

Out[308]: <matplotlib.collections.PathCollection at 0x1a2d4be2d0>



Question 1.2.5 Unproject data back to high dimensions [10pt]

For this question, you need to project the 10 images you plotted in **1.1** on the first 2 principal components, and then unproject the "compressed" 2-D representations back to the original space. Plot the "compressed" digit (the reconstructed digit). Do they look similar to the original images?

No they do no look like the image when we are using only the two largest principa components. We need to use more PCs.

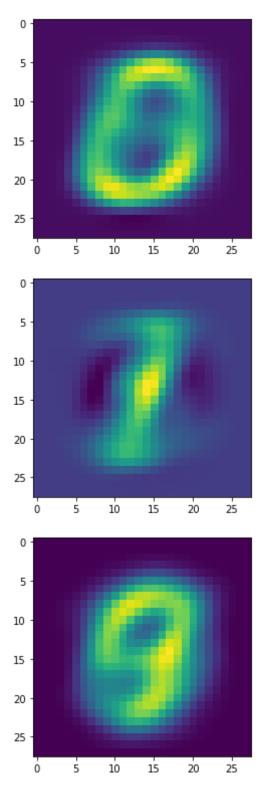
```
In [309]: ### YOUR CODE HERE
X_prime = Z @ new_basis.T
reconstructed_data = []

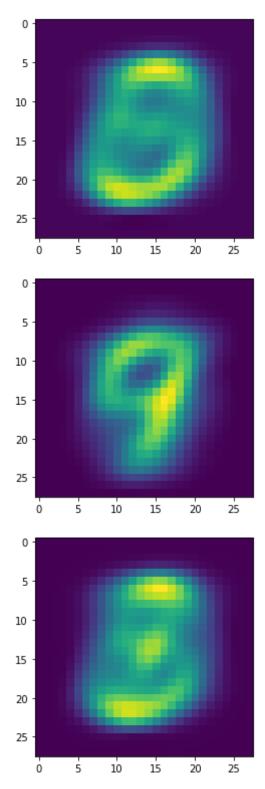
for image in test_images:
    plt.figure()
    current_test_image = data[image].flatten()
    #tranform the test images
    reduced_test_image = (current_test_image - data_mean) @ new_basis

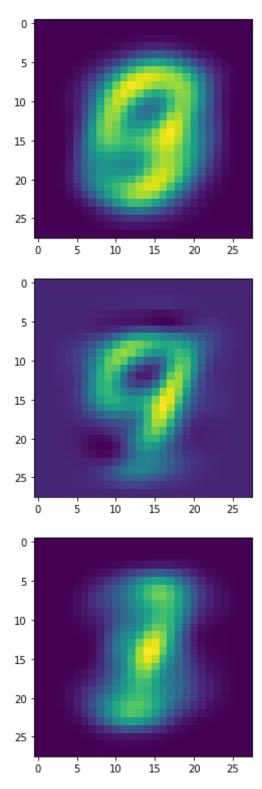
#transform them back
    test_image_prime = (reduced_test_image @ new_basis.T) + data_mean
    test_image_prime = np.array(test_image_prime)
    reconstructed_data.append(test_image_prime)

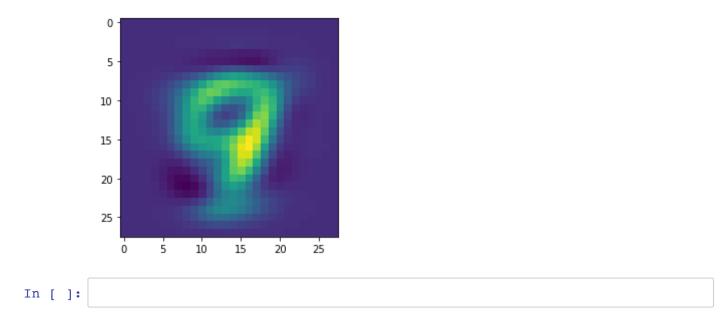
image_shaped = np.reshape(test_image_prime, (28,28))
    plt.imshow(image_shaped)

### END OF CODE
```





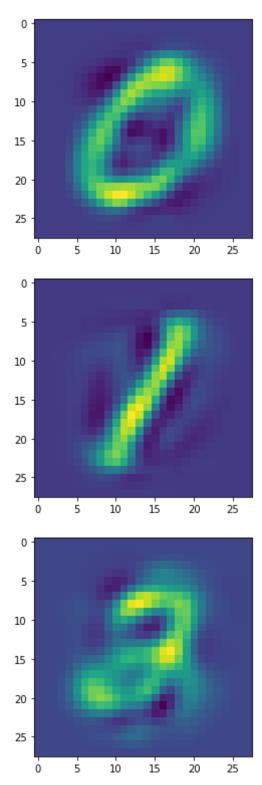


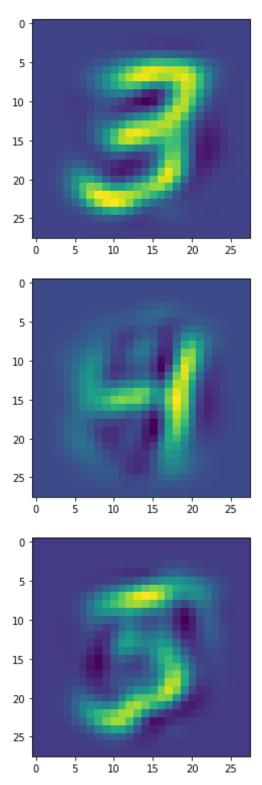


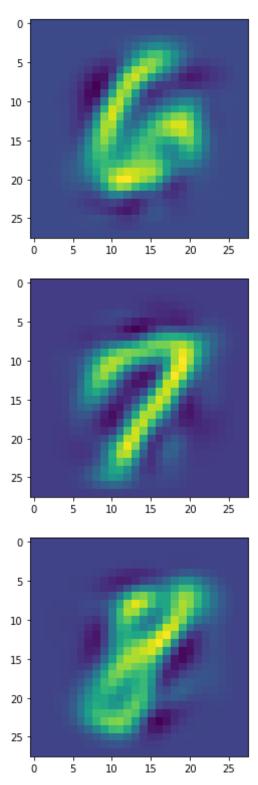
Question 1.2.6 Choose a better low dimension space. [5pt]

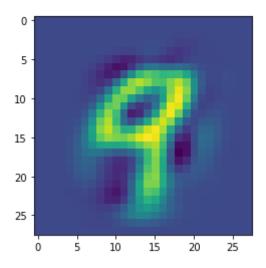
Do the previous problem with more dimensions (e.g. 3, 5, 10, 20, 50, 100). You only need to show results for one of them. Answer the following questinos. How many dimensions are required to represent the digits reasonably well? How are your results related to **question 1.2.3**?

```
In [310]: ### YOUR CODE HERE
          num of pcs = [25]
          for num in num of pcs:
              better_basis = covmat_eig[:,eig_loc[:num]]
              for image in test_images:
                  plt.figure()
                  current_test_image = data[image].flatten()
                  #tranform the test images
                  reduced_test_image = (current_test_image - data_mean) @ better_b
          asis
                  #transform them back
                  test image prime = (reduced test image @ better_basis.T) + data_
          mean
                  test_image_prime = np.array(test_image_prime)
                  reconstructed_data.append(test_image_prime)
                  image shaped = np.reshape(test_image_prime, (28,28))
                  plt.imshow(image_shaped)
          ### END OF CODE
```









(Your explanation)

I used 25 where it was to the point where I had zero issues trying to decipher what the digit was. To the point where i could identify it immediately. Using two principal components, many of digits looked like a 9 or a 0 and where therefore illegible. Using 25 dimensions you can easily identify while decreasing the dimension 30 fold.

Question 1.3 Harris Corner and PCA [10pt]

Recall Harris corner detector algorithm:

- 1. Compute x and y derivatives (I_x, I_y) of an image
- 2. Compute products of derivatives (I_x^2, I_y^2, I_{xy}) at each pixel
- 3. Compute matrix M at each pixel, where \$\$ $M(x_0,y_0) = \sum_{x \in \mathbb{Z}} w(x_x,y) \le \sum_{x \in \mathbb{Z}} w(x_x,y_0) \le \sum_{x \in \mathbb{Z}$

\end{bmatrix} \$\$ Here, we set weight w(x, y) to be a box filter of size 3×3 (the box is placed centered at (x_0, y_0)).

In this problem, you need to show that Harris Corner detector is really just principal component analysis in the gradient space. Your explanation should answer the following quesions.

- 1. As we know, PCA is performed on data points. What are the data points in Harris corner detector when we think of it as a PCA?
- 2. What is the covariance matrix used in Harris corner detector and why it is a covariance matrix?
- 3. What are the principal components in Harris corner detector?
- 4. Briefly explain how principal components imply "cornerness".

(Your proof here)

- 1. The data points are each of the image gradients
- 2. The covariance matrix is the \begin{bmatrix}

\end{bmatrix}

- 3. the principle components are the lambda values (eigen values of M for each x,y). Visualizing the eigenvalues in a ellipse allows us to see the direction of the fastest and slowest change. The higher eigenvalue's eigenvector points in the direction of fast change, and same with the smaller eigenvalue.
- 4. The principle components tell us the points at which the eigenvectors for that M matrix that produce the largest variations when moved in any direction. Based on the C value which is = $(\lambda_1 * \lambda_2) k * (\lambda_1 + \lambda_2)$. The higher this value, the more of a corner it is (highest variation in the directions of the eigenvectors(principle components), the lower the value it is a edge, and if it is a very small value, it is a flat region.

Question 2 KNN, Softmax Regression

In [312]: train X = train X.reshape((train X.shape[0], -1))

test X = test X.reshape((test X.shape[0], -1))

```
In [311]: train_dataset = MNIST(root='.', train=True, transform=transforms.ToTenso
    r, download=True)
    test_dataset = MNIST('.', train=False, transform=transforms.ToTensor())
    train_X = train_dataset.data.numpy() # training data, uint8 type to red
    uce memory and comparison cost
    train_y = train_dataset.targets.numpy() # training label
    test_X = test_dataset.data.numpy() # testing data, uint8 to reduce memor
    y and comparison cost
    test_y = test_dataset.targets.numpy() # testing label
```

Question 2.1 K-Nearest Neighbor [10pt]

In this problem you will be implementing the KNN classifier. Fill in the functions in the starter code below. You are are allowed to use <code>scipy.spatial.KDTree</code> and <code>scipy.stats.mode</code> (in case of a tie, pick any one). Please avoid <code>sklearn.neighbors.KDTree</code> as it appears extremely slow. You are **not** allowed to use a library KNN function that directly solves the problem.

If you do not know what a KD-tree is, please read the documentation for scipy.spatial.KDTree to understand how you can use it.

Note: if you run into memory issues or neighbor queries run for more than 10 minutes, you are allowed to reduce the data size, and explain what you have done to the training data.

```
In [313]: from scipy.spatial import KDTree from scipy.stats import mode
```

```
In [314]: class KNNClassifier:
              def init (self, num neighbors):
                  construct the classifier
                  Args:
                       num centers: number of neighbors
                  ### YOU CODE HERE
                  self.num_centers = num_neighbors
                  self.tree = None
                  self.X = None
                  self.y = None
                  ### END OF CODE
              def fit(self, X, y):
                   train KNN classifier
                  Args:
                       X: training data, numpy array with shape (Nxk) where N is nu
          mber of data points, k is number of features
                      y: training labels, numpy array with shape (N)
                  ### YOU CODE HERE
                  self.X = X
                  self.y = y
                  self.tree = KDTree(X)
                  ### END OF CODE
                  return self
              def predict(self, X):
                  predict labels
                  Args:
                      X: testing data, numpy array with shape (Mxk) where M is num
          ber of data points, k is number of features
                  Return:
                      y: predicted labels, numpy array with shape (N)
                  pred = []
                  ### YOU CODE HERE
                   for x in X:
                       d, i = self.tree.query([x], k=self.num centers)
                      neighbors labels = []
                       for index in i[0]:
                           neighbors labels.append(self.y[index])
                      most_common, _ = mode(neighbors_labels)
                       pred.append(most common[0])
                  ### END OF CODE
                  return pred
```

```
In [315]: from sklearn.metrics import accuracy_score
knn = KNNClassifier(6).fit(train_X, train_y)
pred_y = knn.predict(test_X)
# print(pred_y)
print('KNN accuracy:', accuracy_score(test_y, pred_y))
```

/Users/joshuakang/opt/anaconda3/lib/python3.7/site-packages/scipy/spati al/kdtree.py:388: RuntimeWarning: overflow encountered in ubyte_scalars sd[node.split_dim] = np.abs(node.split-x[node.split_dim])**p

KNN accuracy: 0.5727

Question 2.2 Softmax Regression

In this problm, you will be implementing the softmax regression(multi-class logistic regression). Here is a brief recap of several important concepts. In the following explanation, I will use x for data vector, y' for ground truth label, and y for predicted label.

Suppose we have a problem where we need to classify data points into m classes.

1. Softmax function S normalize a vector to have sum 1. (it turns any vector into a probability distribution)

$$S(x) = \left[\frac{e^{x_1}}{\sum_{j=1}^m e^{x_j}}, \frac{e^{x_2}}{\sum_{j=1}^m e^{x_j}}, \dots, \frac{e^{x_m}}{\sum_{j=1}^m e^{x_j}}\right]$$

2. Cross entropy loss J is the multiclass logistic regression loss.

$$J(y', y) = -\sum_{i=1}^{m} y_i' \log y_i$$

where y' is the one-hot ground truth label and y is the predicted label distribution.

3. Softmax regression is the following optimization problem.

$$\min_{W,b} \sum_{(X,y') \in \{\text{training set}\}} J(y', S(Wx+b))$$

where W has shape $(m \times k)$ where k is the number of features in a data point; b is a m dimensional vector.

4. This objective is optimized with gradient descent. Let

$$L = \sum_{(x,y') \in \{\text{training set}\}} J(y', S(Wx + b))$$

Update W and b with $\frac{\partial L}{\partial W}$ and $\frac{\partial L}{\partial b}$.

Question 2.2.1 Compute the gradients [10pt]

In this question, you need to do the following:

1. Compute the gradient $\frac{\partial J}{\partial y}$. i.e. compute

 $\frac{\partial J}{\partial v_i}$

Express it in terms of y'_i and y_i .

$$\frac{\partial J}{\partial y'} = -\sum_{i=1}^{m} \frac{y_i}{y_i'}$$

2. Let u = Wx + b, $y_i = S_i(u_i)$ Compute

$$\frac{\partial y_i}{\partial u_i}$$

Express it in terms of y_i , y_j and δ_{ij} , where

$$\delta_{ij} = \begin{cases} 1 & i = j \\ 0 & i \neq j \end{cases}$$

We have the case in which when we are getting the derivative of e^{a_j} (the numerator) that we need to check if the S_i' is i == j. To get the derivative of the fraction, we use the quotient rule. So first when i == j:

$$= \frac{(e^{a_i} \sum_{k=1}^{N} e^{a_k}) - e^{a_j} e^{a_i}}{(\sum_{k=1}^{N} e^{a_k})^2}$$

$$= \frac{e^{a_i}}{\sum_{k=1}^{N} e^{a_k}} * \frac{(\sum_{k=1}^{N} e^{a_k}) - e^{a_i}}{\sum_{k=1}^{N} e^{a_k}}$$

$$= S_i (1 - S_i)$$

and if i = j, then this is equivalent to

$$S_j(1-S_j) = S_j - S_j^2$$

Now in the second case that i != j, and using the same quotient rule :

$$= -\frac{\frac{0 - e^{a_j} e^{a_i}}{(\sum_{k=1}^{N} e^{a_k})^2}}{\sum_{k=1}^{N} e^{a_k}} \frac{e^{a_i}}{\sum_{k=1}^{N} e^{a_k}}$$
$$= -S_i S_i$$

Instead of leaving it as a system of equations, we can introduce a new variable to make it a single equation, δ :

$$\delta_{ij} = \begin{cases} 1 & i = j \\ 0 & i \neq j \end{cases}$$

Where

$$= \frac{\partial y_i}{\partial u_i} = S_i(\delta_{ij} - S_j)$$

1. Compute

$$rac{\partial J}{\partial W_{jk}}$$
 and $rac{\partial J}{\partial b_j}$

Express them in terms of y_j , y_j' , x_k . Explain your results in an intuitive way. Hint: the results should have a very simple form that makes sense intuitively. In order to calculate the $\frac{\partial J}{\partial W_{jk}}$, we must use chain rule, therefore:

$$\frac{\partial J}{\partial W_{jk}} = \frac{\partial J}{\partial y} \frac{\partial y}{\partial a} \frac{\partial a}{\partial W_{jk}}$$

Where a is the weighted sum of the inputs. This then results in:

$$(\sum_{i!=k}^{m} \frac{-y'_{k}}{y_{k}}(-y_{k}y_{i}) + \sum_{i=k}^{m} y'_{k} + y'_{k}y_{i})$$

$$= (\sum_{i!=k}^{m} y'_{k}y_{i} + \sum_{i=k}^{m} -y'_{k} + y'_{k}y_{i})$$

$$= (\sum_{i!=k}^{m} y'_{k}y_{i} - \sum_{i=k}^{m} y'_{k} + \sum_{i=k}^{m} y'_{k}y_{i})$$

$$= (\sum_{k}^{m} y'_{k}y_{i} - \sum_{k=i}^{m} y'_{k})$$

$$= (1 * y_{i} - y'_{i})$$

$$= (y_{i} - y'_{i})$$

The final term of the chain rule is the partial of the weighted sum with respect to the weight that we are updating. In this case, the result is simply the x_i (the input that we multiply by the w_{ij} with to add to the sum of the total weighted sum.

So putting them together, we get for a weight w_{ij} the gradient is :

$$(y_j - y_j')x_i$$

For the bias, it is simply:

$$(y_j - y_i') * 1$$

since the input for the bias is always 1.

1. Compute

$$\frac{\partial J}{\partial W}$$

in the matrix form. It should be a matrix with the same shape as W, and entry jk is $\frac{\partial J}{\partial W_{jk}}$. Similarly, compute

$$\frac{\partial J}{\partial h}$$

Given the input to the layer is X such that the shape is $N \times d$ ($N : number of datapoints, d : number of dimensions). Also, the gradient is of shape : <math>N \times o$ (o : number of output units of layer). The matrix multiplication for the regular inputs is X.T @ -Gradient, and for the bias it is a [1 x n] @ -Gradient.

(Your proof here)

Question 2.2.2 Stochastic Gradient Descent [10pt]

In gradient descent algorithm, we update W and b with $\partial L/\partial W$ and $\partial L/\partial b$. However, this requires the gradient w.r.t. the whole dataset. Computing such gradient is very slow. Instead, we can update the weights with perdata gradient. This is known as the SGD algorithm, which runs much faster. You need to take the following steps.

- 1. Implement softmax function S. We need to take special care in this function since e^x tends to overflow easily. However, we observe that S(x) = S(x m) for any constant vector m. We can stabilize softmax using $S(x) = S(x \max(x))$.
- 2. Implement function J (loss) and dJ (loss gradient). Note: J is not required to run the algorithm, but you may want to implement it for debug purposes.
- 3. Implement the SGD algorithm.
- 4. Run the algorithm for 20 epochs (each epoch iterates the whole data set once) with learning rate 1e-3 and report accuracy on test set. You may use sklearn.metrics.accuracy_score. You need to achieve accuracy > 90%. You are allowed to experiment with different epoch numbers and learning rates (even learning rate decay) to achieve this accuracy, but they are not required.

You may use print (or progress bar packages) to track the training progress since it might take several minutes.

In [197]: train_dataset = MNIST(root='.', train=True, transform=transforms.ToTenso
r, download=True)
 test_dataset = MNIST('.', train=False, transform=transforms.ToTensor())
 train_X = train_dataset.data.numpy() / 255. # normalize data to 0-1
 train_y = train_dataset.targets.numpy() # training label
 test_X = test_dataset.data.numpy() / 255. # normalize data to 0-1
 test_y = test_dataset.targets.numpy() # testing label
 train_X = train_X.reshape((train_X.shape[0], -1)) # flatten the image
 test_X = test_X.reshape((test_X.shape[0], -1)) # flatten the image

```
In [316]: def softmax(x):
              softmax function
              Args:
                  x: a 1-d numpy array
              Return:
                  results of softmax(x)
              ### YOUR CODE HERE
              exp_x = np.exp(x)
              softmax_output = exp_x / (np.sum(exp_x))
              return softmax_output
                return softmax output - np.amax(softmax output)
              ### END OF CODE
          def J(W, b, y_true, x):
              Softmax Loss function
              Args:
                  W: weights (num classes x num features)
                  b: bias (num classes)
                  y true: ground truth 1-hot label (num classes)
                  x: input data
              Return:
                  J(y', y)
              ### YOUR CODE HERE
              weighted sum = W @ x + b
              activation = softmax(weighted sum)
              error = 0
              for m in x.shape[0]:
                  error += y_true * np.log(activation[m])
              return -error
              ### END OF CODE
          def dJ(W, b, y_true, x):
              Softmax Loss gradient
              Args:
                  W: weights (num classes x num features)
                  b: bias (num features)
                  y_true: ground truth 1-hot label (num_classes)
                  x: input data (num features)
              Return:
                   (dW, db): gradient w.r.t. W and b
              ### YOUR CODE HERE
              weighted sum = W @ x + b
                print('weighted sum')
                print(weighted sum)
```

```
# print('activations')
activation = softmax(weighted_sum)
delta = activation - y_true
    print(activation)
dW = delta[np.newaxis].T @ x[np.newaxis]
db = delta[np.newaxis].T @ np.ones((1,1))

return dW, np.reshape(db, (10,))

#get weighted sums
#put it in the softmax, and get the y_pred
#do y - t and that is the upstream gradient
#dJ/dW is x.T @ (y-t)
#dJ/db is (ones column) @ (y-t)

#Then W = W - lr * dJ/dW
#and b = b - lr * dJ/db

### END OF CODE
```

```
In [317]: from tqdm import tqdm notebook
```

```
In [217]: def SGD(f, df, Xs, ys, n_classes=10, lr=1e-3, max_epoch=20):
              Args:
                  f: function to optimize
                  df: the gradient of the function
                  Xs: input data, numpy array with shape (num_data x num_features)
                  ys: true label, numpy array with shape (num data x num classes)
                  lr: learning rate
                  max epoch: maximum epochs to run SGD
              Return:
                  optimal weights and biases
               .....
              N, m = Xs.shape
              W = np.random.rand(n_classes, m) - 0.5 # you do not need to change
           random initialization
              b = np.random.rand(n_classes) - 0.5
              for epoch in range(max_epoch):
                  for x,y in zip(Xs, ys):
                      dW, db = dJ(W, b, y, x)
                      W = lr * dW
                      b = 1r * db
              ### YOUR CODE HERE
              ### END OF CODE
              return W, b
```

```
In [218]: train_y_onehot = np.zeros((train_y.shape[0], 10))
    train_y_onehot[np.arange(len(train_y)), train_y] = 1
    W, b = SGD(J, dJ, train_X, train_y_onehot, 10, max_epoch=20)
    accuracy_score(test_y, np.argmax(test_X @ W.T + b, axis=1))
Out[218]: 0.9182
```

Question 3 Convolutional Neural Networks

This question requires you to use the PyTorch framework for neural network training. You will not need GPU to train the networks for this problem.

The following is a code sample for training a simple multi-layer perceptron neural network using PyTorch. Running it should give you about 98% testing accuracy.

Since network training takes long, I recommend installing the tqdm package for progress tracking.

```
from tqdm import tqdm notebook
In [318]:
In [319]: train dataset = MNIST(root='.', train=True, transform=transforms.ToTenso
          r(), download=True)
          test dataset = MNIST('.', train=False, transform=transforms.ToTensor())
In [320]: class MLP(nn.Module):
              def init (self, input size, hidden size, num classes):
                  """init function builds the required layers"""
                  super(MLP, self). init () # This line is always required
                  # Hidden layer
                  self.layer1 = nn.Linear(input size, hidden size)
                  # activation
                  self.relu = nn.ReLU()
                  # output layer
                  self.layer2 = nn.Linear(hidden size, num classes)
              def forward(self, x):
                  """forward function describes how input tensor is transformed to
          output tensor"""
                  # flatten the input from (Nx1x28x28) to (Nx784)
                  torch.flatten(x, 1)
                  x = self.layer1(x)
                  x = self.relu(x)
                  x = self.layer2(x)
                  # Note we do not need softmax layer, since this layer is include
          d in the CrossEntropyLoss provided by torch
                  return x
```

```
model = MLP(784, 1024, 10)
In [321]:
          model
Out[321]: MLP(
            (layer1): Linear(in_features=784, out_features=1024, bias=True)
            (relu): ReLU()
            (layer2): Linear(in_features=1024, out_features=10, bias=True)
          )
In [322]: opts = {
              'lr': 5e-4,
              'epochs': 5,
              'batch_size': 64
          }
In [323]: optimizer = torch.optim.Adam(model.parameters(), opts['lr']) # Adam is
           a much better optimizer compared to SGD
          criterion = torch.nn.CrossEntropyLoss() # loss function
          train_loader = torch.utils.data.DataLoader(dataset=train_dataset, batch_
          size=opts['batch_size'], shuffle=True)
          test_loader = torch.utils.data.DataLoader(dataset=test_dataset, batch_si
          ze=opts['batch_size'], shuffle=True)
```

```
In [324]: for epoch in range(opts['epochs']):
              train loss = []
              for i, (data, labels) in tqdm notebook(enumerate(train loader), tota
          l=len(train_loader)):
                  # reshape data
                  data = data.reshape([-1, 784])
                  # pass data through network
                  outputs = model(data)
                  loss = criterion(outputs, labels)
                  optimizer.zero_grad() # Important! Otherwise the optimizer will
          accumulate gradients from previous runs!
                  loss.backward()
                  optimizer.step()
                  train loss.append(loss.item())
              test loss = []
              test_accuracy = []
              for i, (data, labels) in enumerate(test_loader):
                  # reshape data
                  data = data.reshape([-1, 784])
                  # pass data through network
                  outputs = model(data)
                  _, predicted = torch.max(outputs.data, 1)
                  loss = criterion(outputs, labels)
                  test loss.append(loss.item())
                  test accuracy.append((predicted == labels).sum().item() / predic
          ted.size(0))
              print('epoch: {}, train loss: {}, test loss: {}, test accuracy: {}'.
          format(epoch, np.mean(train loss), np.mean(test loss), np.mean(test accu
          racy)))
          epoch: 0, train loss: 0.28823375489030567, test loss: 0.136084674033010
          08, test accuracy: 0.960390127388535
          epoch: 1, train loss: 0.11209188347289176, test loss: 0.090257487675044
          57, test accuracy: 0.971437101910828
          epoch: 2, train loss: 0.07314085077667144, test loss: 0.081120974455432
          15, test accuracy: 0.9755175159235668
```

epoch: 3, train loss: 0.05211033039951701, test loss: 0.061676299605803

epoch: 4, train loss: 0.037828655509865965, test loss: 0.06651330238006

314, test accuracy: 0.980593152866242

486, test accuracy: 0.9786027070063694

Question 3.1 Implementing CNN [15pt]

You need to implement a convolutional neural network for the same task as above. You may find the PyTorch documentation helpful. https://pytorch.org/docs/stable/nn.html (https://pytorch.org/nn.html (<a href="https://py

We provide a working network structure below. You can adjust the network size and training options for better performance, but a correct implementation of the provided network should give you the required accuracy. For convolutional layers, (conv MxM, N) means the layer has kernel size M by M and N output channels; for pooling layers, (maxpool MxM) means doing max pooling with kernel size M by M.

(conv 5x5, 32) -> (relu) -> (maxpool 2x2) -> (conv 5x5, 64) -> (relu) -> (maxpool 2x2) -> (flatten) -> (linear 10) -> (output)

For full score, you need to achieve 99% testing accuracy. Also, plot the hand-written digits that your network got wrong.

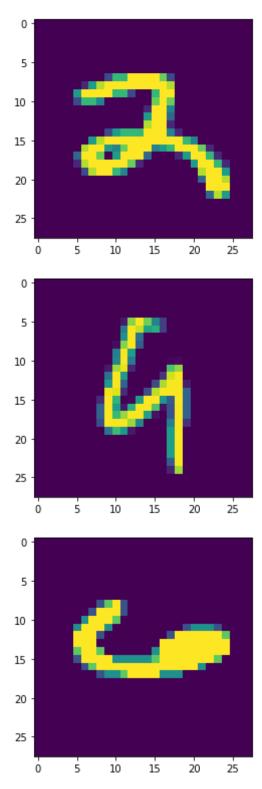
```
In [362]: class CNN(nn.Module):
              def __init__(self, input size, num classes):
                   init convolution and activation layers
                   Args:
                       input size: (1,28,28)
                       num classes: 10
                   super(CNN, self).__init__()
                   ### YOUR CODE HERE
                   self.first = nn.Conv2d(1, 32 , 5, stride=1, padding=2, bias=Fals
          e)
                   self.main = nn.Sequential(
                      nn.ReLU(inplace=True),
                      nn.MaxPool2d(kernel_size=2, stride=2),
                       nn.Conv2d(32 , 64, 5, stride=1, padding=2, bias=False),
                      nn.ReLU(inplace=True),
                      nn.MaxPool2d(kernel_size=2, stride=2),
                  self.fc = nn.Sequential(
                       nn.Linear(3136, num_classes)
                   ### END OF CODE
              def forward(self, x):
                   forward function describes how input tensor is transformed to ou
           tput tensor
                  Args:
                      x: (Nx1x28x28) tensor
                  ### YOUR CODE HERE
                  x = self.first(x)
                  x = self.main(x)
                  x = x.reshape(x.size(0), -1)
                  x = self.fc(x)
                   ### END OF CODE
                  return x
```

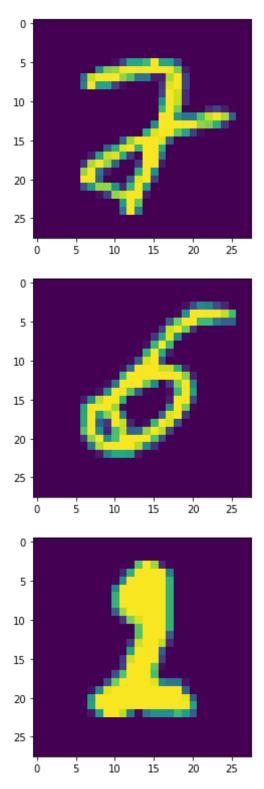
```
In [363]:
          model = CNN((1, 28, 28), 10)
          model
Out[363]: CNN(
            (first): Conv2d(1, 32, kernel_size=(5, 5), stride=(1, 1), padding=(2,
          2), bias=False)
            (main): Sequential(
              (0): ReLU(inplace=True)
              (1): MaxPool2d(kernel_size=2, stride=2, padding=0, dilation=1, ceil
          mode=False)
              (2): Conv2d(32, 64, kernel_size=(5, 5), stride=(1, 1), padding=(2,
          2), bias=False)
              (3): ReLU(inplace=True)
              (4): MaxPool2d(kernel_size=2, stride=2, padding=0, dilation=1, ceil
          _mode=False)
            )
            (fc): Sequential(
              (0): Linear(in_features=3136, out_features=10, bias=True)
            )
          )
In [369]:
          ### You may (and should) change these
          opts = {
               'lr': 1e-3,
               'epochs': 10,
              'batch size': 64
          }
          ### if you cannot get 99% with SGD, Adam optimizer can help you
          optimizer = torch.optim.Adam(model.parameters(), opts['lr'])
In [370]: criterion = torch.nn.CrossEntropyLoss() # loss function
          train loader = torch.utils.data.DataLoader(dataset=train dataset, batch
          size=opts['batch size'], shuffle=True)
          test loader = torch.utils.data.DataLoader(dataset=test dataset, batch si
          ze=opts['batch size'], shuffle=True)
```

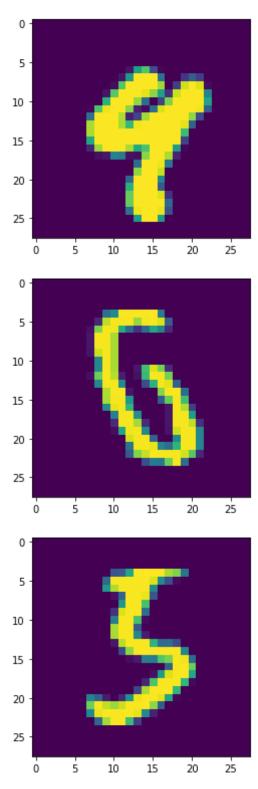
```
In [ ]: for epoch in range(opts['epochs']):
            train loss = []
            for i, (data, labels) in tqdm_notebook(enumerate(train_loader), tota
        l=len(train_loader)):
                # pass data through network
                outputs = model(data)
                loss = criterion(outputs, labels)
                optimizer.zero grad()
                loss.backward()
                optimizer.step()
                train loss.append(loss.item())
            test_loss = []
            test_accuracy = []
            for i, (data, labels) in enumerate(test loader):
                 # pass data through network
                outputs = model(data)
                _, predicted = torch.max(outputs.data, 1)
                loss = criterion(outputs, labels)
                test_loss.append(loss.item())
                wrong = []
                for label_i in range(len(labels)):
                     if predicted[label_i] != labels[label_i]:
                         wrong.append(np.squeeze(data[label i]))
                         plt.figure()
                        plt.imshow(wrong[-1])
                test accuracy.append((predicted == labels).sum().item() / predic
        ted.size(0))
            print('epoch: {}, train loss: {}, test loss: {}, test accuracy: {}'.
        format(epoch, np.mean(train loss), np.mean(test loss), np.mean(test accu
        racy)))
```

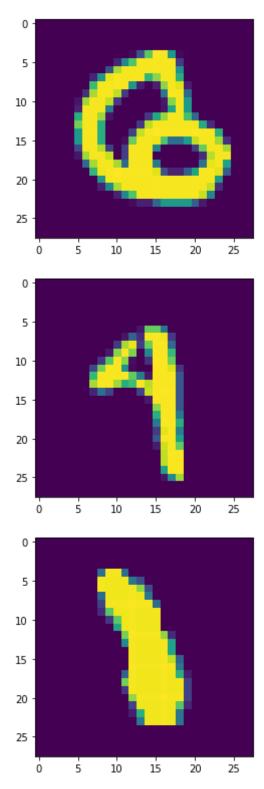
```
/Users/joshuakang/opt/anaconda3/lib/python3.7/site-packages/ipykernel 1
auncher.py:24: RuntimeWarning: More than 20 figures have been opened. F
igures created through the pyplot interface (`matplotlib.pyplot.figure
`) are retained until explicitly closed and may consume too much memor
y. (To control this warning, see the rcParam `figure.max open warning
`).
epoch: 0, train loss: 0.019554508298107145, test loss: 0.02285140537961
5997, test accuracy: 0.9931329617834395
epoch: 1, train loss: 0.013642958154125895, test loss: 0.03457956339269
142, test accuracy: 0.9898487261146497
                                          Traceback (most recent call 1
KeyboardInterrupt
ast)
<ipython-input-371-502b2b3d73b1> in <module>
      6
                loss = criterion(outputs, labels)
      7
                optimizer.zero grad()
---> 8
                loss.backward()
      9
                optimizer.step()
     10
                train_loss.append(loss.item())
~/opt/anaconda3/lib/python3.7/site-packages/torch/tensor.py in backward
(self, gradient, retain_graph, create_graph)
    148
                        products. Defaults to `False`.
    149
--> 150
                torch.autograd.backward(self, gradient, retain graph, c
reate graph)
    151
    152
            def register hook(self, hook):
~/opt/anaconda3/lib/python3.7/site-packages/torch/autograd/ init .py
 in backward(tensors, grad tensors, retain graph, create graph, grad va
riables)
     97
            Variable. execution engine.run backward(
                tensors, grad_tensors, retain_graph, create graph,
     98
---> 99
                allow unreachable=True) # allow unreachable flag
    100
    101
```

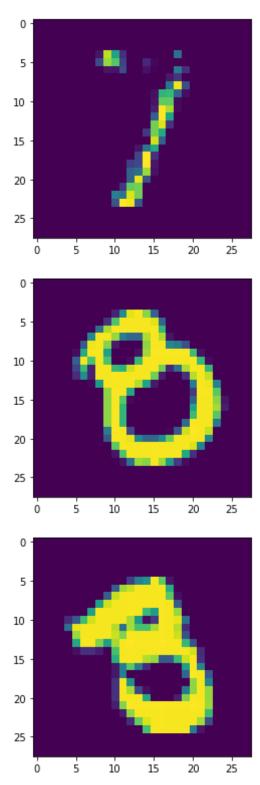
KeyboardInterrupt:

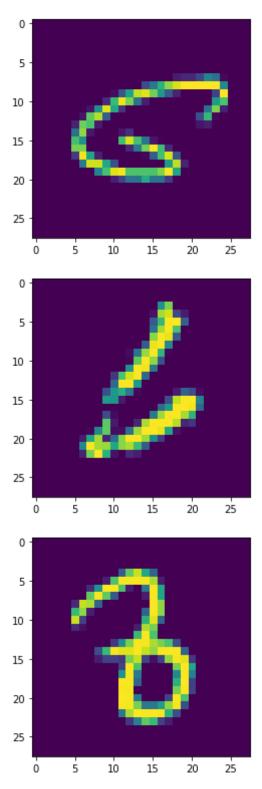


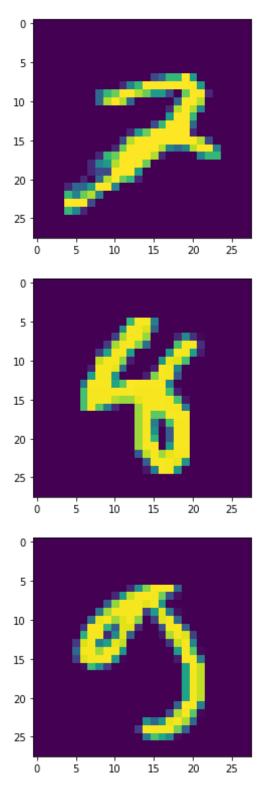


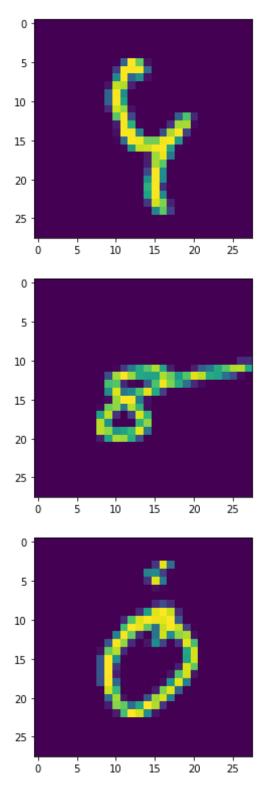


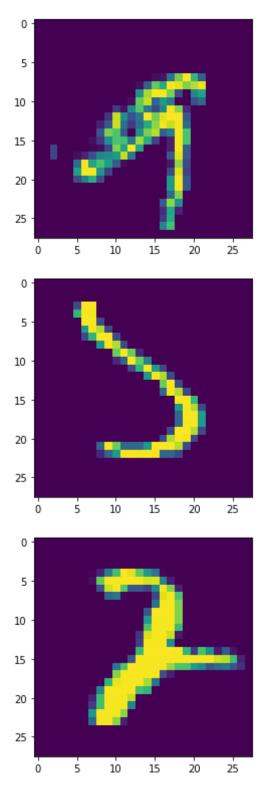


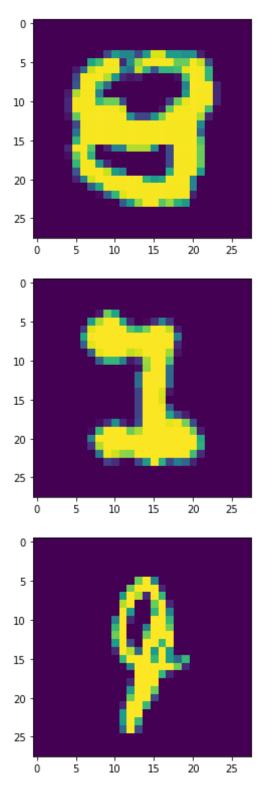


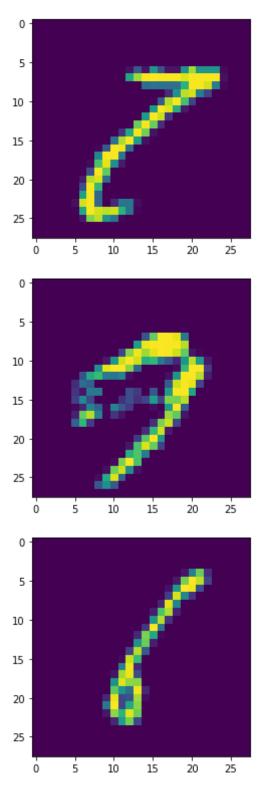


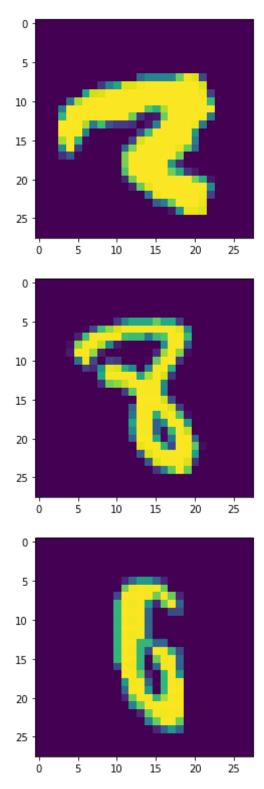


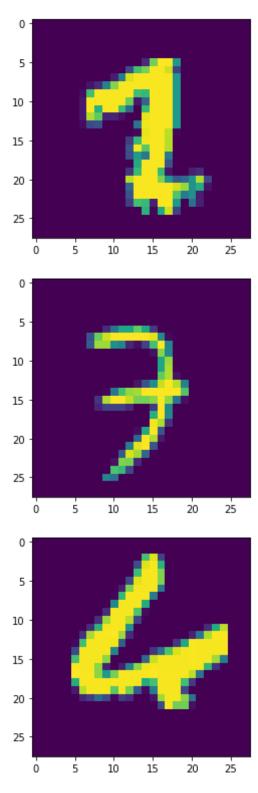


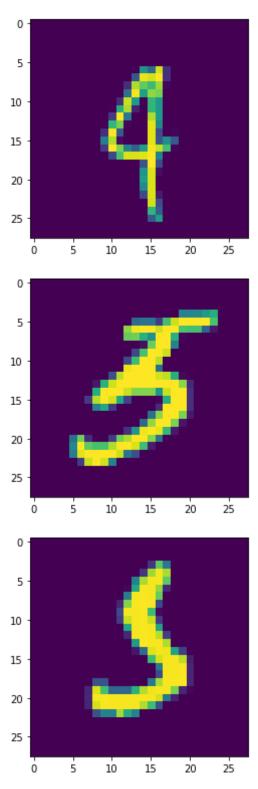


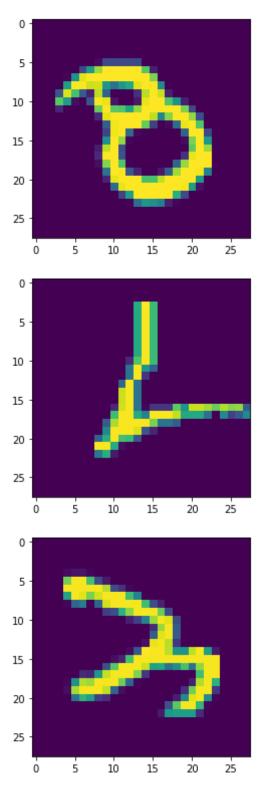


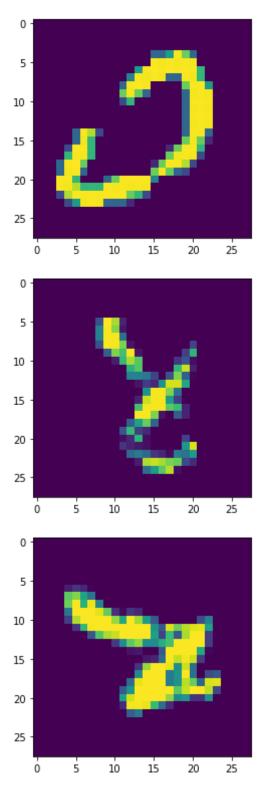


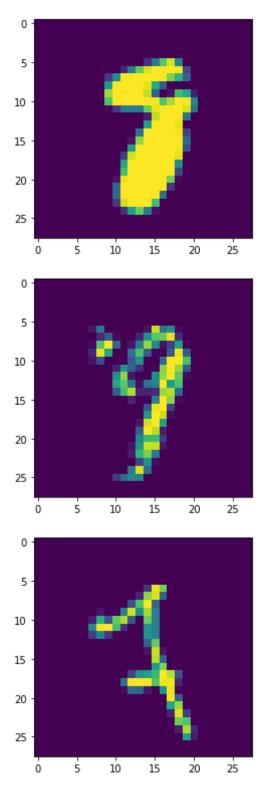


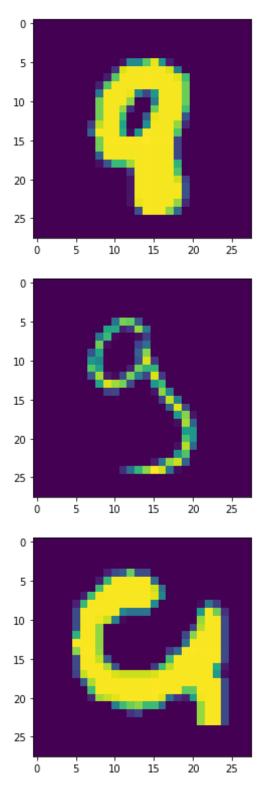


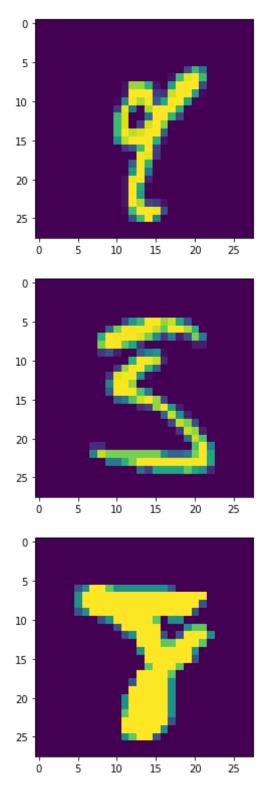


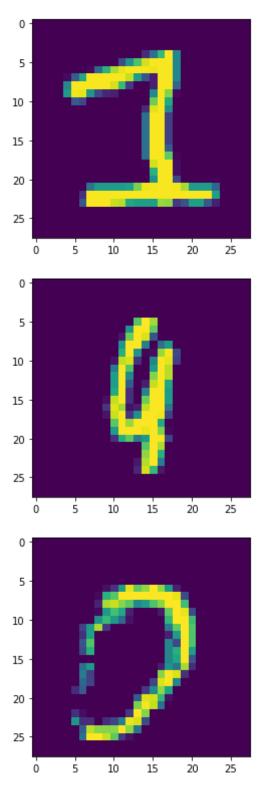


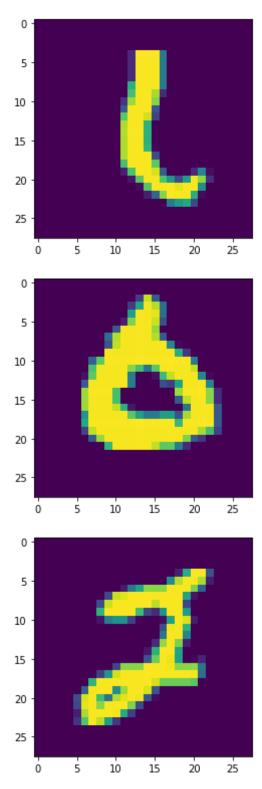


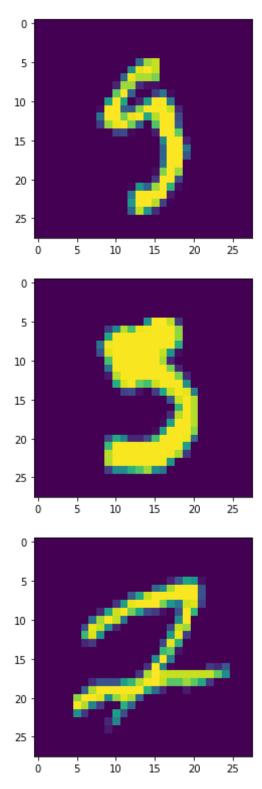


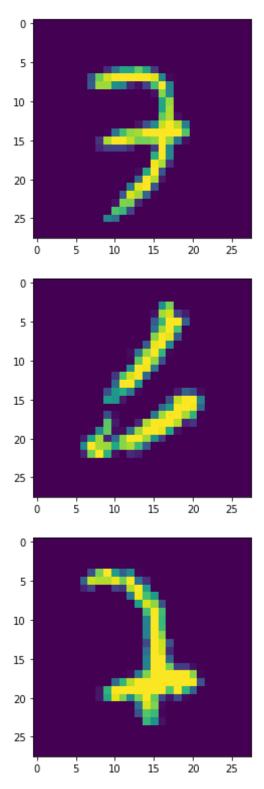


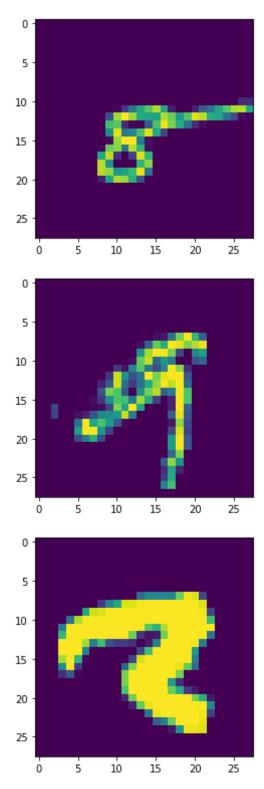


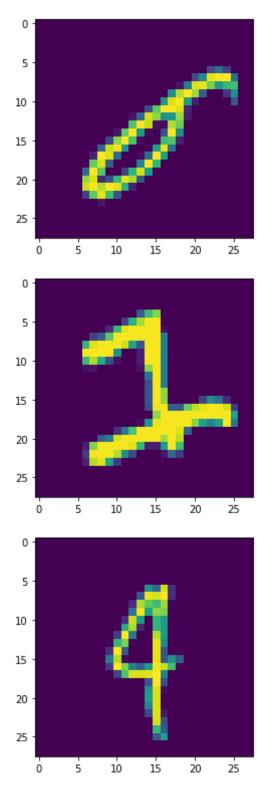


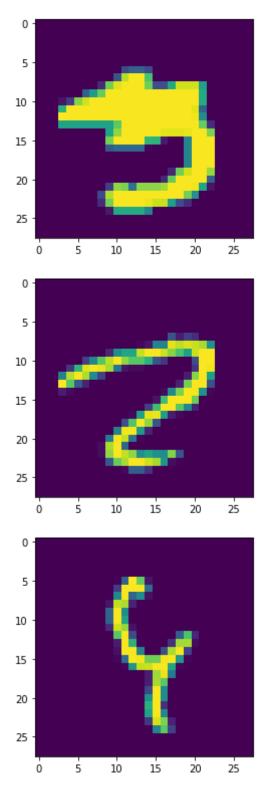


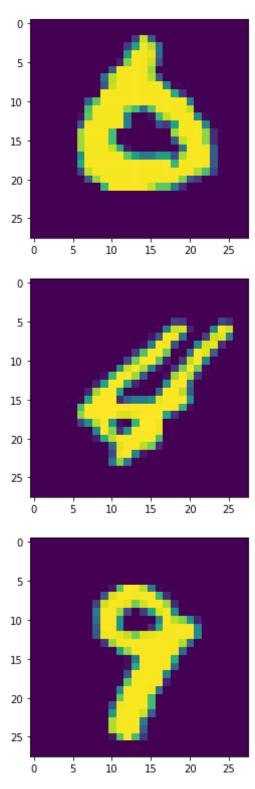


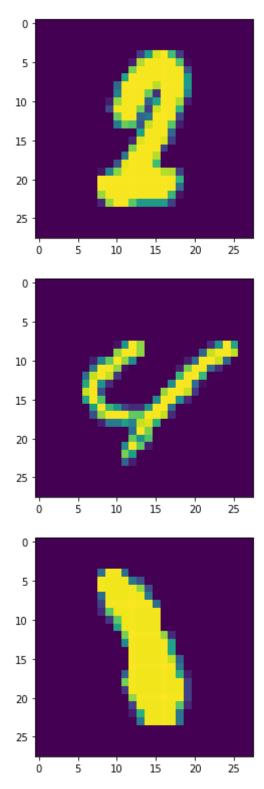


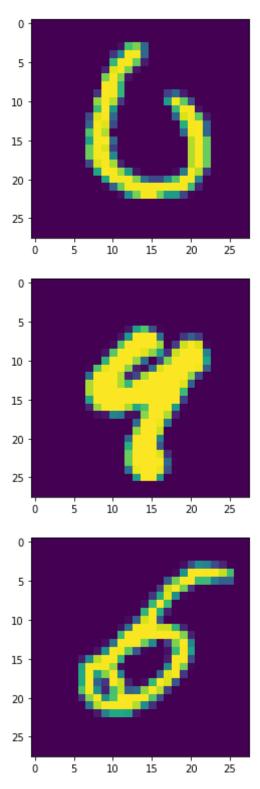


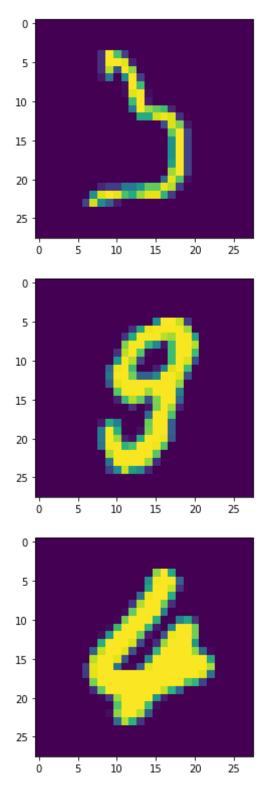


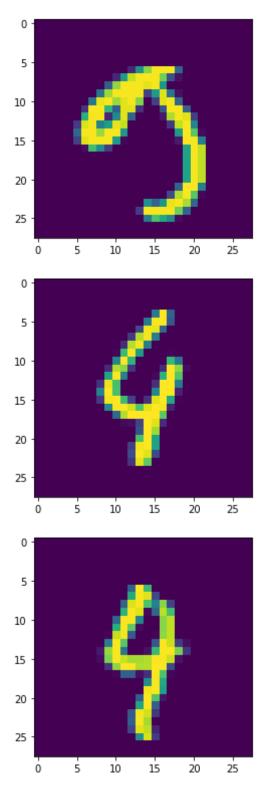


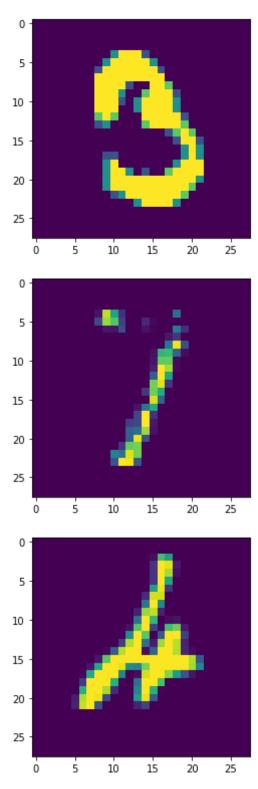


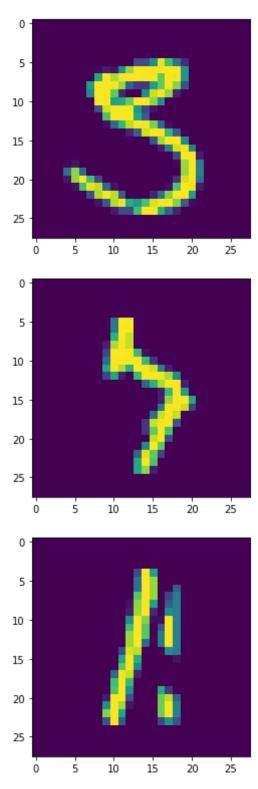


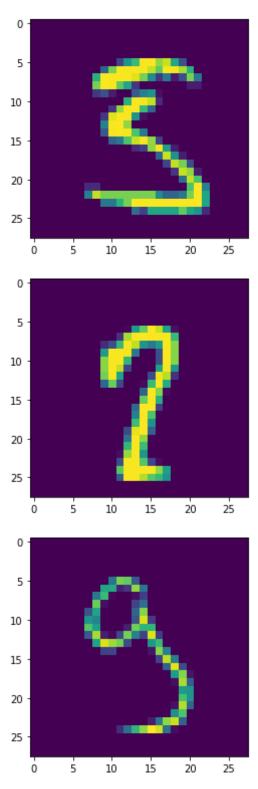


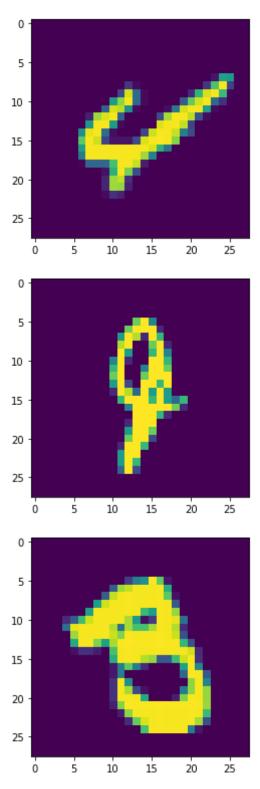


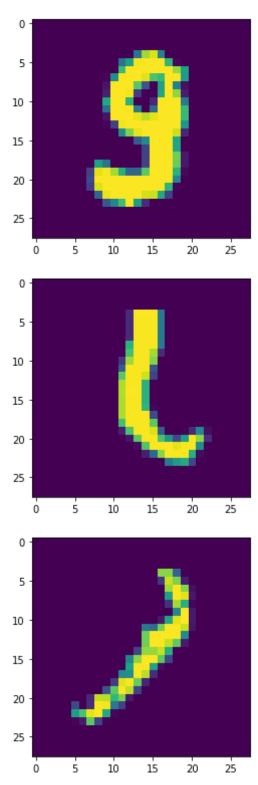


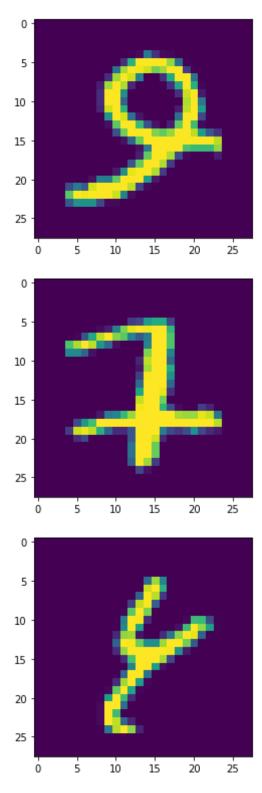


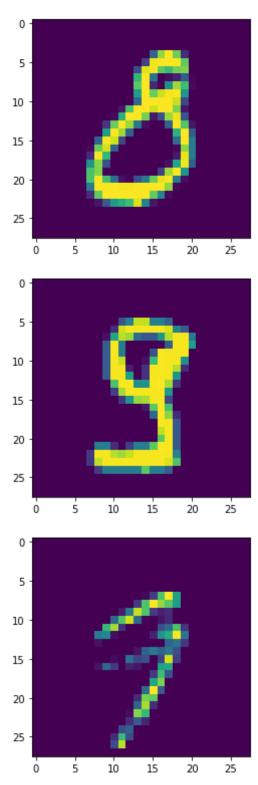


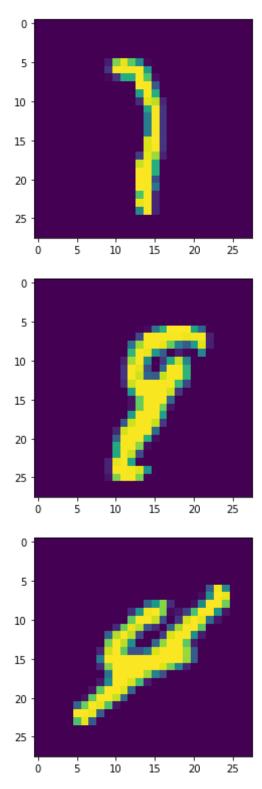


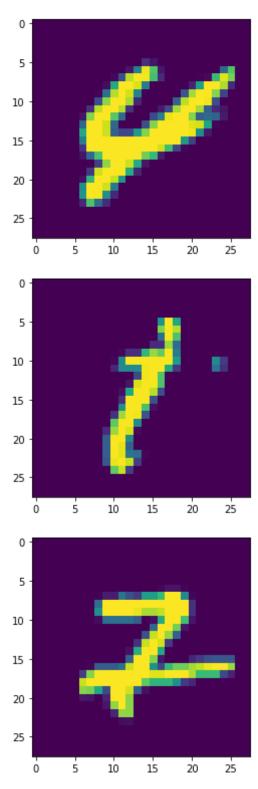


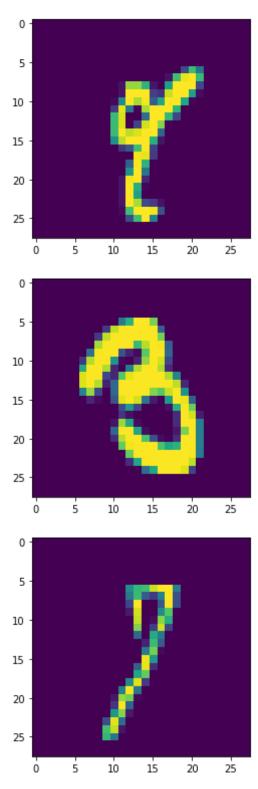


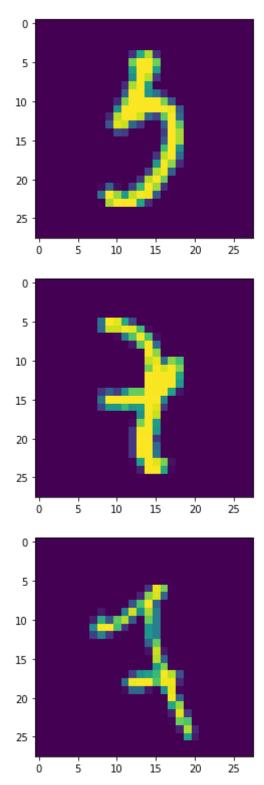


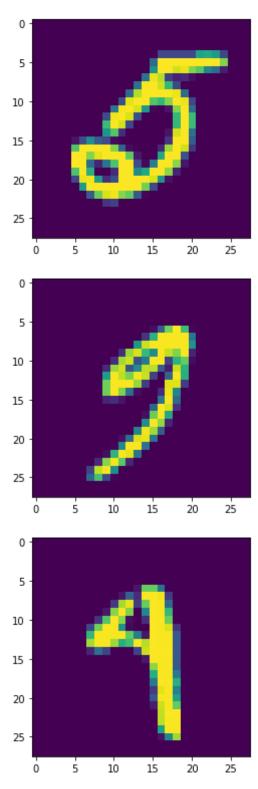


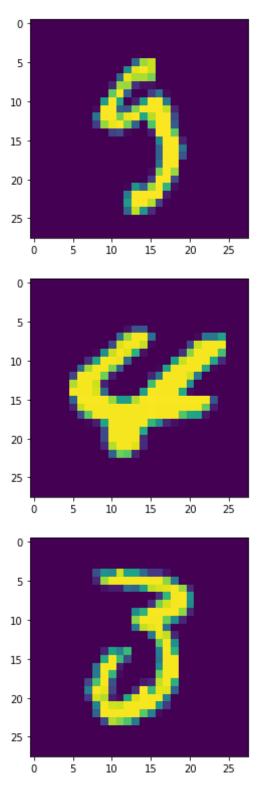


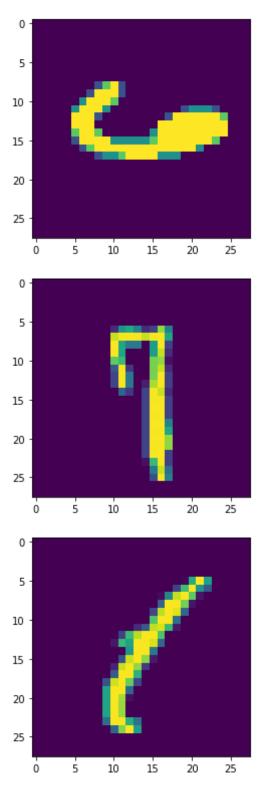


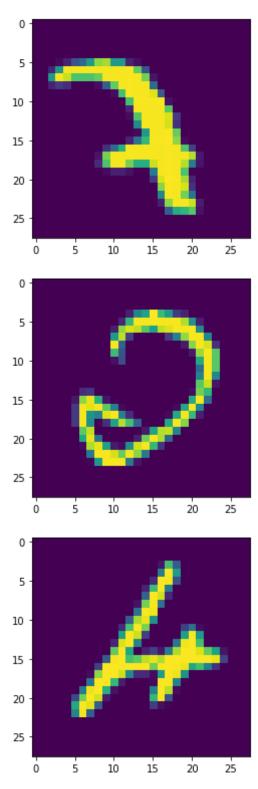


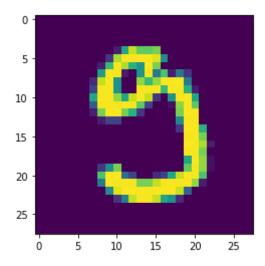












Don't forget plotting the digits that the network got wrong.

Question 3.2 Kernel weights visualization [5pt]

For this question, you need to visualize the kernel weights for your first convolutional layer. Suppose you have 5x5 kernels with 32 output channels. You will plot 32 5x5 images.

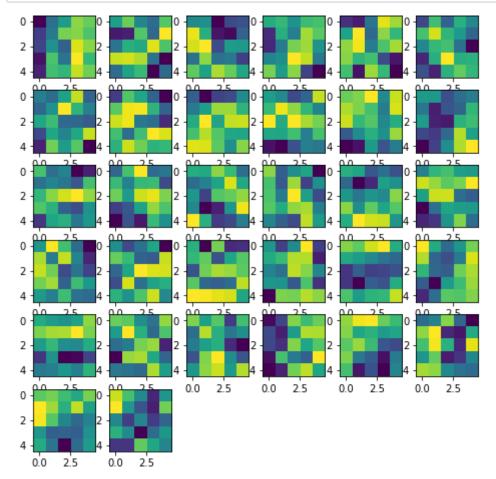
hint: You might need to look at PyTorch documentation (or play with the PyTorch model) to figure out how to get the weights.

```
In [367]: ### YOUR CODE HERE
   weights = model.first.weight
   weights.shape

   fig=plt.figure(figsize=(8, 8))

   count = 1
   for i in range(weights.shape[0]):
        kernel = weights[i]
        kernel = np.squeeze(kernel)
        fig.add_subplot(6, 6, count)
        plt.imshow(kernel.detach().numpy())

        count+= 1
        ### END OF CODE
```



In []: