Applying different Convolutional Neural Networks on MNIST data

ConvNets are designed specifically for image datasets. It is inspired from Biological visual cortex.

Steps followed in this assignment:-

- 1. Load libraries and MNIST data
- 2. Defing parameters
- 3. Normalizing the data
- 4. Convert class label values to one-hot encoded values
- 5. Implementing CNN with 3 Convolution layers with Kernel size of 2X2, 5 Convolution layers with Kernel size of 5X5 and 7 Convolution layers with Kernel size of 3X3.
- 6. Implementing max-pooling, Padding, Strides, Dropouts and Batch Normalization in each model.
- 7. Compiling the model
- 8. Plotting Categorical Crossentropy Loss vs Epochs plot
- 9. Plotting Violin plots to check weights distribution.

1. Loading Libraries

```
In [0]: from __future__ import print_function
    import keras
    from keras.datasets import mnist
    from keras.models import Sequential
    from keras.layers import Dense, Dropout, Flatten
    from keras.layers import Conv2D, MaxPooling2D
    from keras import backend as K
    import numpy as np
    import matplotlib.pyplot as plt
    %matplotlib inline
```

Using TensorFlow backend.

1.1 Loading MNIST data

```
In [0]: # Defing some parameters
    batch_size = 128
    num_classes = 10
    epochs = 12

# input image dimensions
    img_rows, img_cols = 28, 28

# Loading MNIST data
# the data, split between train and test sets
(X_train, y_train), (X_test, y_test) = mnist.load_data()
Downloading data from https://s3.amazonaws.com/img_datasets/mpist.nnz
```

1.2 Normalizing the data

```
In [0]: if K.image data format() == 'channels first':
            X_train = X_train.reshape(X_train.shape[0], 1, img_rows, img_cols)
            X_test = X_test.reshape(X_test.shape[0], 1, img_rows, img_cols)
            input_shape = (1, img_rows, img_cols)
        else:
            X_train = X_train.reshape(X_train.shape[0], img_rows, img_cols, 1)
            X_test = X_test.reshape(X_test.shape[0], img_rows, img_cols, 1)
            input_shape = (img_rows, img_cols, 1)
        X train = X train.astype('float32')
        X_test = X_test.astype('float32')
        # Normalizing the data
        X train /= 255
        X_test /= 255
        print('X_train shape:', X_train.shape)
        # Printing to see how many data points are there for train and test and to see
        each image size
        print("Number of Train data points :", X train.shape[0], "and each image is of
        shape (%d, %d)"%(X train.shape[1], X train.shape[2]))
        print("Number of Test data points :", X_test.shape[0], "and each image is of s
        hape (%d, %d)"%(X test.shape[1], X test.shape[2]))
```

```
X_train shape: (60000, 28, 28, 1)
Number of Train data points: 60000 and each image is of shape (28, 28)
Number of Test data points: 10000 and each image is of shape (28, 28)
```

1.3 Converting to One-Hot encoding

```
In [0]: # Here in this dataset, class labels for each image are numbers (0,1,2,3,...),
        so I want to convert them into one-hot encoded vectors
        print("Class label of first image :", y_train[0])
        # lets convert this into a 10 dimensional vector
        # ex: consider an image with class label of 5 and convert it into one-hot enco
        ded vector of 0's and 1's - 5 => [0, 0, 0, 0, 0, 1, 0, 0, 0]
        # this conversion is needed for MLPs
        # convert class vectors to binary class matrices
        Y_train = keras.utils.to_categorical(y_train, num_classes)
        Y_test = keras.utils.to_categorical(y_test, num_classes)
        print("After converting the output into a vector : ",Y train[0])
        Class label of first image : 5
        After converting the output into a vector : [0. 0. 0. 0. 0. 1. 0. 0. 0. 0.]
In [0]: # this function is used draw Categorical Crossentropy Loss vs No. of epochs pl
        import numpy as np
        import time
        def plt dynamic(x, vy, ty):
            plt.figure(figsize=(10,6))
            plt.plot(x, ty, 'b', label="Train Loss")
            plt.plot(x, vy, 'r', label="Validation/Test Loss")
            plt.title('\nCategorical Crossentropy Loss VS Epochs')
            plt.xlabel('Epochs')
            plt.ylabel('Categorical Crossentropy Loss-Train and Test loss')
            plt.legend()
            plt.grid()
            plt.show()
```

2. CNN with 3 Convolution layers with Kernel size of 2X2

2.1 Building the model

```
In [0]: %matplotlib inline
    import warnings
    warnings.filterwarnings("ignore")
```

```
In [0]: from keras.layers.normalization import BatchNormalization
        from keras.layers import Dropout
        from keras.models import Sequential
        from keras.layers import Dense, Activation
        from keras.initializers import he normal
        %matplotlib inline
        import warnings
        warnings.filterwarnings("ignore")
        # start building a model
        # Initializing the sequential model as the layers are in sequential
        model 1 = Sequential()
        # Adding first Convolution layer
        model_1.add(Conv2D(32, kernel_size=(2, 2), activation='relu', input_shape=inpu
        t shape))
        # Adding Maxpooling layer
        model 1.add(MaxPooling2D(pool size=(2, 2)))
        # Adding Dropout
        model_1.add(Dropout(0.25))
        # Adding second Convolution layer
        model_1.add(Conv2D(64, (2, 2), activation='relu'))
        # Adding Batch Normalization
        model 1.add(BatchNormalization())
        # Adding Maxpooling Layer
        model 1.add(MaxPooling2D(pool size=(2, 2)))
        # Adding Dropout
        model_1.add(Dropout(0.25))
        # Adding third Convolution layer
        model_1.add(Conv2D(128, (2, 2), activation='relu'))
        # Adding Maxpooling layer
        model 1.add(MaxPooling2D(pool size=(2, 2)))
        # Adding Dropout
        model_1.add(Dropout(0.25))
        # Adding flatten layer
        model_1.add(Flatten())
        # Adding a dense layer
        model_1.add(Dense(256, activation='relu', kernel_initializer=he_normal(seed=No
        ne)))
        # Adding Batch Normalization
        model 1.add(BatchNormalization())
        # Adding Dropout
        model 1.add(Dropout(0.5))
        # Output softmax layer
        model 1.add(Dense(num classes, activation='softmax'))
        # Printing model summary
        model_1.summary()
```

Layer (type)	Output	Shape	Param #
conv2d_356 (Conv2D)	(None,	27, 27, 32)	160
max_pooling2d_325 (MaxPoolin	(None,	13, 13, 32)	0
dropout_337 (Dropout)	(None,	13, 13, 32)	0
conv2d_357 (Conv2D)	(None,	12, 12, 64)	8256
batch_normalization_179 (Bat	(None,	12, 12, 64)	256
max_pooling2d_326 (MaxPoolin	(None,	6, 6, 64)	0
dropout_338 (Dropout)	(None,	6, 6, 64)	0
conv2d_358 (Conv2D)	(None,	5, 5, 128)	32896
max_pooling2d_327 (MaxPoolin	(None,	2, 2, 128)	0
dropout_339 (Dropout)	(None,	2, 2, 128)	0
flatten_31 (Flatten)	(None,	512)	0
dense_67 (Dense)	(None,	256)	131328
batch_normalization_180 (Bat	(None,	256)	1024
dropout_340 (Dropout)	(None,	256)	0
dense_68 (Dense)	(None,	10)	2570
Total params: 176.490	=====:	=======================================	=======

Total params: 176,490 Trainable params: 175,850 Non-trainable params: 640

2.2 Compiling the model

In [0]: # Compiling the model

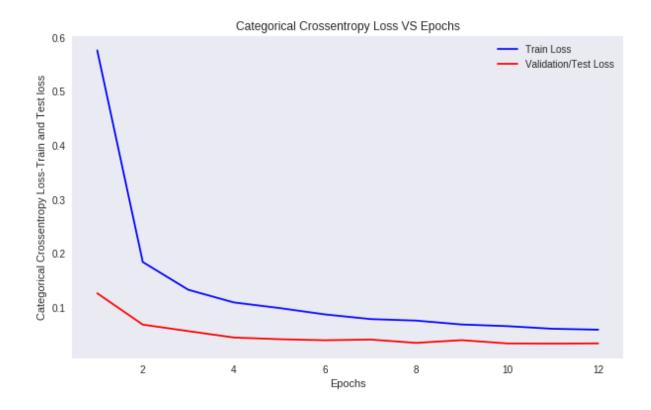
```
model 1.compile(optimizer='adam', loss='categorical crossentropy', metrics=['a
ccuracy'])
# Fitting the model
# To fit the model, give input data, batch_size, number of epochs and validati
on/test data
history = model 1.fit(X train, Y train, batch size=batch size, epochs=epochs,
verbose=1, validation_data=(X_test, Y_test))
Train on 60000 samples, validate on 10000 samples
Epoch 1/12
- acc: 0.8230 - val_loss: 0.1265 - val_acc: 0.9602
Epoch 2/12
- acc: 0.9416 - val loss: 0.0685 - val acc: 0.9784
Epoch 3/12
acc: 0.9582 - val_loss: 0.0565 - val_acc: 0.9813
Epoch 4/12
- acc: 0.9665 - val loss: 0.0447 - val acc: 0.9855
Epoch 5/12
60000/60000 [============== ] - 9s 158us/step - loss: 0.0992 -
acc: 0.9689 - val loss: 0.0416 - val acc: 0.9870
acc: 0.9729 - val loss: 0.0396 - val acc: 0.9873
Epoch 7/12
- acc: 0.9752 - val loss: 0.0410 - val acc: 0.9874
Epoch 8/12
acc: 0.9765 - val loss: 0.0348 - val acc: 0.9889
Epoch 9/12
- acc: 0.9784 - val loss: 0.0398 - val acc: 0.9872
Epoch 10/12
- acc: 0.9792 - val loss: 0.0338 - val acc: 0.9892
acc: 0.9804 - val loss: 0.0335 - val acc: 0.9897
Epoch 12/12
acc: 0.9808 - val loss: 0.0338 - val acc: 0.9886
```

2.3 Plotting the Loss vs Epoch Plot

```
In [0]:
        # Evaluating the model on test data
        score_1 = model_1.evaluate(X_test, Y_test, verbose=0)
        print('Test score:', score_1[0])
        print('Test accuracy:', score_1[1])
        # Plotting the results
        # list of epoch numbers
        x = list(range(1, epochs+1))
        # we will get val_loss and val_acc only when you pass the paramter validation_
        data
        # val_loss : validation loss
        vy = history.history['val_loss']
        # Training loss
        ty = history.history['loss']
        # calling the dynamic function to draw the plot
        plt_dynamic(x, vy, ty)
```

Test score: 0.03383973254601297

Test accuracy: 0.9886

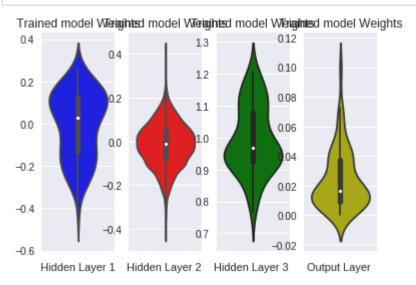


Observation:-

- 1. In this plot both Train and Test loss are significantly reducing/converging and there is no divergence / increasing between them .
- 2. This means model is not overfitting and it is working well.

2.4 Plotting Violin plots of hidden and output layers to see weights distribution

```
In [0]:
        import seaborn as sns
        w_after = model_1.get_weights()
        # weights of convolution layer 1
        c1_w = w_after[0].flatten().reshape(-1,1)
        # weights of convolution layer 2
        c2 w = w after[2].flatten().reshape(-1,1)
        # weights of convolution layer 3
        c3_w = w_after[4].flatten().reshape(-1,1)
        # weights of output layer
        out_w = w_after[6].flatten().reshape(-1,1)
        fig = plt.figure()
        plt.title("Weight matrices after model trained\n")
        # Hidden layer 1
        plt.subplot(1, 4, 1)
        plt.title("Trained model Weights")
        ax = sns.violinplot(y=c1 w,color='b')
        plt.xlabel('Hidden Layer 1')
        # Hidden Layer 2
        plt.subplot(1, 4, 2)
        plt.title("Trained model Weights")
        ax = sns.violinplot(y=c2 w, color='r')
        plt.xlabel('Hidden Layer 2 ')
        # Hidden Layer 3
        plt.subplot(1, 4, 3)
        plt.title("Trained model Weights")
        ax = sns.violinplot(y=c3 w, color='g')
        plt.xlabel('Hidden Layer 3 ')
        # Output Layer
        plt.subplot(1, 4, 4)
        plt.title("Trained model Weights")
        ax = sns.violinplot(y=out_w,color='y')
        plt.xlabel('Output Layer ')
        plt.show()
```



Observation:-

For all the layers, I got good gaussian structures and weights are nicely distributed . For some layers it is like combination of 2 gaussian structures.

3. CNN with 5 Convolution layers with Kernel size of 5X5

```
In [0]: | %matplotlib inline
        import warnings
        warnings.filterwarnings("ignore")
        # start building a model
        # Initializing the sequential model as the layers are in sequential
        model 2 = Sequential()
        # Adding first Convolution layer
        model 2.add(Conv2D(16, kernel size=(5, 5), activation='relu', padding='same',
        input shape=input shape))
        # Adding Maxpooling layer
        model 2.add(MaxPooling2D(pool size=(2, 2)))
        # Adding Dropout
        model_2.add(Dropout(0.25))
        # Adding second Convolution layer
        model_2.add(Conv2D(32, (5, 5), activation='relu', padding='same'))
        # Adding Batch Normalization
        model 2.add(BatchNormalization())
        # Adding Maxpooling Layer
        model 2.add(MaxPooling2D(pool size=(2, 2), strides=(1, 1)))
        # Adding Dropout
        model 2.add(Dropout(0.25))
        # Adding third Convolution layer
        model_2.add(Conv2D(32, (5, 5), activation='relu', padding='same'))
        # Adding Maxpooling Layer
        model 2.add(MaxPooling2D(pool size=(2, 2)))
        # Adding Dropout
        model_2.add(Dropout(0.25))
        # Adding fourth Convolution layer
        model_2.add(Conv2D(64, (5, 5), activation='relu', padding='same'))
        # Adding Batch Normalization
        model 2.add(BatchNormalization())
        # Adding Maxpooling layer
        model 2.add(MaxPooling2D(pool size=(2, 2)))
        # Adding Dropout
        model_2.add(Dropout(0.25))
        # Adding fifth Convolution layer
        model_2.add(Conv2D(64, (5, 5), activation='relu', padding='same'))
        # Adding Maxpooling Layer
        model_2.add(MaxPooling2D(pool_size=(2, 2), strides=(1, 1)))
        # Adding Dropout
        model 2.add(Dropout(0.25))
        # Adding flatten layer
        model 2.add(Flatten())
        # Adding a dense Layer
        model_2.add(Dense(128, activation='relu', kernel_initializer=he_normal(seed=No
        ne)))
        # Adding Batch Normalization
        model 2.add(BatchNormalization())
```

```
# Adding Dropout
model_2.add(Dropout(0.5))

# Output softmax Layer
model_2.add(Dense(num_classes, activation='softmax'))

# Printing model summary
model_2.summary()
```

Layer (type)	Output	Shape	Param #
conv2d_359 (Conv2D)	(None,	28, 28, 16)	416
max_pooling2d_328 (MaxPoolin	(None,	14, 14, 16)	0
dropout_341 (Dropout)	(None,	14, 14, 16)	0
conv2d_360 (Conv2D)	(None,	14, 14, 32)	12832
batch_normalization_181 (Bat	(None,	14, 14, 32)	128
max_pooling2d_329 (MaxPoolin	(None,	13, 13, 32)	0
dropout_342 (Dropout)	(None,	13, 13, 32)	0
conv2d_361 (Conv2D)	(None,	13, 13, 32)	25632
max_pooling2d_330 (MaxPoolin	(None,	6, 6, 32)	0
dropout_343 (Dropout)	(None,	6, 6, 32)	0
conv2d_362 (Conv2D)	(None,	6, 6, 64)	51264
batch_normalization_182 (Bat	(None,	6, 6, 64)	256
max_pooling2d_331 (MaxPoolin	(None,	3, 3, 64)	0
dropout_344 (Dropout)	(None,	3, 3, 64)	0
conv2d_363 (Conv2D)	(None,	3, 3, 64)	102464
max_pooling2d_332 (MaxPoolin	(None,	2, 2, 64)	0
dropout_345 (Dropout)	(None,	2, 2, 64)	0
flatten_32 (Flatten)	(None,	256)	0
dense_69 (Dense)	(None,	128)	32896
batch_normalization_183 (Bat	(None,	128)	512
dropout_346 (Dropout)	(None,	128)	0
dense_70 (Dense)	(None,	10)	1290

Total params: 227,690 Trainable params: 227,242 Non-trainable params: 448

3.1 Compiling the model

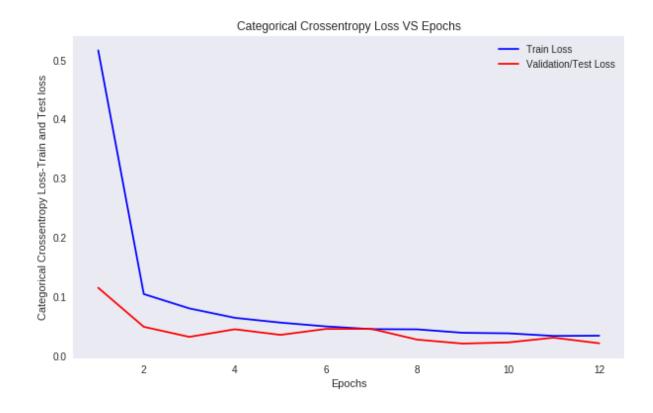
```
Train on 60000 samples, validate on 10000 samples
Epoch 1/12
- acc: 0.8433 - val_loss: 0.1152 - val_acc: 0.9660
- acc: 0.9700 - val_loss: 0.0493 - val_acc: 0.9841
Epoch 3/12
- acc: 0.9763 - val_loss: 0.0322 - val_acc: 0.9906
Epoch 4/12
60000/60000 [============ ] - 12s 205us/step - loss: 0.0644
- acc: 0.9819 - val loss: 0.0451 - val acc: 0.9868
Epoch 5/12
- acc: 0.9838 - val_loss: 0.0356 - val_acc: 0.9899
Epoch 6/12
- acc: 0.9860 - val loss: 0.0457 - val acc: 0.9876
Epoch 7/12
- acc: 0.9868 - val_loss: 0.0458 - val_acc: 0.9872
60000/60000 [=======================] - 12s 206us/step - loss: 0.0450
- acc: 0.9869 - val loss: 0.0276 - val acc: 0.9920
Epoch 9/12
60000/60000 [============= ] - 12s 207us/step - loss: 0.0392
- acc: 0.9891 - val loss: 0.0210 - val acc: 0.9941
Epoch 10/12
- acc: 0.9887 - val loss: 0.0229 - val acc: 0.9923
Epoch 11/12
- acc: 0.9899 - val_loss: 0.0309 - val_acc: 0.9913
Epoch 12/12
60000/60000 [============ ] - 12s 206us/step - loss: 0.0343
- acc: 0.9903 - val loss: 0.0214 - val acc: 0.9940
```

3.2 Plotting the Loss vs Epochs plot

```
In [0]:
        # Evaluating the model on test data
        score_2 = model_2.evaluate(X_test, Y_test, verbose=0)
        print('Test score:', score 2[0])
        print('Test accuracy:', score_2[1])
        # Plotting the results
        # list of epoch numbers
        x = list(range(1, epochs+1))
        # we will get val_loss and val_acc only when you pass the paramter validation_
        data
        # val_loss : validation loss
        vy = history.history['val_loss']
        # Training loss
        ty = history.history['loss']
        # calling the dynamic function to draw the plot
        plt_dynamic(x, vy, ty)
```

Test score: 0.02143788941586681

Test accuracy: 0.994



Observation:-

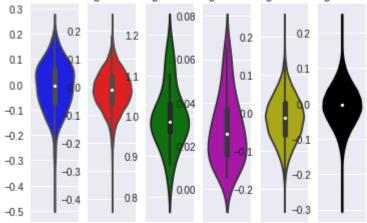
In this plot both Train and Test loss are significantly reducing/converging and there is no divergence / increasing between them . This means model is working well without overfitting .

3.3 Plotting Violin plots of hidden and output layers to see weights distribution

In [0]: import seaborn as sns w_after = model_2.get_weights() # weights of convolution layer 1 c1_w = w_after[0].flatten().reshape(-1,1) # weights of convolution layer 2 c2 w = w after[2].flatten().reshape(-1,1) # weights of convolution layer 3 c3_w = w_after[4].flatten().reshape(-1,1) # weights of convolution layer 4 c4_w = w_after[6].flatten().reshape(-1,1) # weights of convolution layer 5 c5 w = w after[8].flatten().reshape(-1,1) # weights of output layer out w = w after[10].flatten().reshape(-1,1) fig = plt.figure() plt.title("Weight matrices after model trained\n") # Hidden layer 1 plt.subplot(1, 6, 1) plt.title("Trained model Weights") ax = sns.violinplot(y=c1_w,color='b') plt.xlabel('Hidden Layer 1') # Hidden Layer 2 plt.subplot(1, 6, 2) plt.title("Trained model Weights") ax = sns.violinplot(y=c2_w, color='r') plt.xlabel('Hidden Layer 2 ') # Hidden Layer 3 plt.subplot(1, 6, 3) plt.title("Trained model Weights") ax = sns.violinplot(y=c3_w, color='g') plt.xlabel('Hidden Layer 3 ') # Hidden layer 4 plt.subplot(1, 6, 4) plt.title("Trained model Weights") ax = sns.violinplot(y=c4_w, color='m') plt.xlabel('Hidden Layer 4 ') # Hidden Layer 5 plt.subplot(1, 6, 5) plt.title("Trained model Weights") ax = sns.violinplot(y=c5 w, color='y') plt.xlabel('Hidden Layer 5 ') # Output Layer plt.subplot(1, 6, 6) plt.title("Trained model Weights") ax = sns.violinplot(y=out w,color='k')

plt.xlabel('Output Layer ')

Piłrainselig Wichelin wedigholiste in wedighol



Hidden Layelidden Laye

4. CNN with 7 Convolution layers with Kernel size of 3X3

```
In [0]: | %matplotlib inline
        import warnings
        warnings.filterwarnings("ignore")
        # start building a model
        # Initializing the sequential model as the layers are in sequential
        model 3 = Sequential()
        # Adding first Convolution layer
        model 3.add(Conv2D(16, kernel size=(3, 3), activation='relu', padding='same',
        input shape=input shape))
        # Adding second Convolution Layer
        model 3.add(Conv2D(16, (3, 3), activation='relu', padding='same'))
        # Adding Batch Normalization
        model 3.add(BatchNormalization())
        # Adding Maxpooling layer
        model_3.add(MaxPooling2D(pool_size=(2, 2), strides=(1,1)))
        # Adding Dropout
        model 3.add(Dropout(0.25))
        # Adding third Convolution layer
        model_3.add(Conv2D(32, (3, 3), activation='relu', padding='same'))
        # Adding fourth Convolution layer
        model 3.add(Conv2D(32, (3, 3), activation='relu', padding='same'))
        # Adding Batch Normalization
        model 3.add(BatchNormalization())
        # Adding Maxpooling Layer
        model 3.add(MaxPooling2D(pool size=(2, 2)))
        # Adding Dropout
        model 3.add(Dropout(0.25))
        # Adding fifth Convolution layer
        model_3.add(Conv2D(64, (3, 3), activation='relu', padding='same'))
        # Adding Maxpooling Layer
        model 3.add(MaxPooling2D(pool size=(2, 2)))
        # Adding Dropout
        model_3.add(Dropout(0.25))
        # Adding sixth Convolution Layer
        model 3.add(Conv2D(64, (3, 3), activation='relu', padding='same'))
        # Adding Batch Normalization
        model 3.add(BatchNormalization())
        # Adding Maxpooling Layer
        model 3.add(MaxPooling2D(pool size=(2, 2), strides=(1,1)))
        # Adding Dropout
        model 3.add(Dropout(0.25))
        # Adding seventh Convolution layer
        model_3.add(Conv2D(64, (3, 3), activation='relu', padding='same'))
        # Adding Maxpooling Layer
        model 3.add(MaxPooling2D(pool size=(2, 2), strides=(1,1)))
        # Adding Dropout
```

```
model_3.add(Dropout(0.25))
# Adding flatten layer
model_3.add(Flatten())
# Adding a dense layer
model_3.add(Dense(128, activation='relu', kernel_initializer=he_normal(seed=No
ne)))
# Adding Batch Normalization
model_3.add(BatchNormalization())
# Adding Dropout
model_3.add(Dropout(0.5))
# Adding a dense Layer
model_3.add(Dense(128, activation='relu', kernel_initializer=he_normal(seed=No
ne)))
# Adding Dropout
model_3.add(Dropout(0.5))
# Output softmax layer
model 3.add(Dense(num classes, activation='softmax'))
# Printing model summary
model_3.summary()
```

Lavon (type)	0	Chana	
Layer (type)	Output =====	Shape =========	Param # =======
conv2d_385 (Conv2D)	(None,	28, 28, 16)	160
conv2d_386 (Conv2D)	(None,	28, 28, 16)	2320
batch_normalization_190 (Bat	(None,	28, 28, 16)	64
max_pooling2d_345 (MaxPoolin	(None,	27, 27, 16)	0
dropout_364 (Dropout)	(None,	27, 27, 16)	0
conv2d_387 (Conv2D)	(None,	27, 27, 32)	4640
conv2d_388 (Conv2D)	(None,	27, 27, 32)	9248
batch_normalization_191 (Bat	(None,	27, 27, 32)	128
max_pooling2d_346 (MaxPoolin	(None,	13, 13, 32)	0
dropout_365 (Dropout)	(None,	13, 13, 32)	0
conv2d_389 (Conv2D)	(None,	13, 13, 64)	18496
max_pooling2d_347 (MaxPoolin	(None,	6, 6, 64)	0
dropout_366 (Dropout)	(None,	6, 6, 64)	0
conv2d_390 (Conv2D)	(None,	6, 6, 64)	36928
batch_normalization_192 (Bat	(None,	6, 6, 64)	256
max_pooling2d_348 (MaxPoolin	(None,	5, 5, 64)	0
dropout_367 (Dropout)	(None,	5, 5, 64)	0
conv2d_391 (Conv2D)	(None,	5, 5, 64)	36928
max_pooling2d_349 (MaxPoolin	(None,	4, 4, 64)	0
dropout_368 (Dropout)	(None,	4, 4, 64)	0
flatten_36 (Flatten)	(None,	1024)	0
dense_80 (Dense)	(None,	128)	131200
batch_normalization_193 (Bat	(None,	128)	512
dropout_369 (Dropout)	(None,	128)	0
dense_81 (Dense)	(None,	128)	16512
dropout_370 (Dropout)	(None,	128)	0
dense_82 (Dense)	(None,	10)	1290

Total params: 258,682 Trainable params: 258,202 Non-trainable params: 480

4.1 Compiling the model

```
In [0]: # Compiling the model
      model 3.compile(optimizer='adam', loss='categorical crossentropy', metrics=['a
      ccuracy'])
      # Fitting the model
      # To fit the model, give input data, batch size, number of epochs and validati
      on/test data
      history = model 3.fit(X train, Y train, batch size=batch size, epochs=epochs,
      verbose=1, validation_data=(X_test, Y_test))
      Train on 60000 samples, validate on 10000 samples
      Epoch 1/12
      - acc: 0.7261 - val_loss: 0.0885 - val_acc: 0.9715
      60000/60000 [================ ] - 20s 340us/step - loss: 0.1660
      - acc: 0.9536 - val loss: 0.0521 - val acc: 0.9836
      Epoch 3/12
      60000/60000 [========================] - 20s 340us/step - loss: 0.1141
      - acc: 0.9685 - val loss: 0.0511 - val acc: 0.9863
      Epoch 4/12
      60000/60000 [=======================] - 20s 340us/step - loss: 0.0871
      - acc: 0.9765 - val loss: 0.0383 - val acc: 0.9881
      Epoch 5/12
      - acc: 0.9806 - val loss: 0.0522 - val acc: 0.9830
      Epoch 6/12
      60000/60000 [========================] - 20s 340us/step - loss: 0.0652
      - acc: 0.9829 - val loss: 0.0376 - val acc: 0.9880
      Epoch 7/12
      - acc: 0.9844 - val_loss: 0.0501 - val_acc: 0.9839
      - acc: 0.9854 - val loss: 0.0340 - val acc: 0.9899
      Epoch 9/12
      - acc: 0.9868 - val loss: 0.0265 - val acc: 0.9934
      Epoch 10/12
      - acc: 0.9871 - val loss: 0.0216 - val acc: 0.9939
      Epoch 11/12
      60000/60000 [========================] - 20s 339us/step - loss: 0.0420
      - acc: 0.9889 - val_loss: 0.0234 - val_acc: 0.9937
      Epoch 12/12
      60000/60000 [============ ] - 20s 339us/step - loss: 0.0417
```

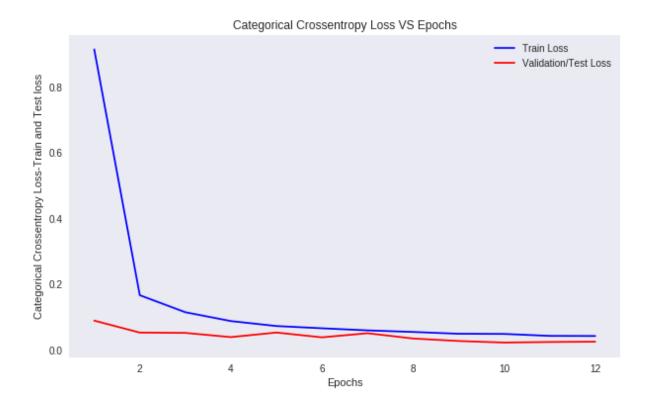
- acc: 0.9886 - val_loss: 0.0243 - val_acc: 0.9932

4.2 Plotting the Loss vs Epoch values

```
In [0]:
        # Evaluating the model on test data
        score_3 = model_3.evaluate(X_test, Y_test, verbose=0)
        print('Test score:', score_3[0])
        print('Test accuracy:', score 3[1])
        # Plotting the results
        # list of epoch numbers
        x = list(range(1, epochs+1))
        # we will get val_loss and val_acc only when you pass the paramter validation_
        data
        # val loss : validation loss
        vy = history.history['val_loss']
        # Training loss
        ty = history.history['loss']
        # calling the dynamic function to draw the plot
        plt_dynamic(x, vy, ty)
```

Test score: 0.02432433779832136

Test accuracy: 0.9932



Observation:-

In this plot both Train and Test loss are significantly reducing/converging and there is no divergence / increasing between them . This means model is working well without overfitting .

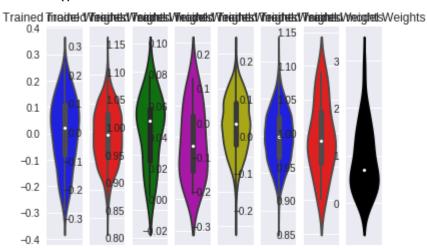
4.3 Plotting Violin plots of hidden and output layers to see weights distribution

```
In [130]:
          import seaborn as sns
          w_after = model_3.get_weights()
          # weights of convolution layer 1
          c1_w = w_after[0].flatten().reshape(-1,1)
          # weights of convolution layer 2
          c2 w = w after[2].flatten().reshape(-1,1)
          # weights of convolution layer 3
          c3_w = w_after[4].flatten().reshape(-1,1)
          # weights of convolution layer 4
          c4_w = w_after[6].flatten().reshape(-1,1)
          # weights of convolution layer 5
          c5 w = w after[8].flatten().reshape(-1,1)
          # weights of convolution layer 6
          c6_w = w_after[10].flatten().reshape(-1,1)
          # weights of convolution layer 7
          c7_w = w_after[12].flatten().reshape(-1,1)
          # weights of output layer
          out w = w after[14].flatten().reshape(-1,1)
          fig = plt.figure()
          plt.title("Weight matrices after model trained\n")
          # Hidden Layer 1
          plt.subplot(1, 8, 1)
          plt.title("Trained model Weights")
          ax = sns.violinplot(y=c1 w,color='b')
          plt.xlabel('Hidden Layer 1')
          # Hidden Layer 2
          plt.subplot(1, 8, 2)
          plt.title("Trained model Weights")
          ax = sns.violinplot(y=c2_w, color='r')
          plt.xlabel('Hidden Layer 2 ')
          # Hidden Layer 3
          plt.subplot(1, 8, 3)
          plt.title("Trained model Weights")
          ax = sns.violinplot(y=c3_w, color='g')
          plt.xlabel('Hidden Layer 3 ')
          # Hidden layer 4
          plt.subplot(1, 8, 4)
          plt.title("Trained model Weights")
          ax = sns.violinplot(y=c4_w, color='m')
          plt.xlabel('Hidden Layer 4 ')
          # Hidden Layer 5
          plt.subplot(1, 8, 5)
          plt.title("Trained model Weights")
          ax = sns.violinplot(y=c5_w, color='y')
          plt.xlabel('Hidden Layer 5 ')
          # Hidden Layer 6
```

```
plt.subplot(1, 8, 6)
plt.title("Trained model Weights")
ax = sns.violinplot(y=c6_w, color='b')
plt.xlabel('Hidden Layer 6 ')

# Hidden Layer 7
plt.subplot(1, 8, 7)
plt.title("Trained model Weights")
ax = sns.violinplot(y=c7_w, color='r')
plt.xlabel('Hidden Layer 7 ')

# Output Layer
plt.subplot(1, 8, 8)
plt.title("Trained model Weights")
ax = sns.violinplot(y=out_w,color='k')
plt.xlabel('Output Layer ')
plt.show()
```



Hidden U-biyateleti Lletiyateleti Lletiyatel

5. Models Summarization

In [132]: Final_conclusions = DataFrame(CNN)
Final_conclusions

Out[132]:

	Activation	CNN Model	Kernel size	Kernel_initializer	Optimizer	Test accuracy	Test loss	Train loss	Training accuracy
0	ReLU	3- Conv layers	2X2	he_normal	Adam	0.988	0.03	0.05	0.980
1	ReLU	5- Conv layers	5X5	he_normal	Adam	0.994	0.02	0.03	0.990
2	ReLU	7- Conv layers	3X3	he_normal	Adam	0.99	0.02	0.04	0.98

6. Conclusions:-

From the above observations we can observe,

- 1. As I had used ReLU activation and Adam optimizer, all the accuracies are good.
- 2. Output layer is softmax layer.
- 3. As I added dropout, Batch normalization, max-pooling, padding, all the accuracies are good.
- 4. Test accuracy is little more than Train accuracy in all models, so the model is working well.
- 5. Test loss is less than train loss for every model.
- 6. In the Categorical Crossentropy Loss VS Epochs plot, train and test loss also converged or decreased gradually, there is no overfitting.
- 7. I have used he_normal as kernel initializer as it have inbuilt mean and std values, without need of defing them explicitly.