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CSE3B

In [1]:

```
# example of loading the fashion mnist dataset
from matplotlib import pyplot
from keras.datasets import fashion mnist
# load dataset
(trainX, trainy), (testX, testy) = fashion mnist.load data()
# summarize loaded dataset
print('Train: X=%s, y=%s' % (trainX.shape, trainy.shape))
print('Test: X=%s, y=%s' % (testX.shape, testy.shape))
# plot first few images
for i in range(9):
 # define subplot
pyplot.subplot(330 + 1 + i)
 # plot raw pixel data
pyplot.imshow(trainX[i], cmap=pyplot.get cmap('gray'))
# show the figure
pyplot.show()
Downloading data from https://storage.googleapis.com/tensorflow/tf-keras-datasets/train-l
abels-idx1-ubyte.gz
Downloading data from https://storage.googleapis.com/tensorflow/tf-keras-datasets/train-i
mages-idx3-ubyte.gz
26427392/26421880 [============ ] - 0s Ous/step
Downloading data from https://storage.googleapis.com/tensorflow/tf-keras-datasets/t10k-la
bels-idx1-ubyte.gz
8192/5148 [=======] - Os Ous/step
Downloading data from https://storage.googleapis.com/tensorflow/tf-keras-datasets/t10k-im
ages-idx3-ubyte.gz
4423680/4422102 [============ ] - Os Ous/step
Train: X=(60000, 28, 28), y=(60000,)
Test: X=(10000, 28, 28), y=(10000,)
 0
10
             10
                          10
20
 0
10
             10
 0
                          10
10
                          20
20
```

In [13]:

```
# baseline cnn model for fashion mnist
from numpy import mean
from numpy import std
from matplotlib import pyplot
from sklearn.model_selection import KFold
from keras.datasets import fashion_mnist
from keras.utils import to_categorical
from keras.models import Sequential
from keras.layers import Conv2D
```

```
from keras.layers import MaxPooling2D
from keras.layers import Dense
from keras.layers import Flatten
from keras.optimizers import SGD
# load train and test dataset
def load dataset():
 # load dataset
 (trainX, trainY), (testX, testY) = fashion mnist.load data()
 # reshape dataset to have a single channel
 trainX = trainX.reshape((trainX.shape[0], 28, 28, 1))
 testX = testX.reshape((testX.shape[0], 28, 28, 1))
 # one hot encode target values
 trainY = to categorical(trainY)
 testY = to categorical(testY)
 return trainX, trainY, testX, testY
# scale pixels
def prep_pixels(train, test):
# convert from integers to floats
train norm = train.astype('float32')
test_norm = test.astype('float32')
 # normalize to range 0-1
train norm = train norm / 255.0
test norm = test norm / 255.0
 # return normalized images
return train norm, test norm
# define cnn model
def define model():
model = Sequential()
model.add(Conv2D(32, (3, 3), activation='relu', kernel initializer='he uniform', input
shape=(28, 28, 1))
model.add(MaxPooling2D((2, 2)))
model.add(Flatten())
model.add(Dense(100, activation='relu', kernel initializer='he uniform'))
model.add(Dense(10, activation='softmax'))
 # compile model
opt = SGD(1r=0.01, momentum=0.9)
model.compile(optimizer=opt, loss='categorical crossentropy', metrics=['accuracy'])
return model
# evaluate a model using k-fold cross-validation
def evaluate model(dataX, dataY, n folds=5):
scores, histories = list(), list()
 # prepare cross validation
kfold = KFold(n folds, shuffle=True, random state=1)
 # enumerate splits
 for train ix, test ix in kfold.split(dataX):
  # define model
 model = define_model()
  # select rows for train and test
  trainX, trainY, testX, testY = dataX[train ix], dataY[train ix], dataX[test ix], dataY
[test ix]
  # fit model
 history = model.fit(trainX, trainY, epochs=10, batch size=32, validation data=(testX,
testY), verbose=0)
 # evaluate model
  , acc = model.evaluate(testX, testY, verbose=0)
 print('> %.3f' % (acc * 100.0))
 # append scores
 scores.append(acc)
 histories.append(history)
return scores, histories
# plot diagnostic learning curves
def summarize diagnostics(histories):
for i in range(len(histories)):
  # plot loss
  pyplot.subplot(211)
  pyplot.title('Cross Entropy Loss')
 pyplot.plot(histories[i].history['loss'], color='blue', label='train')
```

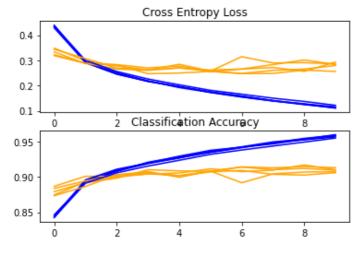
```
pyplot.plot(histories[i].history['val loss'], color='orange', label='test')
  # plot accuracy
 pyplot.subplot(212)
 pyplot.title('Classification Accuracy')
 pyplot.plot(histories[i].history['accuracy'], color='blue', label='train')
 pyplot.plot(histories[i].history['val accuracy'], color='orange', label='test')
pyplot.show()
# summarize model performance
def summarize performance(scores):
 # print summary
print('Accuracy: mean=%.3f std=%.3f, n=%d' % (mean(scores)*100, std(scores)*100, len(sco
res)))
 # box and whisker plots of results
pyplot.boxplot(scores)
pyplot.show()
# run the test harness for evaluating a model
def run test harness():
# load dataset
trainX, trainY, testX, testY = load dataset()
# prepare pixel data
trainX, testX = prep_pixels(trainX, testX)
 # evaluate model
scores, histories = evaluate model(trainX, trainY)
# learning curves
summarize diagnostics(histories)
# summarize estimated performance
summarize performance(scores)
# entry point, run the test harness
run test harness()
```

```
> 90.608
```

> 90.758

/usr/local/lib/python3.7/dist-packages/ipykernel_launcher.py:76: MatplotlibDeprecationWar ning: Adding an axes using the same arguments as a previous axes currently reuses the ear lier instance. In a future version, a new instance will always be created and returned. Meanwhile, this warning can be suppressed, and the future behavior ensured, by passing a unique label to each axes instance.

/usr/local/lib/python3.7/dist-packages/ipykernel_launcher.py:81: MatplotlibDeprecationWar ning: Adding an axes using the same arguments as a previous axes currently reuses the ear lier instance. In a future version, a new instance will always be created and returned. Meanwhile, this warning can be suppressed, and the future behavior ensured, by passing a unique label to each axes instance.



Accuracy: mean=90.933 std=0.239, n=5



> 91.308

> 91.017

> 90.975

In [14]:

```
# example of loading the mnist dataset
from keras.datasets import mnist
from matplotlib import pyplot
# load dataset
(trainX, trainy), (testX, testy) = mnist.load data()
# summarize loaded dataset
print('Train: X=%s, y=%s' % (trainX.shape, trainy.shape))
print('Test: X=%s, y=%s' % (testX.shape, testy.shape))
# plot first few images
for i in range(9):
 # define subplot
pyplot.subplot(330 + 1 + i)
# plot raw pixel data
pyplot.imshow(trainX[i], cmap=pyplot.get cmap('gray'))
# show the figure
pyplot.show()
```

Train: X=(60000, 28, 28), y=(60000,)Test: X=(10000, 28, 28), y=(10000,)

In [15]:

```
# baseline cnn model for mnist
from numpy import mean
from numpy import std
from matplotlib import pyplot
from sklearn.model selection import KFold
from keras.datasets import mnist
from keras.utils import to categorical
from keras.models import Sequential
from keras.layers import Conv2D
from keras.layers import MaxPooling2D
from keras.layers import Dense
from keras.layers import Flatten
from keras.optimizers import SGD
# load train and test dataset
def load dataset():
 # load dataset
```

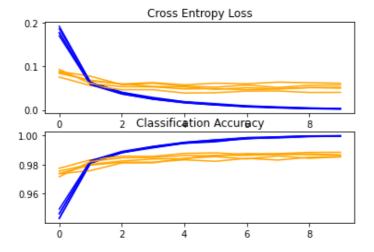
```
(trainX, trainY), (testX, testY) = mnist.load_data()
 # reshape dataset to have a single channel
 trainX = trainX.reshape((trainX.shape[0], 28, 28, 1))
testX = testX.reshape((testX.shape[0], 28, 28, 1))
 # one hot encode target values
trainY = to categorical(trainY)
 testY = to categorical(testY)
return trainX, trainY, testX, testY
# scale pixels
def prep pixels(train, test):
 # convert from integers to floats
 train_norm = train.astype('float32')
test norm = test.astype('float32')
 # normalize to range 0-1
 train norm = train norm / 255.0
 test norm = test norm / 255.0
 # return normalized images
return train norm, test norm
# define cnn model
def define model():
model = Sequential()
model.add(Conv2D(32, (3, 3), activation='relu', kernel initializer='he uniform', input
shape=(28, 28, 1))
model.add(MaxPooling2D((2, 2)))
model.add(Flatten())
model.add(Dense(100, activation='relu', kernel initializer='he uniform'))
model.add(Dense(10, activation='softmax'))
# compile model
opt = SGD(lr=0.01, momentum=0.9)
model.compile(optimizer=opt, loss='categorical crossentropy', metrics=['accuracy'])
 return model
# evaluate a model using k-fold cross-validation
def evaluate model(dataX, dataY, n folds=5):
scores, histories = list(), list()
 # prepare cross validation
kfold = KFold(n_folds, shuffle=True, random_state=1)
 # enumerate splits
for train ix, test ix in kfold.split(dataX):
  # define model
 model = define model()
  # select rows for train and test
 trainX, trainY, testX, testY = dataX[train ix], dataY[train ix], dataX[test ix], dataY
[test ix]
  # fit model
 history = model.fit(trainX, trainY, epochs=10, batch size=32, validation data=(testX,
testY), verbose=0)
  # evaluate model
  _, acc = model.evaluate(testX, testY, verbose=0)
  print('> %.3f' % (acc * 100.0))
 # stores scores
 scores.append(acc)
 histories.append(history)
return scores, histories
# plot diagnostic learning curves
def summarize diagnostics(histories):
for i in range(len(histories)):
  # plot loss
 pyplot.subplot(2, 1, 1)
 pyplot.title('Cross Entropy Loss')
 pyplot.plot(histories[i].history['loss'], color='blue', label='train')
 pyplot.plot(histories[i].history['val loss'], color='orange', label='test')
 # plot accuracy
 pyplot.subplot(2, 1, 2)
  pyplot.title('Classification Accuracy')
 pyplot.plot(histories[i].history['accuracy'], color='blue', label='train')
 pyplot.plot(histories[i].history['val accuracy'], color='orange', label='test')
pyplot.show()
```

```
# summarize model performance
def summarize performance(scores):
# print summary
print('Accuracy: mean=%.3f std=%.3f, n=%d' % (mean(scores)*100, std(scores)*100, len(sco
res)))
# box and whisker plots of results
pyplot.boxplot(scores)
pyplot.show()
# run the test harness for evaluating a model
def run test harness():
# load dataset
trainX, trainY, testX, testY = load dataset()
 # prepare pixel data
trainX, testX = prep_pixels(trainX, testX)
 # evaluate model
scores, histories = evaluate model(trainX, trainY)
# learning curves
summarize diagnostics(histories)
 # summarize estimated performance
summarize performance (scores)
# entry point, run the test harness
run test harness()
```

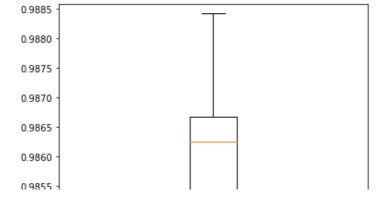
```
> 98.533
```

/usr/local/lib/python3.7/dist-packages/ipykernel_launcher.py:76: MatplotlibDeprecationWar ning: Adding an axes using the same arguments as a previous axes currently reuses the ear lier instance. In a future version, a new instance will always be created and returned. Meanwhile, this warning can be suppressed, and the future behavior ensured, by passing a unique label to each axes instance.

/usr/local/lib/python3.7/dist-packages/ipykernel_launcher.py:81: MatplotlibDeprecationWar ning: Adding an axes using the same arguments as a previous axes currently reuses the ear lier instance. In a future version, a new instance will always be created and returned. Meanwhile, this warning can be suppressed, and the future behavior ensured, by passing a unique label to each axes instance.



Accuracy: mean=98.640 std=0.113, n=5



> 98.667

> 98.533

> 98.842

> 98.625

i