Problem 1: Time Series Forecasting (20 points) In this problem, we want to create time series model to predict monthly car sales for a company. Download the zip file for homework assignment #5, and use the CarSales dataset which is a standard univariate time series dataset consists of 108 months of car sales in Quebec 1960-1968. The first column is the date and the second is the number of sales.

```
In [1]:
         #Import necessary packages to the Jupyter notebook
         import pandas as pd
         from pandas import read_csv
         from matplotlib import pyplot
         from pandas.plotting import autocorrelation plot
         from statsmodels.tsa.stattools import adfuller
         from random import randrange
         from statsmodels.tsa.seasonal import seasonal_decompose
         from sklearn.linear model import LinearRegression
         import numpy as np
         from pandas import DataFrame
         from statsmodels.tsa.arima_model import ARIMA
         from sklearn.metrics import mean squared error
         from math import sqrt
         filename ='Dataset5/CarSales.csv'
```

(a) Load time series data: Use read csv() function to load your time series datasets as a Series object, instead of DataFrame. Use the following arguments to the read csv() function to ensure the data is loaded as a Series.

```
pd1=pd.read_csv(filename,header=0,parse_dates=True,index_col=0,squeeze=True)
```

(b) Exploring time series data: Use the head() function to peek at the first 10 records of your data.

```
In [3]:
    pd3=pd.read_csv(filename,parse_dates=True)
    pd3.head(10)
```

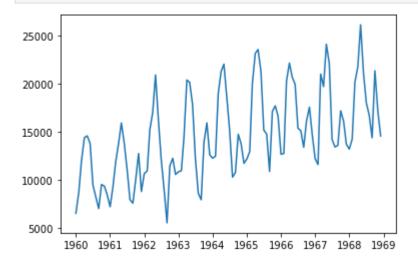
Out[3]:		Month	Sales
	0	1960-01	6550
	1	1960-02	8728
	2	1960-03	12026
	3	1960-04	14395
	4	1960-05	14587
	5	1960-06	13791
	6	1960-07	9498
	7	1960-08	8251
	8	1960-09	7049
	9	1960-10	9545

```
In [4]: pd3.columns
```

Out[4]: Index(['Month', 'Sales'], dtype='object')

(c) Line plot: Use the plotting functions(.plot() and .show()) from Matplotlib to visualize your Series of the monthly car sales dataset as a line plot.

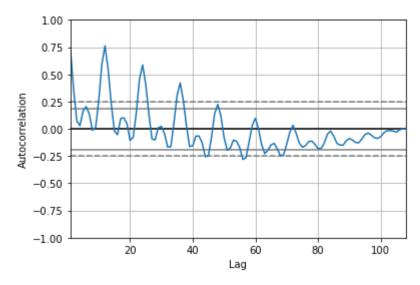
```
import matplotlib.pyplot as plt
plt.plot(pd1)
plt.show()
```



(d) Autocorrelation plot: Use Pandas plotting function autocorrelation plot() to create an autocorrelation plot for your series.

```
In [6]: autocorrelation_plot(pd1)
```

Out[6]: <AxesSubplot:xlabel='Lag', ylabel='Autocorrelation'>



(e) Stationarity in time series data: Use the adfuller() function from the Statsmodels library to perform Dickey-Fuller test to check if your time series is stationary or non-stationary. Interpret the results of the test.

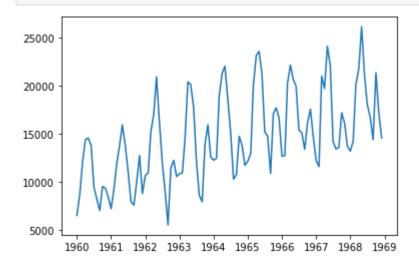
```
# Augmented Dickey Fuller test
          adfuller(pd1)
        (-1.223812766175286,
Out[7]:
          0.663269104983286,
          12,
          95,
          {'1%': -3.5011373281819504,
           '5%': -2.8924800524857854,
           '10%': -2.5832749307479226},
          1671.1995896872572)
        We see the p-value is greater then 0.05 so the series is non stationary
        (f) Automatic time series decomposition: Use seasonal decompose() function from Statsmodels
        library. Specify your model as 'additive' and use the .plot() function to visualize the four resulting
        series. Interpret the results (Do you observe any trend; if yes, what kind of trend; Is there any
        seasonality in your data).
In [8]:
         X=seasonal_decompose(pd1,period=4)
         print(X.seasonal)
         Month
         1960-01-01 -513.270433
         1960-02-01 264.806490
         1960-03-01
                     505.965144
         1960-04-01 -257.501202
         1960-05-01 -513.270433
                           . . .
         1968-08-01 -257.501202
         1968-09-01 -513.270433
                     264.806490
         1968-10-01
         1968-11-01
                       505.965144
                     -257.501202
         1968-12-01
         Name: seasonal, Length: 108, dtype: float64
In [9]:
         plt.plot(X.seasonal)
         plt.show()
          400
          200
            0
```

```
In [10]: plt.plot(X.observed)
    plt.show()
```

1960 1961 1962 1963 1964 1965 1966 1967 1968

-200

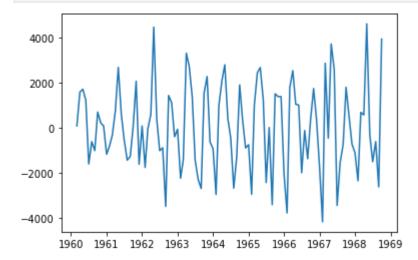
-400



In [11]: plt.plot(X.trend)
 plt.show()

22000 -18000 -16000 -14000 -12000 -1960 1961 1962 1963 1964 1965 1966 1967 1968 1969

In [12]: plt.plot(X.resid)
 plt.show()



(g) Detrend by model fitting: Use a linear model to detrend your time series data:

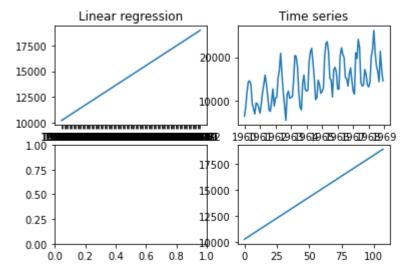
- i. Use the scikit-learn LinearRegression model to fit a linear model on your data.
- ii. Use .predict() function to calculate the trend.
- iii. Use .plot() and .show() functions to visualtize the trend and the series data on the same plot.
- iv. Deterend your sereis by subtracting the trend values from the original values of the series, and plot the resulting detrented series in a separate plot.

Out[13]: 0.3158854329342885

```
In [14]:
    Y2=reg.predict(np.array(X).reshape(-1,1))
    import matplotlib.pyplot as plt

figure, axis = plt.subplots(2, 2)
    axis[0, 0].plot(X1,Y2)
    axis[0, 0].set_title("Linear regression")

# For Cosine Function
    axis[0, 1].plot(pd1)
    axis[0, 1].set_title("Time series")
    plt.plot(Y2)
    plt.show()
```



ARIMA with Python: To answer the questions for this part, you can take a look at the code from Lab Session 11. (5 points)

- i. Extract the NumPy array of data values and split your data into train and test with a split of 70-30.
- ii. Use the forecast() function to perform a one-step forecast using the model. Use the train set to fit the model, and generate a prediction for each element on the test set.
- iii. Perform a rolling forecast by keeping track of all observations in a list called history that is seeded with the training data and to which new observations are appended each iteration; Print the prediction and expected value each iteration. To define your ARIMA model for this part use the ARIMA function from Statsmodels library, and pass in the parameters p=5, d=1, q=1.
- iv. Calculate a final root mean squared error score (RMSE) for the predictions.
- v. Create a line plot to show the expected values (blue) compared to the rolling forecast predictions (red).

```
In [15]: import warnings

In [28]: # split into train and test sets
    size = int(len(pd3) * 0.66)
    train, test = X[0:size], X[size:len(X)]
    history = [x for x in train]

In [27]: pd1.dropna(inplace=True)
```

model = ARIMA(history, order=(5,1,1)) # model_fit = model.fit() # output = model_fit.forecast() # prepare training dataset train_size = int(len(X) * 0.50) train, test = X[0:train_size], X[train_size:] history = [x for x in train]

```
In [29]: # make prdictions
predictions = list()
# walk-forward validation
for t in range(len(test)):
    model = ARIMA(history, order=(5,1,1))
    model_fit = model.fit()
    output = model_fit.forecast()
    yhat = output[0]
    predictions.append(yhat)
    obs = test[t]
    history.append(obs)
    print('predicted=%f, expected=%f' % (yhat, obs))
```

```
LinAlgError Traceback (most recent call last)

C:\Users\SUMAMA~1\AppData\Local\Temp/ipykernel_59700/4132154632.py in <module>

4 for t in range(len(test)):

5 model = ARIMA(history, order=(5,1,1))

----> 6 model_fit = model.fit()

7 output = model_fit.forecast()

8 yhat = output[0]

~\anaconda3\lib\site-packages\statsmodels\tsa\arima_model.py in fit(self, start_params, trend, method, transparams, solver, maxiter, full_output, disp, callback, start_ar_lag s, **kwargs)
```

```
r, order = 'F')
  1226
   1227
-> 1228
                mlefit = super(ARIMA, self).fit(start_params, trend,
   1229
                                                method, transparams, solver,
   1230
                                                maxiter, full output, disp,
~\anaconda3\lib\site-packages\statsmodels\tsa\arima model.py in fit(self, start params,
trend, method, transparams, solver, maxiter, full_output, disp, callback, start_ar_lag
s, **kwargs)
   1025
                    kwargs.setdefault('m', 12)
  1026
                    kwargs.setdefault('approx_grad', True)
-> 1027
                mlefit = super(ARMA, self).fit(start_params, method=solver,
  1028
                                               maxiter=maxiter,
   1029
                                               full_output=full_output, disp=disp,
~\anaconda3\lib\site-packages\statsmodels\base\model.py in fit(self, start_params, metho
d, maxiter, full_output, disp, fargs, callback, retall, skip_hessian, **kwargs)
                warn_convergence = kwargs.pop('warn_convergence', True)
    517
    518
                optimizer = Optimizer()
                xopt, retvals, optim settings = optimizer. fit(f, score, start params,
--> 519
    520
                                                                fargs, kwargs,
    521
                                                                hessian=hess,
~\anaconda3\lib\site-packages\statsmodels\base\optimizer.py in fit(self, objective, gra
dient, start params, fargs, kwargs, hessian, method, maxiter, full output, disp, callbac
k, retall)
    222
    223
                func = fit funcs[method]
--> 224
                xopt, retvals = func(objective, gradient, start params, fargs, kwargs,
    225
                                    disp=disp, maxiter=maxiter, callback=callback,
    226
                                    retall=retall, full output=full output,
~\anaconda3\lib\site-packages\statsmodels\base\optimizer.py in fit lbfgs(f, score, star
t params, fargs, kwargs, disp, maxiter, callback, retall, full output, hess)
    627
                func = f
    628
--> 629
            retvals = optimize.fmin_1_bfgs_b(func, start_params, maxiter=maxiter,
    630
                                             callback=callback, args=fargs,
    631
                                             bounds=bounds, disp=disp,
~\anaconda3\lib\site-packages\scipy\optimize\lbfgsb.py in fmin 1 bfgs b(func, x0, fprim
e, args, approx_grad, bounds, m, factr, pgtol, epsilon, iprint, maxfun, maxiter, disp, c
allback, maxls)
                    'maxls': maxls}
    195
    196
--> 197
            res = _minimize_lbfgsb(fun, x0, args=args, jac=jac, bounds=bounds,
                                   **opts)
    198
    199
            d = {'grad': res['jac'],
~\anaconda3\lib\site-packages\scipy\optimize\lbfgsb.py in _minimize_lbfgsb(fun, x0, arg
s, jac, bounds, disp, maxcor, ftol, gtol, eps, maxfun, maxiter, iprint, callback, maxls,
finite diff rel step, **unknown options)
    304
                    iprint = disp
    305
--> 306
           sf = _prepare_scalar_function(fun, x0, jac=jac, args=args, epsilon=eps,
    307
                                          bounds=new bounds,
                                          finite diff rel step=finite diff rel step)
    308
~\anaconda3\lib\site-packages\scipy\optimize\optimize.py in prepare scalar function(fu
n, x0, jac, args, bounds, epsilon, finite_diff_rel_step, hess)
```

```
# ScalarFunction caches. Reuse of fun(x) during grad
    259
    260
            # calculation reduces overall function evaluations.
--> 261
            sf = ScalarFunction(fun, x0, args, grad, hess,
    262
                                finite_diff_rel_step, bounds, epsilon=epsilon)
    263
~\anaconda3\lib\site-packages\scipy\optimize\_differentiable_functions.py in __init__(se
lf, fun, x0, args, grad, hess, finite_diff_rel_step, finite_diff_bounds, epsilon)
    138
    139
                self. update fun impl = update fun
                self._update_fun()
--> 140
    141
    142
                # Gradient evaluation
~\anaconda3\lib\site-packages\scipy\optimize\ differentiable functions.py in update fun
(self)
            def update fun(self):
    231
    232
                if not self.f_updated:
--> 233
                    self. update fun impl()
                    self.f updated = True
    234
    235
~\anaconda3\lib\site-packages\scipy\optimize\ differentiable functions.py in update fun
()
    135
    136
                def update_fun():
--> 137
                    self.f = fun_wrapped(self.x)
    138
                self. update fun impl = update fun
    139
~\anaconda3\lib\site-packages\scipy\optimize\ differentiable functions.py in fun wrapped
(x)
    132
                    # Overwriting results in undefined behaviour because
                    # fun(self.x) will change self.x, with the two no longer linked.
    133
--> 134
                    return fun(np.copy(x), *args)
    135
    136
                def update fun():
~\anaconda3\lib\site-packages\statsmodels\base\model.py in f(params, *args)
    499
    500
                def f(params, *args):
--> 501
                    return -self.loglike(params, *args) / nobs
    502
    503
                if method == 'newton':
~\anaconda3\lib\site-packages\statsmodels\tsa\arima_model.py in loglike(self, params, se
t_sigma2)
    839
                method = self.method
    840
                if method in ['mle', 'css-mle']:
--> 841
                    return self.loglike_kalman(params, set_sigma2)
    842
                elif method == 'css':
                    return self.loglike css(params, set sigma2)
    843
~\anaconda3\lib\site-packages\statsmodels\tsa\arima model.py in loglike kalman(self, par
ams, set_sigma2)
    849
                Compute exact loglikelihood for ARMA(p,q) model by the Kalman Filter.
    850
--> 851
                return KalmanFilter.loglike(params, self, set_sigma2)
    852
            def loglike_css(self, params, set_sigma2=True):
    853
```

```
~\anaconda3\lib\site-packages\statsmodels\tsa\kalmanf\kalmanfilter.py in loglike(cls, pa
         rams, arma_model, set_sigma2)
                          paramsdtype) = cls._init_kalman_state(params, arma_model)
             216
             217
                         if np.issubdtype(paramsdtype, np.float64):
         --> 218
                             loglike, sigma2 = kalman_loglike.kalman_loglike_double(
             219
                                 y, k, k_ar, k_ma, k_lags, int(nobs),
             220
                                 Z_mat, R_mat, T_mat)
         statsmodels\tsa\kalmanf\kalman loglike.pyx in statsmodels.tsa.kalmanf.kalman loglike.kal
         man loglike double()
         statsmodels\tsa\kalmanf\kalman_loglike.pyx in statsmodels.tsa.kalmanf.kalman_loglike.kal
         man_filter_double()
         <_array_function__ internals> in pinv(*args, **kwargs)
         ~\anaconda3\lib\site-packages\numpy\linalg\linalg.py in pinv(a, rcond, hermitian)
            2000
                         return wrap(res)
            2001
                   a = a.conjugate()
         -> 2002
                   u, s, vt = svd(a, full_matrices=False, hermitian=hermitian)
            2003
            2004
                     # discard small singular values
         <__array_function__ internals> in svd(*args, **kwargs)
         ~\anaconda3\lib\site-packages\numpy\linalg\linalg.py in svd(a, full_matrices, compute_u
         v, hermitian)
            1658
            1659
                        signature = 'D->DdD' if isComplexType(t) else 'd->ddd'
         -> 1660
                         u, s, vh = gufunc(a, signature=signature, extobj=extobj)
            1661
                        u = u.astype(result_t, copy=False)
            1662
                         s = s.astype( realType(result t), copy=False)
         ~\anaconda3\lib\site-packages\numpy\linalg\linalg.py in _raise_linalgerror_svd_nonconver
         gence(err, flag)
              95
              96 def raise linalgerror svd nonconvergence(err, flag):
                   raise LinAlgError("SVD did not converge")
              98
              99 def _raise_linalgerror_lstsq(err, flag):
         LinAlgError: SVD did not converge
In [30]:
          import math
          from sklearn.metrics import mean_squared_error
          # evaluate forecasts
          rmse = math.sqrt(mean squared error(test, predictions))
          print("Test RMSE: {}".format(rmse))
          # plot forecasts against actual outcomes
          plt.plot(test)
          plt.plot(predictions, color='red')
          plt.show()
         ValueError
                                                   Traceback (most recent call last)
         C:\Users\SUMAMA~1\AppData\Local\Temp/ipykernel 59700/1095995198.py in <module>
```

3

```
4 # evaluate forecasts
        ----> 5 rmse = math.sqrt(mean squared error(test, predictions))
              6 print("Test RMSE: {}".format(rmse))
              7 # plot forecasts against actual outcomes
        ~\anaconda3\lib\site-packages\sklearn\metrics\_regression.py in mean_squared_error(y_tru
        e, y pred, sample weight, multioutput, squared)
            436
                    0.825...
                    0.00
            437
        --> 438
                    y type, y true, y pred, multioutput = check reg targets(
                        y_true, y_pred, multioutput
            439
            440
        ~\anaconda3\lib\site-packages\sklearn\metrics\_regression.py in _check_reg_targets(y_tru
        e, y_pred, multioutput, dtype)
             92
                        the dtype argument passed to check_array.
             93
        ---> 94
                    check_consistent_length(y_true, y_pred)
             95
                    y_true = check_array(y_true, ensure_2d=False, dtype=dtype)
                    y_pred = check_array(y_pred, ensure_2d=False, dtype=dtype)
        ~\anaconda3\lib\site-packages\sklearn\utils\validation.py in check_consistent_length(*ar
        rays)
            330
                    uniques = np.unique(lengths)
            331
                    if len(uniques) > 1:
        --> 332
                        raise ValueError(
            333
                            "Found input variables with inconsistent numbers of samples: %r"
            334
                            % [int(l) for l in lengths]
        ValueError: Found input variables with inconsistent numbers of samples: [37, 0]
In [ ]:
In [ ]:
```

Problem 2: Multilayer Perceptron for Binary Classification (15 points) In this problem you will create a classification neural network in Keras using the KerasClassifier wrapper. For this problem, from the zip file you downloaded for homework assignment #5, use the pima-indians-diabetes dataset. The objective of the dataset is to diagnostically predict whether or not a patient has diabetes, based on certain diagnostic measurements included in the dataset. Several constraints were placed on the selection of these instances from a larger database. In particular, all patients here are females at least 21 years old of Pima Indian heritage. The datasets consists of several medical predictor variables and one target variable. Predictor variables includes the number of pregnancies the patient has had, their BMI, insulin level, age, and so on.

!pip install tensorflow

```
import numpy as np
import tensorflow as tf
import random as python_random
# fix random seed for reproducibility
def reset_seeds ():
    np.random.seed ( 123 )
    python_random.seed ( 123 )
```

```
tf.random.set seed ( 1234 )
    reset_seeds ()
from numpy import loadtxt
from keras.models import Sequential
from keras.layers import Dense,InputLayer
from keras.wrappers. scikit_learn import KerasClassifier
from sklearn.model selection import KFold
from sklearn.model_selection import cross_val_score
from sklearn.model_selection import GridSearchCV
import numpy
from pandas import read csv
# Load the dataset
filename = "Dataset5/pima-indians-diabetes.csv"
names = ['preg', 'plas', 'pres', 'skin', 'test', 'mass', 'pedi', 'age', 'class']
dataset = read_csv ( filename , names = names )
# split into input (X) and output (y) variables
array = dataset . values
X = array [:,0:8]
y = array [:,8]
```

(a) Define your Keras model

```
In [32]:
          def create model():
              model = tf.keras.models.Sequential()
              model.add(tf.keras.layers.Dense(12,input_dim=8 ,activation='relu'))
          # Now the model will take as input arrays of shape (None, 16)
          # and output arrays of shape (None, 32).
          # Note that after the first layer, you don't need to specify
          # the size of the input anymore:
              model.add(tf.keras.layers.Dense(8,activation='relu'))
              model.add(tf.keras.layers.Dense(1, activation='sigmoid'))
              model.compile(loss='binary_crossentropy',optimizer='adam',metrics=['accuracy'])
              return model
          #m1=create model(X,y)
          m2=KerasClassifier(build fn=create model,epochs=10,batch size=16,verbose=0)
         C:\Users\SUMAMA~1\AppData\Local\Temp/ipykernel_59700/291242446.py:14: DeprecationWarnin
         g: KerasClassifier is deprecated, use Sci-Keras (https://github.com/adriangb/scikeras) i
         nstead. See https://www.adriangb.com/scikeras/stable/migration.html for help migrating.
           m2=KerasClassifier(build fn=create model,epochs=10,batch size=16,verbose=0)
In [33]:
          print(m2)
```

<keras.wrappers.scikit_learn.KerasClassifier object at 0x0000027FB39876A0>

(b) Create model using KerasClassifier wrapper class: Wrap your deep learning model using KerasClassifier wrapper:

^{&#}x27;\nkfold=KFold(n splits=4)\ncv results=cross val score(clf,X,y,cv=kfold,verbose=1)\n'

```
Out[34]:
```

(c) Evaluate model with cross-validation

```
std fit time [0.04970124 0.06568983 0.08453245 0.00267233 0.01904517 0.07985517]
mean score time [0.08825666 0.08235198 0.07608241 0.07194698 0.072833
                                                                        0.073623541
std score time [0.00861939 0.00559923 0.00145848 0.00149754 0.00072484 0.00268758]
param_batch_size [16 16 16 32 32 32]
param epochs [20 50 70 20 50 70]
params [{'batch_size': 16, 'epochs': 20}, {'batch_size': 16, 'epochs': 50}, {'batch_siz
e': 16, 'epochs': 70}, {'batch_size': 32, 'epochs': 20}, {'batch_size': 32, 'epochs': 5
0}, {'batch_size': 32, 'epochs': 70}]
split0_test_score [0.640625
                             0.703125    0.703125    0.60416669    0.66666669    0.69270831]
split1 test score [0.67708331 0.69791669 0.63541669 0.58333331 0.66666669 0.671875
split2 test score [0.6875 0.79166669 0.76041669 0.60416669 0.671875
                                                                          0.76041669]
                                                                          0.66666691
split3_test_score [0.61458331 0.63541669 0.65104169 0.65625
                                                              0.65625
mean_test_score [0.65494791 0.70703126 0.68750001 0.61197917 0.66536459 0.69791667]
std test score [0.02908634 0.0556555 0.0489971 0.02693771 0.00567565 0.03737683]
rank test score [5 1 3 6 4 2]
```

(d) Grid search parameters: Using GridSearchCV class, perform a grid search on the number of epochs [20,50,70], and the batch sizes of [5,10,20].

```
In [37]: dict1={'epochs':[20,50,70],'batch_size':[16,32]}
    clf = GridSearchCV(m2, dict1)
In []:
```

Problem 3: CNN Model for Photo Classification (15 points) Define a simple CNN network and evaluate how well it performs on the problem of CIFAR-10 Photo Classification. CIFAR is an acronym that stands for the Canadian Institute For Advanced Research and the CIFAR-10 dataset was developed along with the CIFAR-100 dataset by researchers at the CIFAR institute. The dataset is comprised of 60,000 32×32 pixel color photographs of objects from 10 classes, such as frogs, birds, cats, ships, etc. Part of the code below loads the CIFAR-10 train and test dataset using the Keras API.

```
from keras.datasets import cifar10
from keras.models import Sequential
from keras.layers import Dense
from keras.layers import Dropout
from keras.layers import Flatten
from keras.constraints import maxnorm
from tensorflow.keras.optimizers import SGD
from keras.layers.convolutional import Conv2D
from keras.layers.convolutional import MaxPooling2D
from keras.utils import np_utils
from tensorflow.keras.utils import to_categorical
# Load data
```

```
( X_train,y_train),(X_test ,y_test)=cifar10.load_data ()
# normalize inputs from 0-255 to 0.0-1.0
X_train = X_train.astype('float32')
X_test = X_test.astype('float32')
X_train = X_train / 255.0
X_test = X_test / 255.0
# one hot encode outputs
y_train = np_utils.to_categorical ( y_train )
y_test = np_utils.to_categorical ( y_test )
num_classes = y_test.shape [1]
```

- (a) Create the CNN model: Use a structure with two convolutional layers followed by max pooling and a flattening out of the network to fully connected layers to make predictions.
- (b) Compile the model: Compile your model with an SGD optimizer that uses learning rate schedule

```
In [39]:
          from keras.layers import Input,Conv2D,MaxPooling2D,UpSampling2D,Flatten,Conv2DTranspose
          from keras.models import Model, Sequential
          from keras.layers.core import Dense, Dropout
          from keras.layers.advanced activations import LeakyReLU
          def create_model():
              input1 = Input(shape=(32,32,3))
              first conv = Conv2D(32,(3,3),activation='relu',padding='same')(input1)
              #sec_max=MaxPooling2D((2,2))(first_conv)
              first conv=(BatchNormalization(momentum=0.8))(first conv)
              first_conv=Dropout(rate=0.3)(first_conv) # apply 30% dropout to the next layer
              first_conv = Conv2D(32,(3,3),activation='relu',padding='same')(input1)
              #sec max=MaxPooling2D((2,2))(first conv)
              first_conv=Dropout(rate=0.3)(first_conv) # apply 30% dropout to the next Layer
              first conv=(BatchNormalization(momentum=0.8))(first conv)
              sec_max=MaxPooling2D((2,2))(first_conv)
              flat = Flatten()(sec max)
              dense1=Dense(512,activation='relu')(flat)
              dense2=Dense(10, activation='softmax')(dense1)
              model = Model(inputs=input1, outputs=dense2, name="classify_model")
              model.compile(loss='categorical_crossentropy',optimizer='sgd',metrics=['accuracy'])
              return model
          mod1=create model()
```

In [40]:

mod1.summary()

Model: "classify_model"

Layer (type)	Output Shape	Param #
input_1 (InputLayer)	[(None, 32, 32, 3)]	0
conv2d_1 (Conv2D)	(None, 32, 32, 32)	896
dropout_1 (Dropout)	(None, 32, 32, 32)	0
<pre>batch_normalization_1 (Batc hNormalization)</pre>	(None, 32, 32, 32)	128
<pre>max_pooling2d (MaxPooling2D)</pre>	(None, 16, 16, 32)	0

(c) Fit the model

```
In [41]: __________
```

```
mod1.fit(np.array(X train),np.array(y train),epochs=25,batch size=32)
Epoch 1/25
4745
Epoch 2/25
Epoch 3/25
6645
Epoch 4/25
7175
Epoch 5/25
7607
Epoch 6/25
Epoch 7/25
8450
Epoch 8/25
8807
Epoch 9/25
9090
Epoch 10/25
9292
Epoch 11/25
1563/1563 [================ ] - 55s 35ms/step - loss: 0.1659 - accuracy: 0.
9459
Epoch 12/25
9569
Epoch 13/25
1563/1563 [================ ] - 50s 32ms/step - loss: 0.1061 - accuracy: 0.
9676
Epoch 14/25
9736
```

```
Epoch 15/25
  9790
  Epoch 16/25
  Epoch 17/25
  9841
  Epoch 18/25
  9852
  Epoch 19/25
  9870
  Epoch 20/25
  9900
  Epoch 21/25
  9898
  Epoch 22/25
  9916
  Epoch 23/25
  9918
  Epoch 24/25
  9929
  Epoch 25/25
  <keras.callbacks.History at 0x27fbcafee80>
Out[41]:
  (d) Evaluate the model
In [42]:
  x=[1,-1,-1,-1,7]
  f=np.array(x)
  print(f)
  print(np.gradient(f))
  [ 1 -1 -1 -1 7]
  [-2. -1. 0. 4. 8.]
In [ ]:
```