```
In [1]: import numpy as np
   import matplotlib.pyplot as plt
   import statsmodels as ss
   import numpy as np
   import matplotlib
   import pandas as pd
```

Exercise #1.1

In this exercise, your task is two-fold:

- 1. Determine if **Dataset_A** and **Dataset_B** are additive or multiplicative time series.
- 2. Determine the frequency of the seasonal component.

Set Path / Load Datasets

```
In [2]: # get data
path_to_file = "./" # Modify if data are in a different directory

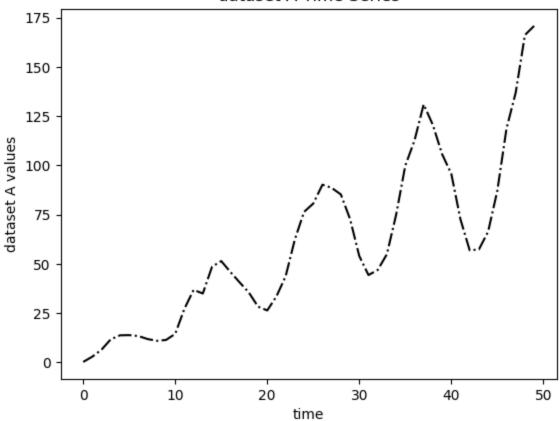
time = np.arange(0, 50)
dataset_A = np.load(path_to_file + "dataset_A.npy")
dataset_B = np.load(path_to_file + "dataset_B.npy")
```

Plot Dataset_A

```
In [3]: # insert code here

plt.plot(time, dataset_A, 'k-.')
plt.title("dataset A Time Series")
plt.xlabel("time")
plt.ylabel("dataset A values");
```

dataset A Time Series



Additive or Multiplicative?

multiplicative

Frequency of Seasonal Component?

10

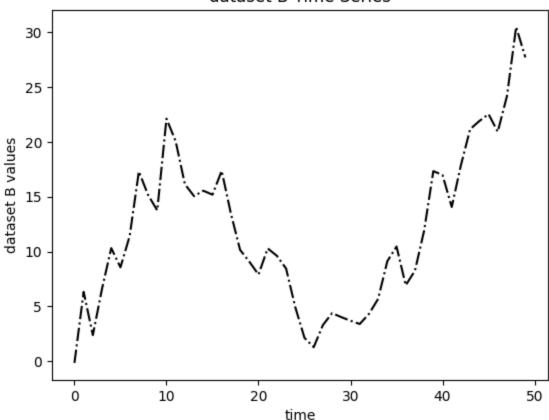
Plot Dataset_B

```
In [4]: # insert code here

#plt.plot(dataset_B)

plt.plot(time, dataset_B, 'k-.')
plt.title("dataset B Time Series")
plt.xlabel("time")
plt.ylabel("dataset B values");
```

dataset B Time Series



Additive or Multiplicative?

Additive

Frequency of Seasonal Component?

5

Exercise #1.2

In this exercise, your task is decompose **Dataset_A** and **Dataset_B**. You should first create a decomposition model in Python. Then you should plot the original series, the trend, seasonality, and residuals, in that order.

Decomposition Models

```
In [5]: from statsmodels.tsa.seasonal import seasonal_decompose
    from statsmodels.tsa.seasonal import seasonal_decompose

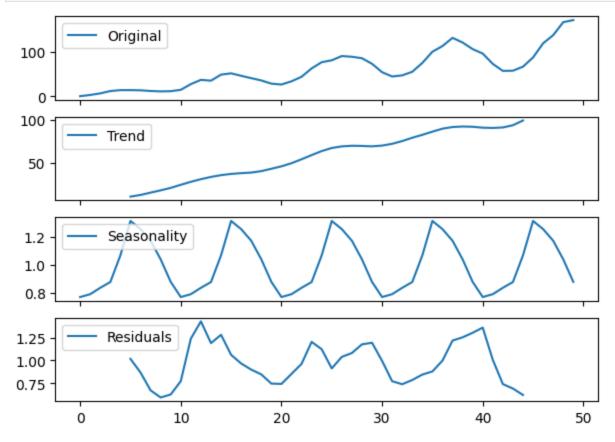
def ses_add(data, p):
    ss_decomposition = seasonal_decompose(x=data, model='additive', period=p)
    return ss_decomposition

def ses_mul(data, p):
```

```
ss_decomposition = seasonal_decompose(x=data, model='multiplicative', period=p)
return ss_decomposition
```

Dataset_A Plot

```
In [6]:
        ss_decomposition = ses_mul(dataset_A, 10)
        estimated_trend = ss_decomposition.trend
        estimated_seasonal = ss_decomposition.seasonal
        estimated_residual = ss_decomposition.resid
        fig, axes = plt.subplots(4, 1, sharex=True, sharey=False)
        fig.set_figheight(5)
        fig.set_figwidth(7)
        axes[0].plot(dataset_A, label='Original')
        axes[0].legend(loc='upper left');
        axes[1].plot(estimated_trend, label='Trend')
        axes[1].legend(loc='upper left');
        axes[2].plot(estimated_seasonal, label='Seasonality')
        axes[2].legend(loc='upper left');
        axes[3].plot(estimated_residual, label='Residuals')
        axes[3].legend(loc='upper left');
```



Dataset_B Plot

```
In [7]: ss_decomposition = ses_add(dataset_B, 5)
    estimated_trend = ss_decomposition.trend
    estimated_seasonal = ss_decomposition.seasonal
    estimated_residual = ss_decomposition.resid

fig, axes = plt.subplots(4, 1, sharex=True, sharey=False)
    fig.set_figheight(5)
```

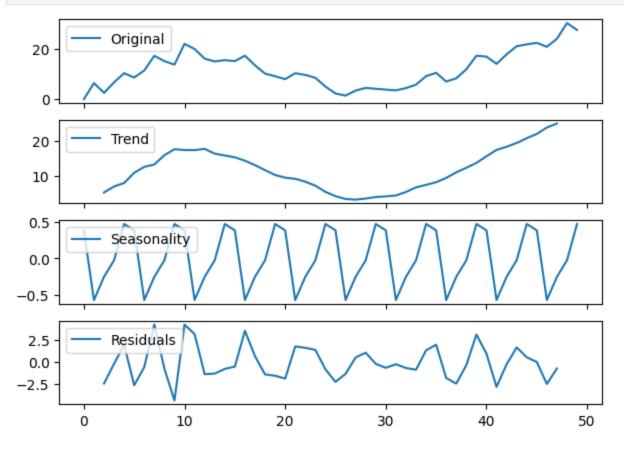
```
fig.set_figwidth(7)

axes[0].plot(dataset_B, label='Original')
axes[0].legend(loc='upper left');

axes[1].plot(estimated_trend, label='Trend')
axes[1].legend(loc='upper left');

axes[2].plot(estimated_seasonal, label='Seasonality')
axes[2].legend(loc='upper left');

axes[3].plot(estimated_residual, label='Residuals')
axes[3].legend(loc='upper left');
```



Exercise #2.1

In this exercise, your task is to:

- 1. Create a time variable called **mytime** that is composed of the integers from 0 to 99 inclusive.
- 2. Read in dataset_SNS_1.npy and dataset_SNS_2.npy as dataset_SNS_1 and dataset_SNS_2, respectively.
- 3. Plot each time series dataset.
- 4. Start thinking about whether each is stationary or nonstationary.

```
In [8]: # create time variable
    time = np.arange(0, 100)

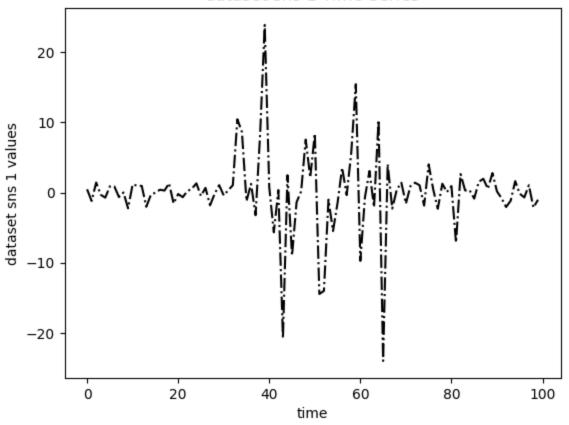
In [9]: # get data
    path_to_file = "./" # Modify if data are in a different directory
```

```
dataset_SNS_1 = np.load(path_to_file + "dataset_SNS_1.npy")
dataset_SNS_2 = np.load(path_to_file + "dataset_SNS_2.npy")

In [10]: # plot dataset_SNS_1

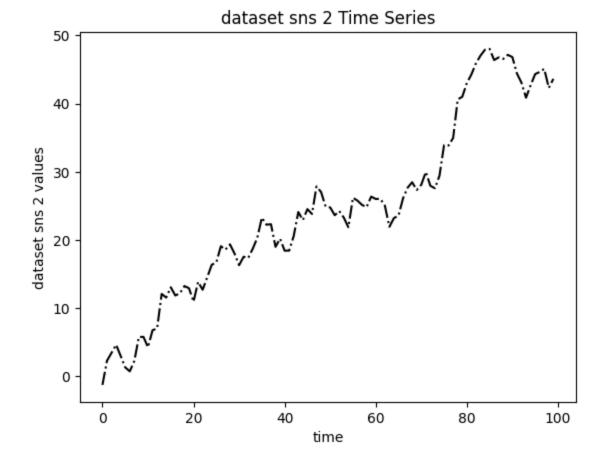
plt.plot(time, dataset_SNS_1, 'k-.')
plt.title("dataset sns 1 Time Series")
plt.xlabel("time")
plt.ylabel("dataset sns 1 values");
```

dataset sns 1 Time Series



```
In [11]: # plot dataset_SNS_2

plt.plot(time, dataset_SNS_2, 'k-.')
plt.title("dataset sns 2 Time Series")
plt.xlabel("time")
plt.ylabel("dataset sns 2 values");
```



Your Preliminary Thoughts

Are both datasets stationary or is one stationary and one nonstationary or are both nonstationary?

dataset 2 appears most definitely to be nonstationary because of clear trend, while dataset 1 appears to have heteroscedacity. It also appears to be non stationary

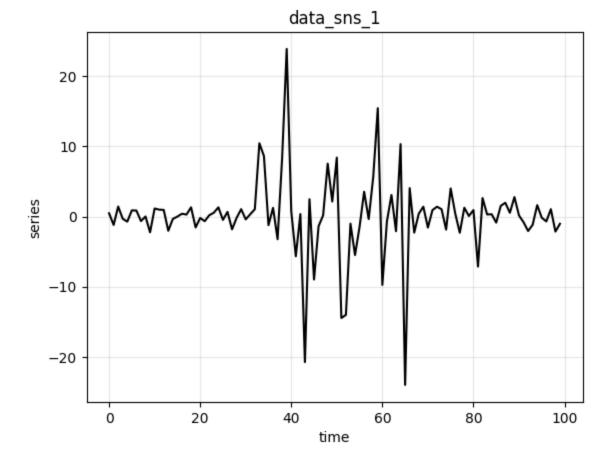
Exercise #2.2

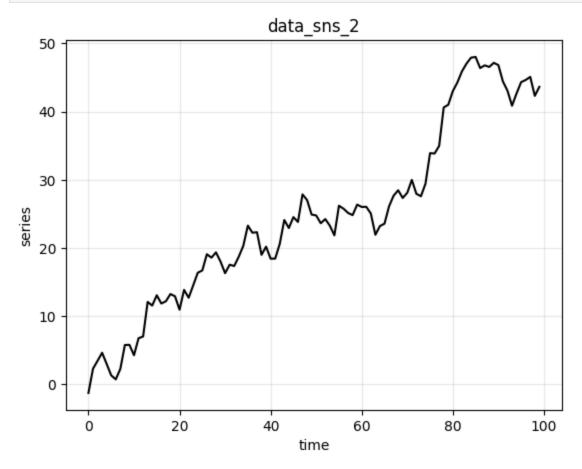
Think back to the two datasets from Exercise #2.1.

You should have the tools to answer whether each is stationary or not.

Provide your answers and explanations below.

```
In [12]: def run_sequence_plot(x, y, title, xlabel="time", ylabel="series"):
    plt.plot(x, y, 'k-')
    plt.title(title)
    plt.xlabel(xlabel)
    plt.ylabel(ylabel)
    plt.grid(alpha=0.3);
```

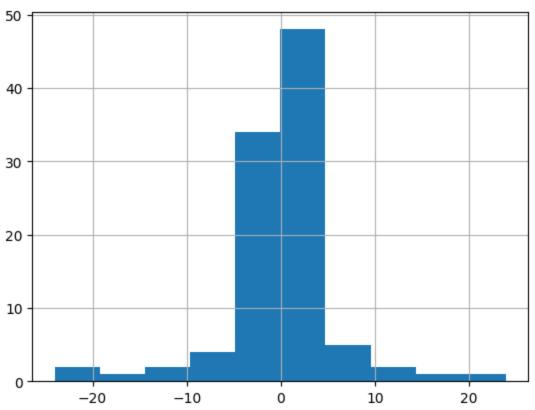


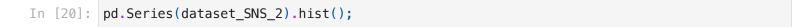


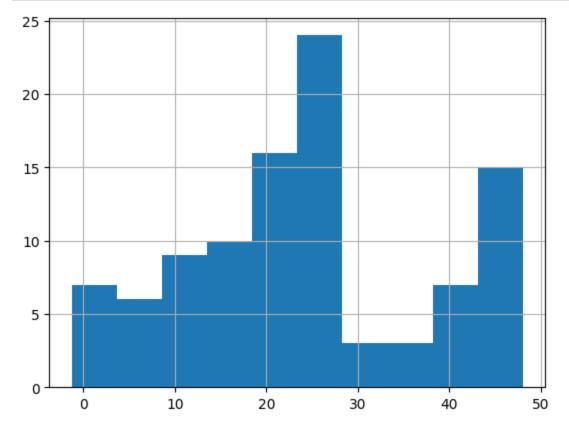
Explanation:

the observation is similar to the preliminary observation recorded in the first exercise, that being dataset 2 appearing as non stationary and dataset 1 appearing to be non-stationary due to it not having constant variance although appearing to have constant mean.

```
In [15]: # chunked stats
         # split data into 10 chunks
         chunks_1 = np.split(dataset_SNS_1, indices_or_sections=10)
         chunks_2 = np.split(dataset_SNS_2, indices_or_sections=10)
         np.mean(chunks_1, axis=1)
Out[15]: array([-0.14368349, 0.121089
                                            0.04714784, 4.92083495, -2.32626967,
                -0.3706503 , -1.95084875,
                                                         0.30483126, -0.51907842
                                            0.34634898,
In [16]: np.var(chunks_1, axis=1)
Out[16]: array([ 1.1064868 , 1.16521661,
                                            0.76635153, 59.91993023, 55.94682032,
                79.168351 , 77.81561226,
                                            3.18839451, 7.29302955, 1.34694601])
In [17]:
         np.mean(chunks_2, axis=1)
Out[17]: array([ 2.8186155 , 10.51083038, 16.02252814, 19.73251898, 23.26775272,
                24.60790371, 25.54323527, 32.73577811, 46.28942898, 43.77348672])
In [18]:
         np.var(chunks_2, axis=1)
Out[18]: array([ 4.50258195,
                              9.30343813, 7.51911006,
                                                         4.97894604,
                                                                      9.42707844,
                 1.77171835,
                              3.96080036, 22.71299455,
                                                         2.22467478,
                                                                      2.47819651])
         varying means and variances from both the datasets makes them out to be non stationary
In [19]:
         # histograms
         pd.Series(dataset_SNS_1).hist();
```







Explanation:

the second dataset is clearly not a normal distribution proving its non stationary property, while the first data set appears to be closely normal distributed suggesting it may be stationary.

```
In [21]: # ADF tests
    from statsmodels.tsa.stattools import adfuller
    adf, pvalue, usedlag, nobs, critical_values, icbest = adfuller(dataset_SNS_1)
    print("ADF: ", adf)
    print("p-value for 1st data set:", pvalue)

ADF: -3.032415903501602
    p-value for 1st data set: 0.03197606455861599

In [22]: adf, pvalue, usedlag, nobs, critical_values, icbest = adfuller(dataset_SNS_2)
    print("ADF: ", adf)
    print("p-value for 2nd data set:", pvalue)

ADF: -1.3222642986946496
```

Explanation:

p-value for 2nd data set: 0.6189258221979334

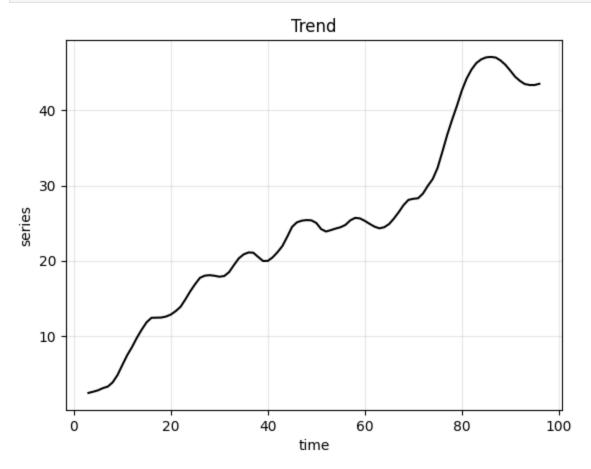
In the second dataset, p-value being well over 0.05 implies that we cannot reject the null hypthothesis. Hence dataset 2 is non stationary, on the other we we can reject the null hypthesis in favor of alternate in case of the forst data set as the p-value is below 0.05. Thus the first dataset can be considered stationary.

Exercise #2.3

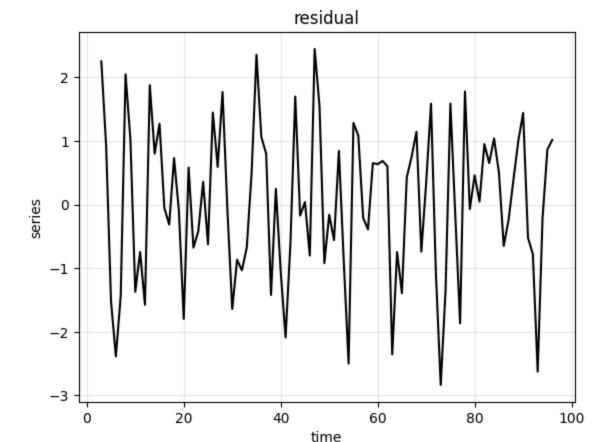
If either or both datasets from exercises one and two are nonstationary, apply the transformations you learned in this section to make them so. Then apply the methods you learned to ensure stationarity.

```
In [23]: #removing heteroscedasticity in dataset 1 by taking log
    #removing trend in dataset 2 by statsmodel
    from statsmodels.tsa.seasonal import seasonal_decompose
    ss_decomposition = seasonal_decompose(x=dataset_SNS_2, model='additive', period=6)
    est_trend = ss_decomposition.trend
    est_seasonal = ss_decomposition.seasonal
    est_residual = ss_decomposition.resid
```

```
In [24]: run_sequence_plot(time, est_trend, title="Trend", ylabel="series")
```



```
In [25]: run_sequence_plot(time, est_residual, title="residual", ylabel="series")
```



```
In [26]: print(est_residual)
```

```
0.91311651 -1.5222694
       nan
                    nan
                                nan
                                     2.25164332
-2.3861046 -1.43438458
                         2.04765693
                                     1.02213979 -1.36989103 -0.74622427
-1.57415546 1.87792254
                         0.8004592
                                     1.27164413 -0.05283683 -0.31420964
 0.7321111
          -0.09920165 -1.79714322
                                     0.58406459 -0.67627894 -0.41738726
 0.3586735
          -0.62448903
                         1.44611184
                                     0.59194224
                                                 1.77082168 -0.07267433
-1.64137035 -0.86710074 -1.03296816 -0.67411847
                                                 0.50378415
                                                             2.35538619
 1.06488261 0.79967166 -1.4207001
                                     0.2479742 - 1.05510569 - 2.08826227
-0.61832126 1.69836598 -0.17246994 0.03988933 -0.79967962 2.44734969
 1.54748355 -0.92071178 -0.15927568 -0.55751889
                                                 0.84458616 -0.88716597
                         1.08726871 -0.20703547 -0.39028846
-2.49946731 1.28485174
                                                             0.65156252
 0.63522219 0.68562812
                         0.60399827 -2.3539276
                                                -0.74505928 -1.39172098
 0.42817036 0.75334466
                         1.14428859 -0.73939069
                                                 0.34680138
                                                             1.58671918
-1.07113773 -2.83438054 -1.34636137
                                     1.58845154 -0.16128733 -1.86664715
                         0.45967517
                                     0.04808054
 1.77905834 -0.07086648
                                                 0.95101104
                                                             0.65333646
 1.03926848 0.50875097 -0.64876726 -0.2378658
                                                 0.39098803
                                                             0.99824046
 1.44094873 -0.52168038 -0.77605248 -2.62453756 -0.21924656
                                                             0.86540196
 1.01617305
                    nan
                                nan
                                            nan]
```

```
In [27]: adf_after, pvalue_after, usedlag_, nobs_, critical_values_, icbest_ = adfuller(est_resid
    print("ADF: ", adf_after)
    print("p-value: ", pvalue_after)
```

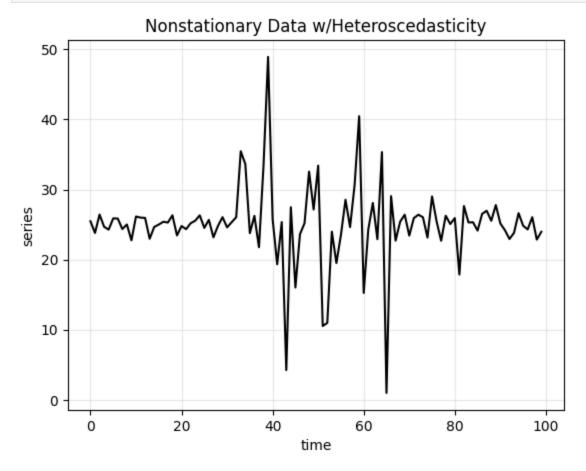
ADF: -6.238390718569909

p-value: 4.763184090747469e-08

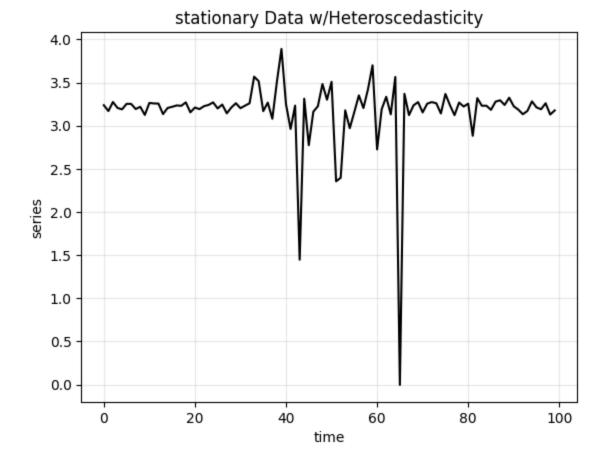
the p-value is sufficiently close to zero to reject the null hypothesis and consider the resiodual data as stationary.

removing heteroscedasticity in dataset 1 by taking log

```
In [28]: positive_data = dataset_SNS_1 + 25
```



ADF: -10.750411172845046 p-value: 2.674288989362179e-19



the first data set has relatively less heteroscedascity.

Exercise #3.1

You have been provided two datasets:

- 1. smooth_1.npy
- 2. smooth_2.npy

Your task is to leverage what you've learned in this and previous courses.

More specifically, you will do the following:

- 1. Read in **smooth_1.npy** and **smooth_2.npy**.
- 2. Create a time variable called **mytime** that starts at 0 and is as long as each dataset.
- 3. Split each dataset into train and test sets (test set is last 5 observations).
- 4. Identify trend and seasonality, if present.
- 5. Identify if trend and/or seasonality are additive or multiplicative, if present.
- 6. Create smoothed model on the train set and use to forecast on the test set.
- 7. Calculate MSE on test data.
- 8. Plot training data, test data, and your model's forecast for each dataset.

1. Get Data

```
In [30]: # get data
path_to_file = "./"
```

```
smooth_1 = np.load(path_to_file + "smooth_1.npy")
smooth_2 = np.load(path_to_file + "smooth_2.npy")

# Find the length of smooth_1 array
length_smooth_1 = len(smooth_1)

# Find the length of smooth_2 array
length_smooth_2 = len(smooth_2)

print("Length of smooth_1:", length_smooth_1)
print("Length of smooth_2:", length_smooth_2)
```

Length of smooth_1: 144 Length of smooth_2: 1000

2. Create mytime

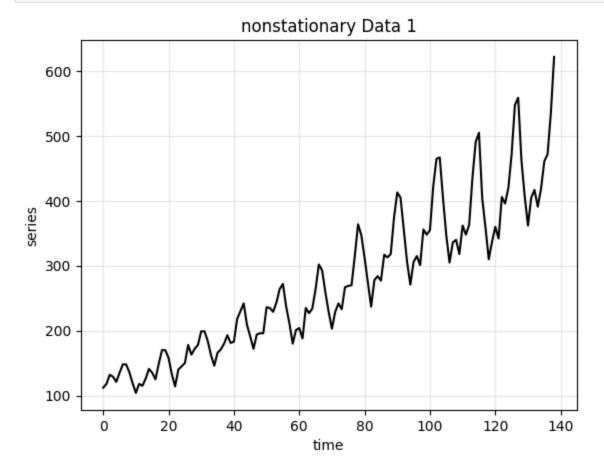
```
In [31]: mytime = np.arange(0, length_smooth_1)
   mytime_2 = np.arange(0, length_smooth_2)
```

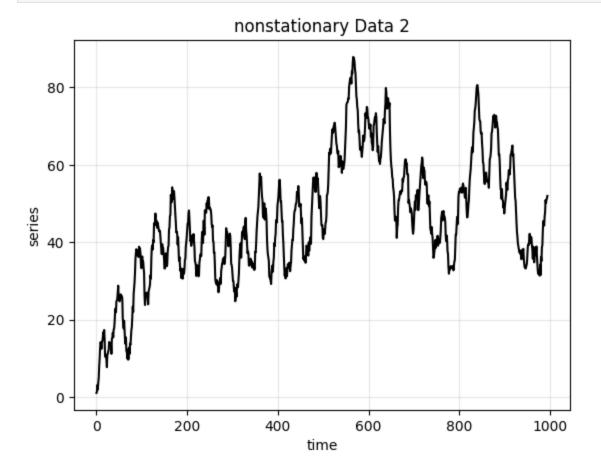
3. Train/Test Split

```
In [32]: train_1 = smooth_1[:-5]
    test_1 = smooth_1[-5:]

    train_2 = smooth_2[:-5]
    test_2 = smooth_2[-5:]
```

4. ID Trend / Seasonality





5. ID Additive vs Multicplicative

from the plots, by running a visual plot, the first time series appears to be multiplicative model.

from the plots, by running a visual plot, the second time series appears to be additive model.

6. Create Smoothed Models

7. Calculate MSE

```
In [37]: def mse(observations, estimates):

INPUT:

observations - numpy array of values indicating observed values
estimates - numpy array of values indicating an estimate of values

OUTPUT:

Mean Square Error value
```

```
# check arg types
assert type(observations) == type(np.array([])), "'observations' must be a numpy arr
assert type(estimates) == type(np.array([])), "'estimates' must be a numpy array"
# check length of arrays equal
assert len(observations) == len(estimates), "Arrays must be of equal length"

# calculations
difference = observations - estimates
sq_diff = difference ** 2
mse = sum(sq_diff)

return mse

In [38]: triple_preds_1 = triple_1.forecast(len(test_1))
triple_mse_1 = mse(test_1, triple_preds_1)
```

```
triple_mse_1 = mse(test_1, triple_preds_1)

print("MSE for first dataset : ", triple_mse_1)

MSE for first dataset : 633.505320669449

In [39]: triple_preds_2 = triple_2.forecast(len(test_2))
    triple_mse_2 = mse(test_2, triple_preds_2)

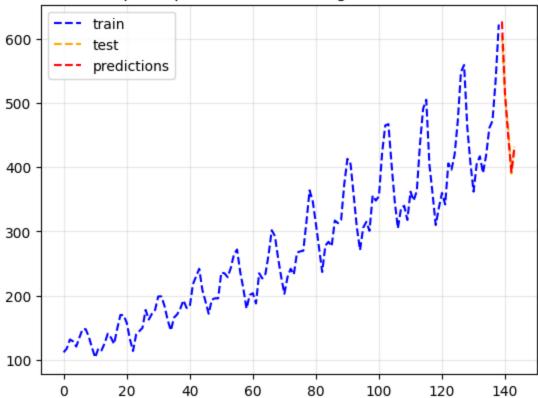
print("MSE for second dataset : ", triple_mse_2)
```

MSE for second dataset : 68.93466739372548

8. Plot Train, Test, Forecast

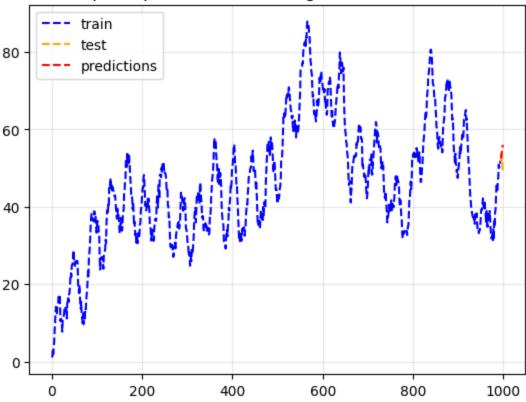
```
In [40]: plt.plot(mytime[:-5], train_1, 'b--', label="train")
    plt.plot(mytime[-5:], test_1, color='orange', linestyle="--", label="test")
    plt.plot(mytime[-5:], triple_preds_1, 'r--', label="predictions")
    plt.legend(loc='upper left')
    plt.title("Triple Exponential Smoothing -- 1st data set ")
    plt.grid(alpha=0.3);
```

Triple Exponential Smoothing -- 1st data set



```
In [41]: plt.plot(mytime_2[:-5], train_2, 'b--', label="train")
   plt.plot(mytime_2[-5:], test_2, color='orange', linestyle="--", label="test")
   plt.plot(mytime_2[-5:], triple_preds_2, 'r--', label="predictions")
   plt.legend(loc='upper left')
   plt.title("Triple Exponential Smoothing -- for 2nd data set ")
   plt.grid(alpha=0.3);
```

Triple Exponential Smoothing -- for 2nd data set



Exercise #4.1

You have been provided two datasets:

```
    auto_1.npy
    auto_2.npy
```

Your task is to leverage what you've learned in this and previous lectures.

More specifically, you will do the following:

- 1. Read in auto_1.npy and auto_2.npy.
- 2. Create a time variable called **mytime** that starts at 0 and is as long as both datasets.
- 3. Generate run-sequence plots of auto_1 and auto_2.
- 4. Determine the order of p and q.

1. Get Data

```
In [42]: # get data
path_to_file = "./"
auto_1 = np.load(path_to_file + "auto_1.npy")
auto_2 = np.load(path_to_file + "auto_2.npy")

# Find the length of smooth_1 array
length_auto_1 = len(auto_1)

# Find the length of smooth_2 array
length_auto_2 = len(auto_2)

print("Length of auto_1:", length_auto_1)
print("Length of auto_2:", length_auto_2)

Length of auto_1: 100
```

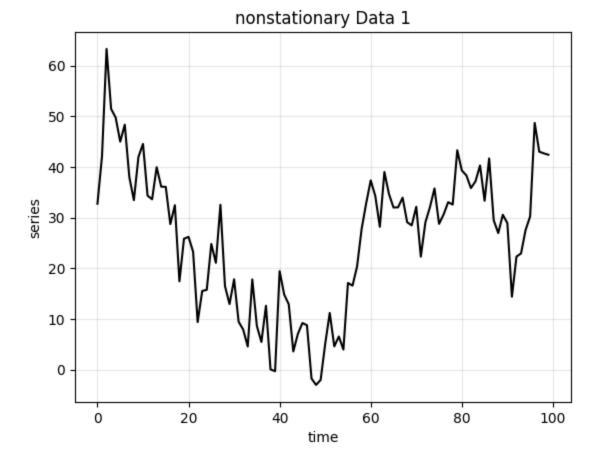
Length of auto_2: 100

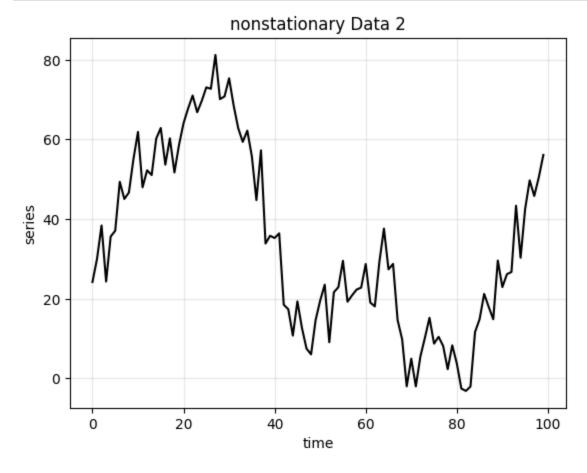
Create mytime

```
In [43]: # time component

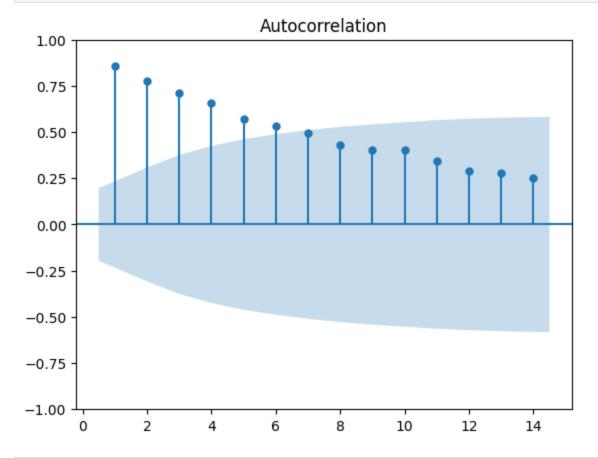
mytime = np.arange(0, length_auto_1)
mytime_2 = np.arange(0, length_auto_2)
```

Run-Sequence Plots

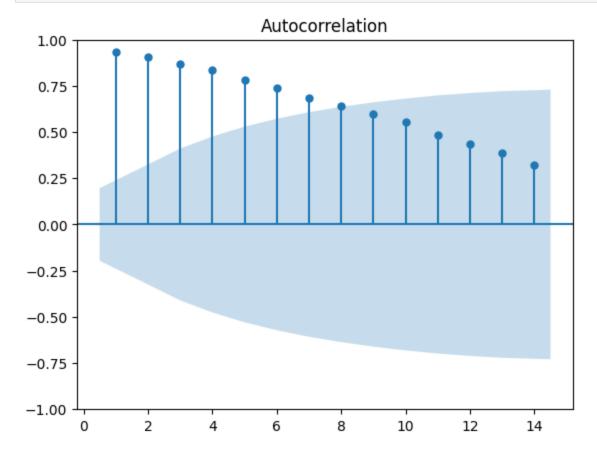




Determine Order (p & q)



In [47]: fig = plot_acf(auto_2, lags=range(1,15), alpha=0.05)



In [48]: from statsmodels.tsa.stattools import adfuller
 difference_1 = auto_1[:-1] - auto_1[1:]

```
print(pvalue)

p-value: 0.000
3.816622935918145e-24

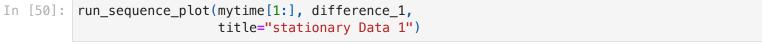
In [49]: difference_2 = auto_2[:-1] - auto_2[1:]

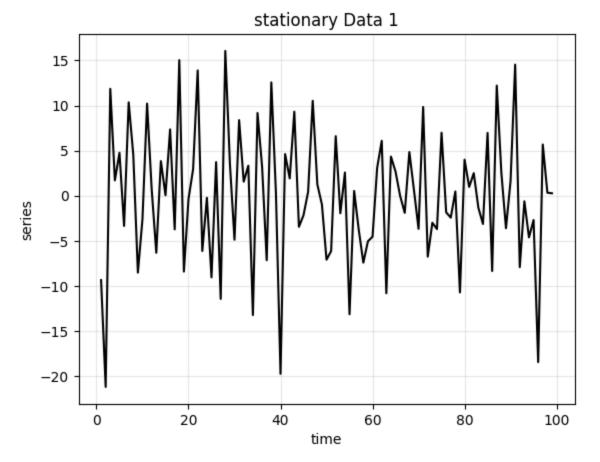
_, pvalue, _, _, _, _ = adfuller(difference_2)
    print(f"p-value: {pvalue:.3f}")
    print(pvalue)

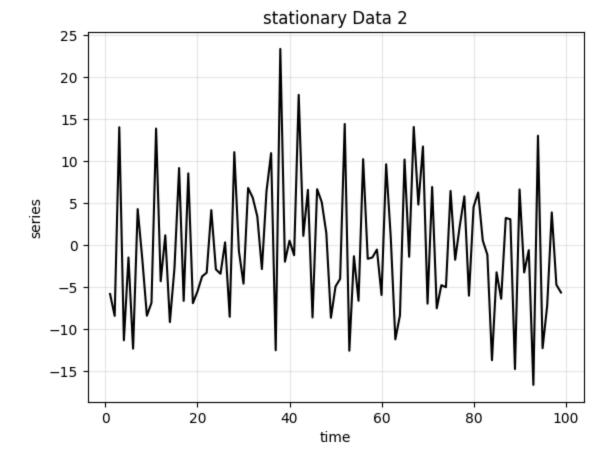
p-value: 0.000
4.667882463338721e-25
In [50]: run sequence plot(mytime[1:] difference_1)
```

= adfuller(difference_1)

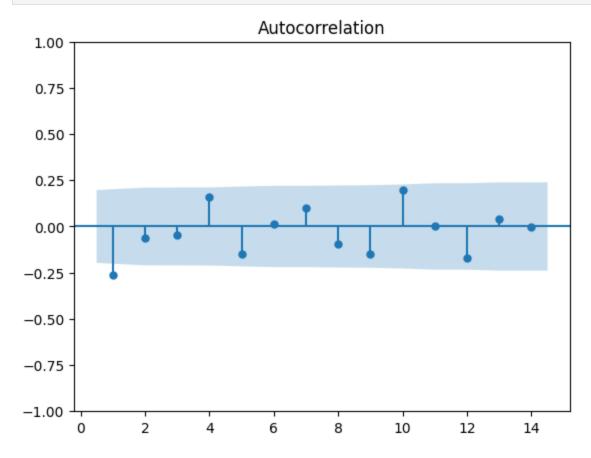
print(f"p-value: {pvalue:.3f}")



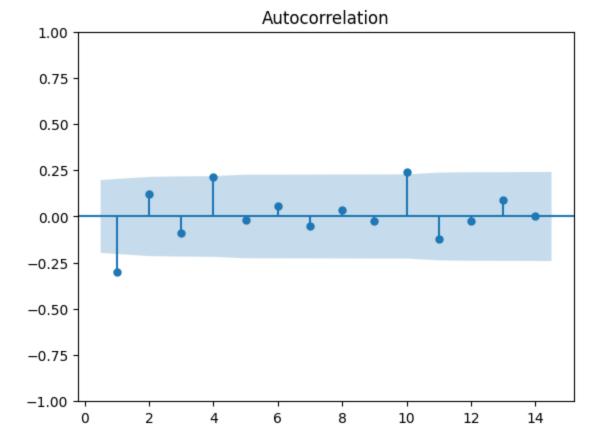




In [52]: fig = plot_acf(difference_1, lags=range(1,15), alpha=0.05)

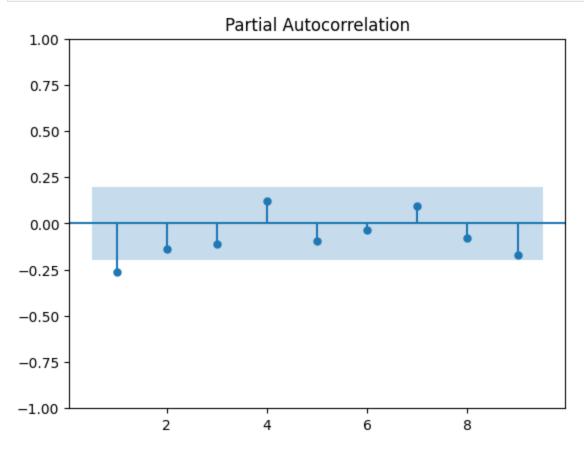


In [53]: fig = plot_acf(difference_2, lags=range(1,15), alpha=0.05)

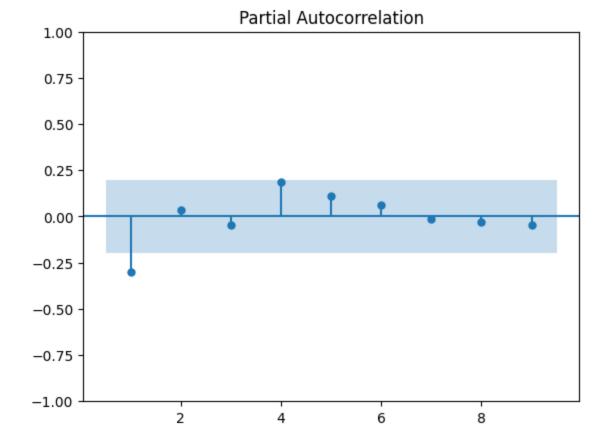


thus q value is chosen to be 1 in both the data sets as all the other branches in the above autocorrelation plot os almost zero

In [54]: from statsmodels.graphics.tsaplots import plot_pacf
fig = plot_pacf(difference_1, lags=range(1,10), alpha=0.05)



In [55]: fig = plot_pacf(difference_2, lags=range(1,10), alpha=0.05)



thus deciding the p value to be 1 as all the branches in the partial auto correlation plot drops to almost 0 after that \blacksquare

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