## **EXERCISE 3**

The data of ESE06\_ex3.csv report the daily changes of the General Motors Co. closing prices since September 4, 1998 to November 27, 1998.

- 1. Design a suitable quality control tool by assuming the existence of an assignable cause for the OOC observations if any.
- 2. Determine if the values reported in the array new\_obs are IC (use the previously designed control chart point 1). new\_obs = np.array((1.327, 1.594, 0.716, 1.767, 0.915, 2.524, 0.563, 2.053))

```
In []: # Import the necessary libraries
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
from scipy import stats
import qda

# Import the dataset
data = pd.read_csv('ESE06_ex3.csv')

# Inspect the dataset
data.head()
```

```
Out[]: GM

0 -0.875

1 2.437

2 -1.187

3 -2.063

4 0.938
```

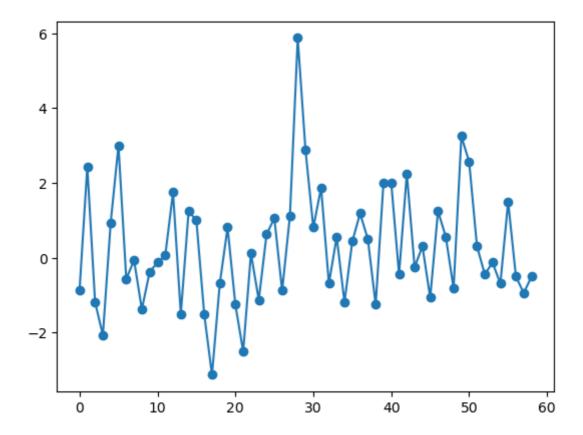
## Point 1

Design a suitable quality control tool (by using run rules too) by assuming the existence of an assignable cause for the OOC observations (if any).

### **Solution**

Let's plot the data first.

```
In [ ]: # Plot the data
plt.plot(data, 'o-')
plt.show()
```



Looks like there's one point with a value much higher than the others. But let's test all assumptions first.

Perform the runs test to check if the data are random. Use the runstest\_1samp function from the statsmodels package.

```
In []: # Verify if the data are random with runs test
     # Import the necessary Libraries for the runs test
     from statsmodels.sandbox.stats.runs import runstest_1samp

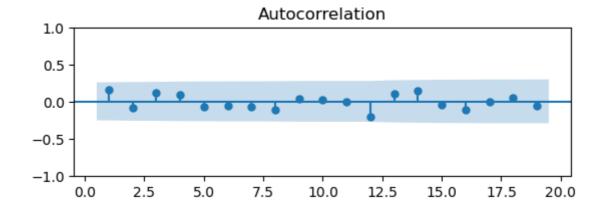
_, pval_runs = runstest_1samp(data['GM'], correction=False)
     print('Runs test p-value = {:.3f}'.format(pval_runs))
```

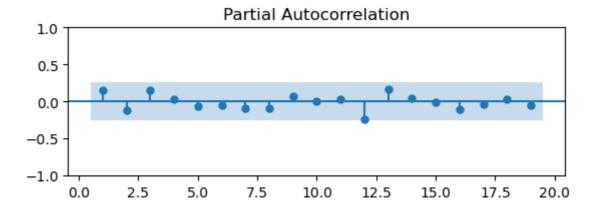
Runs test p-value = 0.879

Plot the autocorrelation and partial autocorrelation functions of the data. Use the plot\_acf and plot\_pacf functions from the statsmodels package.

```
In []: # Plot the acf and pacf using the statsmodels library
import statsmodels.graphics.tsaplots as sgt

fig, ax = plt.subplots(2, 1)
    sgt.plot_acf(data['GM'], lags = int(len(data)/3), zero=False, ax=ax[0])
    fig.subplots_adjust(hspace=0.5)
    sgt.plot_pacf(data['GM'], lags = int(len(data)/3), zero=False, ax=ax[1], method = plt.show()
```





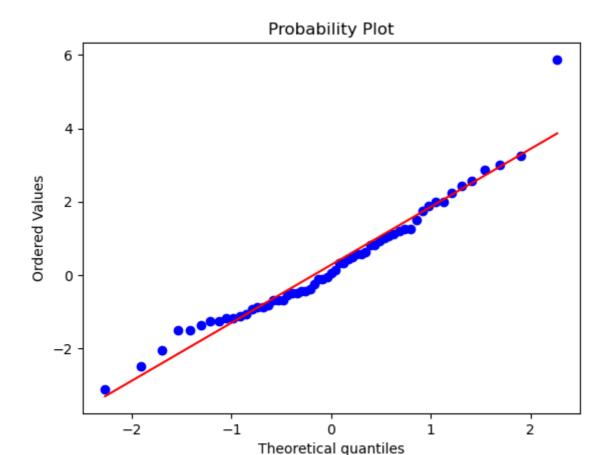
From the results of the autocorrelation and the runs tests, there is no statistical evidence to assume non randomness of the process.

Now let's verify the normality assumption.

```
In [ ]: # Perform the Shapiro-Wilk test
_, pval_SW = stats.shapiro(data['GM'])
print('Shapiro-Wilk test p-value = %.3f' % pval_SW)

# Plot the qaplot
stats.probplot(data['GM'], dist="norm", plot=plt)
plt.show()
```

Shapiro-Wilk test p-value = 0.068



We cannot reject the null hypothesis that the data are normally distributed with confidence 95%. However, one point deserves attention, as it is responsible for borderline normality.

Let's go ahead with the design of the I-MR control chart.

Remember, the computation of moving ranges MR:

- 1. Compute the differences between consecutive observations (lag = 1):  $D_i = X_{i+1} X_i. \label{eq:Discrete}$
- 2. Compute the absolute values of the differences:  $MR = |D_i|$ .

```
In [ ]: # Compute the moving ranges using the diff function
    data['MR'] = data['GM'].diff().abs()

# Print out descriptive statistics of MR and time
    data.describe()
```

Out[ ]:		GM	MR				
	count	59.000000	58.000000				
	mean	0.275424	1.743603				
	std	1.581153	1.091674				
	min	-3.125000	0.000000				
	25%	-0.844000	0.749500				
	50%	0.062000	1.718500				
	<b>75</b> %	1.156500	2.484500				
	max	5.875000	4.750000				

Now let's make the control chart for the mean of the moving ranges.

Remember the formulas for the control limits.

#### I chart:

- $egin{aligned} ullet & UCL = ar{x} + 3\left(rac{ar{MR}}{d_2}
  ight) \ ullet & CL = ar{x} \end{aligned}$
- ullet  $LCL=ar{x}-3\left(rac{ar{MR}}{d_2}
  ight)$

#### MR chart:

- $UCL = D_4 \bar{M}R$
- $CL = M\overline{R}$
- LCL = 0

### Factors for contructing variable control charts

	Char	Chart for Averages			Chart for Standard Deviations					Chart for Ranges						
bservations in	-	Factors for Control Limits		Factors for Center Line		Facto	Factors for Control Limits			Factors for Center Line		Factors for Control Limits				
Sample, n-	Λ	A A2		c <sub>4</sub>	1/c <sub>4</sub>	$B_3$	B4 B5	$B_5$	B4	$d_2$	$1/d_2$	$d_3$	$D_1$	$D_2$	$D_3$	$D_4$
2	2.121	1.880	2.659	0.7979	1.2533	0	3.267	0	2.606	1.128	0.8865	0.853	0	3.686	0	3.267
3	1.732	1.023	1.954	0.8862	1.1284	0	2.568	0	2.276	1.693	0.5907		Û	4.338	O .	2.574
4 -	1,500	0.729	1.628	0.9213	1.0854	0	.2.266	0	2.088	2.059	0.4857	0.880	0	4.698	0	2.282
5	1.342	0.577	1.427	0.9400	1.0638	0	2.089	0	1.964	2.326	0.4299	0.864	0	4.918	0	2.114
6	1.225	0.483	1.287	0.9515	1.0510	0.030	1.970	0.029	1.874	2,534	0.3946	0.848	D	5.078	0	2.004
7	1.134	0.419	1.182	0.9594	1.0423	0.118	1.882	0.113	1.806	2.704	0.3698	0.833	0.204	5.204	0.076	1.924
8	1.061	0.373	1.099	0.9650	1.0363	0.185	1.815	0.179	1.751	2.847	0.3512	0.820	0.388	5.306	0.136	1.864
9	1.000	0.337	1.032	0.9693	1.0317	0.239	1.761	0.232	1.707	2.970	0.3367	0.808	0.547	5.393	0.184	1.816
10	0.949	0.308	0.975	0.9727	1.0281	0.284	1.716	0.276	1.669	3.078	0.3249	0.797	0.687	5.469	0.223	1.777
.11	0.905	0.285	0.927	0.9754	1.0252	0.321	1.679	0.313	1.637	3.173	0.3152	0.787	0.811	5.535	0.256	1.744
12	0.866	0.266	0.886	0.9776	: 1.0229	0.354	1.646	0.346	1.610	3.258	0.3069	0.778	0.922	5.594	0.283	1.717
13	0.832	0.249	0.850	0.9794	1.0210	0.382	1.618	0.374	1.585	3.336	0.2998	0.770	1.025	5.647	0.307	1.693
14	0.802	0.235	0.817	0.9810	1.0194	0.406	1.594	0.399	1.563	3.407	0.2935	0.763	1.118	5.696	0.328	1.672
15	0.775	0.223	0.789	0.9823	1.0180	0.428	1.572	0.421	1.544	3.472	0.2880	0.756	1.203	5.741	0.347	1.653
16	0.750	0.212	0.763	0.9835	1.0168	0.448	1.552	0.440	1.526	3.532	0.2831	0.750	1.282	5.782	0.363	1.637
17	0.728	0.203	0.739	0.9845	1.0157	0.466	1.534	0.458	1.511	3.588	0.2787	0.744	1.356	5.820	0.378	1.622
18	0.707	0.194	0.718	0.9854	1.0148	0.482	1.518	0.475	1.496	3.640	0.2747	0.739	1.424	5.856	0.391	1.608
19	0.688	0.187	0.698	0.9862	1.0140	0.497	1.503	0.490	1.483	3.689	0.2711	0.734	1.487	5.891	0.403	1.597
20	0.671	0.180	0.680	0.9869	1.0133	0.510	1.490	0.504	1.470	3.735	0.2677	0.729	1.549	5.921	0.415	1.585
21	0.655	0.173	0.663	0.9876	1.0126	0.523	1.477	0.516	1.459	3.778	0.2647	0.724	1.605	5.951	0.425	1.575
22	0.640	0.167	0.647	0.9882	1.0119	0.534	1.466	0.528	1.448	3.819	0.2618	0.720	1.659	5.979	0.434	1.566
23	0.626	0.162	0.633	0.9887	1.0114	0.545	1.455	0.539	1.438	3.858	0.2592	0.716	1.710	6.006	0.443	1.557
24	0.612	0.157	0.619	0.9892	1.0109	0.555	1.445	0.549	1.429	3.895	0.2567	0.712	1.759	6.031	0.451	1.548
25	0.600	0.153	0.606	0.9896	1.0105	0.565	1.435	0.559	1.420	3.931	0.2544	0.708	1.806	6.056	0.459	1.541

```
D4 = 3.267
         # make a copy of the data
         df = data.copy()
         # change the name of the column time to I
         df.rename(columns={'GM':'I'}, inplace=True)
         # Print the first 5 rows of the new dataframe
         df.head()
                   MR
Out[ ]:
         0 -0.875 NaN
         1 2.437 3.312
         2 -1.187 3.624
        3 -2.063 0.876
         4 0.938 3.001
In [ ]: # Create columns for the upper and lower control limits
         df['I\_UCL'] = df['I'].mean() + (3*df['MR'].mean()/d2)
         df['I_CL'] = df['I'].mean()
         df['I_LCL'] = df['I'].mean() - (3*df['MR'].mean()/d2)
         df['MR\_UCL'] = D4 * df['MR'].mean()
         df['MR_CL'] = df['MR'].mean()
         df['MR_LCL'] = 0
         # Print the first 5 rows of the new dataframe
         df.head()
                                    I_CL
                                            I_LCL MR_UCL MR_CL MR_LCL
Out[ ]:
               1
                   MR
                          I_UCL
         0 -0.875 NaN 4.912667 0.275424 -4.361819 5.696352 1.743603
                                                                         0
         1 2.437 3.312 4.912667 0.275424 -4.361819 5.696352 1.743603
                                                                         0
         2 -1.187 3.624 4.912667 0.275424 -4.361819 5.696352 1.743603
                                                                         0
        3 -2.063 0.876 4.912667 0.275424 -4.361819 5.696352 1.743603
                                                                         0
         4 0.938 3.001 4.912667 0.275424 -4.361819 5.696352 1.743603
                                                                         0
In [ ]: # Define columns for possible violations of the control limits
         df['I_TEST1'] = np.where((df['I'] > df['I_UCL']) |
                         (df['I'] < df['I_LCL']), df['I'], np.nan)</pre>
         df['MR_TEST1'] = np.where((df['MR'] > df['MR_UCL']) |
                         (df['MR'] < df['MR_LCL']), df['MR'], np.nan)</pre>
```

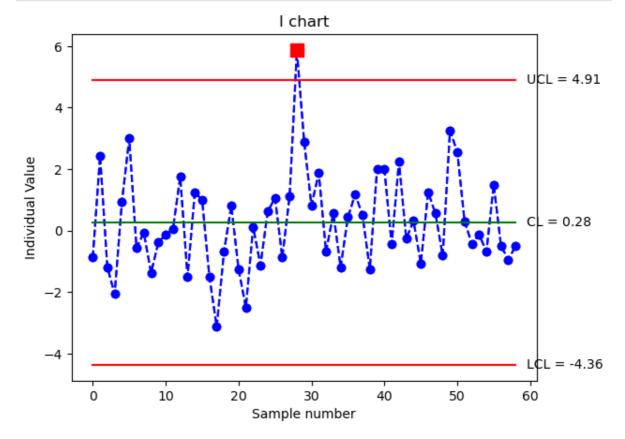
# Print the first 5 rows of the new dataframe

df.head()

```
Out[ ]:
                     MR
                             I UCL
                                        I CL
                                                 I_LCL MR_UCL
                                                                   MR_CL MR_LCL I_TEST1 MR_TEST1
         0 -0.875
                          4.912667
                                   0.275424
                                             -4.361819
                                                       5.696352
                                                                 1.743603
                    NaN
                                                                                       NaN
                                                                                                  NaN
                                                                                 0
             2.437 3.312 4.912667
                                   0.275424
                                             -4.361819
                                                       5.696352
                                                                1.743603
                                                                                       NaN
                                                                                                  NaN
            -1.187
                   3.624
                          4.912667
                                    0.275424
                                             -4.361819
                                                       5.696352
                                                                                 0
                                                                 1.743603
                                                                                       NaN
                                                                                                  NaN
            -2.063 0.876 4.912667
                                   0.275424
                                             -4.361819
                                                       5.696352 1.743603
                                                                                 0
                                                                                       NaN
                                                                                                  NaN
             0.938 3.001 4.912667 0.275424 -4.361819 5.696352 1.743603
                                                                                 0
                                                                                       NaN
                                                                                                  NaN
```

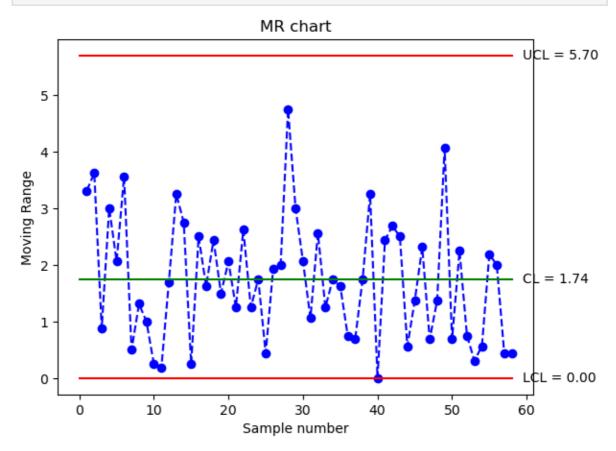
```
4
```

```
In []: # Plot the I chart
plt.title('I chart')
plt.plot(df['I'], color='b', linestyle='--', marker='o')
plt.plot(df['I'], color='b', linestyle='--', marker='o')
plt.plot(df['I_UCL'], color='r')
plt.plot(df['I_UCL'], color='g')
plt.plot(df['I_LCL'], color='r')
plt.ylabel('Individual Value')
plt.xlabel('Sample number')
# add the values of the control limits on the right side of the plot
plt.text(len(df)+.5, df['I_UCL'].iloc[0], 'UCL = {:.2f}'.format(df['I_UCL'].iloc[0]),
plt.text(len(df)+.5, df['I_CL'].iloc[0], 'CL = {:.2f}'.format(df['I_LCL'].iloc[0]),
plt.text(len(df)+.5, df['I_LCL'].iloc[0], 'LCL = {:.2f}'.format(df['I_LCL'].iloc[0])
# highlight the points that violate the alarm rules
plt.plot(df['I_TEST1'], linestyle='none', marker='s', color='r', markersize=10)
plt.show()
```



```
In [ ]: plt.title('MR chart')
    plt.plot(df['MR'], color='b', linestyle='--', marker='o')
    plt.plot(df['MR_UCL'], color='r')
    plt.plot(df['MR_CL'], color='g')
    plt.plot(df['MR_LCL'], color='r')
    plt.ylabel('Moving Range')
```

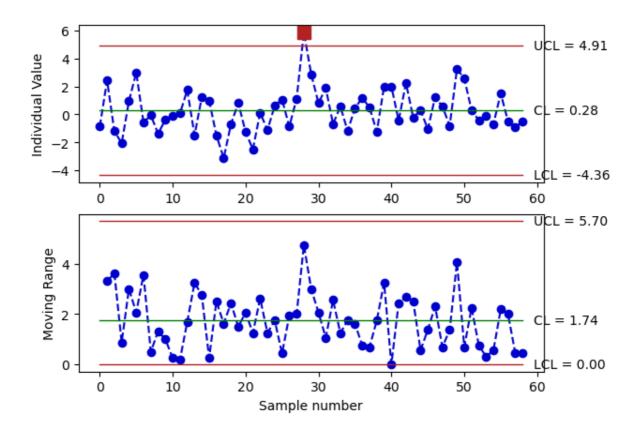
```
plt.xlabel('Sample number')
# add the values of the control limits on the right side of the plot
plt.text(len(df)+.5, df['MR_UCL'].iloc[0], 'UCL = {:.2f}'.format(df['MR_UCL'].iloc
plt.text(len(df)+.5, df['MR_CL'].iloc[0], 'CL = {:.2f}'.format(df['MR_CL'].iloc[0])
plt.text(len(df)+.5, df['MR_LCL'].iloc[0], 'LCL = {:.2f}'.format(df['MR_LCL'].iloc
# highlight the points that violate the alarm rules
plt.plot(df['MR_TEST1'], linestyle='none', marker='s', color='r', markersize=10)
plt.show()
```



There is one point outside the control limits.

You can also design the control chart using the IMR function in qda.ControlCharts package.

```
In [ ]: data_IMR = qda.ControlCharts.IMR(data, 'GM')
```



Let's find the index of the OOC point.

```
In [ ]: # Find the index of the I_TEST1 column different from NaN
OOC_idx = np.where(data_IMR['I_TEST1'].notnull())[0]
# Print the index of the OOC points
print('The index of the OOC point is: {}'.format(OOC_idx))
```

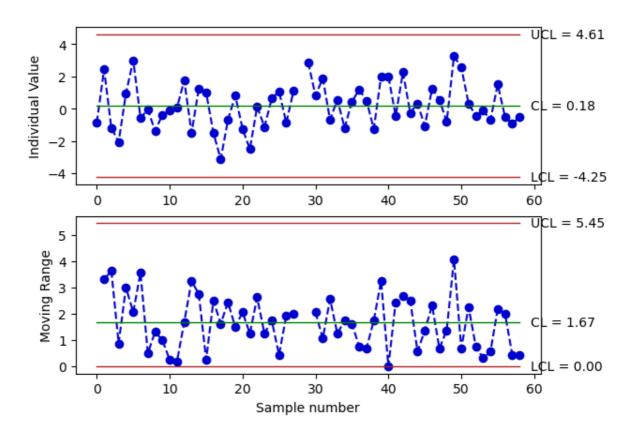
The index of the OOC point is: [28]

Index 28 (i.e. **the 29th** observation) is the one that is out of control.

Assume we found an assignable cause for the OOC point, we have to remove it from the data.

```
In []: # make a copy of the data
data_2 = data.copy()
# replace the OOC point with NaN
data_2['GM'].iloc[OOC_idx] = np.nan

# Plot the new control chart
data_IMR_2 = qda.ControlCharts.IMR(data_2, 'GM')
```



# Point 2

Determine if the values reported in the array new\_obs are IC (use the previously designed control chart point 1).

```
In []: # Import the dataset
data = pd.read_csv('ESE06_ex3.csv')
# replace the OOC point with NaN
data['GM'].iloc[OOC_idx] = np.nan

# Define the array with the new observations
new_obs = np.array((1.327, 1.594, 0.716, 1.767, 0.915, 2.524, 0.563, 2.053))

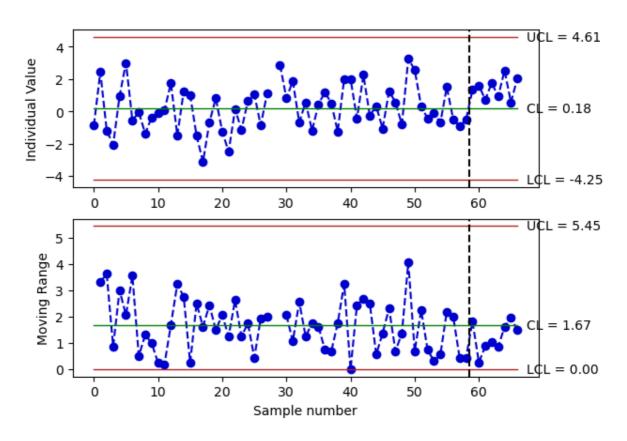
# Add the new observations to the dataset
new_data = data.append(pd.DataFrame(new_obs, columns=['GM']), ignore_index=True)
new_data.tail(10)
```

```
Out[]:
                 GM
               -0.937
          58
               -0.500
                1.327
          59
          60
               1.594
          61
               0.716
          62
               1.767
          63
               0.915
               2.524
          64
          65
               0.563
               2.053
          66
```

Use the parameter subset\_size to specify the number of observations to be used for the computation of the control limits (Phase 1 data).

```
In [ ]: new_data_IMR = qda.ControlCharts.IMR(new_data, 'GM', subset_size=len(data))
```

#### I-MR charts of GM



The new data (Phase II) are in-control