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import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from scipy import stats
from statsmodels.tsa.stattools import acf, pacf, adfuller, kpss
from statsmodels.stats.diagnostic import acorr_ljungbox
from statsmodels.graphics.tsaplots import plot_acf, plot_pacf

# Set random seed for reproducibility
np.random.seed(42)

# Number of observations
T = 200

# Function to generate the time series
def generate_models():
    # Model (a):  $X_t - X_{t-1} = Z_t + 0.3Z_{t-1}$ ,  $Z_t \sim \text{iid Normal}(0, 2)$ 
    # This is an ARIMA(0,1,1) model or IMA(1,1)
    z_a = np.random.normal(0, np.sqrt(2), T+1)
    x_a = np.zeros(T)
    for t in range(1, T):
        x_a[t] = x_a[t-1] + z_a[t] + 0.3 * z_a[t-1]

    # Model (b):  $X_t + 0.5X_{t-1} = Z_t - 0.7Z_{t-1}$ ,  $Z_t \sim \text{iid Normal}(0, 4)$ 
    # This is an ARMA(1,1) model
    z_b = np.random.normal(0, 2, T+1) # std = sqrt(4)
    x_b = np.zeros(T)
    for t in range(1, T):
        x_b[t] = -0.5 * x_b[t-1] + z_b[t] - 0.7 * z_b[t-1]

    # Model (c):  $X_t = 0.5X_{t-1} + 0.3X_{t-2} + Z_t$ ,  $Z_t \sim \text{iid } t\text{-distributed with 5 degrees of freedom}$ 
    # This is an AR(2) model with t-distributed innovations
    z_c = stats.t.rvs(5, size=T)
    x_c = np.zeros(T)
    # Taking X1 and X2 as zero (as specified)
    for t in range(2, T):
        x_c[t] = 0.5 * x_c[t-1] + 0.3 * x_c[t-2] + z_c[t]

    # Model (d):  $X_t = Z_t$ , where  $Z_t$  is iid standard Laplace with pdf
    #  $f(x) = (1/2)\exp(-|x|)$ 
    # This is white noise with Laplace distribution
    u = np.random.uniform(-0.5, 0.5, T)
    x_d = -np.sign(u) * np.log(1 - 2 * np.abs(u))

    return x_a, x_b, x_c, x_d

# Generate the time series
x_a, x_b, x_c, x_d = generate_models()

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# Create DataFrames for each model
df_a = pd.DataFrame({'Model A': x_a})
df_b = pd.DataFrame({'Model B': x_b})
df_c = pd.DataFrame({'Model C': x_c})
df_d = pd.DataFrame({'Model D': x_d})

# Function to analyze time series properties
def analyze_series(series, model_name, max_lag=20):
    print(f"\n===== Analysis for {model_name} =====\n")

    # Summary statistics
    print("Summary Statistics:")
    print(f"Mean: {np.mean(series):.4f}")
    print(f"Variance: {np.var(series):.4f}")
    print(f"Min: {np.min(series):.4f}")
    print(f"Max: {np.max(series):.4f}")

    # ADF Test for stationarity
    adf_result = adfuller(series)
    print("\nAugmented Dickey-Fuller Test:")
    print(f"ADF Statistic: {adf_result[0]:.4f}")
    print(f"p-value: {adf_result[1]:.4f}")
    print("Critical Values:")
    for key, value in adf_result[4].items():
        print(f"    {key}: {value:.4f}")
    print(f"Stationary: {adf_result[1] < 0.05}")

    # KPSS Test for stationarity
    try:
        kpss_result = kpss(series, regression='c')
        print("\nKPSS Test:")
        print(f"KPSS Statistic: {kpss_result[0]:.4f}")
        print(f"p-value: {kpss_result[1]:.4f}")
        print("Critical Values:")
        for key, value in kpss_result[3].items():
            print(f"    {key}: {value:.4f}")
        print(f"Stationary: {kpss_result[1] > 0.05}")
    except:
        print("\nKPSS Test: Could not compute (may be due to perfect stationarity)")

    # Ljung-Box test for white noise
    lb_result = acorr_ljungbox(series, lags=max_lag)
    print("\nLjung-Box Test for White Noise:")
    print(f"Q-statistic at lag {max_lag}: {lb_result.iloc[-1, 0]:.4f}")
    print(f"p-value: {lb_result.iloc[-1, 1]:.4f}")
    print(f"White Noise: {lb_result.iloc[-1, 1] > 0.05}")

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# Jarque-Bera test for normality
jb_result = stats.jarque_bera(series)
print("\nJarque-Bera Test for Normality:")
print(f"JB Statistic: {jb_result[0]:.4f}")
print(f"p-value: {jb_result[1]:.4f}")
print(f"Skewness: {stats.skew(series):.4f}")
print(f"Kurtosis: {stats.kurtosis(series):.4f}")
print(f"Normal Distribution: {jb_result[1] > 0.05}")

# Visualization for all models
def plot_model(series, model_name, max_lag=20):
    plt.figure(figsize=(15, 10))

    # Time Series Plot
    plt.subplot(3, 1, 1)
    plt.plot(series)
    plt.title(f'Time Series - {model_name}')
    plt.xlabel('Time')
    plt.ylabel('Value')
    plt.grid(True)

    # ACF Plot
    plt.subplot(3, 1, 2)
    plot_acf(series, lags=max_lag, ax=plt.gca())
    plt.title(f'Autocorrelation Function (ACF) - {model_name}')

    # PACF Plot
    plt.subplot(3, 1, 3)
    plot_pacf(series, lags=max_lag, ax=plt.gca())
    plt.title(f'Partial Autocorrelation Function (PACF) - {model_name}')

    plt.tight_layout()
    plt.show()

# Analyze each model
analyze_series(x_a, "Model A:  $X_t - X_{t-1} = Z_t + 0.3Z_{t-1}$ ,  $Z_t \sim \text{iid Normal}(0, 2)$ ")
plot_model(x_a, "Model A:  $X_t - X_{t-1} = Z_t + 0.3Z_{t-1}$ ,  $Z_t \sim \text{iid Normal}(0, 2)$ ")

analyze_series(x_b, "Model B:  $X_t + 0.5X_{t-1} = Z_t - 0.7Z_{t-1}$ ,  $Z_t \sim \text{iid Normal}(0, 4)$ ")
plot_model(x_b, "Model B:  $X_t + 0.5X_{t-1} = Z_t - 0.7Z_{t-1}$ ,  $Z_t \sim \text{iid Normal}(0, 4)$ ")

analyze_series(x_c, "Model C:  $X_t = 0.5X_{t-1} + 0.3X_{t-2} + Z_t$ ,  $Z_t \sim \text{iid } t(5)$ ")
plot_model(x_c, "Model C:  $X_t = 0.5X_{t-1} + 0.3X_{t-2} + Z_t$ ,  $Z_t \sim \text{iid } t(5)$ ")

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analyze_series(x_d, "Model D:  $X_t = Z_t$ , where  $Z_t \sim \text{iid Laplace}(0, 1)$ ")
plot_model(x_d, "Model D:  $X_t = Z_t$ , where  $Z_t \sim \text{iid Laplace}(0, 1)$ ")

# Compare distributions using histograms and QQ plots
def compare_distributions(series, model_name, theoretical_dist):
    plt.figure(figsize=(12, 5))

    # Histogram with theoretical PDF
    plt.subplot(1, 2, 1)
    plt.hist(series, bins=30, density=True, alpha=0.7)

    # Add theoretical PDF
    x = np.linspace(min(series), max(series), 1000)
    if theoretical_dist == 'normal':
        mu, sigma = np.mean(series), np.std(series)
        plt.plot(x, stats.norm.pdf(x, mu, sigma), 'r-', lw=2)
        plt.title(f'Histogram vs. Normal PDF - {model_name}')
    elif theoretical_dist == 'laplace':
        mu, b = np.mean(series), np.std(series) / np.sqrt(2) # Laplace scale parameter
        plt.plot(x, stats.laplace.pdf(x, mu, b), 'r-', lw=2)
        plt.title(f'Histogram vs. Laplace PDF - {model_name}')
    elif theoretical_dist == 't':
        # For t-distribution, need to estimate parameters
        params = stats.t.fit(series)
        plt.plot(x, stats.t.pdf(x, *params), 'r-', lw=2)
        plt.title(f'Histogram vs. t-distribution PDF - {model_name}')

    plt.xlabel('Value')
    plt.ylabel('Density')

    # QQ plot
    plt.subplot(1, 2, 2)
    if theoretical_dist == 'normal':
        stats.probplot(series, dist="norm", plot=plt)
        plt.title(f'Q-Q Plot vs. Normal - {model_name}')
    elif theoretical_dist == 'laplace':
        # For Laplace QQ plot, use uniform and transform
        u = np.random.uniform(0, 1, len(series))
        laplace_quantiles = np.sign(u - 0.5) * (-np.log(1 - 2 *
np.abs(u - 0.5)))
        plt.scatter(np.sort(laplace_quantiles), np.sort(series))
        plt.plot([min(laplace_quantiles), max(laplace_quantiles)],
                [min(series), max(series)], 'r-')
        plt.title(f'Q-Q Plot vs. Laplace - {model_name}')
    elif theoretical_dist == 't':
        # Use t-distribution with estimated parameters
        params = stats.t.fit(series)
        stats.probplot(series, dist=stats.t, sparams=params[0],

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plot=plt)
    plt.title(f'Q-Q Plot vs. t-distribution - {model_name}')

    plt.tight_layout()
    plt.show()
```

#### *# Compare distributions*

```
compare_distributions(x_a, "Model A", 'normal')
compare_distributions(x_b, "Model B", 'normal')
compare_distributions(x_c, "Model C", 't')
compare_distributions(x_d, "Model D", 'laplace')
```

===== Analysis for Model A:  $X_t - X_{t-1} = Z_t + 0.3Z_{t-1}$ ,  $Z_t \sim \text{iid Normal}(0, 2)$  =====

#### Summary Statistics:

Mean: -15.2397  
Variance: 54.9160  
Min: -25.5030  
Max: 7.3049

#### Augmented Dickey-Fuller Test:

ADF Statistic: -2.7489  
p-value: 0.0660  
Critical Values:  
1%: -3.4645  
5%: -2.8766  
10%: -2.5748  
Stationary: False

#### KPSS Test:

KPSS Statistic: 0.7057  
p-value: 0.0130  
Critical Values:  
10%: 0.3470  
5%: 0.4630  
2.5%: 0.5740  
1%: 0.7390  
Stationary: False

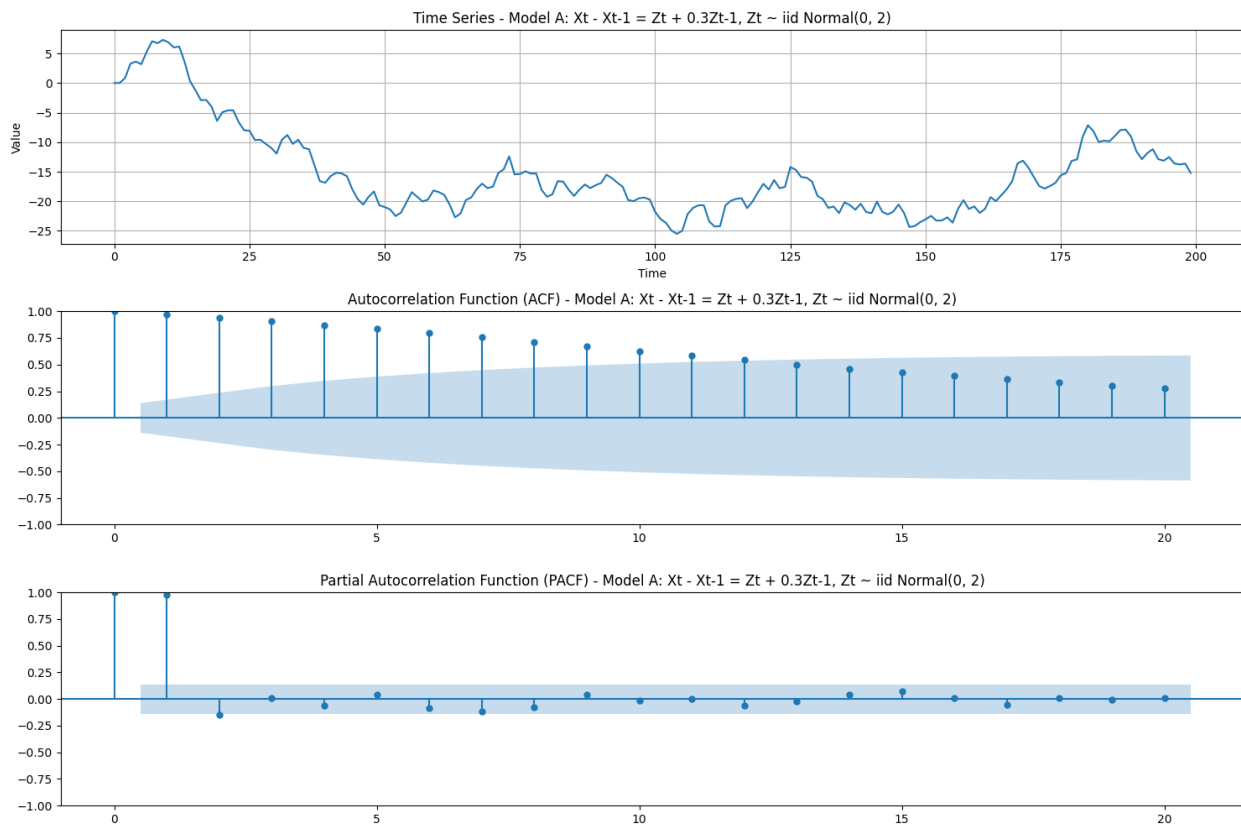
#### Ljung-Box Test for White Noise:

Q-statistic at lag 20: 1778.0986  
p-value: 0.0000  
White Noise: False

#### Jarque-Bera Test for Normality:

JB Statistic: 75.7594  
p-value: 0.0000  
Skewness: 1.3438

Kurtosis: 1.3665  
Normal Distribution: False



==== Analysis for Model B:  $X_t + 0.5X_{t-1} = Z_t - 0.7Z_{t-1}$ ,  $Z_t \sim \text{iid Normal}(0, 4)$  =====

Summary Statistics:

Mean: 0.0067

Variance: 12.3688

Min: -11.3203

Max: 8.8598

Augmented Dickey-Fuller Test:

ADF Statistic: -9.1048

p-value: 0.0000

Critical Values:

1%: -3.4645

5%: -2.8766

10%: -2.5748

Stationary: True

KPSS Test:

KPSS Statistic: 0.1327

p-value: 0.1000  
Critical Values:  
10%: 0.3470  
5%: 0.4630  
2.5%: 0.5740  
1%: 0.7390

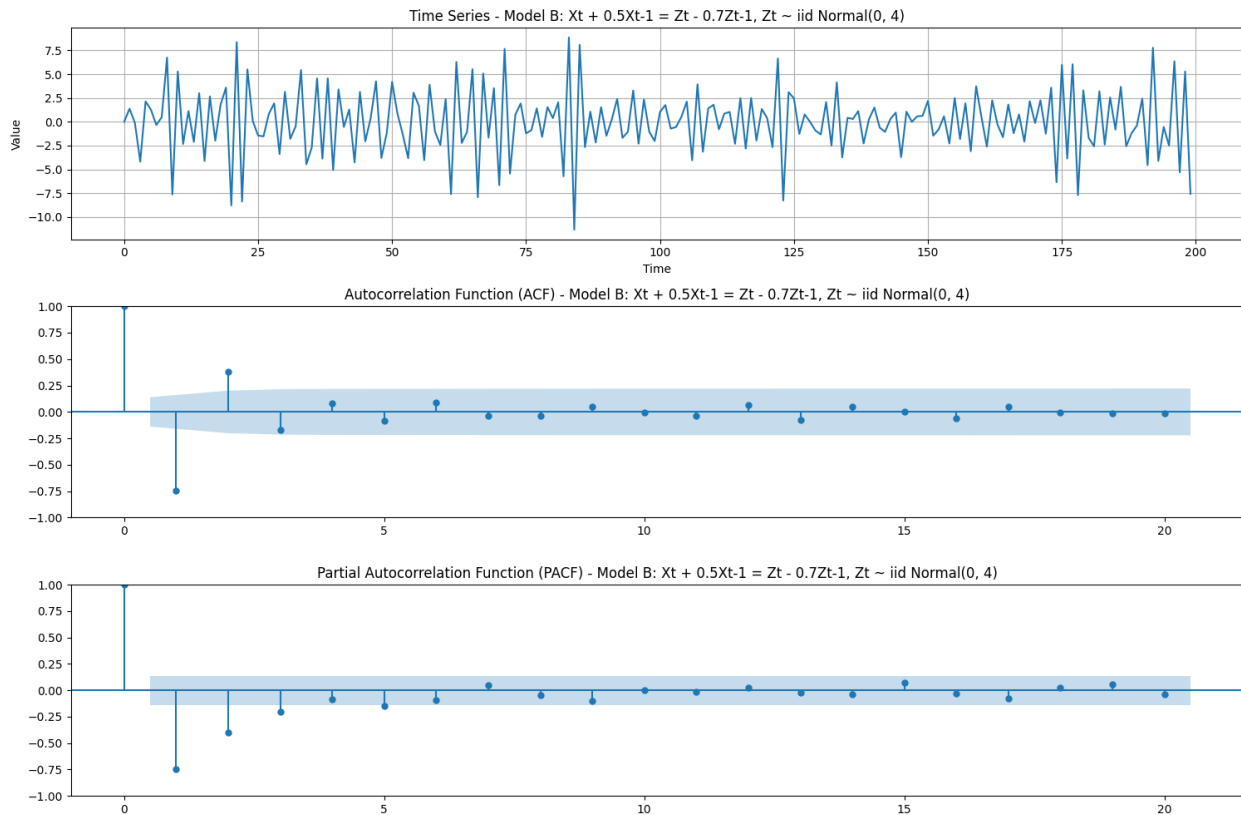
Stationary: True

Ljung-Box Test for White Noise:  
Q-statistic at lag 20: 158.7115  
p-value: 0.0000  
White Noise: False

Jarque-Bera Test for Normality:  
JB Statistic: 2.3503  
p-value: 0.3088  
Skewness: -0.1850  
Kurtosis: 0.3810  
Normal Distribution: True

/var/folders/7z/h\_\_cj55j6hb0wqzy9tb5b92w0000gn/T/  
ipykernel\_78328/935025886.py:78: InterpolationWarning: The test  
statistic is outside of the range of p-values available in the  
look-up table. The actual p-value is greater than the p-value  
returned.

```
kpss_result = kpss(series, regression='c')
```



===== Analysis for Model C:  $X_t = 0.5X_{t-1} + 0.3X_{t-2} + Z_t$ ,  $Z_t \sim \text{iid } t(5)$  =====

Summary Statistics:

Mean: -0.5219

Variance: 2.3341

Min: -4.7641

Max: 4.6653

Augmented Dickey-Fuller Test:

ADF Statistic: -4.9543

p-value: 0.0000

Critical Values:

1%: -3.4638

5%: -2.8763

10%: -2.5746

Stationary: True

KPSS Test:

KPSS Statistic: 0.0813

p-value: 0.1000

Critical Values:

10%: 0.3470

5%: 0.4630



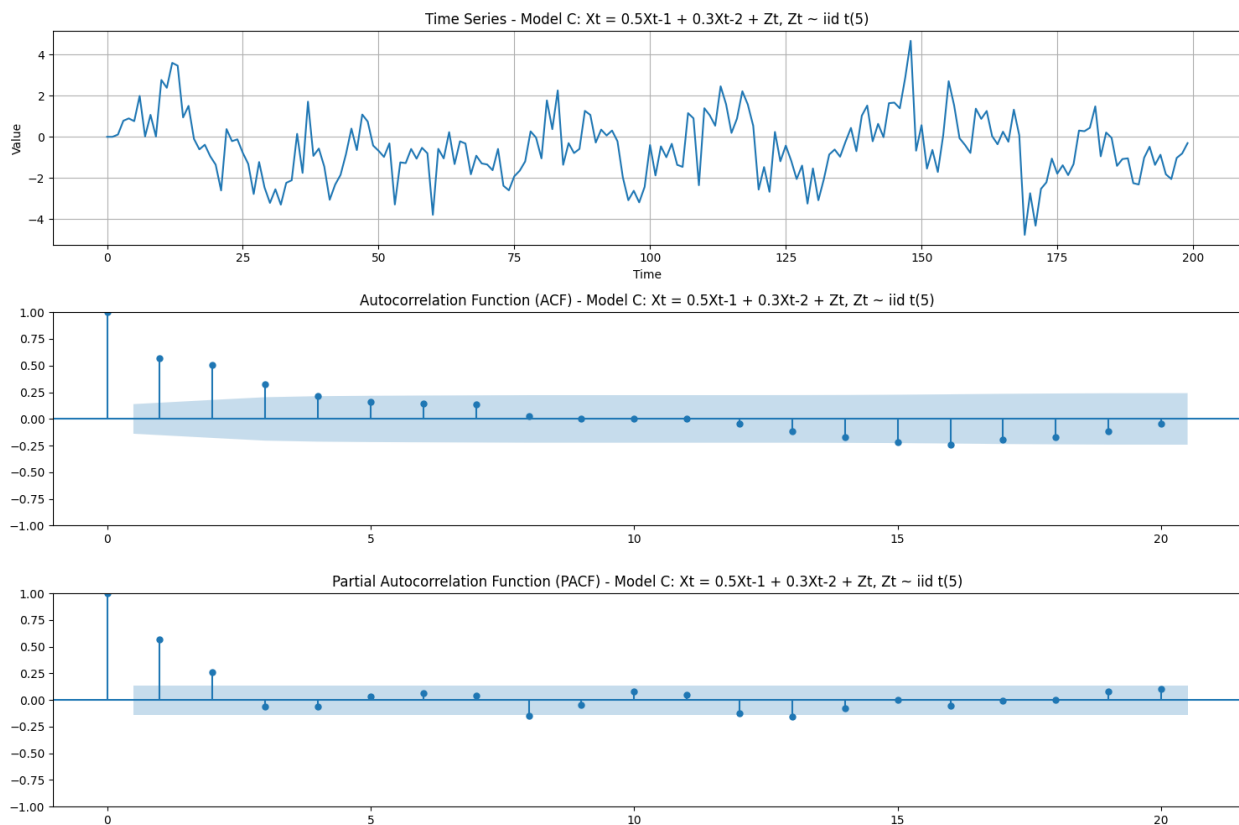
2.5%: 0.5740  
1%: 0.7390  
Stationary: True

Ljung-Box Test for White Noise:  
Q-statistic at lag 20: 211.9966  
p-value: 0.0000  
White Noise: False

Jarque-Bera Test for Normality:  
JB Statistic: 3.1610  
p-value: 0.2059  
Skewness: 0.2444  
Kurtosis: 0.3746  
Normal Distribution: True

/var/folders/7z/h\_\_cj55j6hb0wqzy9tb5b92w0000gn/T/  
ipykernel\_78328/935025886.py:78: InterpolationWarning: The test  
statistic is outside of the range of p-values available in the  
look-up table. The actual p-value is greater than the p-value  
returned.

```
kpss_result = kpss(series, regression='c')
```



```
===== Analysis for Model D:  $X_t = Z_t$ , where  $Z_t \sim \text{iid Laplace}(0, 1)$ 
=====
```

Summary Statistics:

Mean: -0.0866  
Variance: 2.0362  
Min: -4.6172  
Max: 4.9155

Augmented Dickey-Fuller Test:

ADF Statistic: -15.0773  
p-value: 0.0000  
Critical Values:  
1%: -3.4636  
5%: -2.8762  
10%: -2.5746  
Stationary: True

KPSS Test:

KPSS Statistic: 0.3466  
p-value: 0.1000  
Critical Values:  
10%: 0.3470  
5%: 0.4630  
2.5%: 0.5740  
1%: 0.7390  
Stationary: True

Ljung-Box Test for White Noise:

Q-statistic at lag 20: 13.2660  
p-value: 0.8657  
White Noise: True

Jarque-Bera Test for Normality:

JB Statistic: 11.0949  
p-value: 0.0039  
Skewness: 0.1542  
Kurtosis: 1.1119  
Normal Distribution: False

```
/var/folders/7z/h__cj55j6hb0wqzy9tb5b92w0000gn/T/
ipykernel_78328/935025886.py:78: InterpolationWarning: The test
statistic is outside of the range of p-values available in the
look-up table. The actual p-value is greater than the p-value
returned.
```

```
kpss_result = kpss(series, regression='c')
```

