

In [1]: *#Overall IPC Crime Category*

```
In [2]: import pandas as pd
import numpy as np
import statsmodels.api as sm
from statsmodels.stats.outliers_influence import variance_inflation_factor
from linearmodels.panel import PanelOLS
from linearmodels.panel import compare
```

```
In [3]: df = pd.read_csv('panel_data/new_data.csv')
df['l_ipc'] = np.log(df['ipc_cr'])
df.head()
```

```
Out[3]:
```

	s.no.	districts	year	ipc_crimes	pop_in_lak	ipc_cr	avg_temp	tot_rf	l_ipc
0	1	ariyalur	2011	1953	7.52	259.54	28.312353	1103.207404	5.558911
1	1	ariyalur	2012	2480	7.63	324.83	28.777312	973.207972	5.783302
2	1	ariyalur	2013	1991	7.76	256.53	28.730311	870.158045	5.547246
3	1	ariyalur	2014	1615	7.88	204.88	28.536042	1090.802339	5.322424
4	1	ariyalur	2015	1386	8.00	173.00	28.565911	1501.644532	5.153292

```
In [4]: df = df.set_index(['districts','year'])
y = df['l_ipc']
X = df[['avg_temp','tot_rf']]
```

```
In [5]: #PooledOLS Estimation
X = sm.add_constant(X)
pols = PanelOLS(y,X)
pols_result = pols.fit()
print(pols_result.summary)
```

PanelOLS Estimation Summary

```

=====
Dep. Variable:          l_ipc      R-squared:                0.0457
Estimator:              PanelOLS   R-squared (Between):      0.0408
No. Observations:       384        R-squared (Within):       0.0466
Date:                   Tue, Nov 11 2025  R-squared (Overall):      0.0457
Time:                   15:29:21    Log-likelihood            -265.57
Cov. Estimator:         Unadjusted

                               F-statistic:                9.1189
Entities:                32      P-value                0.0001
Avg Obs:                 12.000  Distribution:          F(2,381)
Min Obs:                 12.000
Max Obs:                 12.000  F-statistic (robust):    9.1189
                               P-value                0.0001
Time periods:           12      Distribution:          F(2,381)
Avg Obs:                 32.000
Min Obs:                 32.000
Max Obs:                 32.000

```

Parameter Estimates

```

=====
               Parameter  Std. Err.    T-stat    P-value    Lower CI    Upper CI
-----
const          4.8315      0.2463     19.618    0.0000     4.3473     5.3158
avg_temp       0.0187      0.0078      2.3838    0.0176     0.0033     0.0341
tot_rf         0.0002     5.757e-05    4.2059    0.0000     0.0001     0.0004
=====

```

```

In [6]: #FE Model Estimation
X = sm.add_constant(X)
FEmodel = PanelOLS(y,X,entity_effects=True)
feresult = FEmodel.fit()
print(feresult.summary)

```

PanelOLS Estimation Summary

```

=====
Dep. Variable:          l_ipc      R-squared:                0.0474
Estimator:              PanelOLS   R-squared (Between):      0.0275
No. Observations:       384        R-squared (Within):      0.0474
Date:                   Tue, Nov 11 2025  R-squared (Overall):     0.0444
Time:                   15:29:21    Log-likelihood            -233.43
Cov. Estimator:         Unadjusted

                               F-statistic:                8.7101
Entities:                32      P-value                0.0002
Avg Obs:                 12.000  Distribution:         F(2,350)
Min Obs:                 12.000
Max Obs:                 12.000  F-statistic (robust):   8.7101
                               P-value                0.0002
Time periods:           12      Distribution:         F(2,350)
Avg Obs:                 32.000
Min Obs:                 32.000
Max Obs:                 32.000

```

Parameter Estimates

```

=====
               Parameter  Std. Err.    T-stat    P-value    Lower CI    Upper CI
-----
const          4.6751      1.1259     4.1523    0.0000     2.4607     6.8895
avg_temp       0.0229      0.0403     0.5669    0.5712    -0.0564     0.1022
tot_rf         0.0003      7.078e-05   3.9602    0.0001     0.0001     0.0004
=====

```

F-test for Poolability: 2.0570

P-value: 0.0010

Distribution: F(31,350)

Included effects: Entity

```

In [9]: #RE Model Estimation
from linearmodels.panel import RandomEffects
import statsmodels.api as sm
X = sm.add_constant(X)
REmodel = RandomEffects(y,X)
reresult = REmodel.fit()
print(reresult.summary)

```

RandomEffects Estimation Summary

```

=====
Dep. Variable:          l_ipc      R-squared:                0.0463
Estimator:              RandomEffects  R-squared (Between):      0.0362
No. Observations:        384      R-squared (Within):       0.0471
Date:                    Tue, Nov 11 2025  R-squared (Overall):     0.0455
Time:                    15:29:22      Log-likelihood            -249.01
Cov. Estimator:          Unadjusted

                               F-statistic:                9.2495
Entities:                 32      P-value                 0.0001
Avg Obs:                  12.000  Distribution:           F(2,381)
Min Obs:                  12.000
Max Obs:                  12.000  F-statistic (robust):    9.2495
                               P-value                 0.0001
Time periods:             12      Distribution:           F(2,381)
Avg Obs:                  32.000
Min Obs:                  32.000
Max Obs:                  32.000

```

Parameter Estimates

```

=====
               Parameter  Std. Err.    T-stat    P-value    Lower CI    Upper CI
-----
const          4.7907      0.3184     15.046    0.0000     4.1646     5.4168
avg_temp        0.0195      0.0106      1.8415    0.0663    -0.0013     0.0403
tot_rf          0.0003      6.03e-05     4.2891    0.0000     0.0001     0.0004
=====

```

```

In [10]: #Hausman Test
from numpy.linalg import inv
from scipy.stats import chi2

b_FE = feresult.params
b_RE = reresult.params

common_coef = list(set(b_FE.index) & set(b_RE.index))

if 'const' in common_coef:
    common_coef.remove('const')

b_FE = b_FE[common_coef]
b_RE = b_RE[common_coef]

V_FE = feresult.cov
V_RE = reresult.cov

diff = b_FE - b_RE
diff_var = V_FE.loc[common_coef, common_coef] - V_RE.loc[common_coef, common_coef]

hausman_stat = np.dot(np.dot(diff.T, inv(diff_var)), diff)

df_h = len(diff)
p_value = 1 - chi2.cdf(hausman_stat, df_h)

print("Hausman Test Statistic:", round(hausman_stat, 3))
print("Degrees of Freedom:", df_h)

```

```
print("p-value:", round(p_value, 4))
```

Hausman Test Statistic: 0.567

Degrees of Freedom: 2

p-value: 0.753

```
In [11]: #Diagnostic Checks
from statsmodels.stats.diagnostic import het_breuschpagan, het_white
from statsmodels.stats.stattools import durbin_watson
```

```
In [12]: #Test for Heteroskedasticity

#H0: No heteroskedasticity
#H1: Heteroskedasticity exists

#p-value <= 0.05 ---> Heteroskedasticity; p-value > 0.05 ---> Homoskedasticity

print('Breusch-Pagan Test')
residuals = reresult.resids
bp_test = het_breuschpagan(residuals, X)
bp_labels = ['Lagrange multiplier statistic', 'p-value', 'f-value', 'f p-value']
print(dict(zip(bp_labels, bp_test)))
print()
print('White Test')
white_test = het_white(residuals, X)
white_labels = ['LM stat', 'LM p-value', 'F p-value']
print(dict(zip(white_labels, white_test)))
```

Breusch-Pagan Test

```
{'Lagrange multiplier statistic': np.float64(9.98334328950044), 'p-value': np.float64(0.006794297342475483), 'f-value': np.float64(5.084872190924652), 'f p-value': np.float64(0.006616473096637813)}
```

White Test

```
{'LM stat': np.float64(13.545411746797555), 'LM p-value': np.float64(0.018770041534266983), 'F p-value': np.float64(2.7642608852180715)}
```

```
In [13]: #Test for serial correlation (autocorrelation)

#Durbin-Watson statistic ranges between 0 to 4

#DW statistic = 2 ---> No autocorrelation
#DW statistic < 2 ---> Positive autocorrelation
#DW statistic > 2 ---> Negative autocorrelation

print('Durbin-Watson Test')
dw_value = durbin_watson(residuals)
print("Durbin-Watson statistic: ", round(dw_value, 3))
```

Durbin-Watson Test

Durbin-Watson statistic: 1.701

```
In [14]: from scipy import stats

#Test for cross-section dependency

#H0: No cross-section dependency
```

```

#H1: Cross-section dependency exists

print('Breusch-Pagan LM Test')
resid_df = residuals.unstack(level=0)
T = resid_df.shape[0]
N = resid_df.shape[1]

rho = resid_df.corr().values
upper_tri_idx = np.triu_indices(N, k=1)
rho_upper = rho[upper_tri_idx]
LM_stat = T * np.sum(rho_upper**2)
p_value = 1 - stats.chi2.cdf(LM_stat, N*(N-1)/2)

print(f"Breusch-Pagan LM statistic: {LM_stat:.3f}")
print(f"p-value: {p_value:.4f}")
print()

print('Pearson CD Test')
CD_stat = np.sqrt(2 / (N*(N-1))) * np.sum(rho_upper)
p_value_cd = 2 * (1 - stats.norm.cdf(abs(CD_stat)))

print(f"Pesaran CD statistic: {CD_stat:.3f}")
print(f"p-value: {p_value_cd:.4f}")

```

```

Breusch-Pagan LM Test
Breusch-Pagan LM statistic: 3117.798
p-value: 0.0000

```

```

Pearson CD Test
Pesaran CD statistic: 14.825
p-value: 0.0000

```

```
In [15]: #Re-estimate RE Model
```

```

In [16]: #RE with cov.type 'clustered'
re_model_robust1 = REmodel.fit(cov_type='clustered', cluster_entity=True)
print(re_model_robust1.summary)

```

RandomEffects Estimation Summary

```

=====
Dep. Variable:          l_ipc      R-squared:                0.0463
Estimator:              RandomEffects  R-squared (Between):      0.0362
No. Observations:       384      R-squared (Within):       0.0471
Date:                   Tue, Nov 11 2025  R-squared (Overall):     0.0455
Time:                   15:29:22      Log-likelihood            -249.01
Cov. Estimator:         Clustered

                               F-statistic:                9.2495
                               P-value                    0.0001
Entities:                32      Distribution:          F(2,381)
Avg Obs:                 12.000
Min Obs:                 12.000
Max Obs:                 12.000
                               F-statistic (robust):        15.015
                               P-value                    0.0000
Time periods:            12      Distribution:          F(2,381)
Avg Obs:                 32.000
Min Obs:                 32.000
Max Obs:                 32.000

```

Parameter Estimates

```

=====
               Parameter  Std. Err.    T-stat    P-value    Lower CI    Upper CI
-----
const          4.7907      0.4626     10.357    0.0000     3.8812     5.7002
avg_temp       0.0195      0.0160      1.2148    0.2252    -0.0121     0.0510
tot_rf         0.0003      4.723e-05    5.4757    0.0000     0.0002     0.0004
=====

```

```

In [17]: #RE with cov.type 'kernel' (Driscoll-Kraay Method)
re_model_robust2 = REmodel.fit(cov_type='kernel')
print(re_model_robust2.summary)

```

RandomEffects Estimation Summary

```

=====
Dep. Variable:          l_ipc      R-squared:                0.0463
Estimator:              RandomEffects  R-squared (Between):      0.0362
No. Observations:       384      R-squared (Within):       0.0471
Date:                   Tue, Nov 11 2025  R-squared (Overall):     0.0455
Time:                   15:29:22      Log-likelihood            -249.01
Cov. Estimator:         Driscoll-Kraay

                               F-statistic:                9.2495
                               P-value                    0.0001
                               Distribution:                F(2,381)

Entities:                32
Avg Obs:                 12.000
Min Obs:                 12.000
Max Obs:                 12.000
                               F-statistic (robust):        4.6172
                               P-value                    0.0104
                               Distribution:                F(2,381)

Time periods:            12
Avg Obs:                 32.000
Min Obs:                 32.000
Max Obs:                 32.000

```

Parameter Estimates

```

=====
               Parameter  Std. Err.    T-stat    P-value    Lower CI    Upper CI
-----
const          4.7907      0.2095     22.865    0.0000      4.3787      5.2027
avg_temp       0.0195      0.0064      3.0323    0.0026      0.0068      0.0321
tot_rf         0.0003      0.0001      2.1170    0.0349     1.842e-05     0.0005
=====

```

```

In [18]: # Check residuals and fitted values
df['residuals1'] = re_model_robust1.resids
df['fitted1'] = re_model_robust1.fitted_values

import matplotlib.pyplot as plt

plt.scatter(df['fitted1'], df['residuals1'], alpha=0.6)
plt.axhline(0, color='red', linestyle='--')
plt.xlabel('Fitted Values')
plt.ylabel('Residuals')
plt.title('Residuals vs Fitted Values (RE model with Clustered Standard Errors)')
plt.show()

sm.qqplot(df['residuals1'], line='45', fit=True)
plt.title('Q-Q Plot of Residuals')
plt.show()

plt.hist(df['residuals1'], bins=30, edgecolor='black', alpha=0.7)
plt.xlabel('Residuals')
plt.ylabel('Frequency')
plt.title('Distribution of Residuals (RE model)')
plt.show()

resid_df = df['residuals1'].unstack(level=0)
plt.plot(resid_df.mean(axis=1))
plt.title('Average Residuals over Time')
plt.xlabel('Year')
plt.ylabel('Mean Residual')

```

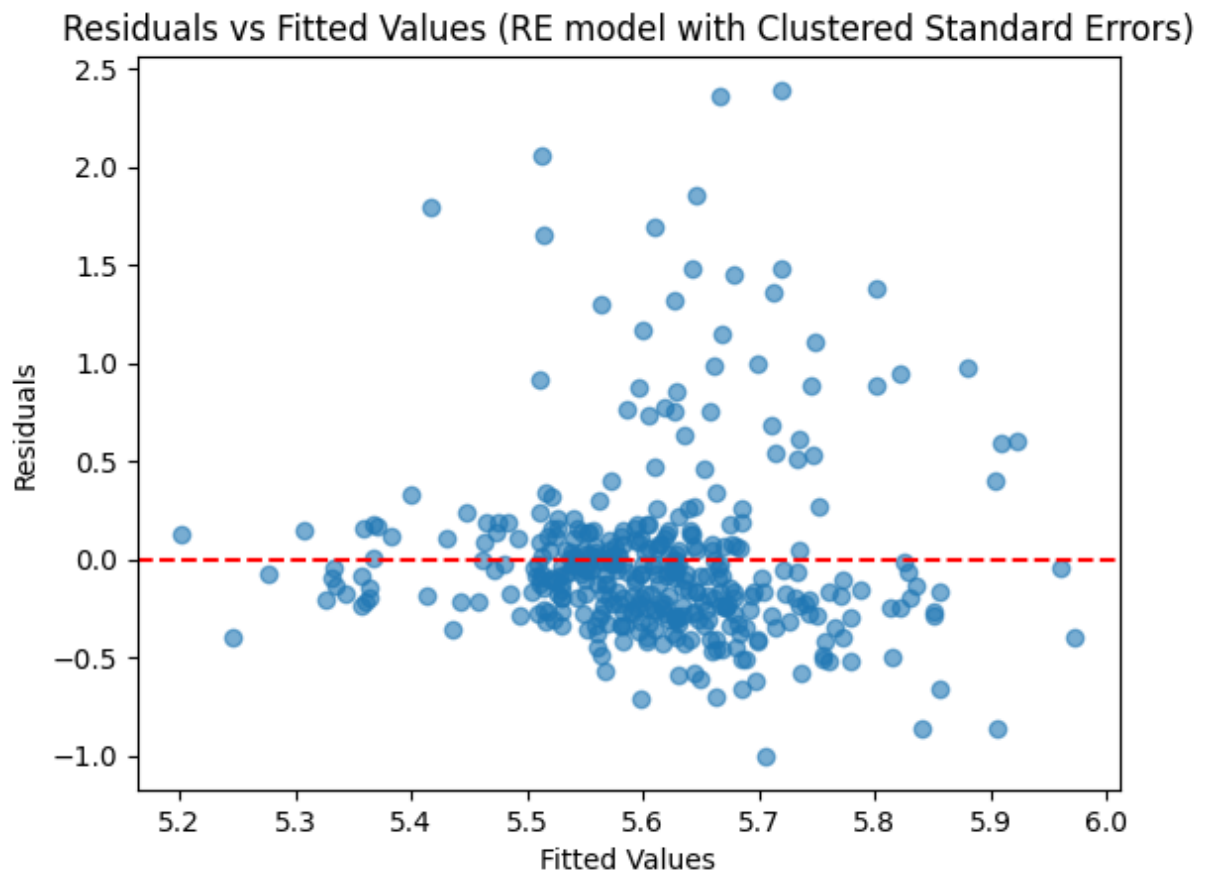


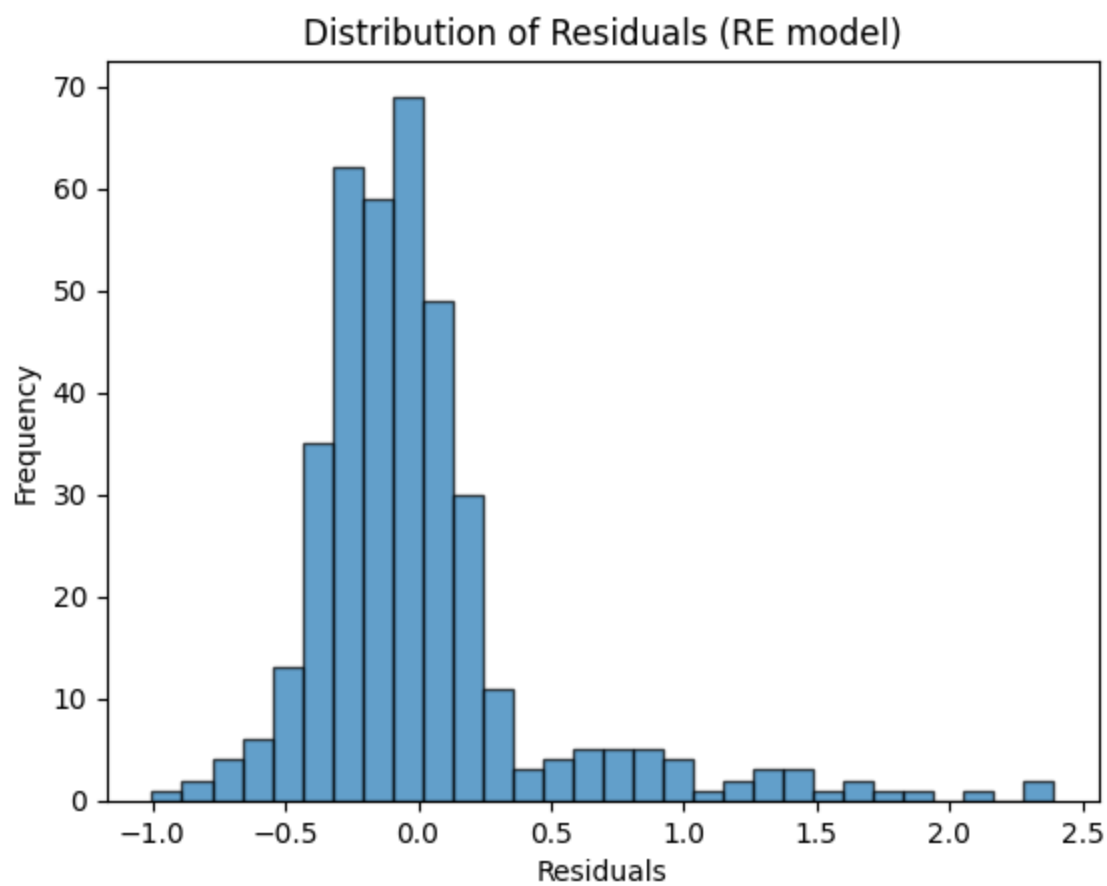
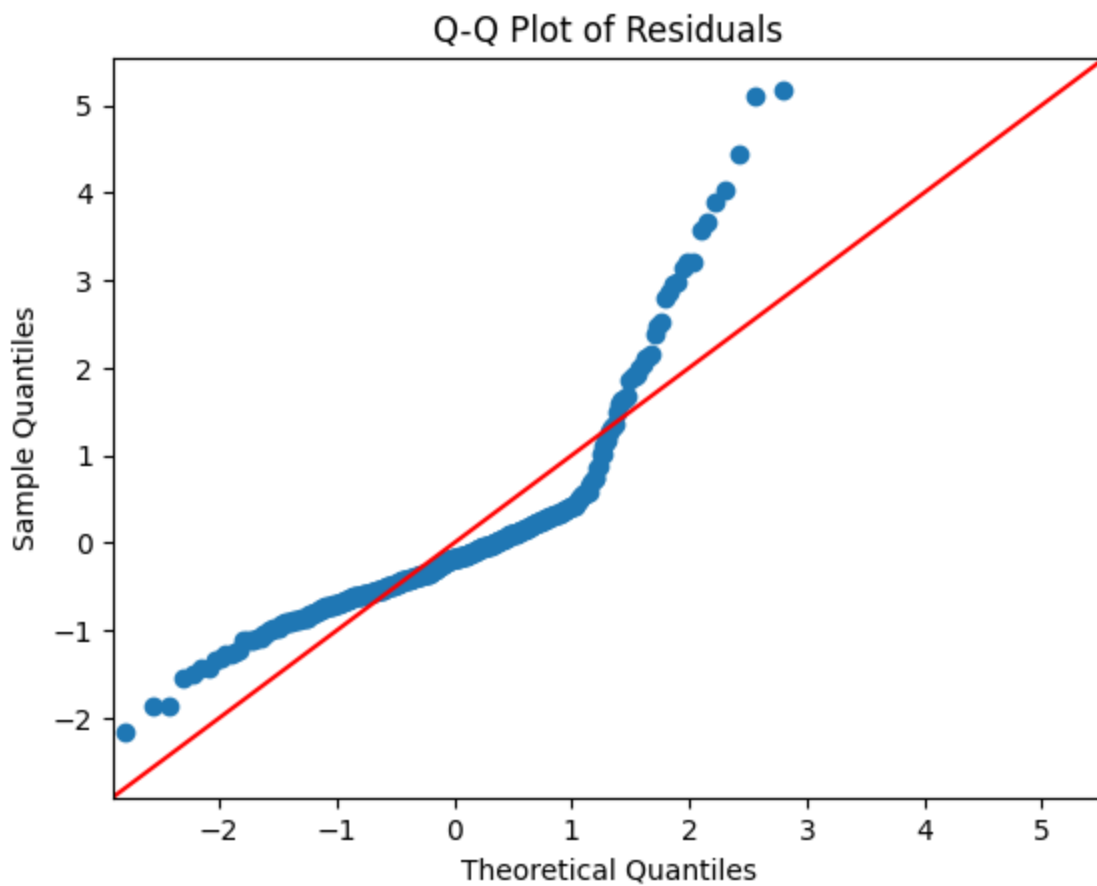
```
plt.show()

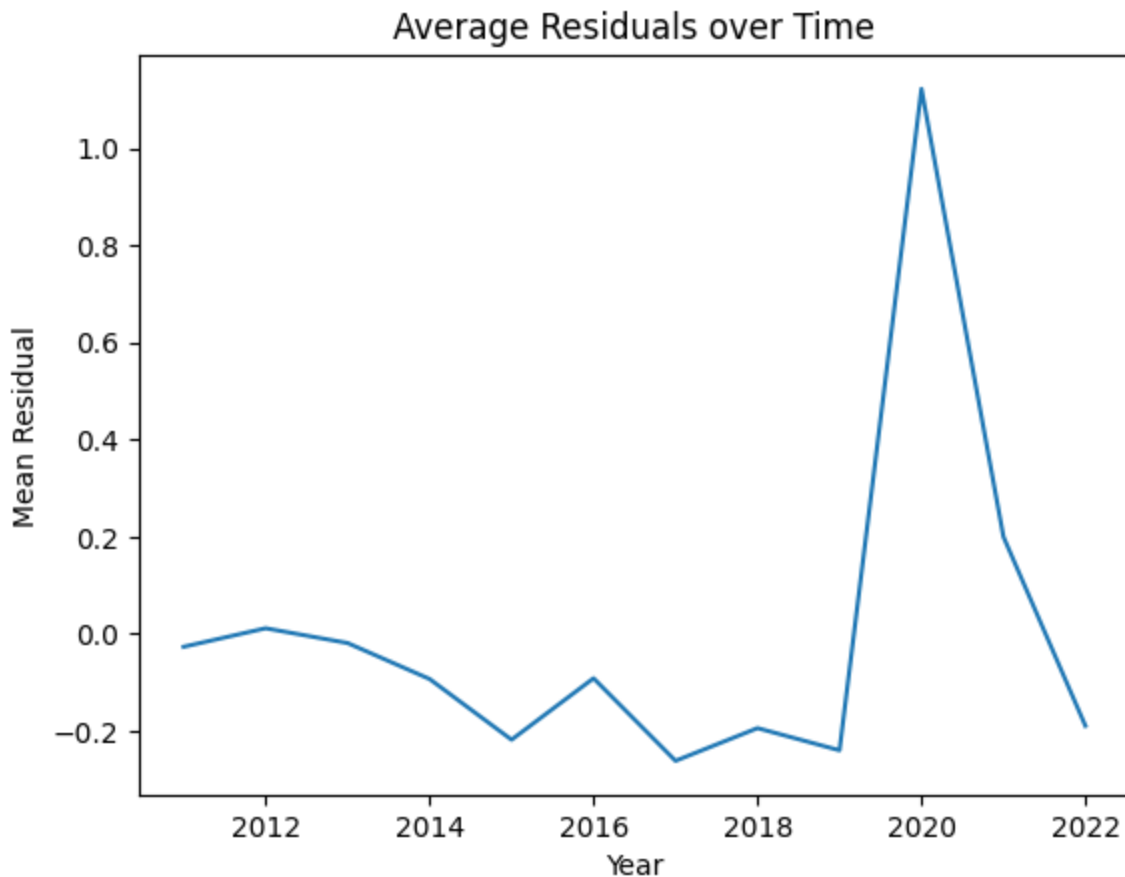
from scipy.stats import shapiro

#Test for normality

stat, p = shapiro(df['residuals1'])
print(f"Shapiro-Wilk Test: Statistic={stat:.3f}, p-value={p:.4f}")
```







Shapiro-Wilk Test: Statistic=0.797, p-value=0.0000

```
In [19]: # Check residuals and fitted values
df['residuals2'] = re_model_robust2.resids
df['fitted2'] = re_model_robust2.fitted_values

import matplotlib.pyplot as plt

plt.scatter(df['fitted2'], df['residuals2'], alpha=0.6)
plt.axhline(0, color='red', linestyle='--')
plt.xlabel('Fitted Values')
plt.ylabel('Residuals')
plt.title('Residuals vs Fitted Values (RE model with Driscoll-Kraay)')
plt.show()

sm.qqplot(df['residuals2'], line='45', fit=True)
plt.title('Q-Q Plot of Residuals')
plt.show()

plt.hist(df['residuals2'], bins=30, edgecolor='black', alpha=0.7)
plt.xlabel('Residuals')
plt.ylabel('Frequency')
plt.title('Distribution of Residuals (RE model)')
plt.show()

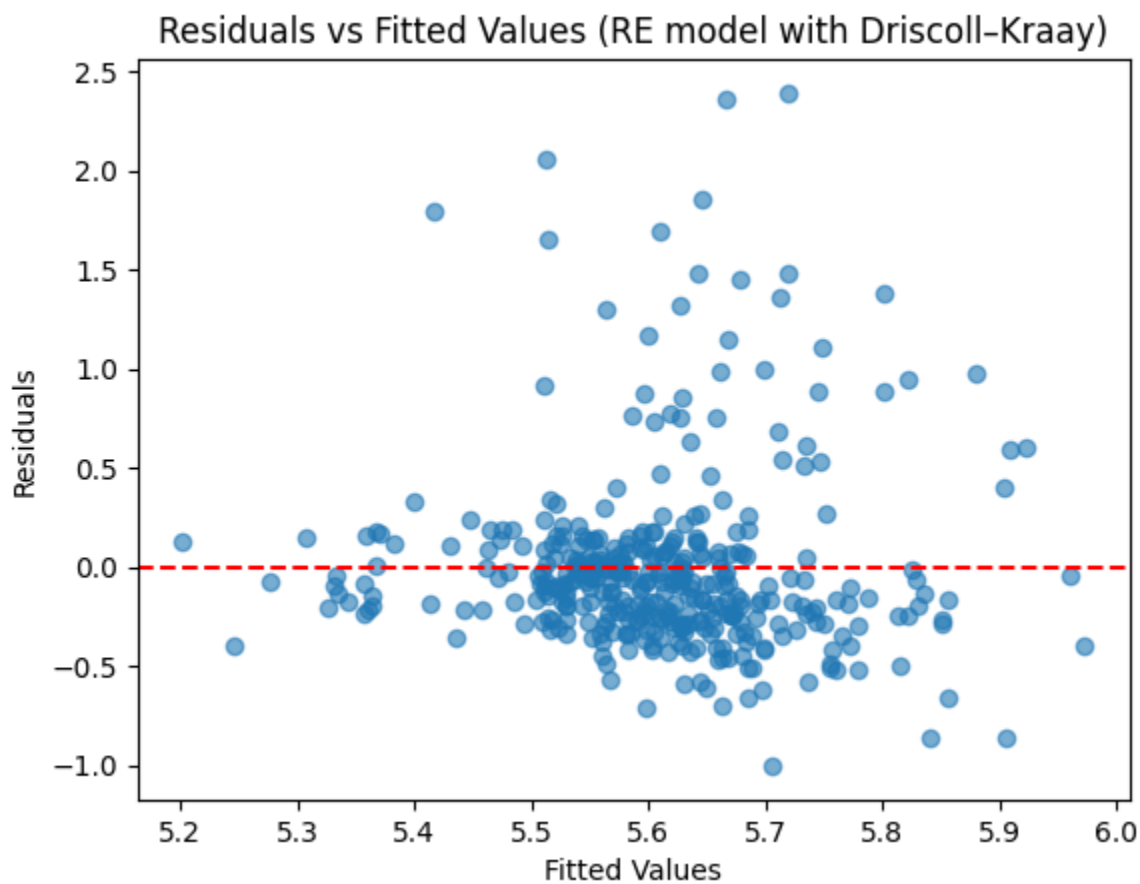
resid_df = df['residuals2'].unstack(level=0)
plt.plot(resid_df.mean(axis=1))
plt.title('Average Residuals over Time')
plt.xlabel('Year')
```

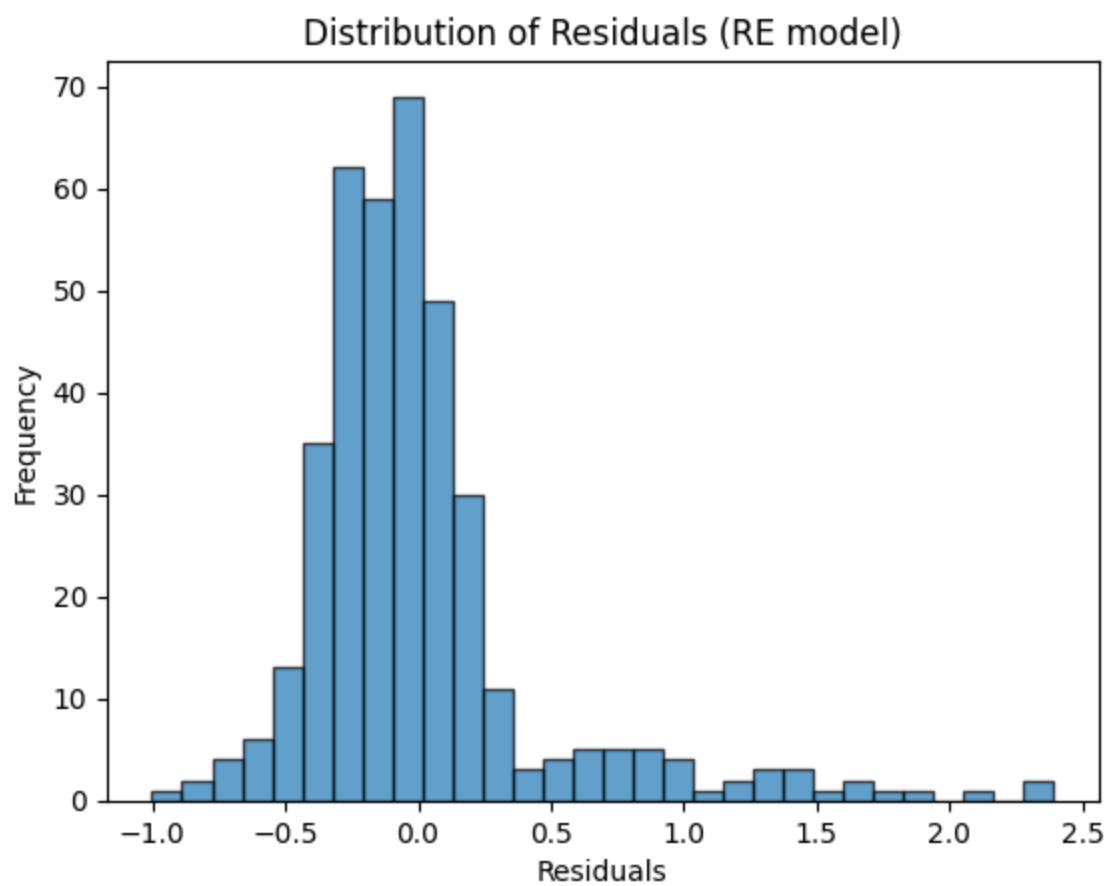
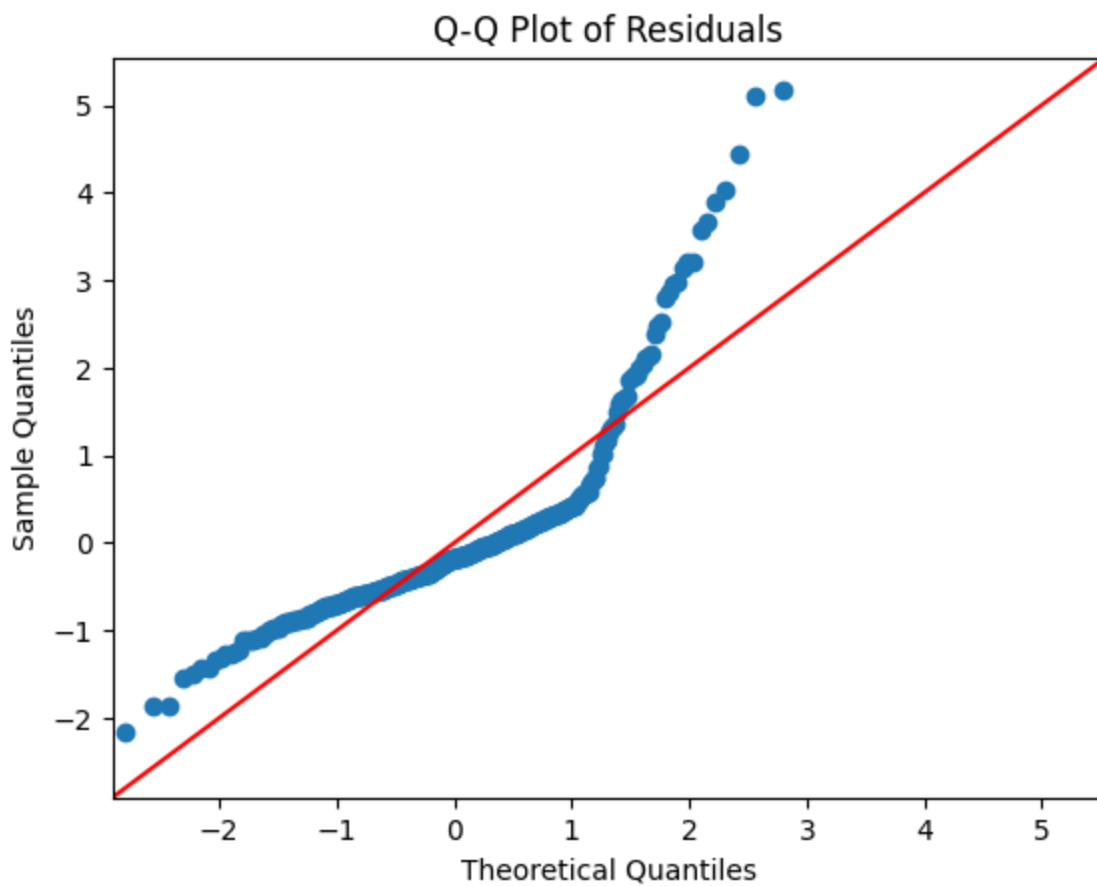
```
plt.ylabel('Mean Residual')
plt.show()

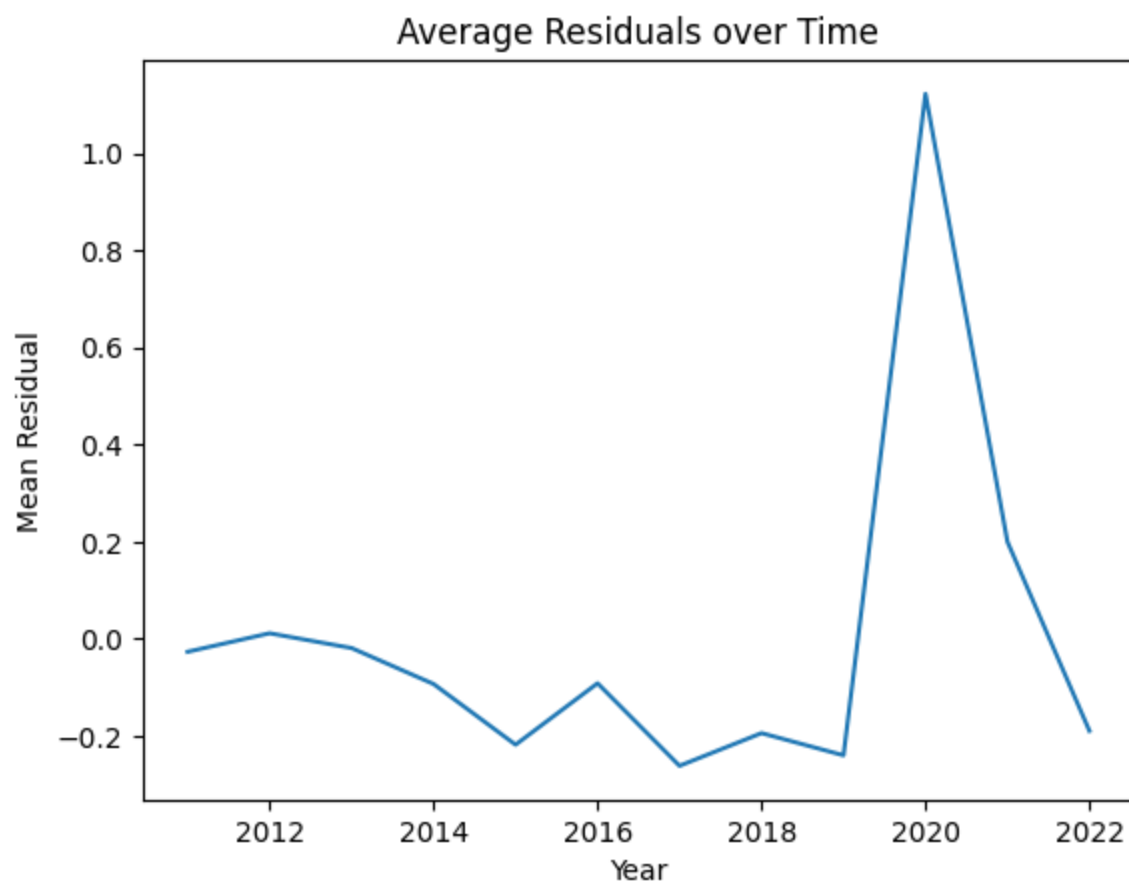
from scipy.stats import shapiro

#Test for normality

stat, p = shapiro(df['residuals2'])
print(f"Shapiro-Wilk Test: Statistic={stat:.3f}, p-value={p:.4f}")
```







Shapiro-Wilk Test: Statistic=0.797, p-value=0.0000

In []: