

```
In [1]: #Crimes Against Women 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/crimes against women new.csv')
#df['L_ipc'] = np.log(df['ipc_cr'])
df.head()
```

```
Out[3]:
```

	s.no.	districts	year	type	crimes_against_women	pop_in_lak	craw_cr	avg_temp	
0	1	ariyalur	2011	crimes against women	99	7.52	13.2	28.312353	11
1	1	ariyalur	2012	crimes against women	117	7.63	15.3	28.777312	9
2	1	ariyalur	2013	crime against women	63	7.76	8.1	28.730311	8
3	1	ariyalur	2014	crime against women	61	7.88	7.7	28.536042	10
4	1	ariyalur	2015	crime against women	32	8.00	4.0	28.565911	15

```
In [4]: df = df.set_index(['districts', 'year'])
y = df['craw_cr']
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:          craw_cr    R-squared:                 0.0272
Estimator:              PanelOLS   R-squared (Between):       0.0752
No. Observations:       384        R-squared (Within):        0.0011
Date:                   Wed, Nov 12 2025  R-squared (Overall):       0.0272
Time:                   18:22:47    Log-likelihood             -1063.2
Cov. Estimator:         Unadjusted

                               F-statistic:                 5.3354
Entities:                32        P-value                  0.0052
Avg Obs:                 12.000    Distribution:            F(2,381)
Min Obs:                 12.000
Max Obs:                 12.000    F-statistic (robust):     5.3354
                               P-value                  0.0052
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          10.588      1.9657     5.3863    0.0000     6.7230     14.453
avg_temp       -0.0992     0.0625    -1.5877    0.1132    -0.2220     0.0237
tot_rf         -0.0015     0.0005    -3.2535    0.0012    -0.0024    -0.0006
=====

```

```

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:          craw_cr    R-squared:                 0.0083
Estimator:              PanelOLS   R-squared (Between):       0.0043
No. Observations:       384        R-squared (Within):        0.0083
Date:                   Wed, Nov 12 2025  R-squared (Overall):       0.0069
Time:                   18:22:47    Log-likelihood             -983.39
Cov. Estimator:         Unadjusted

                               F-statistic:                 1.4639
                               P-value                     0.2327
Entities:                32        Distribution:           F(2,350)
Avg Obs:                  12.000
Min Obs:                  12.000
Max Obs:                  12.000
                               F-statistic (robust):         1.4639
                               P-value                     0.2327
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          5.0647      7.9375     0.6381    0.5238    -10.546     20.676
avg_temp       0.0722      0.2843     0.2540    0.7996    -0.4869     0.6313
tot_rf        -0.0007      0.0005    -1.4059    0.1606    -0.0017     0.0003
=====

```

F-test for Poolability: 5.8183

P-value: 0.0000

Distribution: F(31,350)

Included effects: Entity

```

In [7]: #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:	craw_cr	R-squared:	0.0112
Estimator:	RandomEffects	R-squared (Between):	0.0526
No. Observations:	384	R-squared (Within):	0.0075
Date:	Wed, Nov 12 2025	R-squared (Overall):	0.0234
Time:	18:22:47	Log-likelihood	-999.66
Cov. Estimator:	Unadjusted		
		F-statistic:	2.1663
Entities:	32	P-value	0.1160
Avg Obs:	12.000	Distribution:	F(2,381)
Min Obs:	12.000		
Max Obs:	12.000	F-statistic (robust):	2.1663
		P-value	0.1160
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	8.7975	3.3428	2.6318	0.0088	2.2248	15.370
avg_temp	-0.0562	0.1153	-0.4875	0.6262	-0.2829	0.1705
tot_rf	-0.0009	0.0005	-2.0703	0.0391	-0.0018	-4.717e-05
=====						

```
In [8]: #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: 1.947

Degrees of Freedom: 2

p-value: 0.3777

```
In [9]: #Diagnostic Checks
from statsmodels.stats.diagnostic import het_breuschpagan, het_white
from statsmodels.stats.stattools import durbin_watson
```

```
In [10]: #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(10.555337591621296), 'p-value': np.float64(0.005104316136896624), 'f-value': np.float64(5.384443837638701), 'f p-value': np.float64(0.004943148107568016)}
```

White Test

```
{'LM stat': np.float64(11.496469341991258), 'LM p-value': np.float64(0.042378147203781594), 'F p-value': np.float64(2.333221058923278)}
```

```
In [11]: #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: 1.096

```
In [12]: 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('Pesaran 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: 1764.881
p-value: 0.0000

```

```

Pearson CD Test
Pesaran CD statistic: 10.167
p-value: 0.0000

```

```
In [13]: #Re-estimate RE Model
```

```
In [14]: #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:          craw_cr    R-squared:                0.0112
Estimator:              RandomEffects  R-squared (Between):      0.0526
No. Observations:       384         R-squared (Within):       0.0075
Date:                   Wed, Nov 12 2025  R-squared (Overall):      0.0234
Time:                   18:22:47         Log-likelihood            -999.66
Cov. Estimator:         Clustered

                               F-statistic:                2.1663
                               P-value                    0.1160
Entities:                32         Distribution:          F(2,381)
Avg Obs:                 12.000
Min Obs:                 12.000
Max Obs:                 12.000
                               F-statistic (robust):         2.9307
                               P-value                    0.0546
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          8.7975      3.3391     2.6347    0.0088     2.2321     15.363
avg_temp      -0.0562      0.1164    -0.4829    0.6294    -0.2850     0.1726
tot_rf        -0.0009      0.0004    -2.3546    0.0191    -0.0017    -0.0002
=====

```

```

In [15]: #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:          craw_cr    R-squared:                 0.0112
Estimator:              RandomEffects  R-squared (Between):      0.0526
No. Observations:      384          R-squared (Within):       0.0075
Date:                  Wed, Nov 12 2025  R-squared (Overall):     0.0234
Time:                  18:22:48          Log-likelihood            -999.66
Cov. Estimator:        Driscoll-Kraay

                               F-statistic:          2.1663
                               P-value                0.1160
                               Distribution:           F(2,381)

Avg Obs:                12.000
Min Obs:                12.000
Max Obs:                12.000

                               F-statistic (robust):    0.8908
                               P-value                0.4112
                               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          8.7975      4.0931     2.1493    0.0322     0.7496     16.845
avg_temp      -0.0562      0.0718    -0.7825    0.4344    -0.1974     0.0850
tot_rf        -0.0009      0.0008    -1.2011    0.2304    -0.0025     0.0006
=====

```

```

In [16]: # 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')

```

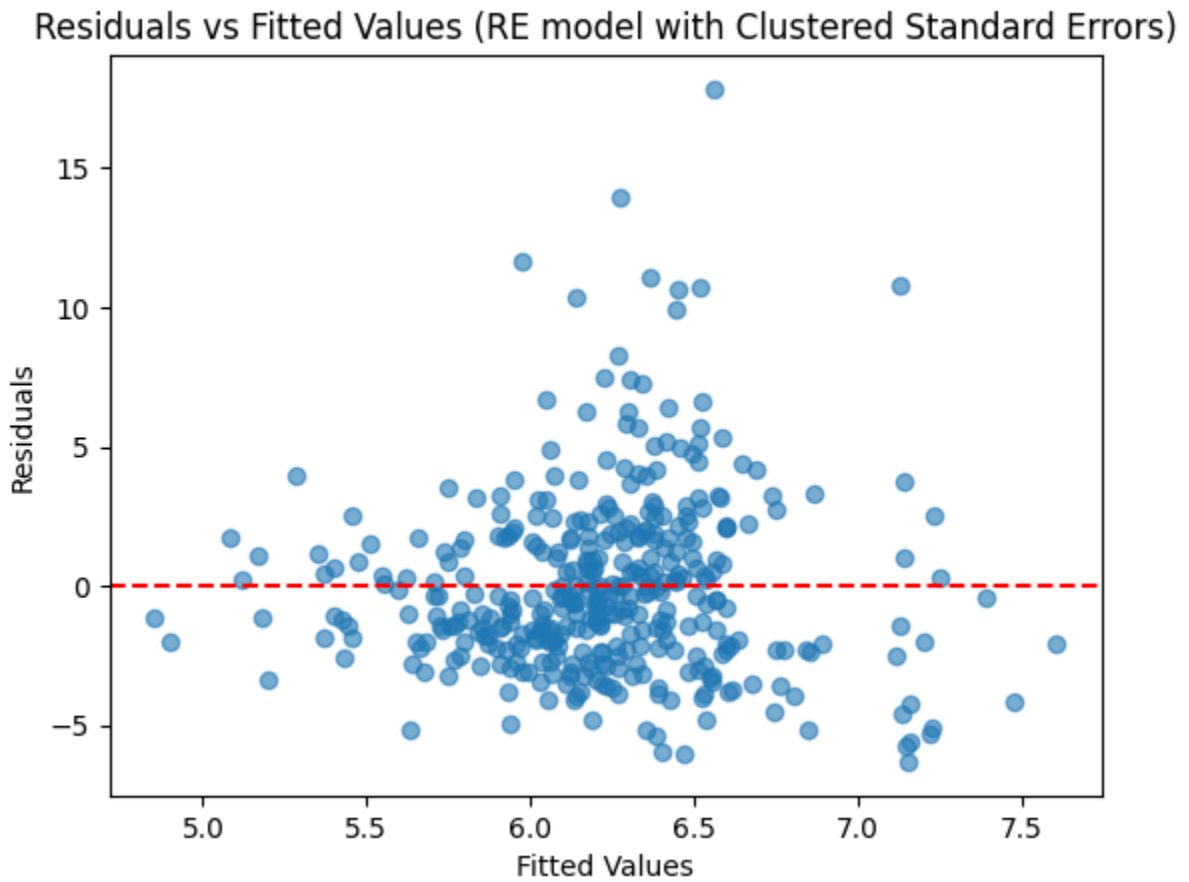


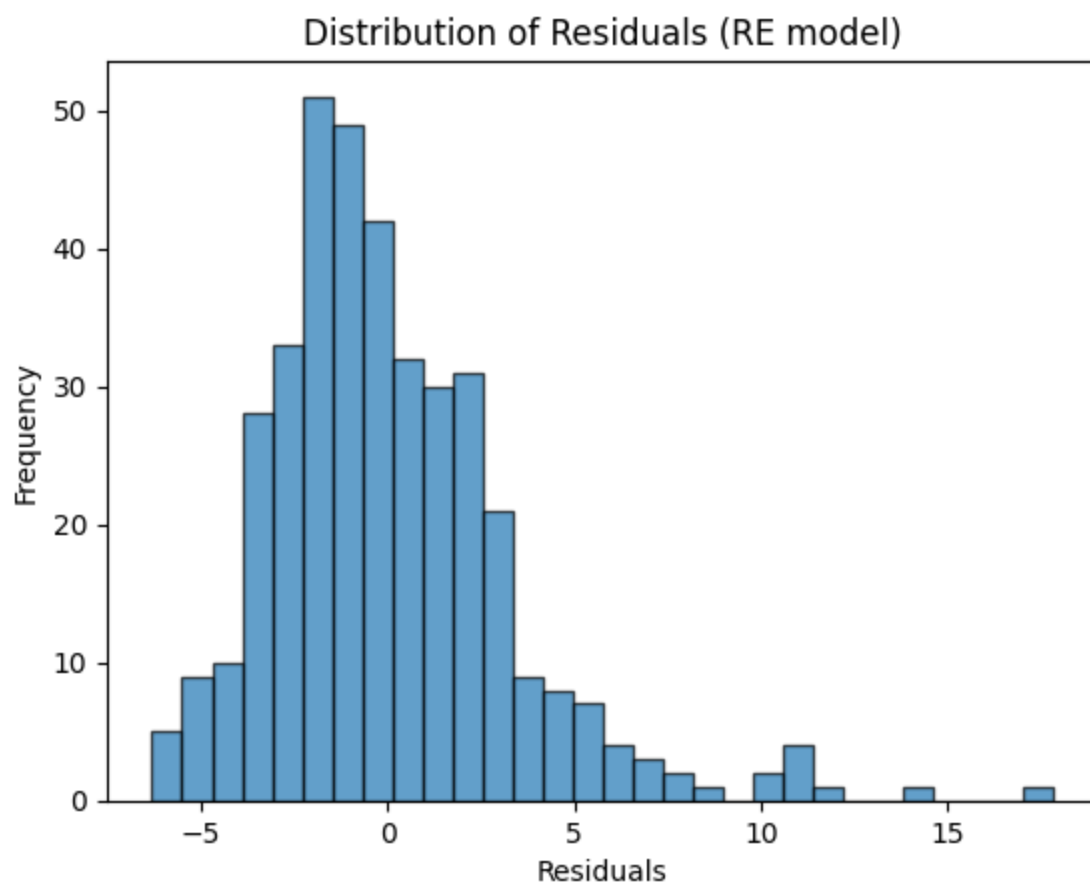
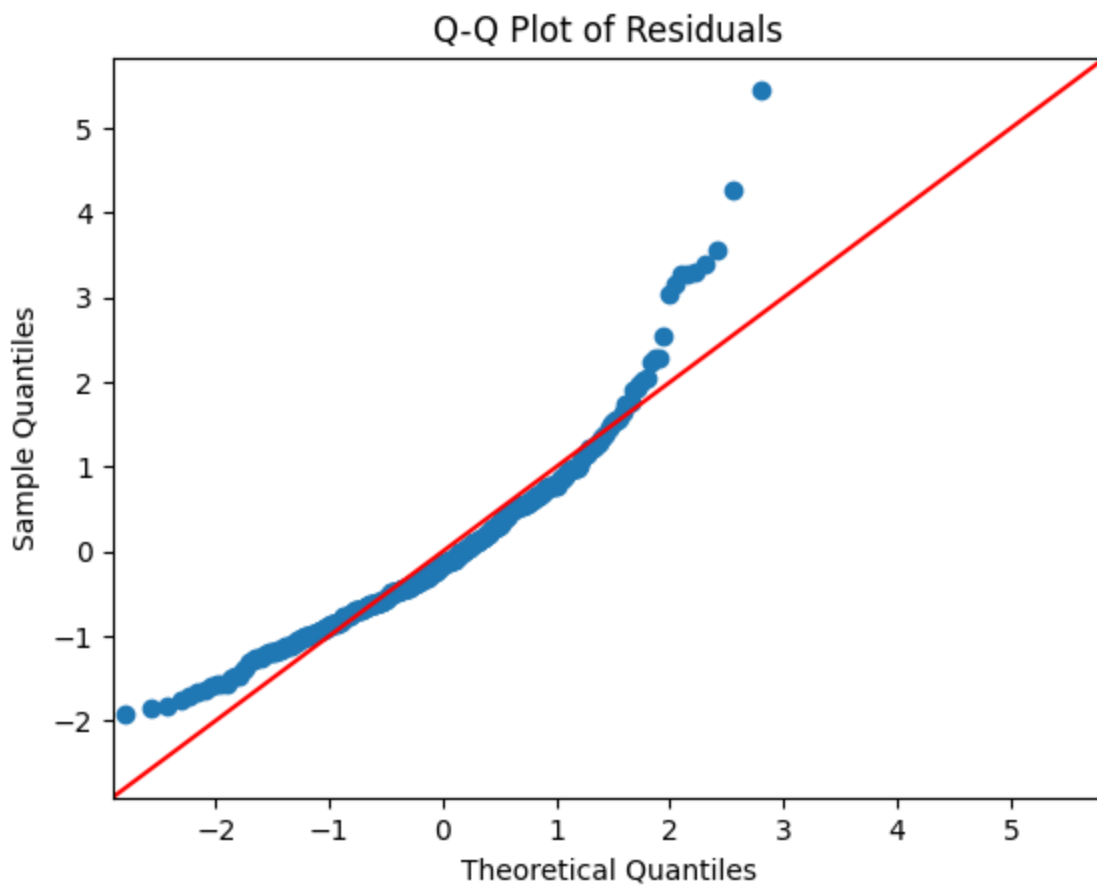
```
plt.show()

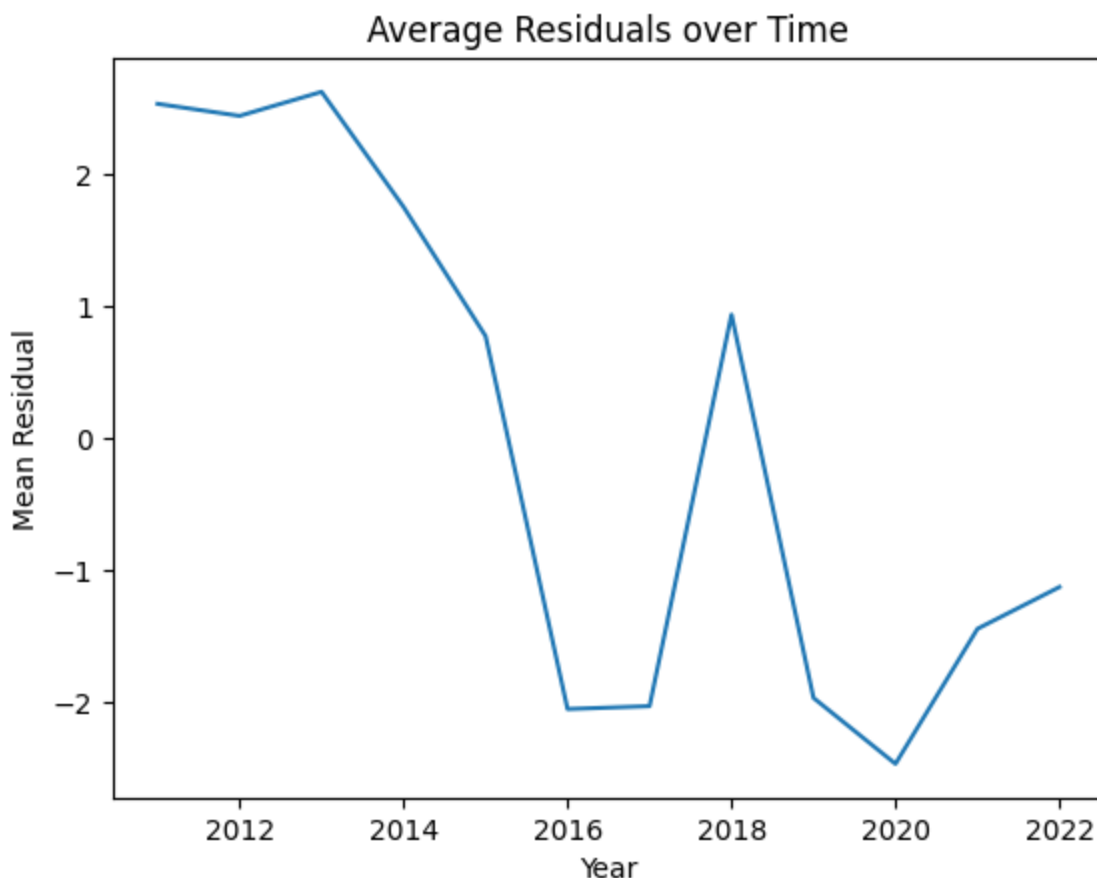
from scipy.stats import shapiro

#Test for normality

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







Shapiro-Wilk Test: Statistic=0.921, p-value=0.0000

```
In [17]: # 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()

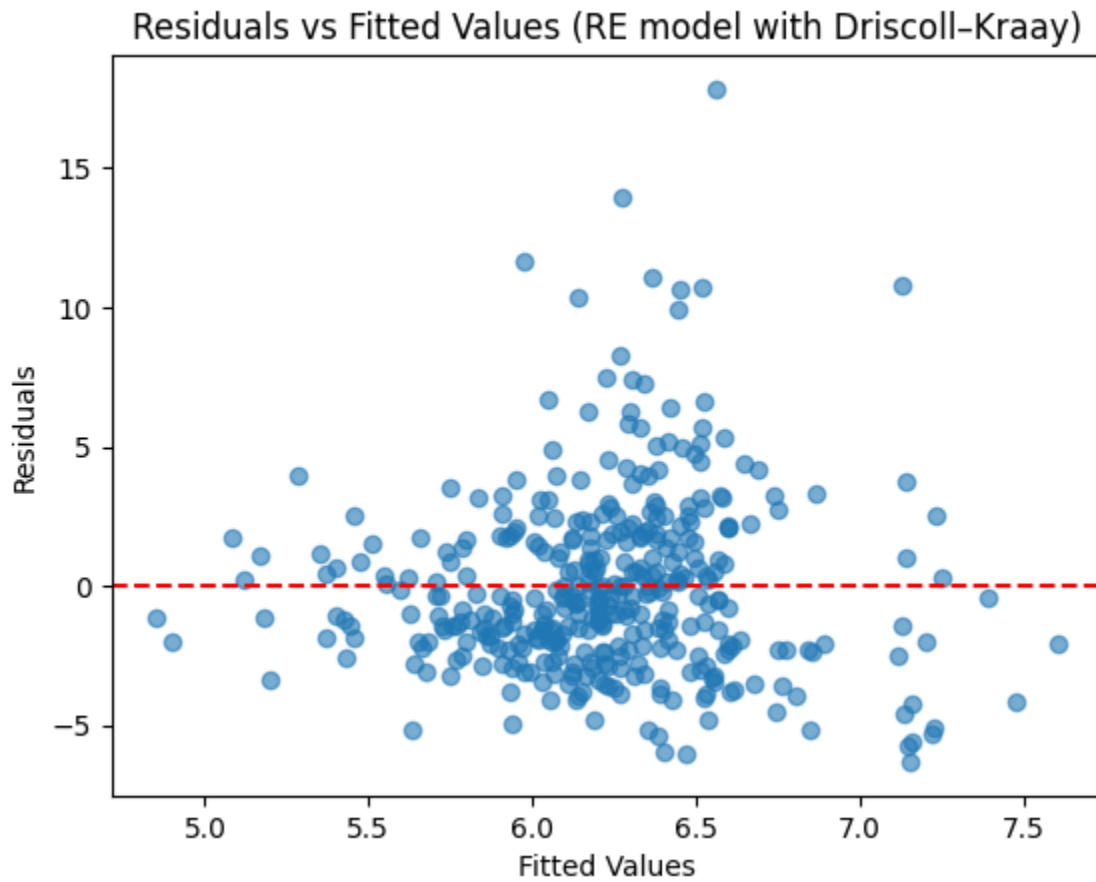
resid_df = df['residuals2'].unstack(level=0)
plt.plot(resid_df.mean(axis=1))
plt.title('Average Residuals over Time')
plt.xlabel('Year')
```

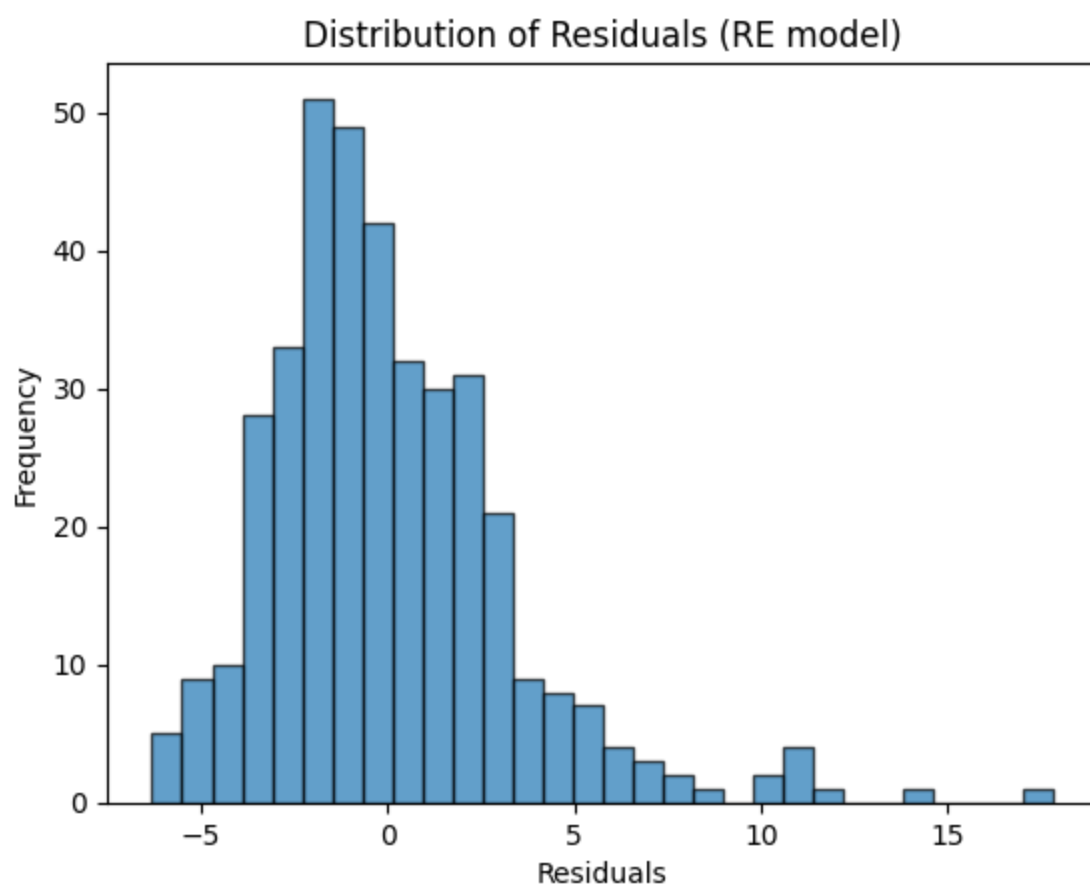
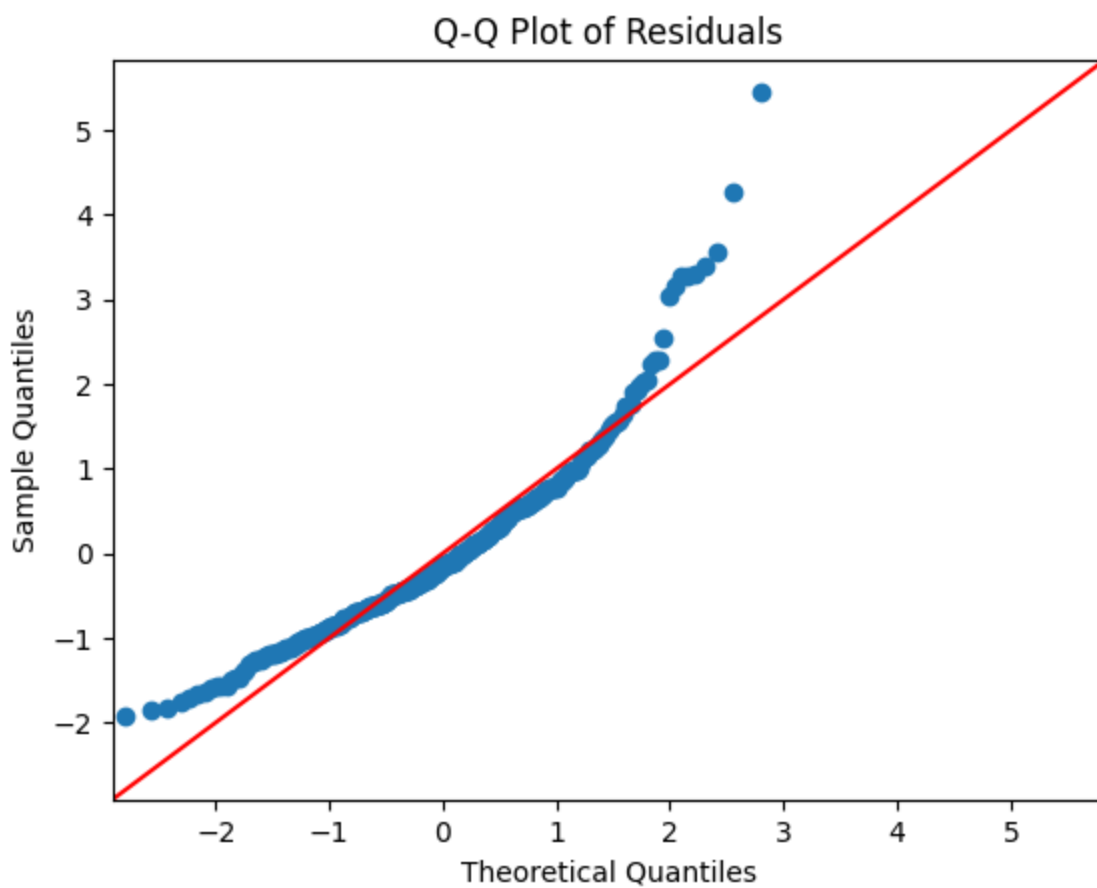
```
plt.ylabel('Mean Residual')
plt.show()

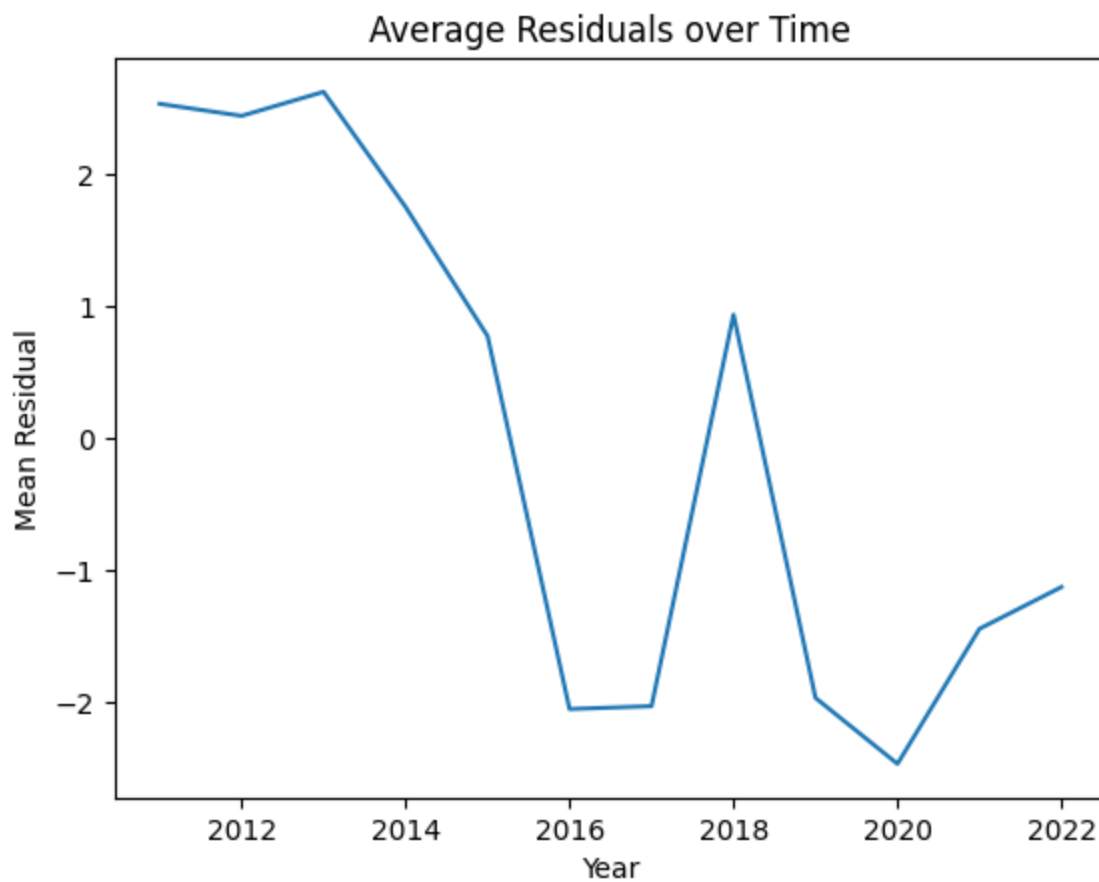
from scipy.stats import shapiro

#Test for normality

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







Shapiro-Wilk Test: Statistic=0.921, p-value=0.0000

In []: