

```
In [11]: import pandas as pd
import matplotlib.pyplot as plt
import statsmodels.formula.api as smf
import numpy as np
import wooldridge as woo
import seaborn as sns
import statsmodels.api as sm
```

```
In [96]: df = pd.read_excel("430_Data_Cleaned.xlsx", index_col = "Country")
df
```

```
Out[96]:
```

	life_expectancy	adult_mortality	infant_deaths	alcohol	percentage_expenditure	hepatitis_b	bmi	under_five_deaths	polio	total_
Country										
Afghanistan	59.5	272	69	0.01	78.184215	67.0	17.6	93	67	
Albania	76.9	86	0	5.14	412.443356	99.0	55.8	1	99	
Algeria	75.1	113	21	0.66	555.926083	95.0	56.1	24	95	
Angola	56.0	358	72	8.24	256.122524	75.0	21.5	110	75	
Antigua and Barbuda	75.9	134	0	8.18	2156.229842	98.0	45.7	0	97	
...
Venezuela (Bolivarian Republic of)	73.7	161	9	6.70	0.000000	81.0	6.4	10	73	
Viet Nam	75.6	13	29	4.12	0.000000	97.0	15.3	36	97	
Yemen	64.7	236	36	0.04	0.000000	67.0	38.7	46	68	
Zambia	59.2	349	29	2.59	196.915250	78.0	21.7	43	7	
Zimbabwe	56.6	429	26	6.09	92.602336	97.0	3.3	39	95	

183 rows × 14 columns

```
In [45]: col = list(df.columns)
col
```

```
Out[45]: ['life_expectancy',
          'adult_mortality',
          'infant_deaths',
          'alcohol',
          'percentage_expenditure',
          'hepatitis_b',
          'bmi',
          'under-five_deaths',
          'polio',
          'total_expenditure',
          'diphtheria',
          'hiv/aids',
          'gdp',
          'schooling']
```

```
In [26]: df.isnull().any()
```

```
Out[26]: life_expectancy      False
adult_mortality      False
infant_deaths        False
alcohol              True
percentage_expenditure False
hepatitis_b          True
bmi                  True
under-five_deaths    False
polio                False
total_expenditure     True
diphtheria           False
hiv/aids             False
gdp                  True
schooling            True
dtype: bool
```

```
In [46]: df.life_expectancy
```

```
Out[46]: Country
Afghanistan      59.5
Albania          76.9
Algeria          75.1
Angola           56.0
Antigua and Barbuda 75.9
...
Venezuela (Bolivarian Republic of) 73.7
Viet Nam        75.6
Yemen           64.7
Zambia          59.2
Zimbabwe        56.6
Name: life_expectancy, Length: 183, dtype: float64
```

```
In [27]: df.isnull().sum()
```

```
Out[27]: life_expectancy      0
adult_mortality      0
infant_deaths        0
alcohol              1
percentage_expenditure  0
hepatitis_b         13
bmi                  2
under-five_deaths    0
polio                0
total_expenditure     2
diphtheria           0
hiv/aids             0
gdp                  29
schooling            10
dtype: int64
```

In [28]: `df.describe()`

Out[28]:

	life_expectancy	adult_mortality	infant_deaths	alcohol	percentage_expenditure	hepatitis_b	bmi	under-five_deaths	polio	1
count	183.000000	183.000000	183.000000	182.000000	183.000000	170.000000	181.000000	183.000000	183.000000	
mean	70.916940	148.792350	26.338798	4.138187	1011.471726	82.635294	40.099448	35.562842	83.939891	
std	8.562151	108.617944	96.811105	4.272119	2273.787938	24.861199	20.775655	127.826312	24.120387	
min	49.700000	2.000000	0.000000	0.010000	0.000000	2.000000	2.100000	0.000000	3.000000	
25%	64.500000	66.500000	0.000000	0.010000	32.597328	81.000000	21.900000	0.000000	83.000000	
50%	73.200000	138.000000	3.000000	2.890000	196.915250	94.000000	45.900000	3.000000	94.000000	
75%	76.500000	215.500000	19.500000	7.785000	847.322649	97.000000	58.500000	23.500000	98.000000	
max	88.000000	513.000000	1100.000000	16.350000	18379.329740	99.000000	76.200000	1400.000000	99.000000	

In [29]: `# do we need harvey collier?`

In [18]: `pip install rpy2`

```
Requirement already satisfied: rpy2 in /Users/rheasethi/opt/anaconda3/lib/python3.9/site-packages (3.5.5)
Requirement already satisfied: Jinja2 in /Users/rheasethi/opt/anaconda3/lib/python3.9/site-packages (from rpy2) (2.11.3)
Requirement already satisfied: Pytz in /Users/rheasethi/opt/anaconda3/lib/python3.9/site-packages (from rpy2) (2021.3)
Requirement already satisfied: tzlocal in /Users/rheasethi/opt/anaconda3/lib/python3.9/site-packages (from rpy2) (4.2)
Requirement already satisfied: cffi>=1.10.0 in /Users/rheasethi/opt/anaconda3/lib/python3.9/site-packages (from rpy2) (1.15.0)
Requirement already satisfied: pycparser in /Users/rheasethi/opt/anaconda3/lib/python3.9/site-packages (from cffi>=1.10.0->rpy2) (2.21)
Requirement already satisfied: MarkupSafe>=0.23 in /Users/rheasethi/opt/anaconda3/lib/python3.9/site-packages (from Jinja2->rpy2) (2.0.1)
Requirement already satisfied: pytz-deprecation-shim in /Users/rheasethi/opt/anaconda3/lib/python3.9/site-packages (from tzlocal->rpy2) (0.1.0.post0)
Requirement already satisfied: tzdata in /Users/rheasethi/opt/anaconda3/lib/python3.9/site-packages (from pytz-deprecation-shim->tzlocal->rpy2) (2022.5)
Note: you may need to restart the kernel to use updated packages.
```

In [19]: `import rpy2`

```
import warnings
warnings.filterwarnings('ignore')

from rpy2.robjects import pandas2ri
import rpy2.rinterface as rinterface
pandas2ri.activate()

%load_ext rpy2.ipython
```

```
In [ ]: # adult_mortality + infant_deaths + alcohol + percentage_expenditure + hepatitis_b + bmi + under-five_de
```

```
In [100]: model = smf.ols('life_expectancy ~ adult_mortality', data = df)
reg = model.fit()
reg.params
```

```
Out[100]: Intercept          79.451273
adult_mortality      -0.057357
dtype: float64
```

```
In [101]: model2 = smf.ols('life_expectancy ~ infant_deaths', data = df)
reg2 = model2.fit()
reg2.params
```

```
Out[101]: Intercept          71.434985
infant_deaths      -0.019669
dtype: float64
```

```
In [102]: model3 = smf.ols('life_expectancy ~ alcohol', data = df)
reg3 = model3.fit()
reg3.params
```

```
Out[102]: Intercept          66.767526
alcohol          1.022519
dtype: float64
```

```
In [103]: model4 = smf.ols('life_expectancy ~ percentage_expenditure', data = df)
reg4 = model4.fit()
reg4.params
```

```
Out[103]: Intercept          69.245494
percentage_expenditure      0.001652
dtype: float64
```



```
In [104]: model5 = smf.ols('life_expectancy ~ hepatitis_b', data = df)
reg5 = model5.fit()
reg5.params
```

```
Out[104]: Intercept      64.733057
hepatitis_b      0.071774
dtype: float64
```

```
In [105]: model6 = smf.ols('life_expectancy ~ bmi', data = df)
reg6 = model6.fit()
reg6.params
```

```
Out[105]: Intercept      62.278638
bmi      0.218540
dtype: float64
```

```
In [106]: model7 = smf.ols('life_expectancy ~ under_five_deaths', data = df)
reg7 = model7.fit()
reg7.params
```

```
Out[106]: Intercept      71.510258
under_five_deaths     -0.016684
dtype: float64
```

```
In [107]: model8 = smf.ols('life_expectancy ~ polio', data = df)
reg8 = model8.fit()
reg8.params
```

```
Out[107]: Intercept      58.501361
polio      0.147910
dtype: float64
```

```
In [108]: model9 = smf.ols('life_expectancy ~ total_expenditure', data = df)
reg9 = model9.fit()
reg9.params
```

```
Out[108]: Intercept          66.546236
total_expenditure      0.715904
dtype: float64
```

```
In [109]: model10 = smf.ols('life_expectancy ~ diphtheria', data = df)
reg10 = model10.fit()
reg10.params
```

```
Out[109]: Intercept          54.783967
diphtheria          0.187796
dtype: float64
```

```
In [110]: model11 = smf.ols('life_expectancy ~ hiv_aids', data = df)
reg11 = model11.fit()
reg11.params
```

```
Out[110]: Intercept          73.487502
hiv_aids          -2.773660
dtype: float64
```

```
In [111]: model12 = smf.ols('life_expectancy ~ gdp', data = df)
reg12 = model12.fit()
reg12.params
```

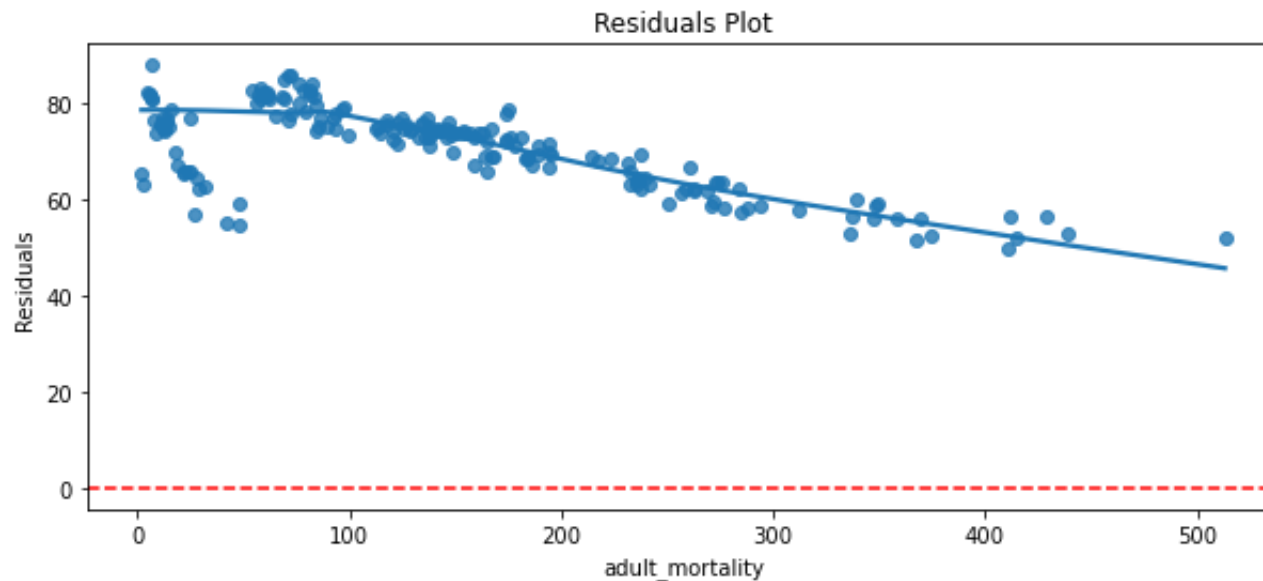
```
Out[111]: Intercept          68.345096
gdp              0.000266
dtype: float64
```

```
In [112]: model13 = smf.ols('life_expectancy ~ schooling', data = df)
reg13 = model13.fit()
reg13.params
```

```
Out[112]: Intercept    41.821061
schooling      2.293171
dtype: float64
```

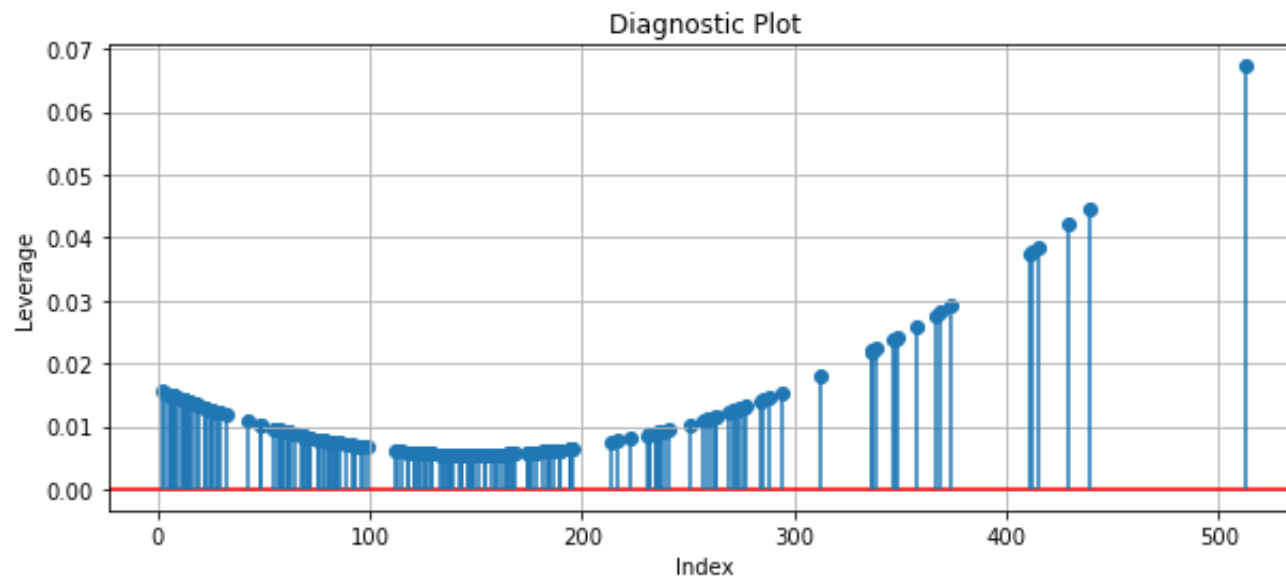
```
In [53]: # model = smf.ols('life_expectancy ~ adult_mortality', data = df)
### reg = model.fit()

plt.figure(figsize = (10, 4))
sns.regplot(x = df.adult_mortality, y = df.life_expectancy, lowess = True)
plt.axhline(0, linestyle = '--', color = "red")
plt.ylabel("Residuals")
plt.title("Residuals Plot")
plt.show()
```



```
In [57]: # leverage = reg.get_influence().hat_matrix_diag

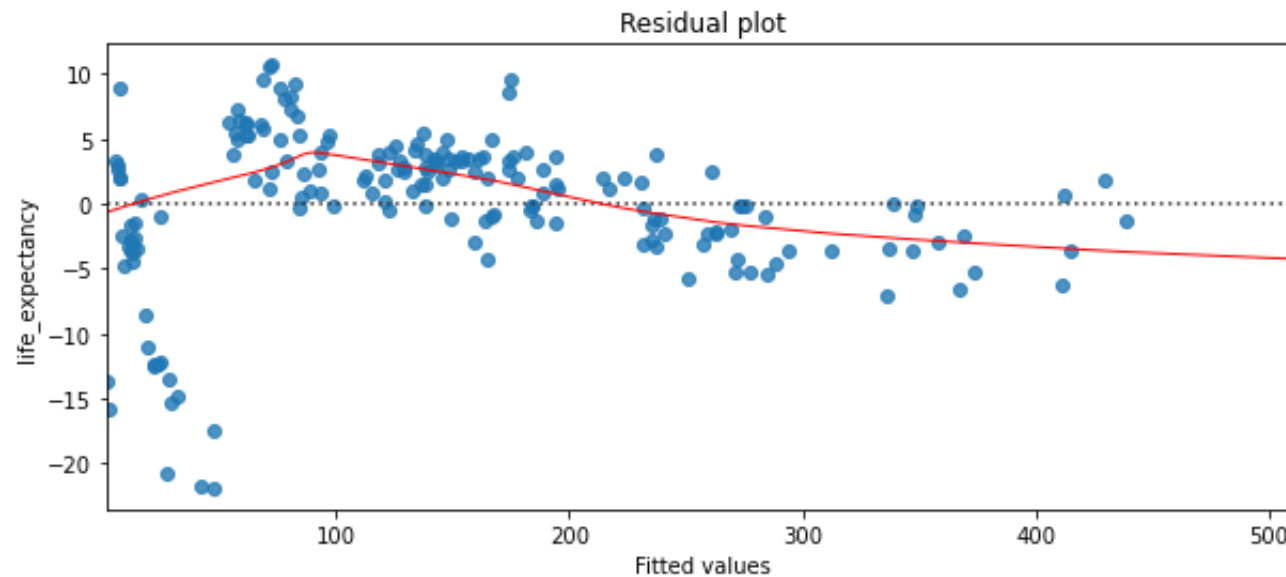
###plt.figure(figsize = (10, 4))
plt.scatter(df.adult_mortality, leverage)
plt.axhline(0, color = 'red')
plt.vlines(x = df.adult_mortality, ymin = 0, ymax = leverage)
plt.xlabel('Index')
plt.ylabel('Leverage')
plt.title("Diagnostic Plot")
plt.grid()
```



```
In [ ]: # adult_mortality + infant_deaths + alcohol + percentage_expenditure + hepatitis_b + bmi + under-five_de
```

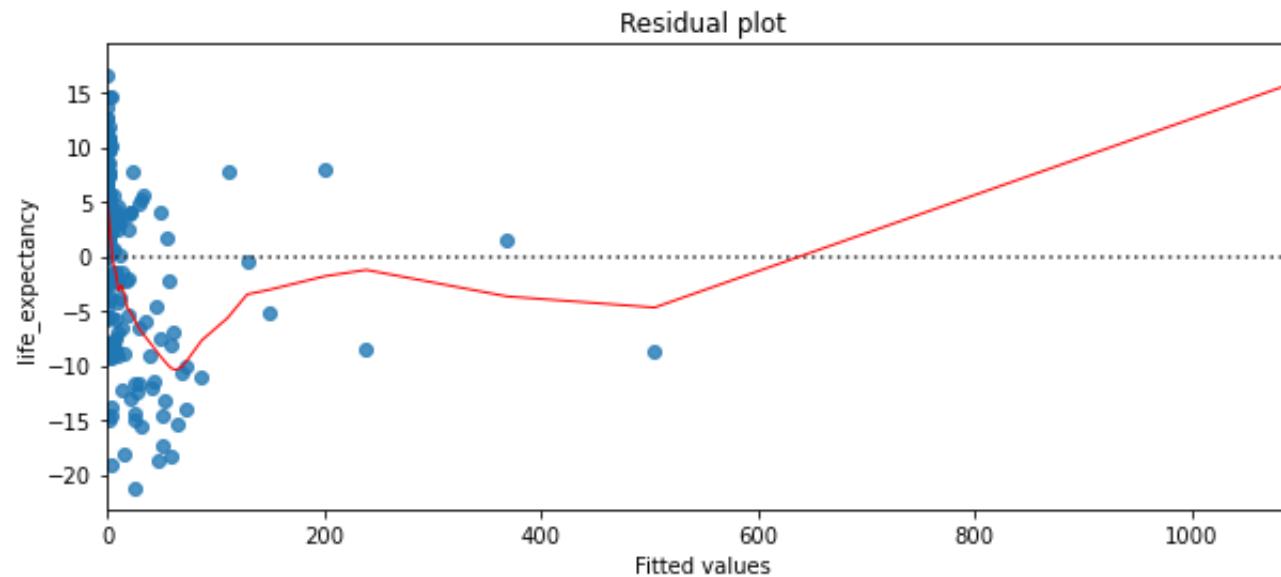
```
In [ ]: # Outliers
```

```
In [59]: plt.figure(figsize = (10, 4))
sns.residplot(x = 'adult_mortality', y = 'life_expectancy', data = df, lowess=True,
              line_kws={'color': 'red', 'lw': 1, 'alpha': 1})
plt.xlabel("Fitted values")
plt.title('Residual plot')
plt.show()
```

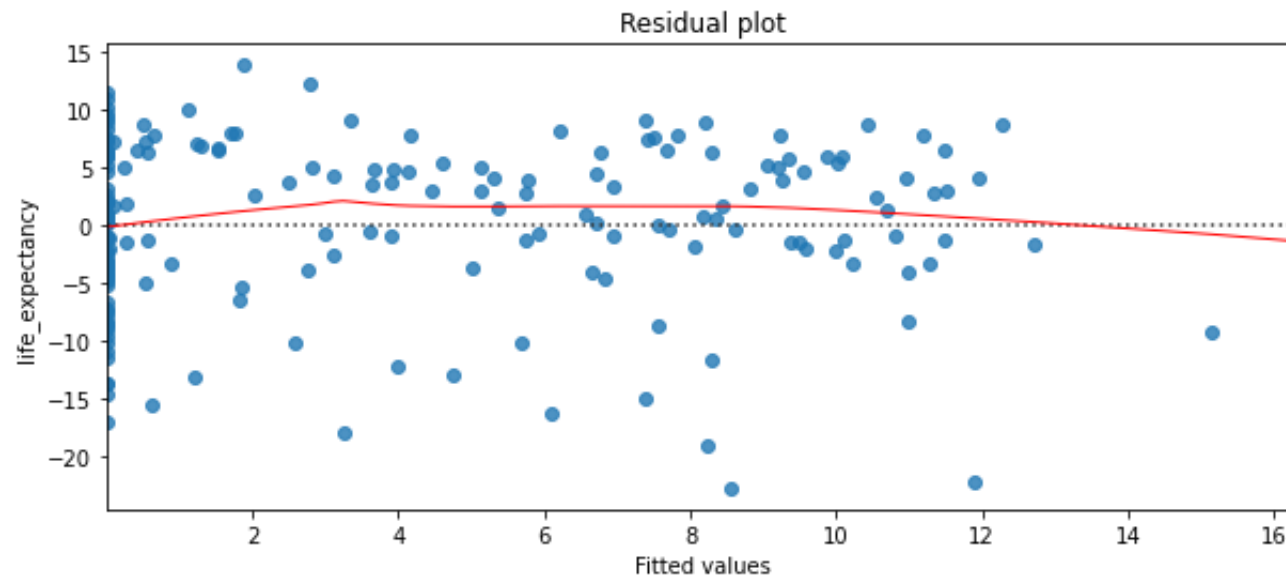


```
In [60]: plt.figure(figsize = (10, 4))
sns.residplot(x = 'infant_deaths', y = 'life_expectancy', data = df, lowess=True,
              line_kws={'color': 'red', 'lw': 1, 'alpha': 1})

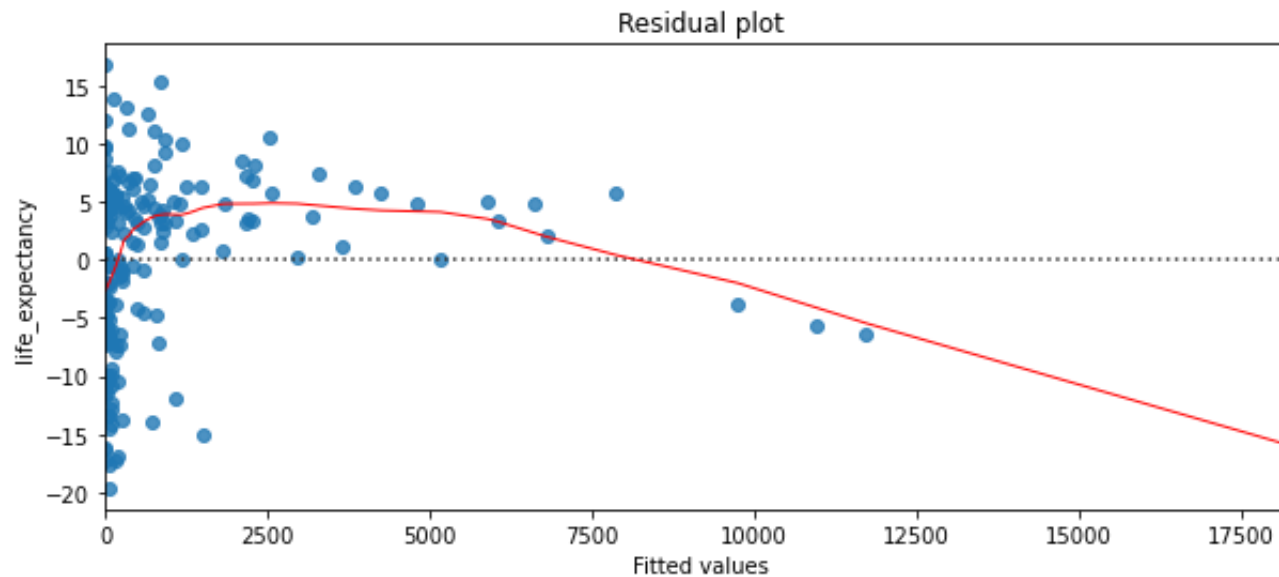
plt.xlabel("Fitted values")
plt.title('Residual plot')
plt.show()
```



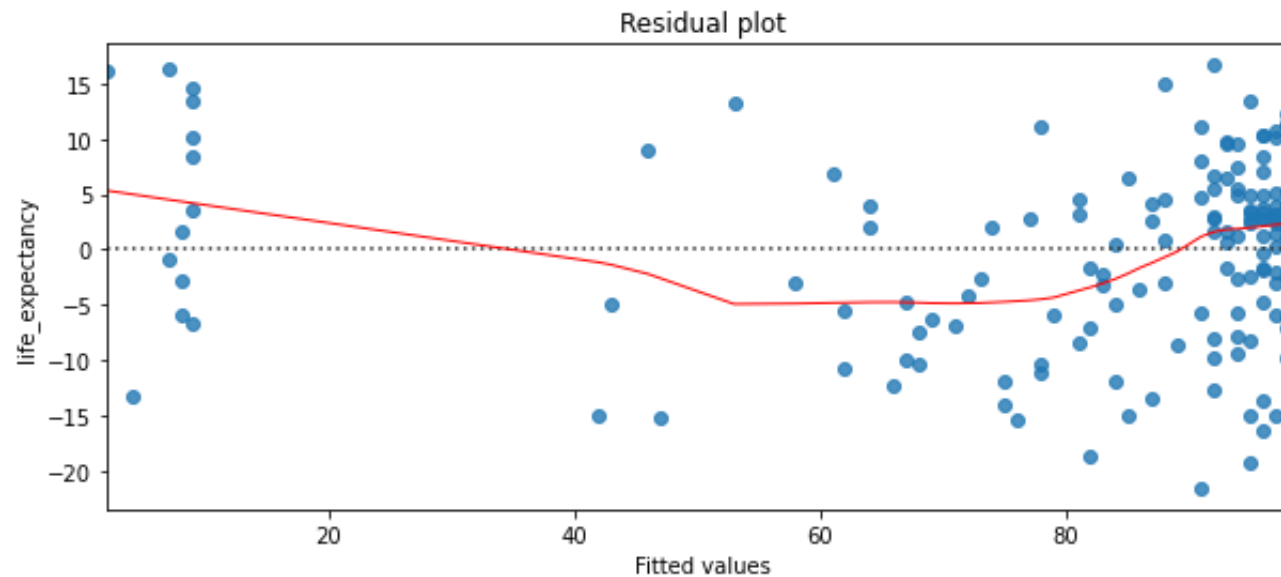
```
In [61]: plt.figure(figsize = (10, 4))
sns.residplot(x = 'alcohol', y = 'life_expectancy', data = df, lowess=True,
              line_kws={'color': 'red', 'lw': 1, 'alpha': 1})
plt.xlabel("Fitted values")
plt.title('Residual plot')
plt.show()
```



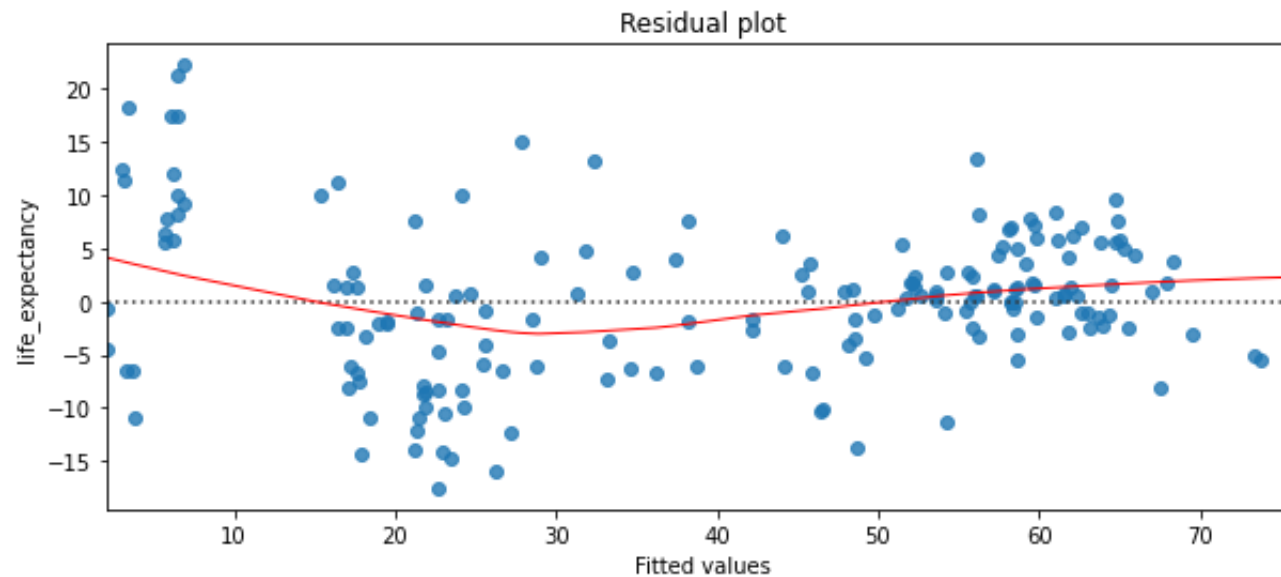
```
In [62]: plt.figure(figsize = (10, 4))
sns.residplot(x = 'percentage_expenditure', y = 'life_expectancy', data = df, lowess=True,
              line_kws={'color': 'red', 'lw': 1, 'alpha': 1})
plt.xlabel("Fitted values")
plt.title('Residual plot')
plt.show()
```



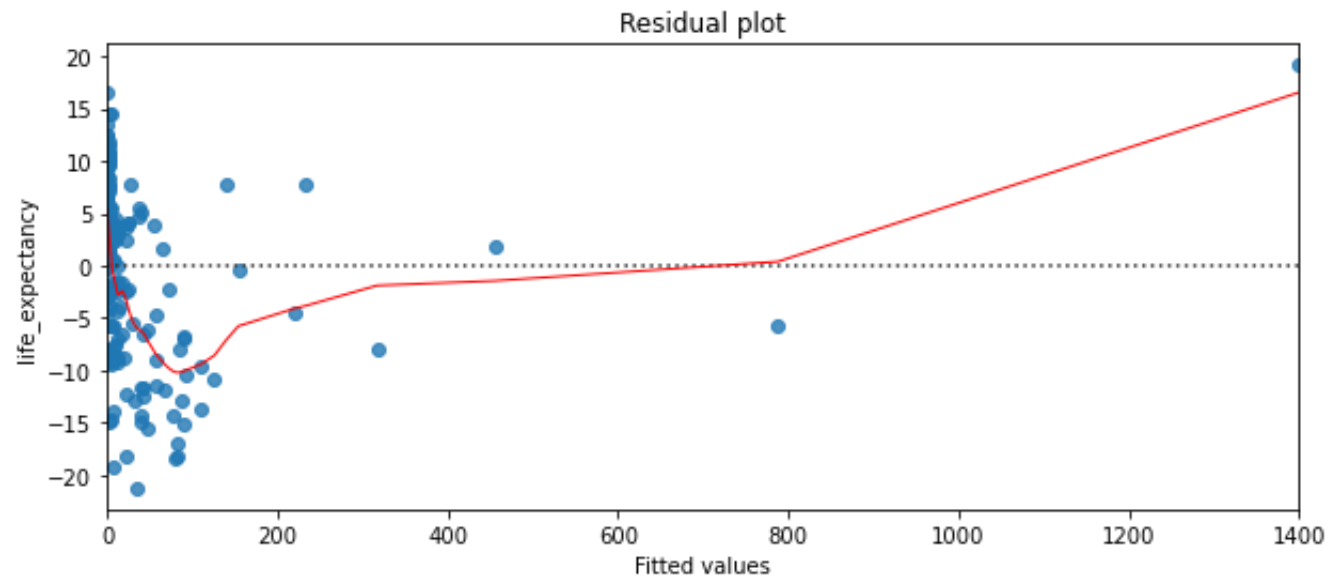

```
In [63]: plt.figure(figsize = (10, 4))
sns.residplot(x = 'hepatitis_b', y = 'life_expectancy', data = df, lowess=True,
              line_kws={'color': 'red', 'lw': 1, 'alpha': 1})
plt.xlabel("Fitted values")
plt.title('Residual plot')
plt.show()
```



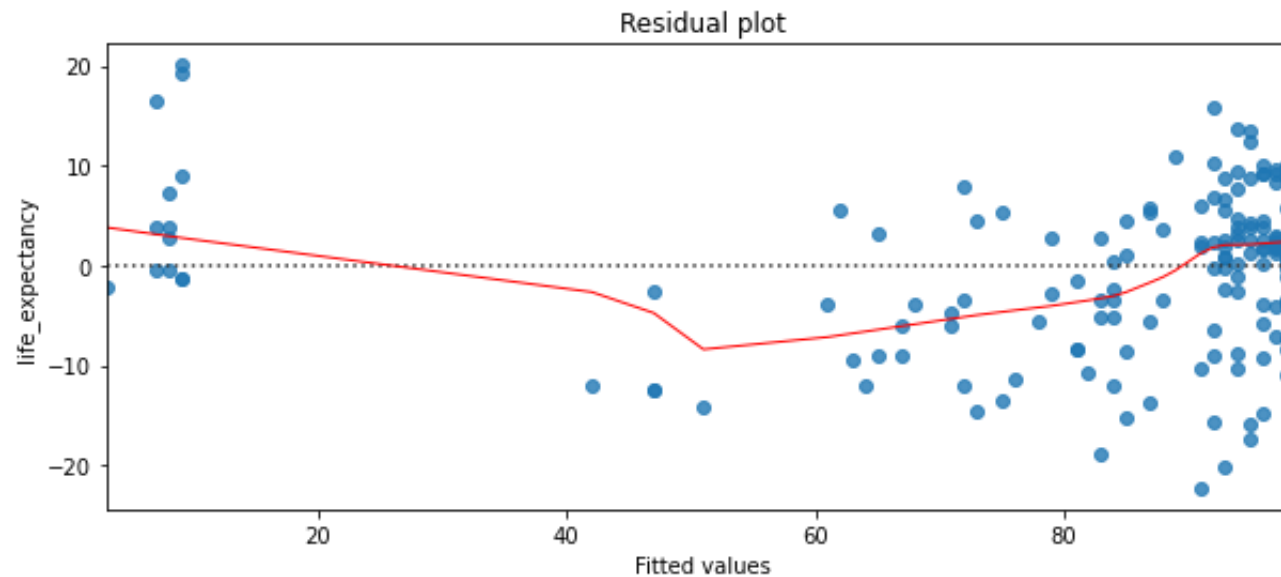
```
In [64]: plt.figure(figsize = (10, 4))
sns.residplot(x = 'bmi', y = 'life_expectancy', data = df, lowess=True,
              line_kws={'color': 'red', 'lw': 1, 'alpha': 1})
plt.xlabel("Fitted values")
plt.title('Residual plot')
plt.show()
```



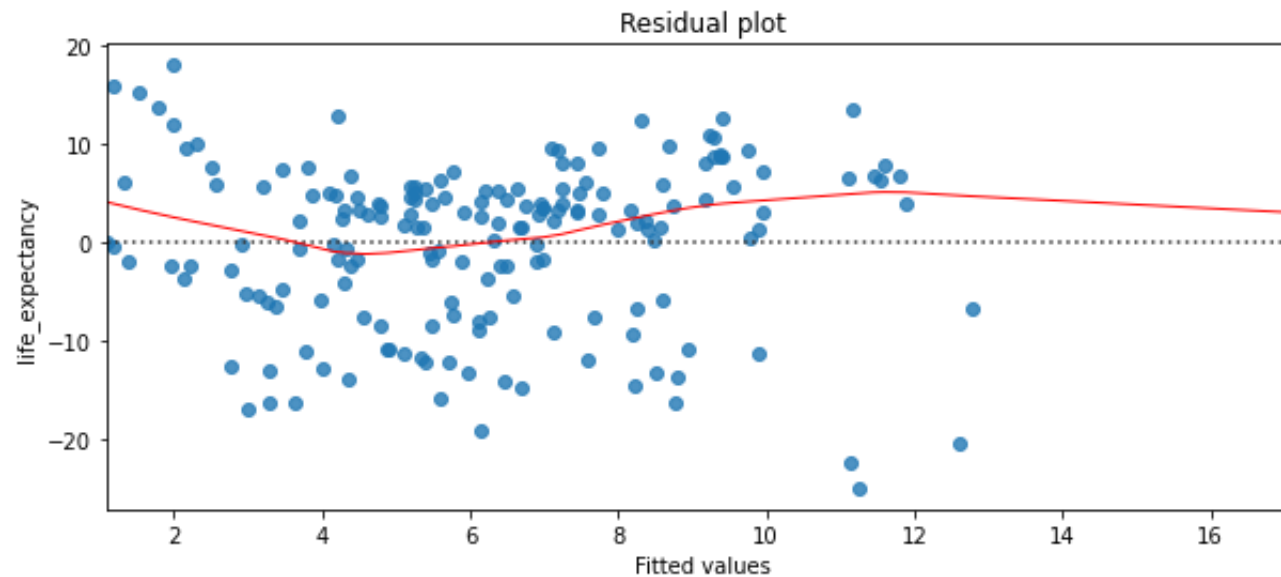
```
In [65]: plt.figure(figsize = (10, 4))
sns.residplot(x = 'under-five_deaths', y = 'life_expectancy', data = df, lowess=True,
              line_kws={'color': 'red', 'lw': 1, 'alpha': 1})
plt.xlabel("Fitted values")
plt.title('Residual plot')
plt.show()
```



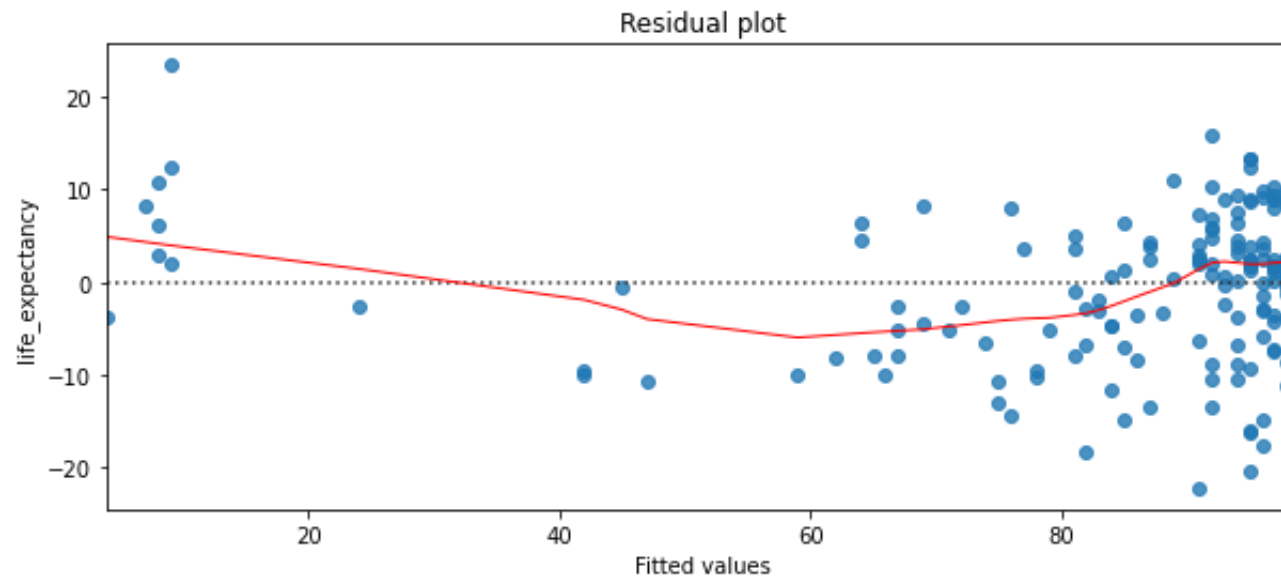
```
In [66]: plt.figure(figsize = (10, 4))
sns.residplot(x = 'polio', y = 'life_expectancy', data = df, lowess=True,
              line_kws={'color': 'red', 'lw': 1, 'alpha': 1})
plt.xlabel("Fitted values")
plt.title('Residual plot')
plt.show()
```



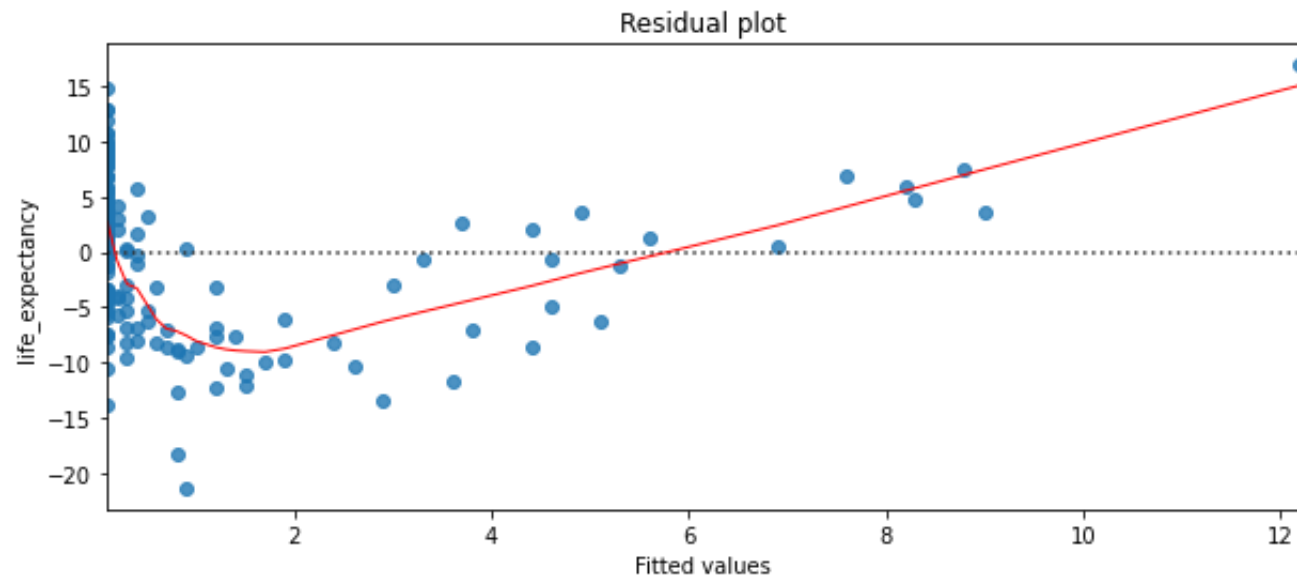
```
In [67]: plt.figure(figsize = (10, 4))
sns.residplot(x = 'total_expenditure', y = 'life_expectancy', data = df, lowess=True,
              line_kws={'color': 'red', 'lw': 1, 'alpha': 1})
plt.xlabel("Fitted values")
plt.title('Residual plot')
plt.show()
```



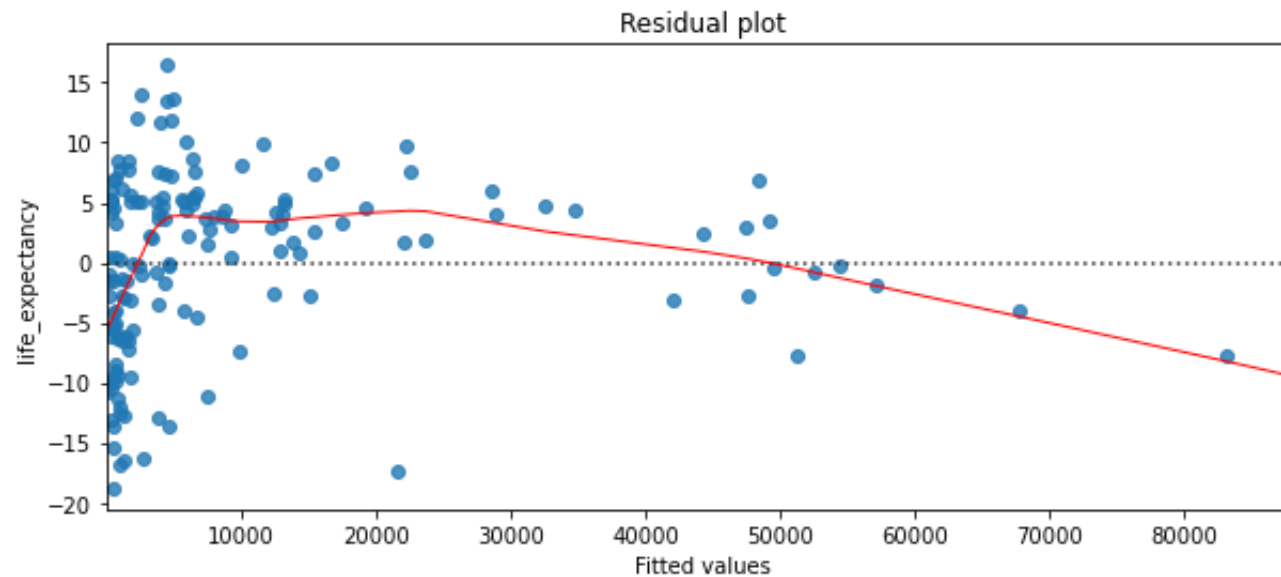
```
In [68]: plt.figure(figsize = (10, 4))
sns.residplot(x = 'diphtheria', y = 'life_expectancy', data = df, lowess=True,
              line_kws={'color': 'red', 'lw': 1, 'alpha': 1})
plt.xlabel("Fitted values")
plt.title('Residual plot')
plt.show()
```



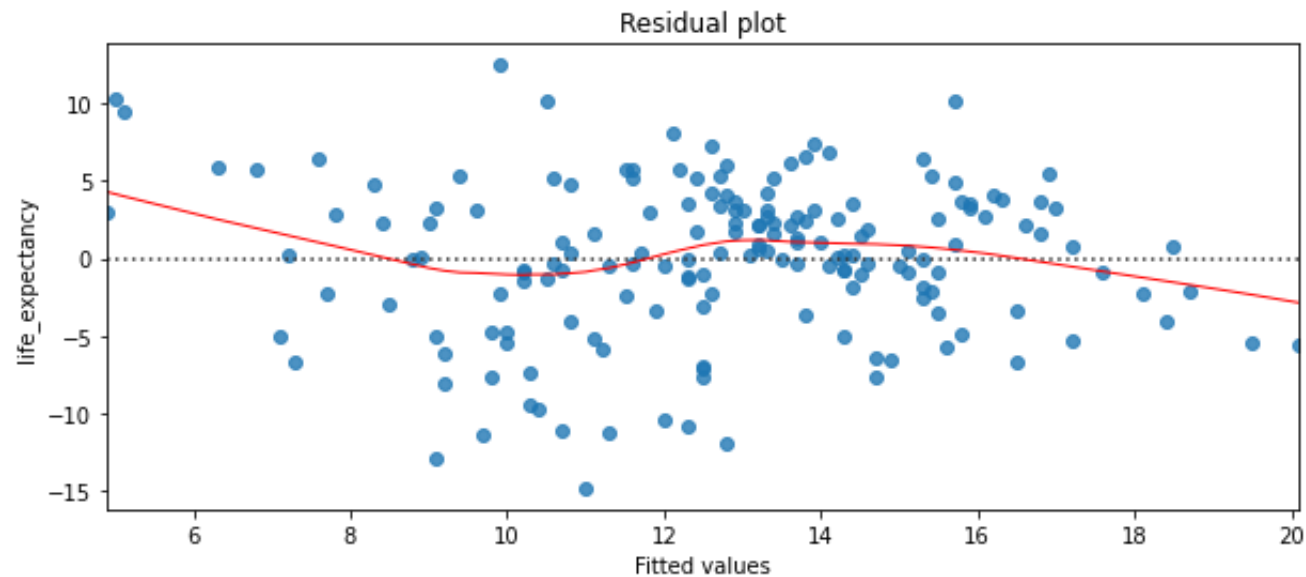
```
In [69]: plt.figure(figsize = (10, 4))
sns.residplot(x = 'hiv/aids', y = 'life_expectancy', data = df, lowess=True,
              line_kws={'color': 'red', 'lw': 1, 'alpha': 1})
plt.xlabel("Fitted values")
plt.title('Residual plot')
plt.show()
```



```
In [70]: plt.figure(figsize = (10, 4))
sns.residplot(x = 'gdp', y = 'life_expectancy', data = df, lowess=True,
              line_kws={'color': 'red', 'lw': 1, 'alpha': 1})
plt.xlabel("Fitted values")
plt.title('Residual plot')
plt.show()
```




```
In [71]: plt.figure(figsize = (10, 4))
sns.residplot(x = 'schooling', y = 'life_expectancy', data = df, lowess=True,
              line_kws={'color': 'red', 'lw': 1, 'alpha': 1})
plt.xlabel("Fitted values")
plt.title('Residual plot')
plt.show()
```

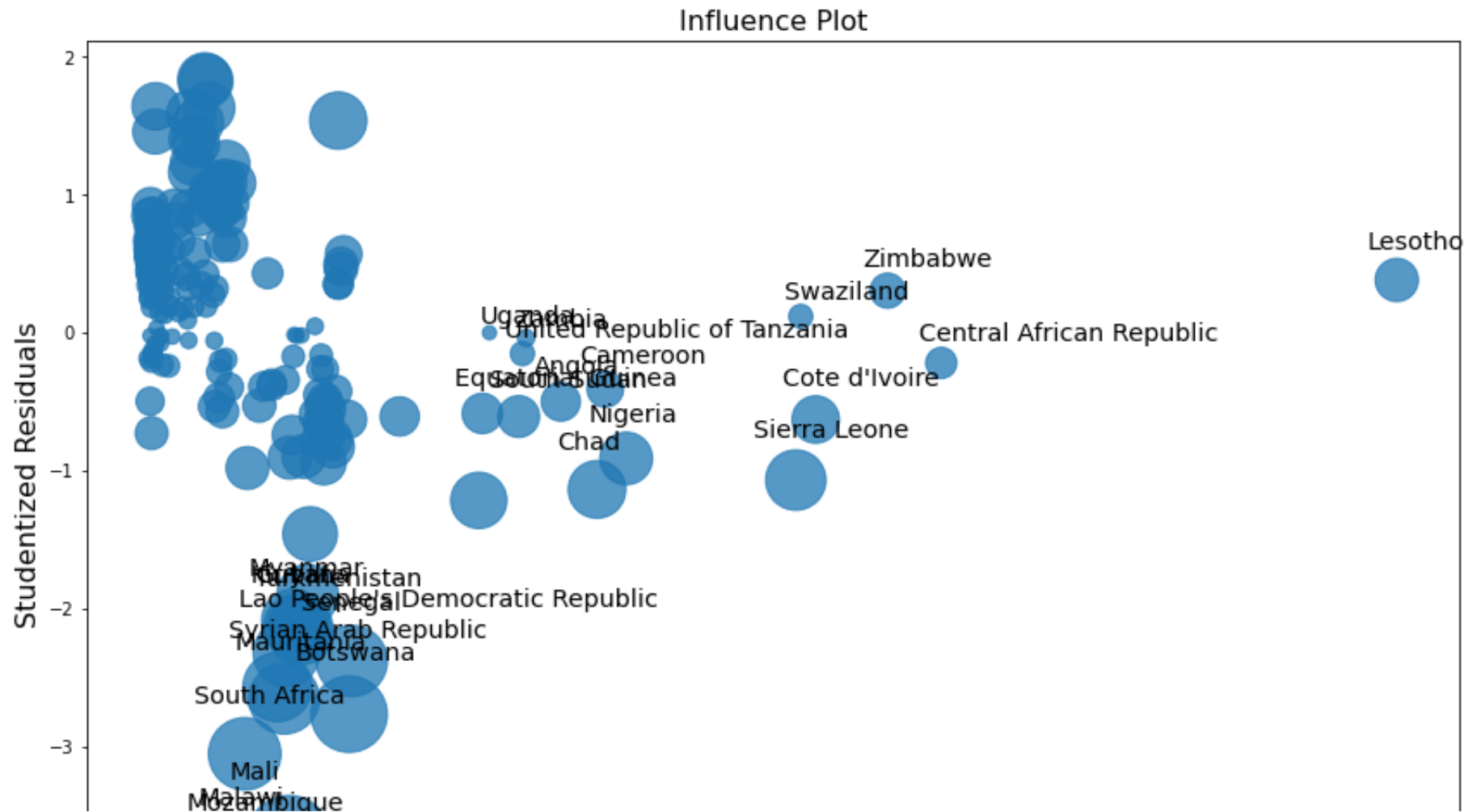


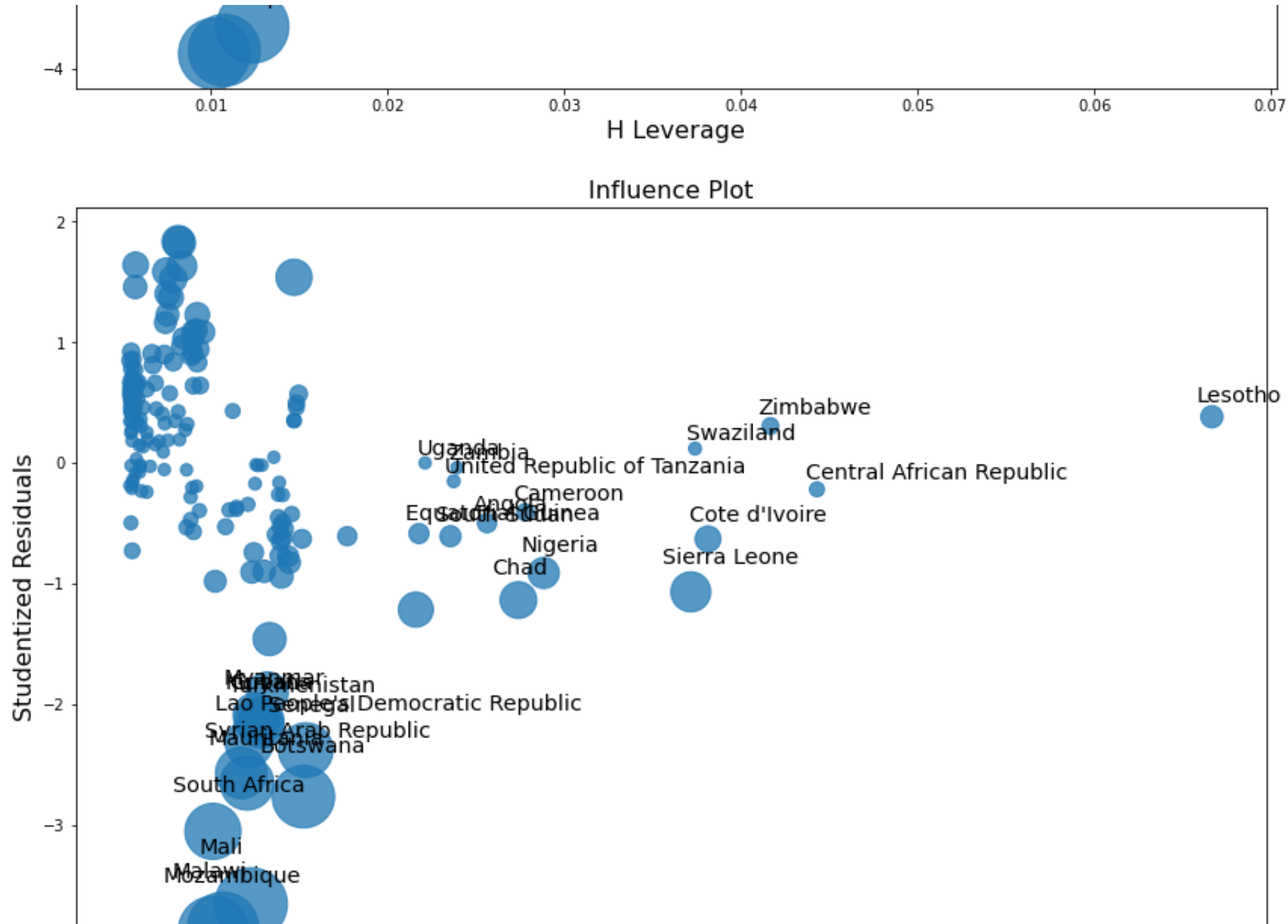
```
In [74]: #plots for all predictors
```

```
In [73]:
```

```
figd, ax = plt.subplots(figsize=(12,8))
figd = sm.graphics.influence_plot(reg, ax = ax, criterion="DFFITS")
figd.tight_layout(pad=1.0)

fige, ax = plt.subplots(figsize=(12,8))
fige = sm.graphics.influence_plot(reg, ax = ax, criterion="cooks")
fige.tight_layout(pad=1.0)
```

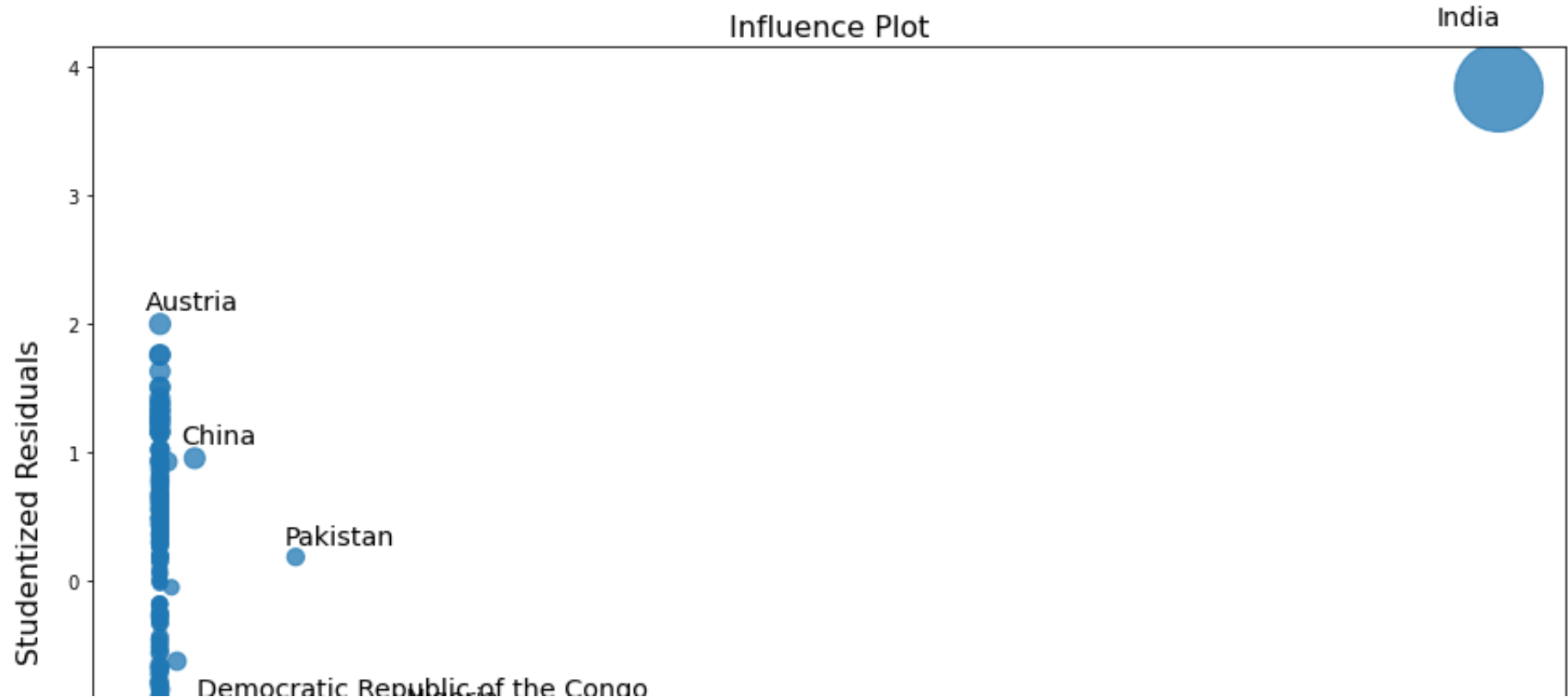


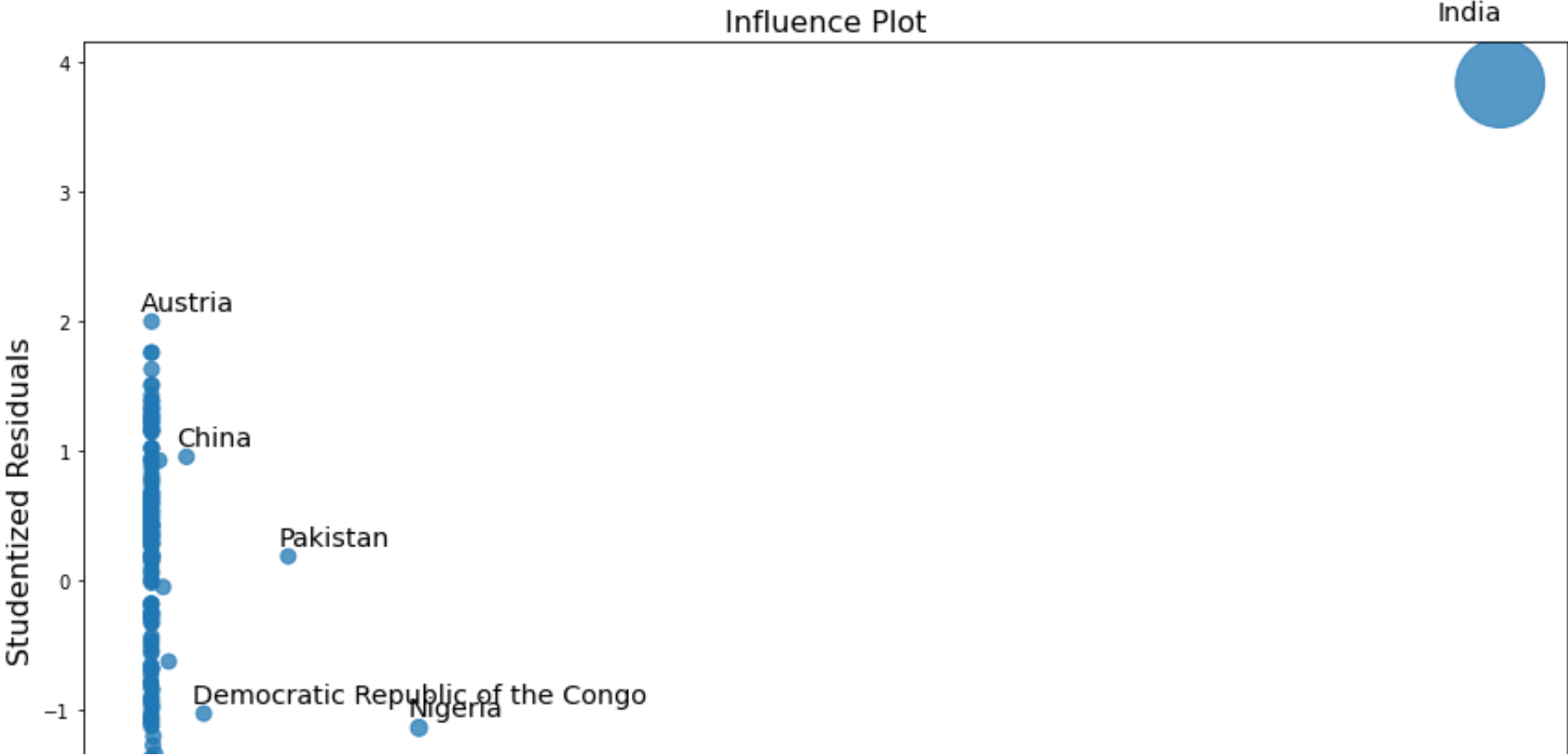
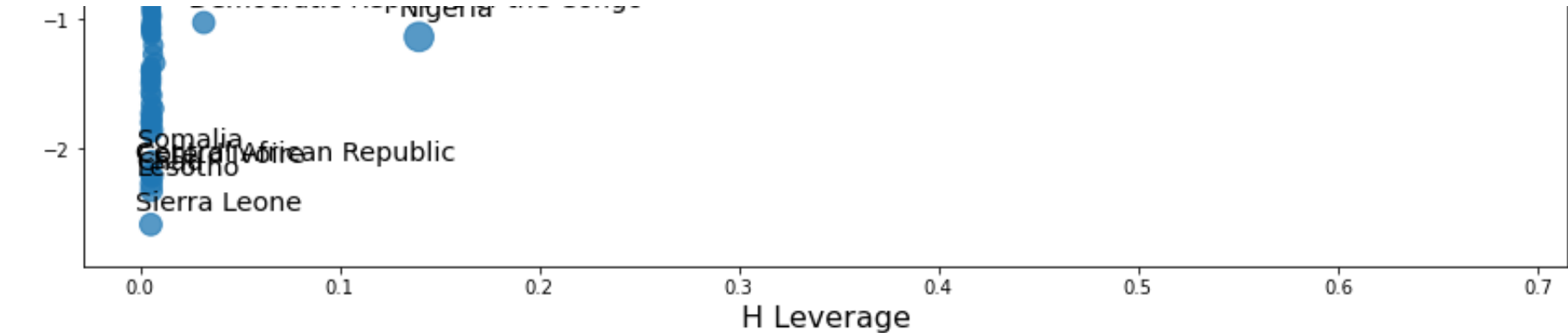


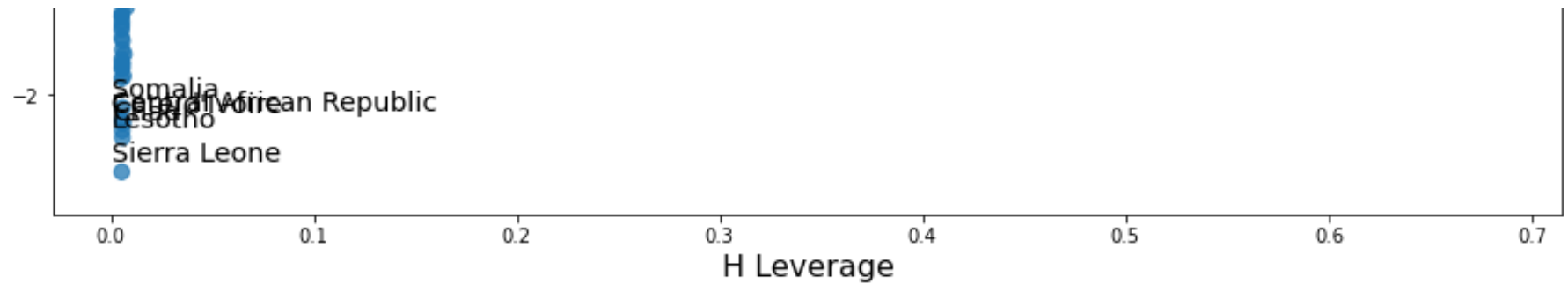


```
In [113]: figd, ax = plt.subplots(figsize=(12,8))
figd = sm.graphics.influence_plot(reg2, ax = ax, criterion="DFFITS")
figd.tight_layout(pad=1.0)

fige, ax = plt.subplots(figsize=(12,8))
fige = sm.graphics.influence_plot(reg2, ax = ax, criterion="cooks")
fige.tight_layout(pad=1.0)
```

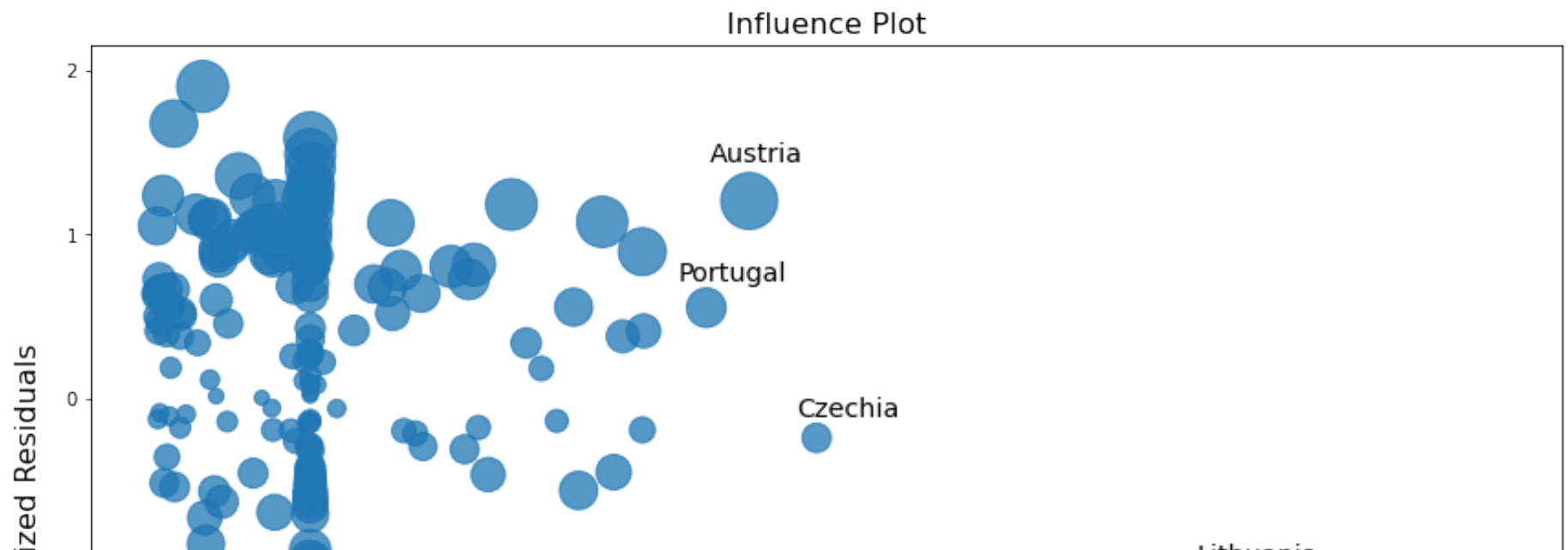


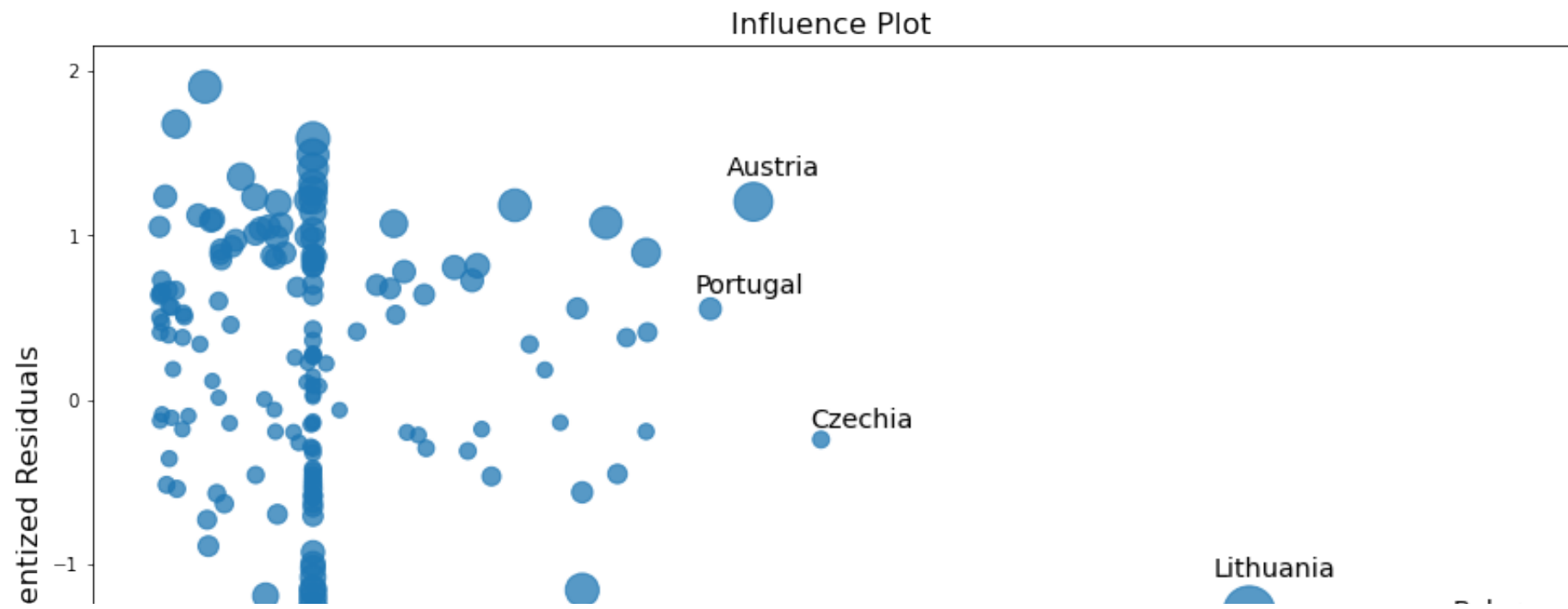
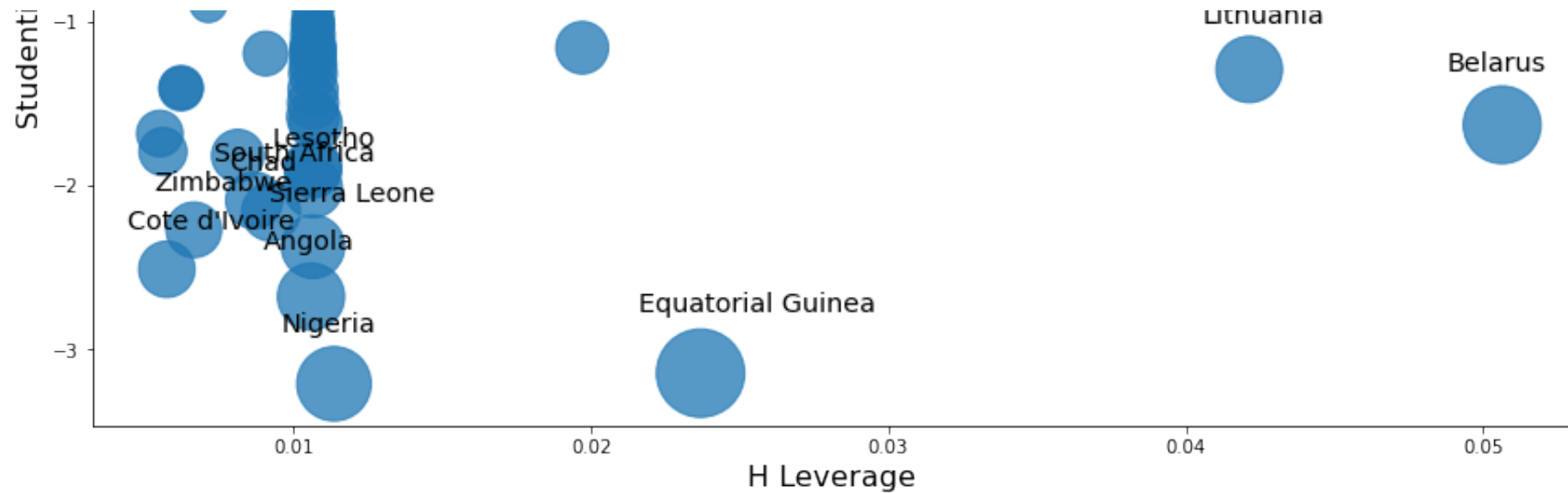


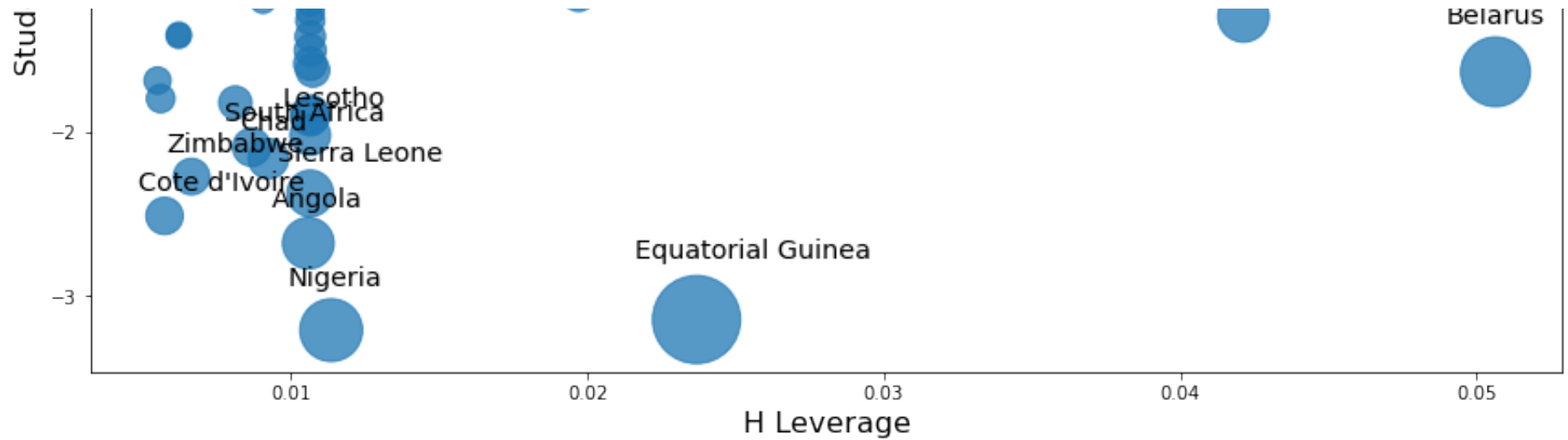


```
In [114]: figd, ax = plt.subplots(figsize=(12,8))
figd = sm.graphics.influence_plot(reg3, ax = ax, criterion="DFFITS")
figd.tight_layout(pad=1.0)

fige, ax = plt.subplots(figsize=(12,8))
fige = sm.graphics.influence_plot(reg3, ax = ax, criterion="cooks")
fige.tight_layout(pad=1.0)
```



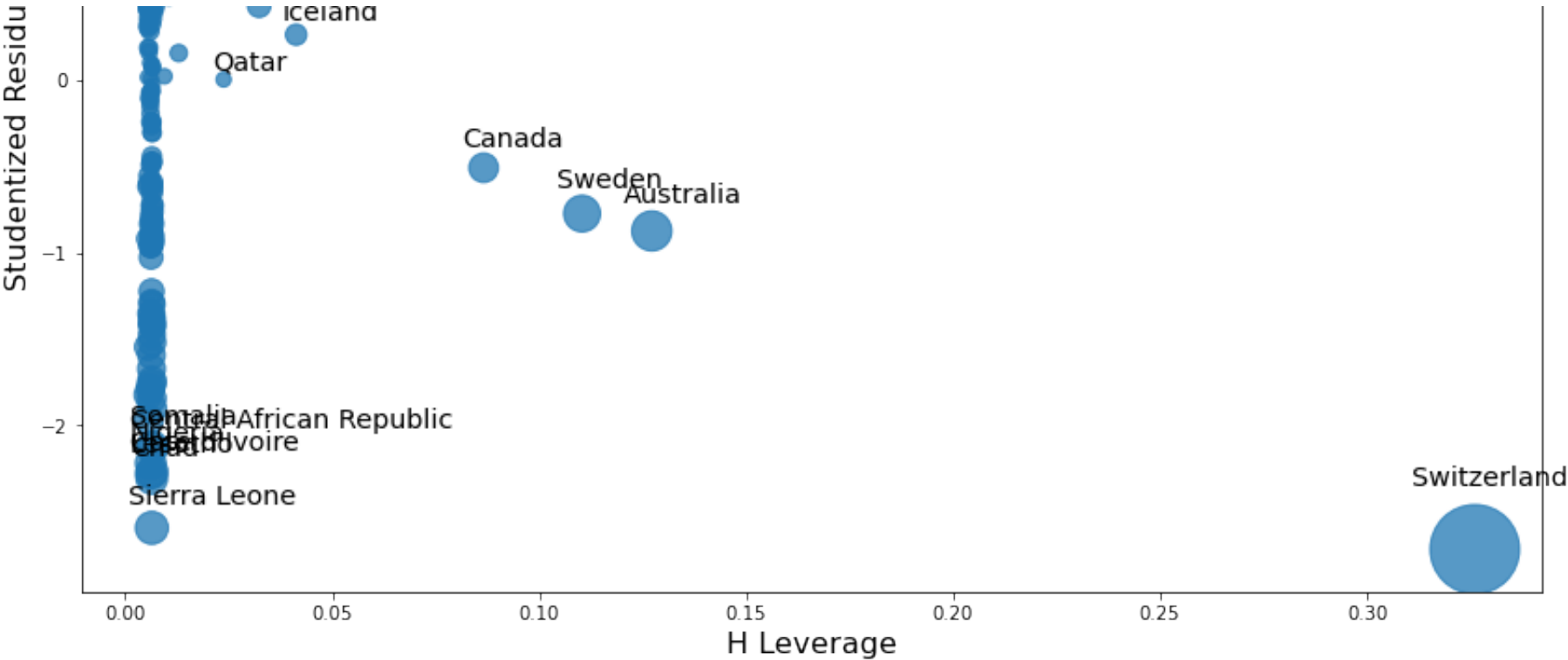


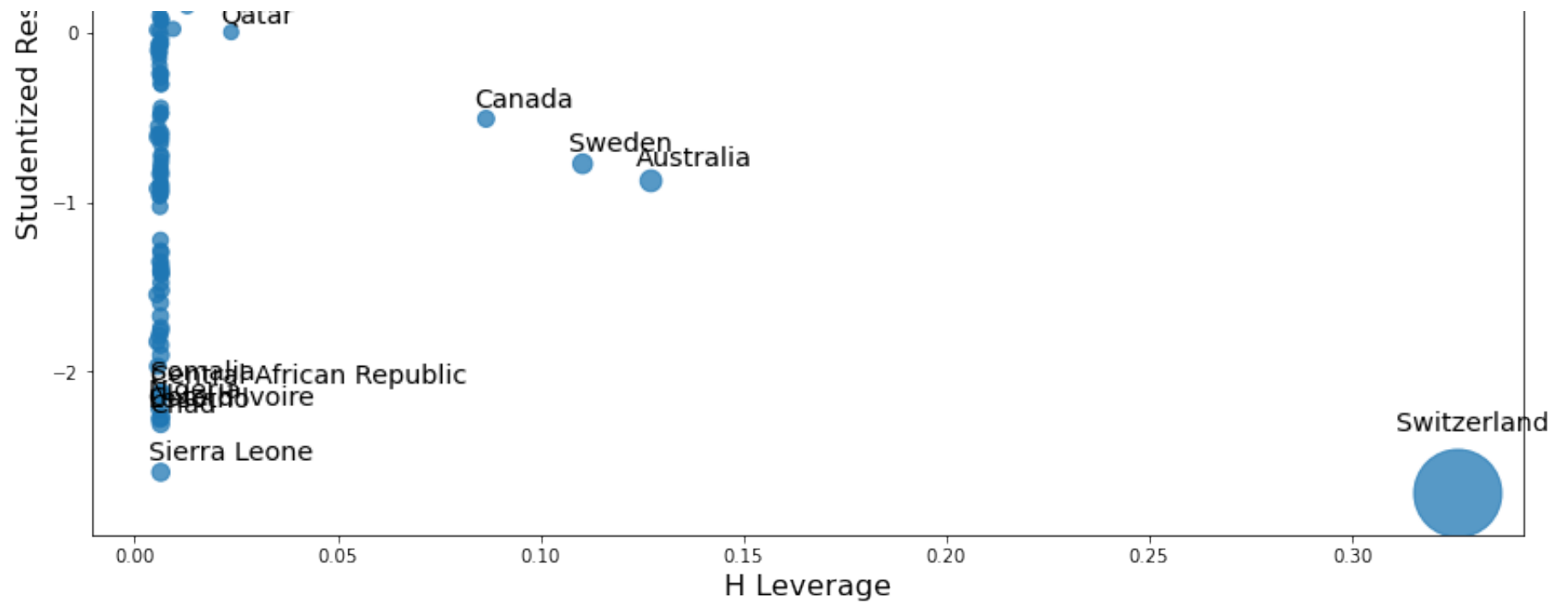


```
In [115]: figd, ax = plt.subplots(figsize=(12,8))
figd = sm.graphics.influence_plot(reg4, ax = ax, criterion="DFFITS")
figd.tight_layout(pad=1.0)

fige, ax = plt.subplots(figsize=(12,8))
fige = sm.graphics.influence_plot(reg4, ax = ax, criterion="cooks")
fige.tight_layout(pad=1.0)
```

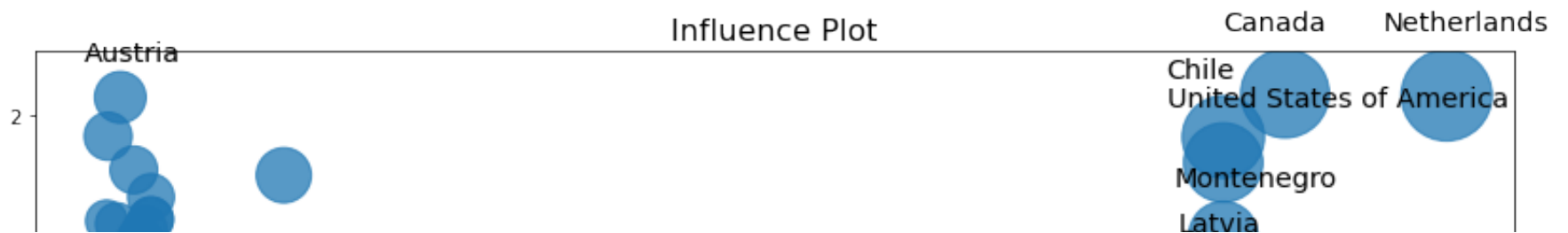


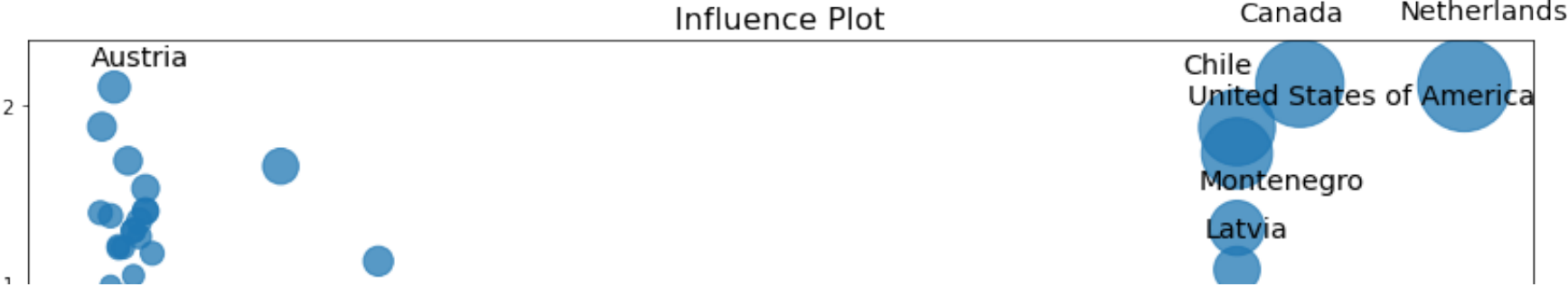
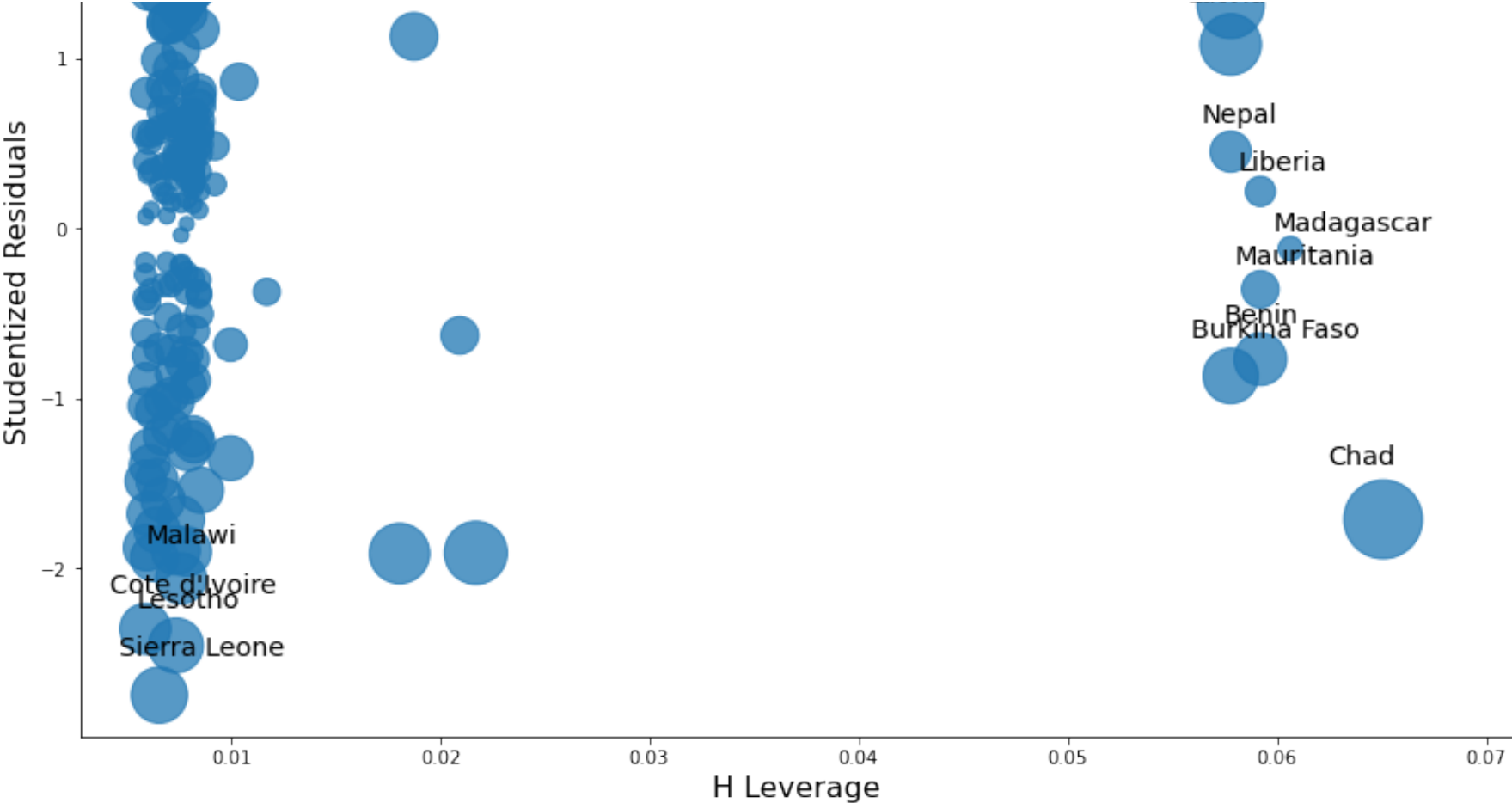


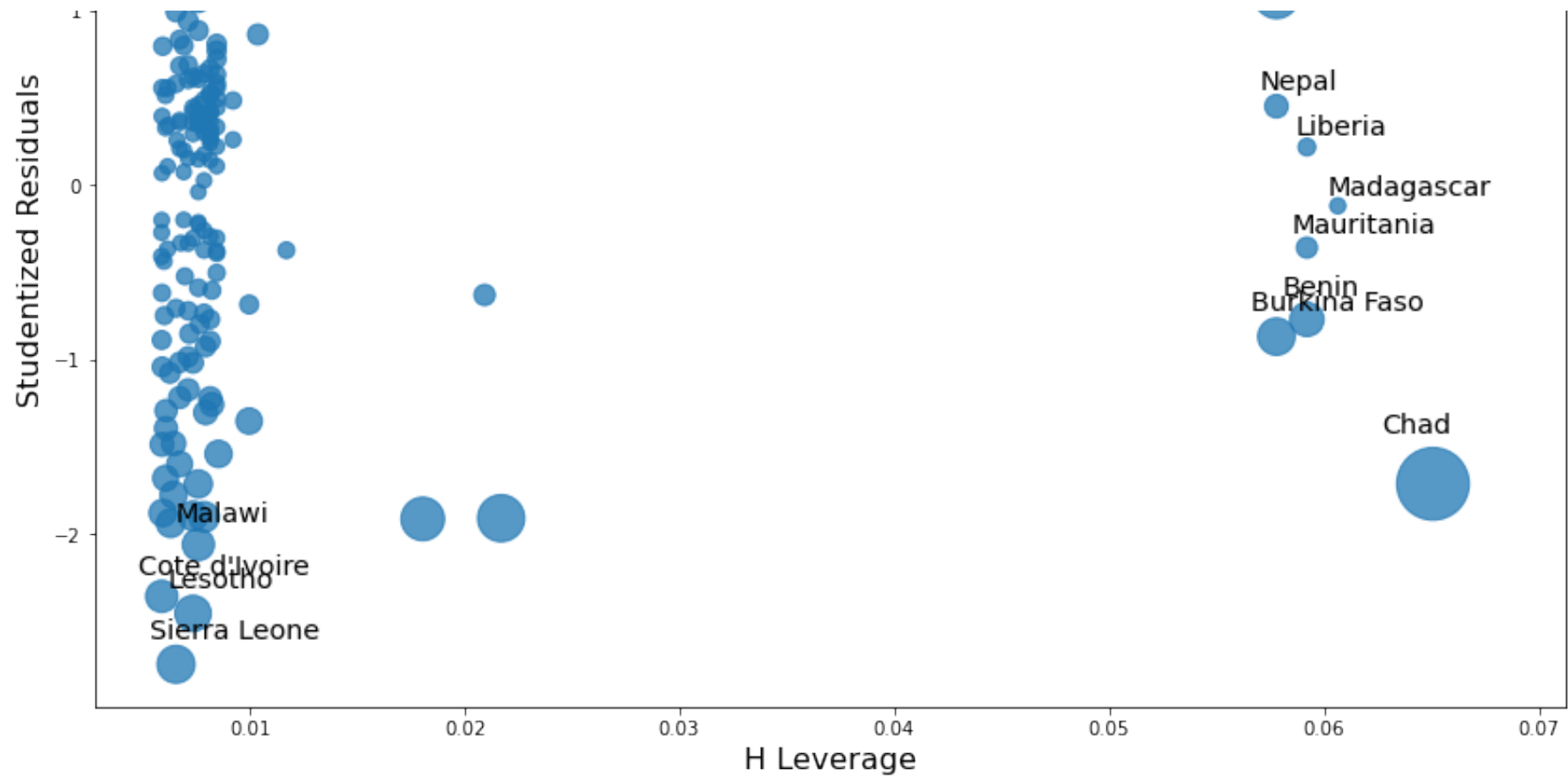


```
In [116]: figd, ax = plt.subplots(figsize=(12,8))
figd = sm.graphics.influence_plot(reg5, ax = ax, criterion="DFFITS")
figd.tight_layout(pad=1.0)

fige, ax = plt.subplots(figsize=(12,8))
fige = sm.graphics.influence_plot(reg5, ax = ax, criterion="cooks")
fige.tight_layout(pad=1.0)
```



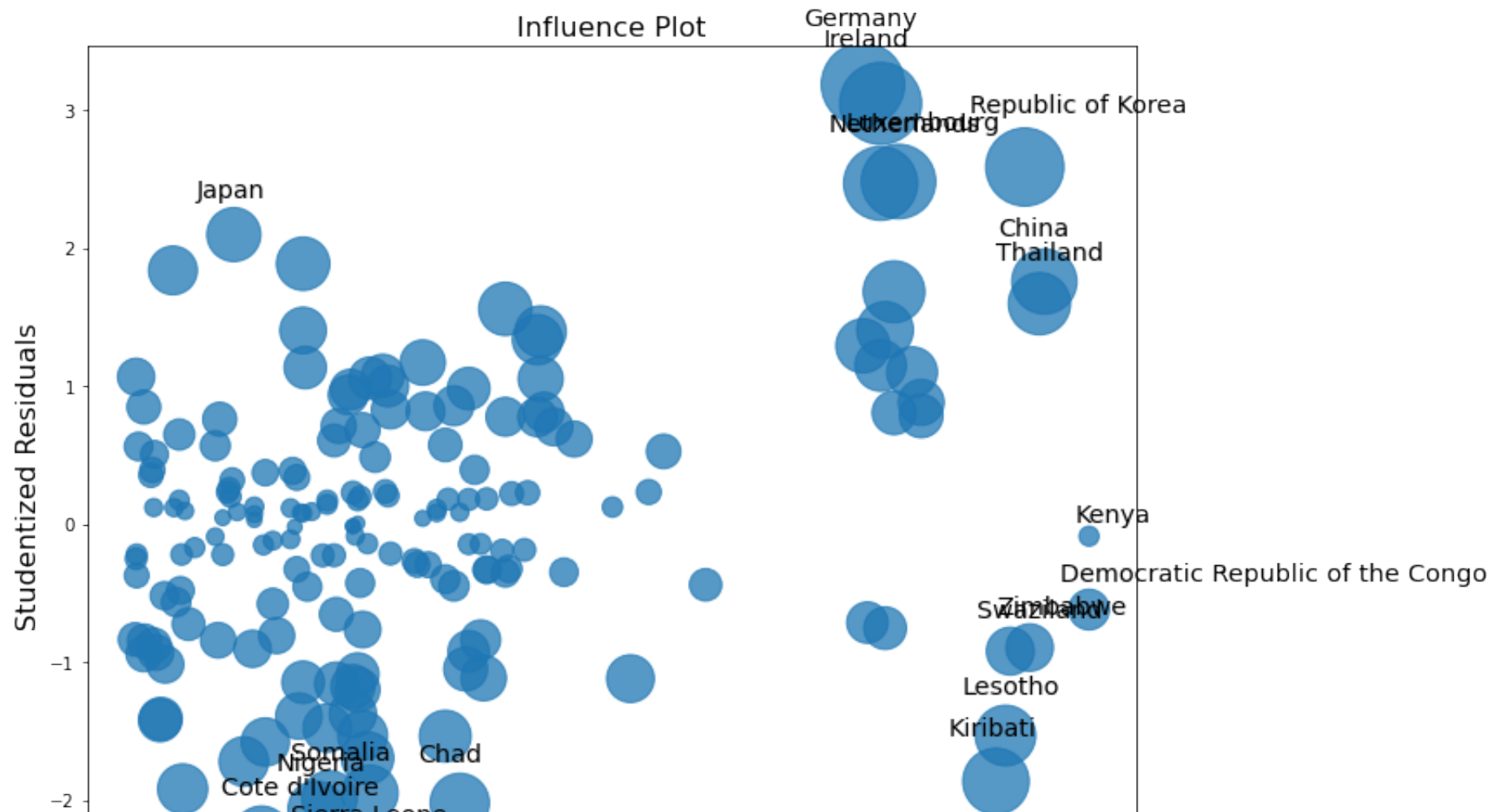


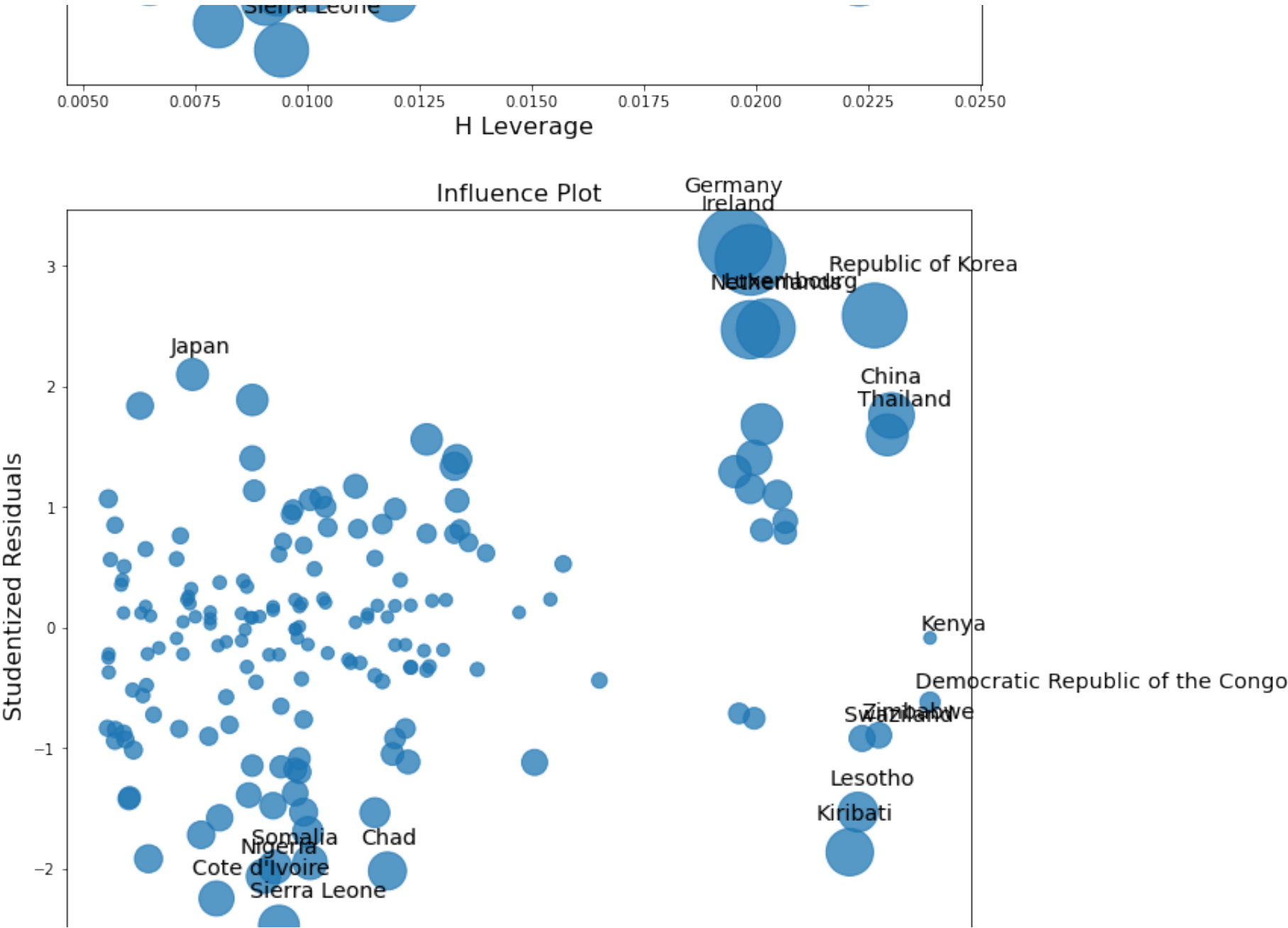


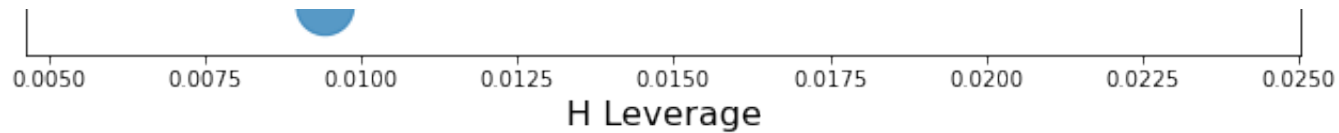
In [117]:

```
figd, ax = plt.subplots(figsize=(12,8))
figd = sm.graphics.influence_plot(reg6, ax = ax, criterion="DFFITS")
figd.tight_layout(pad=1.0)

fige, ax = plt.subplots(figsize=(12,8))
fige = sm.graphics.influence_plot(reg6, ax = ax, criterion="cooks")
fige.tight_layout(pad=1.0)
```

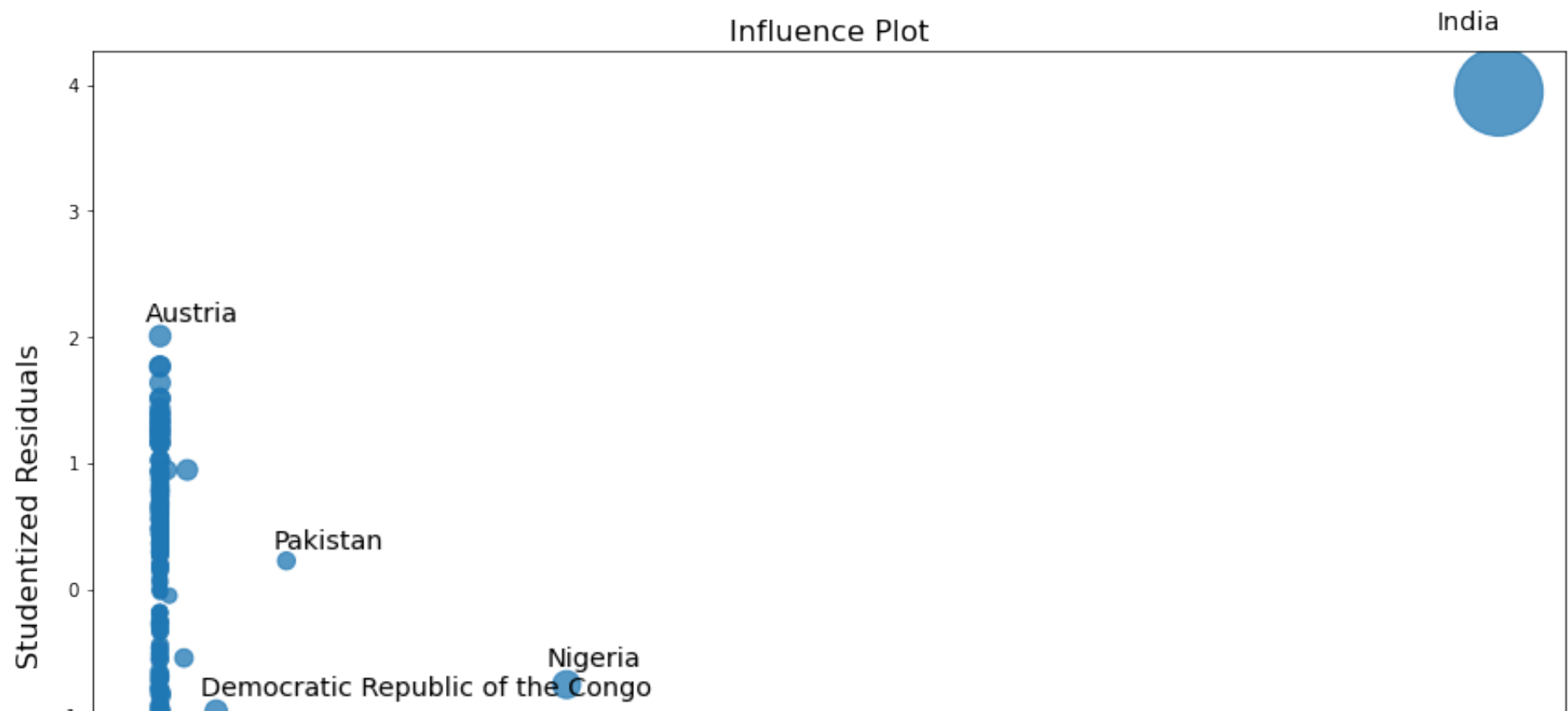


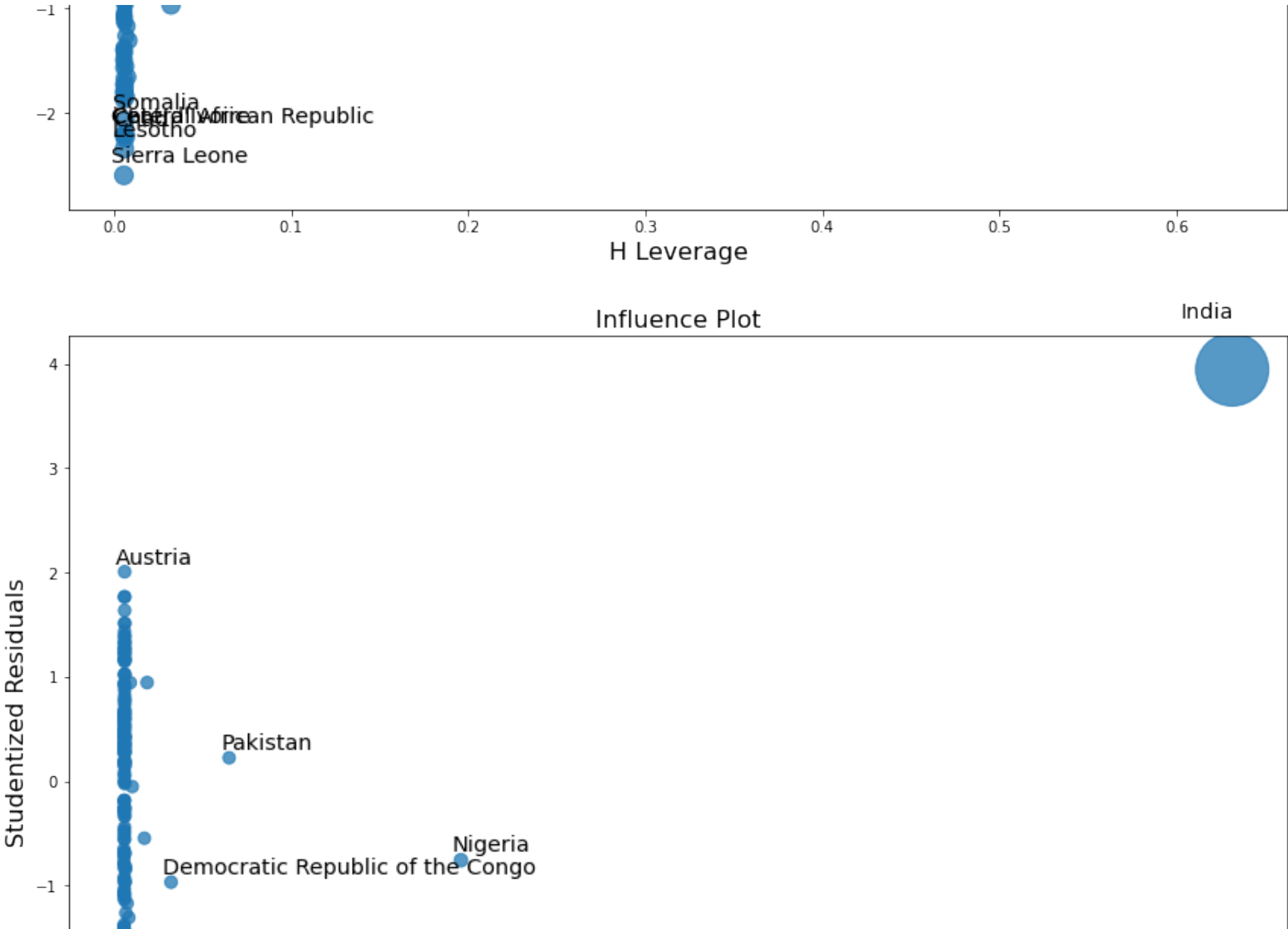


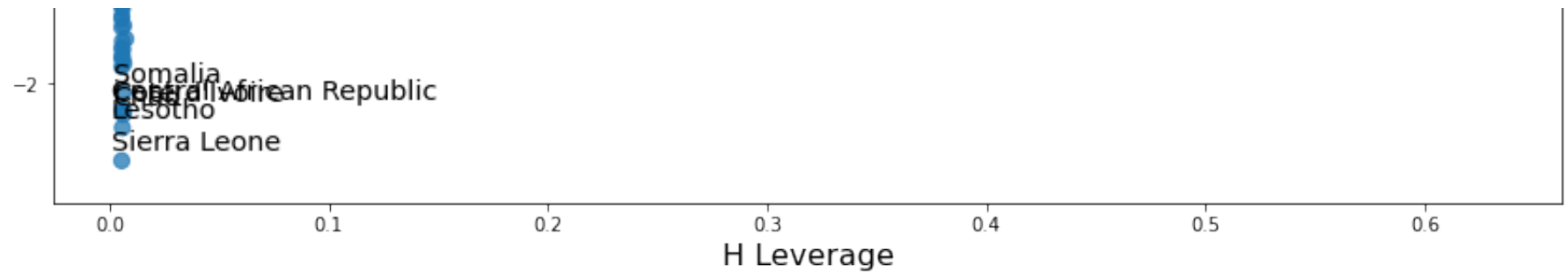


```
In [118]: figd, ax = plt.subplots(figsize=(12,8))
figd = sm.graphics.influence_plot(reg7, ax = ax, criterion="DFFITS")
figd.tight_layout(pad=1.0)

fige, ax = plt.subplots(figsize=(12,8))
fige = sm.graphics.influence_plot(reg7, ax = ax, criterion="cooks")
fige.tight_layout(pad=1.0)
```



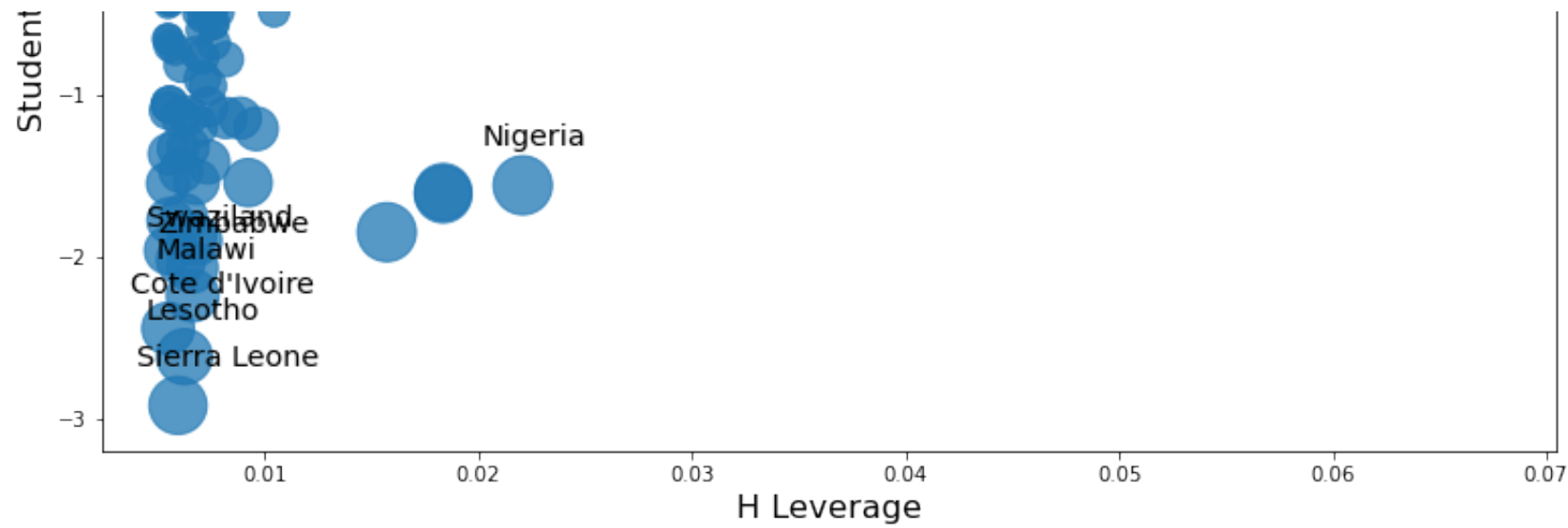


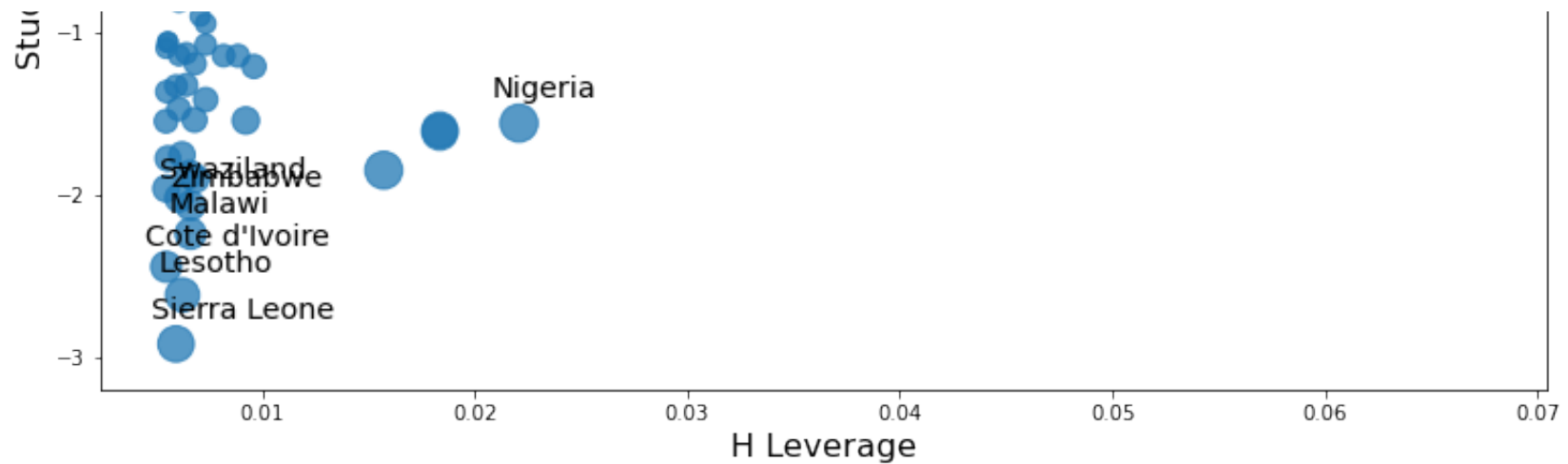


```
In [119]: figd, ax = plt.subplots(figsize=(12,8))
figd = sm.graphics.influence_plot(reg8, ax = ax, criterion="DFFITS")
figd.tight_layout(pad=1.0)

fige, ax = plt.subplots(figsize=(12,8))
fige = sm.graphics.influence_plot(reg8, ax = ax, criterion="cooks")
fige.tight_layout(pad=1.0)
```

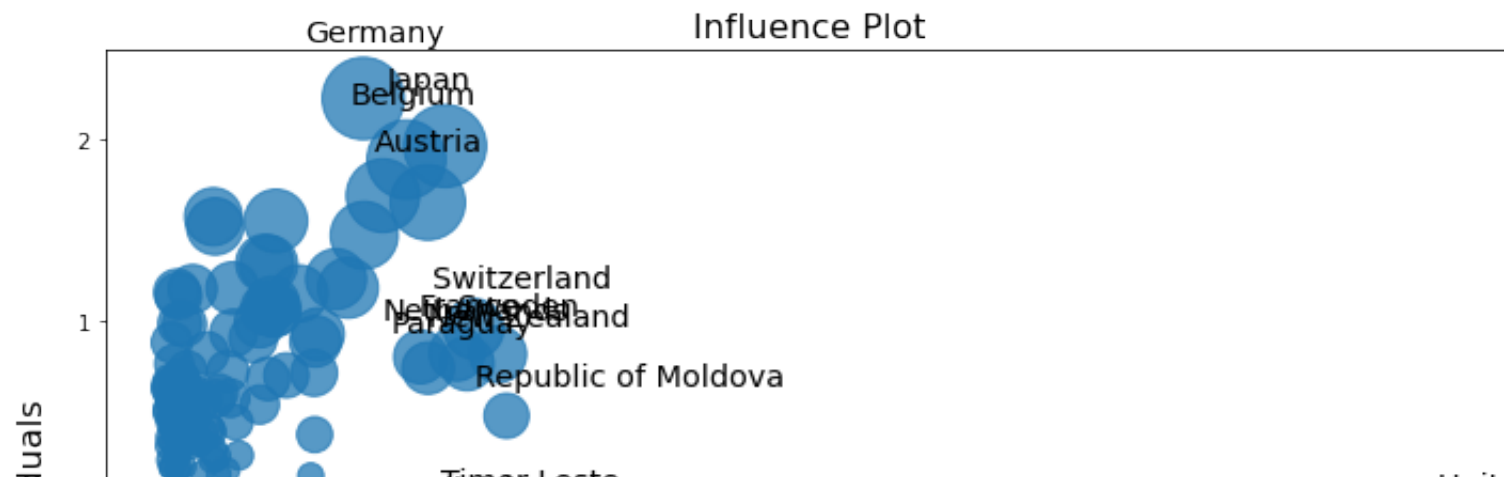


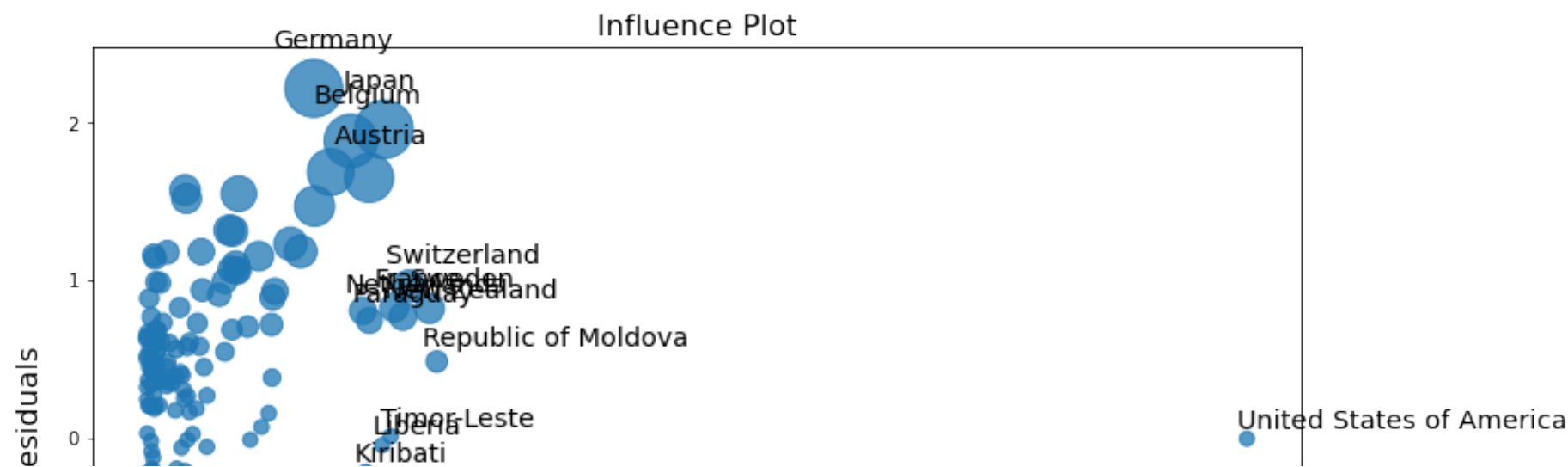
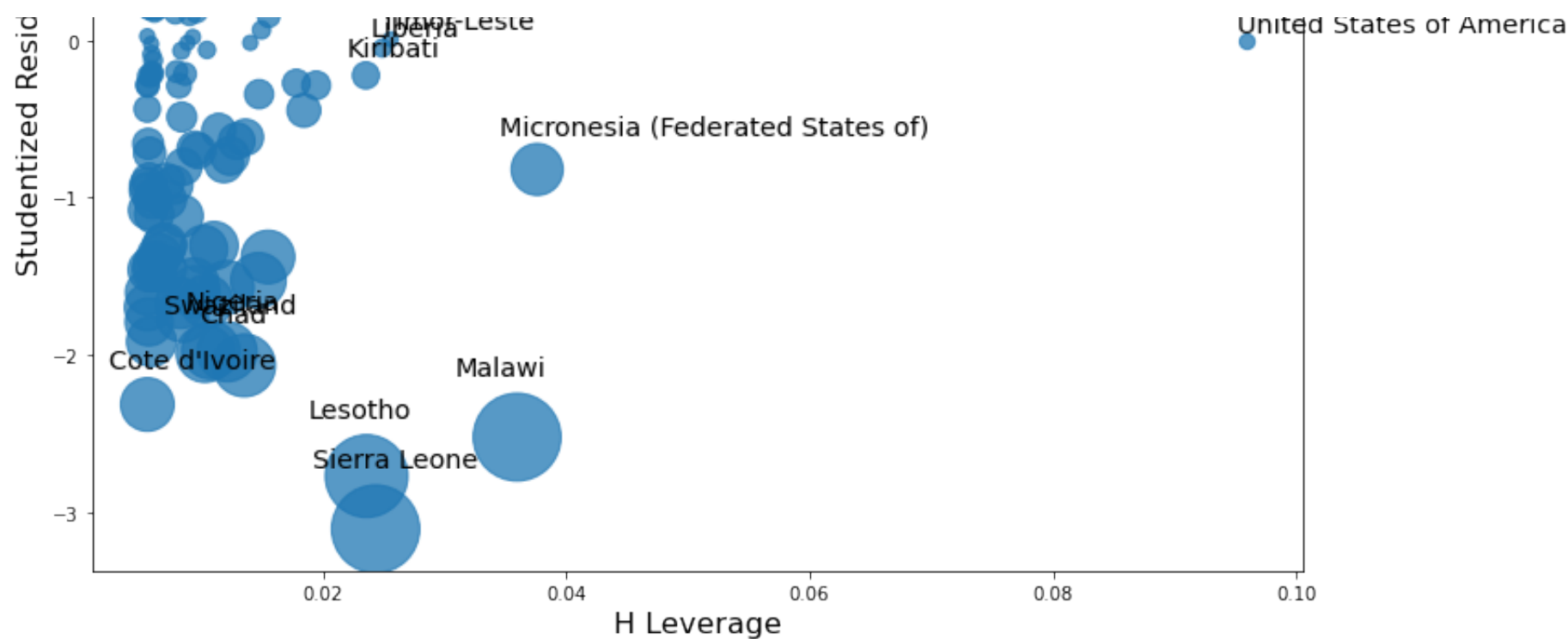


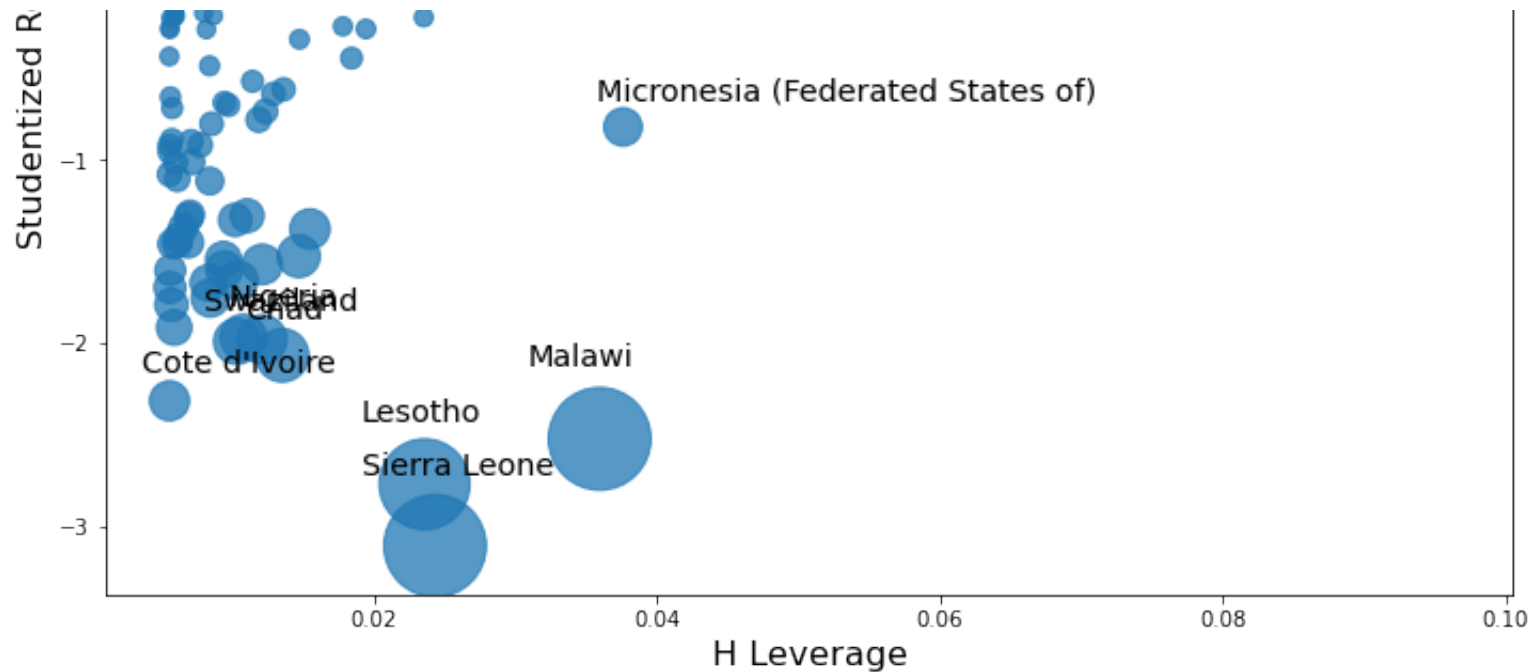


```
In [120]: figd, ax = plt.subplots(figsize=(12,8))
figd = sm.graphics.influence_plot(reg9, ax = ax, criterion="DFFITS")
figd.tight_layout(pad=1.0)

fige, ax = plt.subplots(figsize=(12,8))
fige = sm.graphics.influence_plot(reg9, ax = ax, criterion="cooks")
fige.tight_layout(pad=1.0)
```

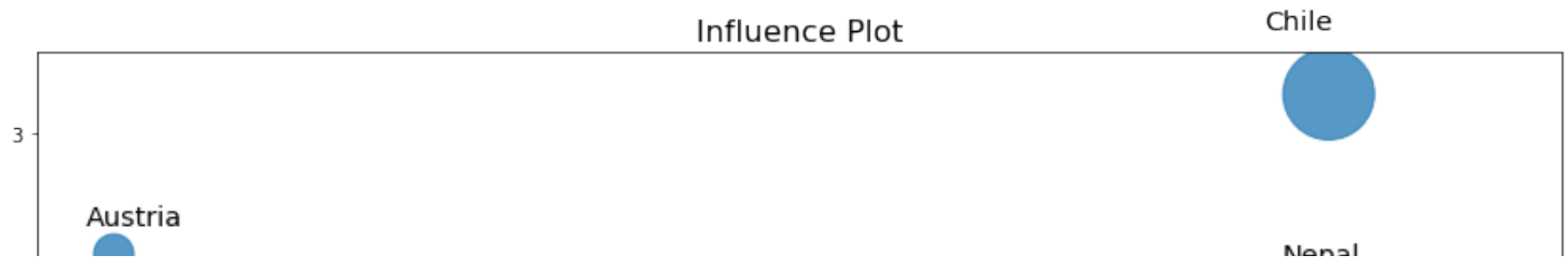


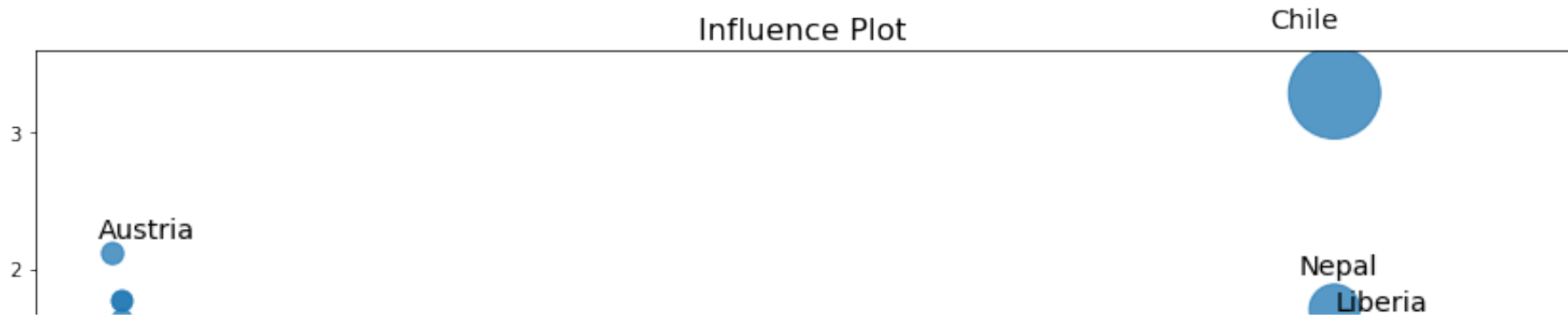
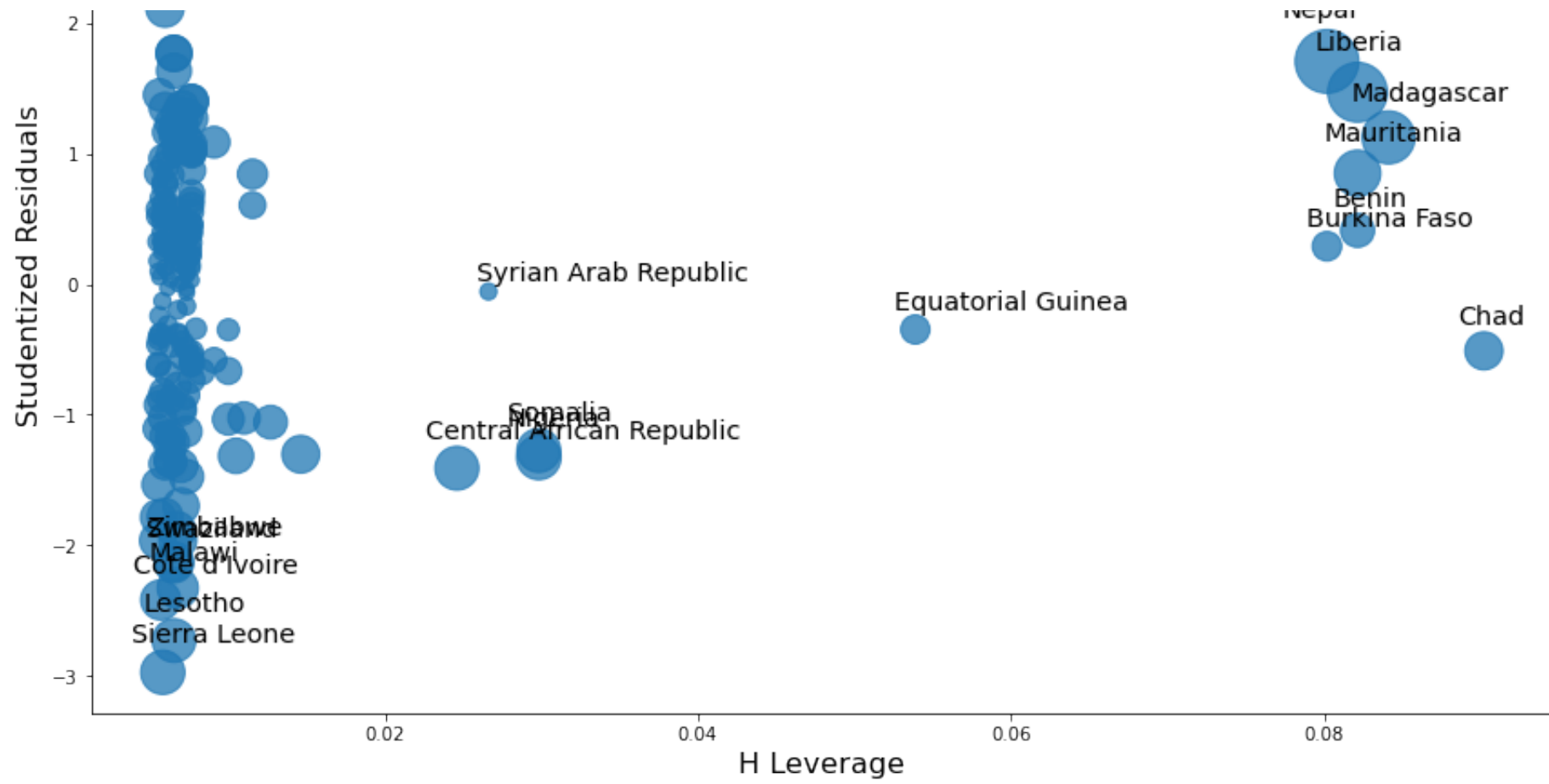


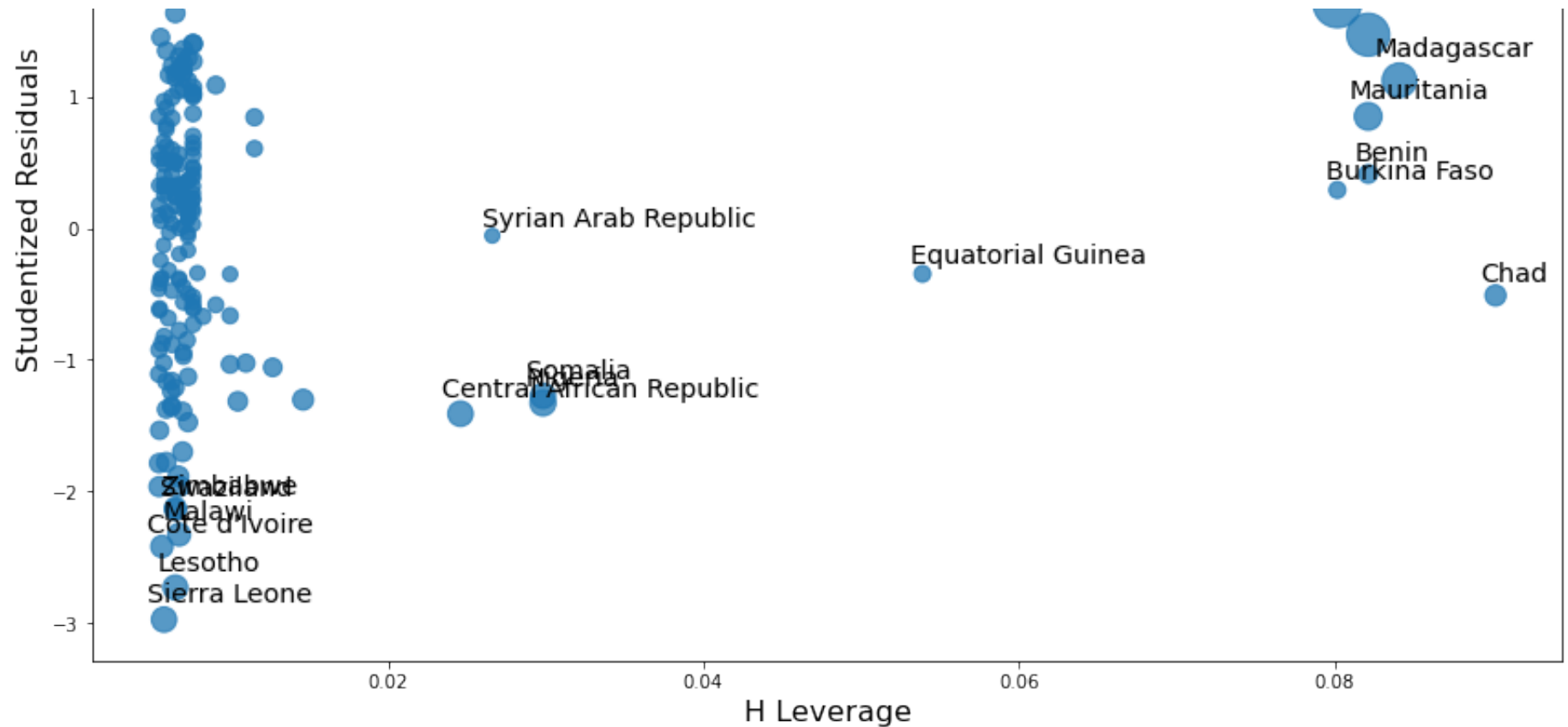


```
In [121]: figd, ax = plt.subplots(figsize=(12,8))
figd = sm.graphics.influence_plot(reg10, ax = ax, criterion="DFFITS")
figd.tight_layout(pad=1.0)

fige, ax = plt.subplots(figsize=(12,8))
fige = sm.graphics.influence_plot(reg10, ax = ax, criterion="cooks")
fige.tight_layout(pad=1.0)
```

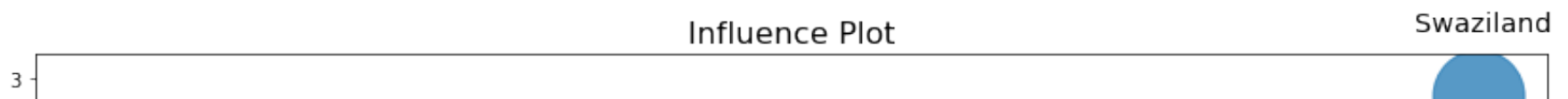


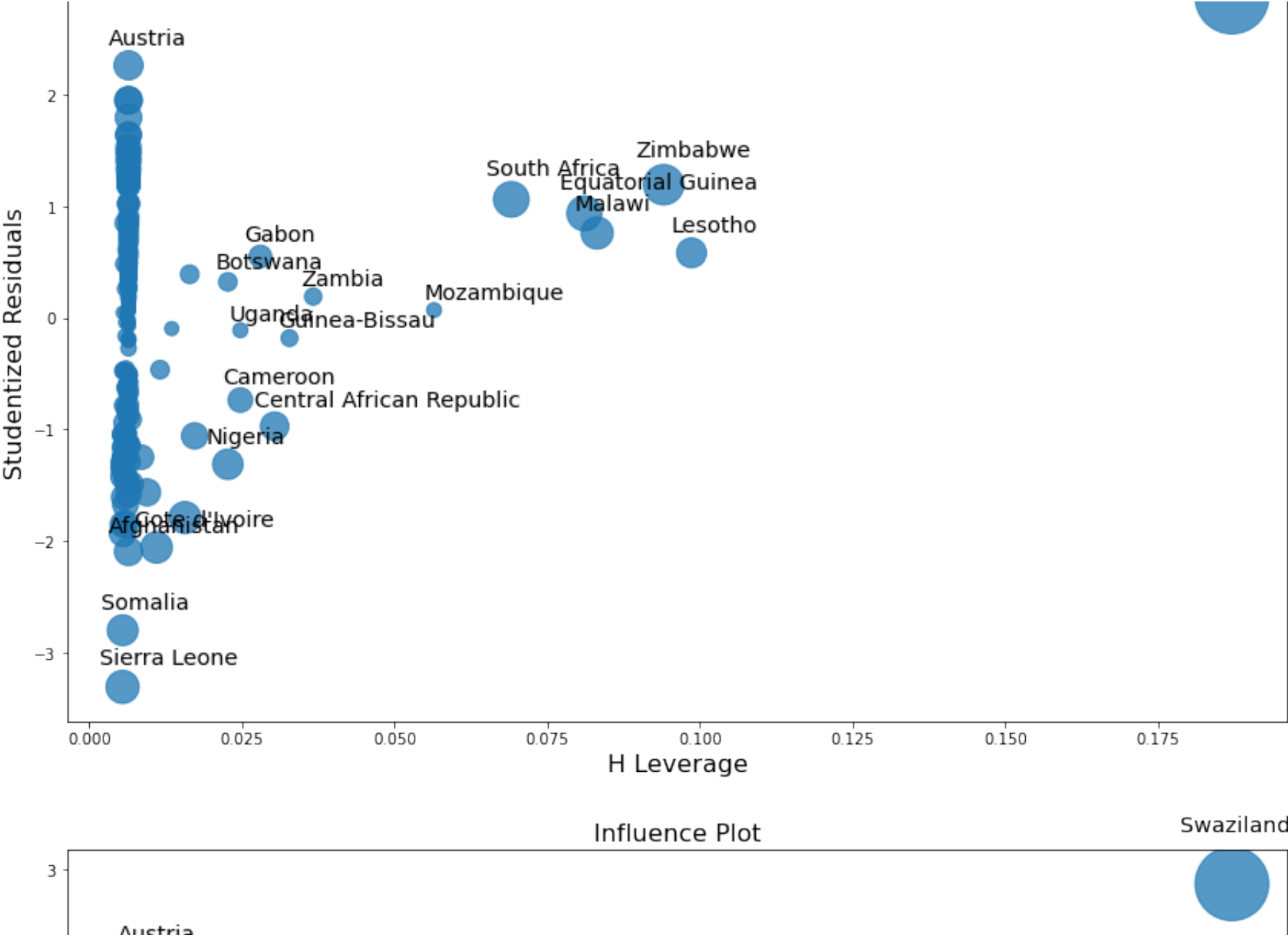


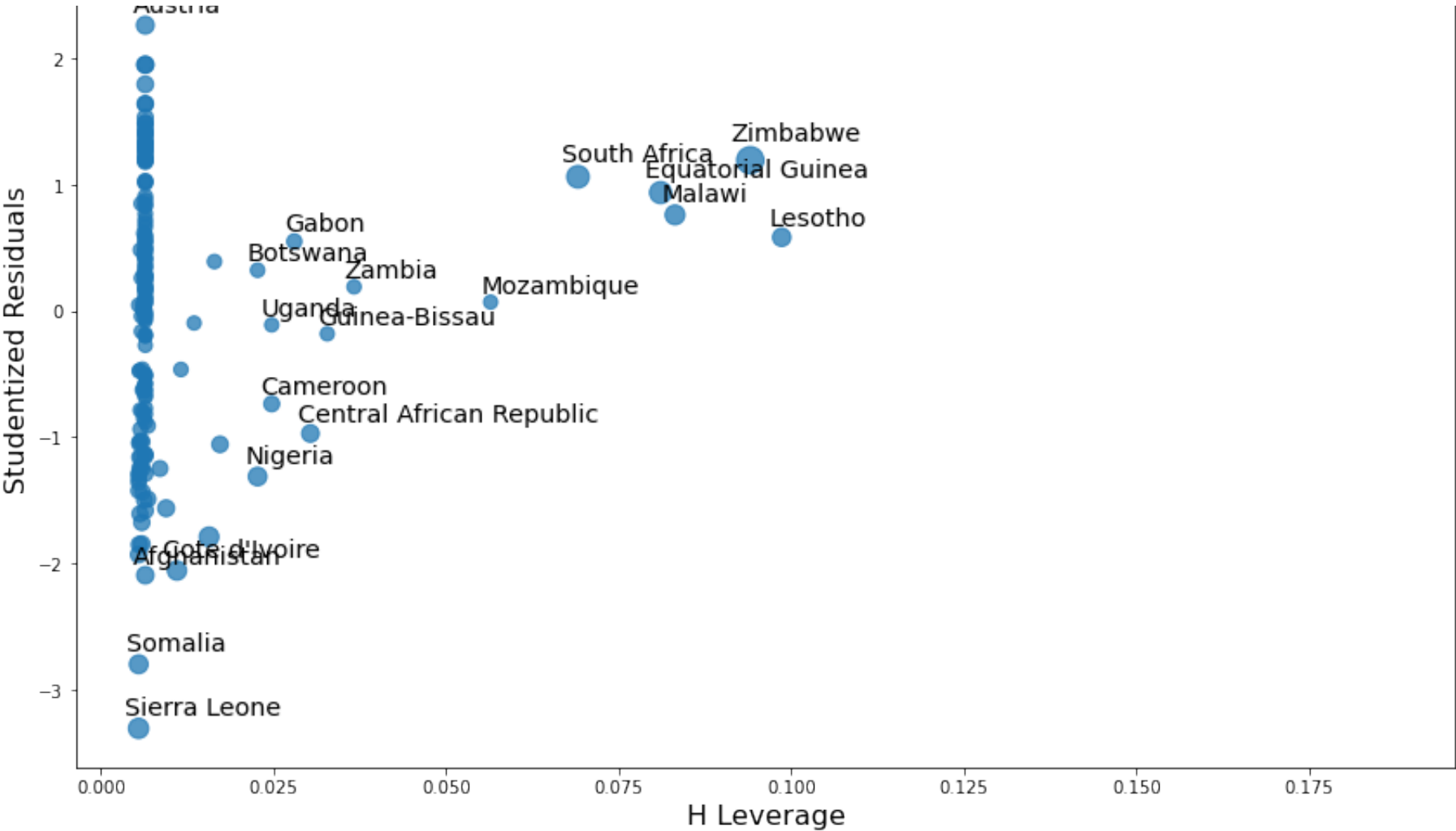


```
In [122]: figd, ax = plt.subplots(figsize=(12,8))
figd = sm.graphics.influence_plot(reg11, ax = ax, criterion="DFFITS")
figd.tight_layout(pad=1.0)

fige, ax = plt.subplots(figsize=(12,8))
fige = sm.graphics.influence_plot(reg11, ax = ax, criterion="cooks")
fige.tight_layout(pad=1.0)
```



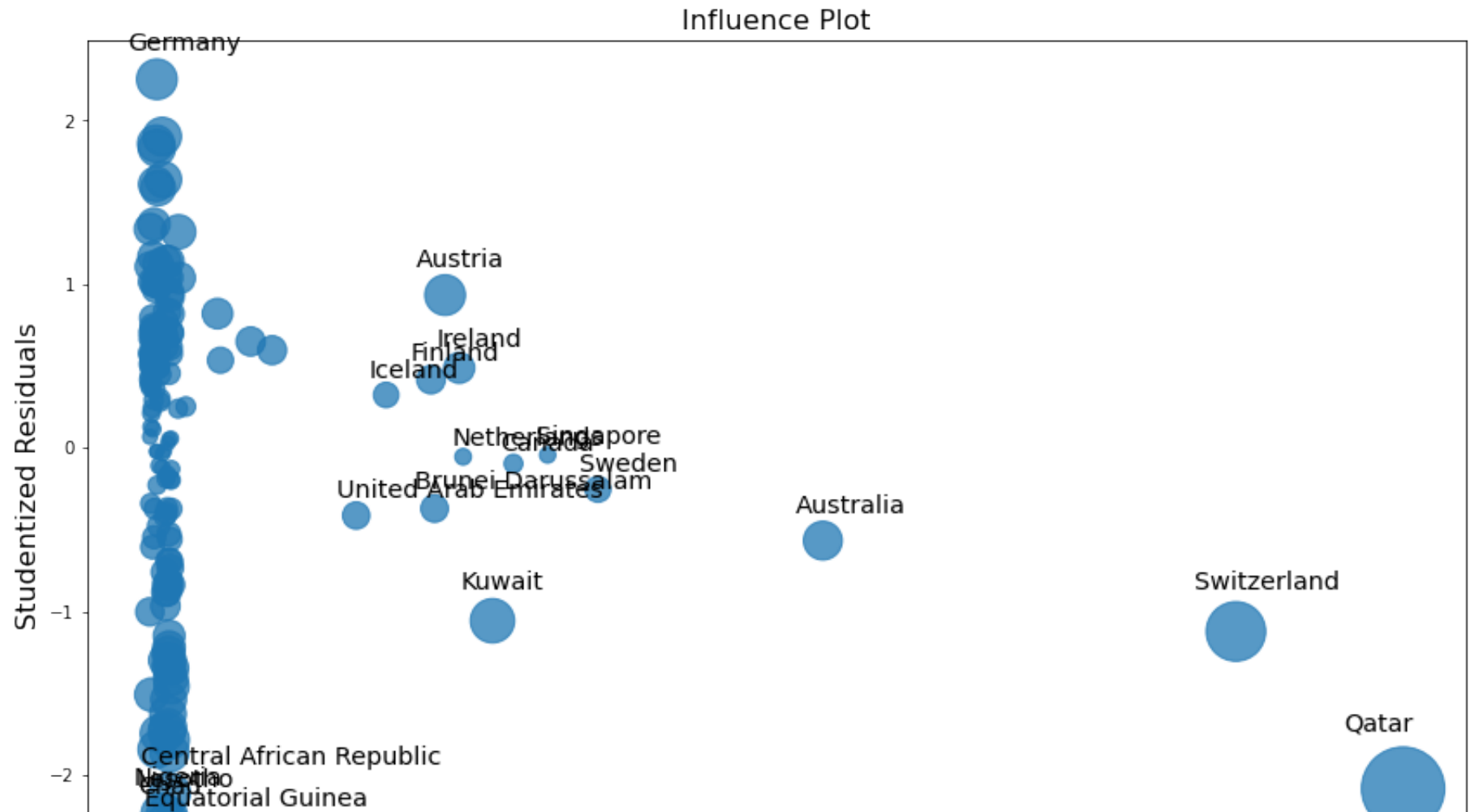


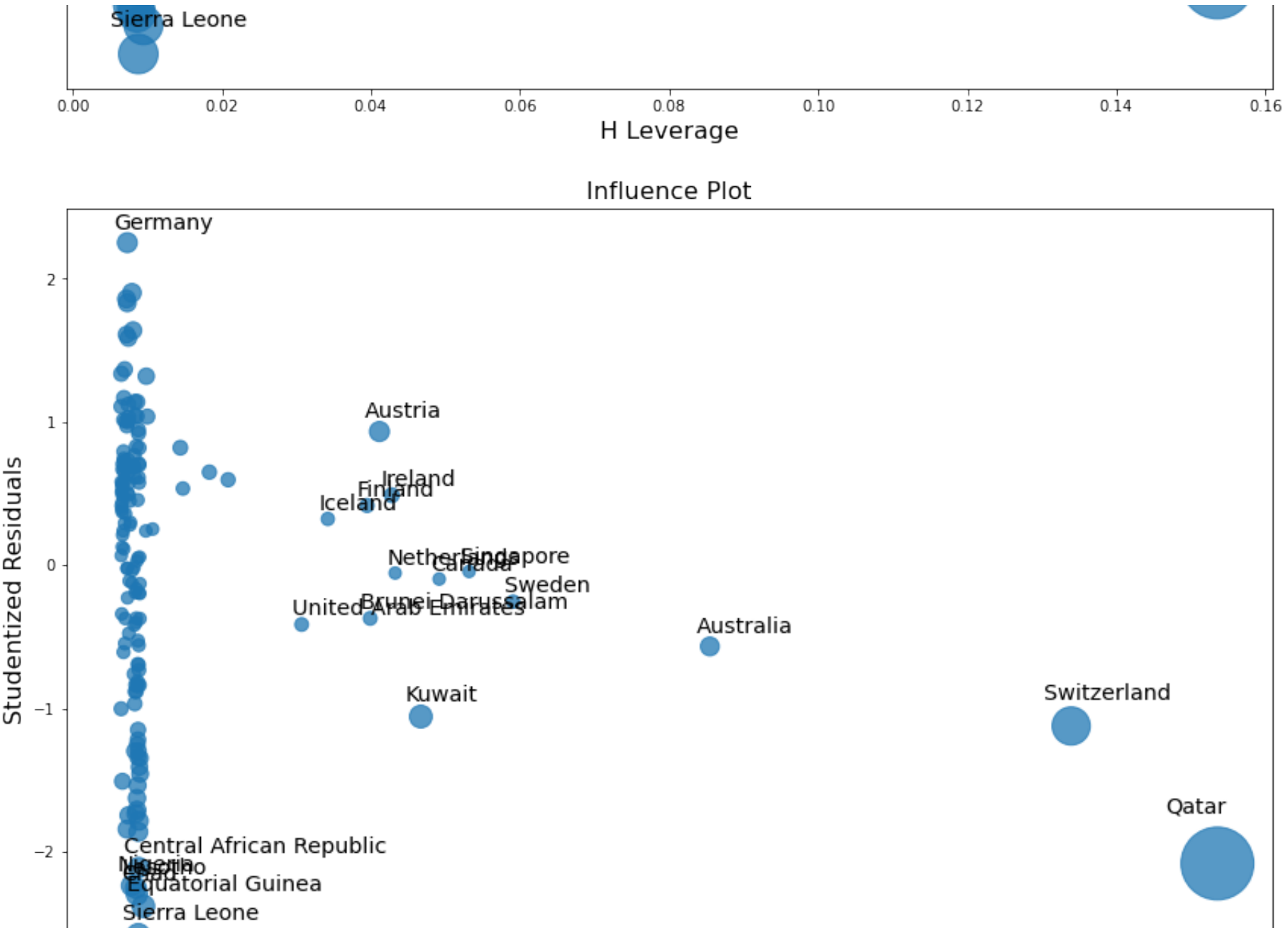


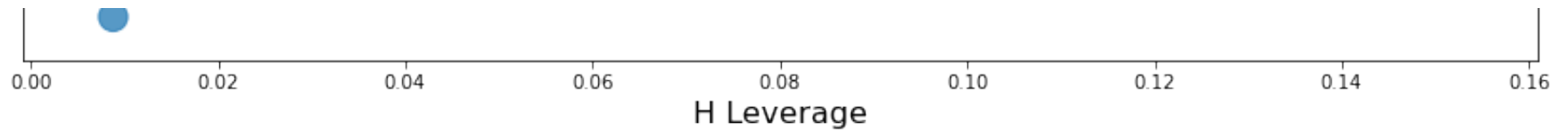
In [123]:

```
figd, ax = plt.subplots(figsize=(12,8))
figd = sm.graphics.influence_plot(reg12, ax = ax, criterion="DFFITS")
figd.tight_layout(pad=1.0)

fige, ax = plt.subplots(figsize=(12,8))
fige = sm.graphics.influence_plot(reg12, ax = ax, criterion="cooks")
fige.tight_layout(pad=1.0)
```







```
In [124]: figd, ax = plt.subplots(figsize=(12,8))
figd = sm.graphics.influence_plot(reg13, ax = ax, criterion="DFFITS")
figd.tight_layout(pad=1.0)

fige, ax = plt.subplots(figsize=(12,8))
fige = sm.graphics.influence_plot(reg13, ax = ax, criterion="cooks")
fige.tight_layout(pad=1.0)
```

