

```
In [2]: import warnings
warnings.filterwarnings("ignore")
import pandas as pd
import numpy as np
import wbapi as wb
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.cluster import AgglomerativeClustering
import matplotlib.pyplot as plt
from kneed import KneeLocator
from sklearn.preprocessing import normalize
import scipy.cluster.hierarchy as shc
```

Clustering (K-Means)

```
In [3]: country_codes = ['CHN', 'IRQ', 'IND']
ind1=["EN.ATM.CO2E.KT"] # CO2 Emission
ind2=["EN.ATM.METH.KT.CE"] # Mithene Emission
```

```
In [4]: df1 = wb.data.DataFrame(ind1, country_codes, mrv=20).T
df1=df1.fillna(df1.mean())
df1.head()
```

```
Out[4]:
```

| economy | CHN | IND | IRQ |
|---------|-----------|-----------|---------|
| YR1999 | 3149200.0 | 904090.0 | 64610.0 |
| YR2000 | 3344090.0 | 940170.0 | 74570.0 |
| YR2001 | 3526750.0 | 953880.0 | 84300.0 |
| YR2002 | 3808330.0 | 987530.0 | 79410.0 |
| YR2003 | 4413300.0 | 1015890.0 | 72080.0 |

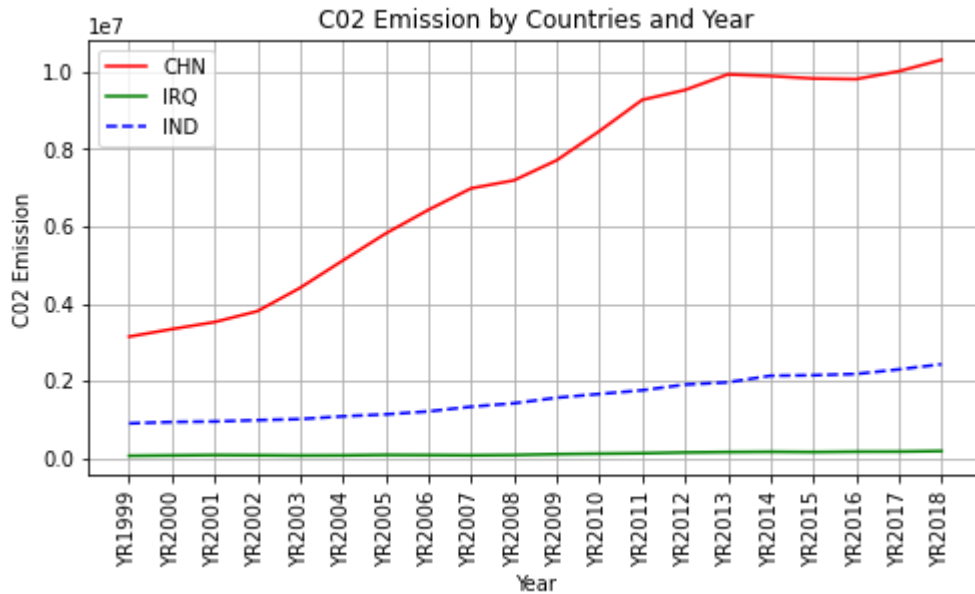
```
In [5]: df2 = wb.data.DataFrame(ind2, country_codes, mrv=20).T
df2=df2.fillna(df2.mean())
df2.head()
```

```
Out[5]:
```

| economy | CHN | IND | IRQ |
|---------|----------|----------|---------|
| YR1999 | 825690.0 | 584490.0 | 10760.0 |
| YR2000 | 829610.0 | 589050.0 | 11050.0 |
| YR2001 | 824170.0 | 594700.0 | 11140.0 |
| YR2002 | 819890.0 | 589390.0 | 11400.0 |
| YR2003 | 815910.0 | 598510.0 | 10830.0 |

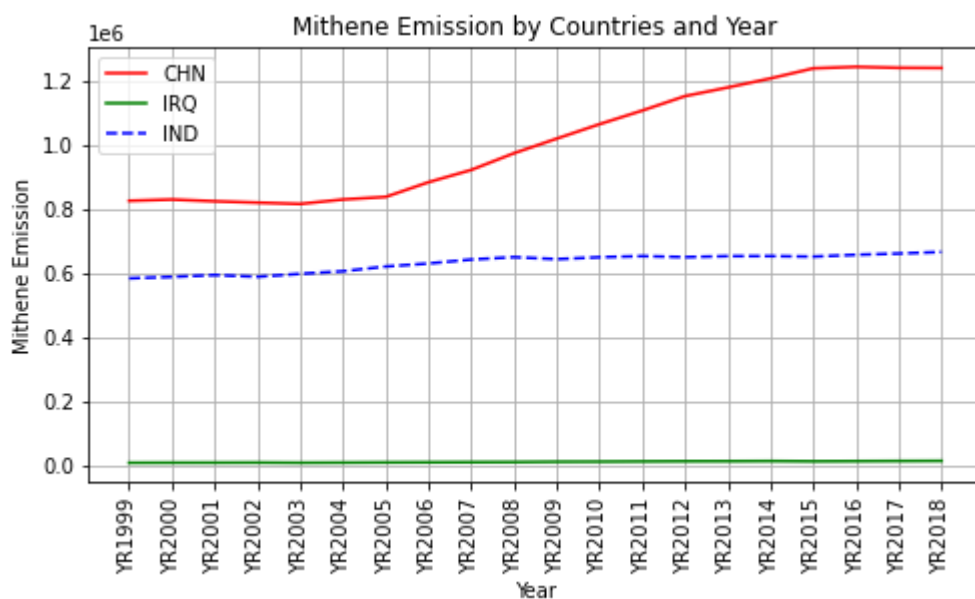
```
In [6]: plt.figure(figsize=(8,4))
plt.title('CO2 Emission by Countries and Year')
plt.plot(df1['CHN'], "r-", label="CHN")
plt.plot(df1['IRQ'], "g-", label="IRQ")
plt.plot(df1['IND'], "b--", label="IND")
plt.xlabel("Year")
```

```
plt.xticks(rotation=90)
plt.ylabel("CO2 Emission")
plt.legend(loc="upper left")
plt.grid()
plt.show()
```



In [7]:

```
plt.figure(figsize=(8,4))
plt.title('Mithene Emission by Countries and Year')
plt.plot(df2['CHN'], "r-", label="CHN")
plt.plot(df2['IRQ'], "g-", label="IRQ")
plt.plot(df2['IND'], "b--", label="IND")
plt.xlabel("Year")
plt.xticks(rotation=90)
plt.ylabel("Mithene Emission")
plt.grid()
plt.legend(loc="best")
plt.show()
```



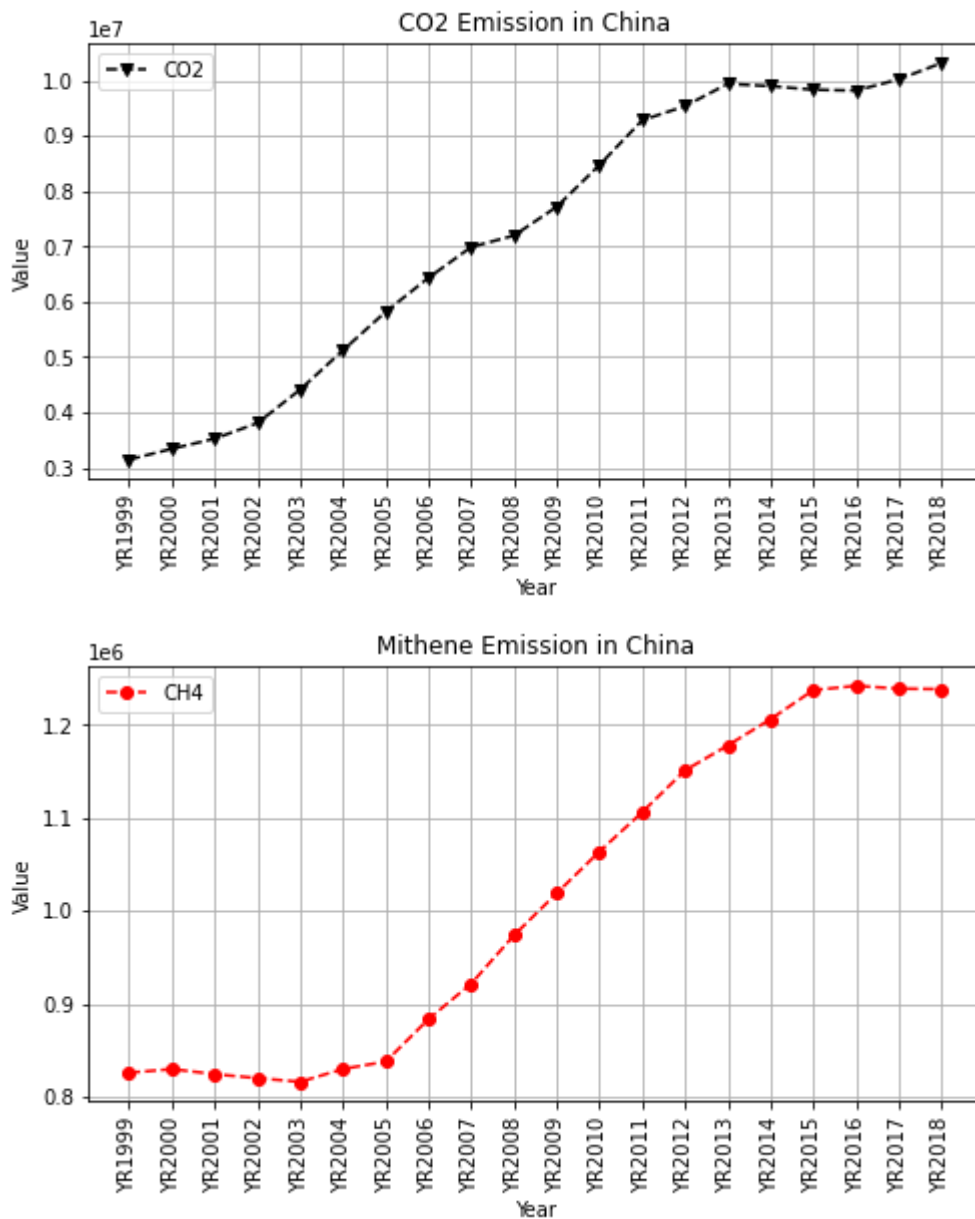
In [8]:

```
plt.figure(figsize=(8,4))
plt.title('CO2 Emission in China')
plt.plot(df1['CHN'], "kv--", label="CO2")
```

```

plt.xlabel("Year")
plt.xticks(rotation=90)
plt.ylabel("Value")
plt.legend(loc="upper left")
plt.grid()
plt.show()
plt.figure(figsize=(8,4))
plt.title('Mithene Emission in China')
plt.plot(df2['CHN'], "ro--", label="CH4")
plt.xlabel("Year")
plt.xticks(rotation=90)
plt.ylabel("Value")
plt.legend(loc="upper left")
plt.grid()
plt.show()

```

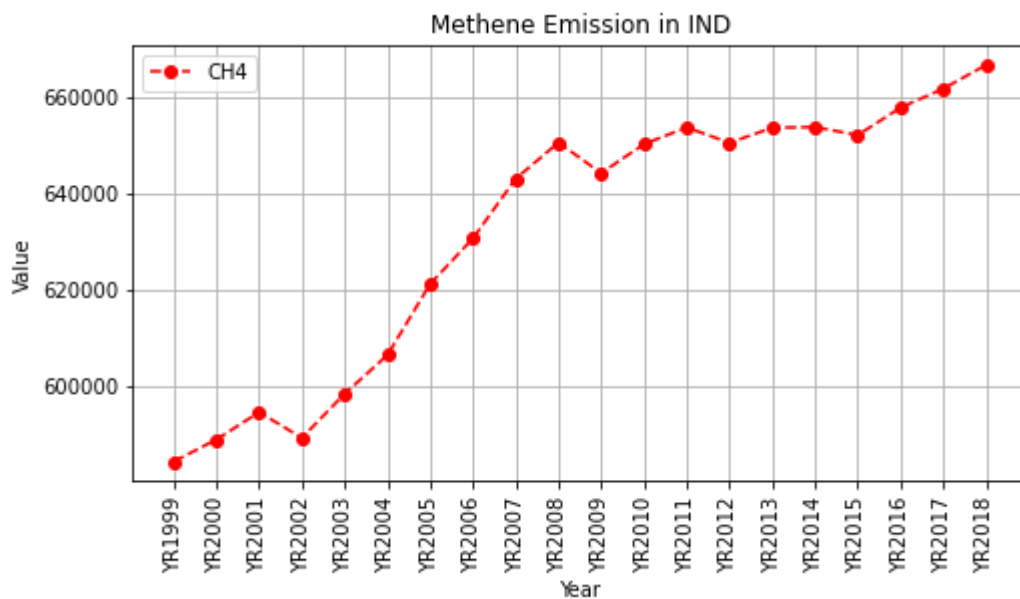
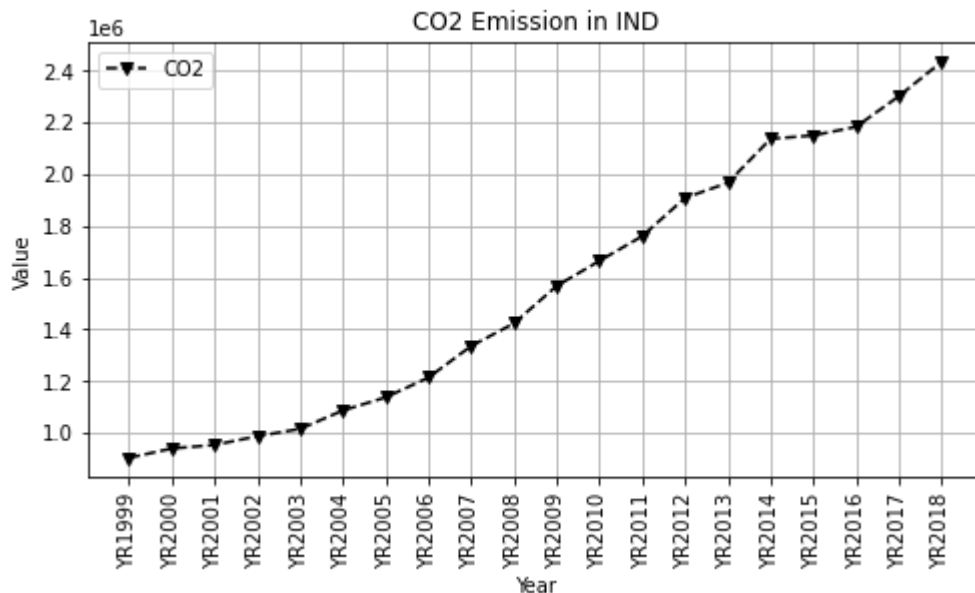


```

In [9]: plt.figure(figsize=(8,4))
plt.title('CO2 Emission in IND')
plt.plot(df1['IND'], "kv--", label="CO2")
plt.xlabel("Year")
plt.xticks(rotation=90)
plt.ylabel("Value")
plt.legend(loc="upper left")

```

```
plt.grid()
plt.show()
plt.figure(figsize=(8,4))
plt.title('Methene Emission in IND')
plt.plot(df2['IND'], "ro--", label="CH4")
plt.xlabel("Year")
plt.xticks(rotation=90)
plt.ylabel("Value")
plt.legend(loc="upper left")
plt.grid()
plt.show()
```

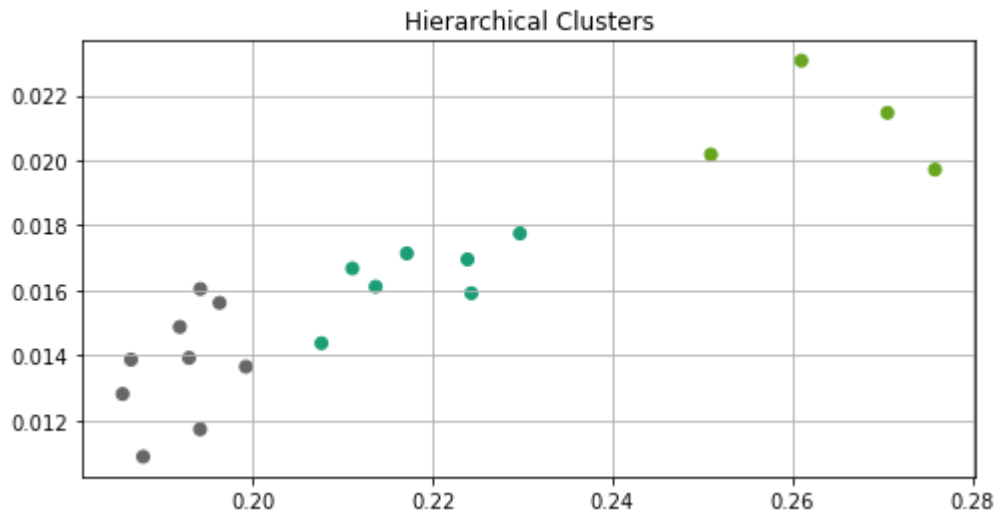


```
In [10]: def normlz(data):
          nm=normalize(data)
          return nm
          data=normlz(df1.values)
```

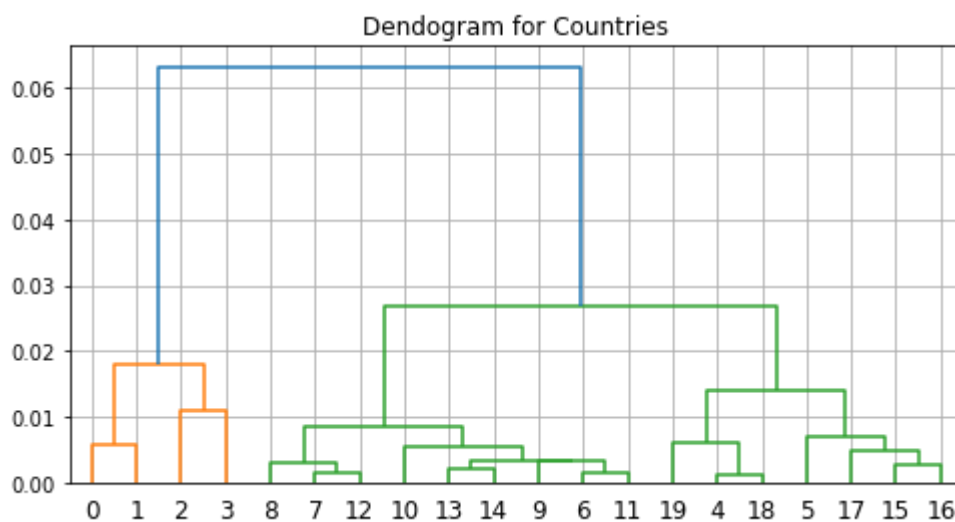
```
In [11]: aggl = AgglomerativeClustering(n_clusters=3, affinity='euclidean', linka
        yhat = aggl.fit_predict(data)
        clusters = np.unique(yhat)
        clusters
```

```
Out[11]: array([0, 1, 2], dtype=int64)
```

```
In [12]: plt.figure(figsize=(8,4))
plt.title('Hierarchical Clusters')
plt.scatter(data[:,1],data[:,2], c=aggl.labels_, cmap='Dark2')
plt.grid()
plt.show()
```



```
In [16]: plt.figure(figsize=(8,4))
plt.title("Dendrogram for Countries")
dend = shc.dendrogram(shc.linkage(data, method='centroid'))
plt.grid()
plt.show()
```



Curve Fitting

```
In [18]: from scipy.optimize import curve_fit
#!pip install lmfit
from lmfit import Model
```

```
In [19]: def gaussian(x, amp, cen, wid):
return (amp / (np.sqrt(2*np.pi) * wid)) * np.exp(-(x-cen)**2 / (2*wid
```

```
In [20]: norml = normlz(df2.values)
y = gaussian(norml[:,1], 2.33, 0.21, 1.51) + np.random.normal(0, 0.2, no
init_vals = [1,0,1]
best_vals, covar = curve_fit(gaussian, norml[:,1], y, p0=init_vals,maxfe
gaus_model = Model(gaussian)
res = gaus_model.fit(y, x=norml[:,2], amp=5, cen=5, wid=1)
print(res.fit_report())
```

```
[[Model]]
      Model(gaussian)
[[Fit Statistics]]
   # fitting method   = leastsq
   # function evals   = 53
   # data points      = 20
   # variables        = 3
   chi-square         = 0.86097405
   reduced chi-square = 0.05064553
   Akaike info crit   = -56.9084637
   Bayesian info crit = -53.9212669
## Warning: uncertainties could not be estimated:
[[Variables]]
   amp: 43449.6344 +/-          nan (nan%) (init = 5)
   cen: -330.729752 +/-         nan (nan%) (init = 5)
   wid: 98.4282254 +/-          nan (nan%) (init = 1)
```

```
In [21]: res
```

Out[21]: **Model**

Model(gaussian)

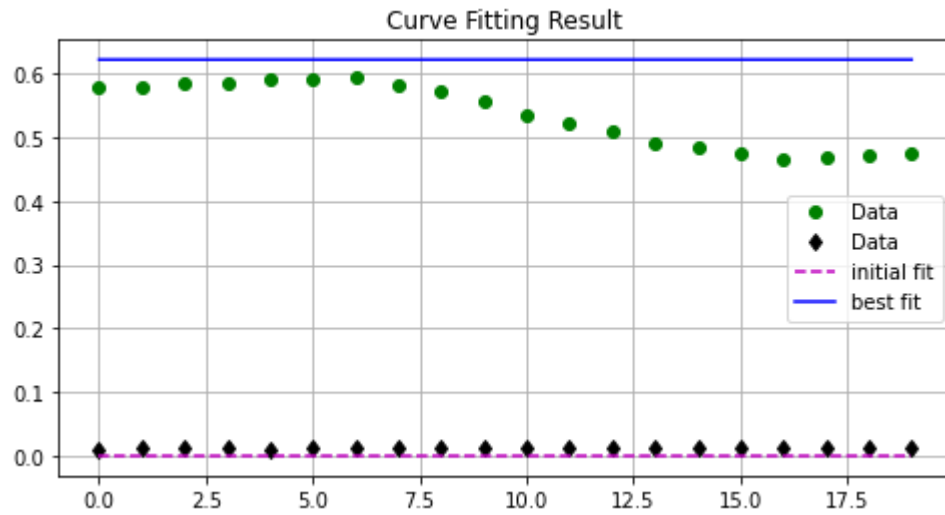
Fit Statistics

| | |
|---------------------|-------------|
| fitting method | leastsq |
| # function evals | 53 |
| # data points | 20 |
| # variables | 3 |
| chi-square | 0.86097405 |
| reduced chi-square | 0.05064553 |
| Akaike info crit. | -56.9084637 |
| Bayesian info crit. | -53.9212669 |

Variables

| name | value | standard error | relative error | initial value | min | max | vary |
|------|-------------|----------------|----------------|---------------|------|-----|------|
| amp | 43449.6344 | nan | (nan%) | 5 | -inf | inf | True |
| cen | -330.729752 | nan | (nan%) | 5 | -inf | inf | True |
| wid | 98.4282254 | nan | (nan%) | 1 | -inf | inf | True |

```
In [22]: plt.figure(figsize=(8,4))
plt.title('Curve Fitting Result')
plt.plot(norml[:,1], "go", label="Data")
plt.plot(norml[:,2], "kd", label="Data")
plt.plot(res.init_fit, 'm--', label='initial fit')
plt.plot(res.best_fit, 'b-', label='best fit')
plt.legend()
plt.grid()
plt.show()
```



```
In [ ]:
```