

MN_median_Clustering

May 3, 2024

```
[1]: !pip install scikit-learn-extra
```

```
Requirement already satisfied: scikit-learn-extra in  
c:\users\tky12\anaconda3\lib\site-packages (0.3.0)  
Requirement already satisfied: numpy>=1.13.3 in  
c:\users\tky12\anaconda3\lib\site-packages (from scikit-learn-extra) (1.26.4)  
Requirement already satisfied: scipy>=0.19.1 in  
c:\users\tky12\anaconda3\lib\site-packages (from scikit-learn-extra) (1.11.4)  
Requirement already satisfied: scikit-learn>=0.23.0 in  
c:\users\tky12\anaconda3\lib\site-packages (from scikit-learn-extra) (1.2.2)  
Requirement already satisfied: joblib>=1.1.1 in  
c:\users\tky12\anaconda3\lib\site-packages (from scikit-learn>=0.23.0->scikit-  
learn-extra) (1.2.0)  
Requirement already satisfied: threadpoolctl>=2.0.0 in  
c:\users\tky12\anaconda3\lib\site-packages (from scikit-learn>=0.23.0->scikit-  
learn-extra) (2.2.0)
```

```
[2]: import pandas as pd  
import seaborn as sns  
import matplotlib.pyplot as plt  
import numpy as np  
from mpl_toolkits.mplot3d import Axes3D  
from scipy.stats import multivariate_normal  
from scipy.stats import norm  
import plotly.express as px  
from sklearn.preprocessing import MinMaxScaler  
from sklearn.metrics import silhouette_score  
import pickle
```

```
[3]: # Surpress warnings  
def warn(*args, **kwargs):  
    pass  
import warnings  
warnings.warn = warn
```

```
[4]: pd.DataFrame.iteritems = pd.DataFrame.items
```

```
[5]: # Retrieve dataset and read first 5 rows
macroNutrient_median =pd.read_csv(r"../Dataset/Dataset_for_EDA/
↳macroNutrient_median.csv", encoding= 'unicode_escape')
macroNutrient_median
```

```
[5]:
```

	No.	Description	Category \
0	15155	ABALONE,MIXED SPECIES,RAW	ABALONE
1	15156	ABALONE,MXD SP,CKD,FRIED	ABALONE
2	9427	ABIYUCH,RAW	ABIYUCH
3	9002	ACEROLA JUICE,RAW	ACEROLA JUICE
4	9001	ACEROLA,(WEST INDIAN CHERRY),RAW	ACEROLA
...
8785	1119	YOGURT,VANILLA,LOFAT,11 GRAMS PROT PER 8 OZ	YOGURT
8786	1220	YOGURT,VANILLA,LOFAT,11 GRAMS PROT PER 8 OZ,FO...	YOGURT
8787	1295	YOGURT,VANILLA,NON-FAT	YOGURT
8788	16004	YOKAN,PREP FROM ADZUKI BNS & SUGAR	YOKAN
8789	3217	ZWIEBACK	ZWIEBACK

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g) \
0	105	6.01	17.10	0.76
1	189	11.05	19.63	6.78
2	69	17.60	1.50	0.10
3	23	4.80	0.40	0.30
4	32	7.69	0.40	0.30
...
8785	85	13.80	4.93	1.25
8786	85	13.80	4.93	1.25
8787	78	17.04	2.94	0.00
8788	260	60.72	3.29	0.12
8789	426	74.20	10.10	9.70

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g) \
0	0.1070	0.104
1	2.7410	1.676
2	1.9385	0.696
3	0.0820	0.090
4	0.0820	0.090
...
8785	0.3430	0.036
8786	0.3430	0.036
8787	0.0000	0.000
8788	0.0110	0.026
8789	4.2440	2.073

	Saturated Fatty Acids(g)
0	0.149
1	1.646

```

2          0.014
3          0.068
4          0.068
...
8785       0.806
8786       0.806
8787       0.000
8788       0.043
8789       2.525

```

[8790 rows x 10 columns]

```
[6]: macroNutrient_median.info()
```

```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 8790 entries, 0 to 8789
Data columns (total 10 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   No.                                    8790 non-null   int64
1   Description                            8790 non-null   object
2   Category                              8790 non-null   object
3   Energy (Kcal)                         8790 non-null   int64
4   Carbohydrate(g)                      8790 non-null   float64
5   Protein(g)                           8790 non-null   float64
6   Total Lipid(g)                       8790 non-null   float64
7   Monounsaturated Fatty Acids(g)        8790 non-null   float64
8   Polyunsaturated Fatty Acids(g)        8790 non-null   float64
9   Saturated Fatty Acids(g)              8790 non-null   float64
dtypes: float64(6), int64(2), object(2)
memory usage: 686.8+ KB

```

```
[7]: macroNutrient_median.describe()
```

```

[7]:
count      No.  Energy (Kcal)  Carbohydrate(g)  Protein(g)  \
count      8790.000000      8790.000000      8790.000000      8790.000000
mean      15663.495222      226.317634       22.127710       11.342849
std       9251.413586      169.877539       27.270822       10.530474
min       1001.000000        0.000000        0.000000        0.000000
25%       9086.250000        91.000000        0.050000        2.380000
50%      14427.500000       191.000000        9.340000        8.000000
75%      20142.750000       337.000000       34.910000       19.880000
max      93600.000000       902.000000      100.000000       88.320000

count      Total Lipid(g)  Monounsaturated Fatty Acids(g)  \
count      8790.000000      8790.000000
mean       10.553725        3.918673
std        15.814842        6.763709

```

min	0.000000	0.000000
25%	0.950000	0.304000
50%	5.140000	1.938500
75%	13.720000	4.759000
max	100.000000	83.689000

	Polyunsaturated Fatty Acids(g)	Saturated Fatty Acids(g)
count	8790.000000	8790.000000
mean	2.168267	3.497547
std	5.032118	6.399843
min	0.000000	0.000000
25%	0.249000	0.259000
50%	0.696000	1.592000
75%	1.880000	4.173750
max	74.623000	95.600000

```
[8]: macroNutrient_median.head()
```

```
[8]:
```

	No.	Description	Category	Energy (Kcal)	\
0	15155	ABALONE,MIXED SPECIES,RAW	ABALONE	105	
1	15156	ABALONE,MXD SP,CKD,FRIED	ABALONE	189	
2	9427	ABIYUCH,RAW	ABIYUCH	69	
3	9002	ACEROLA JUICE,RAW	ACEROLA JUICE	23	
4	9001	ACEROLA,(WEST INDIAN CHERRY),RAW	ACEROLA	32	

	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
0	6.01	17.10	0.76	
1	11.05	19.63	6.78	
2	17.60	1.50	0.10	
3	4.80	0.40	0.30	
4	7.69	0.40	0.30	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
0	0.1070	0.104	
1	2.7410	1.676	
2	1.9385	0.696	
3	0.0820	0.090	
4	0.0820	0.090	

	Saturated Fatty Acids(g)
0	0.149
1	1.646
2	0.014
3	0.068
4	0.068

```
[9]: float_columns = [x for x in macroNutrient_median.columns if x not in ['No.', 'Description', 'Category']]
```

```
[10]: skew_columns = (macroNutrient_median[float_columns]
                      .skew()
                      .sort_values(ascending=False))

skew_columns = skew_columns.loc[skew_columns > 0.75]
print("{} of the 7 columns are skewed with the vast majority being heavily_
skewed".format(len(skew_columns)))
skew_columns
```

7 of the 7 columns are skewed with the vast majority being heavily skewed

```
[10]: Saturated Fatty Acids(g)          6.722778
Polyunsaturated Fatty Acids(g)        6.401464
Monounsaturated Fatty Acids(g)        4.764472
Total Lipid(g)                        3.309724
Protein(g)                            1.166368
Energy (Kcal)                         1.148610
Carbohydrate(g)                       1.127598
dtype: float64
```

```
[11]: # Perform log transform on skewed columns
for col in skew_columns.index.tolist():
    macroNutrient_median[col] = np.log1p(macroNutrient_median[col])
```

```
[12]: macroNutrient_median[float_columns]
```

```
[12]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
0	4.663439	1.947338	2.895912	0.565314	
1	5.247024	2.489065	3.026746	2.051556	
2	4.248495	2.923162	0.916291	0.095310	
3	3.178054	1.757858	0.336472	0.262364	
4	3.496508	2.162173	0.336472	0.262364	
...	
8785	4.454347	2.694627	1.780024	0.810930	
8786	4.454347	2.694627	1.780024	0.810930	
8787	4.369448	2.892592	1.371181	0.000000	
8788	5.564520	4.122608	1.456287	0.113329	
8789	6.056784	4.320151	2.406945	2.370244	
	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)			\
0	0.101654	0.098940			
1	1.319353	0.984323			
2	1.077899	0.528273			
3	0.078811	0.086178			
4	0.078811	0.086178			

...
8785	0.294906	0.035367
8786	0.294906	0.035367
8787	0.000000	0.000000
8788	0.010940	0.025668
8789	1.657085	1.122654

	Saturated Fatty Acids(g)
0	0.138892
1	0.973049
2	0.013903
3	0.065788
4	0.065788

...	...
8785	0.591114
8786	0.591114
8787	0.000000
8788	0.042101
8789	1.259880

[8790 rows x 7 columns]

```
[13]: macroNutrient_median.describe()
```

```
[13]:
```

	No.	Energy (Kcal)	Carbohydrate(g)	Protein(g)	\
count	8790.000000	8790.000000	8790.000000	8790.000000	
mean	15663.495222	5.065342	2.137177	2.047259	
std	9251.413586	0.997539	1.615109	1.068937	
min	1001.000000	0.000000	0.000000	0.000000	
25%	9086.250000	4.521789	0.048790	1.217876	
50%	14427.500000	5.257495	2.336020	2.197225	
75%	20142.750000	5.823046	3.581016	3.038792	
max	93600.000000	6.805723	4.615121	4.492225	

	Total Lipid(g)	Monounsaturated Fatty Acids(g)	\
count	8790.000000	8790.000000	
mean	1.770891	1.126758	
std	1.183944	0.895498	
min	0.000000	0.000000	
25%	0.667829	0.265436	
50%	1.814823	1.077899	
75%	2.689207	1.750764	
max	4.615121	4.438986	

	Polyunsaturated Fatty Acids(g)	Saturated Fatty Acids(g)
count	8790.000000	8790.000000
mean	0.752763	1.054167

std	0.733516	0.871130
min	0.000000	0.000000
25%	0.222343	0.230318
50%	0.528273	0.952430
75%	1.057790	1.643598
max	4.325760	4.570579

```
[14]: scaler = MinMaxScaler()
macroNutrient_median[float_columns] = scaler.
      ↪ fit_transform(macroNutrient_median[float_columns])

macroNutrient_median.describe()
```

```
[14]:
```

	No.	Energy (Kcal)	Carbohydrate(g)	Protein(g)	\
count	8790.000000	8790.000000	8790.000000	8790.000000	
mean	15663.495222	0.744277	0.463082	0.455734	
std	9251.413586	0.146574	0.349960	0.237953	
min	1001.000000	0.000000	0.000000	0.000000	
25%	9086.250000	0.664410	0.010572	0.271107	
50%	14427.500000	0.772511	0.506167	0.489117	
75%	20142.750000	0.855610	0.775931	0.676456	
max	93600.000000	1.000000	1.000000	1.000000	

	Total Lipid(g)	Monounsaturated Fatty Acids(g)	\
count	8790.000000	8790.000000	
mean	0.383715	0.253832	
std	0.256536	0.201735	
min	0.000000	0.000000	
25%	0.144705	0.059797	
50%	0.393234	0.242826	
75%	0.582695	0.394406	
max	1.000000	1.000000	

	Polyunsaturated Fatty Acids(g)	Saturated Fatty Acids(g)
count	8790.000000	8790.000000
mean	0.174019	0.230642
std	0.169569	0.190595
min	0.000000	0.000000
25%	0.051400	0.050391
50%	0.122122	0.208383
75%	0.244533	0.359604
max	1.000000	1.000000

```
[15]: X = macroNutrient_median[float_columns]
      X
```

```
[15]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
0	0.685223	0.421947	0.644650	0.122492	
1	0.770972	0.539328	0.673774	0.444529	
2	0.624253	0.633388	0.203973	0.020652	
3	0.466968	0.380891	0.074901	0.056849	
4	0.513760	0.468498	0.074901	0.056849	
...	
8785	0.654500	0.583869	0.396246	0.175712	
8786	0.654500	0.583869	0.396246	0.175712	
8787	0.642026	0.626764	0.305234	0.000000	
8788	0.817624	0.893283	0.324179	0.024556	
8789	0.889955	0.936086	0.535802	0.513582	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
0	0.022900	0.022872	
1	0.297219	0.227549	
2	0.242826	0.122122	
3	0.017754	0.019922	
4	0.017754	0.019922	
...	
8785	0.066435	0.008176	
8786	0.066435	0.008176	
8787	0.000000	0.000000	
8788	0.002465	0.005934	
8789	0.373303	0.259528	

	Saturated Fatty Acids(g)
0	0.030388
1	0.212894
2	0.003042
3	0.014394
4	0.014394
...	...
8785	0.129330
8786	0.129330
8787	0.000000
8788	0.009211
8789	0.275650

[8790 rows x 7 columns]

0.1 Dimensionality Reduction

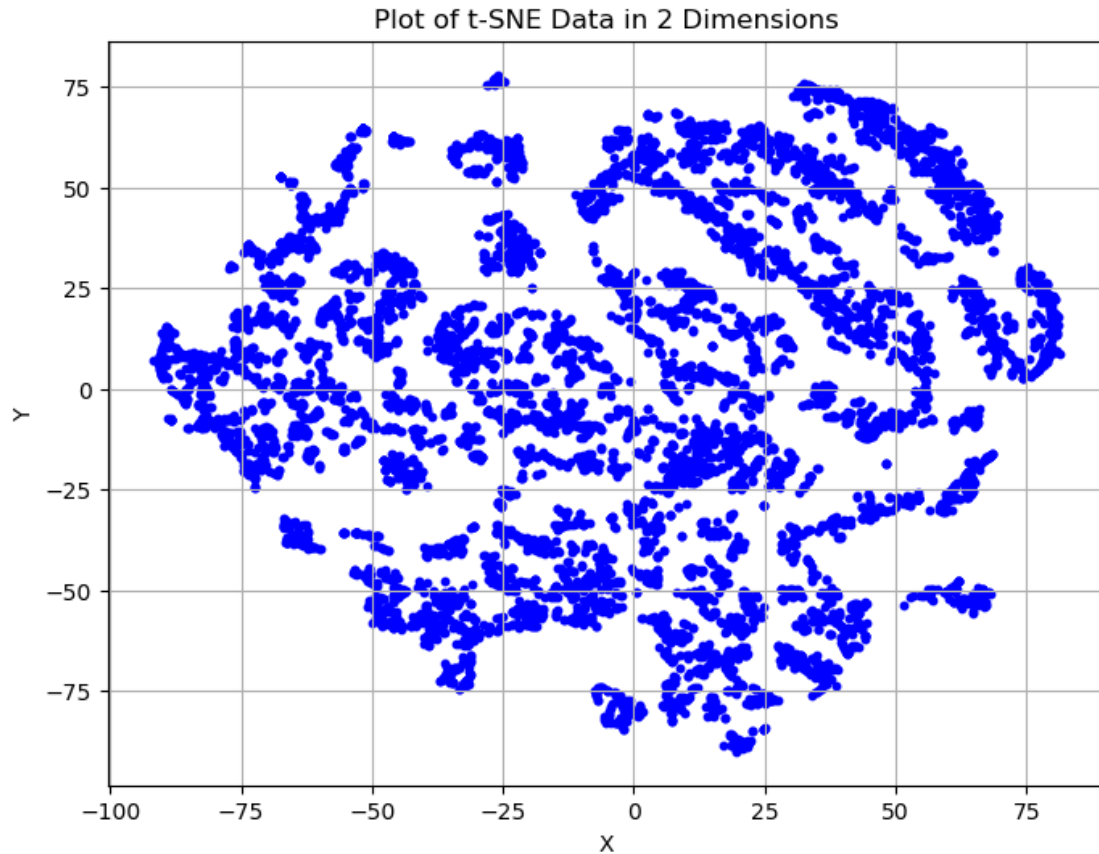
```
[16]: from clustering_function import dimensionality_reduction, scatter_plot_clustering
```

```
[17]: # Reduced to 2 dimensions using tsne
```



```
X_tsne_reduced, tsne_x_data, tsne_y_data, reduction_method =  
↳ dimensionality_reduction(X, 't-SNE', n_components=2)
```

```
[18]: scatter_plot_clustering(tsne_x_data, tsne_y_data, reduction_method)
```



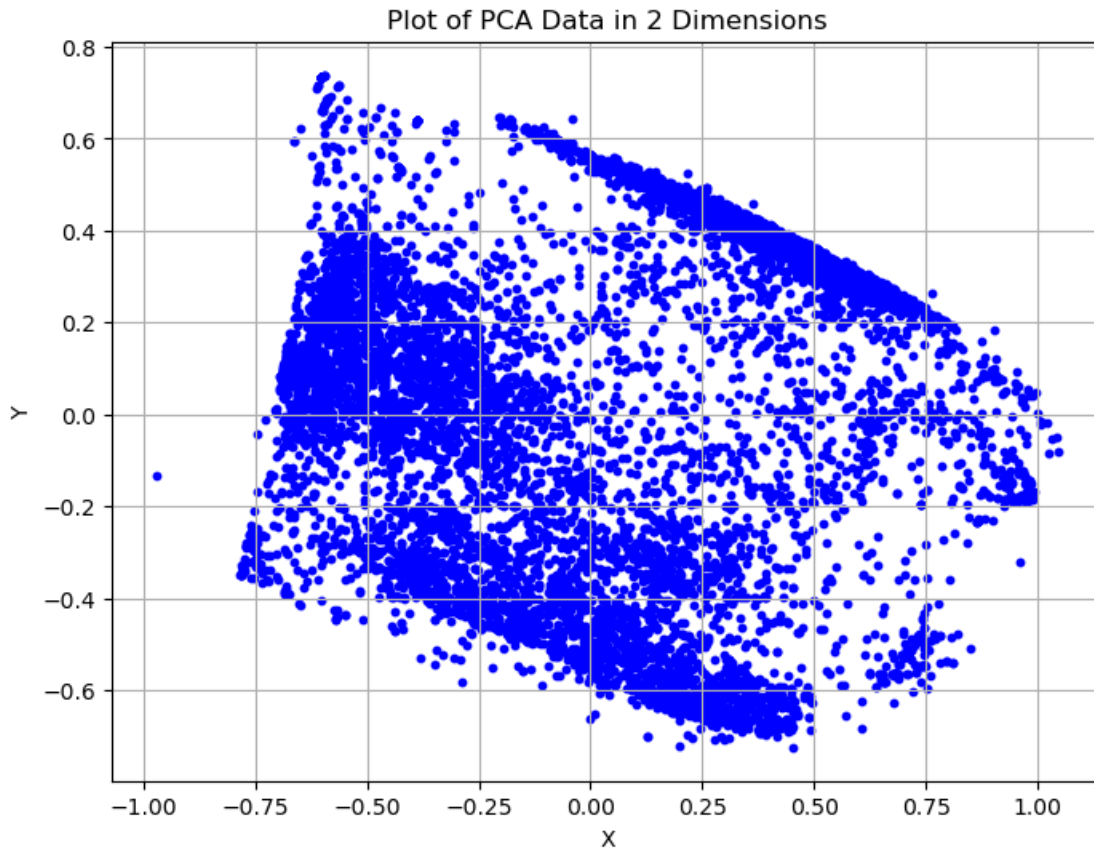
```
[19]: X_tsne_reduced
```

```
[19]: array([[ -2.8866012 ,  25.033136  ],  
          [ 10.356084 ,   0.31811064],  
          [-28.60716 ,  33.482193  ],  
          ...,  
          [-60.97991 ,  -6.97665  ],  
          [-50.850197 , -45.544563  ],  
          [  2.5995357 , -35.31806   ]], dtype=float32)
```

```
[20]: pd.DataFrame(X_tsne_reduced).to_csv("../Dataset/Dataset_for_reduced_data/  
↳ median_reduced.csv")
```

```
[21]: # Reduced to 2 dimensions using pca
X_pca_reduced, pca_x_data, pca_y_data, reduction_method = ↳
dimensionality_reduction(X, 'PCA', n_components=2)
```

```
[22]: scatter_plot_clustering(pca_x_data, pca_y_data, reduction_method)
```



```
[23]: X_pca_reduced
```

```
[23]: array([[ -0.28583447,  0.23920712],
 [  0.11334472, -0.07742649],
 [ -0.48422218, -0.01451935],
 ...,
 [ -0.58623631,  0.0718748 ],
 [ -0.62047369, -0.2098833 ],
 [  0.04574319, -0.51445591]])
```

0.2 K Means: Optimal number of clusters

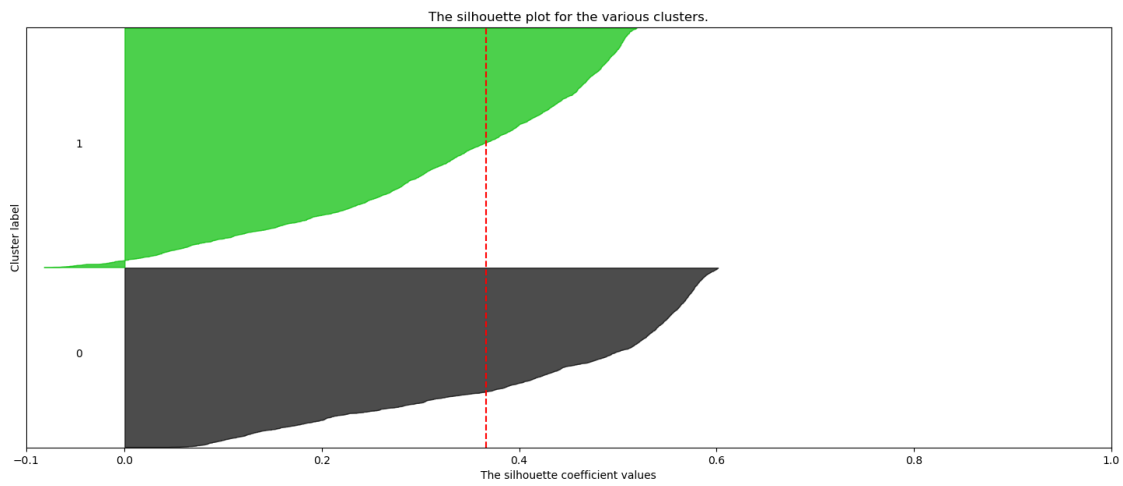
Silhouette Analysis

```
[24]: from clustering_function import plot_silhouette_analysis
```

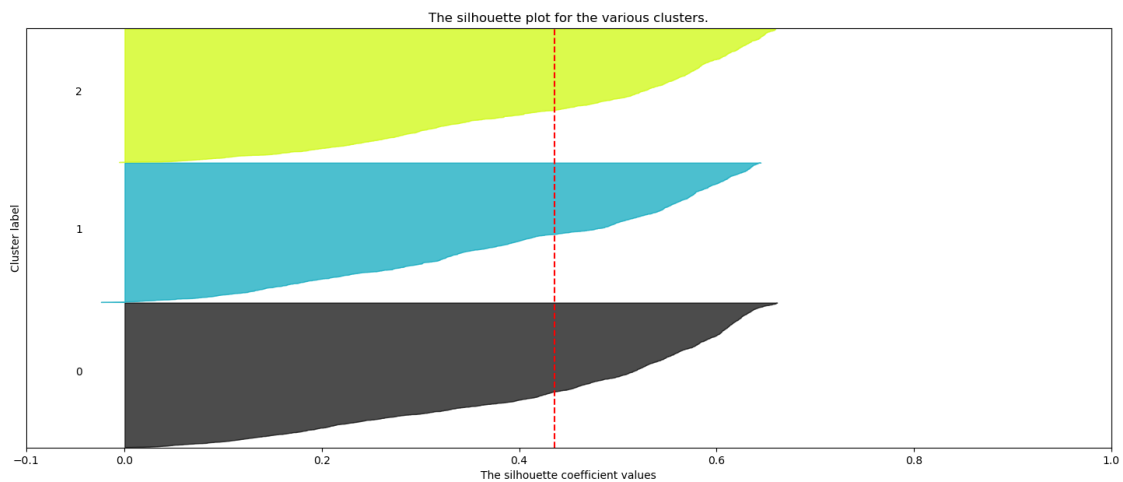
```
[25]: tsne_list_num_clusters = list(range(2,8))
      plot_silhouette_analysis(X_tsne_reduced, tsne_list_num_clusters)
```

For n_clusters = 2 The average silhouette_score is : 0.36614758
 For n_clusters = 3 The average silhouette_score is : 0.43591416
 For n_clusters = 4 The average silhouette_score is : 0.3885924
 For n_clusters = 5 The average silhouette_score is : 0.36350286
 For n_clusters = 6 The average silhouette_score is : 0.39561477
 For n_clusters = 7 The average silhouette_score is : 0.39939517

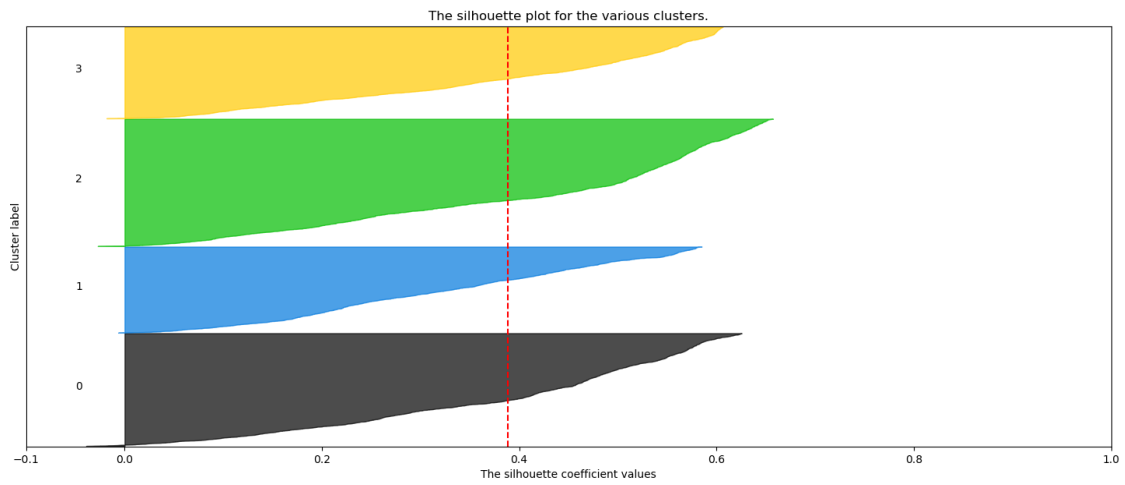
Silhouette analysis for KMeans clustering on sample data with n_clusters = 2



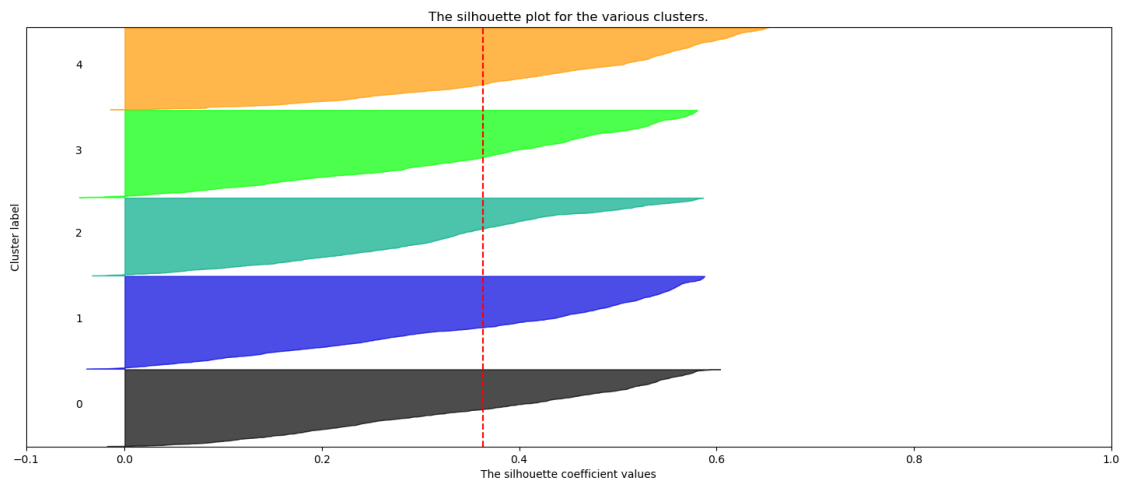
Silhouette analysis for KMeans clustering on sample data with n_clusters = 3

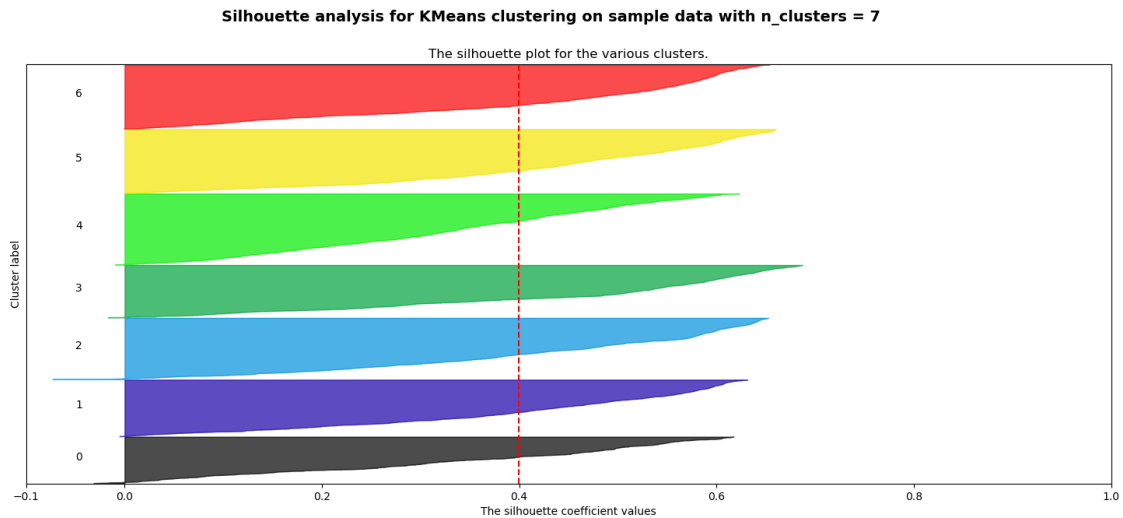
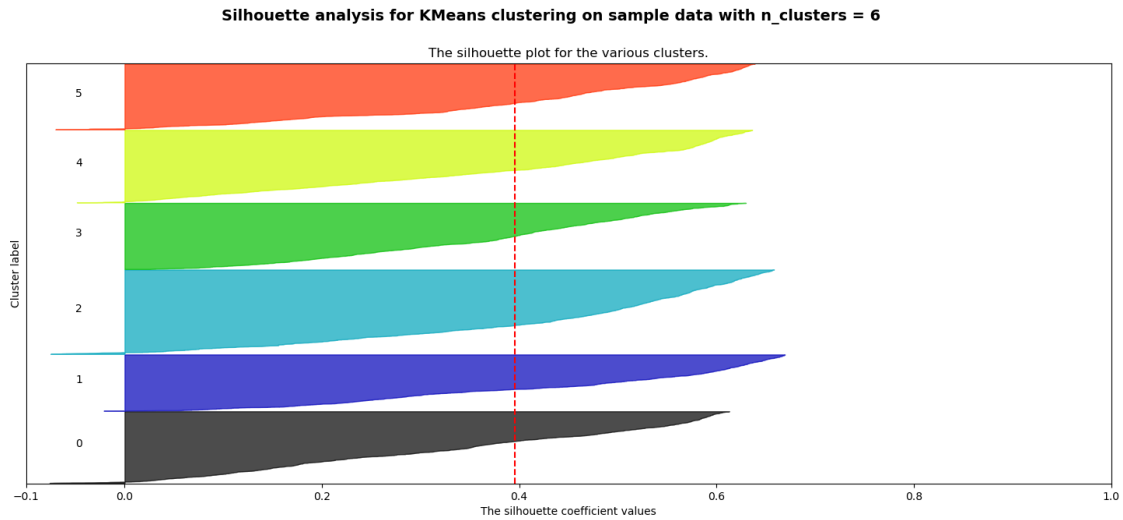


Silhouette analysis for KMeans clustering on sample data with $n_clusters = 4$



Silhouette analysis for KMeans clustering on sample data with $n_clusters = 5$



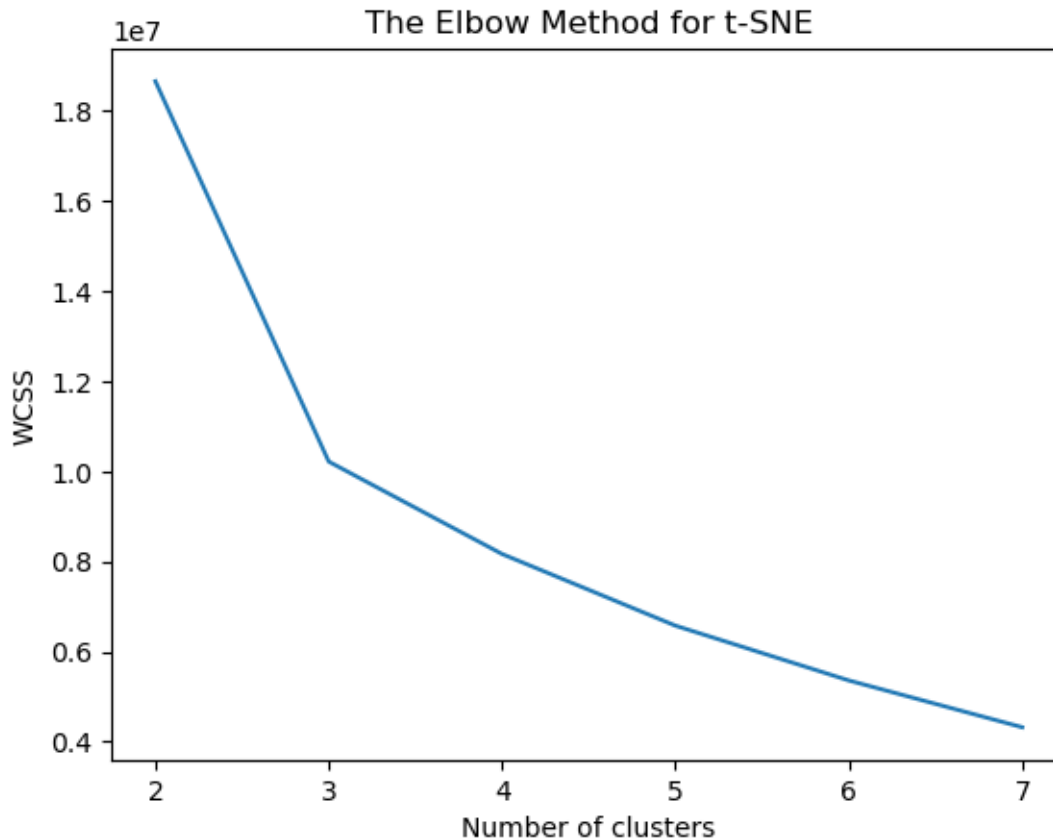


3 clusters seems to be the best number of cluster according to Silhouette Analysis.

Elbow Method

```
[26]: from clustering_function import elbow_method_analysis
```

```
[27]: elbow_method_analysis(X_tsne_reduced)
```



Given that the elbow point falls at 3, the elbow method also suggests that 3 clusters will be the ideal number of clusters for K-Means.

0.3 Clustering

```
[28]: # Trackers throughout each model
scores = {} # to track the silhouette score of the tuned model
food_groups = {} # to track the counts of each group
```

```
[29]: X_with_labels = X.copy()
X_with_labels
```

```
[29]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
0	0.685223	0.421947	0.644650	0.122492	
1	0.770972	0.539328	0.673774	0.444529	
2	0.624253	0.633388	0.203973	0.020652	
3	0.466968	0.380891	0.074901	0.056849	
4	0.513760	0.468498	0.074901	0.056849	
...	
8785	0.654500	0.583869	0.396246	0.175712	

8786	0.654500	0.583869	0.396246	0.175712
8787	0.642026	0.626764	0.305234	0.000000
8788	0.817624	0.893283	0.324179	0.024556
8789	0.889955	0.936086	0.535802	0.513582

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
0	0.022900		0.022872
1	0.297219		0.227549
2	0.242826		0.122122
3	0.017754		0.019922
4	0.017754		0.019922
...
8785	0.066435		0.008176
8786	0.066435		0.008176
8787	0.000000		0.000000
8788	0.002465		0.005934
8789	0.373303		0.259528

	Saturated Fatty Acids(g)
0	0.030388
1	0.212894
2	0.003042
3	0.014394
4	0.014394
...	...
8785	0.129330
8786	0.129330
8787	0.000000
8788	0.009211
8789	0.275650

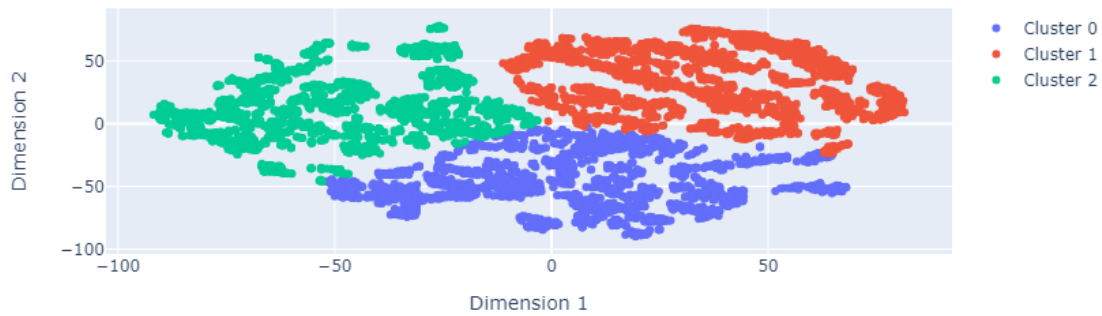
[8790 rows x 7 columns]

0.4 K Means Clustering

```
[30]: from clustering_function import perform_kmeans_clustering, plot_clusters
```

```
[31]: tsne_clust_labels, kmean_model = perform_kmeans_clustering(X_tsne_reduced, 3)
plot_clusters(X_tsne_reduced, tsne_clust_labels, 'K-Means')
```

t-SNE with K-Means



```
[32]: # export model
with open('../Model_fitted/Kmeans_model_median.pkl', 'wb') as files:
    pickle.dump(kmean_model, files)
```

```
[33]: X_with_labels['kmeans_tsne'] = tsne_clust_labels
scores['kmeans_tsne'] = (silhouette_score(X_tsne_reduced, X_with_labels['kmeans_tsne']))

X_with_labels
```

```
[33]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
0	0.685223	0.421947	0.644650	0.122492	
1	0.770972	0.539328	0.673774	0.444529	
2	0.624253	0.633388	0.203973	0.020652	
3	0.466968	0.380891	0.074901	0.056849	
4	0.513760	0.468498	0.074901	0.056849	
...	
8785	0.654500	0.583869	0.396246	0.175712	
8786	0.654500	0.583869	0.396246	0.175712	
8787	0.642026	0.626764	0.305234	0.000000	
8788	0.817624	0.893283	0.324179	0.024556	
8789	0.889955	0.936086	0.535802	0.513582	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
0	0.022900	0.022872	
1	0.297219	0.227549	
2	0.242826	0.122122	
3	0.017754	0.019922	
4	0.017754	0.019922	
...	

8785	0.066435	0.008176
8786	0.066435	0.008176
8787	0.000000	0.000000
8788	0.002465	0.005934
8789	0.373303	0.259528

	Saturated Fatty Acids(g)	kmeans_tsne
0	0.030388	1
1	0.212894	1
2	0.003042	2
3	0.014394	2
4	0.014394	2
...
8785	0.129330	2
8786	0.129330	2
8787	0.000000	2
8788	0.009211	0
8789	0.275650	0

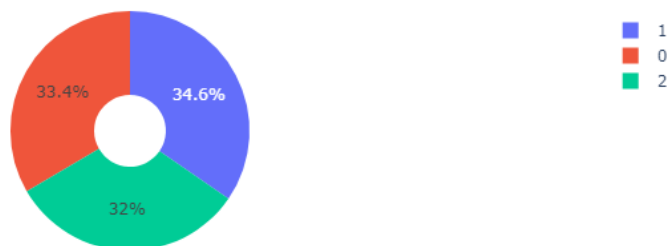
[8790 rows x 8 columns]

```
[34]: from clustering_function import get_food_groups, plot_cluster_distribution
```

```
[35]: food_groups, value_counts = get_food_groups(X_with_labels['kmeans_tsne'],
        ↪ 'KMeans_tsne', food_groups)
```

```
[36]: plot_cluster_distribution(value_counts, 'K-Means')
```

K-Means using t-SNE classes distribution



```
[37]: kmeans_tsne = X_with_labels.groupby('kmeans_tsne')
```

```
[38]: n = kmeans_tsne['kmeans_tsne'].count().count() # number of cluster
      for i in range(0,n):
          display(kmeans_tsne.get_group(i))
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
5	0.913731	0.870851	0.476131	0.745258	
7	0.916054	0.866962	0.491577	0.753711	
8	0.875881	0.808581	0.437893	0.696246	
12	0.904365	0.529205	0.512571	0.822403	
13	0.862406	0.577933	0.329815	0.756303	
...	
8725	0.859472	0.591069	0.577721	0.718901	
8745	0.859894	0.897421	0.719463	0.181414	
8748	0.858196	0.937465	0.463454	0.219193	
8788	0.817624	0.893283	0.324179	0.024556	
8789	0.889955	0.936086	0.535802	0.513582	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
5	0.676104	0.443583	
7	0.684742	0.451553	
8	0.626129	0.398092	
12	0.632893	0.671998	
13	0.615989	0.517992	
...	
8725	0.444714	0.657667	
8745	0.024320	0.103539	
8748	0.080576	0.140080	
8788	0.002465	0.005934	
8789	0.373303	0.259528	

	Saturated Fatty Acids(g)	kmeans_tsne
5	0.348734	0
7	0.355775	0
8	0.308817	0
12	0.494853	0
13	0.473315	0
...
8725	0.360711	0
8745	0.063870	0
8748	0.000000	0
8788	0.009211	0
8789	0.275650	0

[2937 rows x 8 columns]

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
0	0.685223	0.421947	0.644650	0.122492	
1	0.770972	0.539328	0.673774	0.444529	

14	0.816493	0.139076	0.695060	0.644735
100	0.786376	0.000000	0.756312	0.513785
101	0.717455	0.000000	0.681411	0.382380
...
8749	0.769417	0.000000	0.762047	0.442852
8750	0.733270	0.000000	0.708751	0.396735
8771	0.632419	0.346116	0.532774	0.232190
8772	0.601603	0.330664	0.537600	0.071353
8773	0.673693	0.347863	0.512571	0.388237

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
0	0.022900	0.022872	
1	0.297219	0.227549	
14	0.511857	0.352783	
100	0.351960	0.293664	
101	0.175770	0.224155	
...	
8749	0.242826	0.122122	
8750	0.246740	0.204303	
8771	0.089229	0.016934	
8772	0.011634	0.002758	
8773	0.257480	0.088905	

	Saturated Fatty Acids(g)	kmeans_tsne
0	0.030388	1
1	0.212894	1
14	0.395637	1
100	0.254692	1
101	0.180514	1
...
8749	0.208383	1
8750	0.180322	1
8771	0.175470	1
8772	0.024208	1
8773	0.267429	1

[3041 rows x 8 columns]

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
2	0.624253	0.633388	0.203973	0.020652	
3	0.466968	0.380891	0.074901	0.056849	
4	0.513760	0.468498	0.074901	0.056849	
6	0.670663	0.503637	0.457547	0.324453	
9	0.721842	0.757620	0.153183	0.055176	
...	
8783	0.556031	0.463708	0.351950	0.035864	
8784	0.656199	0.583869	0.396246	0.175712	
8785	0.654500	0.583869	0.396246	0.175712	

8786	0.654500	0.583869	0.396246	0.175712
8787	0.642026	0.626764	0.305234	0.000000

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
2	0.242826	0.122122	
3	0.017754	0.019922	
4	0.017754	0.019922	
6	0.222082	0.060118	
9	0.242826	0.122122	
...	
8783	0.010777	0.001153	
8784	0.061346	0.007953	
8785	0.066435	0.008176	
8786	0.066435	0.008176	
8787	0.000000	0.000000	

	Saturated Fatty Acids(g)	kmeans_tsne
2	0.003042	2
3	0.014394	2
4	0.014394	2
6	0.180322	2
9	0.208383	2
...
8783	0.024012	2
8784	0.122689	2
8785	0.129330	2
8786	0.129330	2
8787	0.000000	2

[2812 rows x 8 columns]

0.4.1 Insights

```
[39]: kmeans_tsne.get_group(0).describe()
```

```
[39]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
count	2937.000000	2937.000000	2937.000000	2937.000000	
mean	0.863727	0.805301	0.460011	0.526370	
std	0.053829	0.193784	0.165270	0.210419	
min	0.656199	0.000000	0.000000	0.000000	
25%	0.829515	0.733767	0.383500	0.377123	
50%	0.869298	0.876045	0.478997	0.544657	
75%	0.901848	0.929653	0.560456	0.674634	
max	1.000000	0.986273	0.964519	1.000000	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
count	2937.000000	2937.000000	

mean	0.329947	0.299584
std	0.199535	0.199596
min	0.000000	0.000000
25%	0.192959	0.138181
50%	0.317862	0.255735
75%	0.448099	0.421787
max	0.984335	1.000000

	Saturated Fatty Acids(g)	kmeans_tsne
count	2937.000000	2937.0
mean	0.302195	0.0
std	0.180595	0.0
min	0.000000	0.0
25%	0.142837	0.0
50%	0.307050	0.0
75%	0.430294	0.0
max	0.903884	0.0

Cluster 0: It appears that this cluster contains foods heavy in lipids and energy. The energy value minimum of 0.656 is the greatest minimum value observed across all clusters.

```
[40]: kmeans_tsne.get_group(1).describe()
```

```
[40]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
count	3041.000000	3041.000000	3041.000000	3041.000000	
mean	0.773320	0.078428	0.679336	0.500426	
std	0.068658	0.149217	0.116332	0.179250	
min	0.509238	0.000000	0.000000	0.000000	
25%	0.729216	0.000000	0.660090	0.380891	
50%	0.770972	0.000000	0.698659	0.496306	
75%	0.810708	0.079011	0.739935	0.618189	
max	1.000000	0.751628	1.000000	1.000000	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
count	3041.000000	3041.000000	
mean	0.344084	0.163226	
std	0.166106	0.114097	
min	0.000000	0.000000	
25%	0.233872	0.079265	
50%	0.335200	0.124698	
75%	0.454211	0.220889	
max	1.000000	0.649514	

	Saturated Fatty Acids(g)	kmeans_tsne
count	3041.000000	3041.0
mean	0.316879	1.0
std	0.169445	0.0
min	0.000000	1.0

25%	0.200476	1.0
50%	0.294946	1.0
75%	0.419823	1.0
max	1.000000	1.0

Cluster 1: This cluster predominantly consists of low-carb foods, with approximately half of the data falling into this category. The statistical analysis reveals that this cluster exhibits the highest Q1 range for protein, reaching 0.66. Consequently, it can be inferred that Cluster 1 is primarily composed of protein-rich foods.

```
[41]: kmeans_tsne.get_group(2).describe()
```

```
[41]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
count	2812.000000	2812.000000	2812.000000	2812.000000	
mean	0.588109	0.521628	0.209455	0.108504	
std	0.137224	0.183761	0.138160	0.114048	
min	0.000000	0.000000	0.000000	0.000000	
25%	0.526545	0.416072	0.097559	0.020652	
50%	0.604032	0.527693	0.209686	0.063415	
75%	0.669125	0.633970	0.306923	0.168863	
max	0.881455	1.000000	0.715473	0.606244	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
count	2812.000000	2812.000000	
mean	0.076733	0.054543	
std	0.097881	0.057750	
min	0.000000	0.000000	
25%	0.002687	0.009289	
50%	0.014187	0.030898	
75%	0.143674	0.112947	
max	0.565610	0.363535	

	Saturated Fatty Acids(g)	kmeans_tsne
count	2812.000000	2812.0
mean	0.062648	2.0
std	0.082691	0.0
min	0.000000	2.0
25%	0.003903	2.0
50%	0.016230	2.0
75%	0.104264	2.0
max	0.373856	2.0

Cluster 2: The maximum values for the fatty acids in this cluster were the lowest among all the cluster. Hence can conclude that this cluster is food that low in fatty acids. This suggests that this cluster may consists of healthy food.

0.5 Agglomerative Clustering

```
[42]: from clustering_function import tuning_agglomerative, perform_agg_clustering
```

```
[43]: agg_param_grid = {  
    'n_clusters': [2, 3, 4, 5, 6],  
    'linkage': ['ward'],  
    'affinity': ['euclidean']  
}
```

```
[44]: tuning_agglomerative(X_tsne_reduced, agg_param_grid)
```

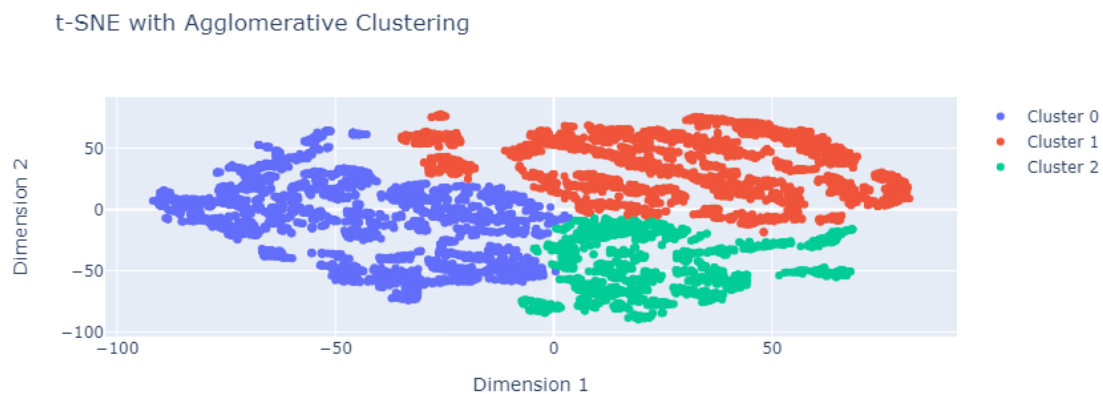
Best silhouette score: 0.38080552

Best parameters: {'affinity': 'euclidean', 'linkage': 'ward', 'n_clusters': 3}

```
[45]: Agg_tsne_clust_labels,agg_model = perform_agg_clustering(X_tsne_reduced, 3)
```

```
[46]: # export model  
with open('../Model_fitted/Agg_model_median.pkl', 'wb') as files:  
    pickle.dump(agg_model, files)
```

```
[47]: plot_clusters(X_tsne_reduced, Agg_tsne_clust_labels, "Agglomerative Clustering")
```

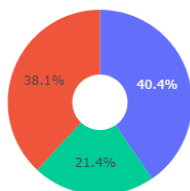


```
[48]: X_with_labels['Agg_tsne'] = Agg_tsne_clust_labels  
scores['Agg_tsne'] = (silhouette_score(X_tsne_reduced, X_with_labels['Agg_tsne']))
```

```
[49]: food_groups, value_counts = get_food_groups(X_with_labels['Agg_tsne'],  
    'Agglomerative_tsne', food_groups)
```

```
[50]: plot_cluster_distribution(value_counts, 'Agglomerative Clustering')
```

Agglomerative Clustering using t-SNE classes distribution



0
1
2

```
[51]: Aggtsne = X_with_labels.groupby('Agg_tsne')
```

```
[52]: n = Aggtsne['Agg_tsne'].count().count() # number of cluster
for i in range(0,n):
    display(Aggtsne.get_group(i))
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
3	0.466968	0.380891	0.074901	0.056849	
4	0.513760	0.468498	0.074901	0.056849	
6	0.670663	0.503637	0.457547	0.324453	
10	0.857339	0.957418	0.221928	0.113698	
15	0.836114	0.000000	0.000000	0.000000	
...	
8784	0.656199	0.583869	0.396246	0.175712	
8785	0.654500	0.583869	0.396246	0.175712	
8786	0.654500	0.583869	0.396246	0.175712	
8787	0.642026	0.626764	0.305234	0.000000	
8788	0.817624	0.893283	0.324179	0.024556	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
3	0.017754	0.019922	
4	0.017754	0.019922	
6	0.222082	0.060118	
10	0.242826	0.122122	
15	0.000000	0.000000	
...	
8784	0.061346	0.007953	
8785	0.066435	0.008176	
8786	0.066435	0.008176	
8787	0.000000	0.000000	
8788	0.002465	0.005934	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne
3	0.014394	2	0

4	0.014394	2	0
6	0.180322	2	0
10	0.208383	2	0
15	0.000000	2	0
...
8784	0.122689	2	0
8785	0.129330	2	0
8786	0.129330	2	0
8787	0.000000	2	0
8788	0.009211	0	0

[3555 rows x 9 columns]

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
0	0.685223	0.421947	0.644650	0.122492	
1	0.770972	0.539328	0.673774	0.444529	
2	0.624253	0.633388	0.203973	0.020652	
9	0.721842	0.757620	0.153183	0.055176	
11	0.622139	0.616810	0.093208	0.030283	
...	
8749	0.769417	0.000000	0.762047	0.442852	
8750	0.733270	0.000000	0.708751	0.396735	
8771	0.632419	0.346116	0.532774	0.232190	
8772	0.601603	0.330664	0.537600	0.071353	
8773	0.673693	0.347863	0.512571	0.388237	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
0	0.022900	0.022872	
1	0.297219	0.227549	
2	0.242826	0.122122	
9	0.242826	0.122122	
11	0.242826	0.122122	
...	
8749	0.242826	0.122122	
8750	0.246740	0.204303	
8771	0.089229	0.016934	
8772	0.011634	0.002758	
8773	0.257480	0.088905	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne
0	0.030388	1	1
1	0.212894	1	1
2	0.003042	2	1
9	0.208383	2	1
11	0.208383	2	1
...
8749	0.208383	1	1
8750	0.180322	1	1

8771	0.175470	1	1
8772	0.024208	1	1
8773	0.267429	1	1

[3350 rows x 9 columns]

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
5	0.913731	0.870851	0.476131	0.745258	
7	0.916054	0.866962	0.491577	0.753711	
8	0.875881	0.808581	0.437893	0.696246	
12	0.904365	0.529205	0.512571	0.822403	
13	0.862406	0.577933	0.329815	0.756303	
...	
8629	0.807727	0.732959	0.581346	0.493436	
8659	0.882545	0.914725	0.526799	0.531641	
8696	0.883985	0.813507	0.761901	0.617939	
8725	0.859472	0.591069	0.577721	0.718901	
8789	0.889955	0.936086	0.535802	0.513582	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
5	0.676104	0.443583	
7	0.684742	0.451553	
8	0.626129	0.398092	
12	0.632893	0.671998	
13	0.615989	0.517992	
...	
8629	0.340380	0.173271	
8659	0.385713	0.213847	
8696	0.438754	0.386834	
8725	0.444714	0.657667	
8789	0.373303	0.259528	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne
5	0.348734	0	2
7	0.355775	0	2
8	0.308817	0	2
12	0.494853	0	2
13	0.473315	0	2
...
8629	0.327419	0	2
8659	0.356463	0	2
8696	0.261418	0	2
8725	0.360711	0	2
8789	0.275650	0	2

[1885 rows x 9 columns]

0.5.1 Insights

```
[53]: Aggtsne.get_group(0).describe()
```

```
[53]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
count	3555.000000	3555.000000	3555.000000	3555.000000	
mean	0.666548	0.633652	0.295738	0.174720	
std	0.167766	0.234432	0.188526	0.153656	
min	0.000000	0.000000	0.000000	0.000000	
25%	0.568816	0.466745	0.142881	0.039505	
50%	0.669125	0.620550	0.286992	0.134466	
75%	0.823148	0.857315	0.447489	0.304672	
max	0.947569	1.000000	0.964519	0.924243	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
count	3555.000000	3555.000000	
mean	0.082623	0.076877	
std	0.097304	0.080969	
min	0.000000	0.000000	
25%	0.003798	0.011719	
50%	0.035946	0.043492	
75%	0.145836	0.122122	
max	0.565610	0.487236	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne
count	3555.000000	3555.000000	3555.0
mean	0.079875	1.381997	0.0
std	0.100300	0.924141	0.0
min	0.000000	0.000000	0.0
25%	0.006255	0.000000	0.0
50%	0.034351	2.000000	0.0
75%	0.129330	2.000000	0.0
max	0.756575	2.000000	0.0

Cluster 0: Among all the food clusters, this one has the lowest fat content, with lipids and three fatty acids each around value of 0.3 (excluding maximum value). This implies that there may be healthy foods in this food cluster.

```
[54]: Aggtsne.get_group(1).describe()
```

```
[54]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
count	3350.000000	3350.000000	3350.000000	3350.000000	
mean	0.749137	0.123298	0.635710	0.448678	
std	0.087602	0.193661	0.179329	0.205318	
min	0.376881	0.000000	0.000000	0.000000	
25%	0.709449	0.000000	0.635389	0.331135	
50%	0.759726	0.000000	0.692396	0.467998	
75%	0.802205	0.225788	0.736299	0.594092	

max	0.999674	0.795036	1.000000	1.000000
-----	----------	----------	----------	----------

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
count	3350.000000	3350.000000	
mean	0.328621	0.156111	
std	0.152704	0.103982	
min	0.000000	0.000000	
25%	0.242826	0.084296	
50%	0.304128	0.122122	
75%	0.436242	0.204303	
max	0.923391	0.629091	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne
count	3350.000000	3350.000000	3350.0
mean	0.288747	1.104478	1.0
std	0.158963	0.311726	0.0
min	0.000000	0.000000	1.0
25%	0.187612	1.000000	1.0
50%	0.267911	1.000000	1.0
75%	0.390189	1.000000	1.0
max	0.786175	2.000000	1.0

Cluster 1: This cluster seems to be a low-carb meal cluster, as Q1 and Q2 have carbohydrate values of 0. The greatest value of carbohydrates was about 0.75g (after log transform and min max scaled), which is the lowest among all clusters.

```
[55]: Aggtsne.get_group(2).describe()
```

```
[55]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
count	1885.000000	1885.000000	1885.000000	1885.000000	
mean	0.882231	0.745255	0.437626	0.662416	
std	0.054809	0.247276	0.183585	0.141062	
min	0.712934	0.000000	0.000000	0.348732	
25%	0.840996	0.681925	0.367857	0.554266	
50%	0.891665	0.842390	0.462898	0.641729	
75%	0.914315	0.908793	0.566303	0.729697	
max	1.000000	0.979258	0.898049	1.000000	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
count	1885.000000	1885.000000	
mean	0.443810	0.389048	
std	0.170806	0.194710	
min	0.000000	0.000000	
25%	0.328592	0.246133	
50%	0.418181	0.367389	
75%	0.528850	0.506677	
max	1.000000	1.000000	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne
count	1885.000000	1885.000000	1885.0
mean	0.411717	0.027586	2.0
std	0.154498	0.163827	0.0
min	0.000000	0.000000	2.0
25%	0.307695	0.000000	2.0
50%	0.391984	0.000000	2.0
75%	0.497368	0.000000	2.0
max	1.000000	1.000000	2.0

Cluster 2: This cluster may consist of foods high in energy because the first quartile of data was around 0.84, the highest of all the clusters. Based on the three fatty acid compositions, this cluster indicates that the meal in this cluster is maximal in fat. The fact that this food cluster has the highest value of total lipids among all clusters serves as additional support.

0.6 GMM

```
[56]: from clustering_function import gmm_bic_score, perform_gmm_clustering
      from sklearn.mixture import GaussianMixture
      from sklearn.model_selection import GridSearchCV
```

```
[57]: gmm_param_grid = {
      "n_components": range(1, 6),
      "covariance_type": ["spherical", "tied", "diag", "full"],
      }
      grid_search = GridSearchCV(
          GaussianMixture(), param_grid=gmm_param_grid, scoring=gmm_bic_score
      )
```

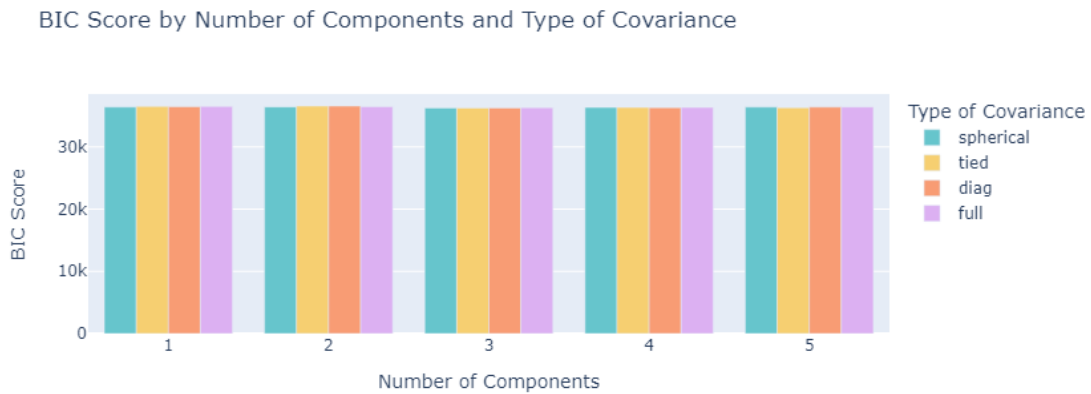
```
[58]: grid_search.fit(X_tsne_reduced)
      Gmm_tsne_results = grid_search.cv_results_
```

```
[59]: Gmm_tsne_df = pd.DataFrame(Gmm_tsne_results)[
      ["param_n_components", "param_covariance_type", "mean_test_score"]
      ]
      Gmm_tsne_df["mean_test_score"] = -Gmm_tsne_df["mean_test_score"]
      Gmm_tsne_df = Gmm_tsne_df.rename(
          columns={
              "param_n_components": "Number of Components",
              "param_covariance_type": "Type of Covariance",
              "mean_test_score": "BIC Score",
          }
      )
      Gmm_tsne_df.sort_values(by="BIC Score").head()
```

```
[59]:   Number of Components  Type of Covariance  BIC Score
      7                  3                  tied  36259.642277
      2                  3                  spherical  36277.982784
```

12	3	diag	36285.380930
9	5	tied	36307.227321
17	3	full	36333.090942

```
[60]: fig = px.bar(Gmm_tsne_df, x="Number of Components", y="BIC Score",
                  color="Type of Covariance", barmode="group",
                  title="BIC Score by Number of Components and Type of
                  ↪Covariance",
                  color_discrete_sequence=px.colors.qualitative.Pastel)
fig.show()
```

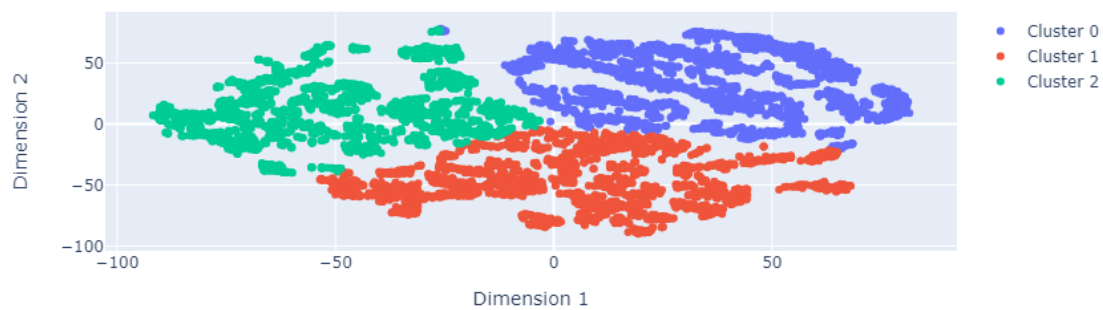


```
[61]: Gmm_tsne_labels, gmm_model = perform_gmm_clustering(X_tsne_reduced, 3, 'tied')
```

```
[62]: # export model
with open('../Model_fitted/Gmm_model_median.pkl', 'wb') as files:
    pickle.dump(gmm_model, files)
```

```
[63]: plot_clusters(X_tsne_reduced, Gmm_tsne_labels, "GMM")
```

t-SNE with GMM

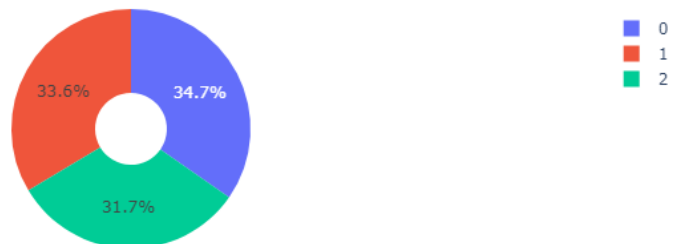


```
[64]: X_with_labels['Gmm_tsne'] = Gmm_tsne_labels
      scores['Gmm_tsne'] = (silhouette_score(X_tsne_reduced,
      ↪ X_with_labels['Gmm_tsne']))

[65]: food_groups, value_counts = get_food_groups(X_with_labels['Gmm_tsne'],
      ↪ 'GMM_tsne', food_groups)

[66]: plot_cluster_distribution(value_counts, 'GMM')
```

GMM using t-SNE classes distribution



```
[67]: GmmTsne = X_with_labels.groupby('Gmm_tsne')

[68]: n = GmmTsne['Gmm_tsne'].count().count() # number of cluster
      for i in range(0,n):
          display(GmmTsne.get_group(i))
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
0	0.685223	0.421947	0.644650	0.122492	
1	0.770972	0.539328	0.673774	0.444529	
14	0.816493	0.139076	0.695060	0.644735	
52	0.654500	0.288696	0.015061	0.000000	
53	0.656199	0.334863	0.015061	0.000000	
...	
8749	0.769417	0.000000	0.762047	0.442852	
8750	0.733270	0.000000	0.708751	0.396735	
8771	0.632419	0.346116	0.532774	0.232190	
8772	0.601603	0.330664	0.537600	0.071353	
8773	0.673693	0.347863	0.512571	0.388237	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
0	0.022900	0.022872	
1	0.297219	0.227549	
14	0.511857	0.352783	
52	0.242826	0.122122	
53	0.242826	0.122122	
...	
8749	0.242826	0.122122	
8750	0.246740	0.204303	
8771	0.089229	0.016934	
8772	0.011634	0.002758	
8773	0.257480	0.088905	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne	Gmm_tsne
0	0.030388	1	1	0
1	0.212894	1	1	0
14	0.395637	1	1	0
52	0.208383	2	1	0
53	0.208383	2	1	0
...
8749	0.208383	1	1	0
8750	0.180322	1	1	0
8771	0.175470	1	1	0
8772	0.024208	1	1	0
8773	0.267429	1	1	0

[3048 rows x 10 columns]

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
5	0.913731	0.870851	0.476131	0.745258	
7	0.916054	0.866962	0.491577	0.753711	
8	0.875881	0.808581	0.437893	0.696246	
12	0.904365	0.529205	0.512571	0.822403	
13	0.862406	0.577933	0.329815	0.756303	
...	

8725	0.859472	0.591069	0.577721	0.718901
8745	0.859894	0.897421	0.719463	0.181414
8748	0.858196	0.937465	0.463454	0.219193
8788	0.817624	0.893283	0.324179	0.024556
8789	0.889955	0.936086	0.535802	0.513582

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
5	0.676104	0.443583	
7	0.684742	0.451553	
8	0.626129	0.398092	
12	0.632893	0.671998	
13	0.615989	0.517992	
...	
8725	0.444714	0.657667	
8745	0.024320	0.103539	
8748	0.080576	0.140080	
8788	0.002465	0.005934	
8789	0.373303	0.259528	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne	Gmm_tsne
5	0.348734	0	2	1
7	0.355775	0	2	1
8	0.308817	0	2	1
12	0.494853	0	2	1
13	0.473315	0	2	1
...
8725	0.360711	0	2	1
8745	0.063870	0	0	1
8748	0.000000	0	0	1
8788	0.009211	0	0	1
8789	0.275650	0	2	1

[2954 rows x 10 columns]

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
2	0.624253	0.633388	0.203973	0.020652	
3	0.466968	0.380891	0.074901	0.056849	
4	0.513760	0.468498	0.074901	0.056849	
6	0.670663	0.503637	0.457547	0.324453	
9	0.721842	0.757620	0.153183	0.055176	
...	
8783	0.556031	0.463708	0.351950	0.035864	
8784	0.656199	0.583869	0.396246	0.175712	
8785	0.654500	0.583869	0.396246	0.175712	
8786	0.654500	0.583869	0.396246	0.175712	
8787	0.642026	0.626764	0.305234	0.000000	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
--	--------------------------------	--------------------------------	---

2	0.242826	0.122122
3	0.017754	0.019922
4	0.017754	0.019922
6	0.222082	0.060118
9	0.242826	0.122122
...
8783	0.010777	0.001153
8784	0.061346	0.007953
8785	0.066435	0.008176
8786	0.066435	0.008176
8787	0.000000	0.000000

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne	Gmm_tsne
2	0.003042	2	1	2
3	0.014394	2	0	2
4	0.014394	2	0	2
6	0.180322	2	0	2
9	0.208383	2	1	2
...
8783	0.024012	2	0	2
8784	0.122689	2	0	2
8785	0.129330	2	0	2
8786	0.129330	2	0	2
8787	0.000000	2	0	2

[2788 rows x 10 columns]

0.6.1 Insights

```
[69]: GmmTsne.get_group(0).describe()
```

```
[69]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
count	3048.000000	3048.000000	3048.000000	3048.000000	
mean	0.771972	0.080391	0.678424	0.496158	
std	0.067737	0.150926	0.118400	0.179918	
min	0.509238	0.000000	0.000000	0.000000	
25%	0.728185	0.000000	0.659228	0.378260	
50%	0.770197	0.000000	0.698562	0.494544	
75%	0.810117	0.088216	0.739855	0.615169	
max	1.000000	0.751628	1.000000	1.000000	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
count	3048.000000	3048.000000	
mean	0.341405	0.161836	
std	0.162391	0.111637	
min	0.000000	0.000000	
25%	0.234012	0.079388	

50%	0.332745	0.124023
75%	0.452313	0.219573
max	0.971727	0.629091

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne	Gmm_tsne
count	3048.000000	3048.000000	3048.000000	3048.0
mean	0.314871	1.003937	1.013123	0.0
std	0.167912	0.076759	0.116669	0.0
min	0.000000	0.000000	0.000000	0.0
25%	0.200410	1.000000	1.000000	0.0
50%	0.292086	1.000000	1.000000	0.0
75%	0.416816	1.000000	1.000000	0.0
max	1.000000	2.000000	2.000000	0.0

Cluster 0: With a maximum value of 1 and the highest reported protein Q1 of 0.659, this cluster appears to have categorized foods based on protein.

```
[70]: GmmTsne.get_group(1).describe()
```

```
[70]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
count	2954.000000	2954.000000	2954.000000	2954.000000	
mean	0.864490	0.803388	0.457639	0.526275	
std	0.053960	0.199434	0.167634	0.214055	
min	0.656199	0.000000	0.000000	0.000000	
25%	0.830033	0.733840	0.382104	0.376456	
50%	0.869694	0.876045	0.478997	0.545270	
75%	0.902165	0.929949	0.560277	0.675738	
max	1.000000	0.986273	0.964519	1.000000	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
count	2954.000000	2954.000000	
mean	0.331457	0.299671	
std	0.202784	0.200190	
min	0.000000	0.000000	
25%	0.192767	0.137161	
50%	0.318110	0.255697	
75%	0.450811	0.422187	
max	1.000000	1.000000	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne	Gmm_tsne
count	2954.000000	2954.000000	2954.000000	2954.0
mean	0.302524	0.012525	1.249492	1.0
std	0.182657	0.145524	0.968016	0.0
min	0.000000	0.000000	0.000000	1.0
25%	0.140432	0.000000	0.000000	1.0
50%	0.307614	0.000000	2.000000	1.0
75%	0.431911	0.000000	2.000000	1.0
max	0.903884	2.000000	2.000000	1.0

Cluster 1: This is a high-energy cluster with the first quartile of the data lying at 0.83 (the highest among all clusters).

```
[71]: GmmTsne.get_group(2).describe()
```

```
[71]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
count	2788.000000	2788.000000	2788.000000	2788.000000	
mean	0.586628	0.520892	0.210258	0.109738	
std	0.136497	0.181437	0.137622	0.114633	
min	0.000000	0.000000	0.000000	0.000000	
25%	0.522406	0.418991	0.101826	0.020652	
50%	0.601603	0.529017	0.211846	0.065026	
75%	0.669125	0.633388	0.307622	0.170102	
max	0.881090	1.000000	0.715473	0.606244	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
count	2788.000000	2788.000000	
mean	0.075846	0.054205	
std	0.097338	0.057715	
min	0.000000	0.000000	
25%	0.002465	0.009067	
50%	0.013975	0.030392	
75%	0.139711	0.108327	
max	0.565610	0.363535	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne	Gmm_tsne
count	2788.000000	2788.000000	2788.000000	2788.0
mean	0.062396	1.997131	0.122310	2.0
std	0.082646	0.075715	0.327702	0.0
min	0.000000	0.000000	0.000000	2.0
25%	0.003903	2.000000	0.000000	2.0
50%	0.016230	2.000000	0.000000	2.0
75%	0.103413	2.000000	0.000000	2.0
max	0.373856	2.000000	1.000000	2.0

Cluster 2: Foods that fall under this category are thought to be lower in fat, or more healthful overall. This is due to the fact that 75% of the nutritional data for fats are less than 0.20 (0.17, 0.13, 0.11, 0.10 for lipid and 3 fatty acids, respectively).

0.7 K-Medoids

```
[72]: from clustering_function import tuning_kmedoids, perform_kmd_clustering
```

```
[73]: kmd_param_grid = {
    'n_clusters': [3, 4],
    'method': ['alternate', 'pam'],
    'init' : ['random', 'heuristic', 'k-medoids++', 'build']
}
```

```
[74]: tuning_kmedoids(X_tsne_reduced, kmd_param_grid)
```

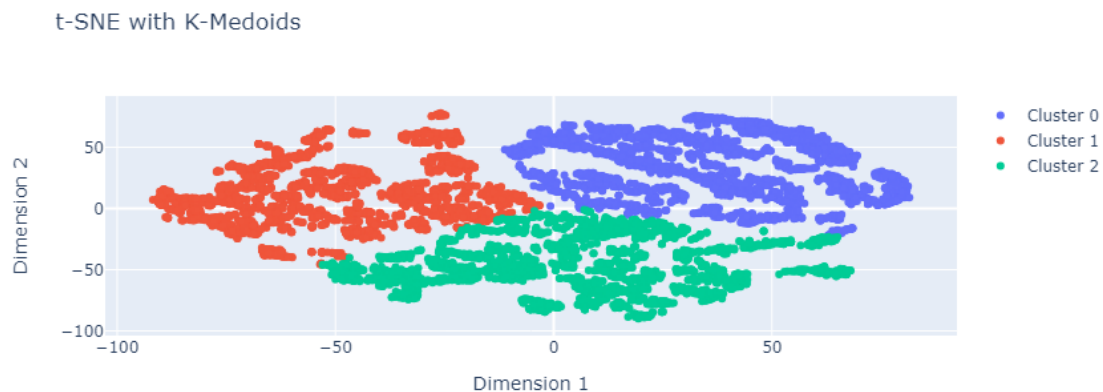
Best silhouette score: 0.4359154

Best parameters: {'init': 'random', 'method': 'alternate', 'n_clusters': 3}

```
[75]: Kmd_tsne_labels,kmd_model = perform_kmd_clustering(X_tsne_reduced, 3, 'random',↵
↪ 'alternate')
```

```
[76]: # export model
with open('../Model_fitted/Kmd_model_median.pkl', 'wb') as files:
    pickle.dump(kmd_model, files)
```

```
[77]: plot_clusters(X_tsne_reduced, Kmd_tsne_labels, "K-Medoids")
```

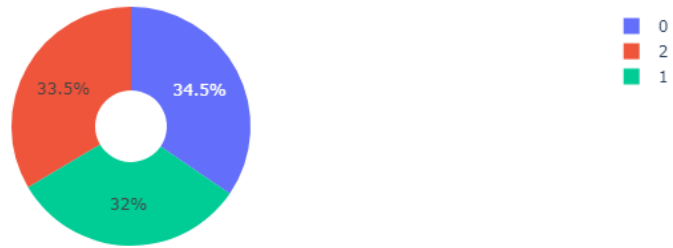


```
[78]: X_with_labels['Kmd_tsne'] = Kmd_tsne_labels
scores['Kmd_tsne'] = (silhouette_score(X_tsne_reduced,↵
↪ X_with_labels['Kmd_tsne']))
```

```
[79]: food_groups, value_counts = get_food_groups(X_with_labels['Kmd_tsne'],↵
↪ 'KMedoids_tsne', food_groups)
```

```
[80]: plot_cluster_distribution(value_counts, 'K-Medoids')
```

K-Medoids using t-SNE classes distribution



```
[81]: Kmdtsne = X_with_labels.groupby('Kmd_tsne')
```

```
[82]: n = Kmdtsne['Kmd_tsne'].count().count() # number of cluster
      for i in range(0,n):
          display(Kmdtsne.get_group(i))
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
0	0.685223	0.421947	0.644650	0.122492	
1	0.770972	0.539328	0.673774	0.444529	
14	0.816493	0.139076	0.695060	0.644735	
100	0.786376	0.000000	0.756312	0.513785	
101	0.717455	0.000000	0.681411	0.382380	
...	
8749	0.769417	0.000000	0.762047	0.442852	
8750	0.733270	0.000000	0.708751	0.396735	
8771	0.632419	0.346116	0.532774	0.232190	
8772	0.601603	0.330664	0.537600	0.071353	
8773	0.673693	0.347863	0.512571	0.388237	
	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\		
0	0.022900	0.022872			
1	0.297219	0.227549			
14	0.511857	0.352783			
100	0.351960	0.293664			
101	0.175770	0.224155			
...			
8749	0.242826	0.122122			
8750	0.246740	0.204303			
8771	0.089229	0.016934			
8772	0.011634	0.002758			
8773	0.257480	0.088905			

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne	Gmm_tsne	Kmd_tsne
0	0.030388	1	1	0	0
1	0.212894	1	1	0	0
14	0.395637	1	1	0	0
100	0.254692	1	1	0	0
101	0.180514	1	1	0	0
...
8749	0.208383	1	1	0	0
8750	0.180322	1	1	0	0
8771	0.175470	1	1	0	0
8772	0.024208	1	1	0	0
8773	0.267429	1	1	0	0

[3035 rows x 11 columns]

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
2	0.624253	0.633388	0.203973	0.020652	
3	0.466968	0.380891	0.074901	0.056849	
4	0.513760	0.468498	0.074901	0.056849	
6	0.670663	0.503637	0.457547	0.324453	
9	0.721842	0.757620	0.153183	0.055176	
...	
8783	0.556031	0.463708	0.351950	0.035864	
8784	0.656199	0.583869	0.396246	0.175712	
8785	0.654500	0.583869	0.396246	0.175712	
8786	0.654500	0.583869	0.396246	0.175712	
8787	0.642026	0.626764	0.305234	0.000000	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
2	0.242826	0.122122	
3	0.017754	0.019922	
4	0.017754	0.019922	
6	0.222082	0.060118	
9	0.242826	0.122122	
...	
8783	0.010777	0.001153	
8784	0.061346	0.007953	
8785	0.066435	0.008176	
8786	0.066435	0.008176	
8787	0.000000	0.000000	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne	Gmm_tsne	Kmd_tsne
2	0.003042	2	1	2	1
3	0.014394	2	0	2	1
4	0.014394	2	0	2	1
6	0.180322	2	0	2	1
9	0.208383	2	1	2	1

...	
8783	0.024012		2	0	2	1
8784	0.122689		2	0	2	1
8785	0.129330		2	0	2	1
8786	0.129330		2	0	2	1
8787	0.000000		2	0	2	1

[2809 rows x 11 columns]

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
5	0.913731	0.870851	0.476131	0.745258	
7	0.916054	0.866962	0.491577	0.753711	
8	0.875881	0.808581	0.437893	0.696246	
12	0.904365	0.529205	0.512571	0.822403	
13	0.862406	0.577933	0.329815	0.756303	
...	
8725	0.859472	0.591069	0.577721	0.718901	
8745	0.859894	0.897421	0.719463	0.181414	
8748	0.858196	0.937465	0.463454	0.219193	
8788	0.817624	0.893283	0.324179	0.024556	
8789	0.889955	0.936086	0.535802	0.513582	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
5	0.676104	0.443583	
7	0.684742	0.451553	
8	0.626129	0.398092	
12	0.632893	0.671998	
13	0.615989	0.517992	
...	
8725	0.444714	0.657667	
8745	0.024320	0.103539	
8748	0.080576	0.140080	
8788	0.002465	0.005934	
8789	0.373303	0.259528	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne	Gmm_tsne	Kmd_tsne
5	0.348734	0	2	1	2
7	0.355775	0	2	1	2
8	0.308817	0	2	1	2
12	0.494853	0	2	1	2
13	0.473315	0	2	1	2
...	
8725	0.360711	0	2	1	2
8745	0.063870	0	0	1	2
8748	0.000000	0	0	1	2
8788	0.009211	0	0	1	2
8789	0.275650	0	2	1	2

[2946 rows x 11 columns]

0.7.1 Insights

```
[83]: Kmdtsne.get_group(0).describe()
```

```
[83]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
count	3035.000000	3035.000000	3035.000000	3035.000000	
mean	0.773084	0.077875	0.680102	0.499991	
std	0.068356	0.147989	0.114427	0.178727	
min	0.509238	0.000000	0.000000	0.000000	
25%	0.729216	0.000000	0.660434	0.380891	
50%	0.770972	0.000000	0.698948	0.496306	
75%	0.810708	0.079011	0.739975	0.617939	
max	1.000000	0.693086	1.000000	1.000000	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
count	3035.000000	3035.000000	
mean	0.343654	0.162658	
std	0.165132	0.113367	
min	0.000000	0.000000	
25%	0.233992	0.079183	
50%	0.335200	0.124428	
75%	0.453866	0.220577	
max	0.971727	0.629151	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne	Gmm_tsne	\
count	3035.000000	3035.0	3035.000000	3035.000000	
mean	0.316777	1.0	1.015815	0.002636	
std	0.169271	0.0	0.127396	0.051282	
min	0.000000	1.0	0.000000	0.000000	
25%	0.200476	1.0	1.000000	0.000000	
50%	0.294946	1.0	1.000000	0.000000	
75%	0.419582	1.0	1.000000	0.000000	
max	1.000000	1.0	2.000000	1.000000	

	Kmd_tsne
count	3035.0
mean	0.0
std	0.0
min	0.0
25%	0.0
50%	0.0
75%	0.0
max	0.0

Cluster 0: This food group is low in carbs and high in protein, as seen by the values of the two nutrients, which are 0.66 and 0.74 for protein and 0 to 0.08 for carbohydrates, respectively.

```
[84]: Kmdtsne.get_group(1).describe()
```

```
[84]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
count	2809.000000	2809.000000	2809.000000	2809.000000	
mean	0.587818	0.521159	0.209288	0.108545	
std	0.137008	0.183297	0.138021	0.114077	
min	0.000000	0.000000	0.000000	0.000000	
25%	0.526545	0.415357	0.097559	0.020652	
50%	0.604032	0.527693	0.209252	0.063415	
75%	0.669125	0.633504	0.306923	0.168863	
max	0.881455	1.000000	0.715473	0.606244	

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
count	2809.000000	2809.000000	
mean	0.076722	0.054554	
std	0.097866	0.057752	
min	0.000000	0.000000	
25%	0.002687	0.009289	
50%	0.014187	0.030898	
75%	0.143644	0.112947	
max	0.565610	0.363535	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne	Gmm_tsne	\
count	2809.000000	2809.0	2809.000000	2809.000000	
mean	0.062696	2.0	0.126735	1.985760	
std	0.082720	0.0	0.332736	0.157245	
min	0.000000	2.0	0.000000	0.000000	
25%	0.003903	2.0	0.000000	2.000000	
50%	0.016230	2.0	0.000000	2.000000	
75%	0.104467	2.0	0.000000	2.000000	
max	0.373856	2.0	1.000000	2.000000	

	Kmd_tsne
count	2809.0
mean	1.0
std	0.0
min	1.0
25%	1.0
50%	1.0
75%	1.0
max	1.0

Cluster 1: Foods in this cluster have the lowest fat content, making them considered healthful.

```
[85]: Kmdtsne.get_group(2).describe()
```

```
[85]:
```

	Energy (Kcal)	Carbohydrate(g)	Protein(g)	Total Lipid(g)	\
count	2946.000000	2946.000000	2946.000000	2946.000000	

mean	0.863782	0.804548	0.459572	0.526301
std	0.053978	0.195272	0.165836	0.211173
min	0.656199	0.000000	0.000000	0.000000
25%	0.829515	0.733345	0.383500	0.377123
50%	0.869298	0.876045	0.478997	0.544305
75%	0.901848	0.929653	0.560591	0.674634
max	1.000000	0.986273	0.964519	1.000000

	Monounsaturated Fatty Acids(g)	Polyunsaturated Fatty Acids(g)	\
count	2946.000000	2946.000000	
mean	0.330171	0.299632	
std	0.200416	0.199654	
min	0.000000	0.000000	
25%	0.192767	0.137799	
50%	0.317862	0.256499	
75%	0.448238	0.421787	
max	1.000000	1.000000	

	Saturated Fatty Acids(g)	kmeans_tsne	Agg_tsne	Gmm_tsne	\
count	2946.000000	2946.000000	2946.000000	2946.000000	
mean	0.302040	0.004073	1.249491	0.999321	
std	0.180888	0.078073	0.966963	0.058268	
min	0.000000	0.000000	0.000000	0.000000	
25%	0.140891	0.000000	0.000000	1.000000	
50%	0.306539	0.000000	2.000000	1.000000	
75%	0.430340	0.000000	2.000000	1.000000	
max	0.903884	2.000000	2.000000	2.000000	

	Kmd_tsne
count	2946.0
mean	2.0
std	0.0
min	2.0
25%	2.0
50%	2.0
75%	2.0
max	2.0

Cluster 3: This food cluster appeared to be food that provides more energy due to high carbohydrate composition (around 0.73 to 0.98) as compared to other food clusters.

[86]: scores

[86]: {'kmeans_tsne': 0.43591416,
 'Agg_tsne': 0.38080552,
 'Gmm_tsne': 0.43544555,
 'Kmd_tsne': 0.4359154}

When the median strategy was used to fill in the missing values in the dataset, it was found that the K-Medoids model was the most successful clustering method, with the highest silhouette score 0.43606 recorded.

[]: