## Preparing Data And Preprocessing

```
In [1]: import pandas as pd
        import numpy as np
        import matplotlib.pyplot as plt
        import seaborn as sns
        from sklearn.ensemble import RandomForestRegressor
        from sklearn.decomposition import PCA
        from sklearn.cluster import DBSCAN
        from sklearn.preprocessing import StandardScaler, PowerTransformer
        from sklearn.feature selection import VarianceThreshold
        from sklearn.model selection import train test split
        from sklearn.metrics import mean squared error, r2 score
        from sklearn.neighbors import LocalOutlierFactor
        from kneed import KneeLocator
        from sklearn.cluster import KMeans
        from sklearn.mixture import GaussianMixture
        import umap.umap as umap
        # ---- STEP 1: LOAD DATA ----
        df = pd.read excel('./data files/Data re.xlsx') # Ensure the file exists
        # ---- STEP 2: FEATURE SELECTION ----
        cols to keep = ['g flux', 'r flux', 'i flux', 'y flux', 'z flux', 'specz
        df selected = df[cols to keep].copy()
        # Compute Flux Color Indices
        df selected['q r'] = df selected['q flux'] - df selected['r flux']
        df selected['r i'] = df selected['r flux'] - df selected['i flux']
        df_selected['i_y'] = df_selected['i_flux'] - df_selected['y_flux']
        df selected['y z'] = df selected['y flux'] - df selected['z flux']
        # ---- STEP 3: REMOVE OUTLIERS ----
        lof = LocalOutlierFactor(n_neighbors=20, contamination=0.02)
        outlier scores = lof.fit predict(df selected.drop(columns=['specz redshif
        df_clean = df_selected[outlier_scores == 1].copy()
        # ---- STEP 4: SCALING & TRANSFORMING ----
        scaler = StandardScaler()
        df scaled = scaler.fit transform(df clean.drop(columns=['specz redshift']
        # Apply Power Transformation (Yeo-Johnson for normalizing skewed data)
        power transformer = PowerTransformer(method='yeo-johnson')
        df_transformed = power_transformer.fit_transform(df_scaled)
        # ---- STEP 5: DIMENSIONALITY REDUCTION WITH UMAP ----
        reducer = umap.UMAP(n components=2, n neighbors=15, min dist=0.1, random
        X umap = reducer.fit transform(df transformed)
        # ---- STEP 6: OPTIONAL PCA FOR EXPLORATION ----
        pca = PCA(n components=3)
        X pca = pca.fit transform(df transformed)
        # ---- STEP 7: CLUSTERING ----
        kmeans = KMeans(n clusters=3, random state=42)
        df_clean['Cluster'] = kmeans.fit_predict(X_umap)
        # ---- PLOT RESULTS ----
```

```
plt.scatter(X_umap[:, 0], X_umap[:, 1], c=df_clean['Cluster'], cmap='viri
plt.xlabel('UMAP Component 1')
plt.ylabel('UMAP Component 2')
plt.title('UMAP Clustering')
plt.show()
```

2025-02-21 19:44:36.244662: E external/local\_xla/xla/stream\_executor/cuda/cuda\_fft.cc:477] Unable to register cuFFT factory: Attempting to register factory for plugin cuFFT when one has already been registered

WARNING: All log messages before absl::InitializeLog() is called are writt en to STDERR

E0000 00:00:1740147276.299220 4415 cuda\_dnn.cc:8310] Unable to register cuDNN factory: Attempting to register factory for plugin cuDNN when one has already been registered

E0000 00:00:1740147276.314946 4415 cuda\_blas.cc:1418] Unable to registe r cuBLAS factory: Attempting to register factory for plugin cuBLAS when on e has already been registered

2025-02-21 19:44:36.430661: I tensorflow/core/platform/cpu\_feature\_guard.c c:210] This TensorFlow binary is optimized to use available CPU instructions in performance-critical operations.

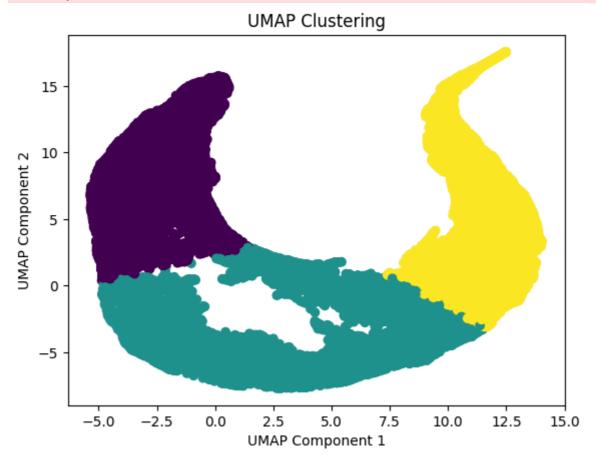
To enable the following instructions: AVX2 FMA, in other operations, rebuild TensorFlow with the appropriate compiler flags.

/home/chloy/miniconda3/lib/python3.10/site-packages/sklearn/utils/deprecat ion.py:151: FutureWarning: 'force\_all\_finite' was renamed to 'ensure\_all\_f inite' in 1.6 and will be removed in 1.8.

warnings.warn(

/home/chloy/miniconda3/lib/python3.10/site-packages/umap/umap\_.py:1952: Us erWarning: n\_jobs value 1 overridden to 1 by setting random\_state. Use no seed for parallelism.

warn(



checking

```
In [2]: print(df.head())
    print(df.info())
    print(df.describe())
```

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    b'(179325.3125, 99694.8046875, -21178.96484375)' -5.893433
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[5 rows x 89 columns]
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 40914 entries, 0 to 40913
Data columns (total 89 columns):
    Column
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g\_major\_axis

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11	g_minor_axis	40914	non-null	float64
12	g_peak_surface_brightness		non-null	float64
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16	i_central_image_pop_10px_rad		non-null	int64
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18	i_central_image_pop_5px_rad		non-null	int64
19	i_cmodel_mag		non-null	float64
20	i_cmodel_magsigma	40914	non-null	float64
21	i_ellipticity	40914	non-null	float64
22	i_half_light_radius	40914	non-null	float64
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25	i minor axis	40914	non-null	float64
26	i peak surface brightness	40914	non-null	float64
27	i_petro_rad		non-null	float64
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36	r_ellipticity		non-null	float64
37	r_half_light_radius	40914	non-null	float64
38	r_isophotal_area	40914	non-null	int64
39	r_major_axis	40914	non-null	float64
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41	r peak surface brightness	40914	non-null	float64
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43	r_pos_angle		non-null	float64
44	r_sersic_index		non-null	float64
45	ra		non-null	float64
46	skymap id		non-null	int64
47	specz dec		non-null	float64
48	specz_dec specz_flag_homogeneous		non-null	bool
49			non-null	float64
	specz_mag_i			
50	specz_name		non-null	object
51	specz_ra		non-null	float64
52	specz_redshift		non-null	float64
53	specz_redshift_err		non-null	float64
54	x_coord		non-null	float64
55	<pre>y_central_image_pop_10px_rad</pre>		non-null	int64
56	<pre>y_central_image_pop_15px_rad</pre>	40914	non-null	int64
57	<pre>y_central_image_pop_5px_rad</pre>	40914	non-null	int64
58	y_cmodel_mag	40914	non-null	float64
59	y_cmodel_magsigma	40914	non-null	float64
60	y coord	40914	non-null	float64
61	y ellipticity	40914	non-null	float64
62	y half light radius		non-null	float64
63	y isophotal area		non-null	int64
64	y_major_axis		non-null	float64
65	y_minor_axis		non-null	float64
66	y_peak_surface_brightness		non-null	float64
67	y_petro_rad		non-null	float64
68	y pos angle		non-null	float64
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70	z_central_image_pop_10px_rad	40914	non-null	int64

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[8 rows x 86 columns]

Clustering (DBScan, K means and Gaussian Mixture Method)

```
In [3]: from sklearn.cluster import DBSCAN, KMeans
    from sklearn.mixture import GaussianMixture
    from sklearn.neighbors import NearestNeighbors
    from sklearn.metrics import silhouette_score
    import matplotlib.pyplot as plt
    import numpy as np
    from kneed import KneeLocator

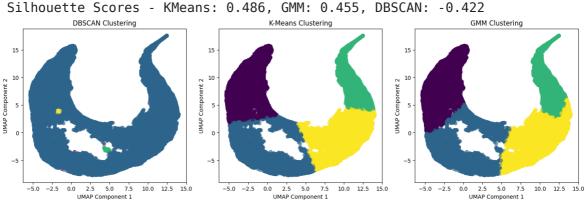
# ---- STEP 6: DETERMINE OPTIMAL DBSCAN EPS ----
k = 5 # Typically, min_samples value
nearest_neighbors = NearestNeighbors(n_neighbors=k)
nearest_neighbors.fit(X_umap)
distances, indices = nearest_neighbors.kneighbors(X_umap)

# Sort distances to find the "knee" point
distances = np.sort(distances[:, -1])
```

```
# Use KneeLocator to find optimal epsilon
knee locator = KneeLocator(range(len(distances)), distances, curve="conve"
optimal eps = distances[knee locator.elbow]
print(f"Optimal eps for DBSCAN: {optimal eps:.3f}")
# ---- STEP 6A: APPLY DBSCAN CLUSTERING WITH OPTIMAL EPS ----
dbscan = DBSCAN(eps=optimal eps, min samples=k, metric='euclidean')
cluster labels dbscan = dbscan.fit predict(X umap)
df_clean.loc[:, 'cluster_dbscan'] = cluster_labels_dbscan
# ---- STEP 6B: APPLY K-MEANS & GMM ----
inertia = []
silhouette scores = []
k range = range(2, 10)
for k in k range:
    kmeans = KMeans(n clusters=k, random state=42, n init='auto')
    labels = kmeans.fit predict(X umap)
    inertia.append(kmeans.inertia )
    score = silhouette score(X umap, labels)
    silhouette scores.append(score)
knee locator = KneeLocator(k range, inertia, curve="convex", direction="d
optimal k = knee locator.elbow
print(f"Optimal k for K-Means/GMM: {optimal k}")
# Apply K-Means with the optimal k
kmeans = KMeans(n clusters=optimal k, random state=42, n init='auto')
df clean.loc[:, 'cluster kmeans'] = kmeans.fit predict(X umap)
# Apply Gaussian Mixture Model (GMM)
gmm = GaussianMixture(n components=optimal k, random state=42)
df_clean.loc[:, 'cluster_gmm'] = gmm.fit_predict(X_umap)
# ---- STEP 7: COMPUTE SILHOUETTE SCORES ----
silhouette kmeans = silhouette score(X umap, df clean['cluster kmeans'])
silhouette_gmm = silhouette_score(X_umap, df_clean['cluster_gmm'])
silhouette_dbscan = silhouette_score(X_umap[df_clean['cluster_dbscan'] !=
                                     df_clean['cluster_dbscan'][df_clean[
print(f"Silhouette Scores - KMeans: {silhouette kmeans:.3f}, GMM: {silhou
# ---- STEP 8: VISUALIZE CLUSTERING METHODS (2D PLOTS) ----
fig, ax = plt.subplots(1, 3, figsize=(18, 5))
# DBSCAN 2D Plot
ax[0].scatter(X umap[:, 0], X umap[:, 1], c=cluster labels dbscan, cmap='
ax[0].set_title("DBSCAN Clustering")
ax[0].set xlabel("UMAP Component 1")
ax[0].set_ylabel("UMAP Component 2")
# K-Means 2D Plot
ax[1].scatter(X_umap[:, 0], X_umap[:, 1], c=df_clean['cluster_kmeans'], c
ax[1].set title("K-Means Clustering")
ax[1].set xlabel("UMAP Component 1")
ax[1].set_ylabel("UMAP Component 2")
# GMM 2D Plot
ax[2].scatter(X_umap[:, 0], X_umap[:, 1], c=df_clean['cluster_gmm'], cmap
```

```
ax[2].set_title("GMM Clustering")
ax[2].set_xlabel("UMAP Component 1")
ax[2].set_ylabel("UMAP Component 2")
plt.show()
```

Optimal eps for DBSCAN: 0.323
Optimal k for K-Means/GMM: 4



## Random Forest

```
In [4]: # ---- STEP 8: RANDOM FOREST REGRESSION ----
        # Train separate Random Forest models for each clustering method using fl
        results = {}
        for cluster type in ['cluster dbscan', 'cluster kmeans', 'cluster qmm']:
            df temp = pd.get dummies(df clean, columns=[cluster type], prefix=[f'
            X = df_temp[['g_flux', 'r_flux', 'i_flux', 'y_flux', 'z_flux'] +
                        [col for col in df temp.columns if col.startswith(f'clust
            y = df temp['specz redshift']
            X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0
            rf = RandomForestRegressor(n_estimators=100, random_state=42)
            rf.fit(X train, y train)
            y_pred = rf.predict(X_test)
            mse = mean_squared_error(y_test, y_pred)
            r2 = r2_score(y_test, y_pred)
            results[cluster_type] = {'MSE': mse, 'R2': r2, 'y_pred': y_pred}
        # ---- STEP 9: VISUALIZE REGRESSION RESULTS ----
        fig, ax = plt.subplots(1, 3, figsize=(18, 6))
        titles = ['DBSCAN', 'K-Means', 'GMM']
        colors = ['blue', 'green', 'red']
        for i, cluster_type in enumerate(results.keys()):
            ax[i].scatter(y test, results[cluster type]['y pred'], alpha=0.6, col
            ax[i].set xlabel('Actual Redshift')
            ax[i].set ylabel('Predicted Redshift')
            ax[i].set title(f'Actual vs Predicted Redshift ({titles[i]})')
            ax[i].plot([y_test.min(), y_test.max()], [y_test.min(), y_test.max()]
        plt.tight layout()
        plt.show()
```

```
---- STEP 10: COMPARE RESULTS ACROSS CLUSTERING METHODS ----
        print("Random Forest Regression Results:")
        for cluster type, metrics in results.items():
            print(f"{cluster_type.upper()} - MSE: {metrics['MSE']:.6f}, R2: {metr
                                       Actual vs Predicted Redshift (K-Means)
                                                                 Actual vs Predicted Redshift (GMM)
       Random Forest Regression Results:
       CLUSTER_DBSCAN - MSE: 0.103494, R2: 0.673538
       CLUSTER KMEANS - MSE: 0.103177, R2: 0.674537
       CLUSTER GMM - MSE: 0.103596, R2: 0.673214
        Checking
In [5]:
        print(df clean.head())
        print(df clean.columns)
                                             i flux
                               r flux
                                                           y flux
                                                                          z flux
                g flux
         7.482335e-09 2.987891e-08 5.117533e-08 8.117985e-08 6.737572e-08
       1 1.297347e-09 5.643225e-09 1.319162e-08 2.141019e-08 1.828079e-08
       2 3.471398e-09 1.571304e-08 3.743521e-08 6.504857e-08 5.585498e-08
       3 4.114510e-08 1.090914e-07 1.841788e-07 3.080686e-07 2.402212e-07
       4 3.987357e-09 1.373544e-08 2.634882e-08 4.449762e-08 3.527128e-08
          specz redshift
                                    g_r
                                                  r_i
                                                                 iу
                                                                               y_z
       \
                 0.31652 -2.239657e-08 -2.129643e-08 -3.000452e-08 1.380413e-08
       0
       1
                 0.56769 -4.345878e-09 -7.548391e-09 -8.218576e-09 3.129404e-09
       2
                 0.53428 -1.224164e-08 -2.172218e-08 -2.761336e-08 9.193588e-09
       3
                 0.11878 -6.794635e-08 -7.508739e-08 -1.238898e-07 6.784739e-08
                 0.23497 -9.748082e-09 -1.261338e-08 -1.814880e-08 9.226337e-09
          Cluster cluster dbscan cluster kmeans cluster gmm
       0
                2
                                                               1
       1
                1
                                 0
                                                 1
       2
                1
                                 0
                                                 3
                                                               3
                                                               2
       3
                2
       Index(['g_flux', 'r_flux', 'i_flux', 'y_flux', 'z_flux', 'specz_redshift',
              'g_r', 'r_i', 'i_y', 'y_z', 'Cluster', 'cluster_dbscan',
              'cluster_kmeans', 'cluster_gmm'],
             dtype='object')
        SVR and UMAP
In [6]:
       # Add-Ons for Photometric Redshift Estimation
        # Fixing UMAP Import Issue and Enhancing Preprocessing
          ---- STEP 11: Install and Import UMAP Properly ----
```

```
# Ensure proper UMAP installation: pip install umap-learn
from umap import UMAP
umap = UMAP(n neighbors=15, min dist=0.1, n components=2, random state=42
X umap = umap.fit transform(df transformed)
# ---- STEP 12: Add More Color Indices (Using Flux Values) ----
df clean['u q'] = df clean['q flux'] - df clean['r flux']
df_clean['g_r'] = df_clean['r_flux'] - df_clean['i_flux']
df_clean['r_i'] = df_clean['i_flux'] - df_clean['y_flux']
df clean['i z'] = df clean['y flux'] - df clean['z flux']
# ---- STEP 13: Additional Clustering (HDBSCAN) for SVR Comparison ----
try:
    from hdbscan import HDBSCAN
except ImportError:
    print("HDBSCAN not installed. Use: pip install hdbscan")
hdbscan = HDBSCAN(min cluster size=8)
df clean['cluster hdbscan'] = hdbscan.fit predict(X umap)
# ---- STEP 14: SVR Regression for Redshift Estimation ----
from sklearn.svm import SVR
svr results = {}
for cluster_type in ['cluster_kmeans', 'cluster_gmm', 'cluster_hdbscan']:
    df temp = pd.get dummies(df clean, columns=[cluster type], prefix=[f'
    X = df temp.drop(columns=['specz redshift'])
    y = df temp['specz redshift']
   X train, X test, y train, y test = train test split(X, y, test size=0
    svr = SVR(kernel='rbf', C=1.0, epsilon=0.1)
    svr.fit(X train, y train)
    y pred = svr.predict(X test)
    mse = mean_squared_error(y_test, y_pred)
    r2 = r2 score(y test, y pred)
    svr results[cluster type] = {'MSE': mse, 'R2': r2, 'y pred': y pred}
# ---- STEP 15: Plot SVR Results ----
fig, ax = plt.subplots(1, 3, figsize=(18, 6))
titles = ['K-Means', 'GMM', 'HDBSCAN']
colors = ['purple', 'orange', 'cyan']
for i, cluster type in enumerate(svr results.keys()):
    ax[i].scatter(y_test, svr_results[cluster_type]['y_pred'], alpha=0.6,
    ax[i].set title(f'SVR Actual vs Predicted ({titles[i]})')
    ax[i].set_xlabel('Actual Redshift')
    ax[i].set_ylabel('Predicted Redshift')
    ax[i].plot([y test.min(), y test.max()], [y test.min(), y test.max()]
plt.tight layout()
plt.show()
# ---- Print SVR Performance Metrics ----
print('SVR Regression Results:')
```

```
for cluster_type, metrics in svr_results.items():
    print(f"{cluster_type.upper()} - MSE: {metrics['MSE']:.6f}, R2: {metr
```

/home/chloy/miniconda3/lib/python3.10/site-packages/sklearn/utils/deprecat ion.py:151: FutureWarning: 'force\_all\_finite' was renamed to 'ensure\_all\_f inite' in 1.6 and will be removed in 1.8.

warnings.warn(

/home/chloy/miniconda3/lib/python3.10/site-packages/umap/umap\_.py:1952: Us erWarning: n\_jobs value 1 overridden to 1 by setting random\_state. Use no seed for parallelism.

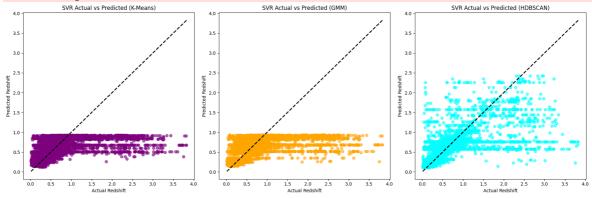
warn(

/home/chloy/miniconda3/lib/python3.10/site-packages/sklearn/utils/deprecat ion.py:151: FutureWarning: 'force\_all\_finite' was renamed to 'ensure\_all\_f inite' in 1.6 and will be removed in 1.8.

warnings.warn(

/home/chloy/miniconda3/lib/python3.10/site-packages/sklearn/utils/deprecat ion.py:151: FutureWarning: 'force\_all\_finite' was renamed to 'ensure\_all\_f inite' in 1.6 and will be removed in 1.8.

warnings.warn(



SVR Regression Results:

CLUSTER\_KMEANS - MSE: 0.227899, R2: 0.281111 CLUSTER\_GMM - MSE: 0.226127, R2: 0.286701 CLUSTER HDBSCAN - MSE: 0.167378, R2: 0.472019