

Lab Notebook

Photonic Lantern Information Determination

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1 Mini Dataset Information Determination

2 The data

2.1 Zernike coefficients dataset

A dataset of 1000 zernike coefficients is created for this report. In particular, each datapoint represents the coefficients of the first 2 Zernike modes, their values ranging between $[-2, -1.8]$ and $[1.8, 2]$.

2.2 PSFs coefficients dataset

A dataset of 1000 PSFs is created using the Zernike coefficients dataset.

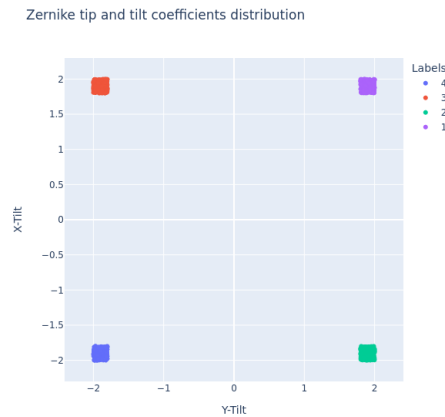
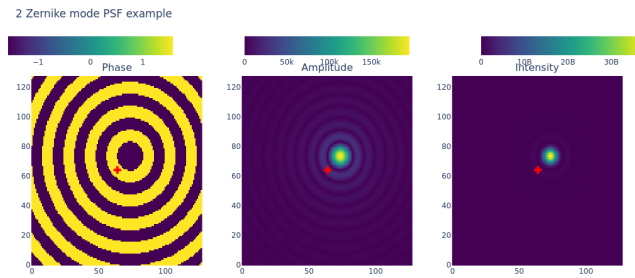
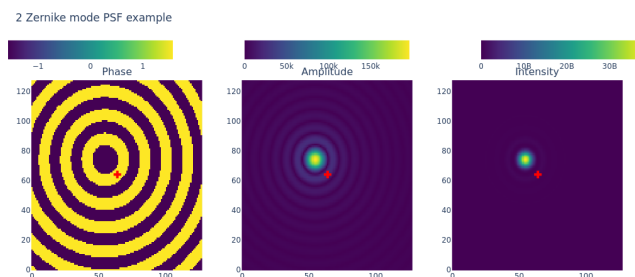


Figure 1: Example original sized PSF

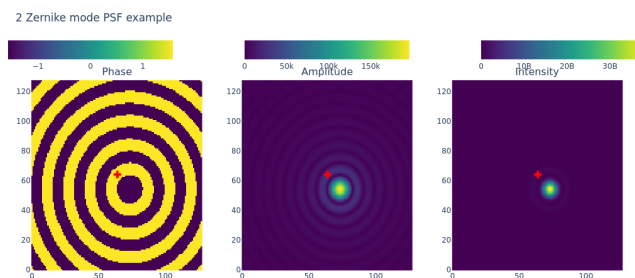
These ranges create 4 original clusters that will be used as reference.



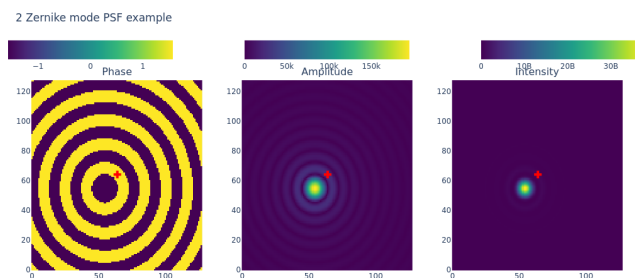
(a) Positive X-tilt and Positive Y-tilt PSF example



(b) Negative X-tilt and Positive Y-tilt PSF example



(c) Positive X-tilt and Negative Y-tilt PSF example



(d) Negative X-tilt and Negative Y-tilt PSF example

Figure 2: 2 Zernike modes PSF examples

2.3 LP mode coefficients dataset

A dataset of 1000 LP mode coefficients obtained from computing the overlap integral of the first 19 LP modes with the PSF dataset.

2.4 LP mode coefficients dataset

A dataset of 1000 PL output fluxes obtained from the PL transfer matrix and LP coefficients.

3 Preprocessing

3.1 PSF Intensities

The 1000x128x128 array is dimensionally reduced using PCA and UMAP both giving an array of 1000x19 projections of the PSF Intensities.

3.2 LP Coefficients

The 1000x19x2 array is dimensionally reduced using PCA and UMAP both giving an array of 1000x2 projections of the original LP coefficients.

3.3 Output fluxes

The 1000x19 array is dimensionally reduced using PCA and UMAP both giving an array of 1000x2 projections of the original LP coefficients.

4 Clustering

A series of different clustering algorithms are used:

- K-Means
- DBSCAN
- HDBSCAN
- Agglomerative clustering

The clusters obtained will be compared the original clusters using NMI

4.1 Zernike coefficients clustering

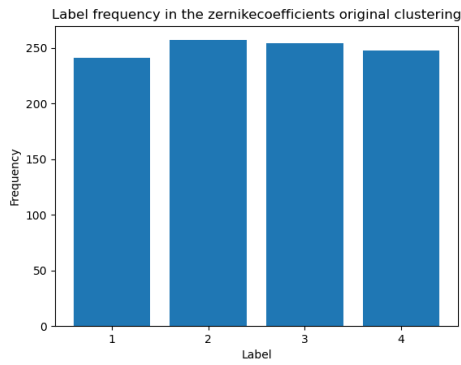
4.1.1 K-Means

As K-Means allows for the number of clusters to define, and we know that there are 4 in the original dataset, K-Means is used to find 4 clusters.

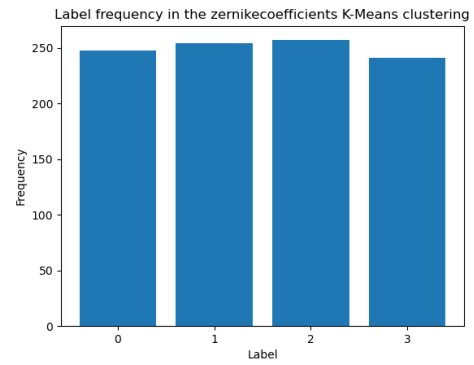
Number of clusters	Number of initializations
4	10

Table 1: K-Means hyperparameter configuration for Zernike coefficients clustering

The results are the following:

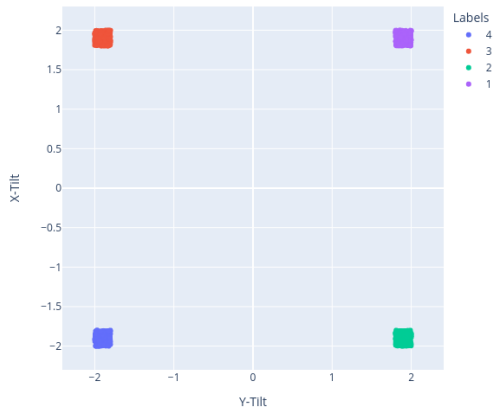


(a) Original cluster densities



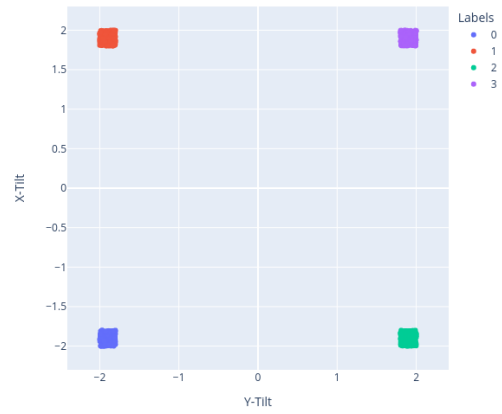
(b) K-Means clusters densities

Zernike tip and tilt original clusters



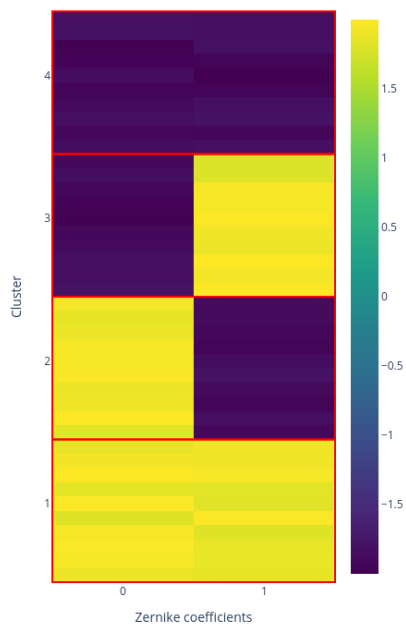
(c) Original clusters

Zernike tip and tilt K-Means clusters



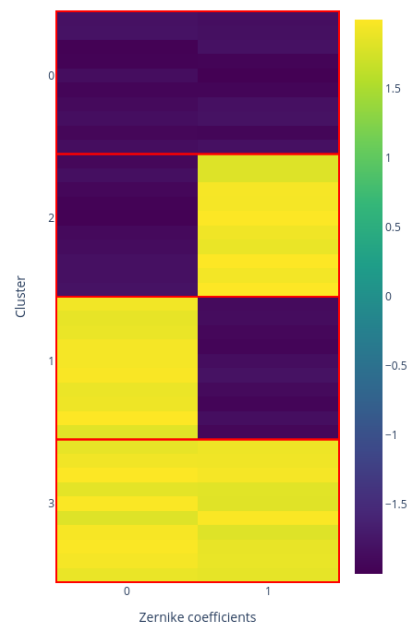
(d) K-Means clusters

Original Zernike coefficients cluster samples



(e) Original cluster samples

K-Means Zernike coefficients cluster samples



(f) K-Means cluster samples

Figure 3: Comparison between original clustering and K-Means clustering

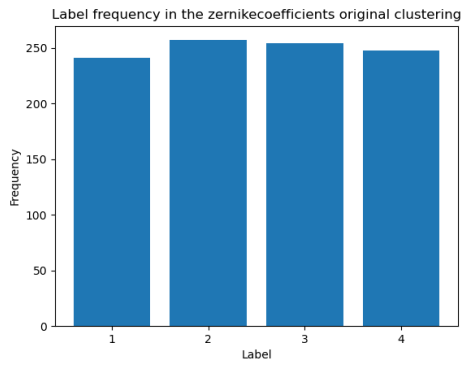
4.1.2 DBSCAN

A configuration that outputs 4 clusters is searched

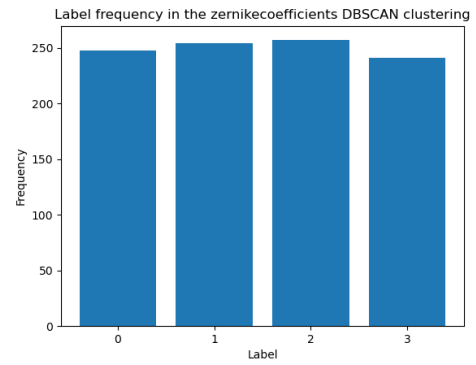
Number of neighbours	Epsilon
5	0.3

Table 2: DBSCAN hyperparameter configuration for Zernike coefficients clustering

The results are the following:

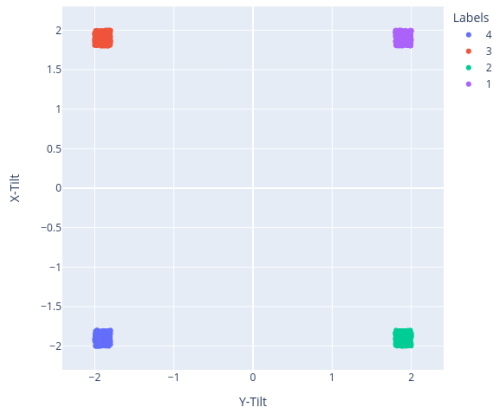


(a) Original cluster densities



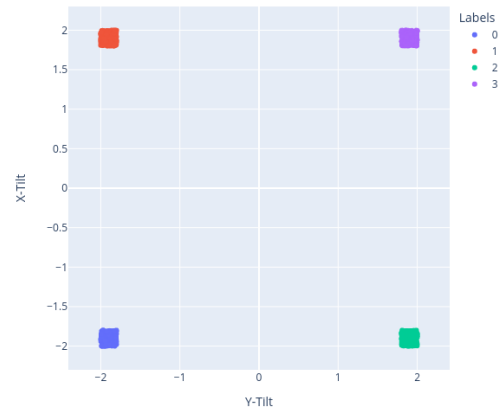
(b) DBSCAN clusters densities

Zernike tip and tilt original clusters



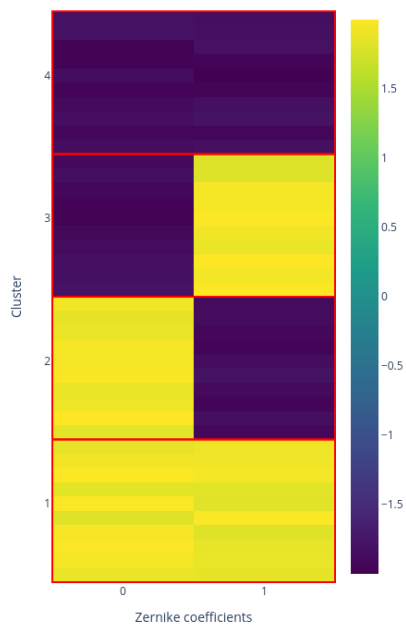
(c) Original clusters

Zernike tip and tilt DBSCAN clusters



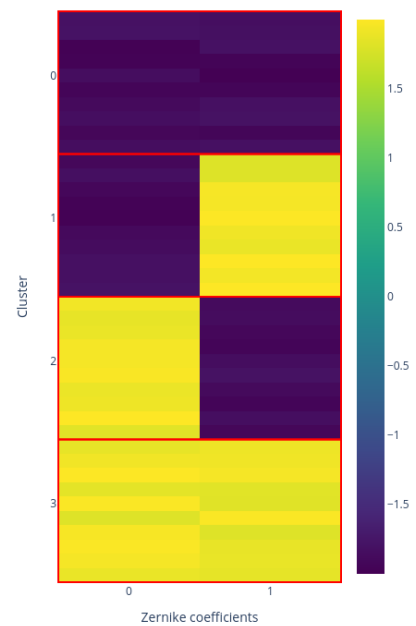
(d) DBSCAN clusters

Original Zernike coefficients cluster samples



(e) Original cluster samples

DBSCAN Zernike coefficients cluster samples



(f) DBSCAN cluster samples

Figure 4: Comparison between original clustering and DBSCAN clustering

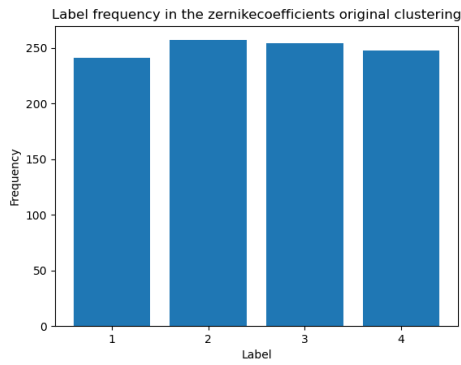
4.1.3 HDBSCAN

A configuration that outputs 4 clusters is searched.

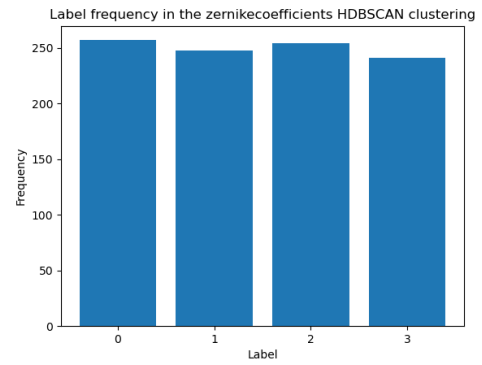
Minimum cluster size
5

Table 3: HDBSCAN hyperparameter configuration for Zernike coefficients clustering

The results are the following:

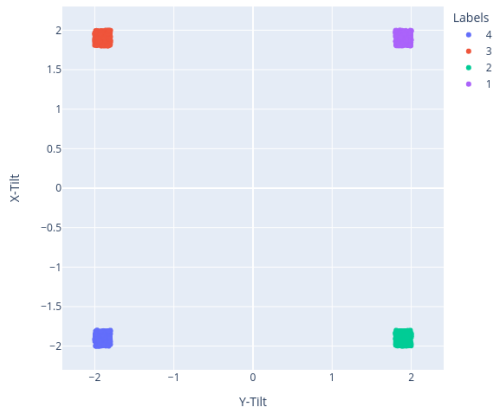


(a) Original cluster densities



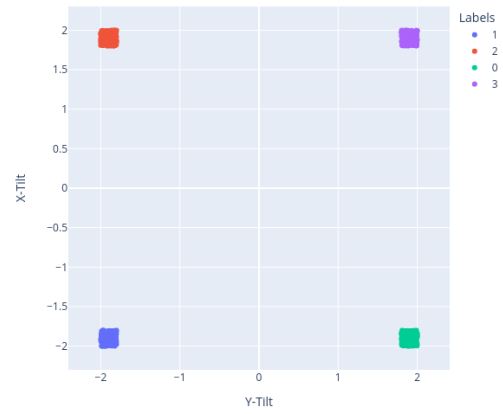
(b) HDBSCAN clusters densities

Zernike tip and tilt original clusters



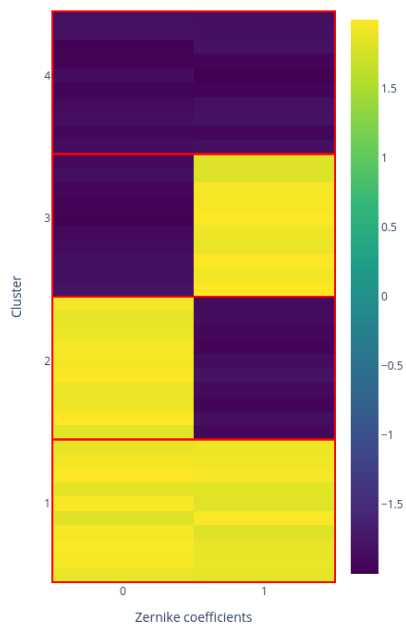
(c) Original clusters

Zernike tip and tilt HDBSCAN clusters



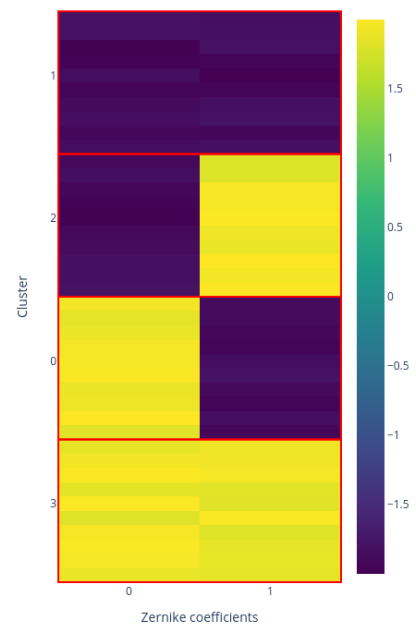
(d) HDBSCAN clusters

Original Zernike coefficients cluster samples



(e) Original cluster samples

HDBSCAN Zernike coefficients cluster samples



(f) HDBSCAN cluster samples

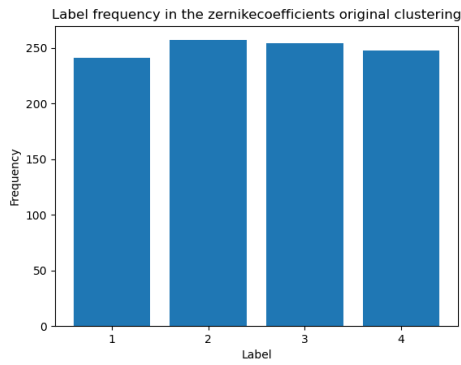
Figure 5: Comparison between original clustering and HDBSCAN clustering

4.1.4 Agglomerative clustering

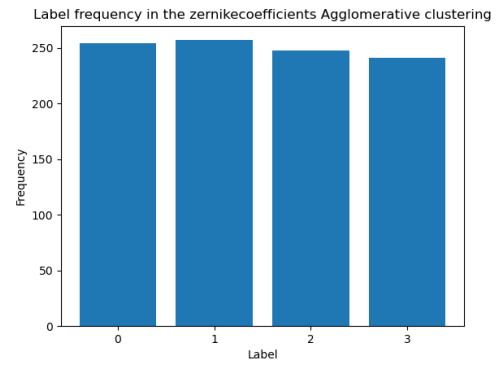
Number of clusters 4
5

Table 4: Agglomerative hyperparameter configuration for Zernike coefficients clustering

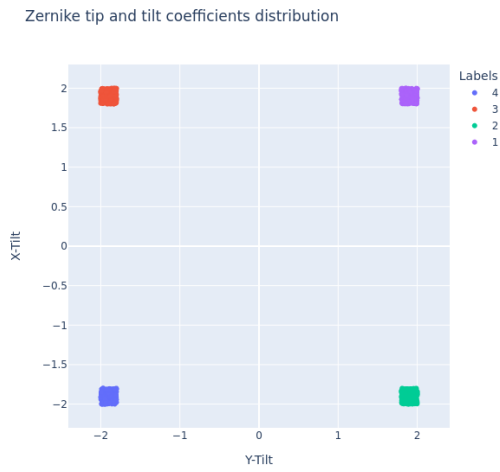
The results are the following:



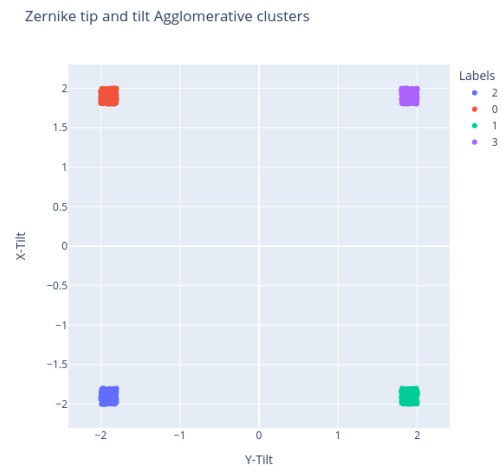
(a) Original cluster densities



(b) Agglomerative clusters densities

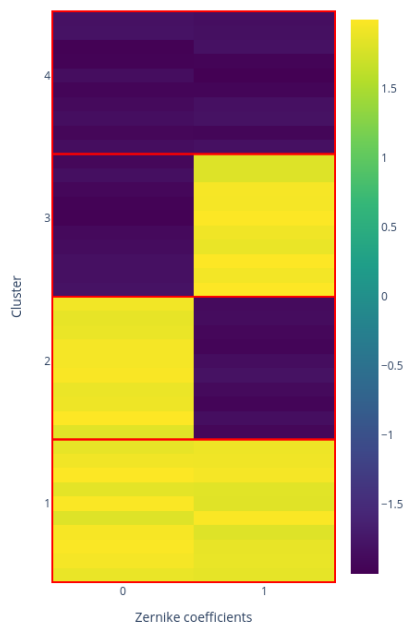


(c) Original clusters



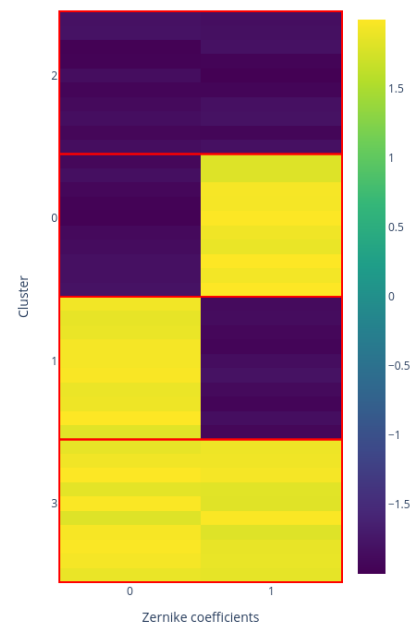
(d) Agglomerative clusters

Original Zernike coefficients cluster samples



(e) Original cluster samples

Agglomerative Zernike coefficients cluster samples



(f) Agglomerative cluster samples

Figure 6: Comparison between original clustering and Agglomerative clustering

4.1.5 Summary

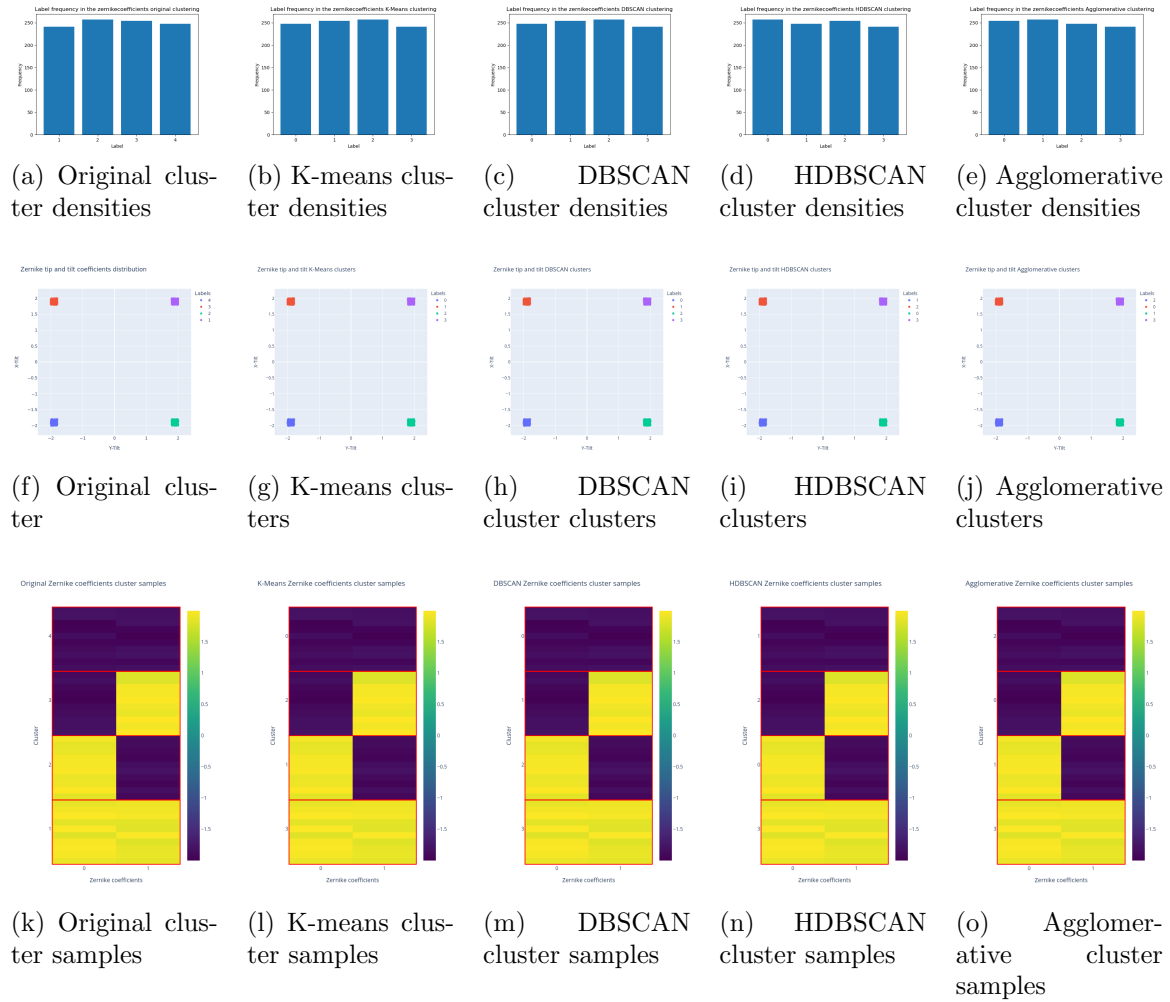


Figure 7: Comparison between clustering algorithms

	Original	K-Means	DBSCAN	HDBSCAN	Agglomerative
Original	\	1	1	1	1
K-Means		\	1	1	1
DBSCAN			\	1	1
HDBSCAN				\	1

Table 5: Normalized Mutual Information between clusters

4.2 LP coefficients clustering

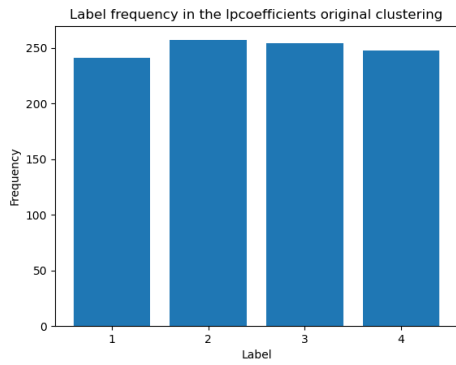
4.2.1 K-Means

As K-Means allows for the number of clusters to be defined, and we know that there are 4 in the original dataset, K-Means is used to find 4 clusters.

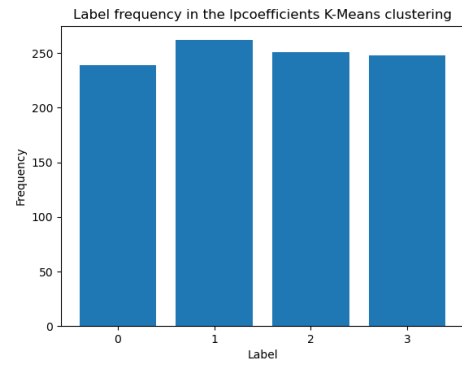
	Number of clusters	Number of initializations
Original LP coefficients	4	10
PCA LP coefficients	4	10
UMAP LP coefficients	4	10

Table 6: K-Means hyperparameter configuration for c coefficients clustering

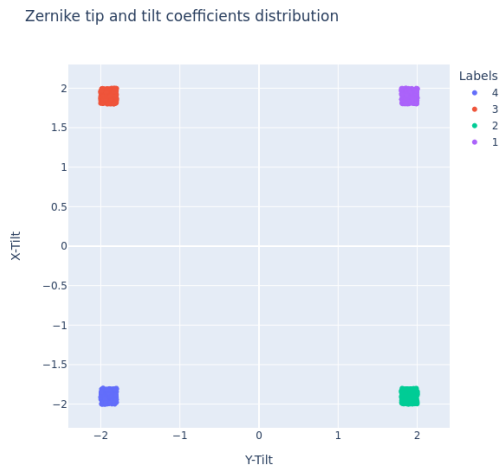
The results are the following:



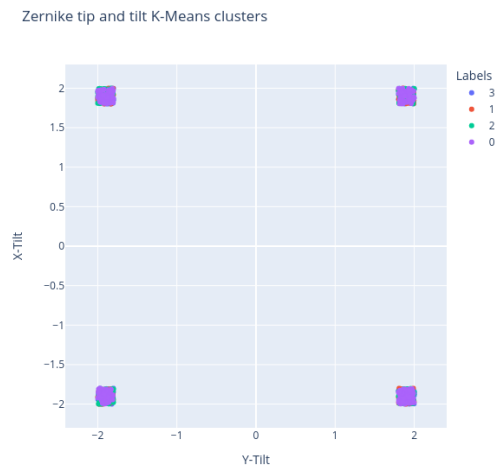
(a) Original cluster densities



(b) K-Means clusters densities

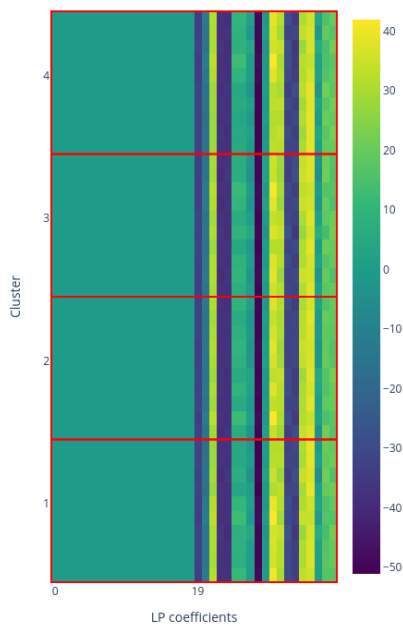


(c) Original clusters



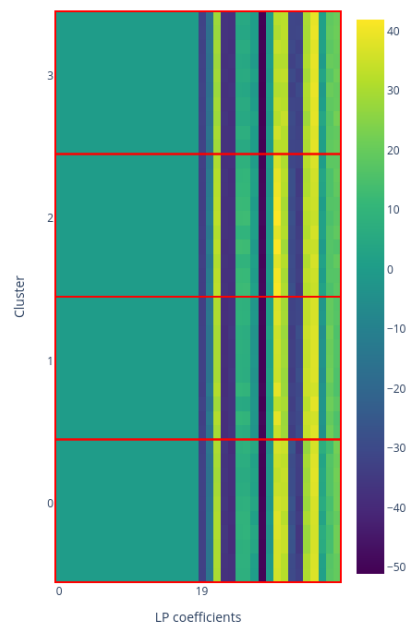
(d) K-Means clusters

Original LP coefficients cluster samples



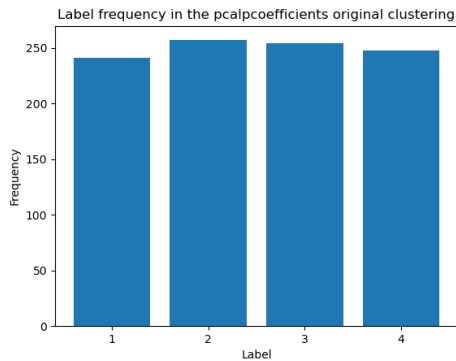
(e) Original cluster samples

K-Means LP coefficients cluster samples

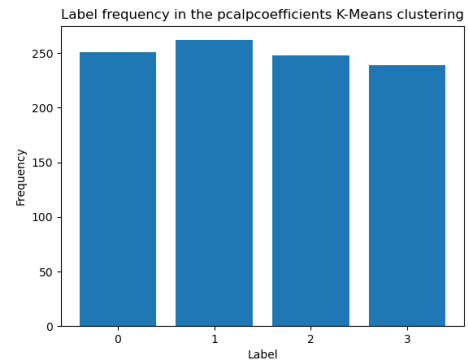


(f) K-Means cluster samples

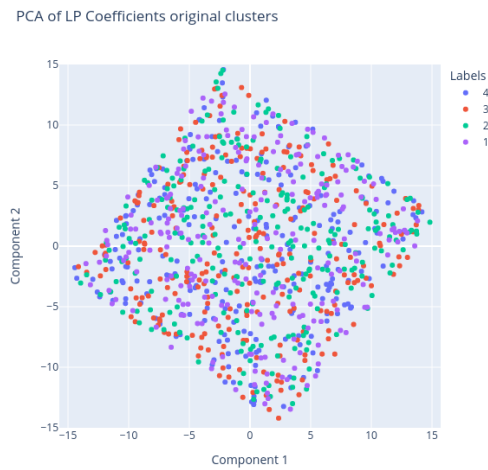
Figure 8: Comparison between original clustering and K-Means clustering from original LP coefficients



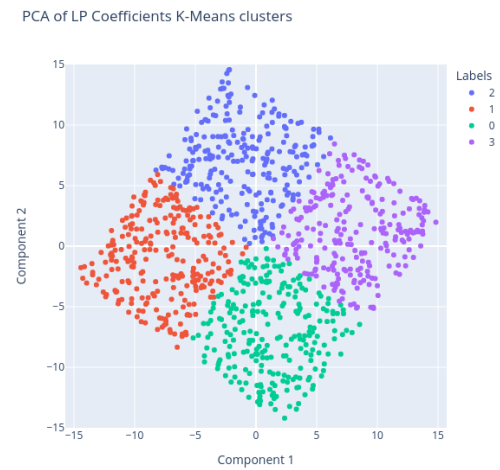
(a) Original cluster densities from PCA



(b) K-Means clusters densities from PCA

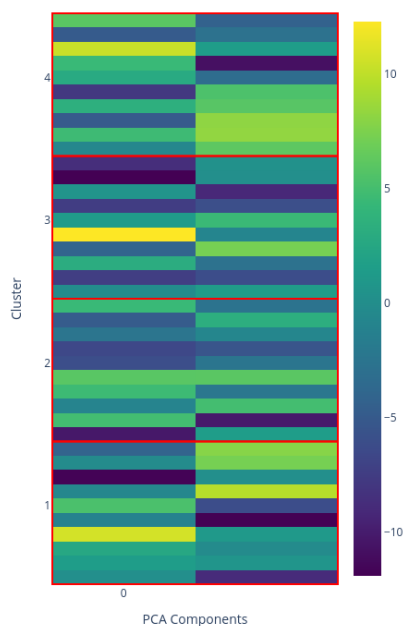


(c) Original clusters



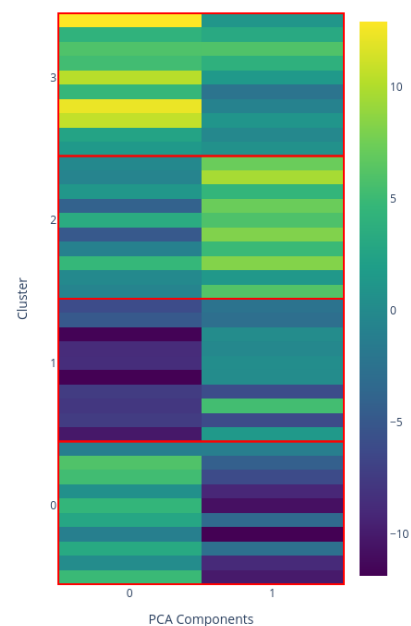
(d) K-Means clusters from PCA

Original PCA LP coefficients cluster samples



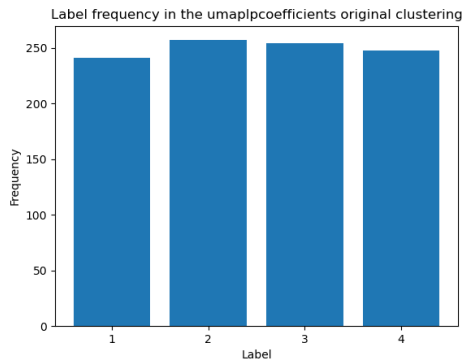
(e) Original cluster samples from PCA

K-Means PCA LP coefficients cluster samples

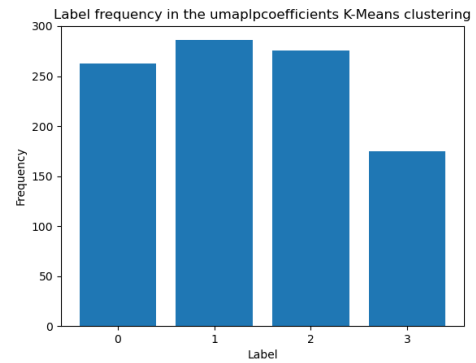


(f) K-Means cluster samples from PCA

Figure 9: Comparison between original clustering and K-Means clustering from PCA of LP coefficients

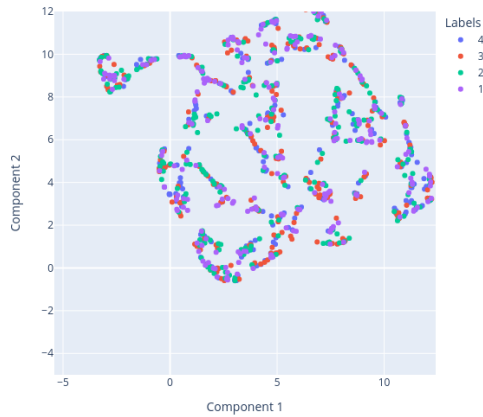


(a) Original cluster densities from UMAP



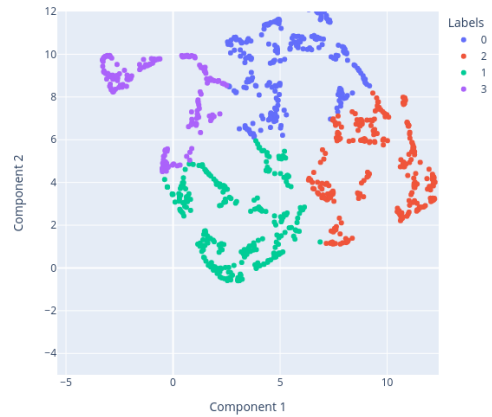
(b) K-Means clusters densities from UMAP

UMAP of LP Coefficients original clusters



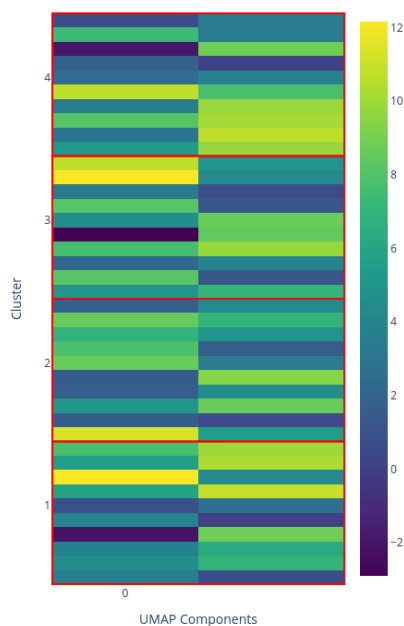
(c) Original clusters from UMAP

UMAP of LP Coefficients K-Means clusters



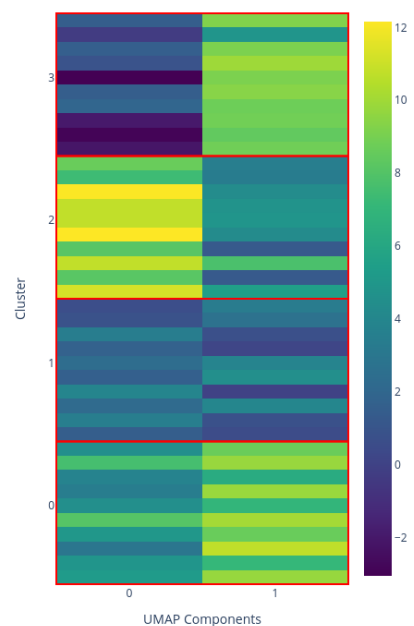
(d) K-Means clusters from UMAP

Original UMAP LP coefficients cluster samples



(e) Original cluster samples from UMAP

K-Means UMAP LP coefficients cluster samples



(f) K-Means cluster samples from UMAP

Figure 10: Comparison between original clustering and K-Means clustering from

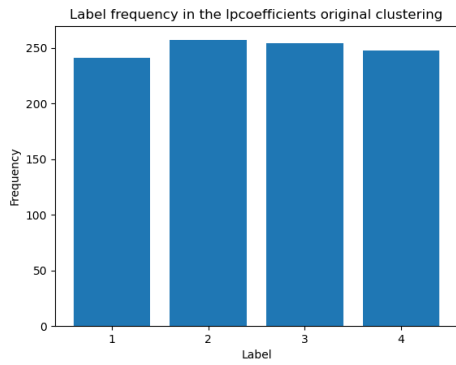
4.2.2 DBSCAN

A configuration that outputs 4 clusters is searched

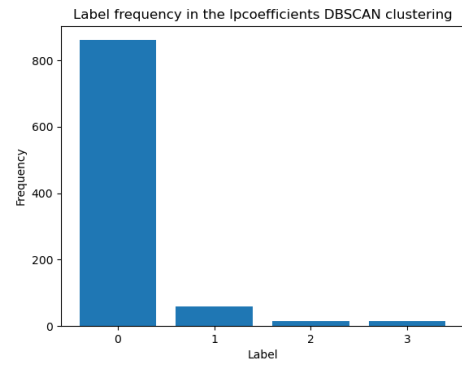
	Number of neighbours	Epsilon
Original LP coefficients	15	1.52
PCA LP coefficients	15	1.52
UMAP LP coefficients	10	0.85

Table 7: DBSCAN hyperparameter configuration for LP coefficients clustering

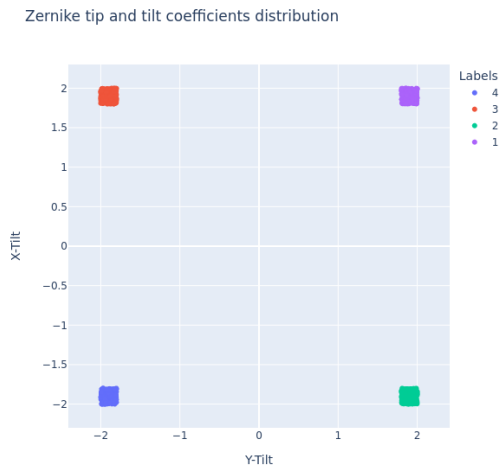
The results are the following:



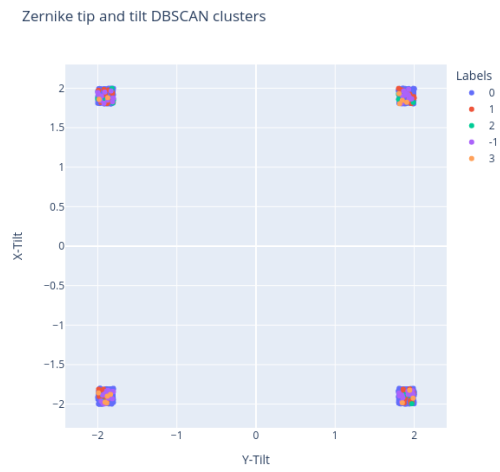
(a) Original cluster densities



(b) DBSCAN clusters densities

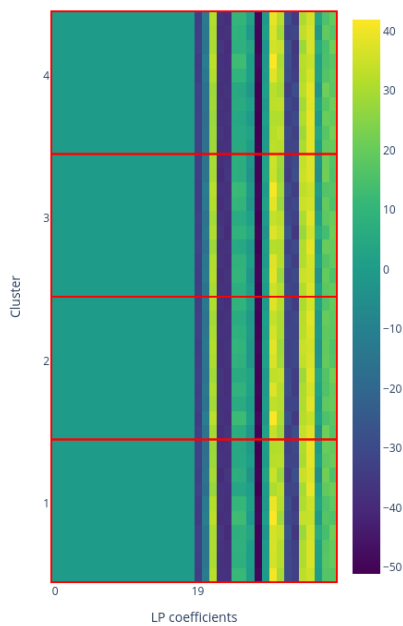


(c) Original clusters



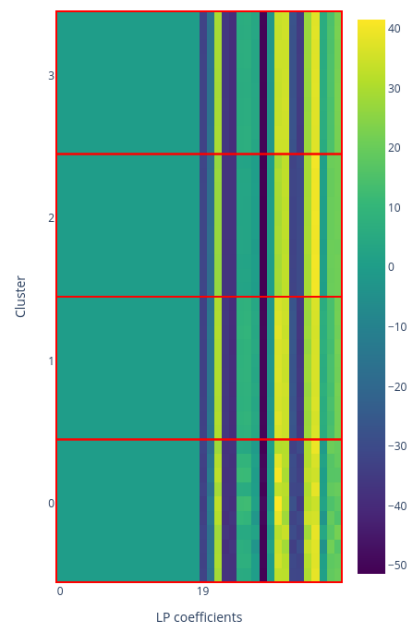
(d) DBSCAN clusters

Original LP coefficients cluster samples



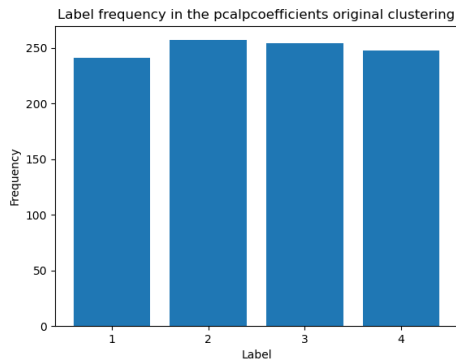
(e) Original cluster samples

DBSCAN LP coefficients cluster samples

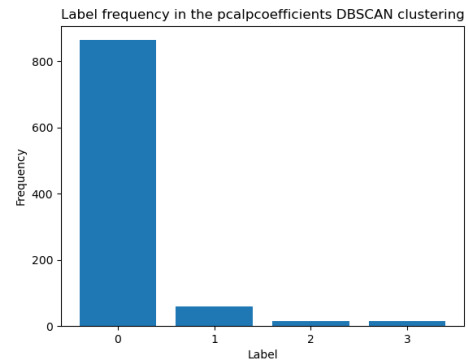


(f) DBSCAN cluster samples

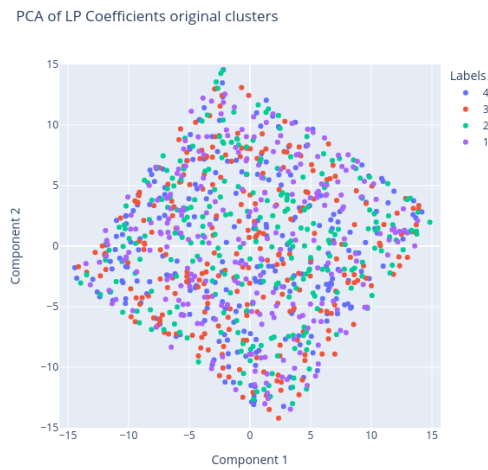
Figure 11: Comparison between original clustering and DBSCAN clustering



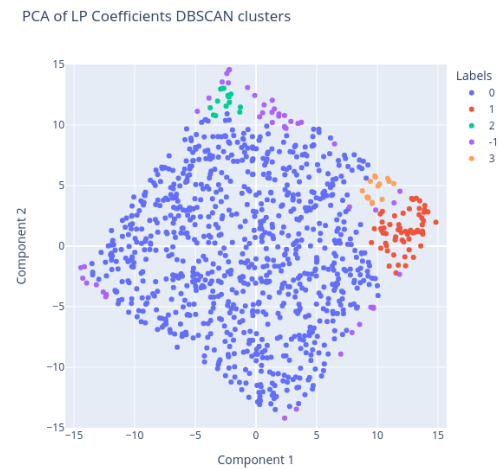
(a) Original cluster densities from PCA



(b) DBSCAN clusters densities from PCA

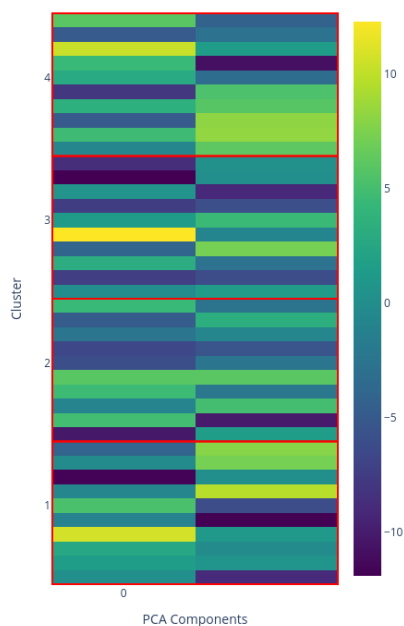


(c) Original clusters from PCA



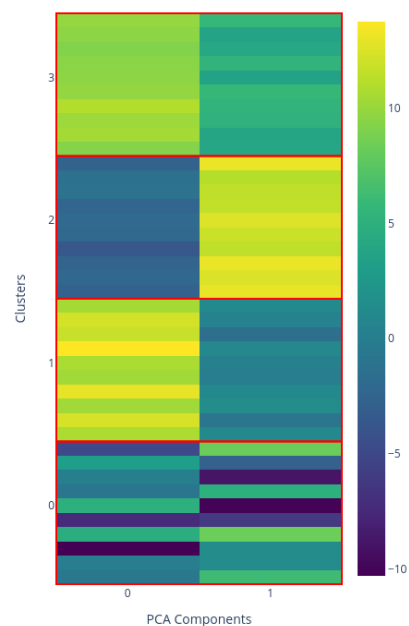
(d) DBSCAN clusters from PCA

Original PCA LP coefficients cluster samples



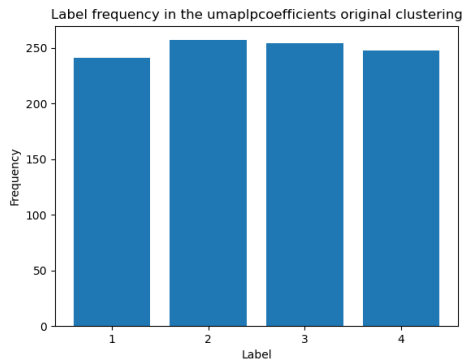
(e) Original cluster samples from PCA

DBSCAN PCA LP coefficients cluster samples

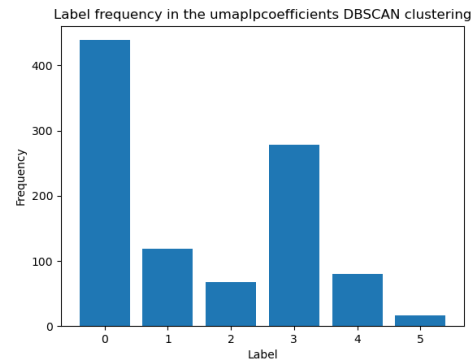


(f) DBSCAN cluster samples from PCA

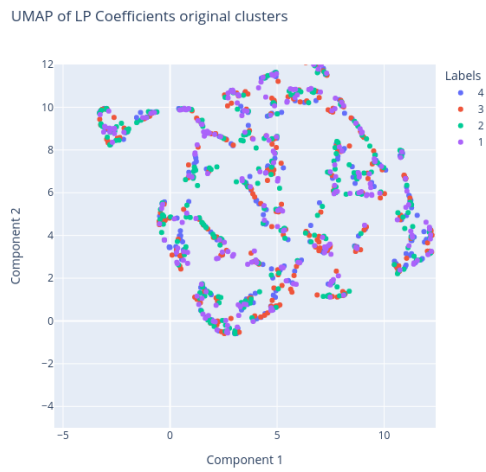
Figure 12: Comparison between original clustering and DBSCAN clustering from PCA of LP coefficients



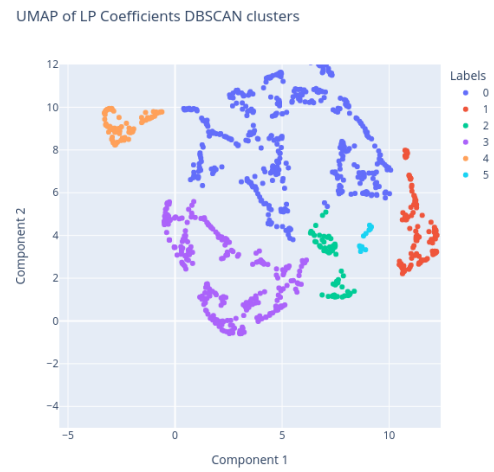
(a) Original cluster densities from UMAP



(b) DBSCAN clusters densities from UMAP

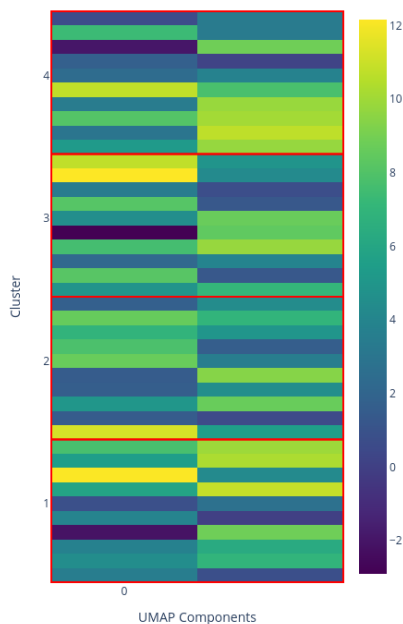


(c) Original clusters from UMAP



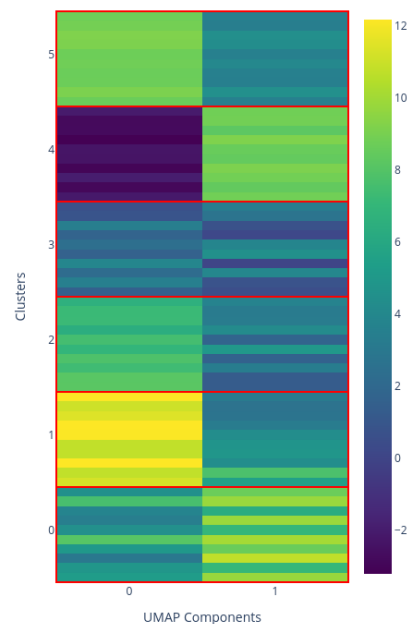
(d) DBSCAN clusters from UMAP

Original UMAP LP coefficients cluster samples



(e) Original cluster samples from UMAP

DBSCAN UMAP LP coefficients cluster samples



(f) DBSCAN cluster samples from UMAP

Figure 13: Comparison between original clustering and DBSCAN clustering from

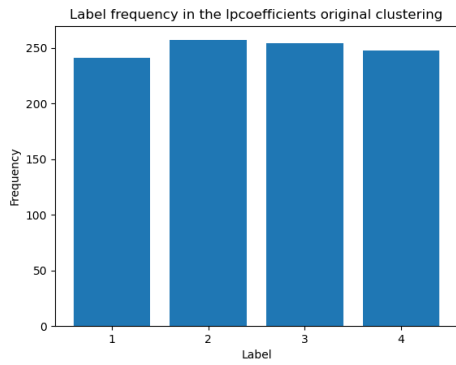
4.2.3 HDBSCAN

A configuration that outputs 4 clusters is searched.

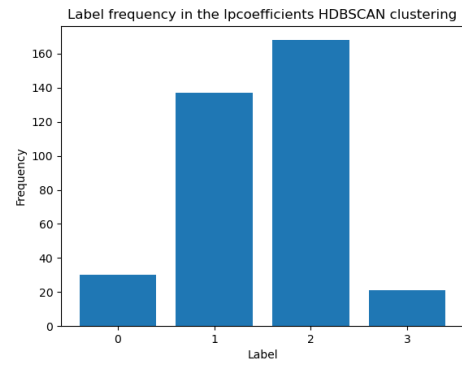
	Minimum cluster size
Original LP coefficients	21
PCA LP coefficients	21
UMAP LP coefficients	25

Table 8: HDBSCAN hyperparameter configuration for LP coefficients clustering

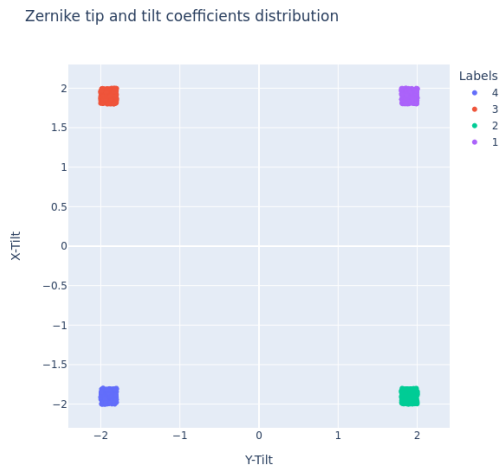
The results are the following:



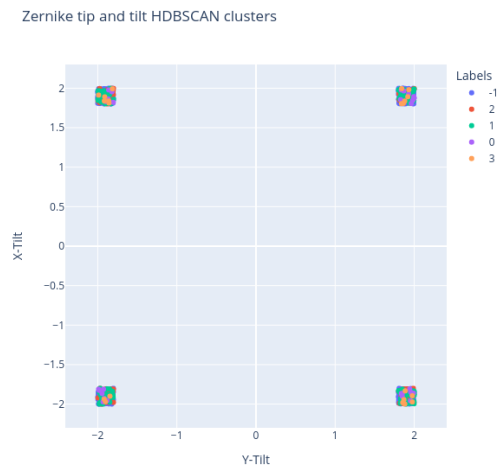
(a) Original cluster densities



(b) HDBSCAN clusters densities

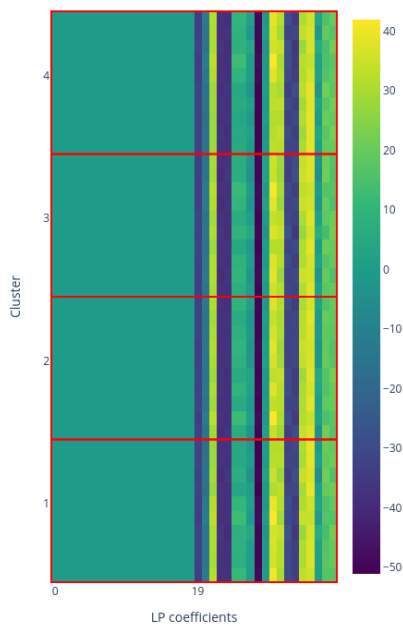


(c) Original clusters



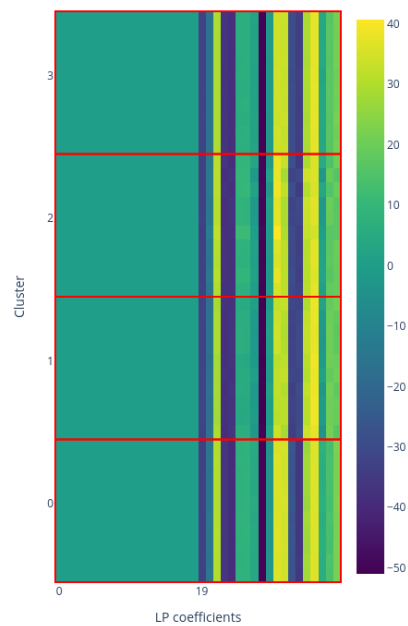
(d) HDBSCAN clusters

Original LP coefficients cluster samples



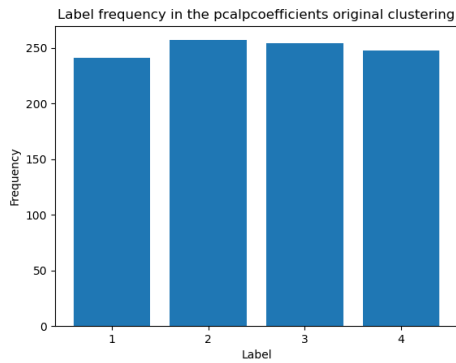
(e) Original cluster samples

HDBSCAN LP coefficients cluster samples

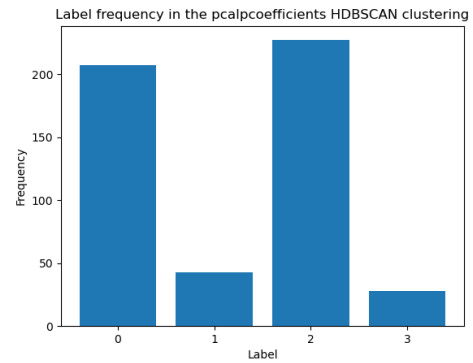


(f) HDBSCAN cluster samples

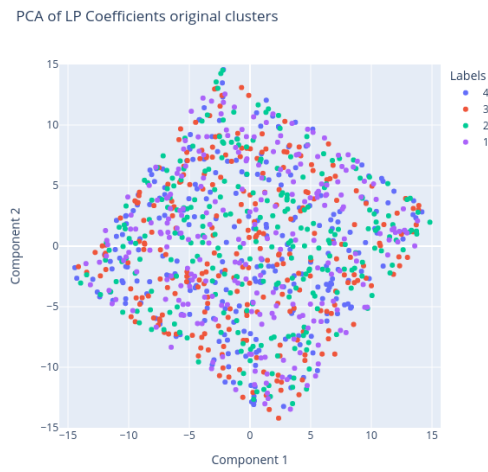
Figure 14: Comparison between original clustering and HDBSCAN clustering



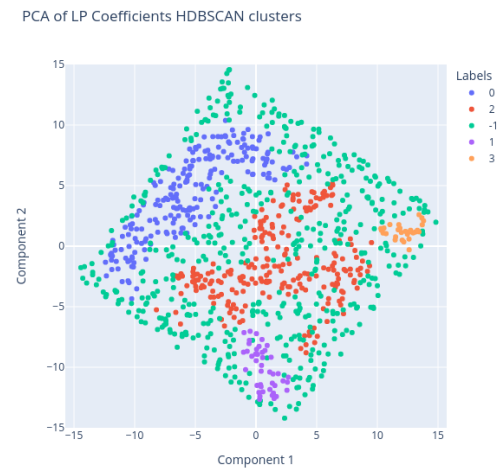
(a) Original cluster densities from PCA



(b) HDBSCAN clusters densities from PCA

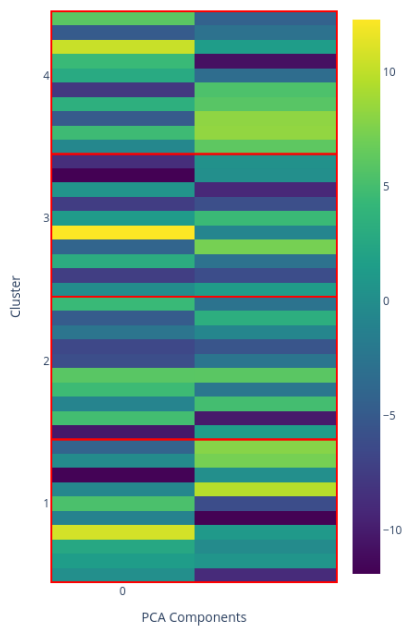


(c) Original clusters from PCA



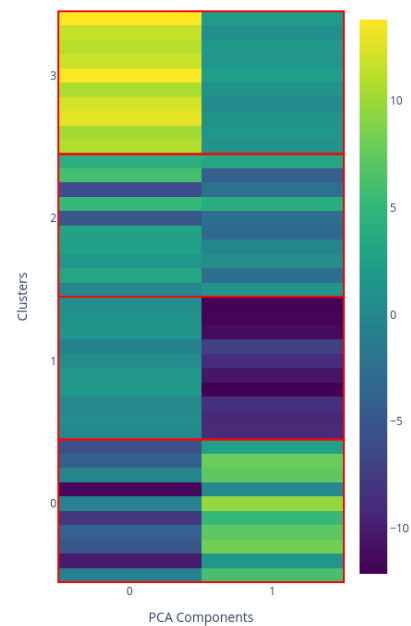
(d) HDBSCAN clusters from PCA

Original PCA LP coefficients cluster samples



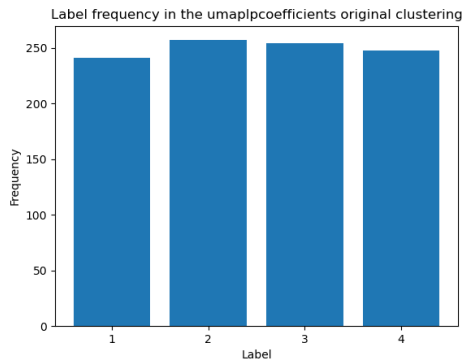
(e) Original cluster samples from PCA

HDBSCAN PCA LP coefficients cluster samples

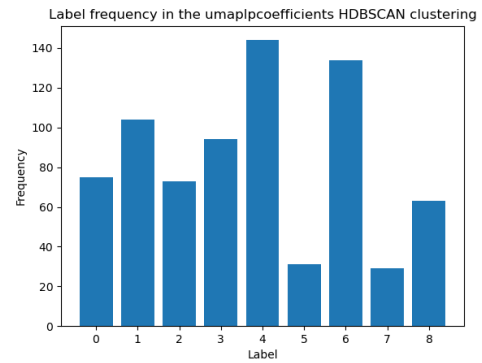


(f) HDBSCAN cluster samples from PCA

Figure 15: Comparison between original clustering and HDBSCAN clustering from

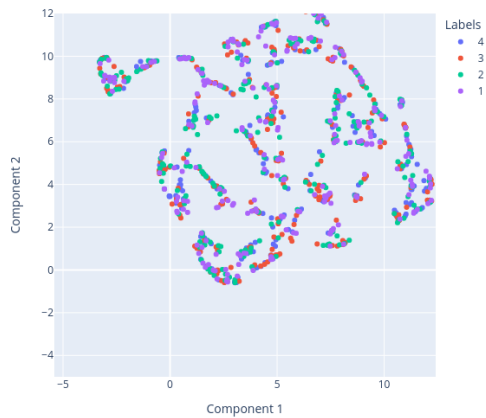


(a) Original cluster densities from UMAP



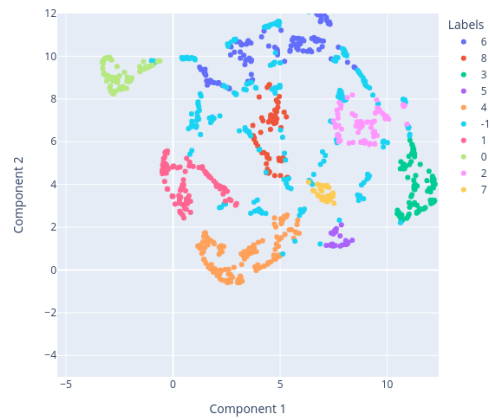
(b) HDBSCAN clusters densities from UMAP

UMAP of LP Coefficients original clusters



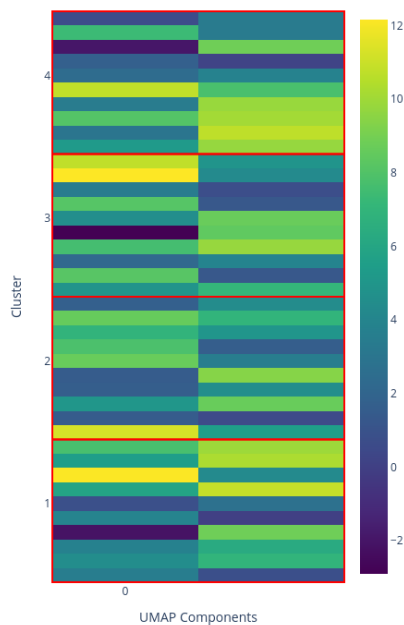
(c) Original clusters from UMAP

UMAP of LP Coefficients HDBSCAN clusters



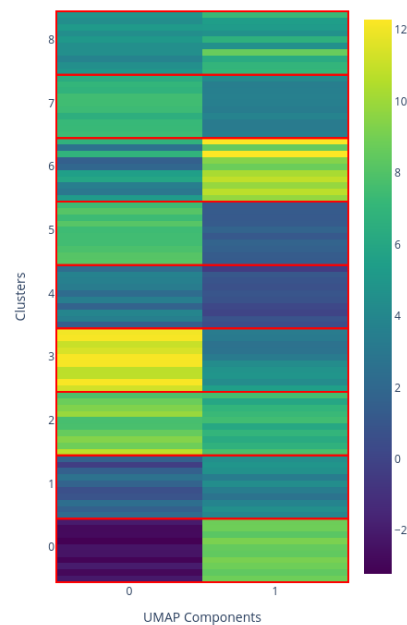
(d) HDBSCAN clusters from UMAP

Original UMAP LP coefficients cluster samples



(e) Original cluster samples from UMAP

HDBSCAN UMAP LP coefficients cluster samples



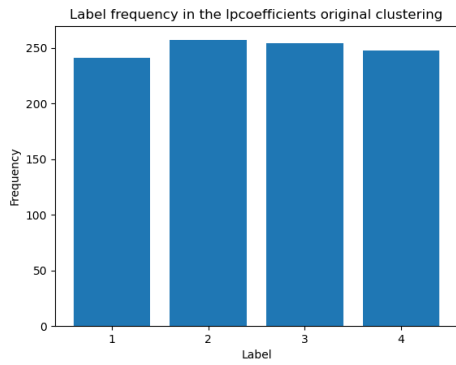
(f) HDBSCAN cluster samples from UMAP

4.2.4 Agglomerative clustering

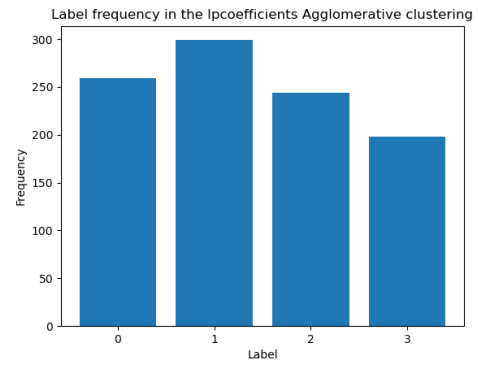
	Number of clusters
Original LP coefficients	4
PCA LP coefficients	4
UMAP LP coefficients	4

Table 9: Agglomerative hyperparameter configuration for LP coefficients clustering

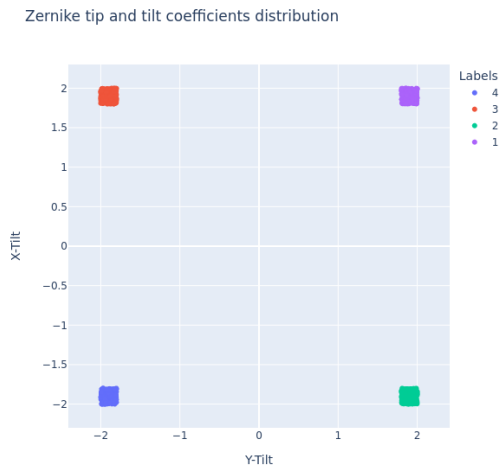
The results are the following:



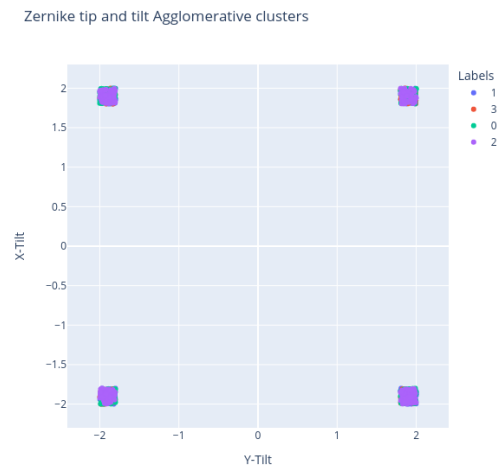
(a) Original cluster densities



(b) Agglomerative clusters densities

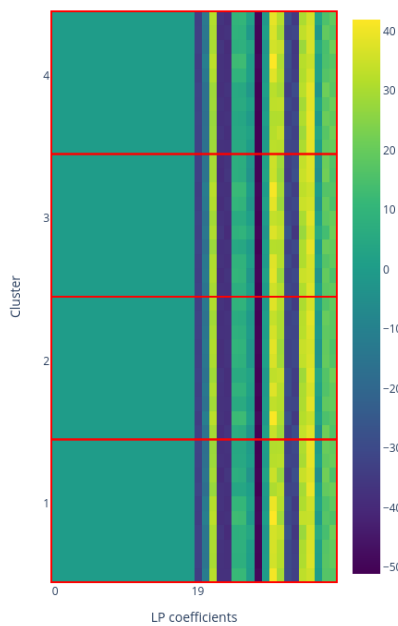


(c) Original clusters



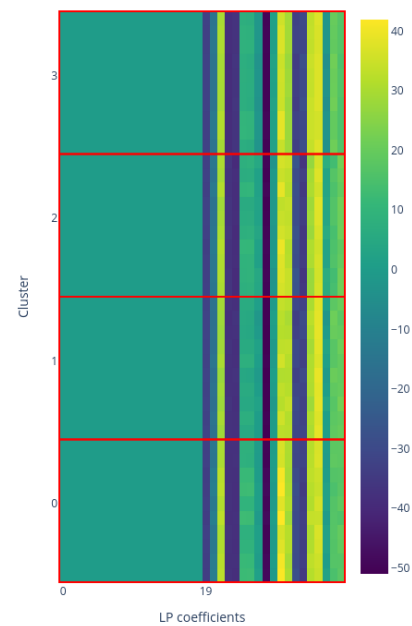
(d) Agglomerative clusters

Original LP coefficients cluster samples



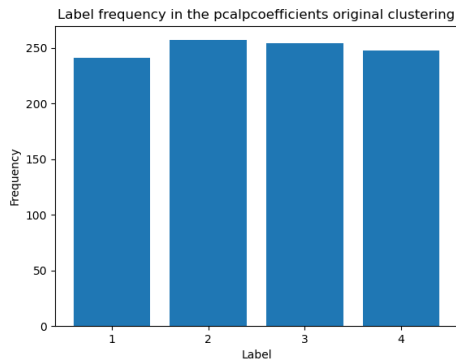
(e) Original cluster samples

Agglomerative LP coefficients cluster samples

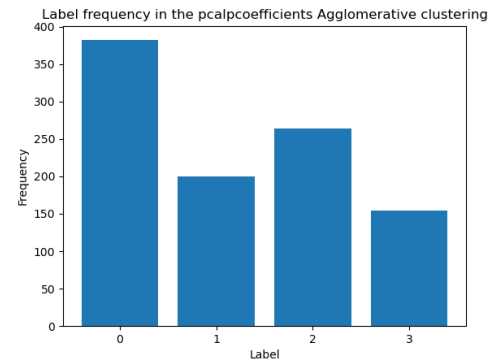


(f) Agglomerative cluster samples

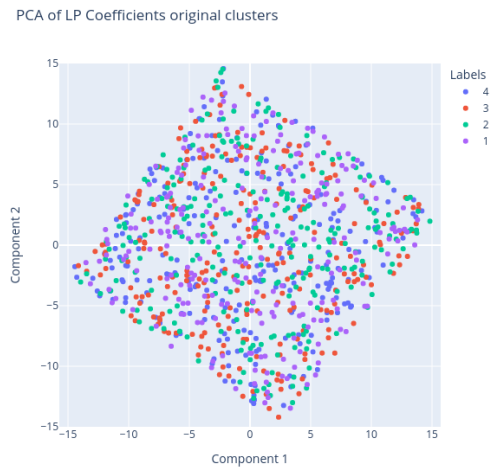
Figure 17: Comparison between original clustering and Agglomerative clustering



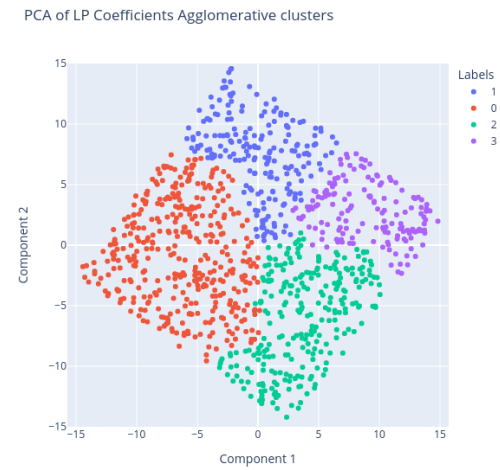
(a) Original cluster densities from PCA



(b) Agglomerative clusters densities from PCA

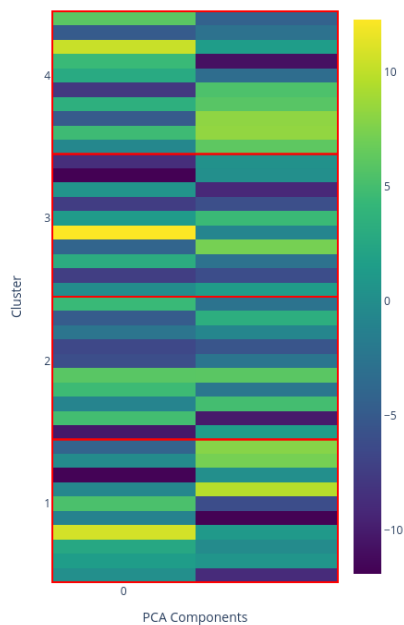


(c) Original clusters from PCA



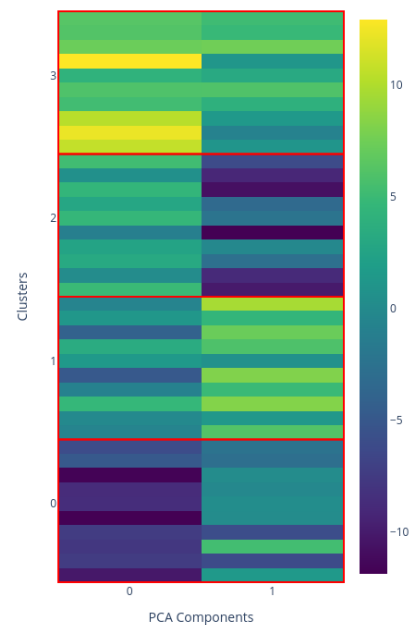
(d) Agglomerative clusters from PCA

Original PCA LP coefficients cluster samples

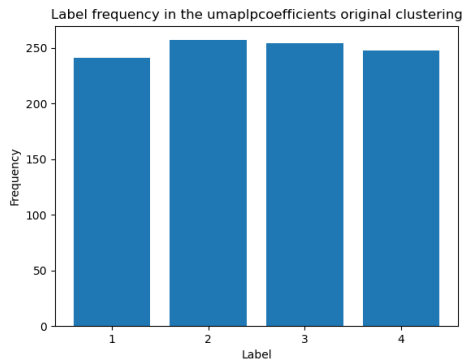


(e) Original cluster samples from PCA

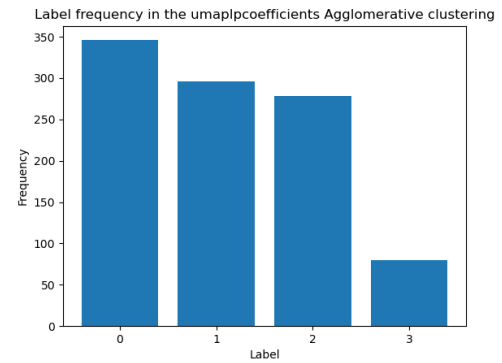
Agglomerative PCA LP coefficients cluster samples



(f) Agglomerative cluster samples from PCA

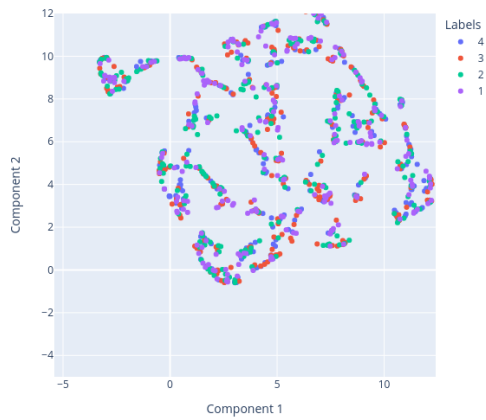


(a) Original cluster densities from UMAP



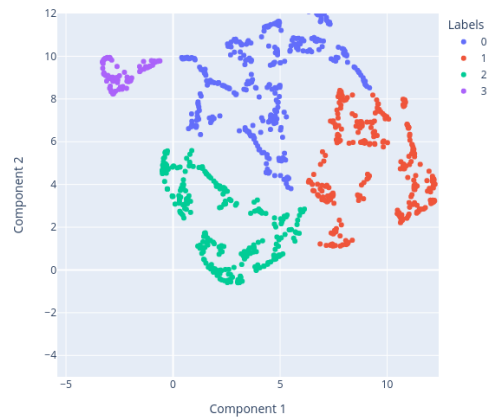
(b) Agglomerative clusters densities from UMAP

UMAP of LP Coefficients original clusters



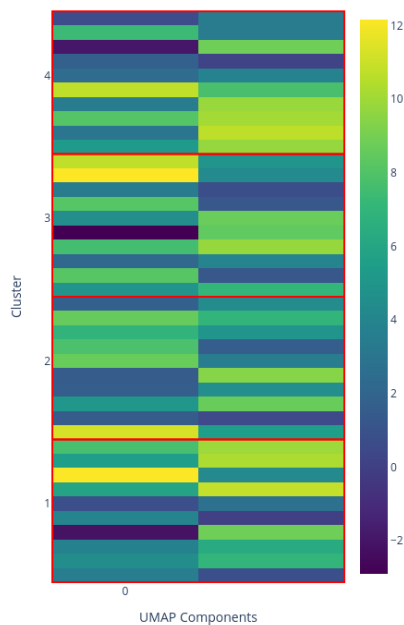
(c) Original clusters from UMAP

UMAP of LP Coefficients Agglomerative clusters



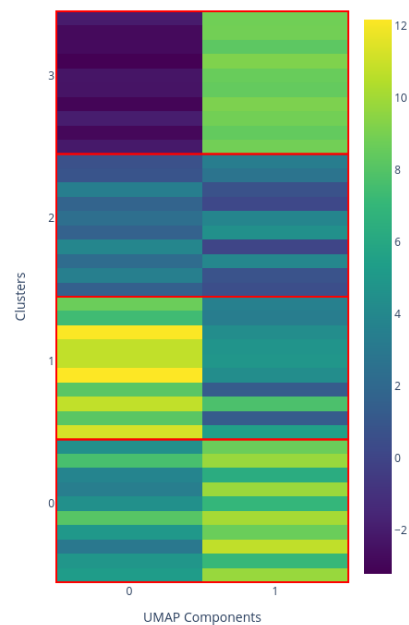
(d) Agglomerative clusters from UMAP

Original UMAP LP coefficients cluster samples



(e) Original cluster samples from UMAP

Agglomerative UMAP LP coefficients cluster samples



(f) Agglomerative cluster samples from UMAP

4.2.5 Summary

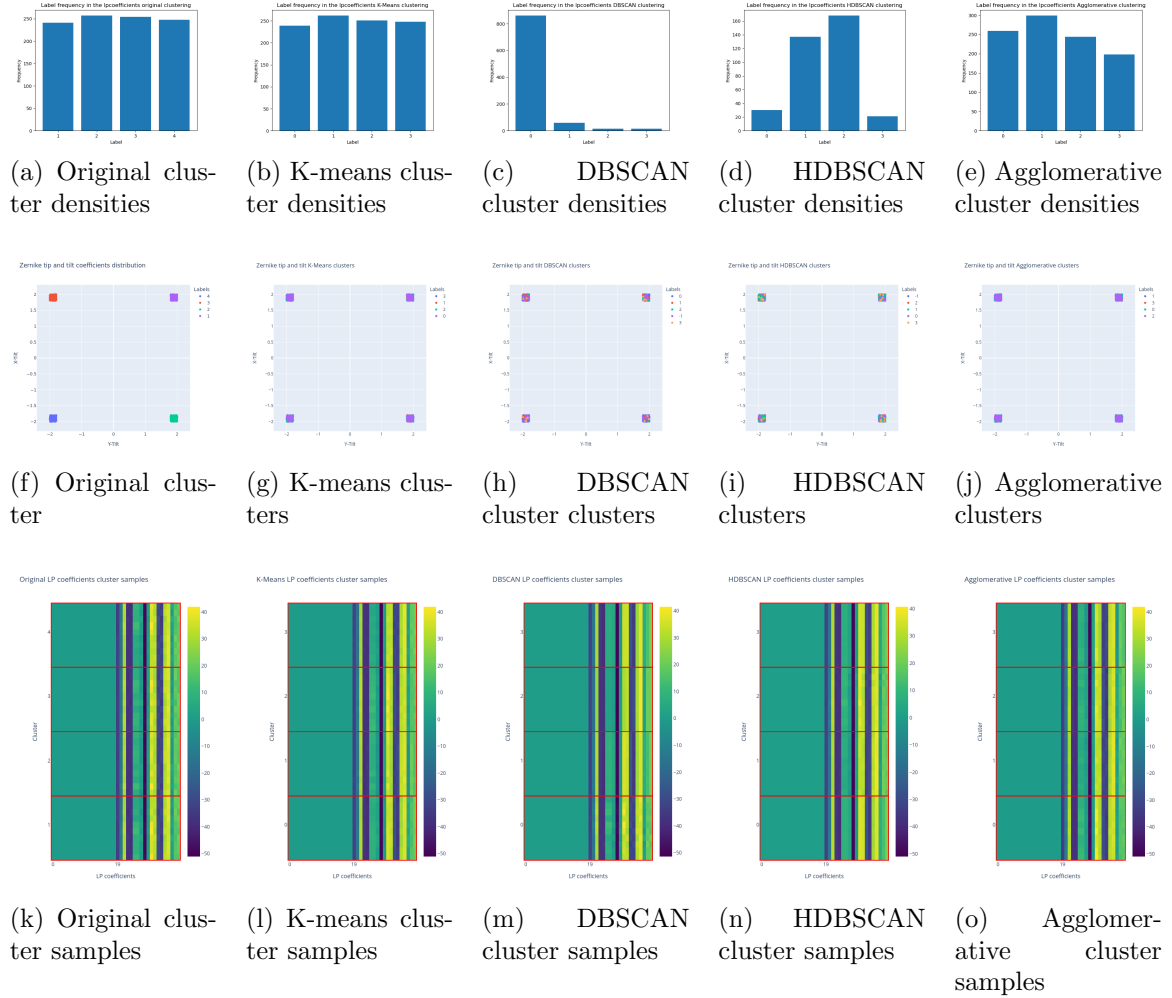


Figure 20: Comparison between clustering LP coefficients algorithms

	Original	K-Means	DBSCAN	HDBSCAN	Agglomerative
Original	\	0.002	0.008	0.004	0.002
K-Means		\	0.154	0.143	0.695
DBSCAN			\	0.175	0.142
HDBSCAN				\	0.002

Table 10: Normalized Mutual Information between original LP coefficients clusters

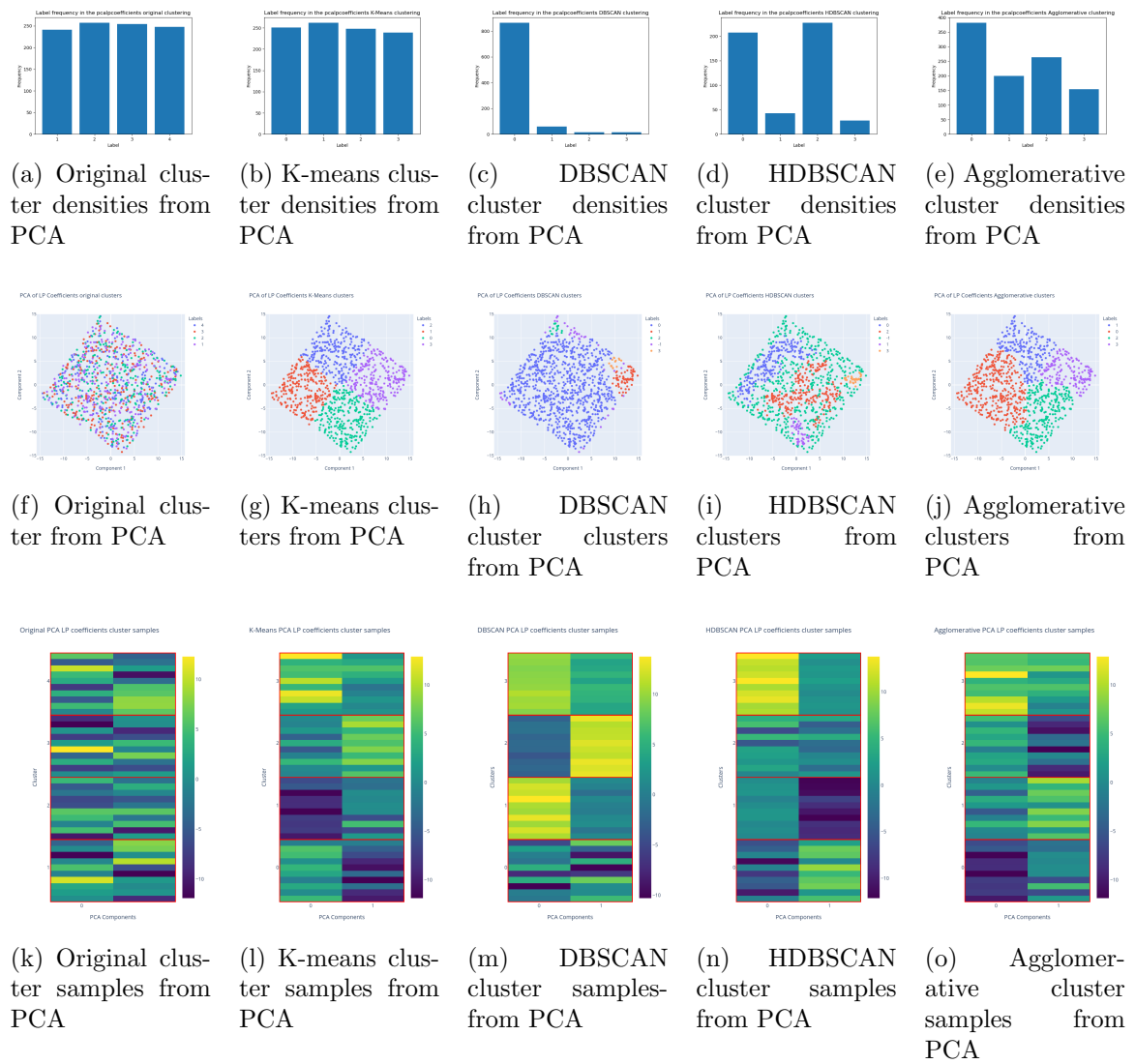


Figure 21: Comparison between clustering PCA LP coefficients algorithms

	Original	K-Means	DBSCAN	HDBSCAN	Agglomerative
Original	\	0.002	0.003	0.004	0.003
K-Means		\	0.150	0.143	0.642
DBSCAN			\	0.174	0.209
HDBSCAN				\	0.003

Table 11: Normalized Mutual Information between PCA LP coefficients clusters

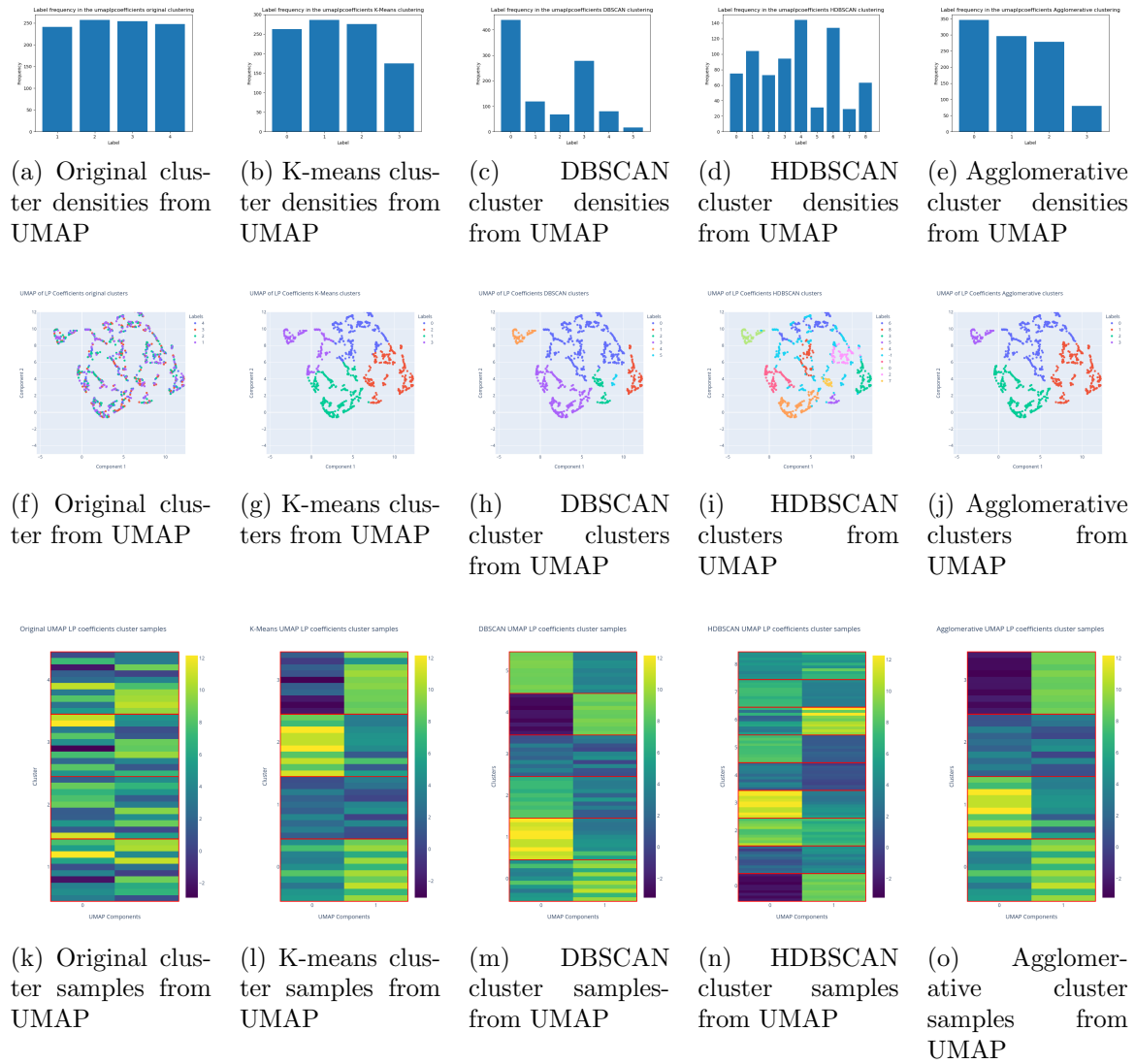


Figure 22: Comparison between clustering UMAP LP coefficients algorithms

	Original	K-Means	DBSCAN	HDBSCAN	Agglomerative
Original	\	0.003	0.001	0.009	0.001
K-Means		\	0.438	0.480	0.667
DBSCAN			\	0.487	0.529
HDBSCAN				\	0.001

Table 12: Normalized Mutual Information between UMAP LP coefficients clusters

4.3 Output fluxes clustering

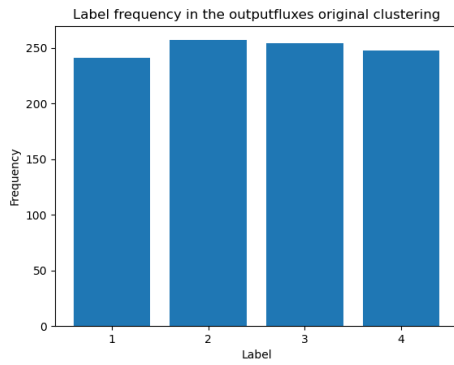
4.3.1 K-Means

As K-Means allows for the number of clusters to be defined, and we know that there are 4 in the original dataset, K-Means is used to find 4 clusters.

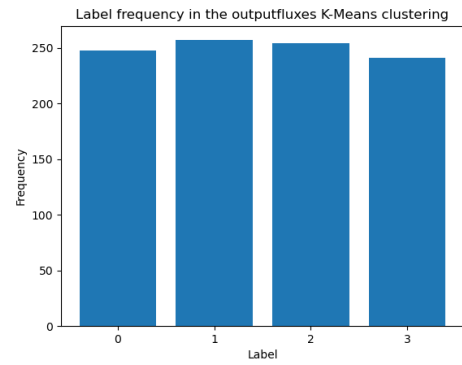
	Number of clusters	Number of initializations
Original Output fluxes	4	10
PCA Output fluxes	4	10
UMAP Output fluxes	4	10

Table 13: K-Means hyperparameter configuration for c coefficients clustering

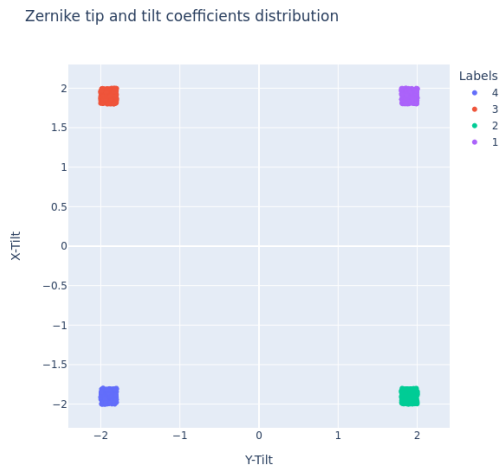
The results are the following:



(a) Original cluster densities



(b) K-Means clusters densities

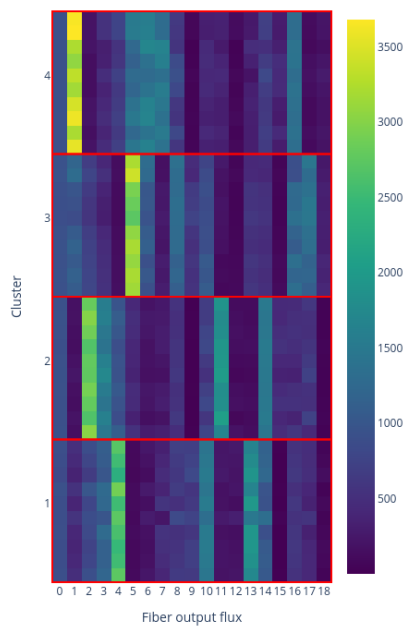


(c) Original clusters



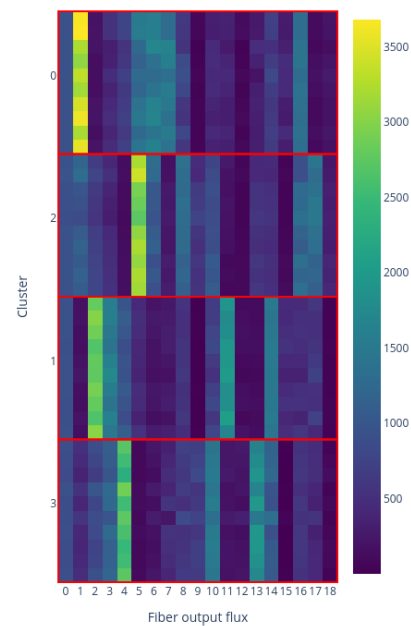
(d) K-Means clusters

Original output fluxes cluster samples



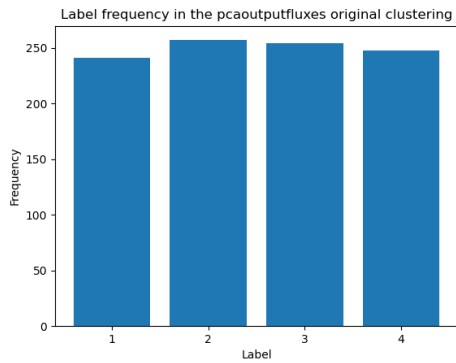
(e) Original cluster samples

K-Means output fluxes cluster samples

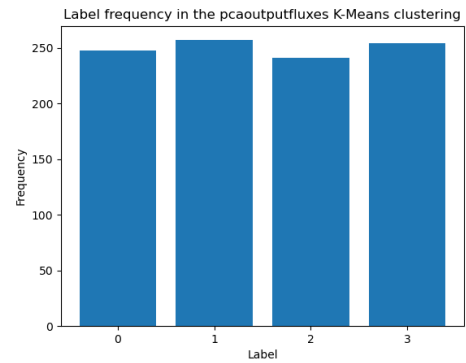


(f) K-Means cluster samples

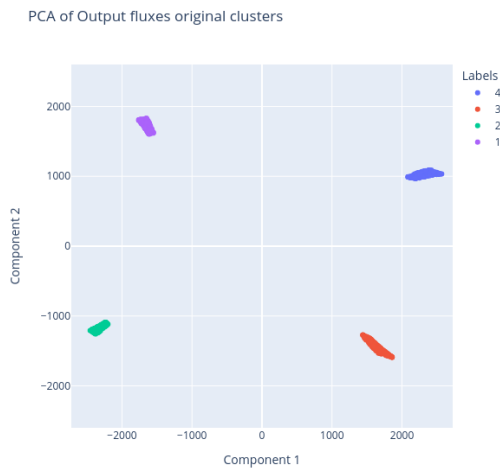
Figure 23: Comparison between original clustering and K-Means clustering from original Output fluxes



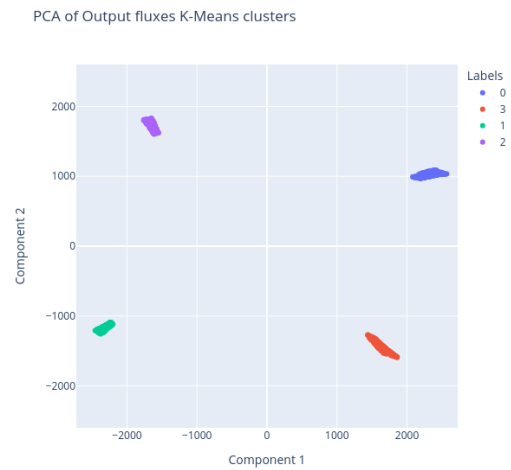
(a) Original cluster densities from PCA



(b) K-Means clusters densities from PCA

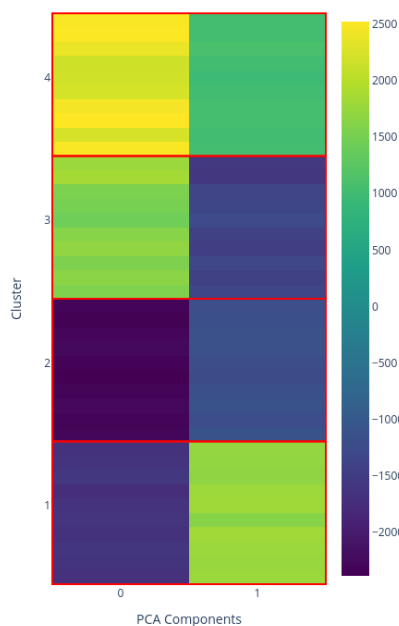


(c) Original clusters



(d) K-Means clusters from PCA

Original PCA Output fluxes cluster samples



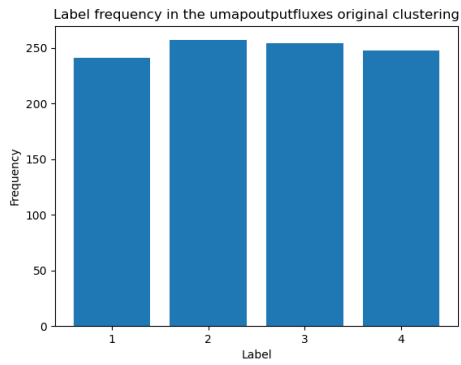
(e) Original cluster samples from PCA

K-Means PCA Output fluxes cluster samples

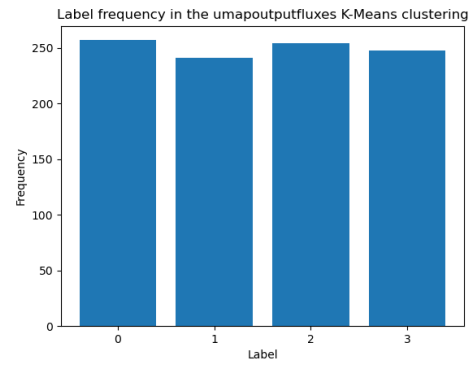


(f) K-Means cluster samples from PCA

Figure 24: Comparison between original clustering and K-Means clustering from PCA of Output fluxes

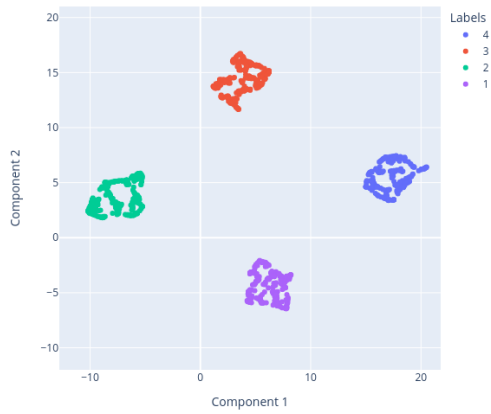


(a) Original cluster densities from UMAP



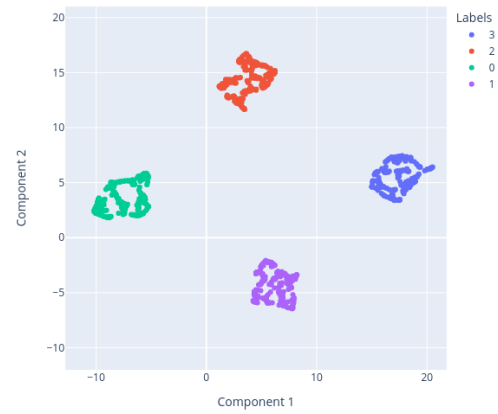
(b) K-Means clusters densities from UMAP

UMAP of Output fluxes original clusters



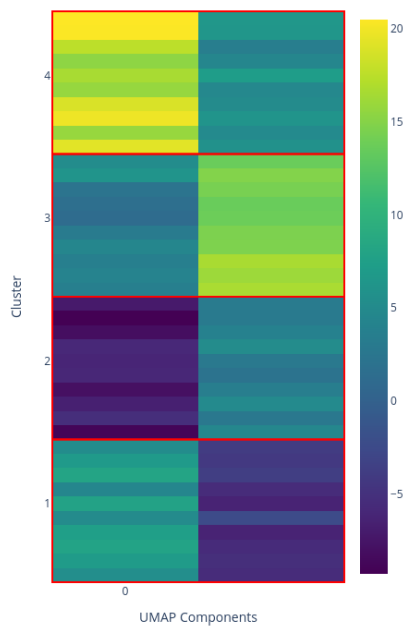
(c) Original clusters from UMAP

UMAP of Output fluxes K-Means clusters



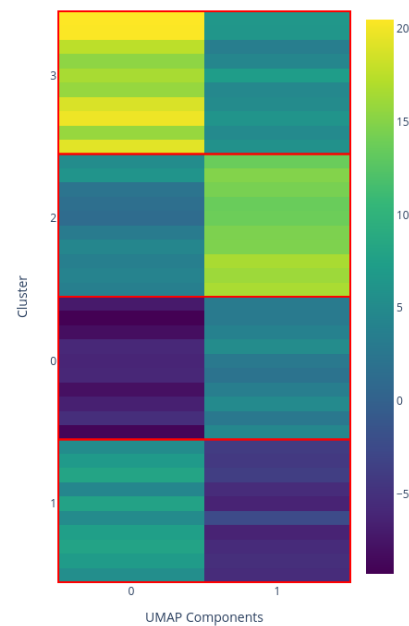
(d) K-Means clusters from UMAP

Original UMAP Output fluxes cluster samples



(e) Original cluster samples from UMAP

K-Means UMAP Output fluxes cluster samples



(f) K-Means cluster samples from UMAP

Figure 25: Comparison between original clustering and K-Means clustering from

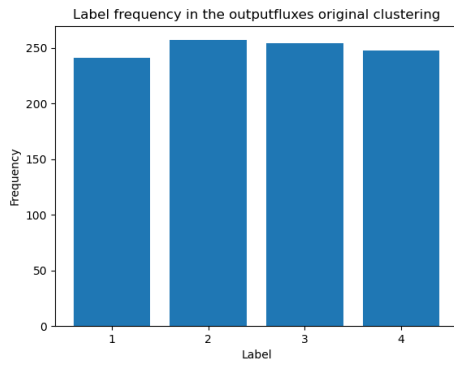
4.3.2 DBSCAN

A configuration that outputs 4 clusters is searched

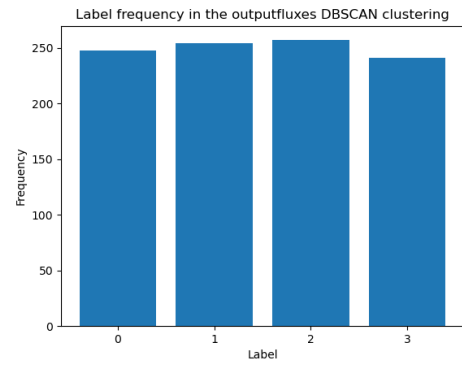
	Number of neighbours	Epsilon
Original Output fluxes	7	120
PCA Output fluxes	15	40
UMAP Output fluxes	10	0.85

Table 14: DBSCAN hyperparameter configuration for Output fluxes clustering

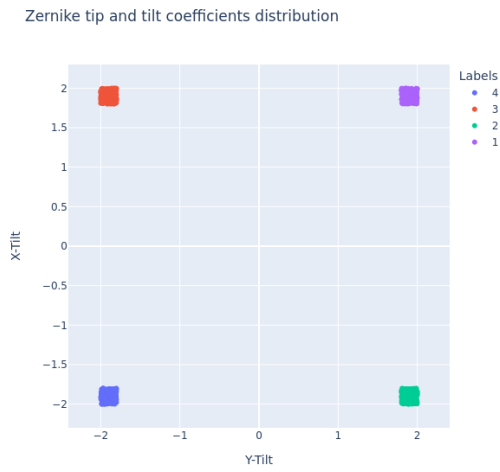
The results are the following:



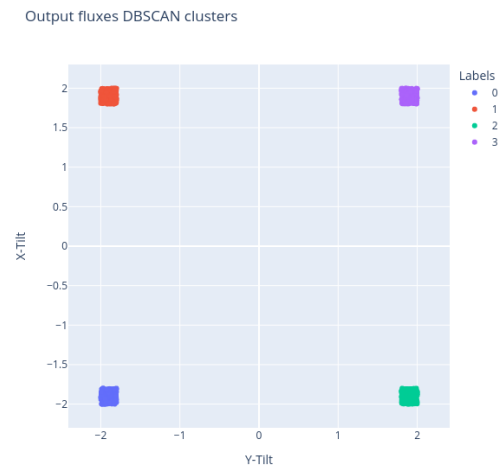
(a) Original cluster densities



(b) DBSCAN clusters densities

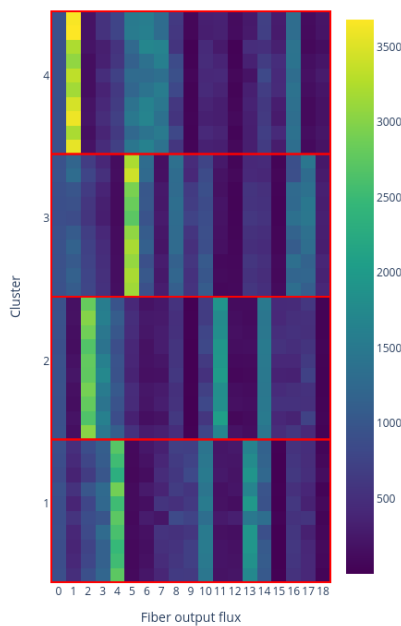


(c) Original clusters



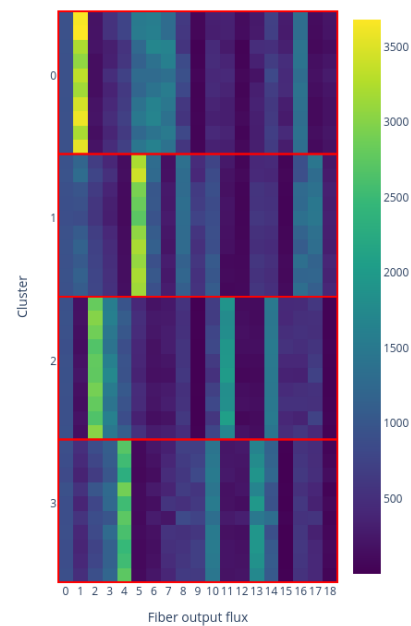
(d) DBSCAN clusters

Original output fluxes cluster samples



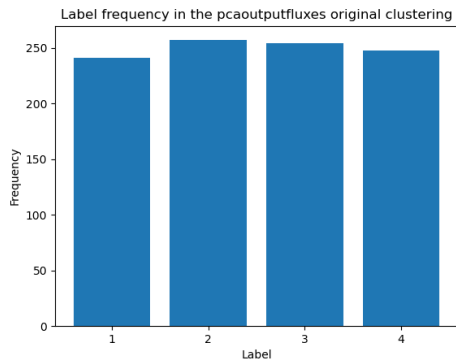
(e) Original cluster samples

DBSCAN output fluxes cluster samples

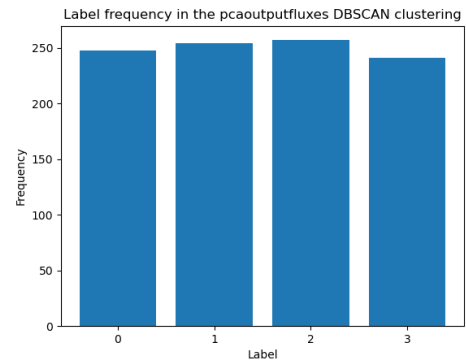


(f) DBSCAN cluster samples

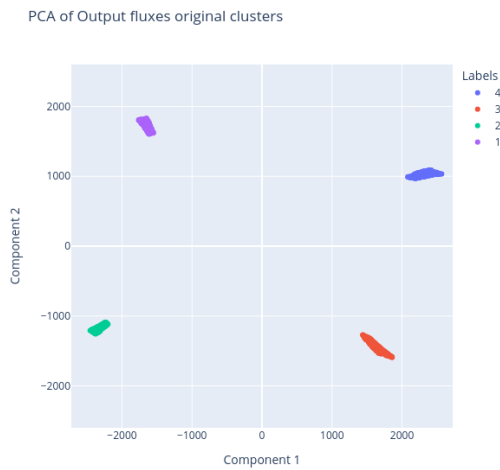
Figure 26: Comparison between original clustering and DBSCAN clustering



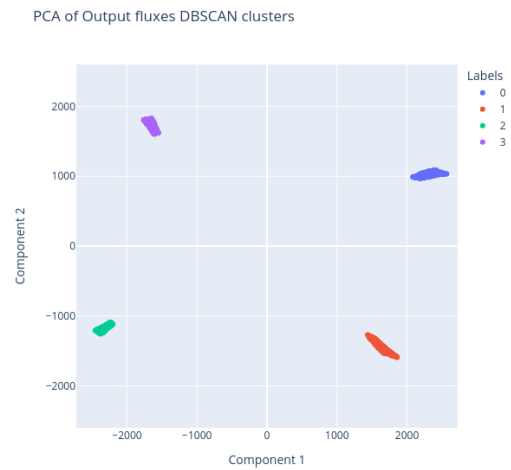
(a) Original cluster densities from PCA



(b) DBSCAN clusters densities from PCA

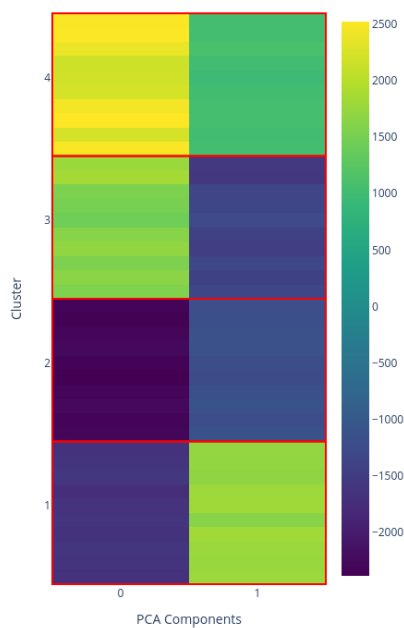


(c) Original clusters from PCA



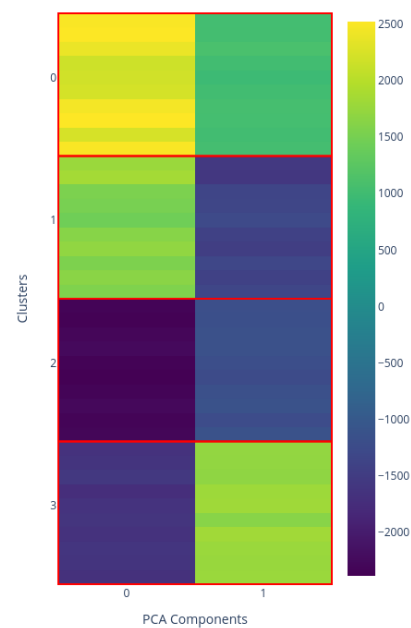
(d) DBSCAN clusters from PCA

Original PCA Output fluxes cluster samples



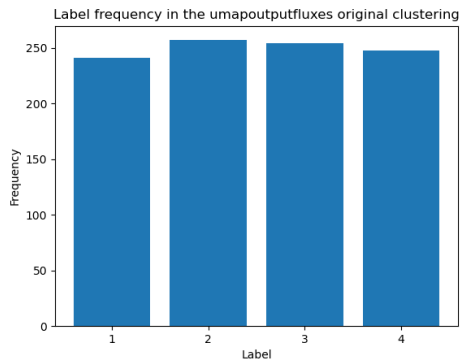
(e) Original cluster samples from PCA

DBSCAN PCA Output fluxes cluster samples

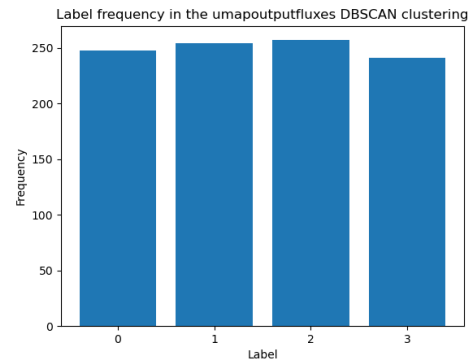


(f) DBSCAN cluster samples from PCA

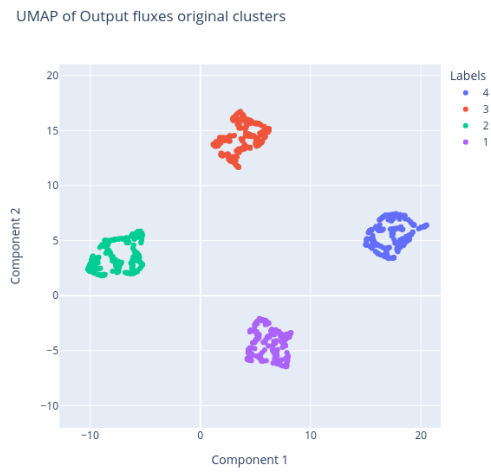
Figure 27: Comparison between original clustering and DBSCAN clustering from PCA of Output fluxes



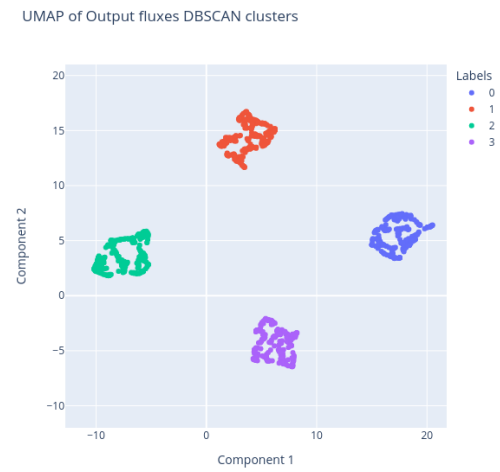
(a) Original cluster densities from UMAP



(b) DBSCAN clusters densities from UMAP

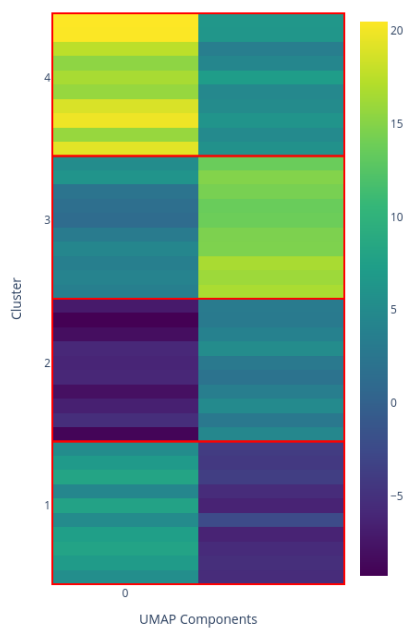


(c) Original clusters from UMAP



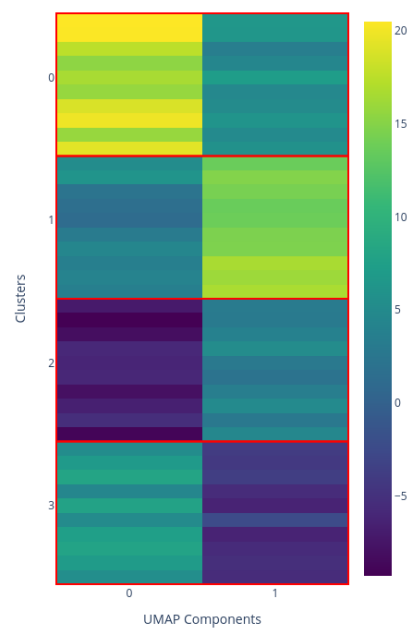
(d) DBSCAN clusters from UMAP

Original UMAP Output fluxes cluster samples



(e) Original cluster samples from UMAP

DBSCAN UMAP LP coefficients cluster samples



(f) DBSCAN cluster samples from UMAP

Figure 28: Comparison between original clustering and DBSCAN clustering from

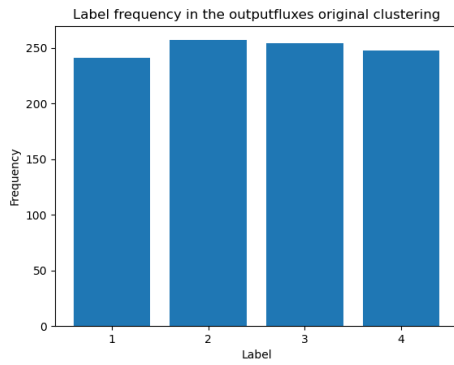
4.3.3 HDBSCAN

A configuration that outputs 4 clusters is searched.

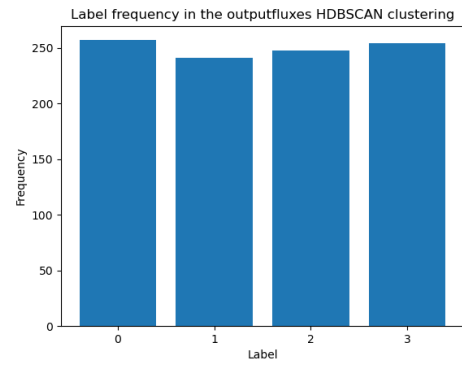
	Minimum cluster size
Original Output fluxes	21
PCA Output fluxes	21
HDBSCAN Output fluxes	25

Table 15: HDBSCAN hyperparameter configuration for Output fluxes clustering

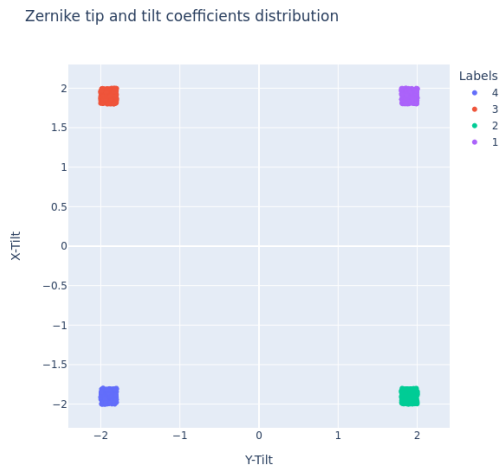
The results are the following:



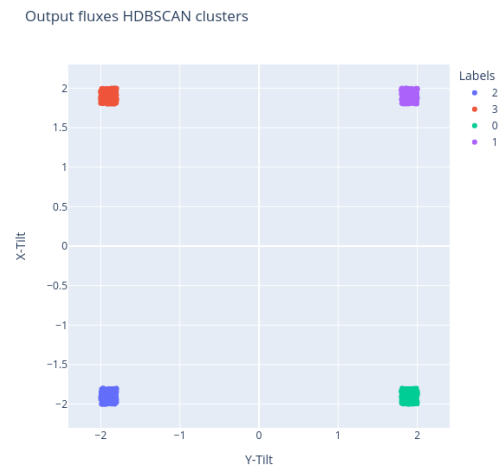
(a) Original cluster densities



(b) HDBSCAN clusters densities

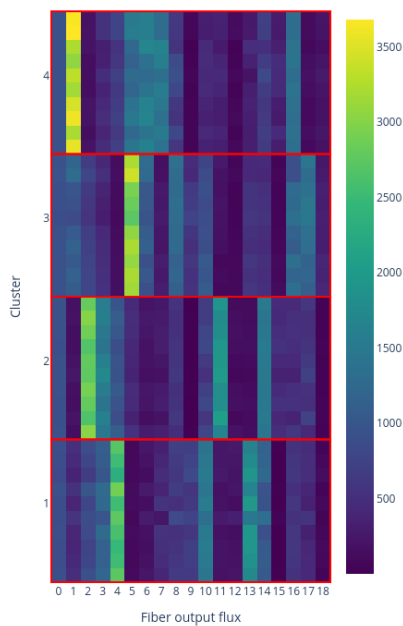


(c) Original clusters



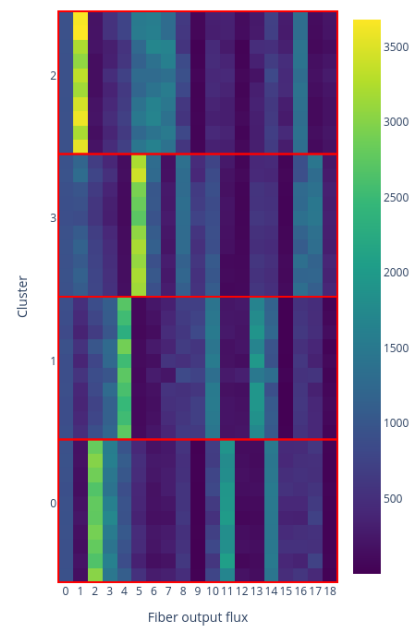
(d) HDBSCAN clusters

Original output fluxes cluster samples



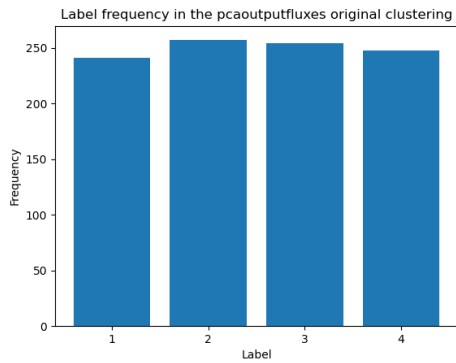
(e) Original cluster samples

HDBSCAN output fluxes cluster samples

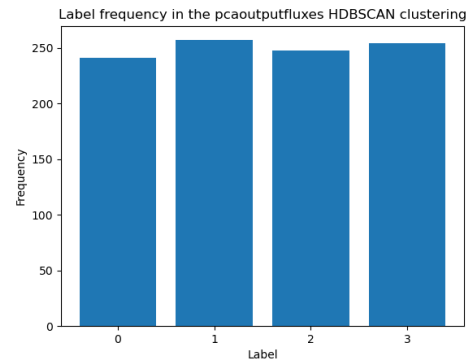


(f) HDBSCAN cluster samples

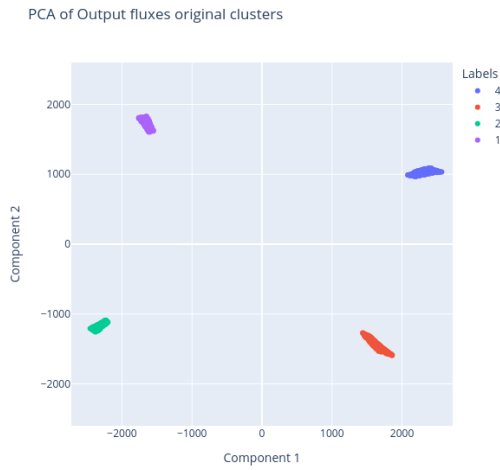
Figure 29: Comparison between original clustering and HDBSCAN clustering



(a) Original cluster densities from PCA



(b) HDBSCAN clusters densities from PCA

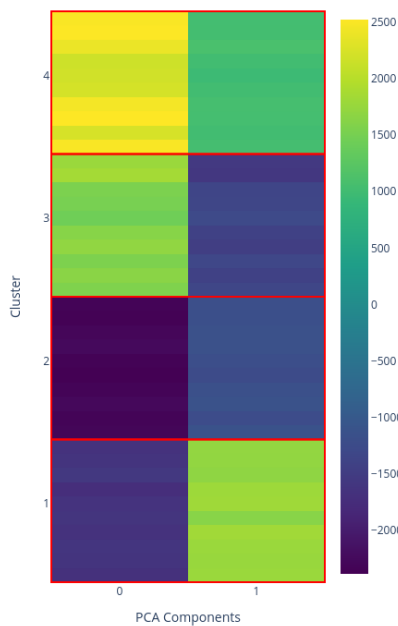


(c) Original clusters from PCA



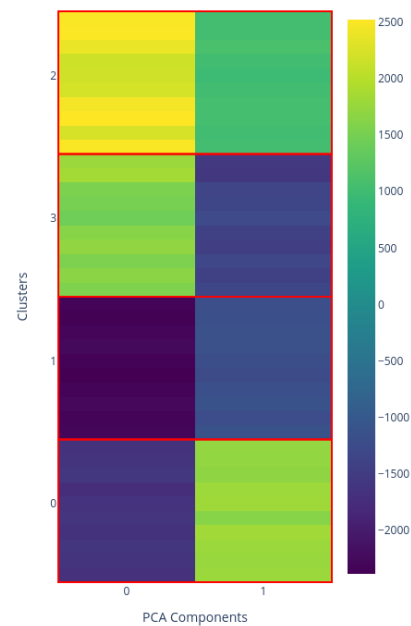
(d) HDBSCAN clusters from PCA

Original PCA Output fluxes cluster samples



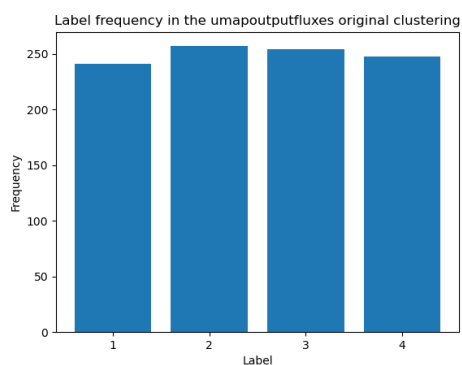
(e) Original cluster samples from PCA

HDBSCAN PCA Output fluxes cluster samples

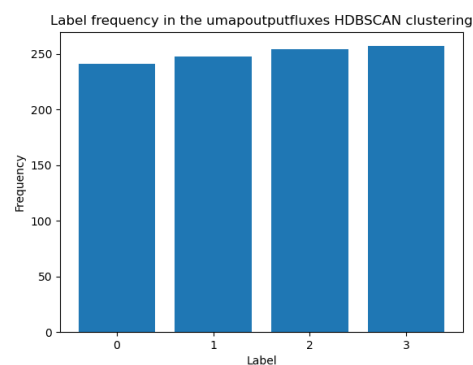


(f) HDBSCAN cluster samples from PCA

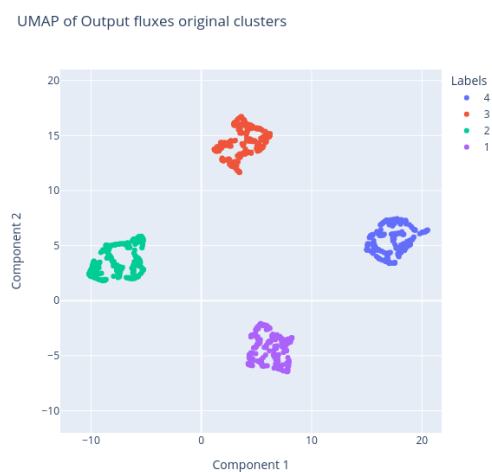
Figure 30: Comparison between original clustering and HDBSCAN clustering from



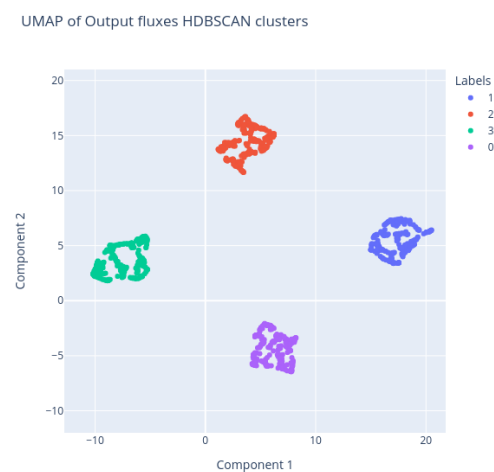
(a) Original cluster densities from UMAP



(b) HDBSCAN clusters densities from UMAP

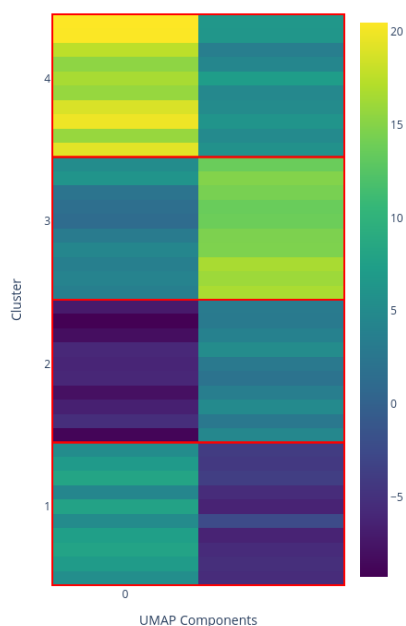


(c) Original clusters from UMAP



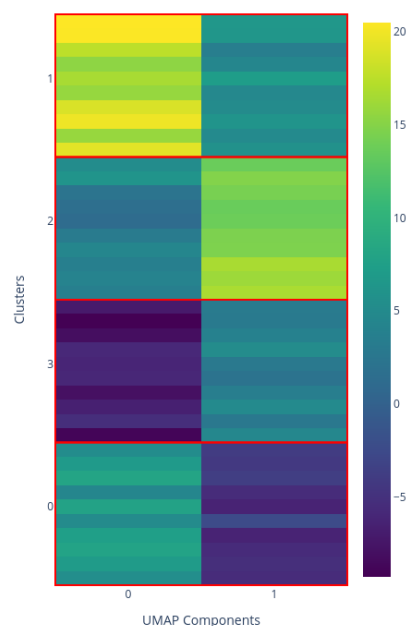
(d) HDBSCAN clusters from UMAP

Original UMAP Output fluxes cluster samples



(e) Original cluster samples from UMAP

HDBSCAN UMAP Output fluxes cluster samples



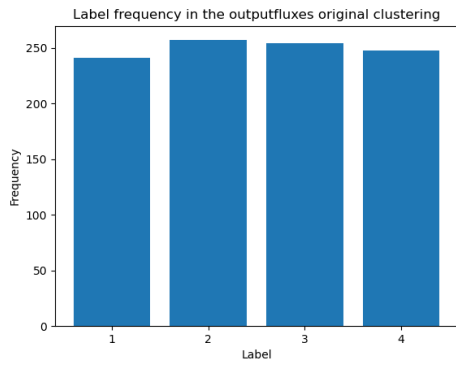
(f) HDBSCAN cluster samples from UMAP

4.3.4 Agglomerative clustering

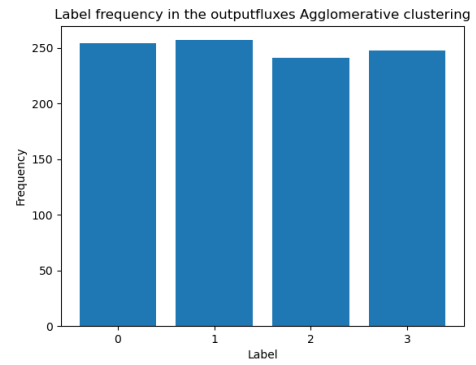
	Number of clusters
Original Output fluxes	4
PCA Output fluxes	4
UMAP Output fluxes	4

Table 16: Agglomerative hyperparameter configuration for Output fluxes clustering

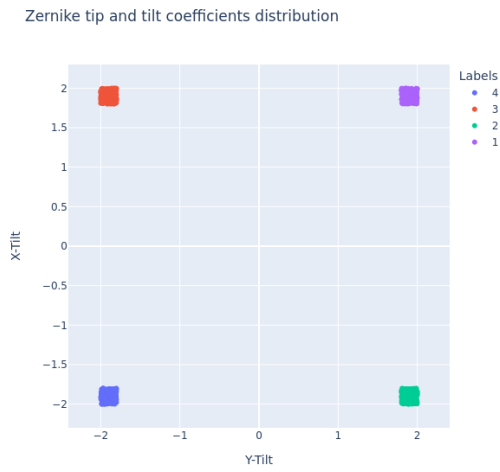
The results are the following:



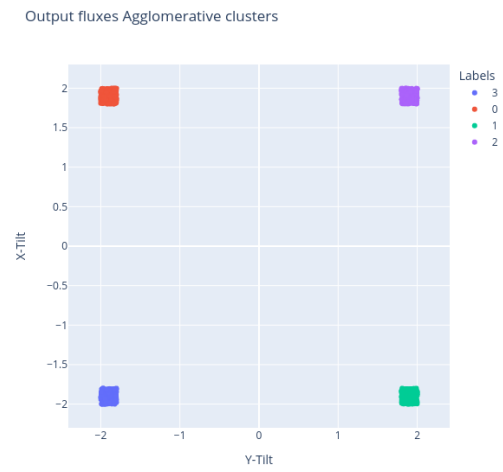
(a) Original cluster densities



(b) Agglomerative clusters densities

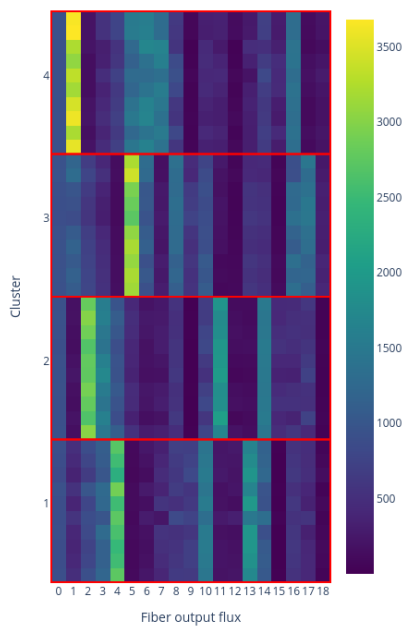


(c) Original clusters



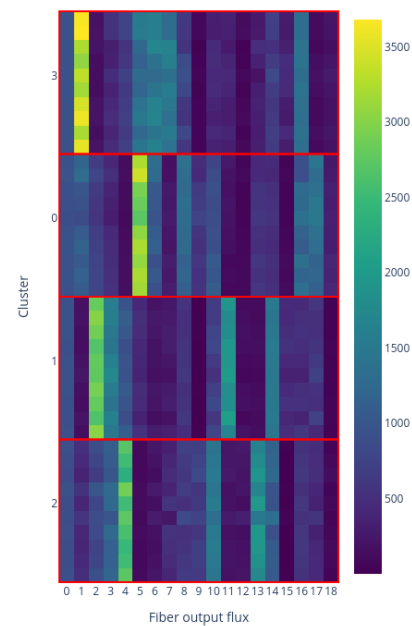
(d) Agglomerative clusters

Original output fluxes cluster samples



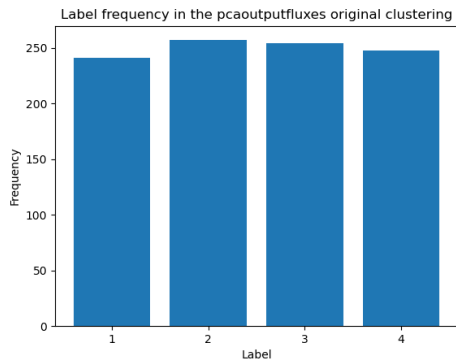
(e) Original cluster samples

Agglomerative output fluxes cluster samples

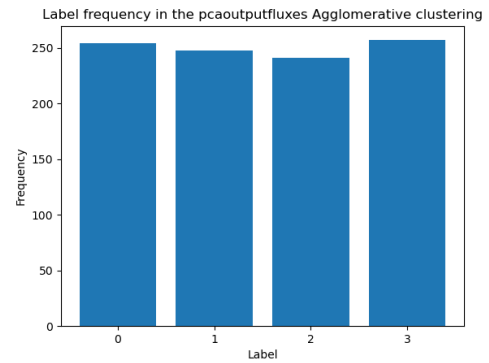


(f) Agglomerative cluster samples

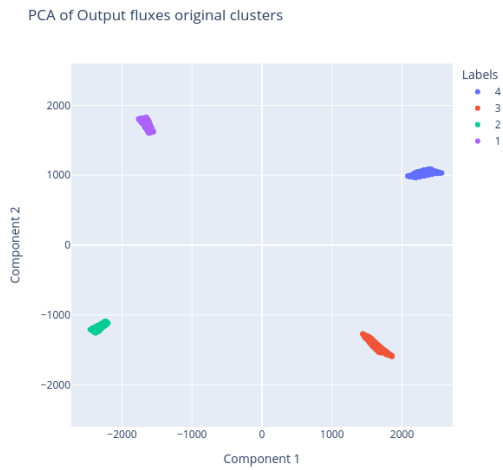
Figure 32: Comparison between original clustering and Agglomerative clustering



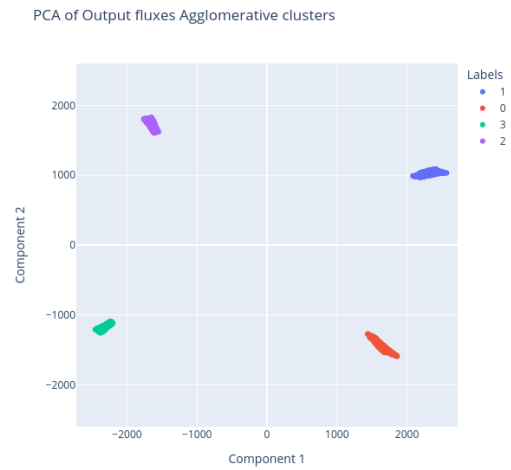
(a) Original cluster densities from PCA



(b) Agglomerative clusters densities from PCA

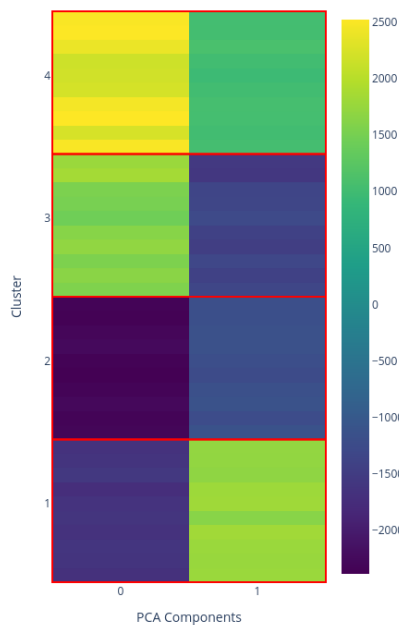


(c) Original clusters from PCA



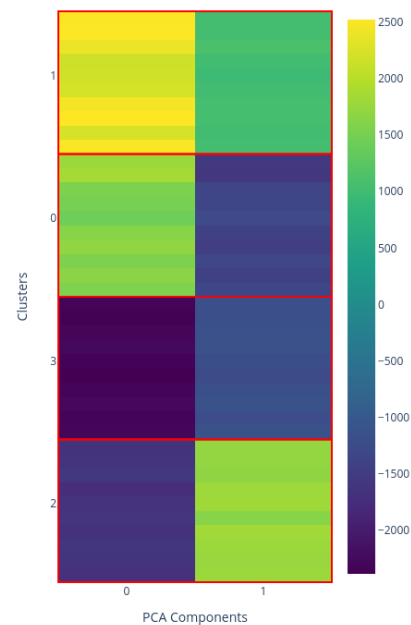
(d) Agglomerative clusters from PCA

Original PCA Output fluxes cluster samples

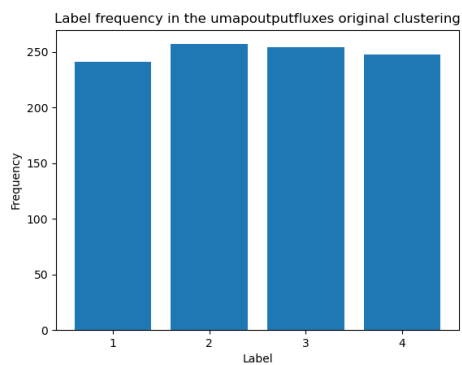


(e) Original cluster samples from PCA

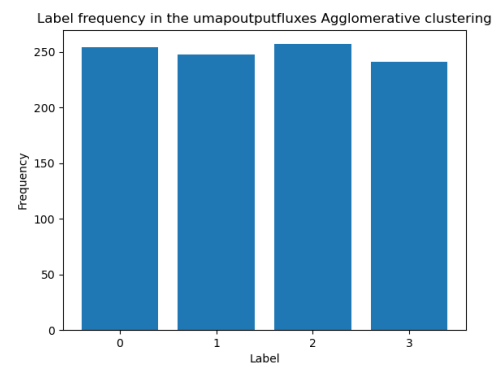
Agglomerative PCA Output fluxes cluster samples



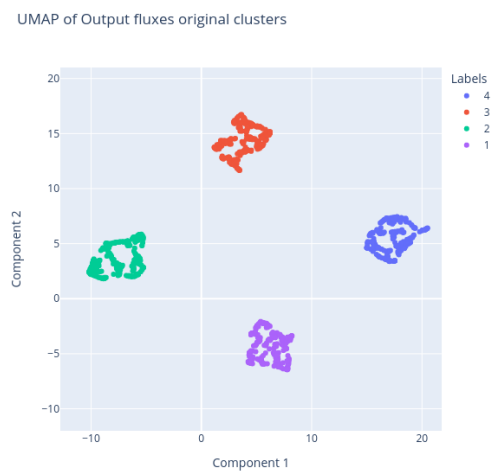
(f) Agglomerative cluster samples from PCA



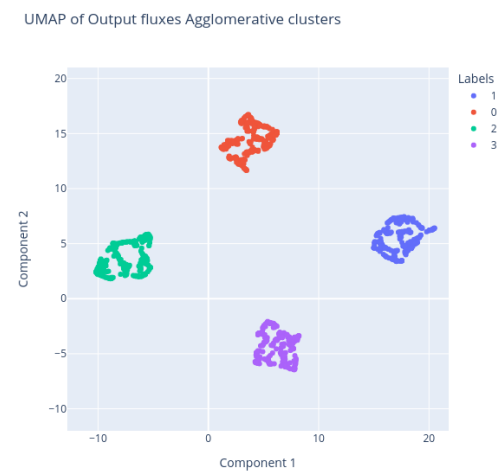
(a) Original cluster densities from UMAP



(b) Agglomerative clusters densities from UMAP

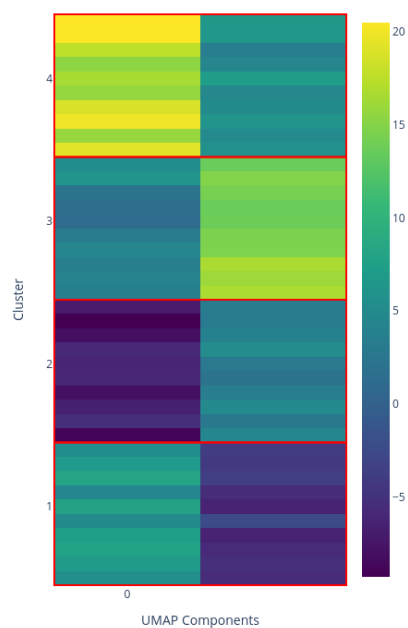


(c) Original clusters from UMAP



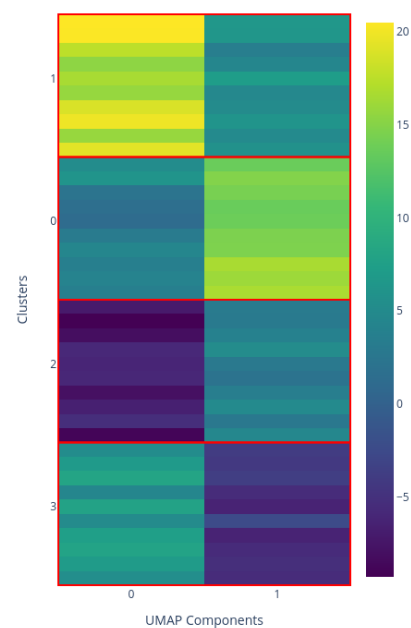
(d) Agglomerative clusters from UMAP

Original UMAP Output fluxes cluster samples



(e) Original cluster samples from UMAP

Agglomerative UMAP Output fluxes cluster samples



(f) Agglomerative cluster samples from UMAP

4.3.5 Summary

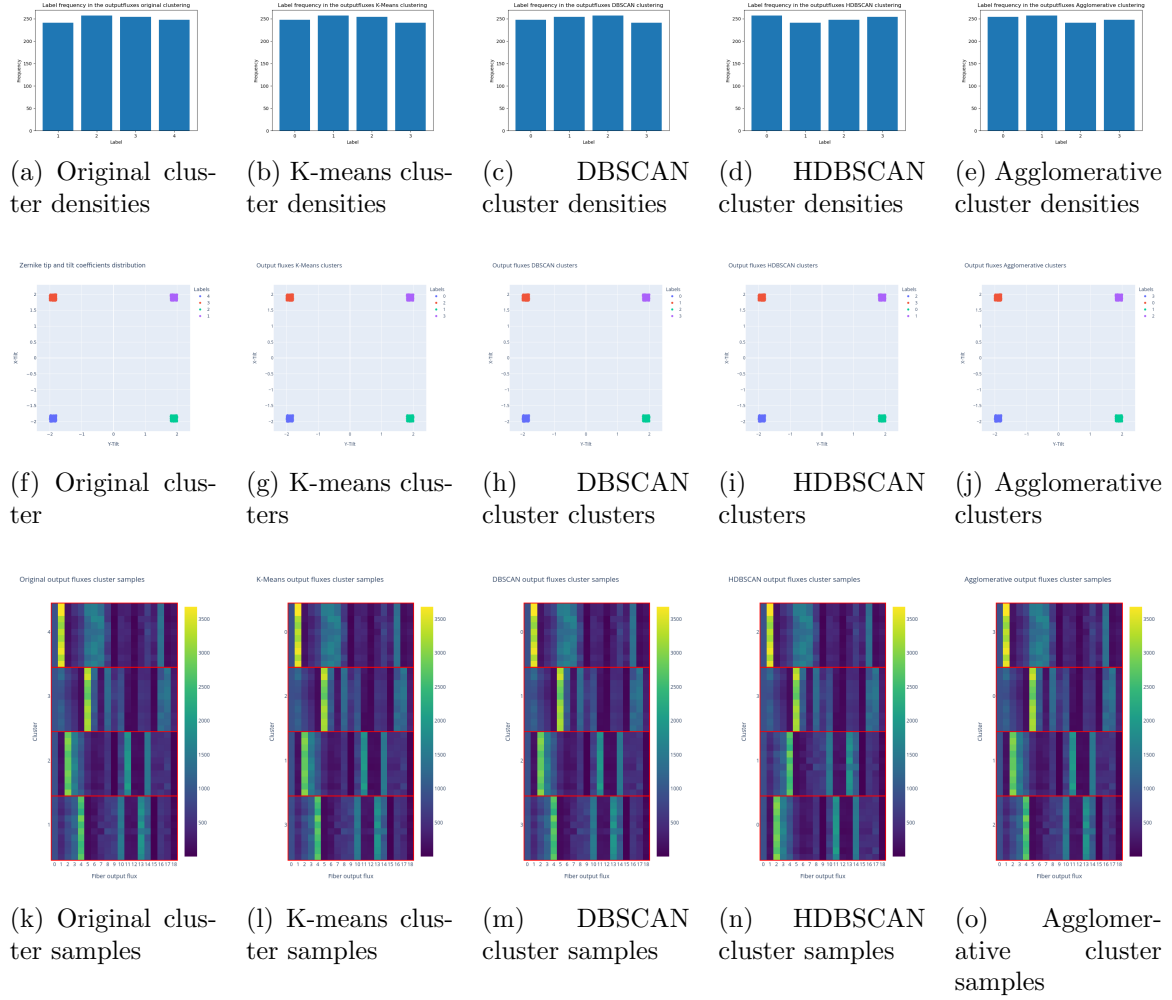


Figure 35: Comparison between clustering Output fluxes algorithms

	Original	K-Means	DBSCAN	HDBSCAN	Agglomerative
Original	\	1	1	1	1
K-Means		\	1	1	1
DBSCAN			\	1	1
HDBSCAN				\	1

Table 17: Normalized Mutual Information between original Output fluxes clusters

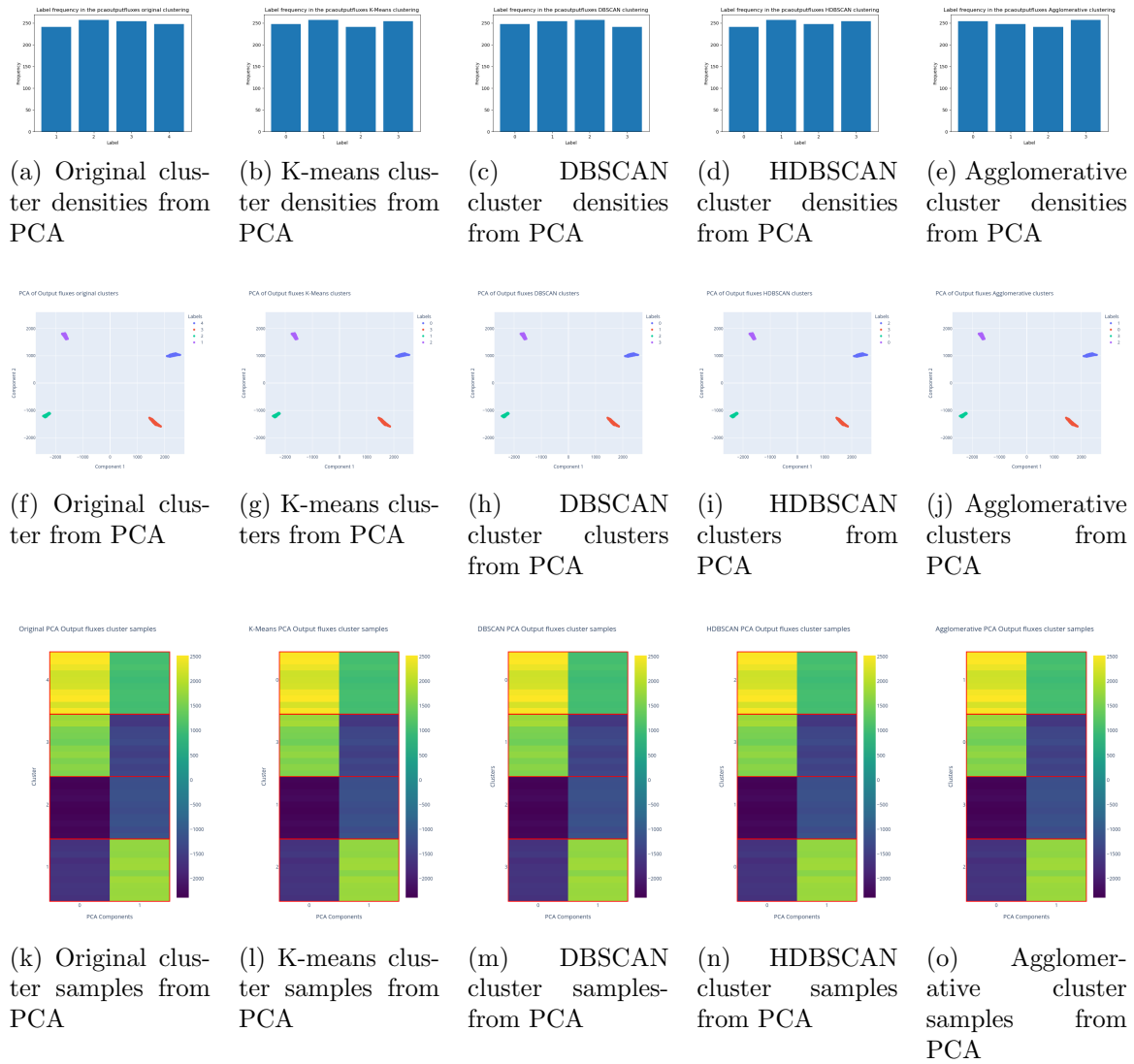


Figure 36: Comparison between clustering PCA Output fluxes algorithms

	Original	K-Means	DBSCAN	HDBSCAN	Agglomerative
Original	—	1	1	1	1
K-Means		—	1	1	1
DBSCAN			—	1	1
HDBSCAN				—	1

Table 18: Normalized Mutual Information between PCA Output fluxes clusters

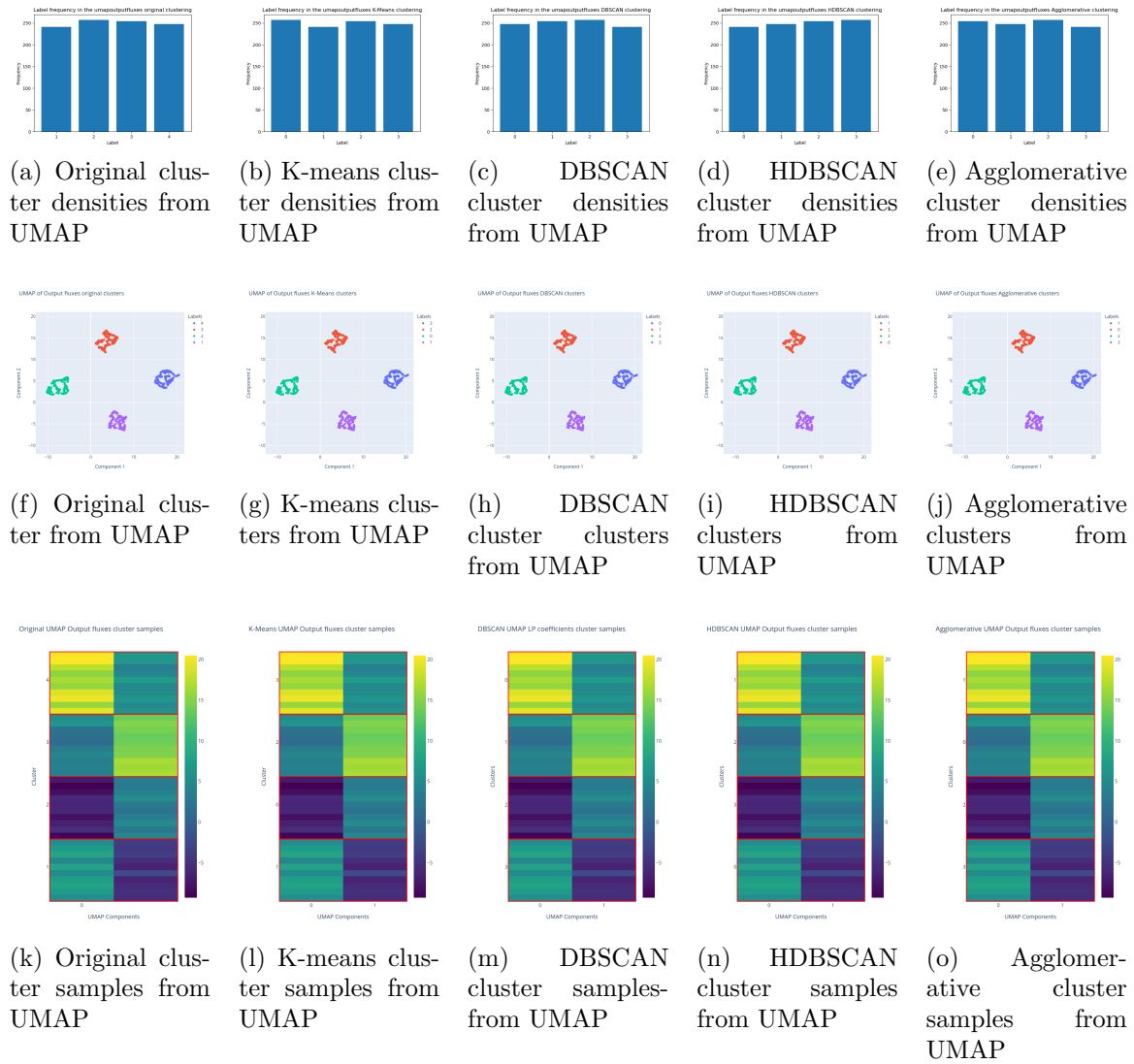


Figure 37: Comparison between clustering UMAP Output fluxes algorithms

	Original	K-Means	DBSCAN	HDBSCAN	Agglomerative
Original	—	1	1	1	1
K-Means		—	1	1	1
DBSCAN			—	1	1
HDBSCAN				—	1

Table 19: Normalized Mutual Information between UMAP Output fluxes clusters

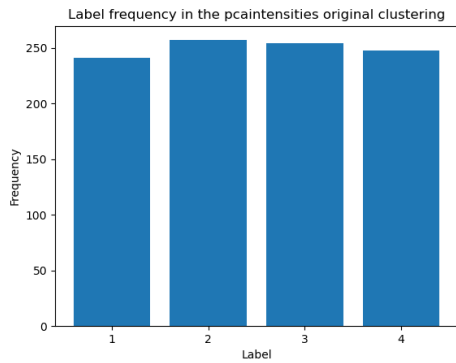
4.4 PSF Intensities clustering

4.4.1 K-Means

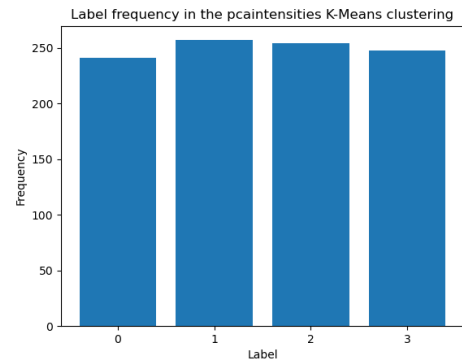
As K-Means allows for the number of clusters to be defined, and we know that there are 4 in the original dataset, K-Means is used to find 4 clusters.

	Number of clusters	Number of initializations
PCA PSF Intensities	4	10
UMAP PSF Intensities	4	10

Table 20: K-Means hyperparameter configuration for c coefficients clustering



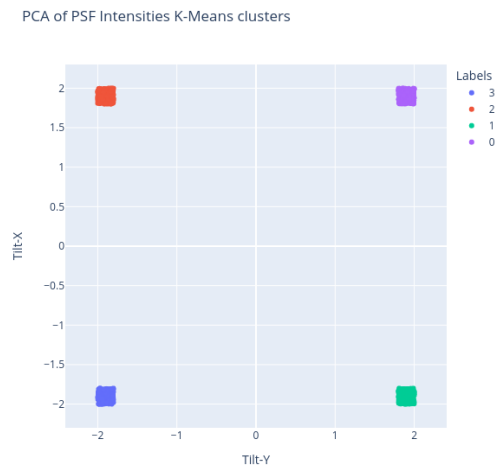
(a) Original cluster densities from PCA



(b) K-Means clusters densities from PCA

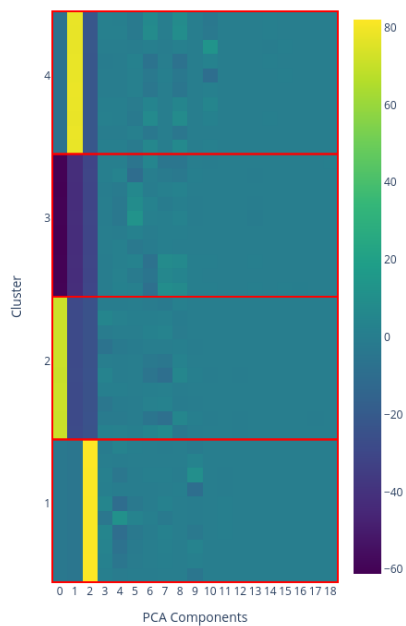


(c) Original clusters



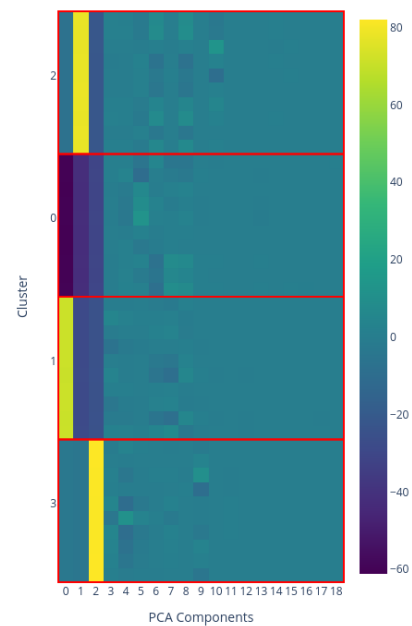
(d) K-Means clusters from PCA

Original PCA PSF intensities cluster samples



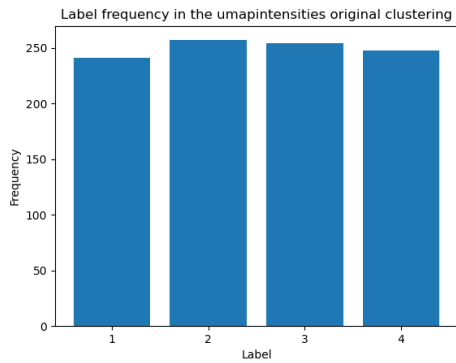
(e) Original cluster samples from PCA

K-Means PCA PSF Intensities cluster samples

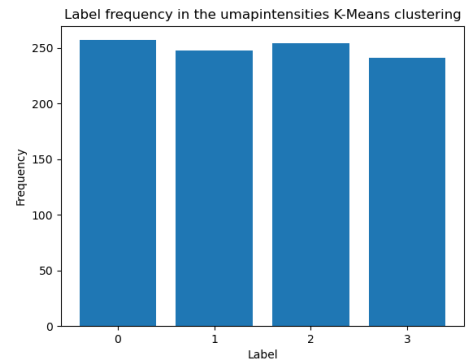


(f) K-Means cluster samples from PCA

Figure 38: Comparison between original clustering and K-Means clustering from PCA of PSF Intensities

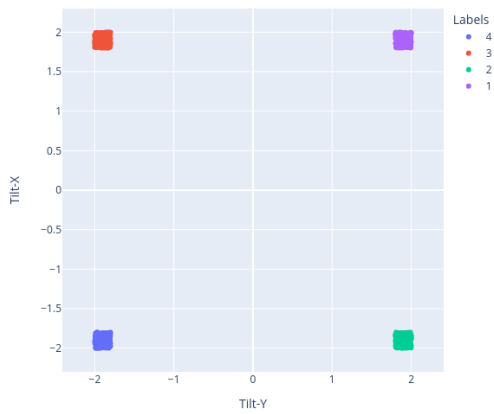


(a) Original cluster densities from UMAP



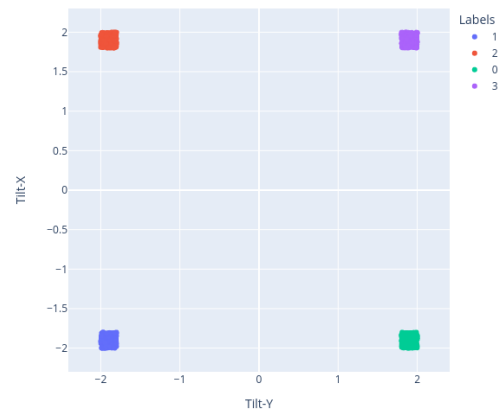
(b) K-Means clusters densities from UMAP

UMAP of PSF Intensities original clusters



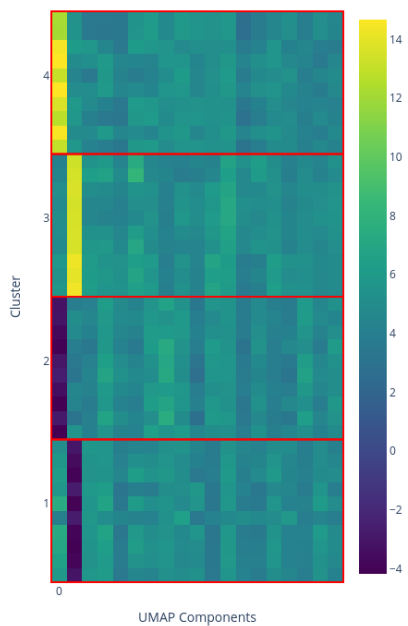
(c) Original clusters from UMAP

UMAP of PSF Intensities K-Means clusters



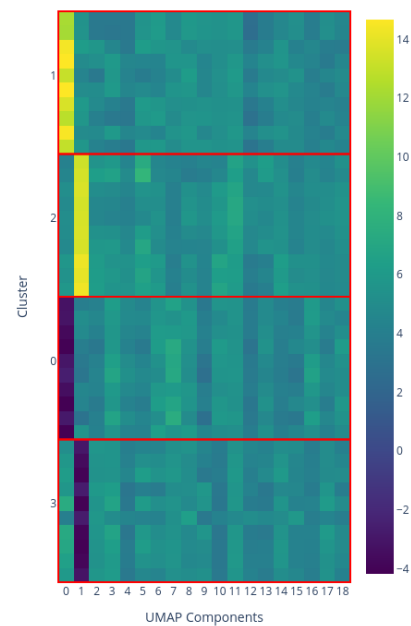
(d) K-Means clusters from UMAP

Original UMAP PSF Intensities cluster samples



(e) Original cluster samples from UMAP

K-Means UMAP PSF Intensities cluster samples



(f) K-Means cluster samples from UMAP

Figure 39: Comparison between original clustering and K-Means clustering from

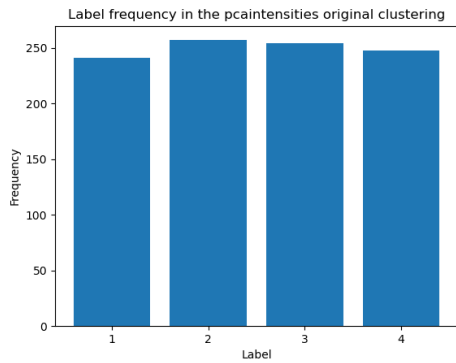
4.4.2 DBSCAN

A configuration that outputs 4 clusters is searched

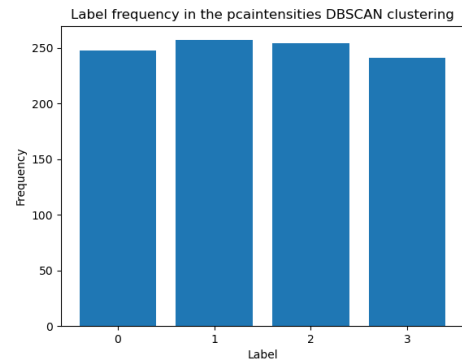
	Number of neighbours	Epsilon
PCA PSF Intensities	15	4.5
UMAP PSF Intensities	10	0.85

Table 21: DBSCAN hyperparameter configuration for PSF Intensities clustering

The results are the following:



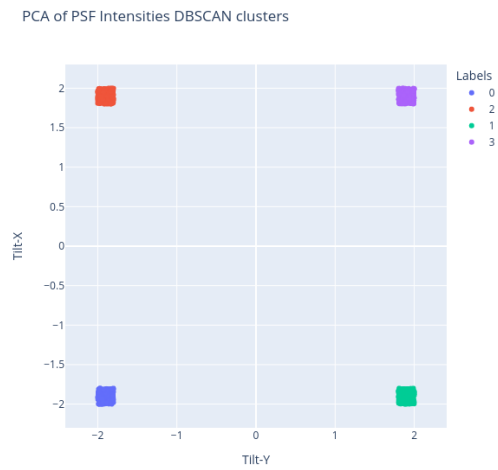
(a) Original cluster densities from PCA



(b) DBSCAN clusters densities from PCA

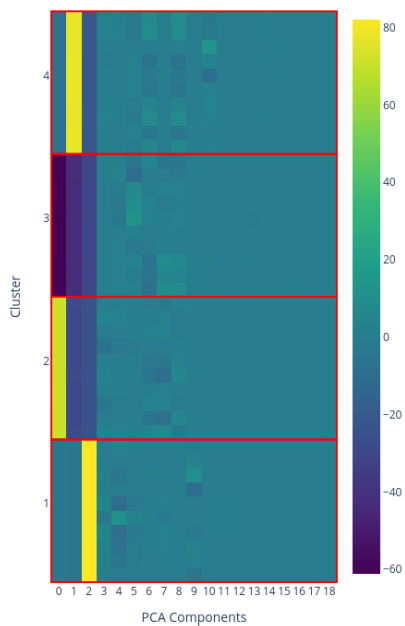


(c) Original clusters from PCA



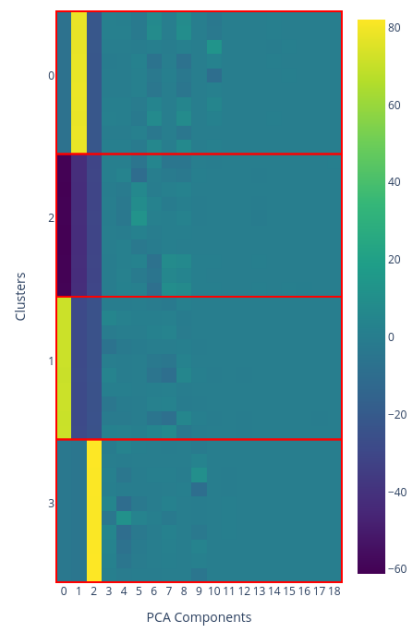
(d) DBSCAN clusters from PCA

Original PCA PSF intensities cluster samples



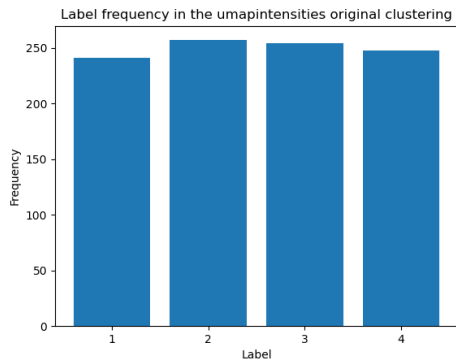
(e) Original cluster samples from PCA

DBSCAN PCA PSF Intensities cluster samples

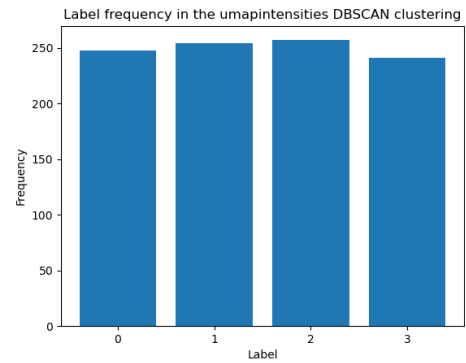


(f) DBSCAN cluster samples from PCA

Figure 40: Comparison between original clustering and DBSCAN clustering from PCA of PSF Intensities

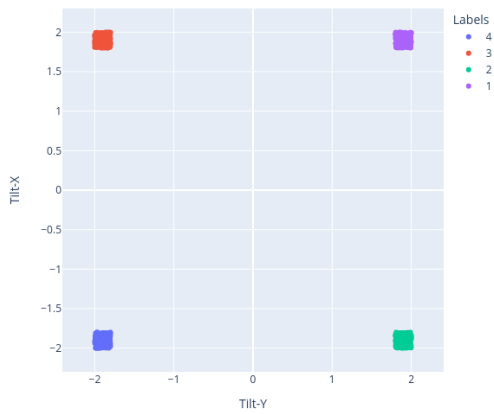


(a) Original cluster densities from UMAP



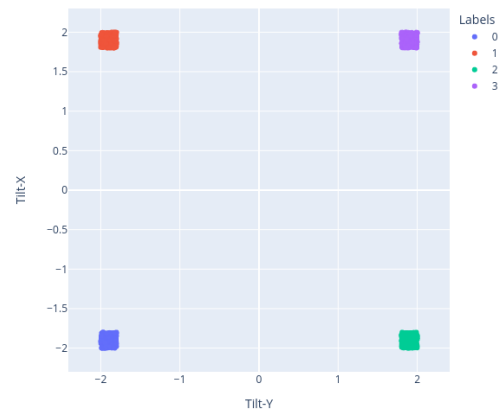
(b) DBSCAN clusters densities from UMAP

UMAP of PSF Intensities original clusters



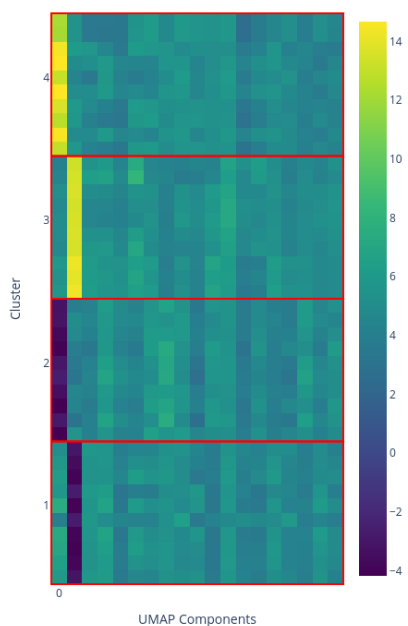
(c) Original clusters from UMAP

UMAP of PSF Intensities DBSCAN clusters



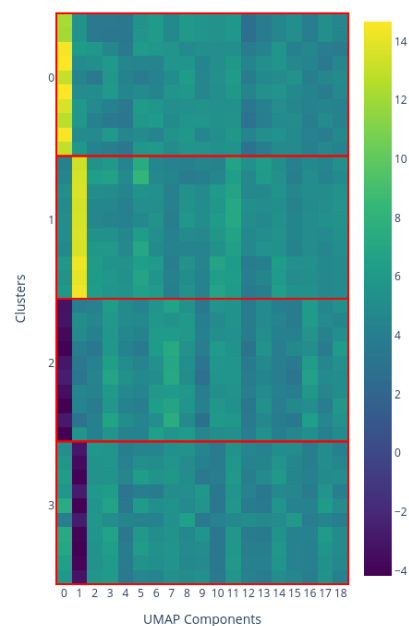
(d) DBSCAN clusters from UMAP

Original UMAP PSF Intensities cluster samples



(e) Original cluster samples from UMAP

DBSCAN UMAP PSF Intensities cluster samples



(f) DBSCAN cluster samples from UMAP

Figure 41: Comparison between original clustering and DBSCAN clustering from

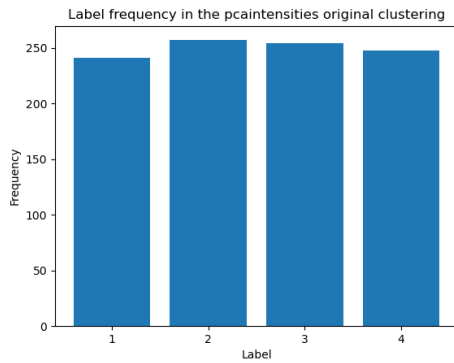
4.4.3 HDBSCAN

A configuration that outputs 4 clusters is searched.

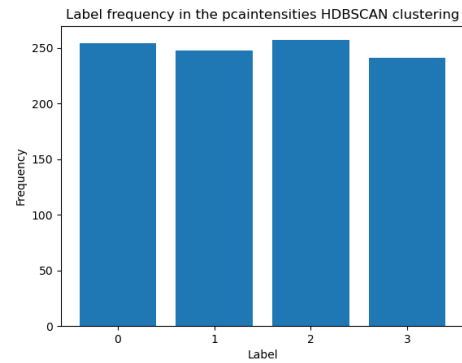
	Minimum cluster size
PCA PSF Intensities	21
UMAP PSF Intensities	25

Table 22: HDBSCAN hyperparameter configuration for PSF Intensities clustering

The results are the following:



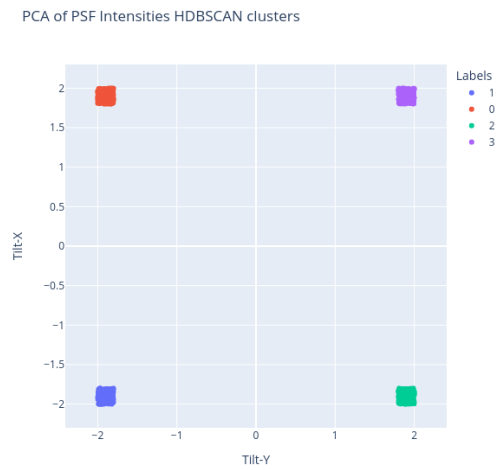
(a) Original cluster densities from PCA



(b) HDBSCAN clusters densities from PCA

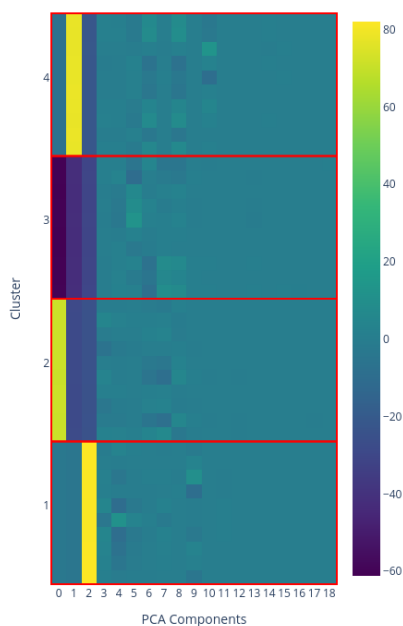


(c) Original clusters from PCA



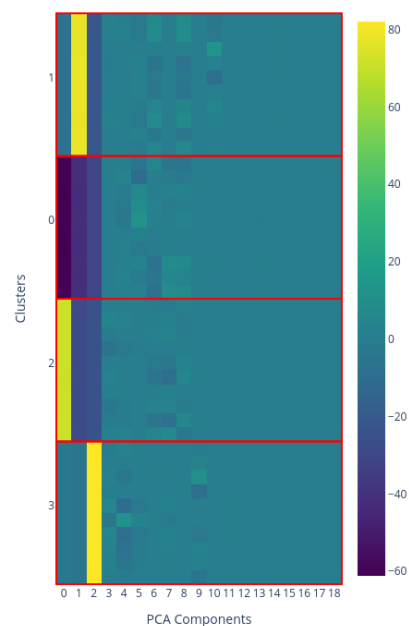
(d) HDBSCAN clusters from PCA

Original PCA PSF intensities cluster samples



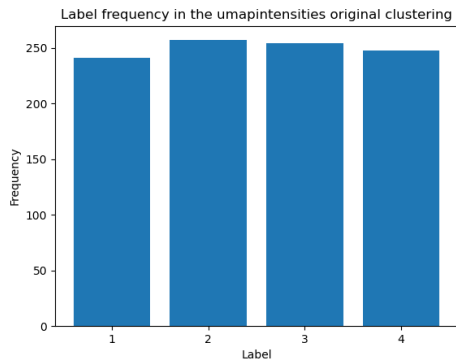
(e) Original cluster samples from PCA

HDBSCAN PCA PSF Intensities cluster samples

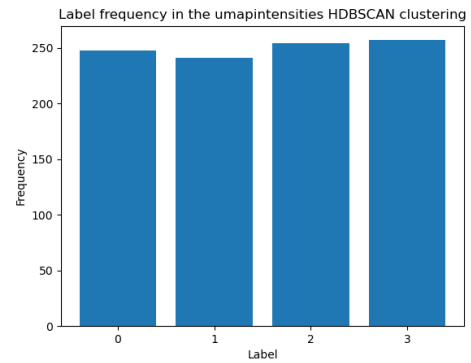


(f) HDBSCAN cluster samples from PCA

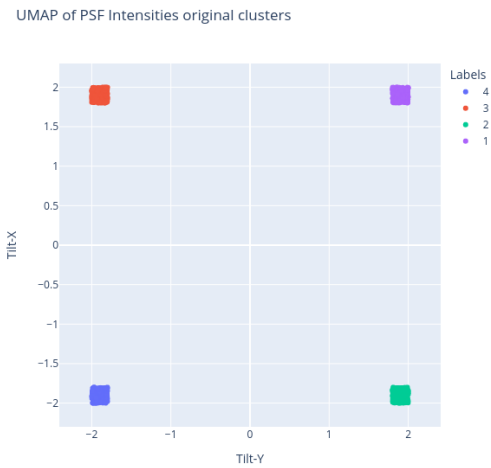
Figure 42: Comparison between original clustering and HDBSCAN clustering from



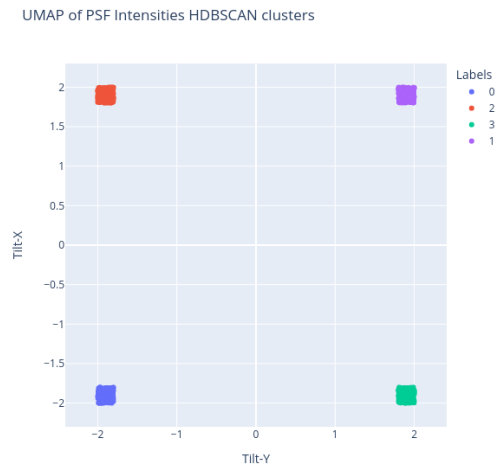
(a) Original cluster densities from UMAP



(b) HDBSCAN clusters densities from UMAP

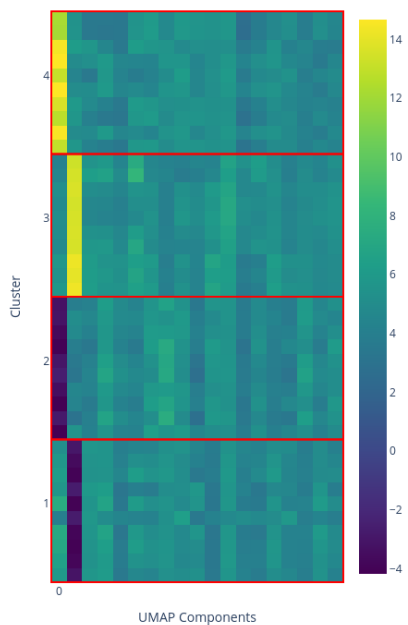


(c) Original clusters from UMAP



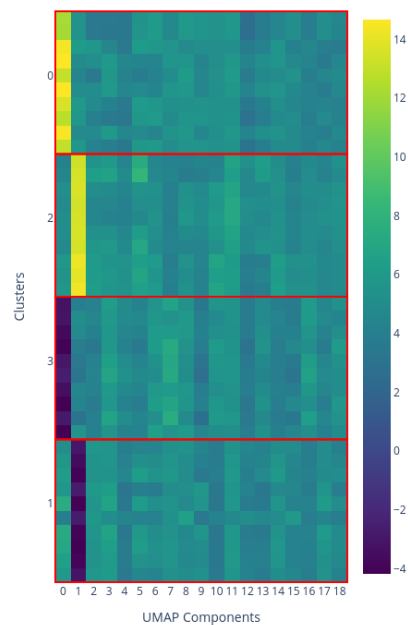
(d) HDBSCAN clusters from UMAP

Original UMAP PSF Intensities cluster samples



(e) Original cluster samples from UMAP

HDBSCAN UMAP PSF Intensities cluster samples



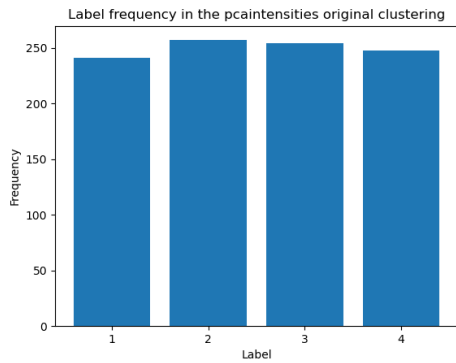
(f) HDBSCAN cluster samples from UMAP

4.4.4 Agglomerative clustering

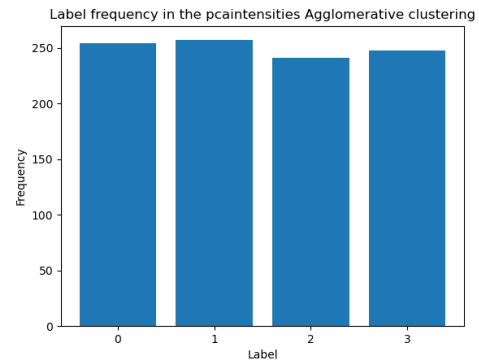
	Number of clusters
PCA PSF Intensities	4
UMAP PSF Intensities	4

Table 23: Agglomerative hyperparameter configuration for PSF Intensities clustering

The results are the following:



(a) Original cluster densities from PCA



(b) Agglomerative clusters densities from PCA

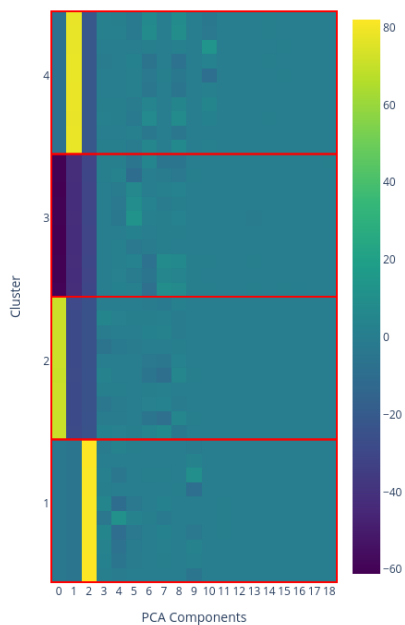


(c) Original clusters from PCA



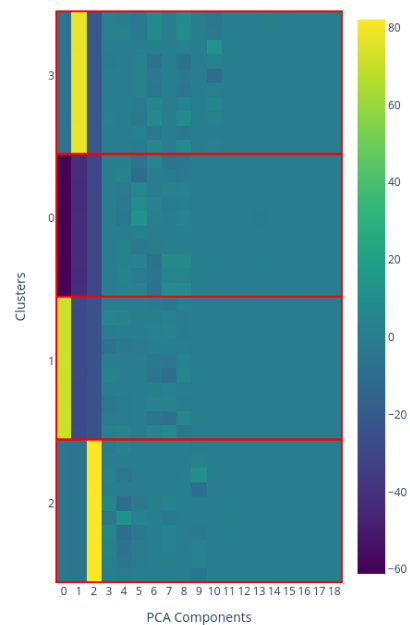
(d) Agglomerative clusters from PCA

Original PCA PSF intensities cluster samples

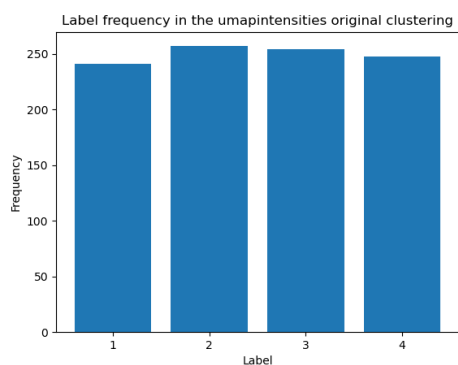


(e) Original cluster samples from PCA

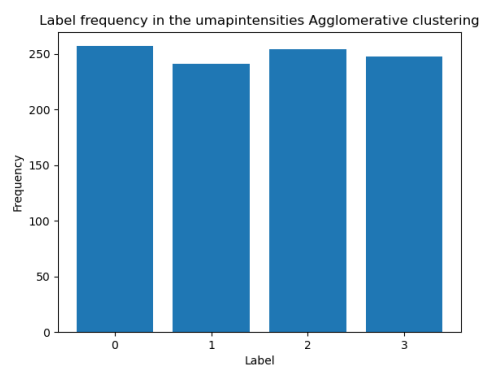
Agglomerative PCA PSF Intensities cluster samples



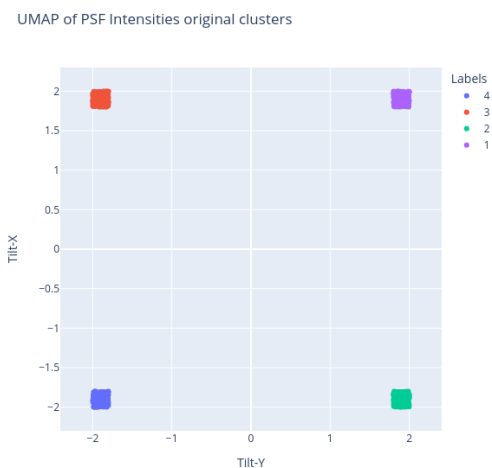
(f) Agglomerative cluster samples from PCA



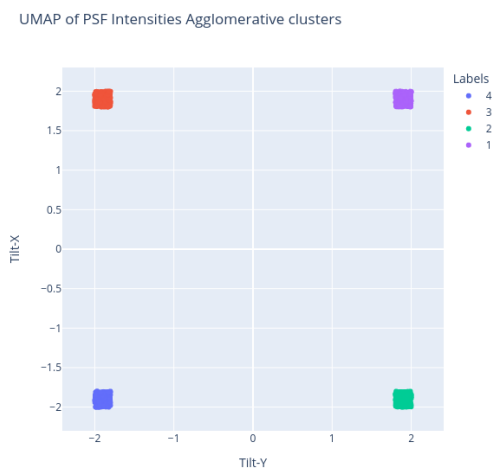
(a) Original cluster densities from UMAP



(b) Agglomerative clusters densities from UMAP

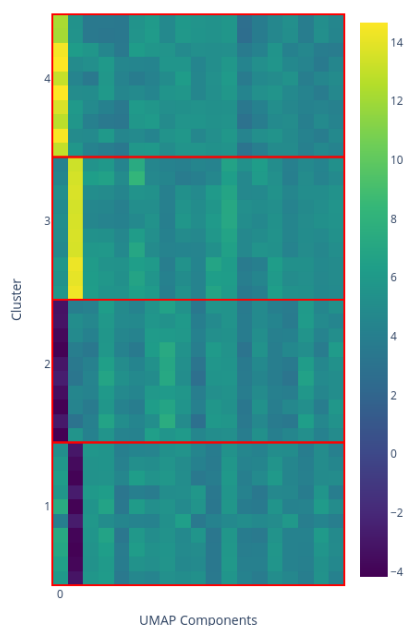


(c) Original clusters from UMAP



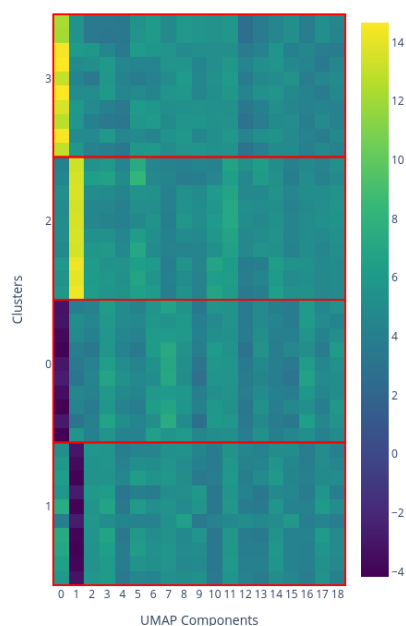
(d) Agglomerative clusters from UMAP

Original UMAP PSF Intensities cluster samples



(e) Original cluster samples from UMAP

Agglomerative UMAP PSF Intensities cluster samples



(f) Agglomerative cluster samples from UMAP

4.4.5 Summary

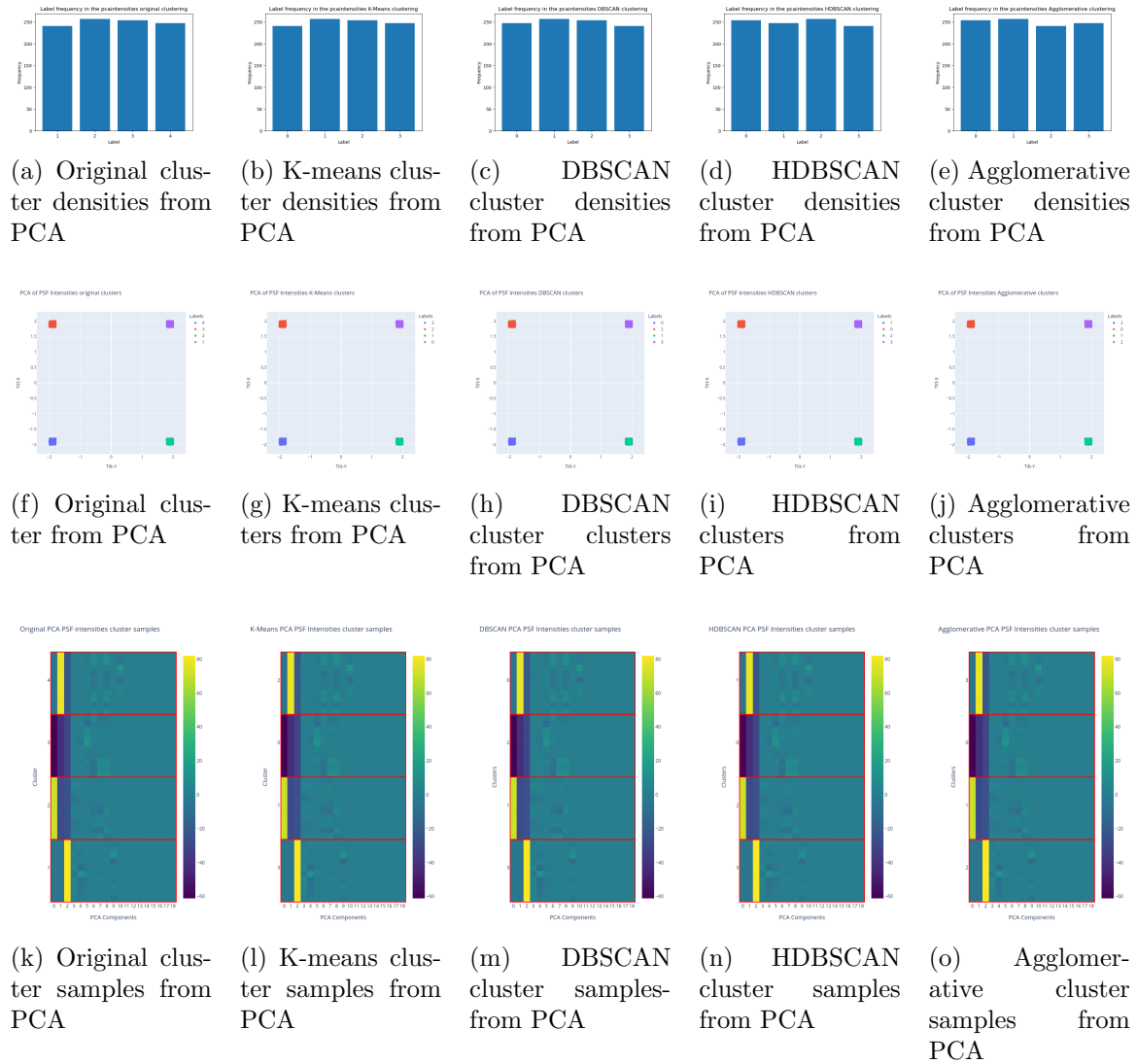


Figure 46: Comparison between clustering PCA PSF Intensities algorithms

	Original	K-Means	DBSCAN	HDBSCAN	Agglomerative
Original	\diagdown	1	1	1	1
K-Means		\diagdown	1	1	1
DBSCAN			\diagdown	1	1
HDBSCAN				\diagdown	1

Table 24: Normalized Mutual Information between PCA PSF Intensities clusters

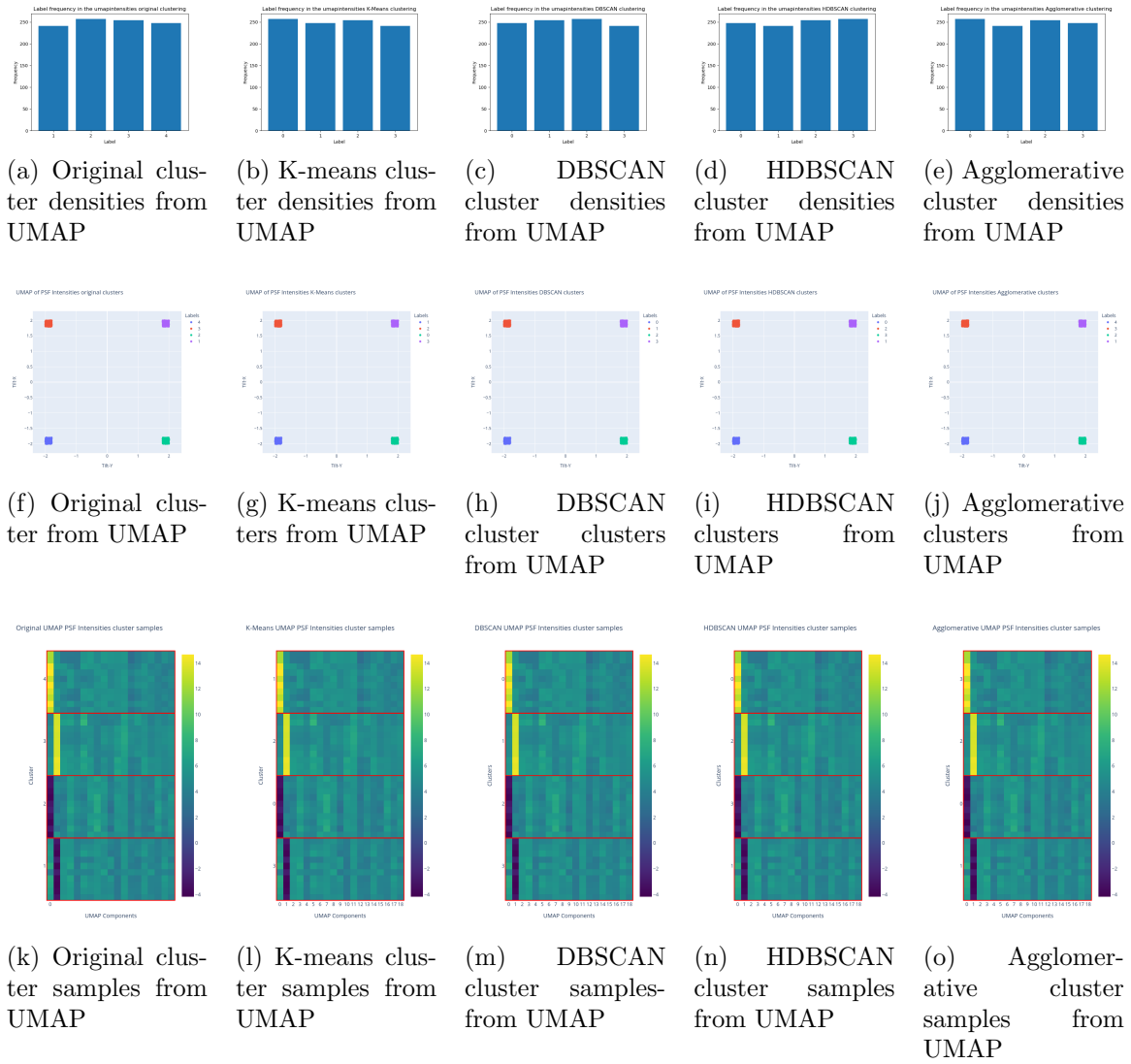


Figure 47: Comparison between clustering UMAP PSF Intensities algorithms

	Original	K-Means	DBSCAN	HDBSCAN	Agglomerative
Original	<div> </div>	1	1	1	1
K-Means		<div> </div>	1	1	1
DBSCAN			<div> </div>	1	1
HDBSCAN				<div> </div>	1

Table 25: Normalized Mutual Information between UMAP PSF Intensities clusters

5 Dataset clusters comparison

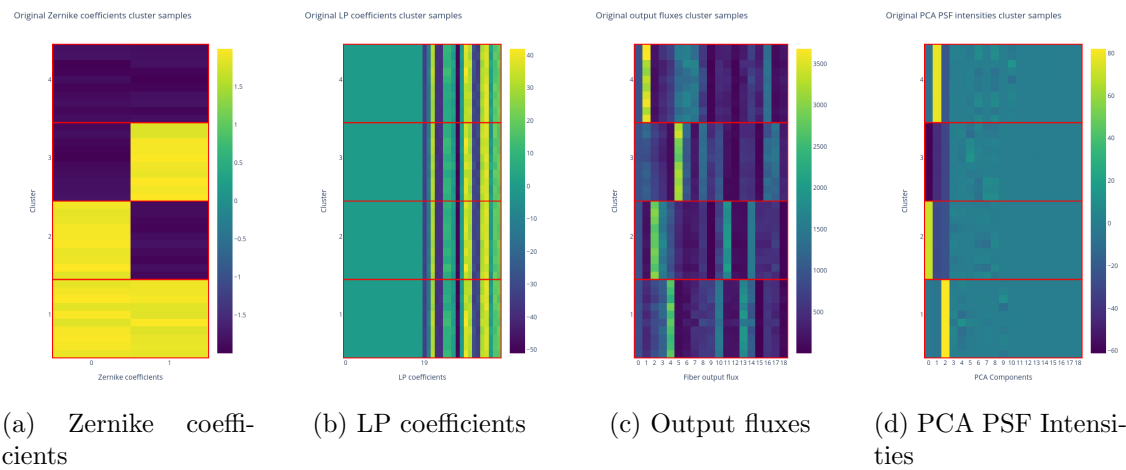


Figure 48: Original clusters from the datasets

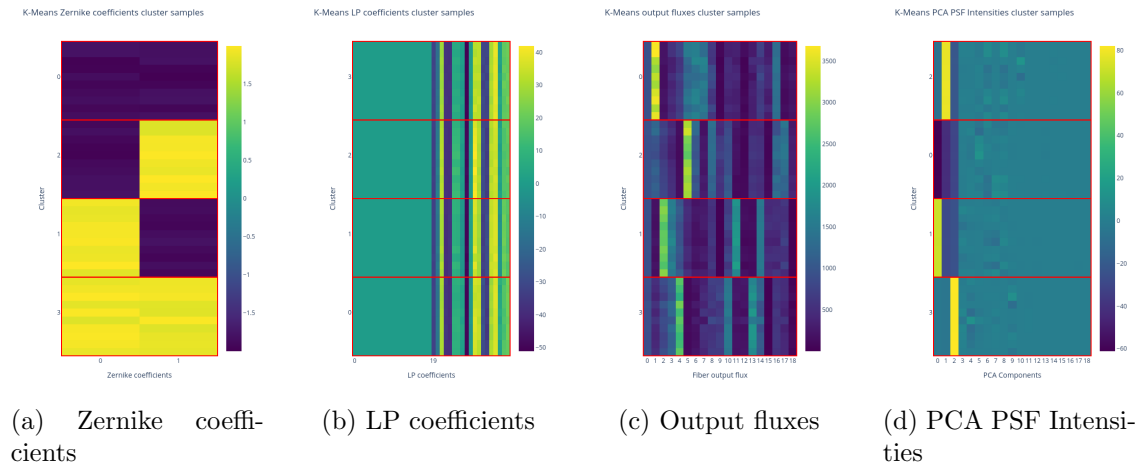


Figure 49: K-Means clusters from the datasets

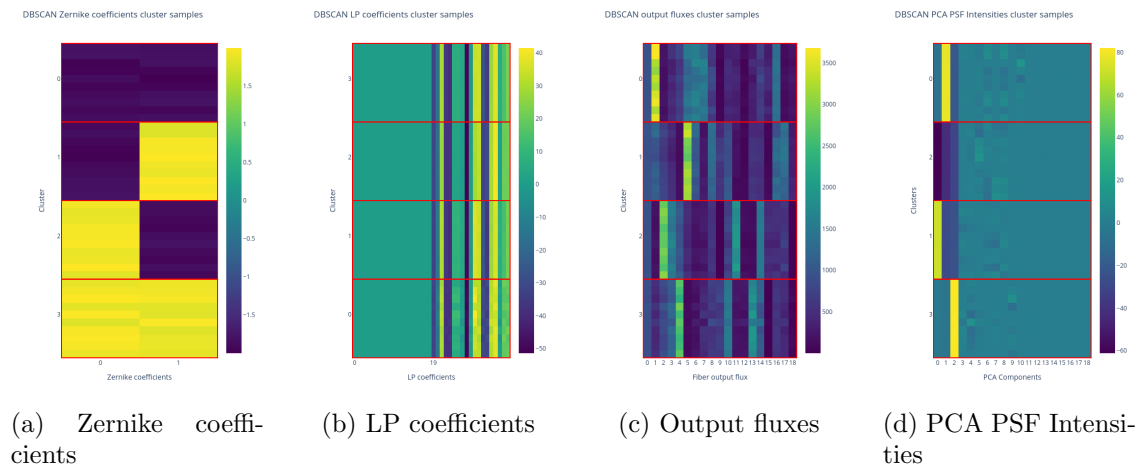


Figure 50: DBSCAN clusters from the datasets

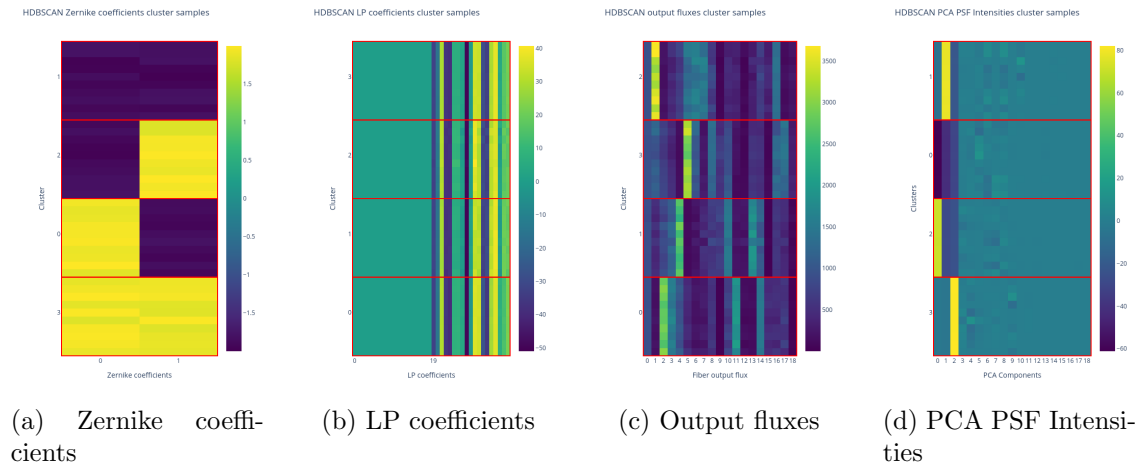


Figure 51: HDBSCAN clusters from the datasets

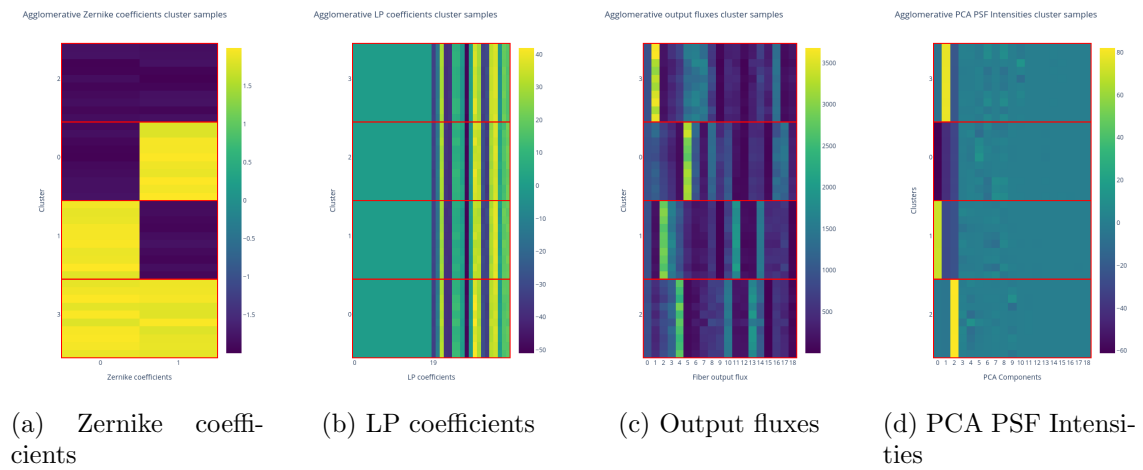


Figure 52: Agglomerative clusters from the datasets