

# Project of Processing Big Data

Group 8:

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# Introduction

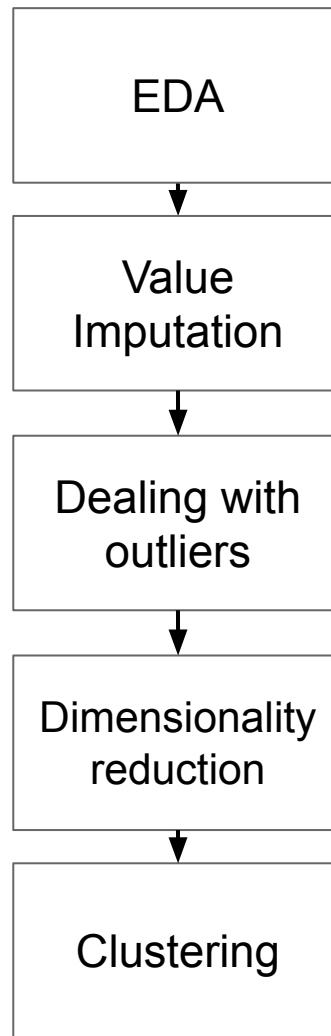


We did it for 3 methods using the dauphine dataset.

We clustered the following datasets:

1. Skeletons
2. Image embeddings
3. Skeletons + images

# Pipeline

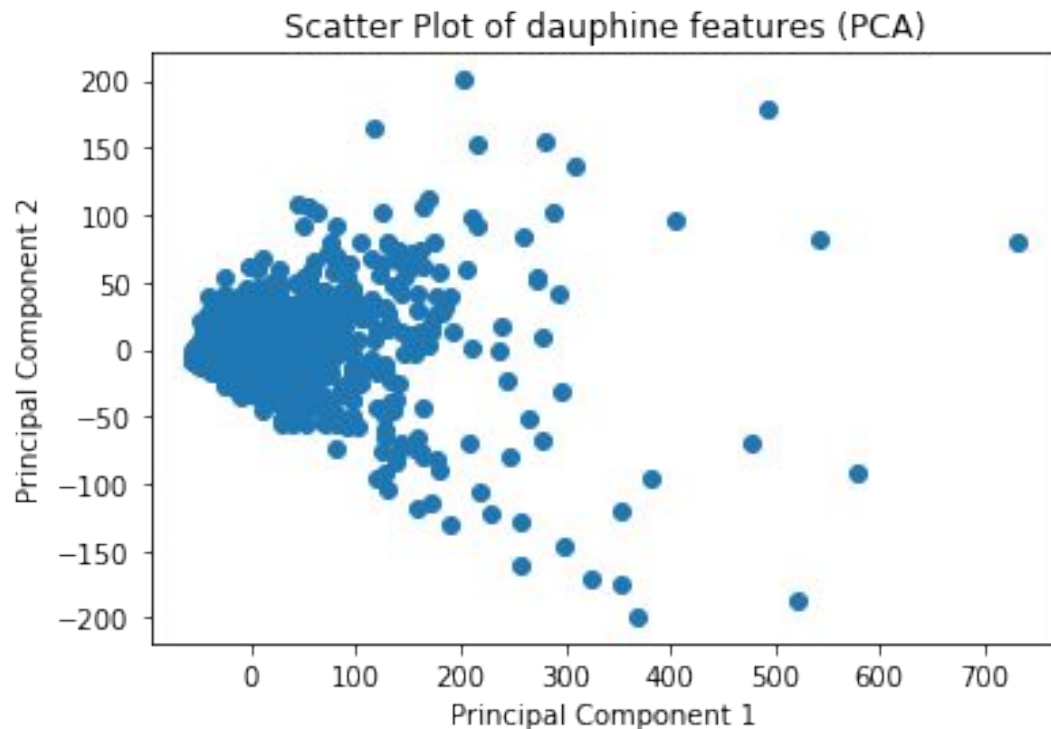


# EDA – Dauphine features

```
In [5]: 1 np.shape(dauphine_features)
```

```
Out[5]: (2048, 10734)
```

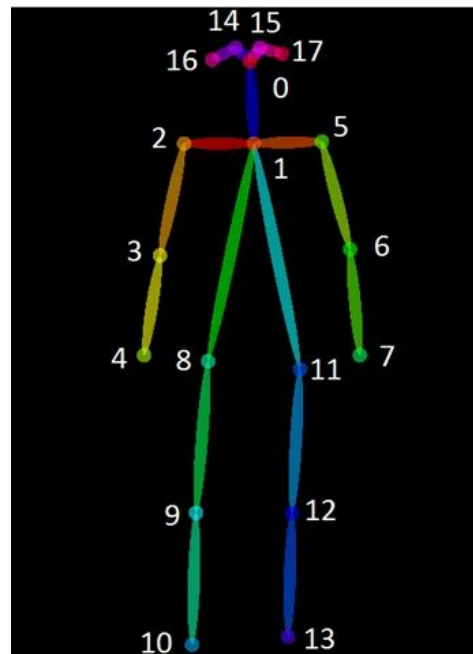
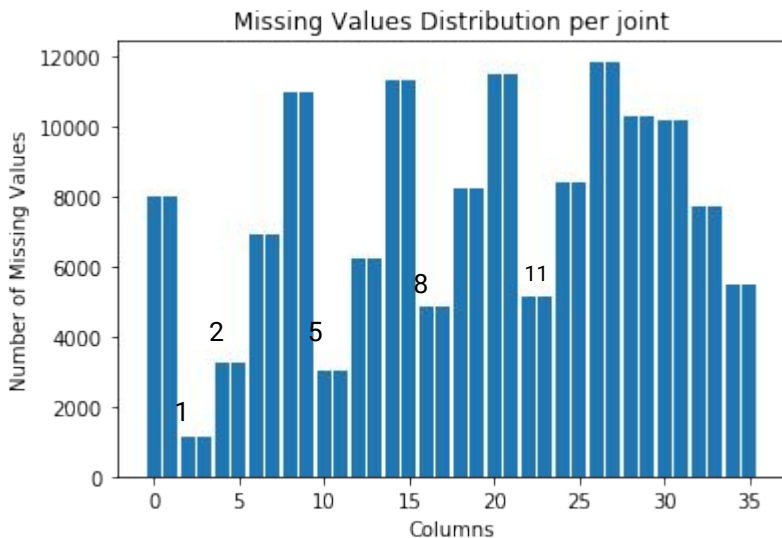
```
threshold for distant points= 400  
percentage of distant points: 0.390625  
Index: 67  
Index: 124  
Index: 138  
Index: 520  
Index: 945  
Index: 1147  
Index: 1222  
Index: 1250
```



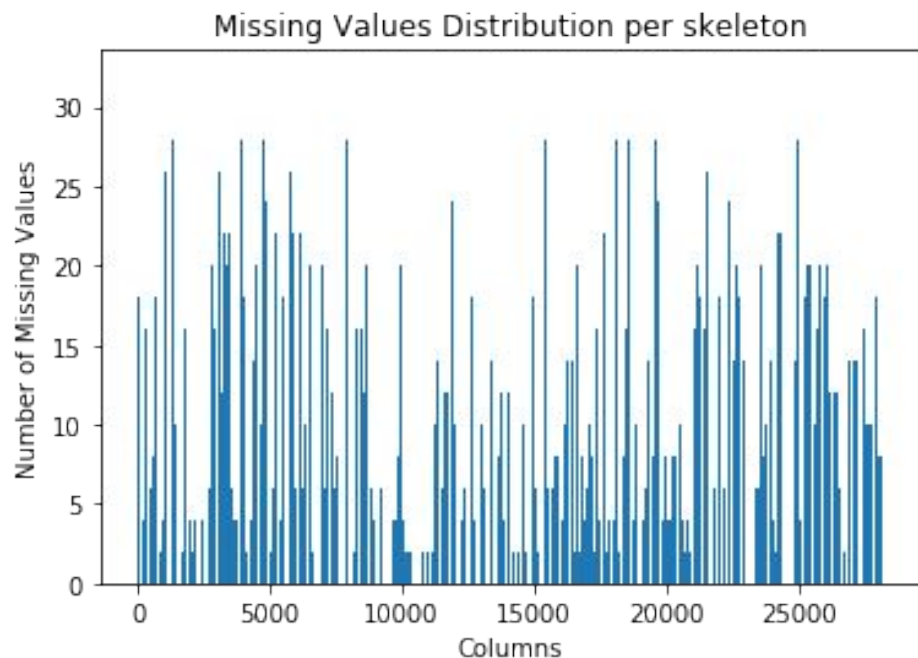
# EDA – Dauphine Incomplete Skeletons

```
1 #without probabilities, only coordinates  
2 np.shape(dauphine_skeletons)
```

(28232, 36)



# EDA – Dauphine Incomplete Skeletons



```
1 #without probabilities, only coordinates  
2 np.shape(dauphine_skeletons)
```

```
(28232, 36)
```

# Missing value imputation

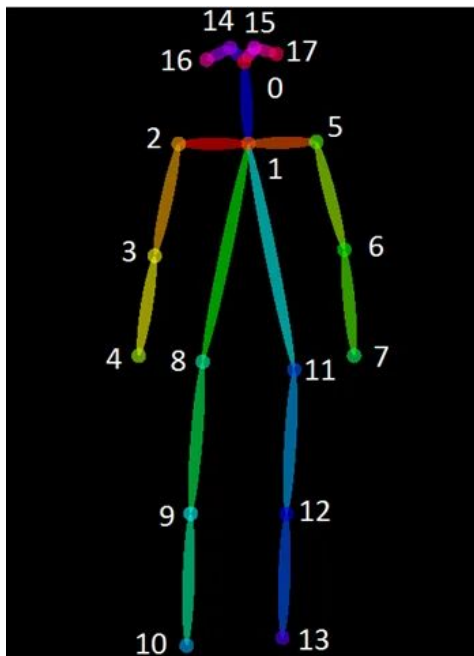


Methods tested:

- ❖ Mean value imputation
- ❖ MICE
- ❖ K-nearest neighbours
- ❖ GLRM usando h2o

# Standardizing skeleton data

Openpose



$$z = (x - \mu) / \sigma$$

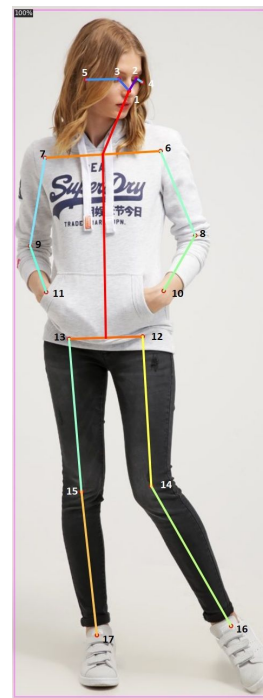
$$\bar{x}_{\text{openpose}} = \frac{x_1 + x_2 + x_5}{3}$$

$$\bar{y}_{\text{openpose}} = \frac{y_1 + y_2 + y_5}{3}$$

$$\bar{x}_{\text{otherpose}} = \frac{x_6 + x_7}{2}$$

$$\bar{y}_{\text{otherpose}} = \frac{y_6 + y_7}{2}$$

Otherpose



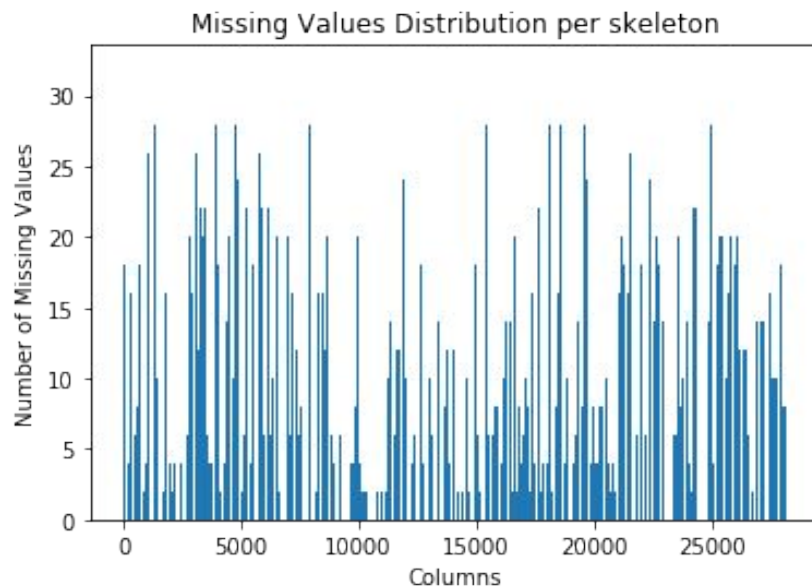
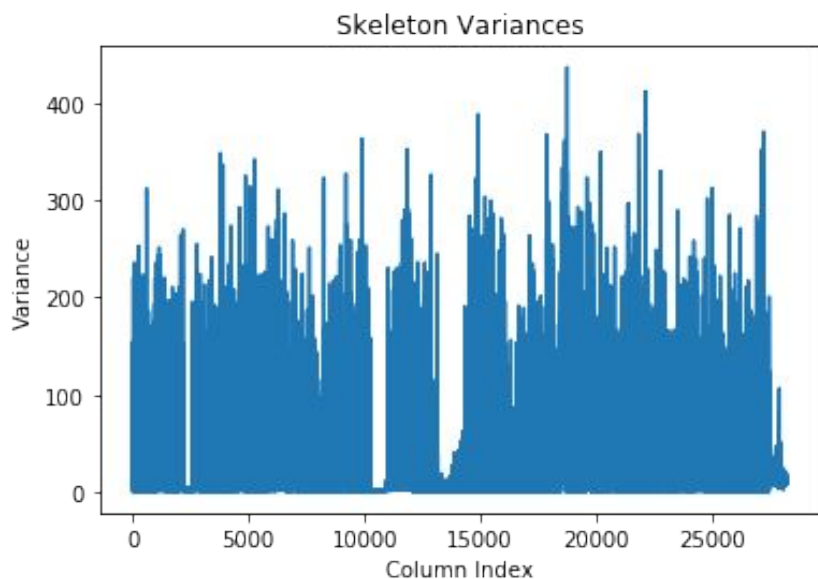


# GLRM

$$m \left\{ \left[ \begin{array}{c} \overbrace{\hspace{1.5cm}}^n \\ A \end{array} \right] \right\} \approx m \left\{ \left[ \begin{array}{c} \overbrace{\hspace{1.5cm}}^k \\ X \end{array} \right] \left[ \begin{array}{c} \overbrace{\hspace{1.5cm}}^n \\ Y \end{array} \right] \right\} k$$

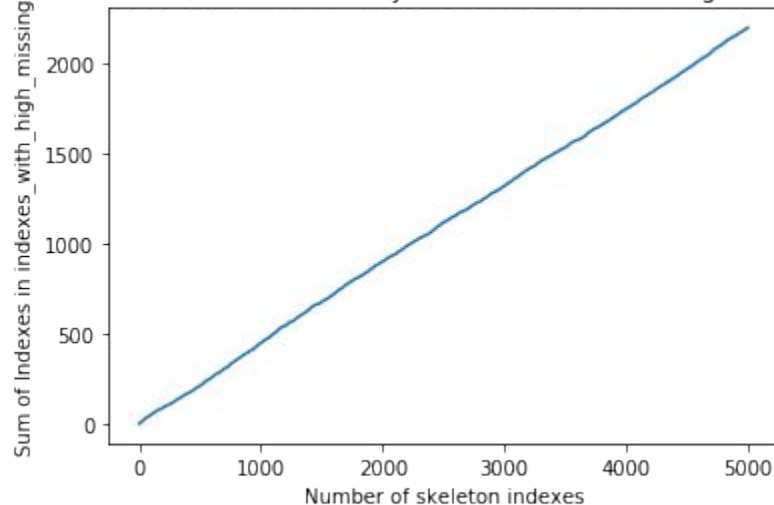
```
# Define and train the GLRM model to impute missing values
glrm_model = H2OGeneralizedLowRankEstimator(k=20,
                                             loss="Quadratic",
                                             regularization_x="l1",
                                             regularization_y="l1",
                                             max_iterations=100,
                                             recover_svd=True)
```

# Dauphine Complete Skeletons

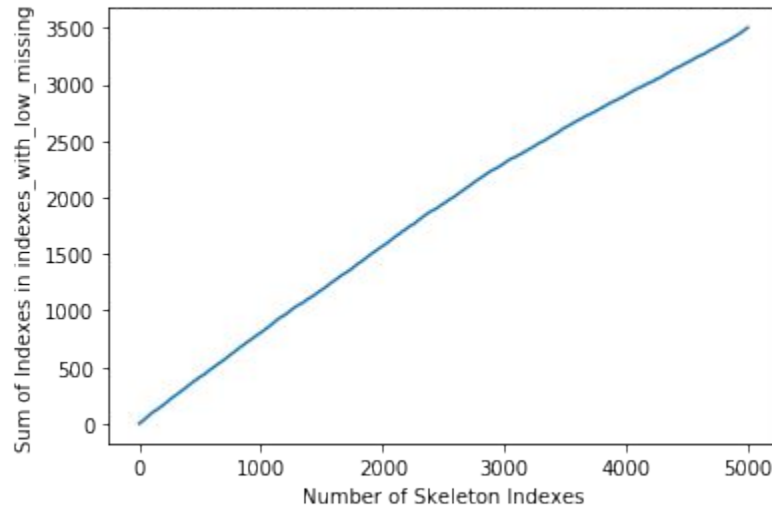


# Dauphine Complete Skeletons

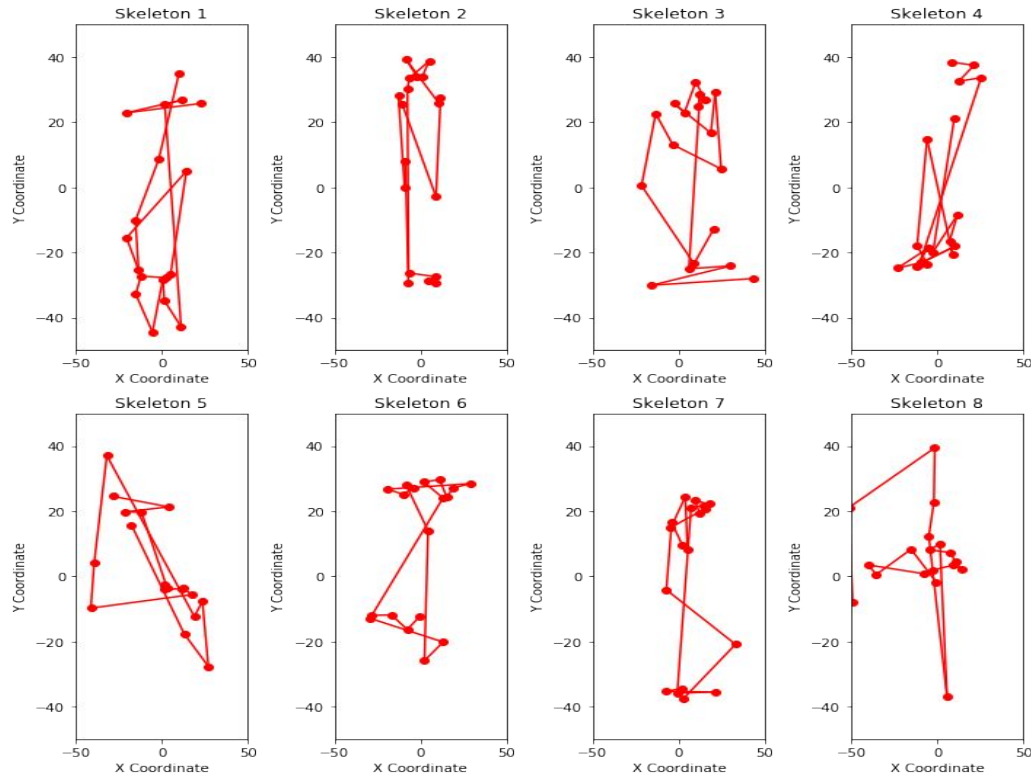
Number of Skeletons with many NAs in skeletons with higher variance



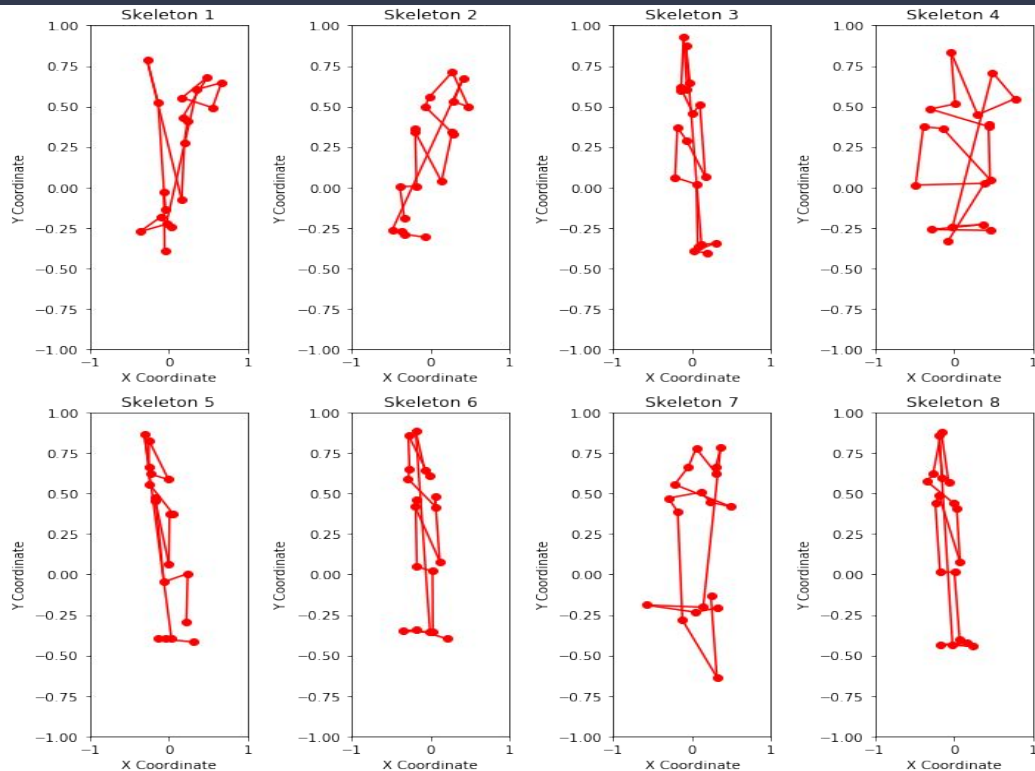
Number of Skeletons with few NAs in skeletons with lower variance



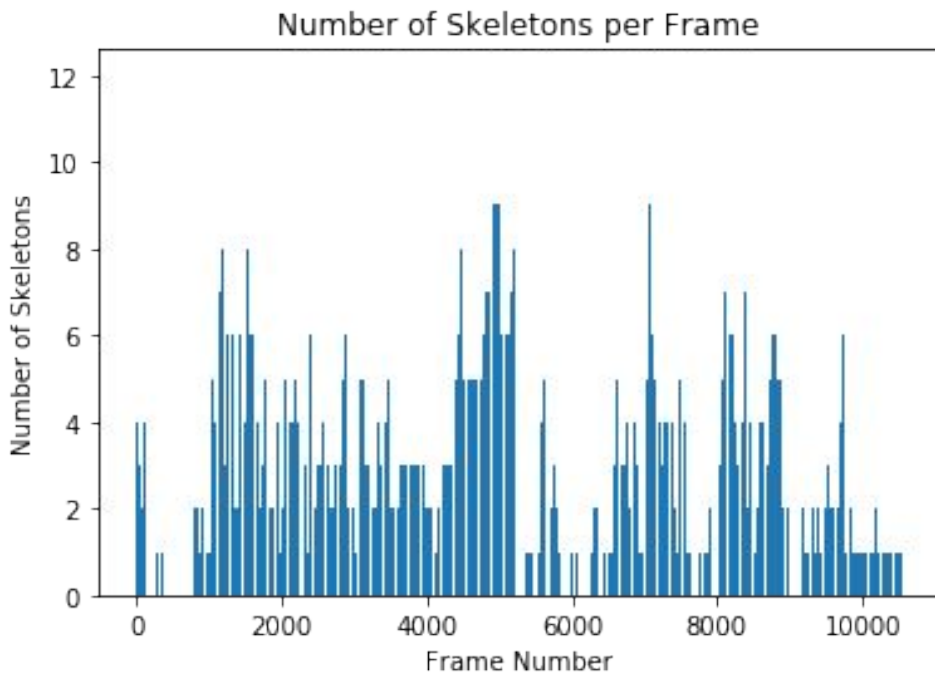
# Dauphine Complete Skeletons



# Dauphine Complete Skeletons



# Dauphine Complete Skeletons



# Detecting rank

- For skeleton and image embeddings data the method is the same
- Perform SVD and  $r$  corresponds to the number of components that explain at least 99.9% of cumulative explained variance

For skeleton the rank is 20.

For image embeddings is 95.

# Removing outliers and reducing dimensionality

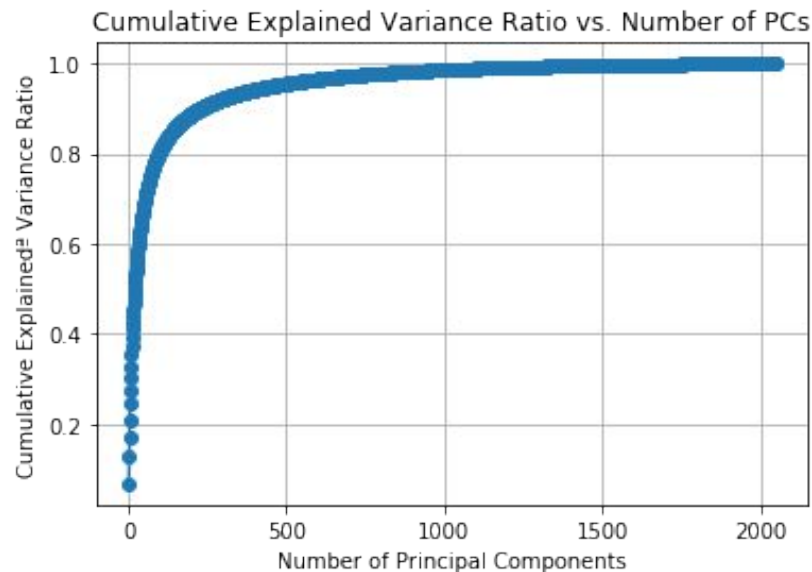
- For skeleton and image embeddings data the method is the same
- We calculate the distance to the Null space

## Reducing dimensionality

- Take the SVD obtained and only retain the most important  $r$  components



# Dimensionality Reduction – Features



Number of components necessary for 80.0 %% cumulative explained percentage is 95  
Shape of reduced\_data: (10734, 95)  
(10734, 95)

# Clustering – The applied methods and datasets

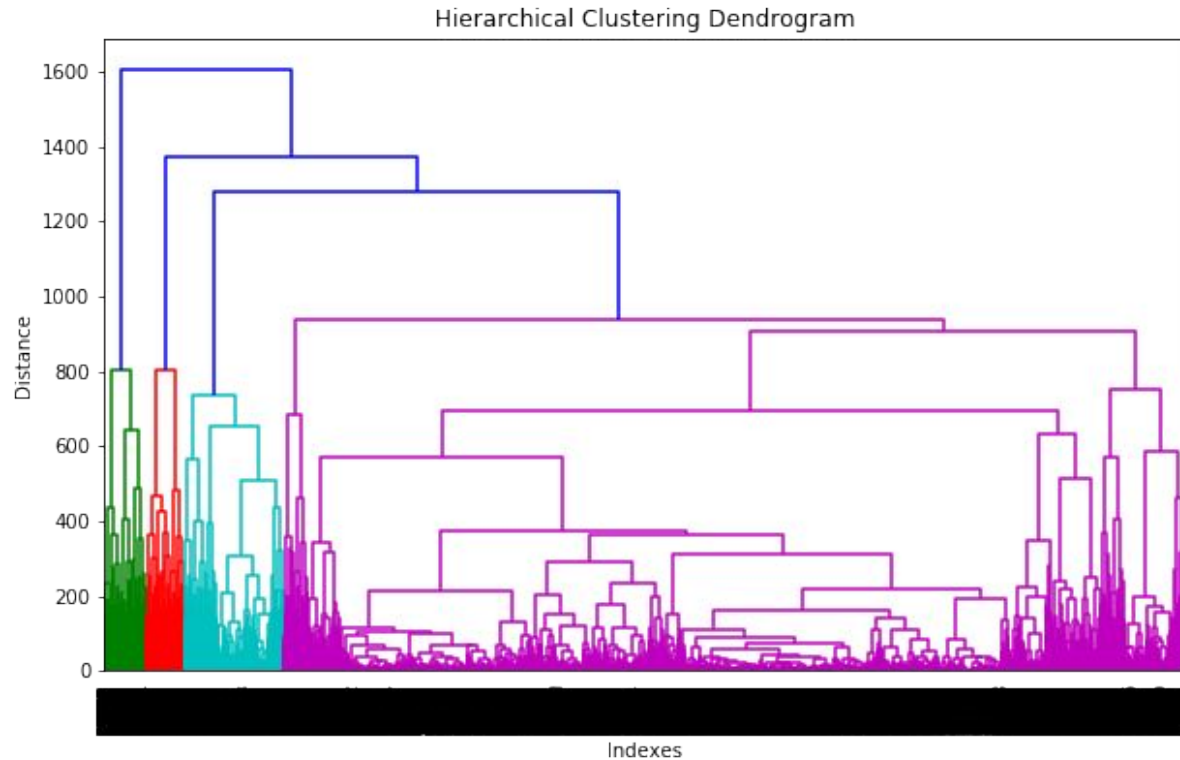
We performed the following clusterings:

- ❖ K-Means
- ❖ Hierarchical Clustering:
  - Complete Linkage
  - Ward Method
  - Centroid Method

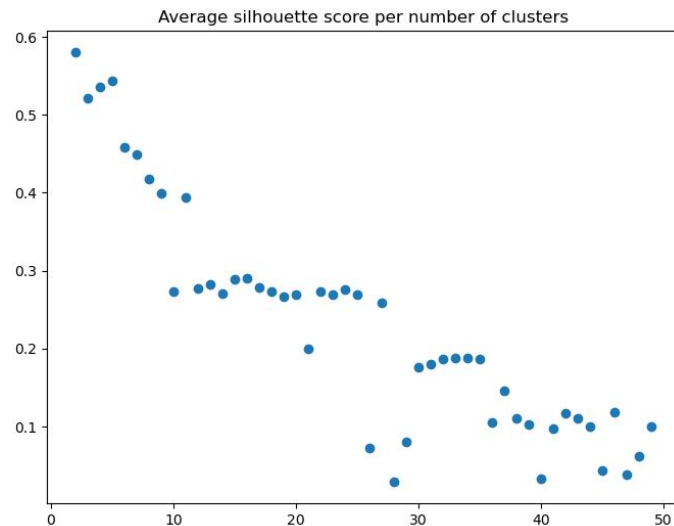
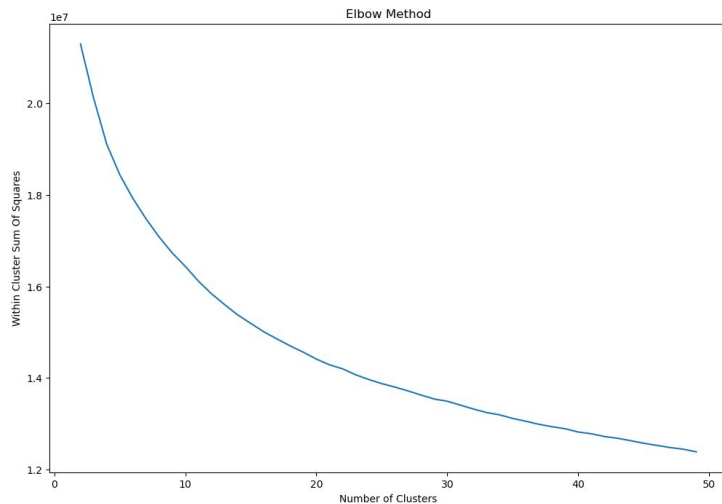
We clustered the following datasets:

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2. Image embeddings
3. Skeletons + images

# 1. Skeletons: Hierarchical Clustering – Ward Method



# 1. Skeletons – Number of clusters



# Clustering visualization only for skeleton data

Cluster

0



1



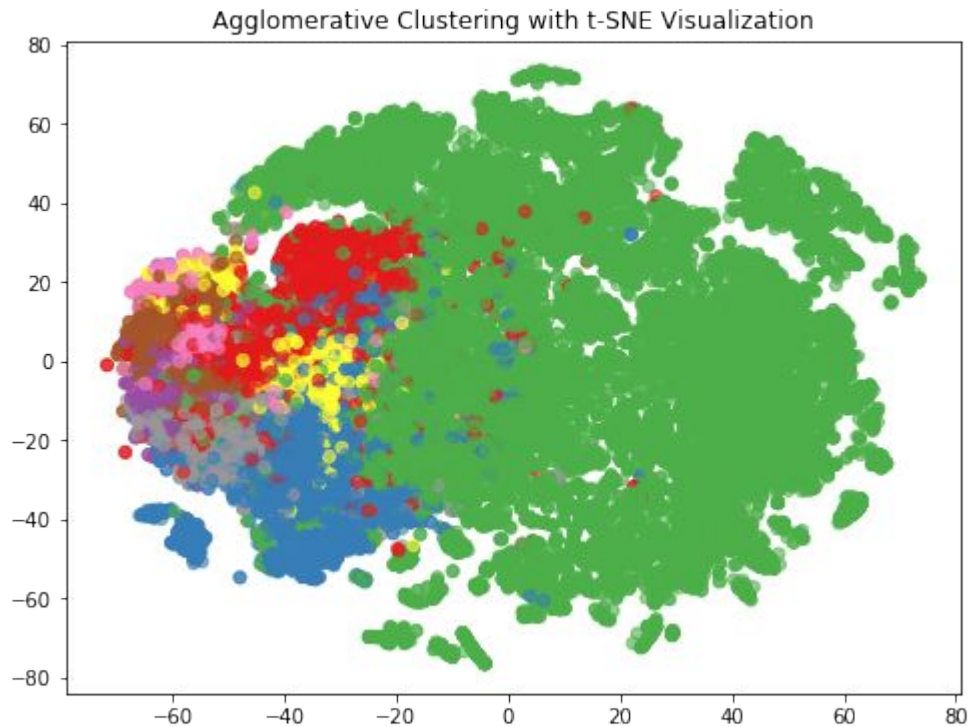
2



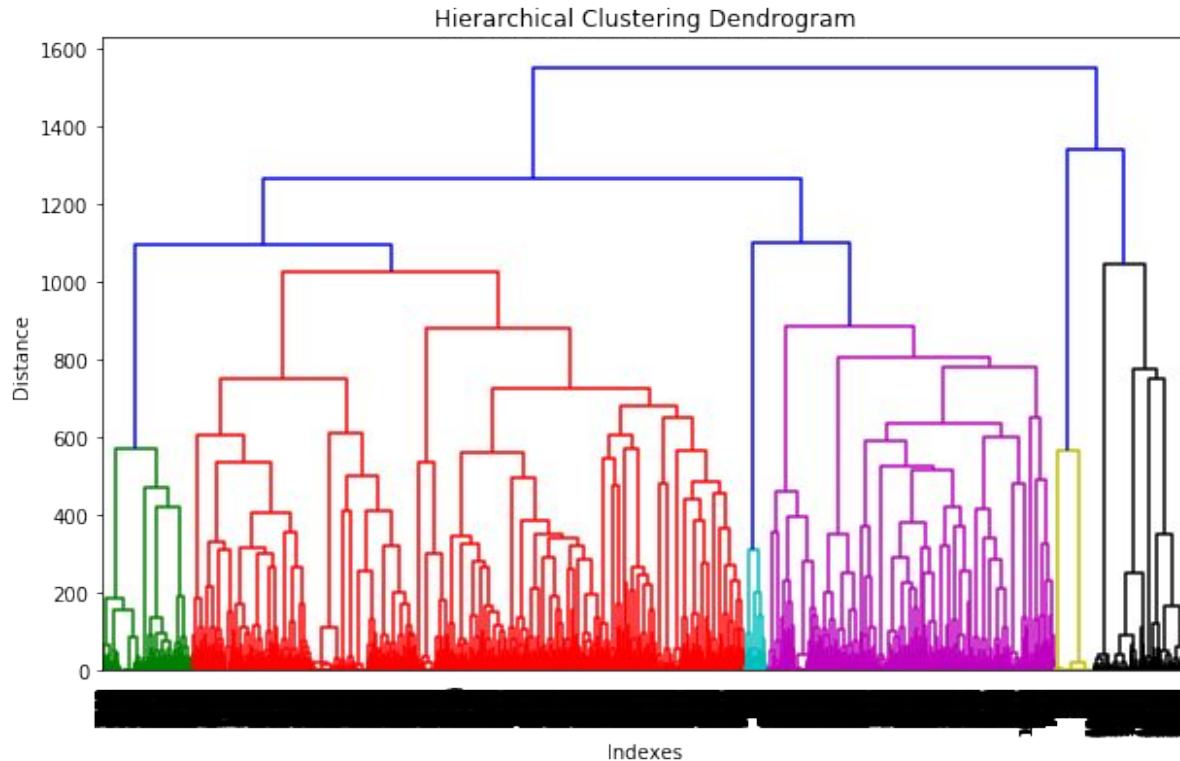
3



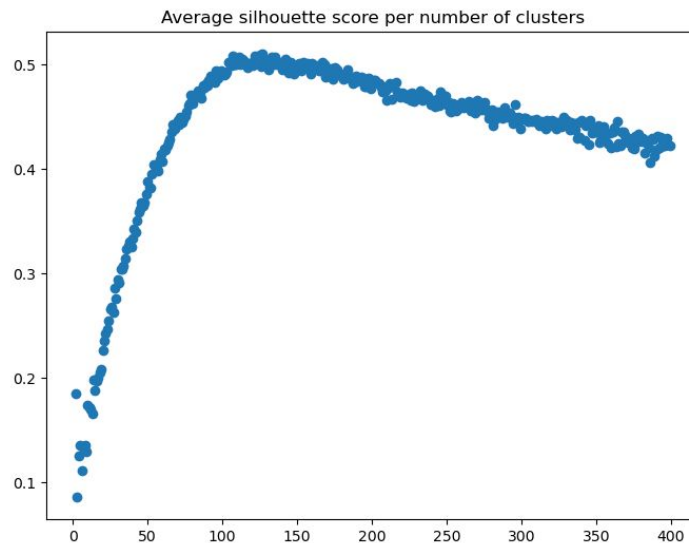
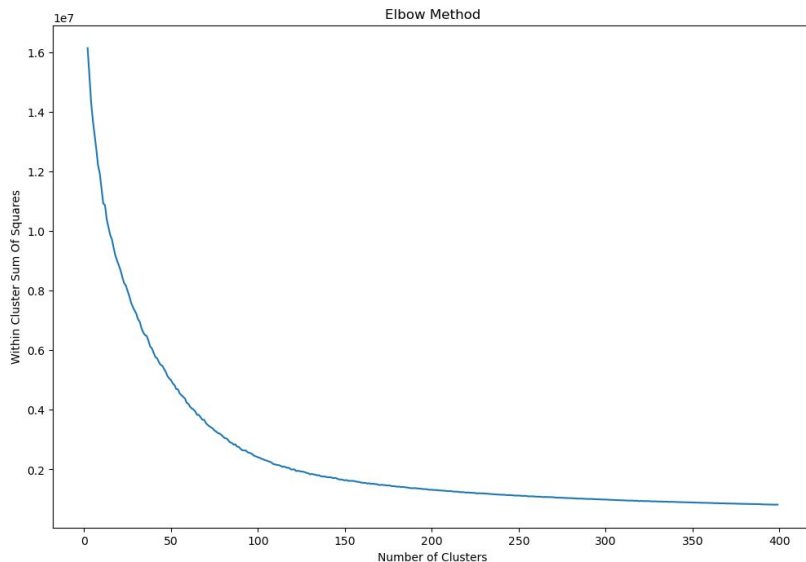
# 1. Skeletons –Clustering Visualization with t-sne



## 2. Images : Hierarchical Clustering – Ward Method



## 2. Images– Number of clusters





# Clustering visualization only for embeddings data

Cluster

4



32



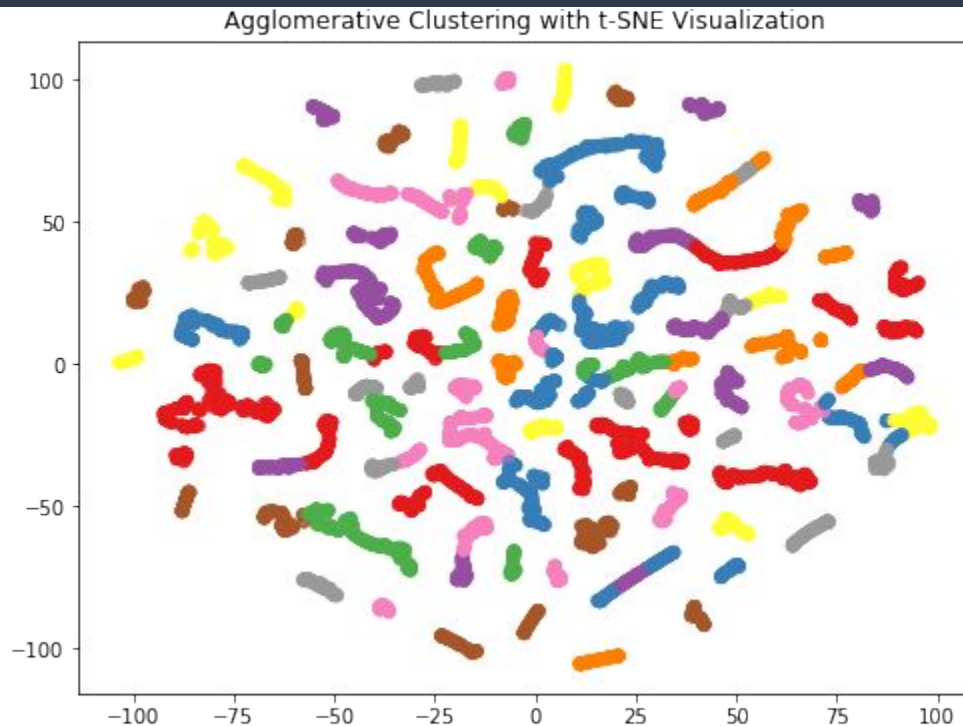
65



78



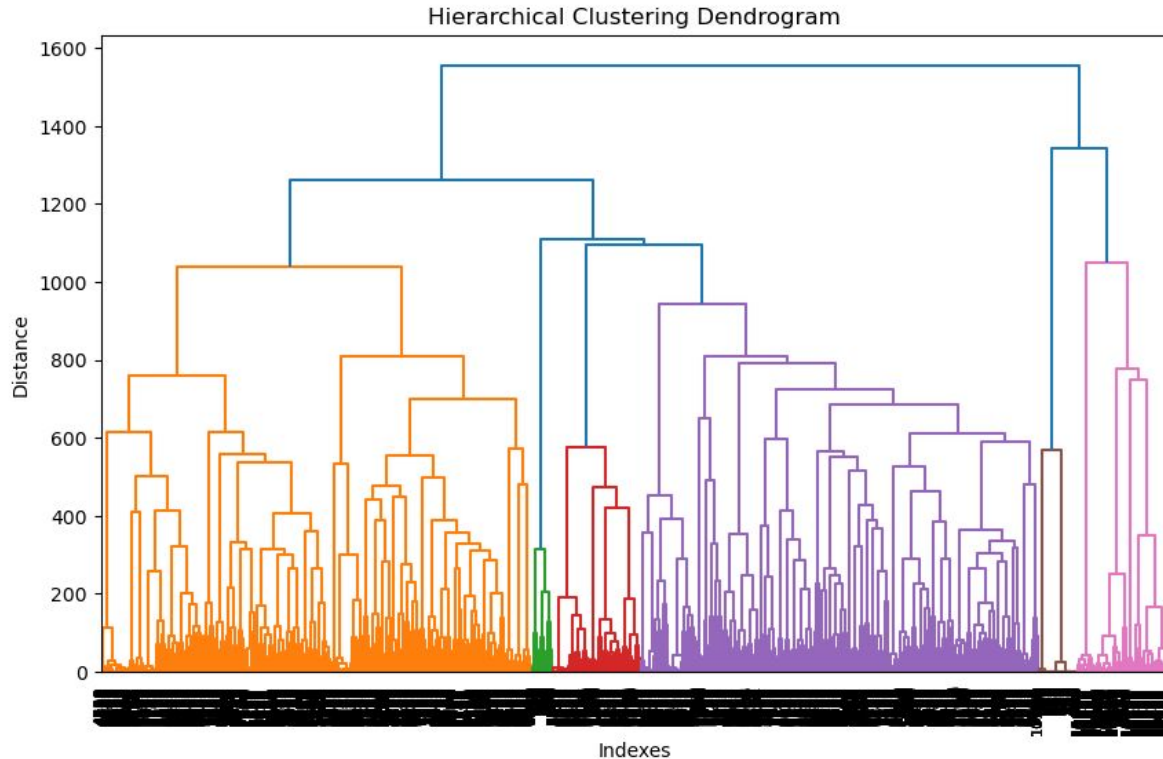
## 2. Images– Clustering Visualization with t-sne



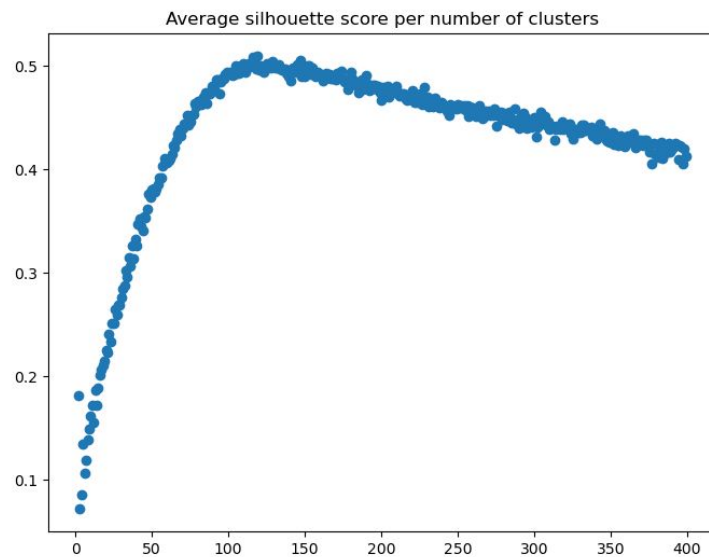
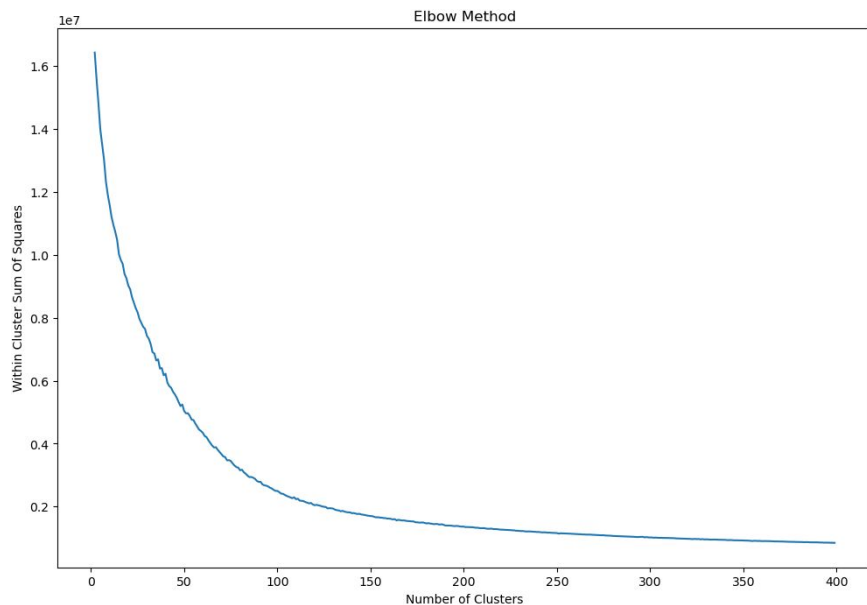
### 3. Skeletons + Images : Dataset Creation

$$\begin{array}{c} \text{Cluster} \end{array} \begin{array}{cccccccc} 0 & 1 & 2 & 3 & 4 & 5 & 6 & 7 \end{array}$$
$$\text{Descriptor} = \begin{bmatrix} 2 & 1 & 0 & 0 & 0 & 2 & 0 & 0 \\ 1 & 0 & 0 & 1 & 1 & 1 & 0 & 0 \\ 1 & 0 & 0 & 1 & 0 & 1 & 0 & 2 \\ 2 & 0 & 0 & 0 & 0 & 2 & 0 & 0 \\ \vdots & \vdots & \vdots & \vdots & \vdots & \vdots & \vdots & \vdots \end{bmatrix}, \quad \text{Descriptor} \in \mathbb{R}^{n \times 8}$$

### 3. Skeletons + Images : Hierarchical Clustering – Ward Method



### 3. Skeletons + Images– Number of clusters



# Clustering visualization for skeleton+embeddings data

Cluster

4



32



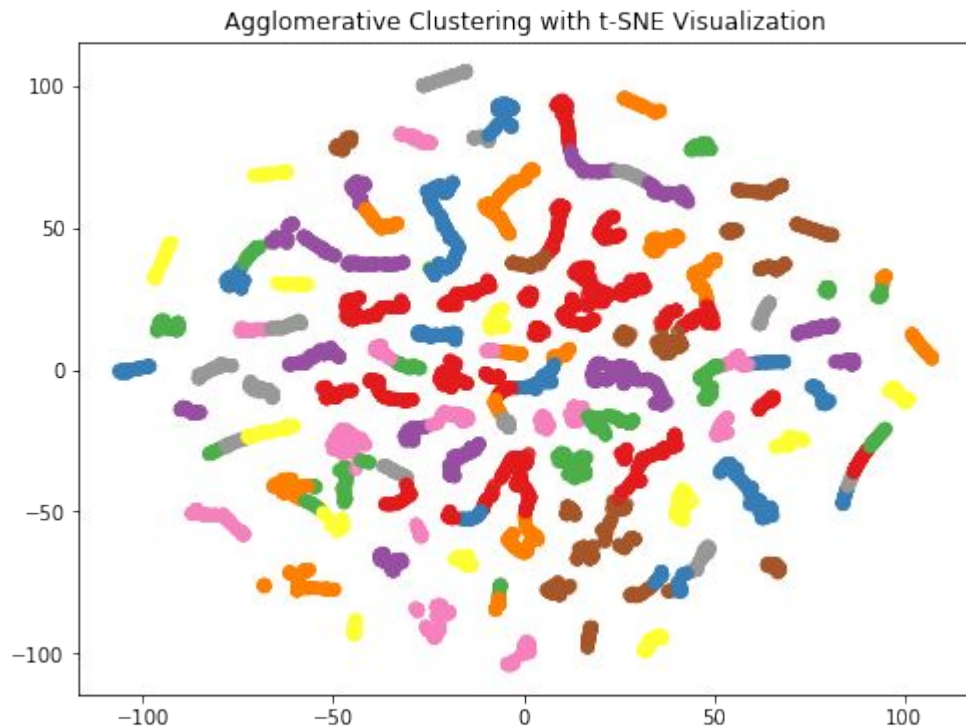
65



81

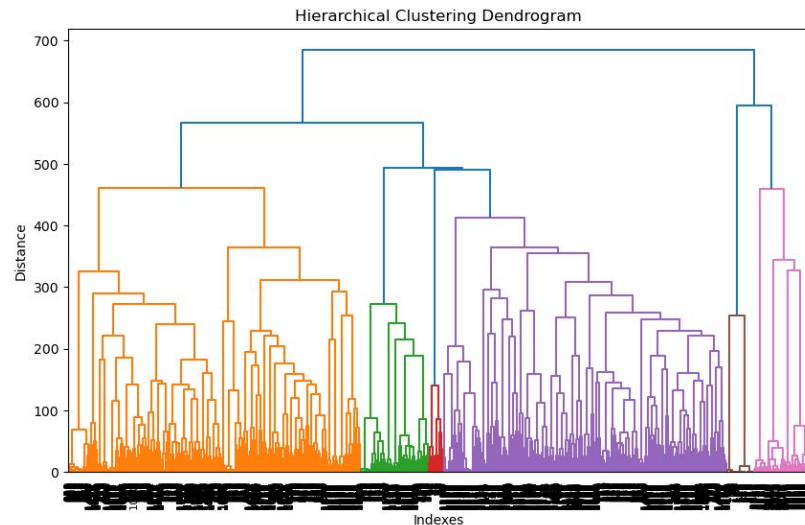
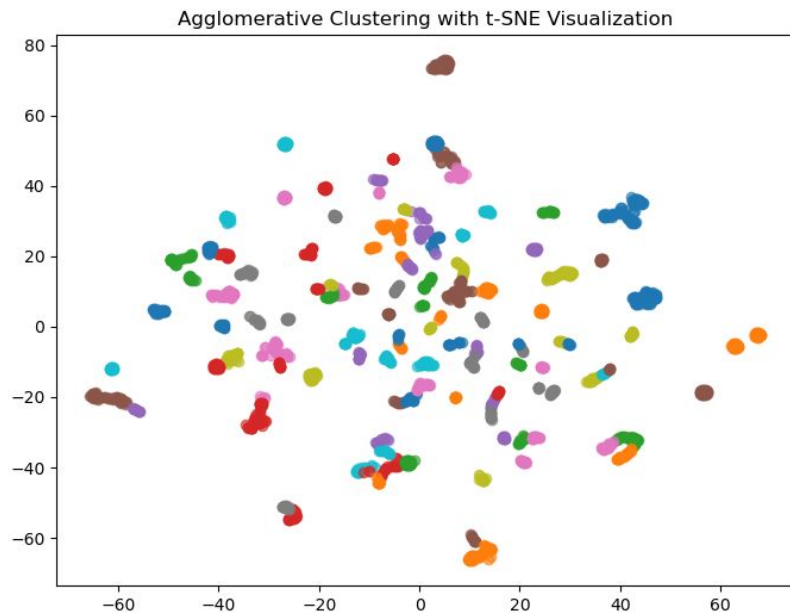


### 3. Skel + Images – Clustering Visualization with t-sne





# Sampling every 5 frames





# Sampling every 5 frames

Cluster

34



44



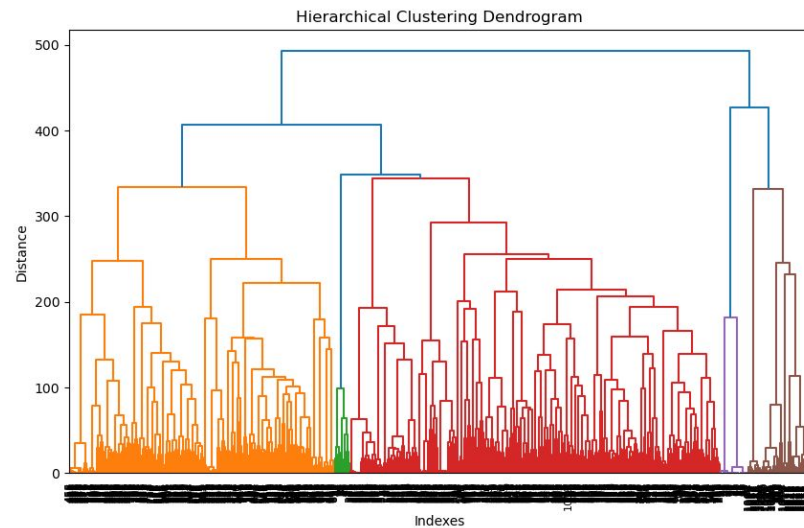
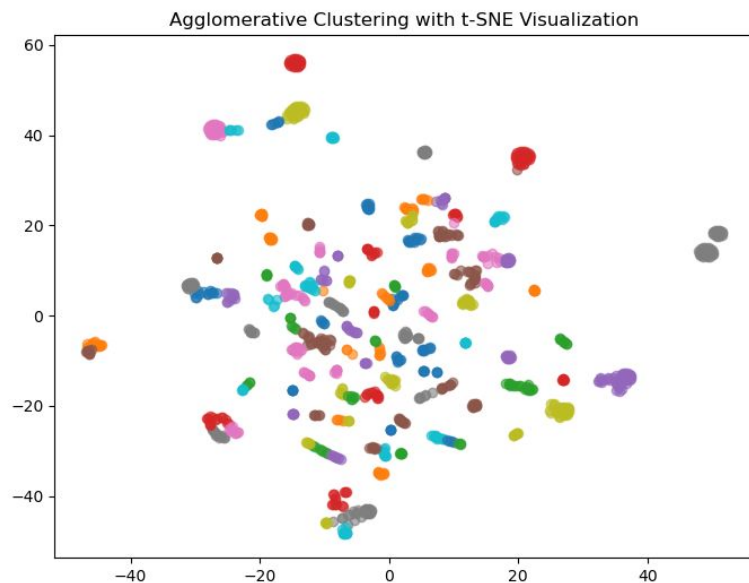
61



75



# Sampling every 10 frames



# Sampling every 10 frames

Cluster

25



55



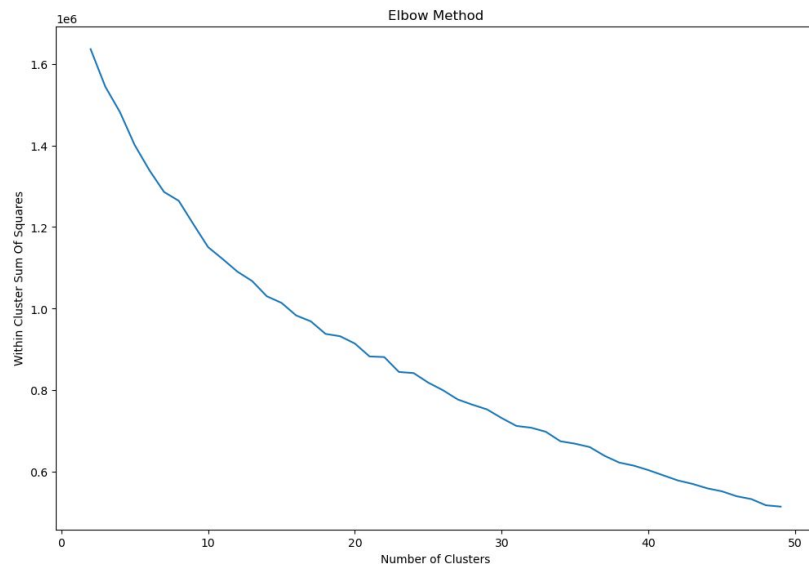
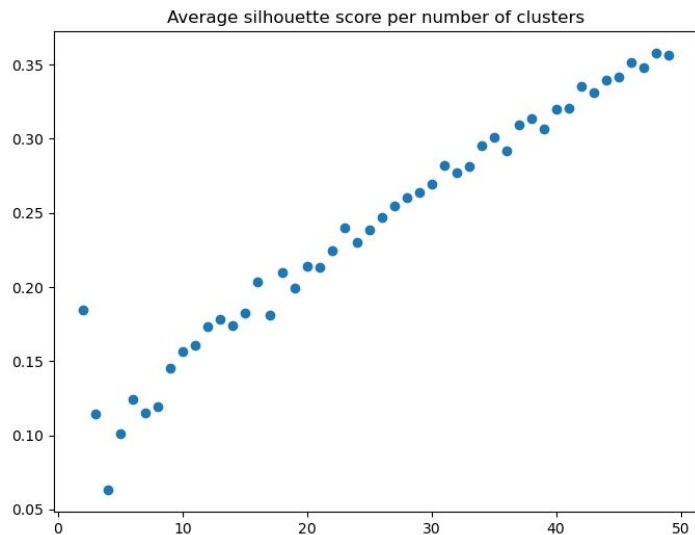
57



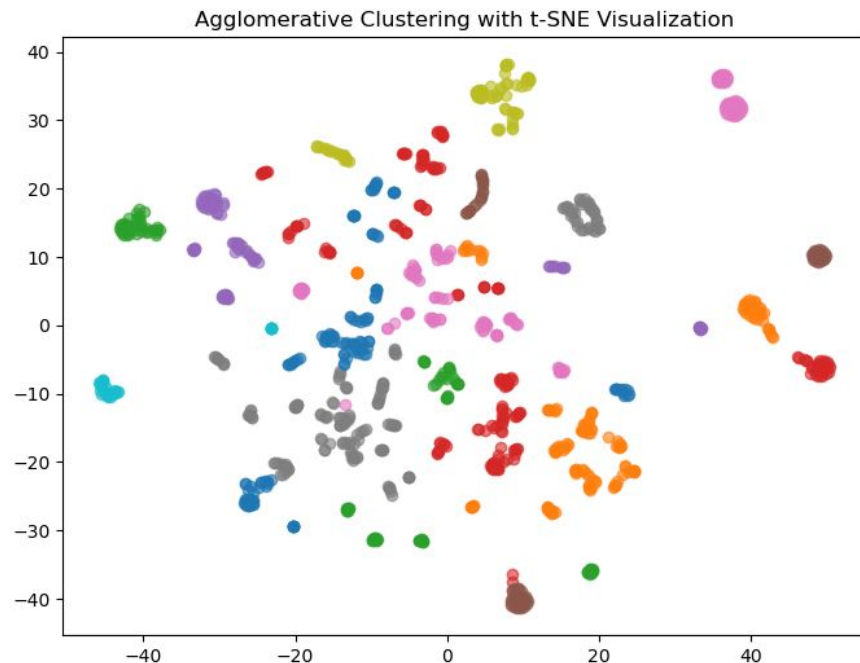
67



# Sampling every 10 frames with fewer clusters



# Sampling every 10 frames with 27 clusters





# Sampling every 10 frames with 27 clusters

Cluster

9



10



16



17



# Conclusions



- Complete pipeline with really satisfactory results for the 3 datasets considered
- There are several methods performed throughout the pipeline which will have a heavy impact on the results

# Future work



- Classify frames based on skeleton data alone
- Test different methods for each pipeline module
- Add extra information to the skeleton and embeddings data
- Further reduce the image embeddings dimensionality



# References

- Udell et al., *Generalized low rank models* 2016
- Scikit-learn: Machine Learning in Python, Pedregosa et al., JMLR 12, pp. 2825-2830, 2011.
- van der Maaten, Laurens & Hinton, Geoffrey. (2008). Visualizing data using t-SNE. Journal of Machine Learning Research. 9. 2579-2605.
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