



## Data Analysis & Visualisation

CSC3062

BEng (CS & SE), MEng (CS & SE), BIT & CIT

Dr Reza Rafiee

Semester 1 2019



#### **Unsupervised clustering**

# Unsupervised learning

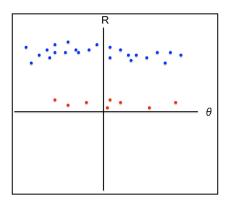
#### K-means clustering

#### Basic algorithm:

- Step 1: select k (number of clusters)
- Step 2: randomly select k initial cluster centers
- Step 3: calculate distance from each data point to each cluster center
  - What type of distance should we use? E.g., Euclidean distance
- Step 4: assign each data point to the closest cluster center (centroid)
- Step 5: calculate new centroids as the mean of the data points that belong to the centroid of the previous step
- Repeat Step 3-5 until a final stop condition (if no data point was reassigned then stop).

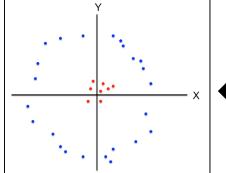
#### K-means clustering

- Strengths
  - Simple & fast and can be applied to high-dimensional large data
  - Finds cluster centres that minimize conditional variance (good representation of data)
  - Easy to implement
- Weaknesses
  - Need to choose k
  - Sensitive to outliers
  - Prone to local minima and no guarantee of optimal solution (local optima)
    - Repeat with different starting values
  - Difficult to guess the correct "k"



Changing features & distance function





K-means algorithm is not able to properly cluster this data points



Assume, we are given a dataset for the purpose of clustering analysis



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- Evaluate your dataset from different aspects
  - What type of features (e.g., numeric or categorical)?
  - The size of the dataset (e.g., large or small)
  - Number of feature (i.e., attributes), Is it a high-dimensional dataset?
  - Assessing <u>outliers</u> and missing
- Consider consensus clustering
- Evaluate the reliability (i.e., consistency/robust) of the clustering result



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#### Consensus clustering

Assume, we are given a dataset for the purpose of clustering analysis

- 1) No knowledge about the number of clusters
- Clustering methods are sensitive to initialisation settings
- The lack of a reliable validation technique when using clustering
  - a) We need a measure of confidence for cluster numbers and cluster assignment

#### Consensus clustering<sup>1</sup>

Assume, we are given a dataset for the purpose of clustering analysis

- 1) Multiple runs of a clustering algorithm
  - a) Determine the number of clusters and assess the stability of the discovered clusters
  - b) In k-means clustering: with using random restart

2) Aggregating the cluster (label) results of different clustering algorithms

<sup>&</sup>lt;sup>1</sup> Ensemble clustering

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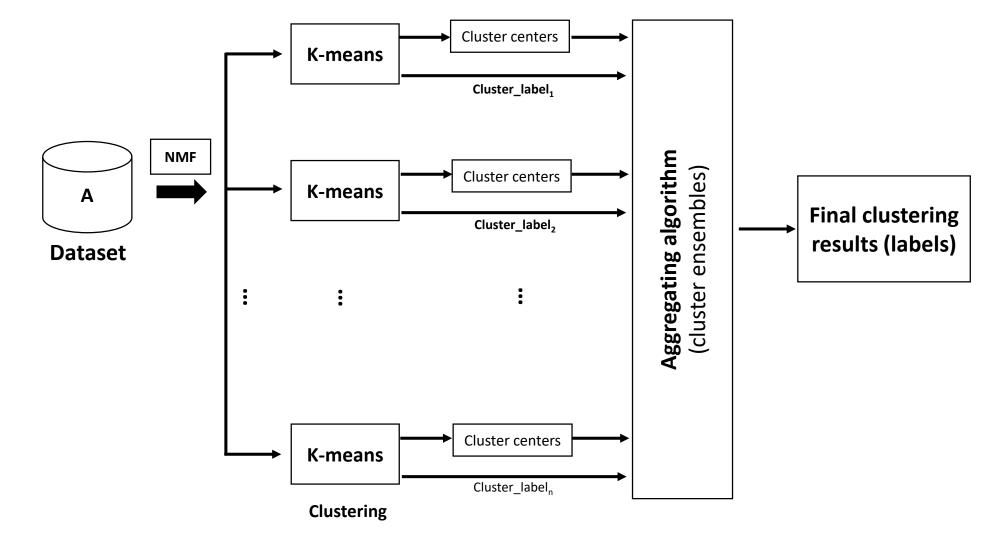
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#### Consensus approach

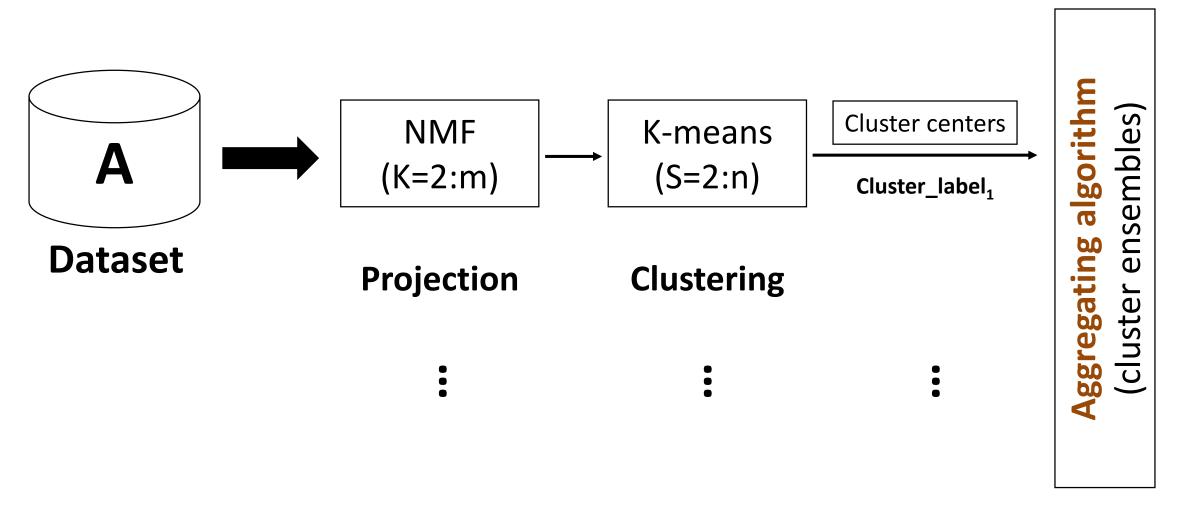
#### 1) Multiple runs of a clustering algorithm



A comprehensive Ensemble approach for unsupervised clustering using NMF projection and k-means clustering

#### Consensus approach

A comprehensive Ensemble approach for unsupervised clustering using NMF projection and k-means clustering



The value of m is dependent on the number of input features

#### n=12 samples with 4 subgroups

After running NMF on our input dataset  $17 \times 220 ===> k=4$ , H matrix Consider only 12 samples of H matrix (for the sake of simplicity)

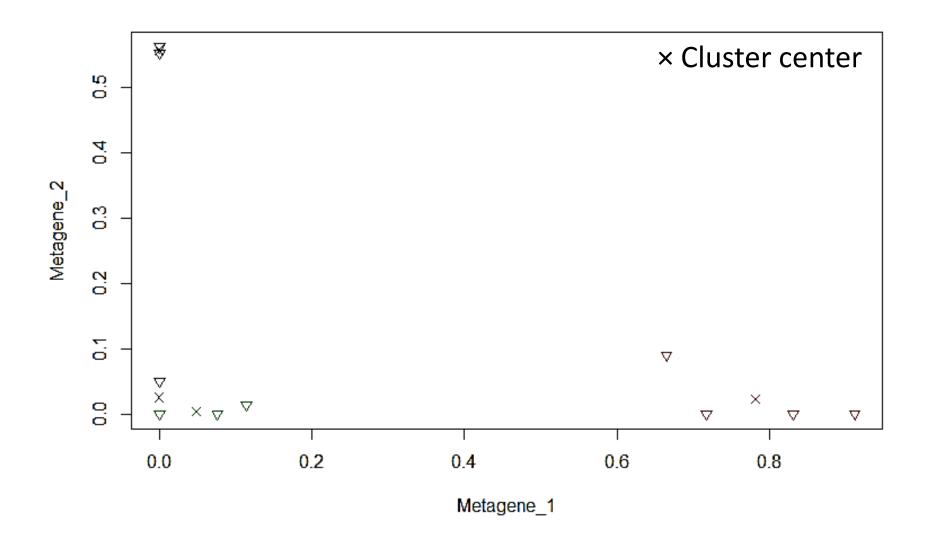
#### n=12 samples with 4 subgroups

<b>^</b>	CSC3062_108_2 <sup>‡</sup>	CSC3062_109_4 <sup>‡</sup>	CSC3062_110_4 <sup>‡</sup>	CSC3062_112_2 <sup>‡</sup>	CSC3062_783_3 <sup>‡</sup>	CSC3062_145_3 <sup>‡</sup>
Metagene_1	1.145277e-01	1.916895e-50	2.654951e-40	7.633172e-02	3.608274e-32	7.042284e-28
Metagene_2	1.338042e-02	5.529235e-01	5.625382e-01	4.172066e-27	5.022959e-02	1.881889e-05
Metagene_3	5.842943e-19	5.115138e-43	1.629874e-28	2.634450e-34	6.117725e-01	6.623634e-01
Metagene_4	9.603256e-01	2.808713e-27	4.787113e-29	9.671474e-01	1.660626e-34	5.350906e-39

•	CSC3062_649_1 <sup>‡</sup>	CSC3062_115_1 <sup>‡</sup>	CSC3062_670_2 <sup>‡</sup>	CSC3062_50080_1 <sup>‡</sup>	CSC3062_436_1 <sup>‡</sup>	CSC3062_674_2 <sup>‡</sup>
Metagene_1	7.176776e-01	9.121094e-01	2.142412e-28	8.314318e-01	6.650897e-01	1.424858e-17
Metagene_2	0.000000e+00	1.312099e-40	2.695954e-17	1.158338e-18	8.997966e-02	3.280249e-12
Metagene_3	1.759033e-70	3.300750e-21	3.208493e-17	1.691378e-40	3.382756e-17	2.059872e-02
Metagene_4	6.929525e-63	3.516017e-59	9.679785e-01	4.684605e-20	1.916895e-23	1.000000e+00

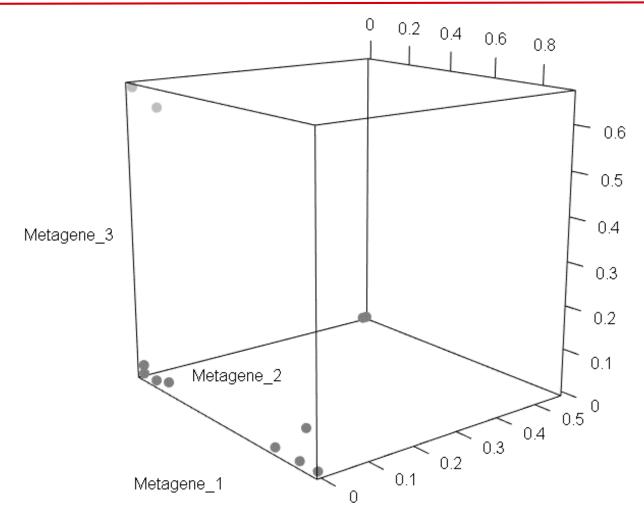
#### After running k-means on these 12 samples

K\_means\_Model <- kmeans(t(Small\_dataset\_cluster\_analysis\_0To1),centers = 4, iter.max = 50,nstart = 5) #</pre>





#### Using plot3d() to visualise samples



Visualising the H matrix using PCA

#### Several runs of k-means

```
# Creating a matrix of all labels of different k-means runnings

Matrix_labels_different_runs <- matrix(nrow = ncol(Small_dataset_cluster_analysis_0To1), ncol = 10,0)
rownames(Matrix_labels_different_runs) <- colnames(Small_dataset_cluster_analysis_0To1)
for (j in 1:10) {
    K_means_Model <- kmeans(t(Small_dataset_cluster_analysis_0To1),centers = 4, iter.max = 50,nstart = 5) #
    #trying several random starts (nstart> 1) is often recommended.
    Matrix_labels_different_runs[,j] <- K_means_Model$cluster
} # for

plot(t(Small_dataset_cluster_analysis_0To1), col = K_means_Model$cluster,pch=6)
points(K_means_Model$centers, col = 1:3, pch = 4, cex = 1)</pre>
```



#### Several runs of k-means

#### Alternative sample names

<b>↓</b> [	_	V1 <sup>‡</sup>	V2 <sup>‡</sup>	V3 <sup>‡</sup>	V4 <sup>‡</sup>	V5 <sup>‡</sup>	V6 <sup>‡</sup>	V7 <sup>‡</sup>	V8 <sup>‡</sup>	V9 <sup>‡</sup>	V10 <sup>‡</sup>
<b>S1</b>	CSC3062_108_2	4	4	4	1	3	1	3	3	4	4
<b>S2</b>	CSC3062_109_4	1	1	1	2	2	2	4	1	1	2
<b>S3</b>	CSC3062_110_4	1	1	1	2	2	2	4	1	1	2
<b>S4</b>	CSC3062_112_2	4	4	4	1	3	1	3	3	4	4
<b>S5</b>	CSC3062_783_3	2	3	3	3	4	3	1	1	3	2
<b>S6</b>	CSC3062_145_3	2	3	3	3	4	3	1	1	3	2
<b>S7</b>	CSC3062_649_1	3	2	2	4	1	4	2	4	2	1
<b>S8</b>	CSC3062_115_1	3	2	2	4	1	4	2	2	2	3
<b>S9</b>	CSC3062_670_2	4	4	4	1	3	1	3	3	4	4
S10	CSC3062_50080_1	3	2	2	4	1	4	2	2	2	3
<b>S11</b>	CSC3062_436_1	3	2	2	4	1	4	2	4	2	1
<b>S12</b>	CSC3062_674_2	4	4	4	1	3	1	3	3	4	4

$$V_1 = (4,1,1,4,2,2,3,3,4,3,3,4)$$

$$V_6 = (1,2,2,1,3,3,4,4,1,4,4,1)$$

$$V_2 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

$$V_7 = (3,4,4,3,1,1,2,2,3,2,2,3)$$

$$V_3 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

$$V_8 = (3,1,1,3,1,1,4,2,3,2,4,3)$$

$$V_4 = (1,2,2,1,3,3,4,4,1,4,4,1)$$

$$V_9 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

$$V_5 = (3,2,2,3,4,4,1,1,3,1,1,3)$$

$$V_{10} = (4,2,2,4,2,2,1,3,4,3,1,4)$$

Which clustering algorithms are creating same/similar cluster labels?

#### Which samples are not confidently clustered?

$$V_1 = (4,1,1,4,2,2,3,3,4,3,3,4)$$

$$V_6 = (1,2,2,1,3,3,4,4,1,4,4,1)$$

$$V_2 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

$$V_7 = (3,4,4,3,1,1,2,2,3,2,2,3)$$

$$V_3 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

$$V_8 = (3,1,1,3,1,1,4,2,3,2,4,3)$$

$$V_4 = (1,2,2,1,3,3,4,4,1,4,4,1)$$

$$V_9 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

$$V_5 = (3,2,2,3,4,4,1,1,3,1,1,3)$$

$$V_{10} = (4,2,2,4,2,2,1,3,4,3,1,4)$$

Clusterings V2 and V3 are identical.

$$V_1 = (4,1,1,4,2,2,3,3,4,3,3,4)$$

$$V_6 = (1,2,2,1,3,3,4,4,1,4,4,1)$$

$$V_2 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

$$V_7 = (3,4,4,3,1,1,2,2,3,2,2,3)$$

$$V_3 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

$$V_8 = (3,1,1,3,1,1,4,2,3,2,4,3)$$

$$V_4 = (1,2,2,1,3,3,4,4,1,4,4,1)$$

$$V_9 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

$$V_5 = (3,2,2,3,4,4,1,1,3,1,1,3)$$

$$V_{10} = (4,2,2,4,2,2,1,3,4,3,1,4)$$

Clusterings V1 and V2 are logically identical.

$$V_1 = (4,1,1,4,2,2,3,3,4,3,3,4)$$
  $V_6 = (1,2,2,1,3,3,4,4,1,4,4,1)$   $V_2 = (4,1,1,4,3,3,2,2,4,2,2,4)$   $V_7 = (3,4,4,3,1,1,2,2,3,2,2,3)$   $V_8 = (4,1,1,4,3,3,2,2,4,2,2,4)$   $V_9 = (4,1,1,4,3,3,2,2,4,2,2,4)$   $V_9 = (4,1,1,4,3,3,2,2,4,2,2,4)$   $V_9 = (4,1,1,4,3,3,2,2,4,2,2,4)$   $V_9 = (4,2,2,4,2,2,1,3,4,3,1,4)$ 

Clusterings V1, V2, V4, V5, V6, V7 and V9 are logically identical.



$$V_1 = (4,1,1,4,2,2,3,3,4,3,3,4)$$

$$V_6 = (1,2,2,1,3,3,4,4,1,4,4,1)$$

$$V_2 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

$$V_7 = (3,4,4,3,1,1,2,2,3,2,2,3)$$

$$V_3 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

$$V_8 = (3, 1, 1, 3, 1, 1, 4, 2, 3, 2, 4, 3)$$

$$V_4 = (1,2,2,1,3,3,4,4,1,4,4,1)$$

$$V_9 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

$$V_5 = (3,2,2,3,4,4,1,1,3,1,1,3)$$

$$V_{10} = (4, 2, 2, 4, 2, 2, 1, 3, 4, 3, 1, 4)$$

#### What about V8 and V10?

Clusterings V8 and V10 are also logically identical.

$$V_1 = (4,1,1,4,2,2,3,3,4,3,3,4)$$

$$V_2 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

$$V_3 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

$$V_4 = (1,2,2,1,3,3,4,4,1,4,4,1)$$

$$V_5 = (3,2,2,3,4,4,1,1,3,1,1,3)$$

$$V_6 = (1,2,2,1,3,3,4,4,1,4,4,1)$$

$$V_7 = (3,4,4,3,1,1,2,2,3,2,2,3)$$

$$V_9 = (4,1,1,4,3,3,2,2,4,2,2,4)$$



Clustering results in case A are logically identical.

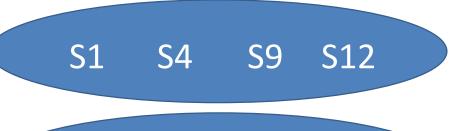
$$V_8 = (3,1,1,3,1,1,4,2,3,2,4,3)$$

$$V_{10} = (4,2,2,4,2,2,1,3,4,3,1,4)$$

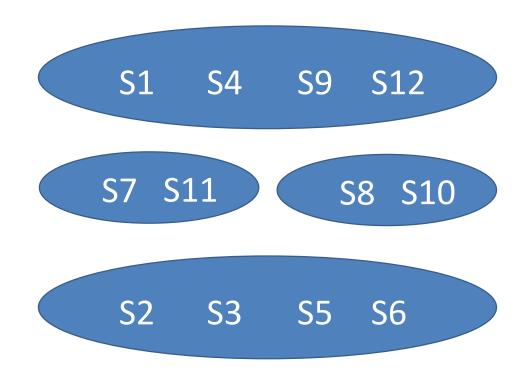


Clustering results in case B are logically identical.









$$V_8 = (3,1,1,3,1,1,4,2,3,2,4,3)$$
  
 $V_{10} = (4,2,2,4,2,2,1,3,4,3,1,4)$ 

$$A = \{V_1, V_2, V_3, V_4, V_5, V_6, V_7, V_9\}$$

$$B = \{V_8, V_{10}\}$$

A and B are two sets each including identical cluster labels or same number of clusters/groups

$$\Delta = \{V_1, V_2, V_3, V_4, V_5, V_6, V_7, V_8, V_9, V_{10}\}$$

 $\Delta$  is a set including all the iterations' cluster label vectors when running multiple k-means algorithms with different initialisation settings

$$P(S_i \in A) = ? where i \in \{1, ..., 12\}, A \subseteq \Delta$$

Probability that sample  $S_i$  belongs to set A

 $A \subseteq \Delta$ : A is a subset of  $\Delta$ 

$$P(S_i \in B) = ? where i \in \{1, ..., 12\}, B \subseteq \Delta$$

Probability that sample  $S_i$  belongs to set B

$$V_8 = (3,1,1,3,1,1,4,2,3,2,4,3)$$
  
 $V_{10} = (4,2,2,4,2,2,1,3,4,3,1,4)$ 

$$A = \{V_1, V_2, V_3, V_4, V_5, V_6, V_7, V_9\}$$

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A and B are two sets each including identical cluster labels or same number of clusters/groups

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$$P(S_i \in A) = \frac{8}{10} = 0.8 \text{ where } i \in \{1, ..., 12\}, A \subseteq \Delta$$

Probability that sample  $S_i$  belongs to set A

 $A \subseteq \Delta$ : A is a subset of  $\Delta$ 

$$P(S_i \in B) = \frac{2}{10} = 0.2 \text{ where } i \in \{1, ..., 12\}, B \subseteq \Delta$$

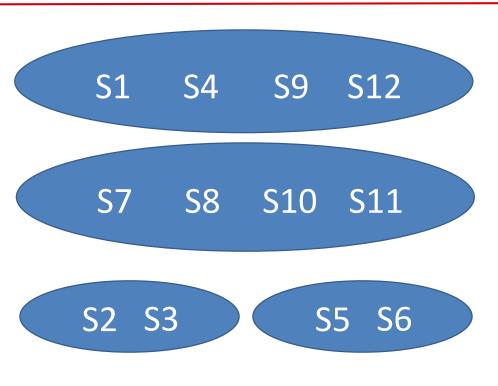
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$$V_8 = (3,1,1,3,1,1,4,2,3,2,4,3)$$
  
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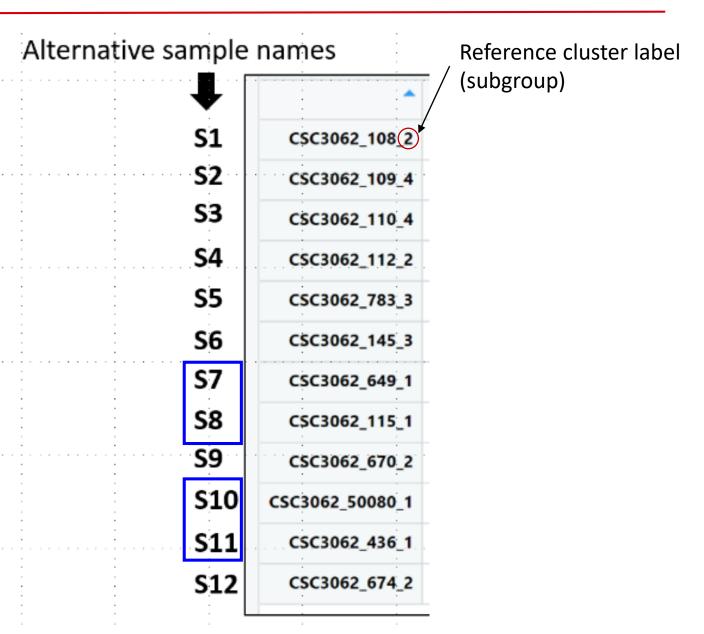
$$A = \{V_1, V_2, V_3, V_4, V_5, V_6, V_7, V_9\}$$

$$B = \{V_8, V_{10}\}$$





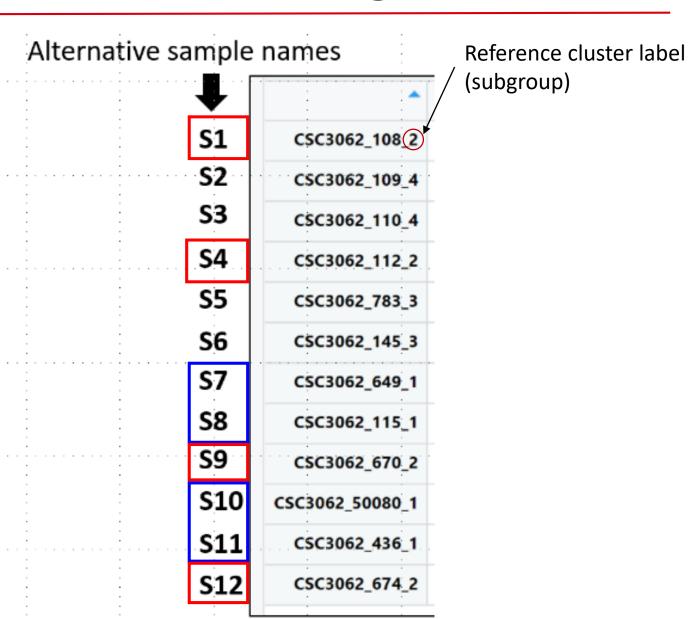
$$V_2 = (4,1,1,4,3,3,2,2,4,2,2,4)$$



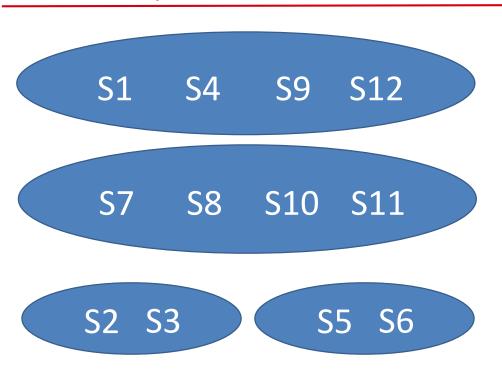




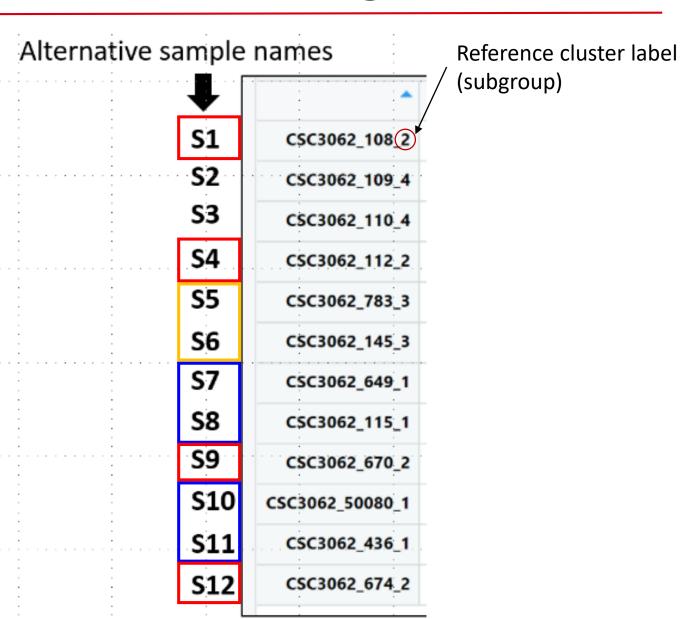
$$V_2 = (4,1,1,4,3,3,2,2,4,2,2,4)$$







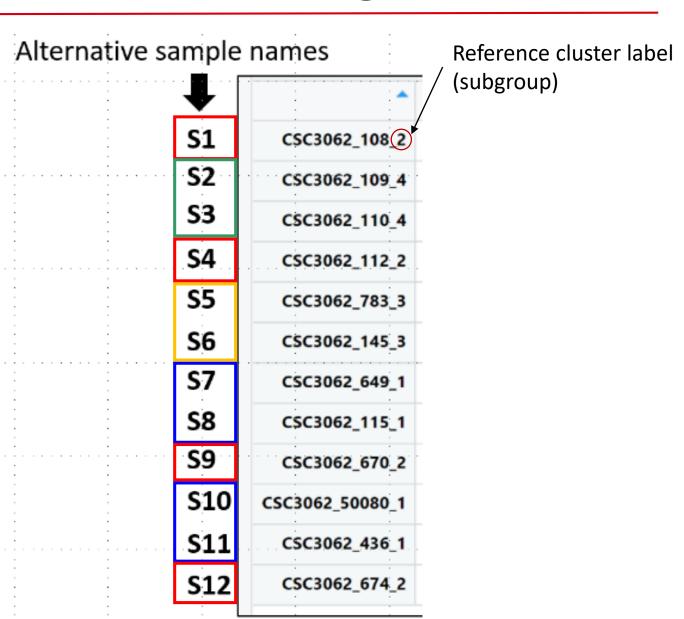
$$V_2 = (4,1,1,4,3,3,2,2,4,2,2,4)$$





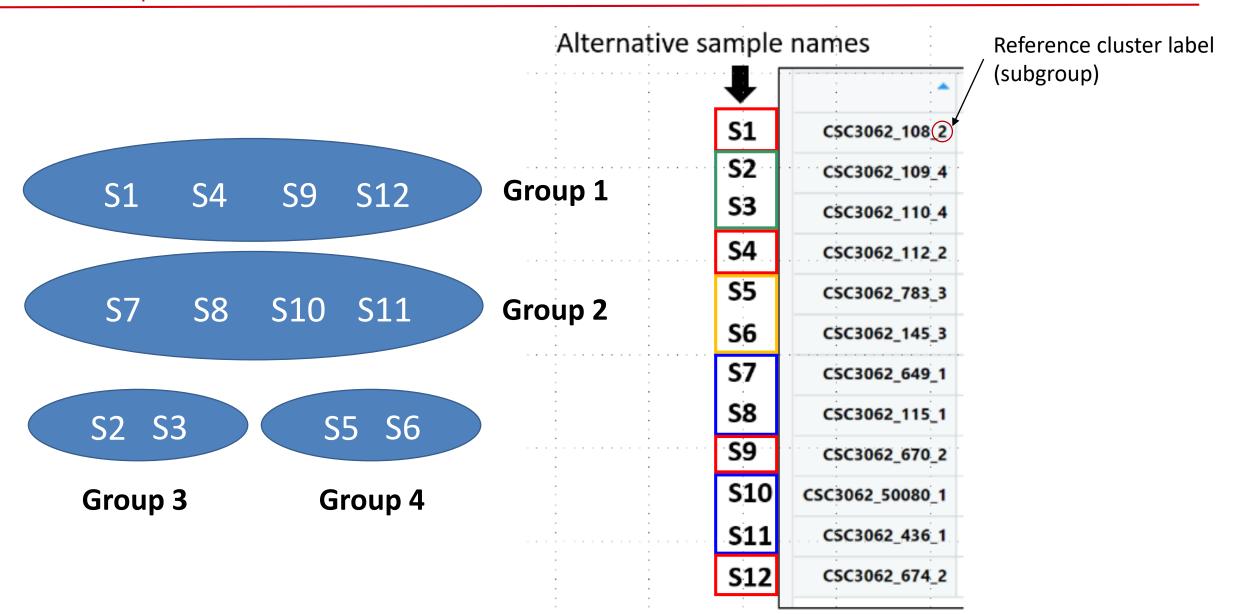


$$V_2 = (4,1,1,4,3,3,2,2,4,2,2,4)$$

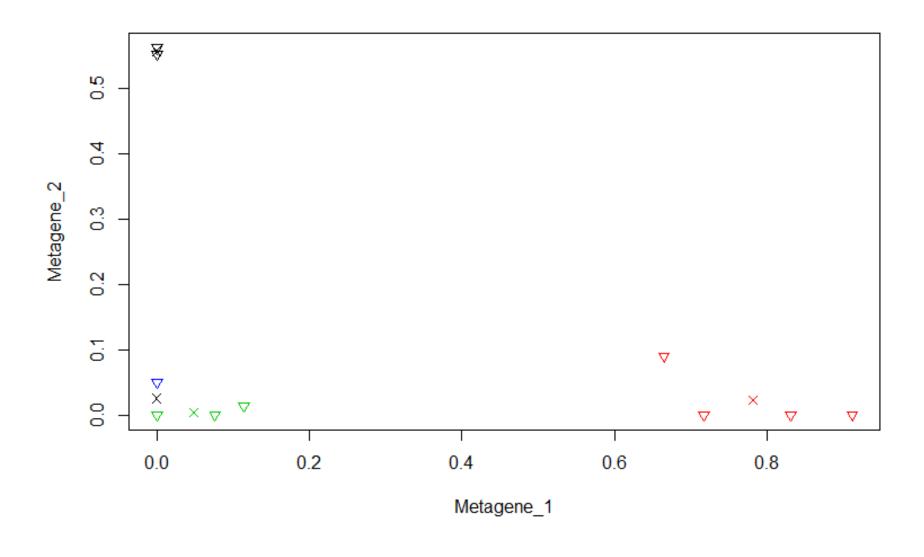




#### Final clustering result from consensus

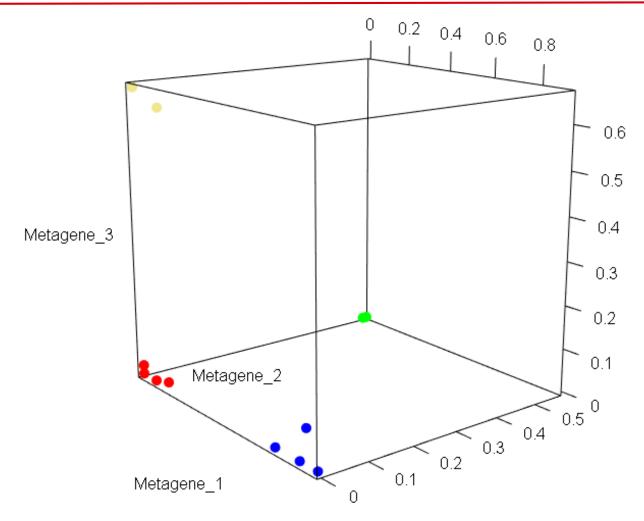


#### After running k-means on these 12 samples





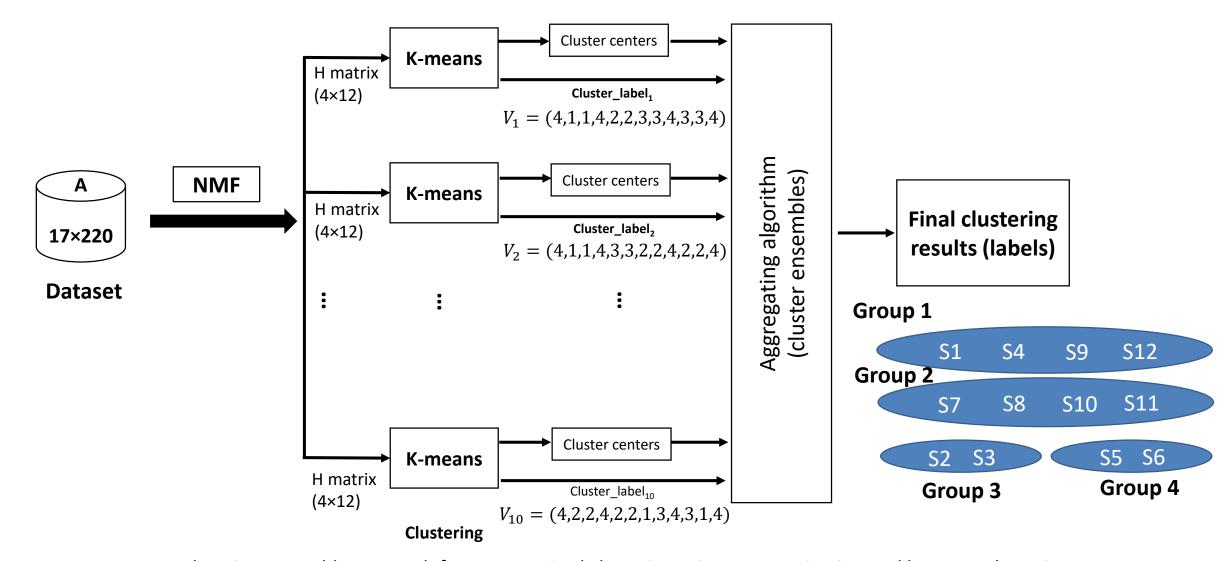
#### Using plot3d() to visualise samples



Visualising the original samples when they coloured by the subgroups obtained from final k-means clustering aggregation (consensus clustering)

#### Summary: consensus approach

#### 1) Multiple runs of a clustering algorithm



A comprehensive Ensemble approach for unsupervised clustering using NMF projection and k-means clustering



### Any Questions?