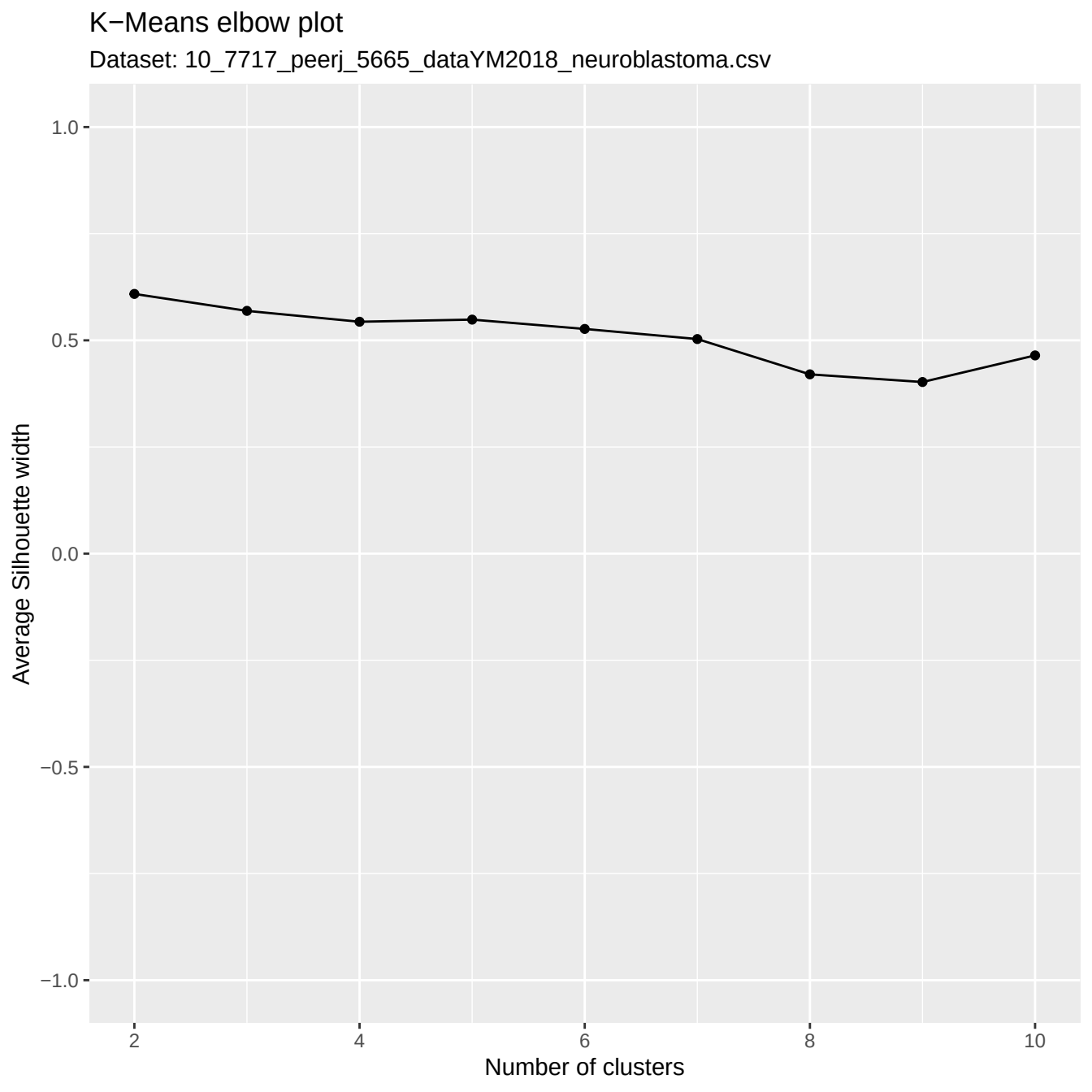
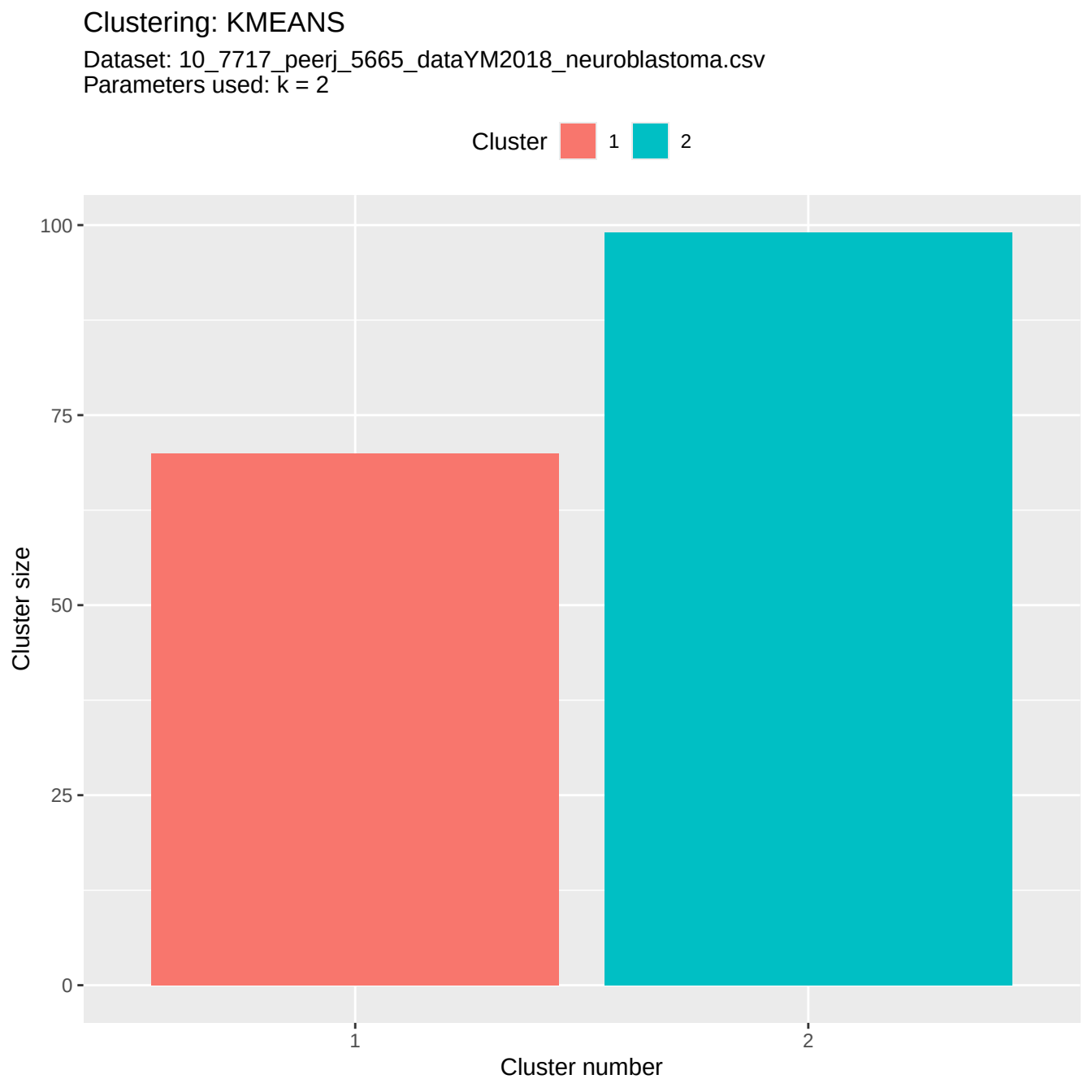
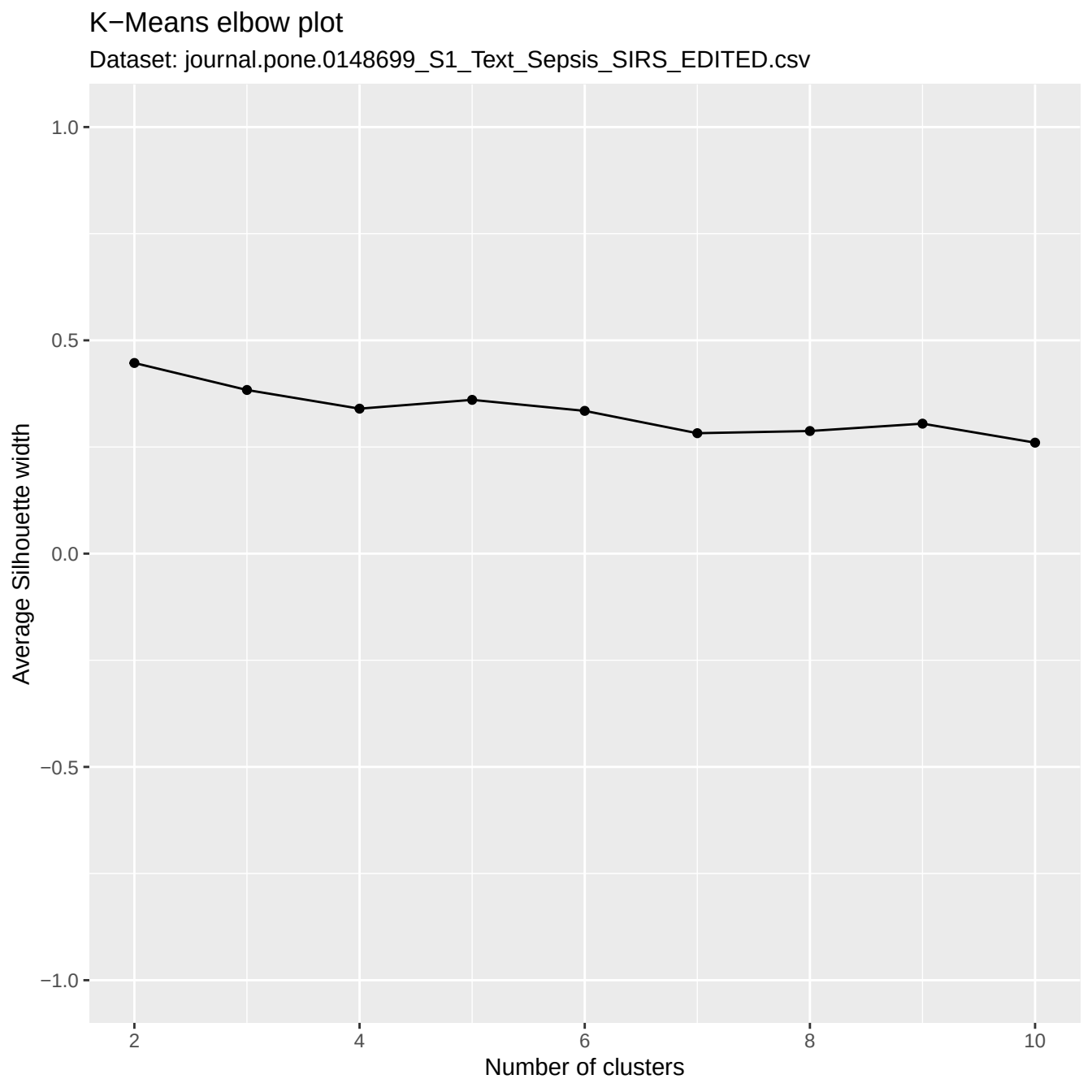
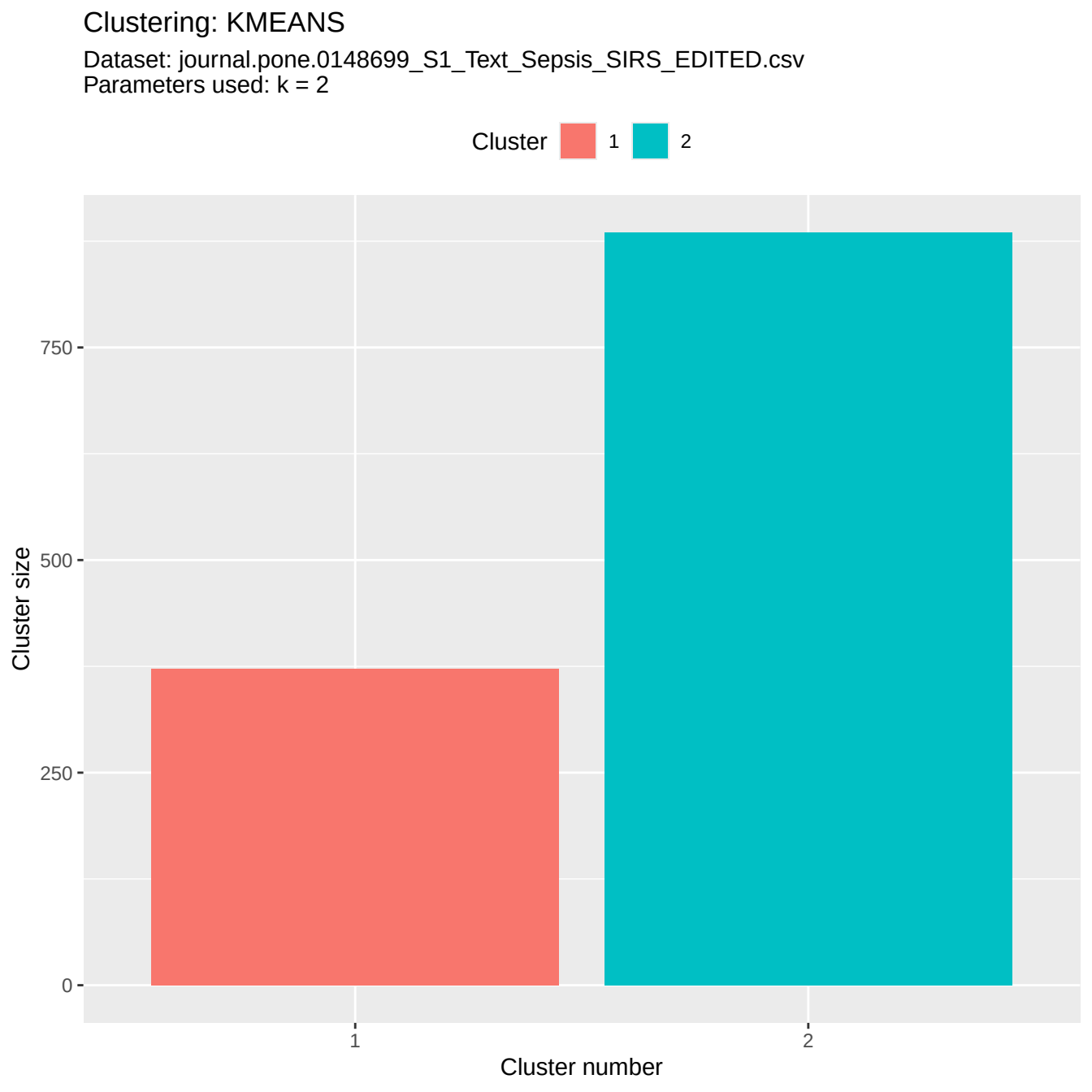
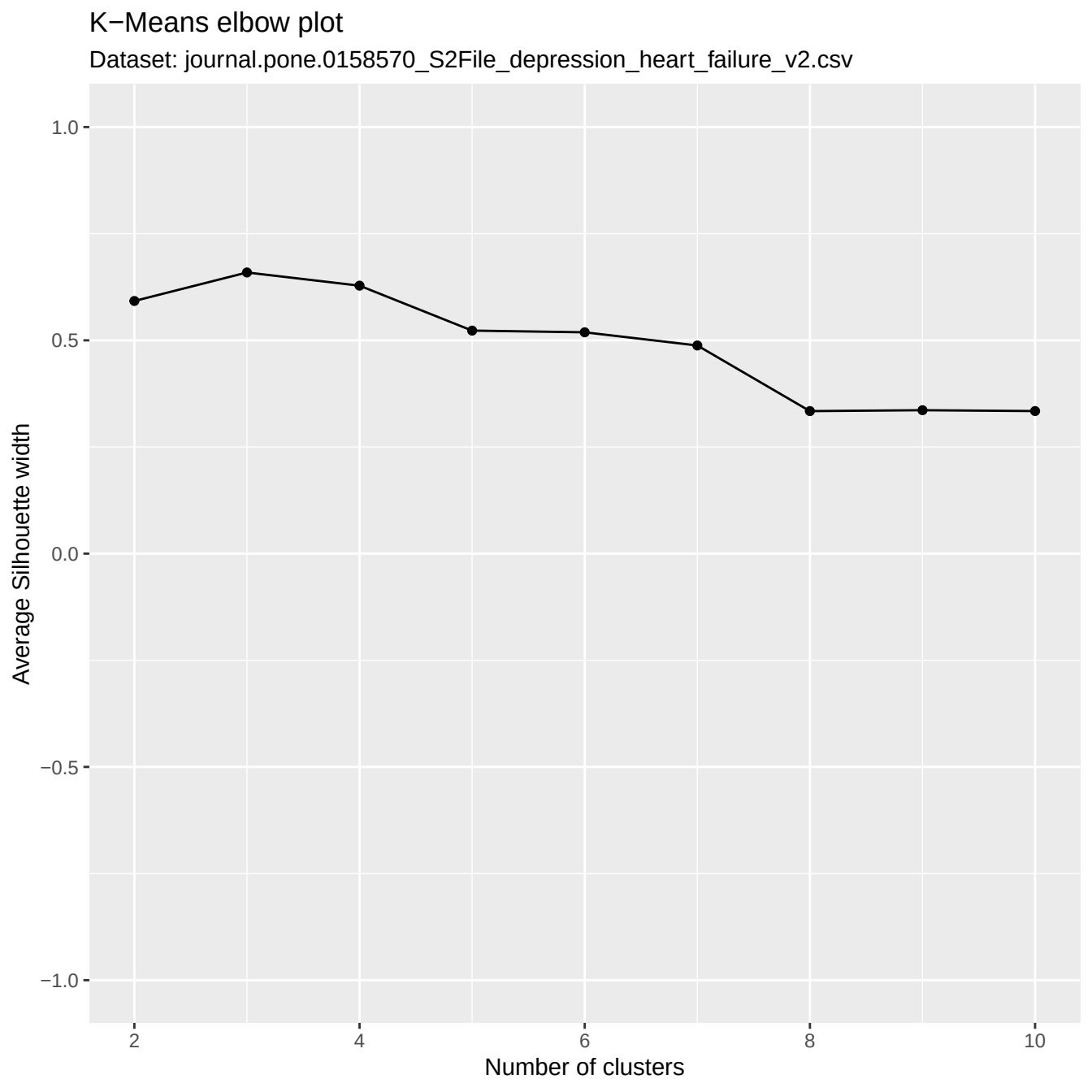
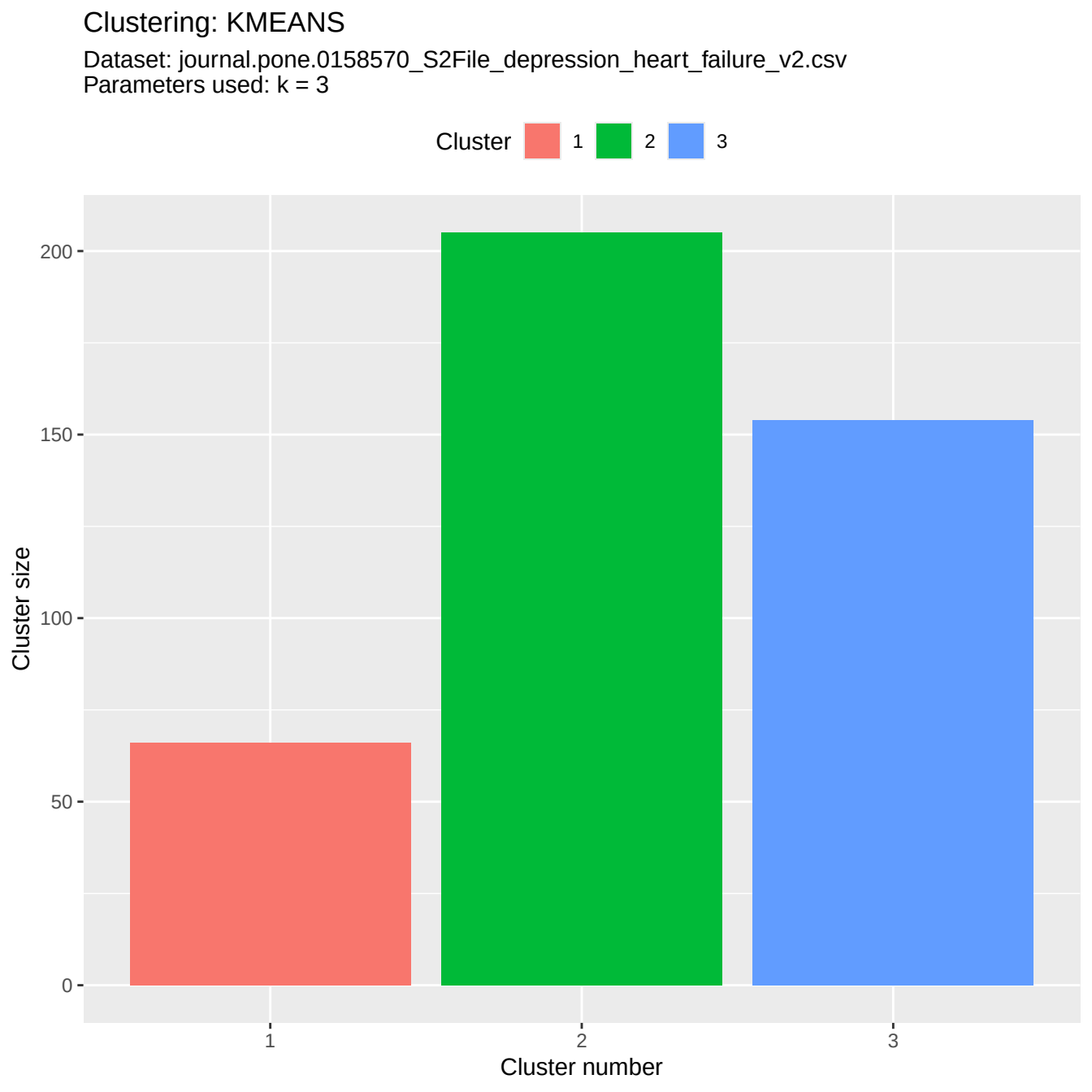

K−Means elbow plot
Dataset: 10_7717_peerj_5665_dataYM2018_neuroblastoma.csv
1.0
0.5
0.0
−0.5
−1.0
Average Silhouette width
2
4
6
8
10
Number of clusters

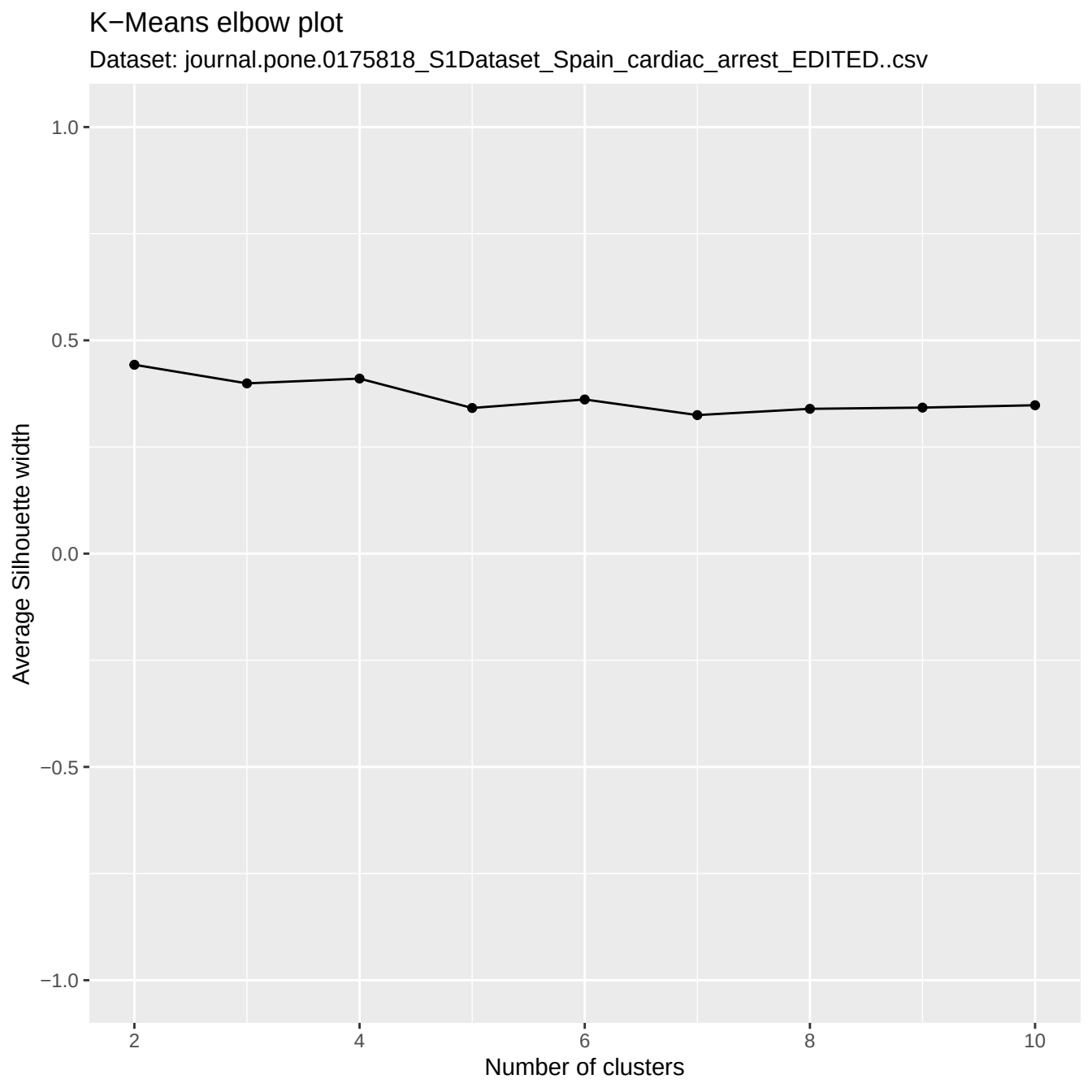

Clustering: KMEANS
Dataset: 10_7717_peerj_5665_dataYM2018_neuroblastoma.csv
Parameters used: k = 2
Cluster
1
2
100
75
50
25
0
Cluster size
1
2
Cluster number

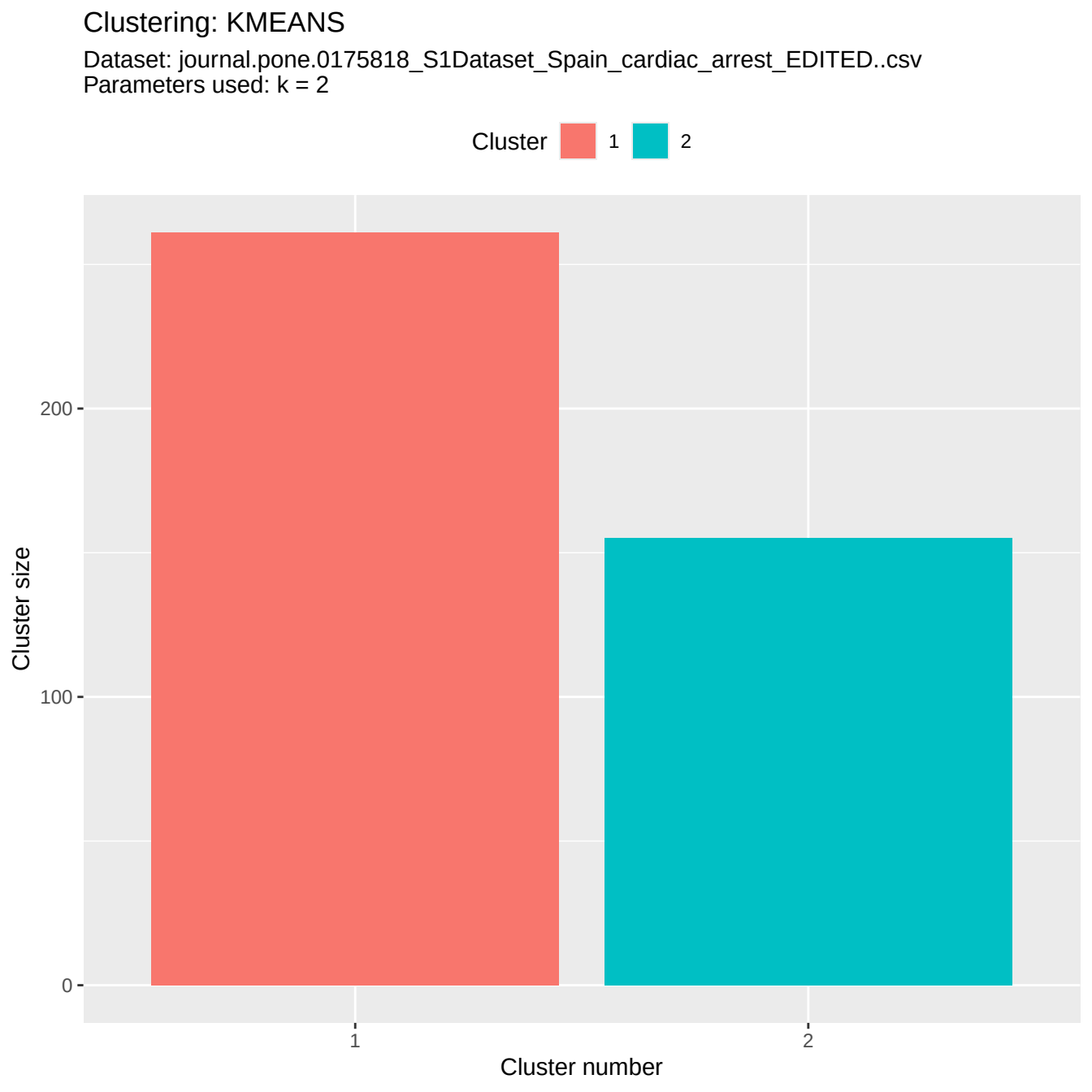

K−Means elbow plot
Dataset: journal.pone.0148699_S1_Text_Sepsis_SIRS_EDITED.csv
1.0
0.5
0.0
−0.5
−1.0
Average Silhouette width
2
4
6
8
10
Number of clusters

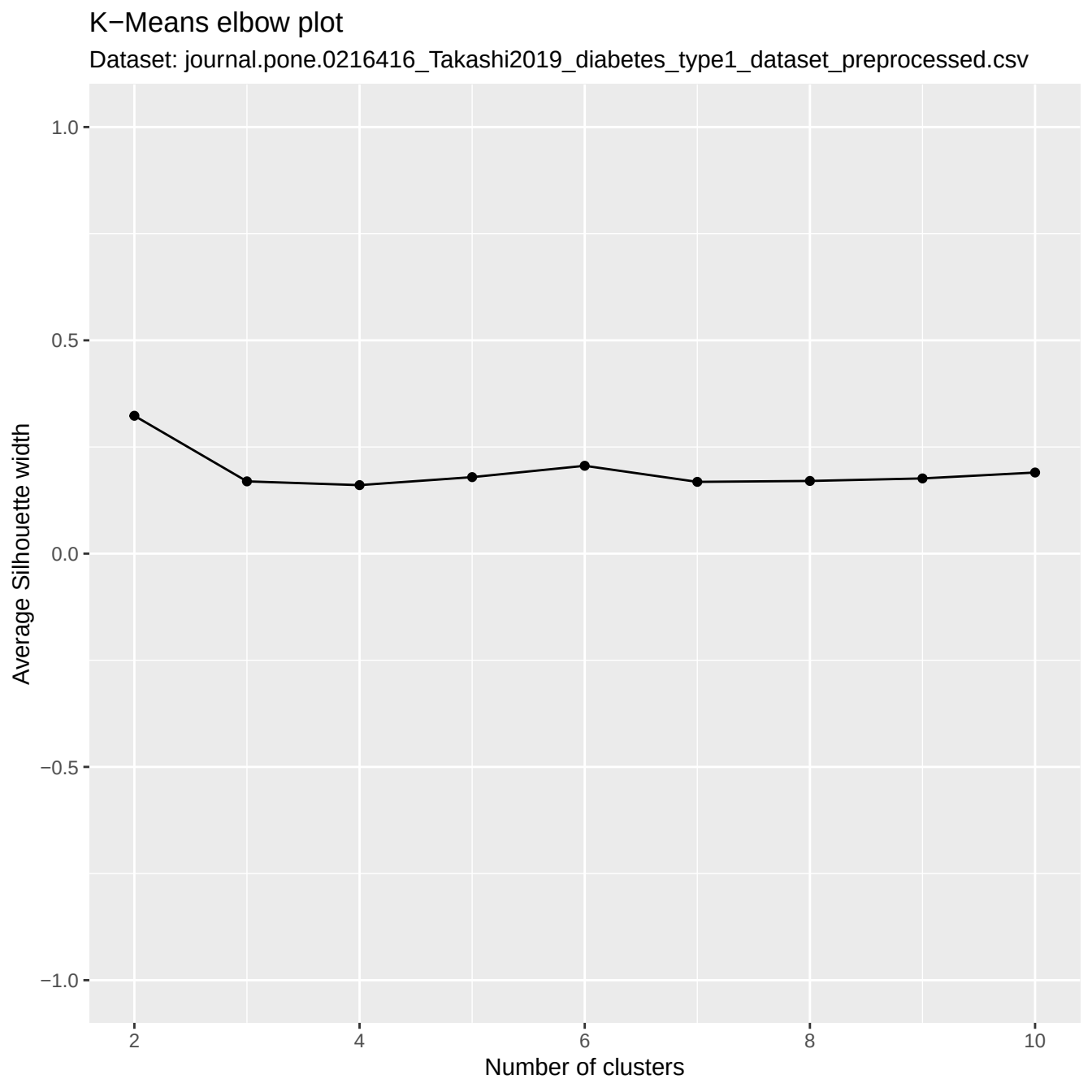

Clustering: KMEANS
Dataset: journal.pone.0148699_S1_Text_Sepsis_SIRS_EDITED.csv
Parameters used: k = 2
Cluster
1
2
750
500
250
0
Cluster size
1
2
Cluster number

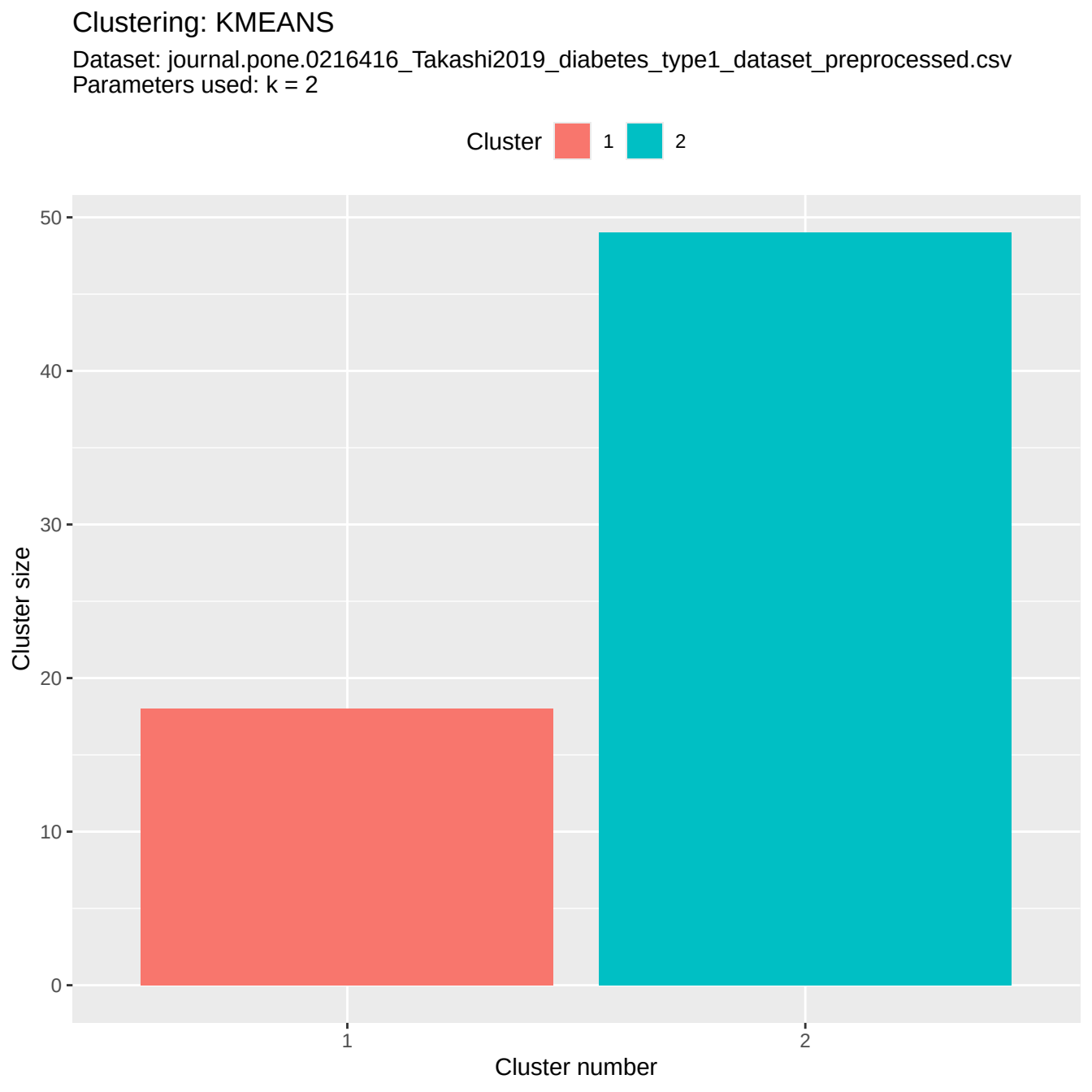

K−Means elbow plot
Dataset: journal.pone.0158570_S2File_depression_heart_failure_v2.csv
1.0
0.5
0.0
−0.5
−1.0
Average Silhouette width
2
4
6
8
10
Number of clusters


Clustering: KMEANS
Dataset: journal.pone.0158570_S2File_depression_heart_failure_v2.csv
Parameters used: k = 3
Cluster
1
2
3
200
150
100
50
0
Cluster size
1
2
3
Cluster number


K−Means elbow plot
Dataset: journal.pone.0175818_S1Dataset_Spain_cardiac_arrest_EDITED..csv
1.0
0.5
0.0
−0.5
−1.0
Average Silhouette width
2
4
6
8
10
Number of clusters


Clustering: KMEANS
Dataset: journal.pone.0175818_S1Dataset_Spain_cardiac_arrest_EDITED..csv
Parameters used: k = 2
Cluster
1
2
200
100
0
Cluster size
1
2
Cluster number


K−Means elbow plot
Dataset: journal.pone.0216416_Takashi2019_diabetes_type1_dataset_preprocessed.csv
Average Silhouette width
1.0
0.5
0.0
−0.5
−1.0
2
4
6
8
10
Number of clusters


Clustering: KMEANS
Dataset: journal.pone.0216416_Takashi2019_diabetes_type1_dataset_preprocessed.csv
Parameters used: k = 2
Cluster
1
2
50
40
30
20
10
0
Cluster size
1
2
Cluster number