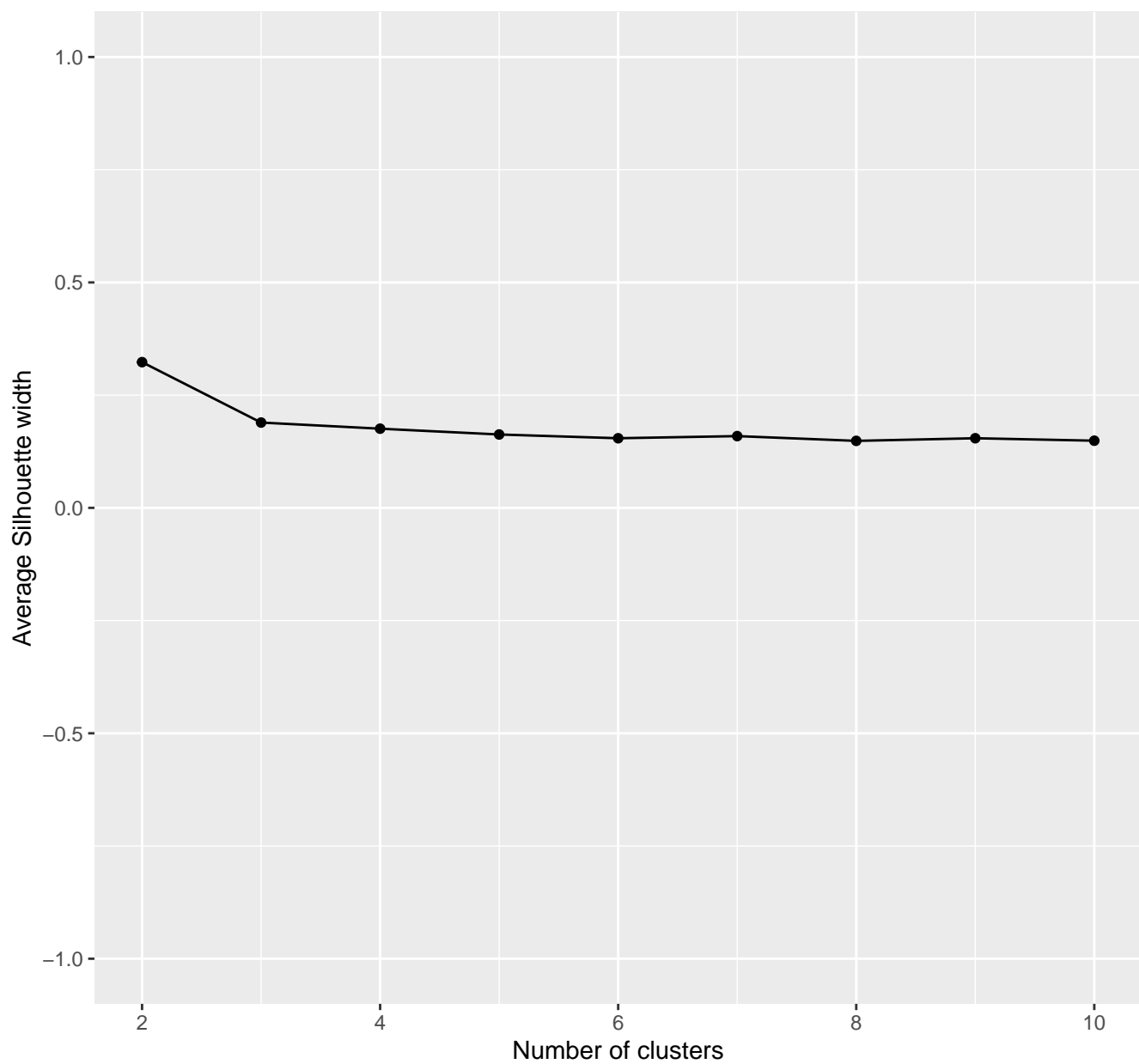


K-Means elbow plot

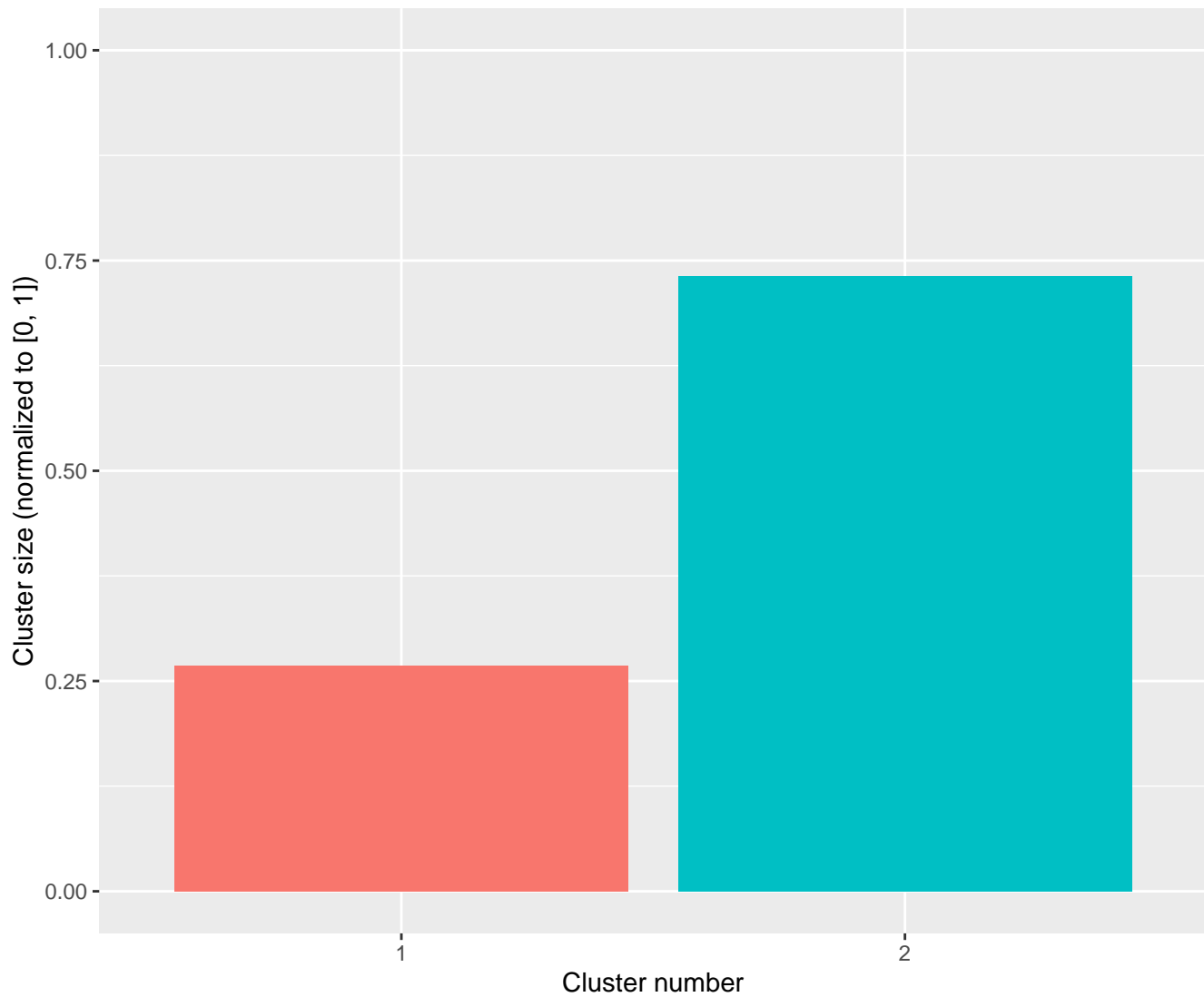


Dataset: journal.pone.0216416_Takashi2019_diabetes_type1_dataset_preproces

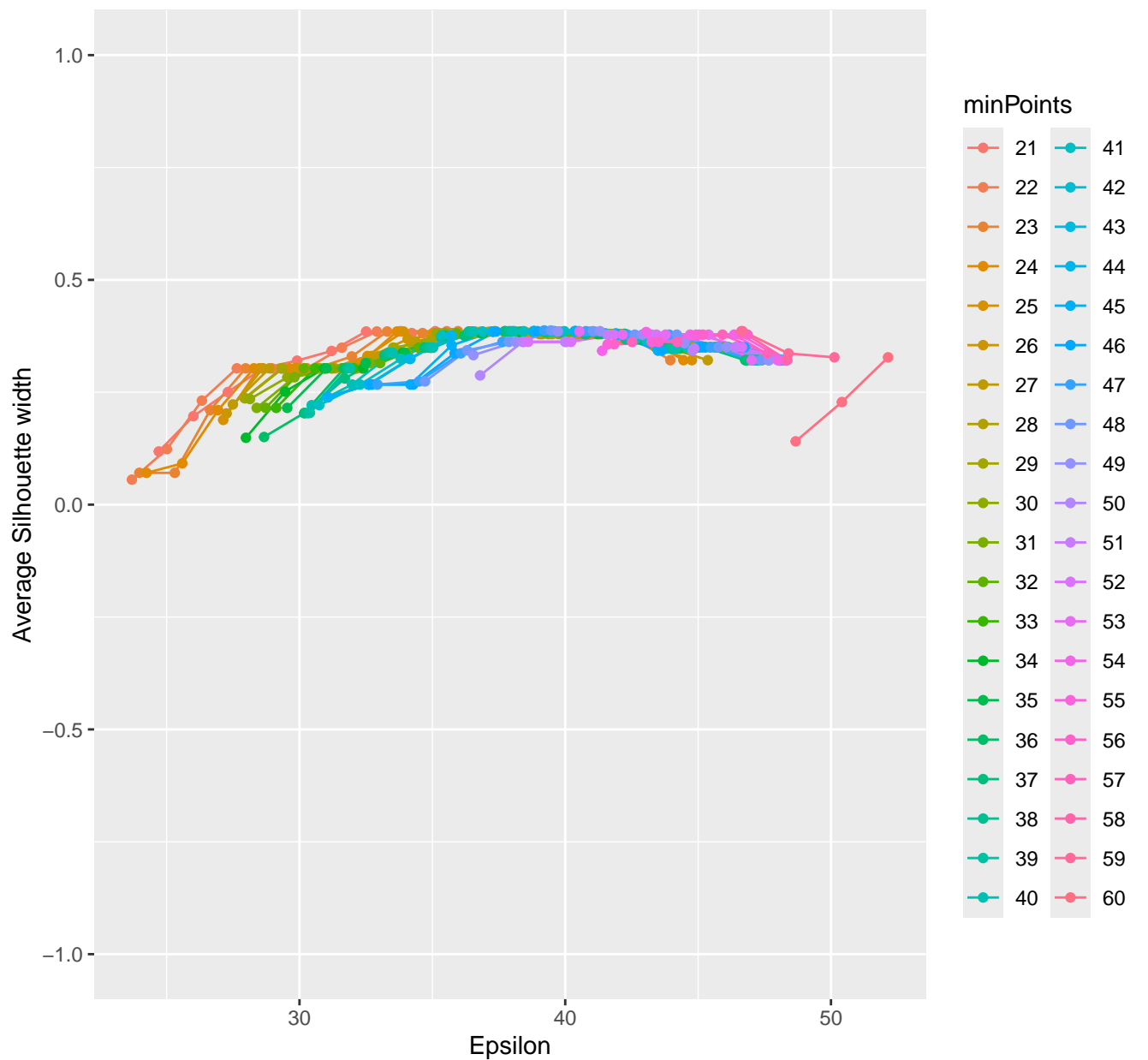
Clustering: KMEANS

Parameters used: k = 2

Cluster 1 2



DBSCAN elbow plot

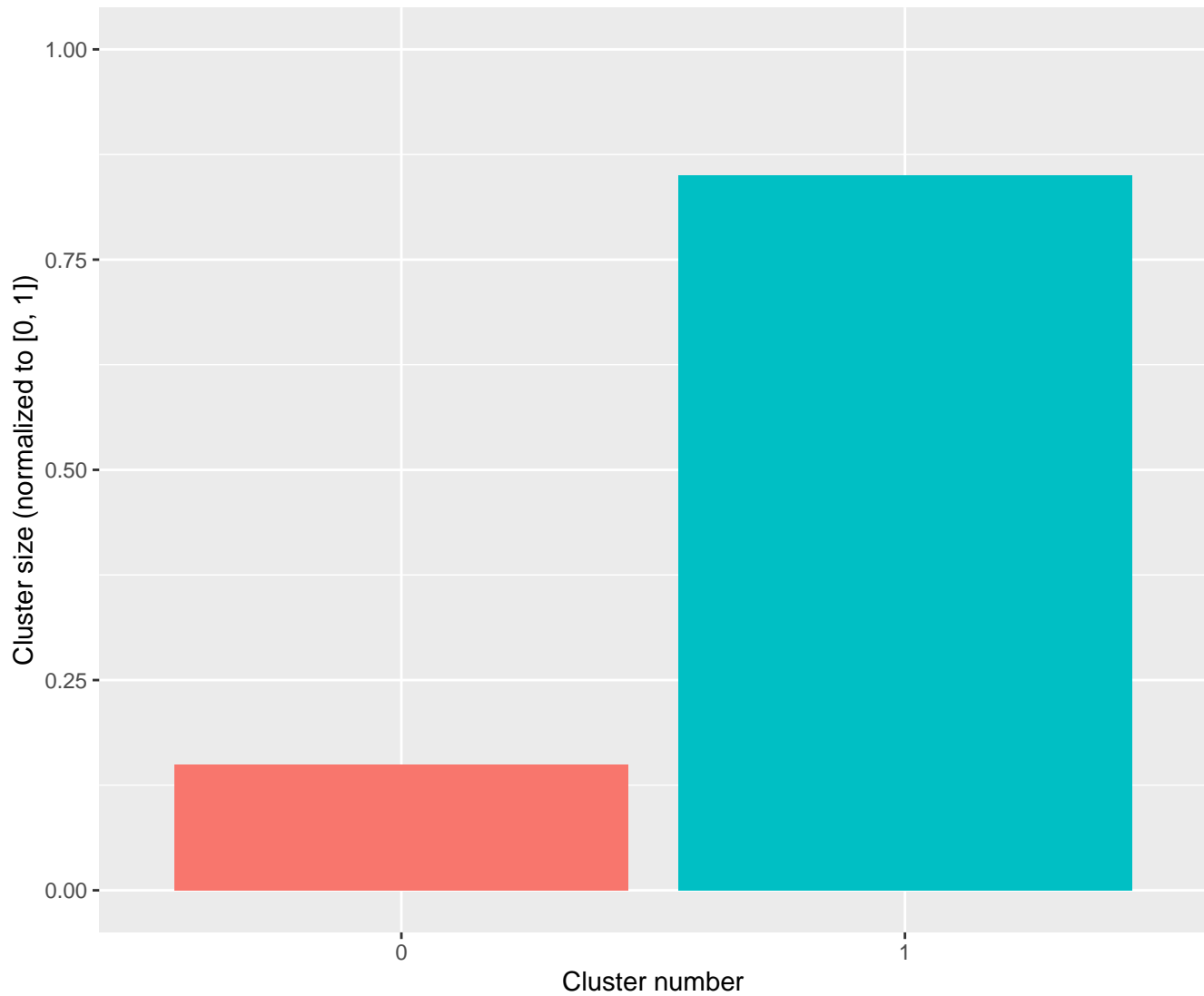


Dataset: journal.pone.0216416_Takashi2019_diabetes_type1_dataset_preproces

Clustering: DBSCAN

Parameters used: minPoints = 47, epsilon = 39.20665

Cluster 0 1

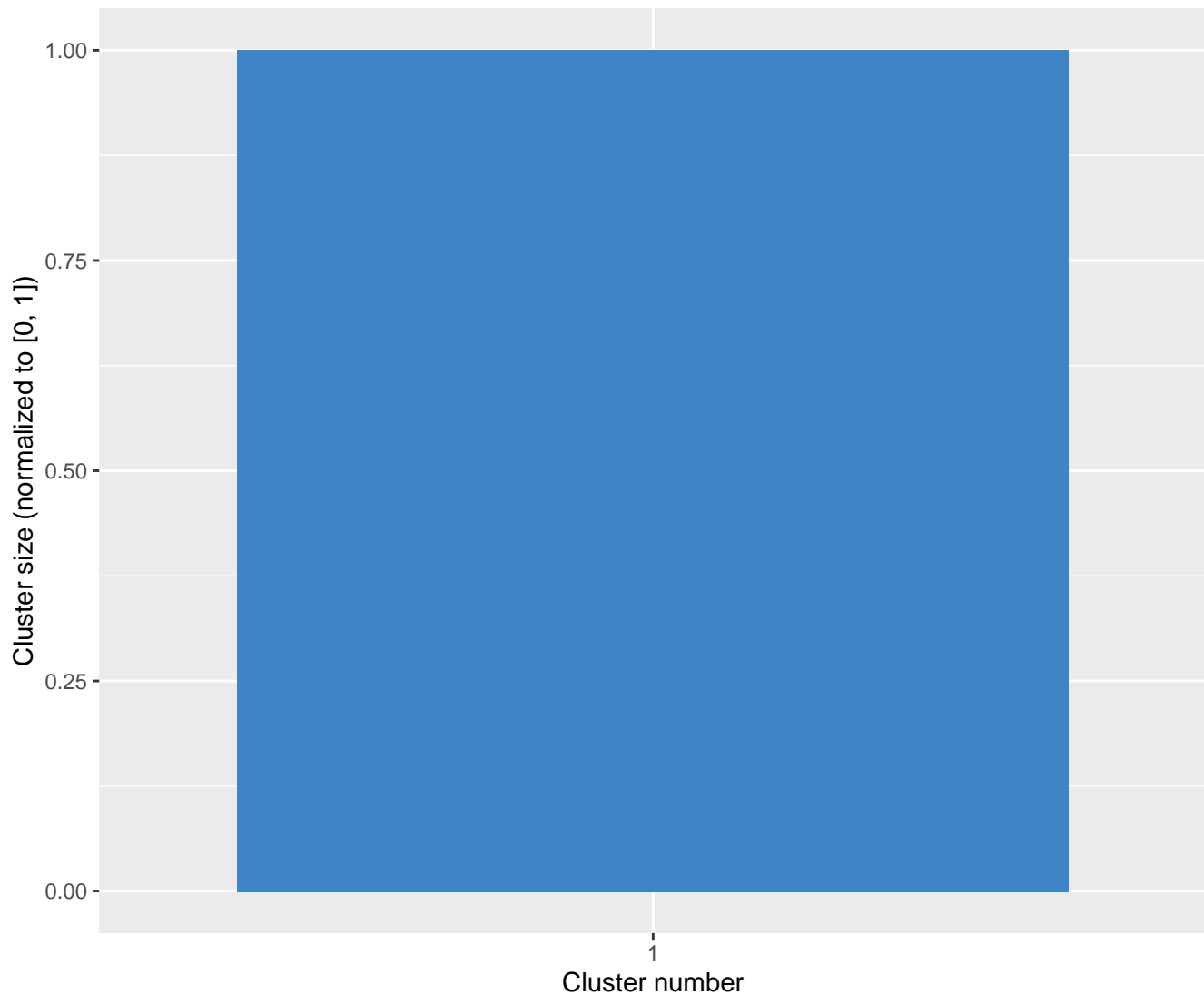


Dataset: journal.pone.0216416_Takashi2019_diabetes_type1_dataset_preproces

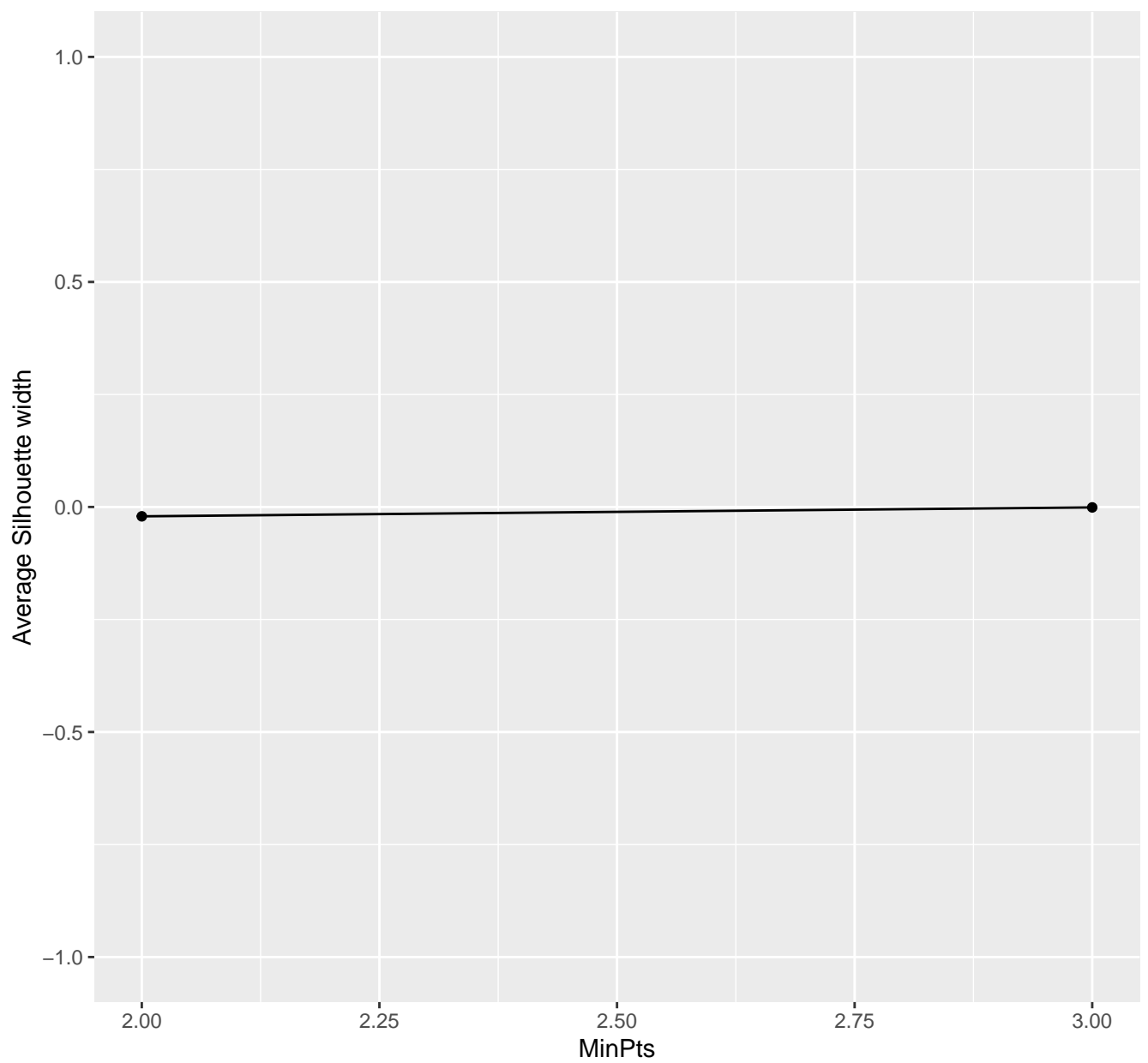
DBSCAN clustering with visual inspection

Parameters used: minPoints = 39, epsilon = 50

Cluster  1



HDBSCAN elbow plot

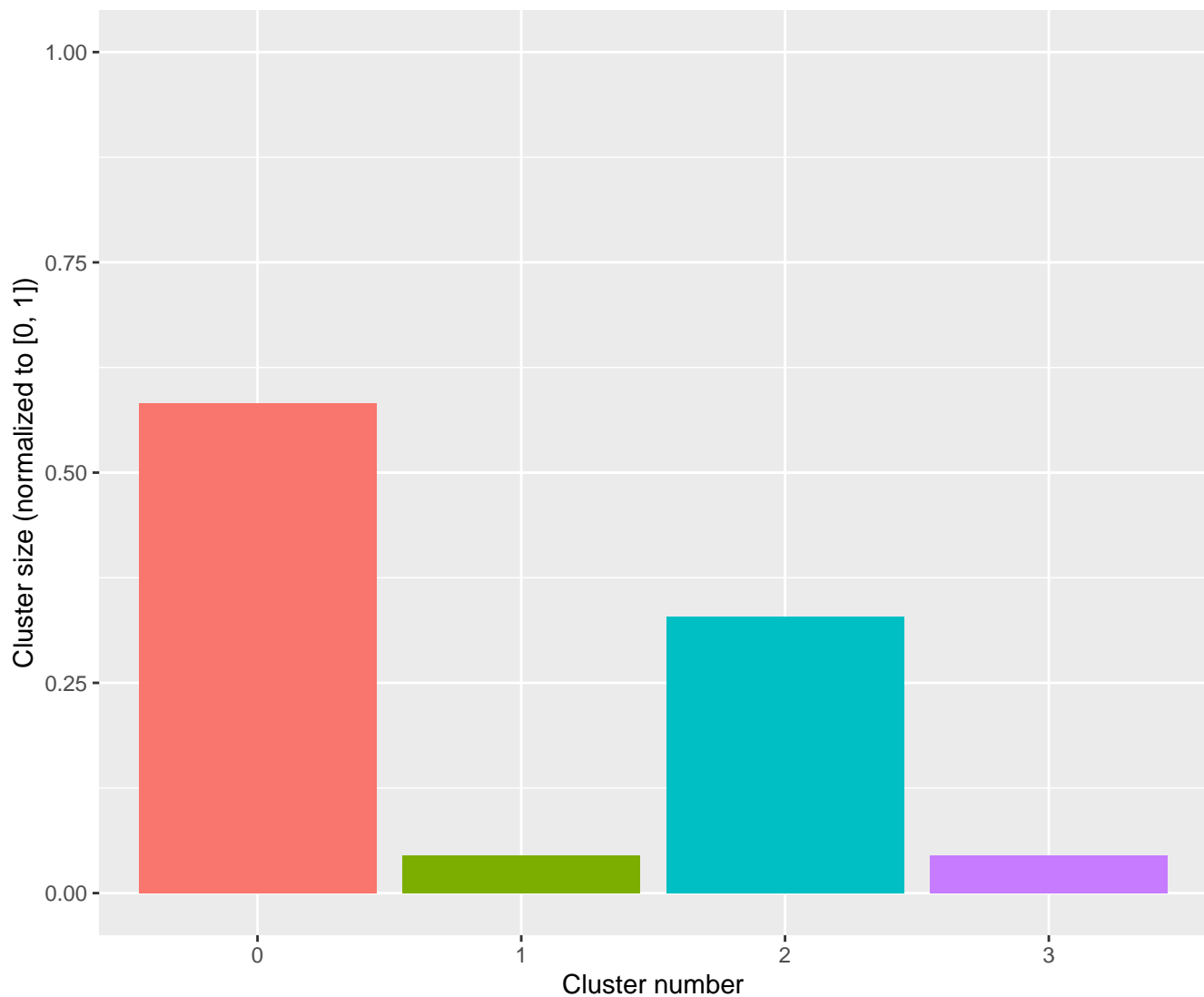


Dataset: journal.pone.0216416_Takashi2019_diabetes_type1_dataset_preproces

Clustering: HDBSCAN

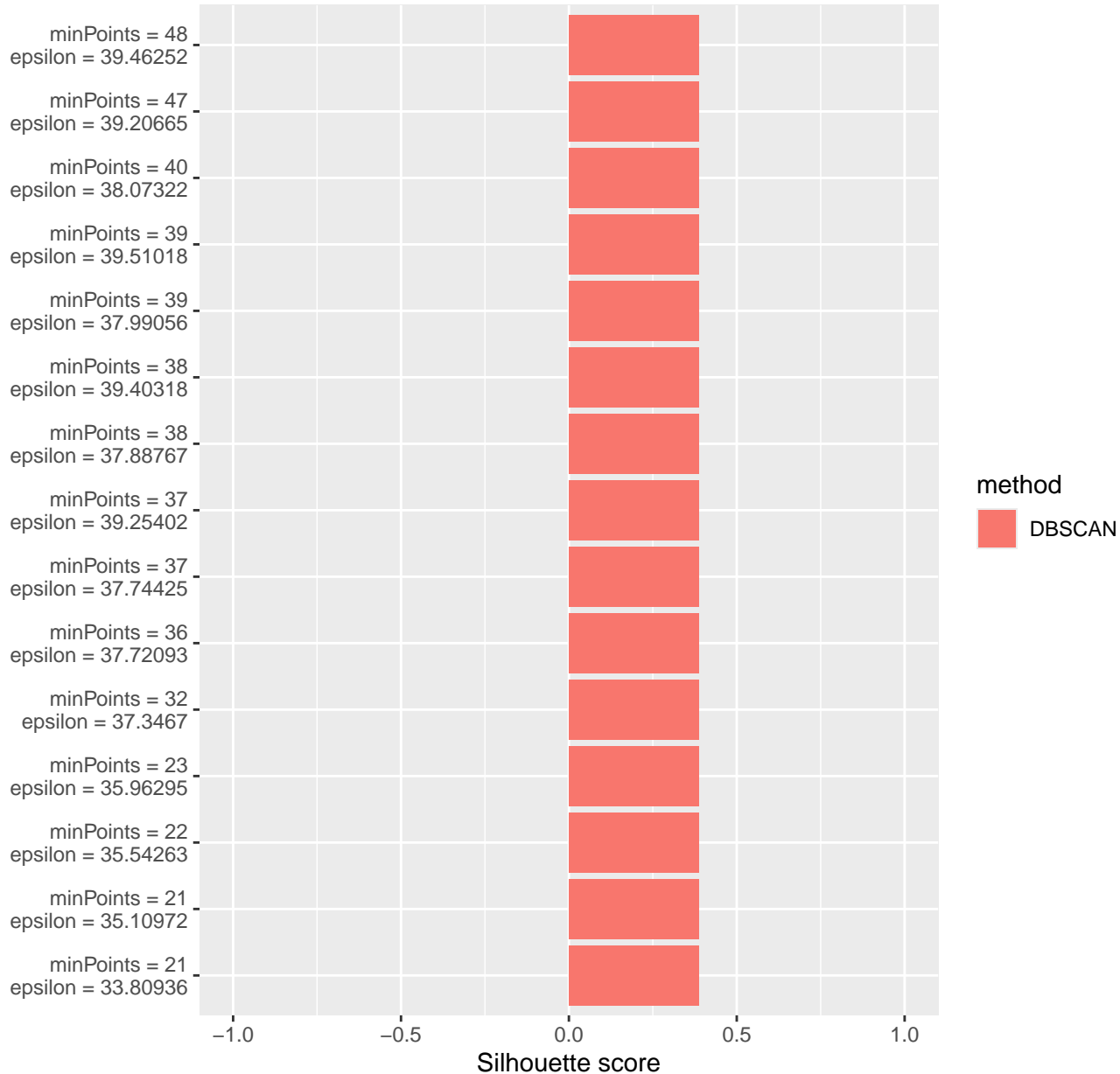
Parameters used: minPoints = 3

Cluster 0 1 2 3



Dataset: journal.pone.0216416_Takashi2019_diabetes_type1_dataset

Hyperparameter combinations ranked by its average Silhouette score



Dataset: journal.pone.0216416_Takashi2019_diabetes_type1_dataset_preproc

Algorithms ranked by their average Silhouette score

