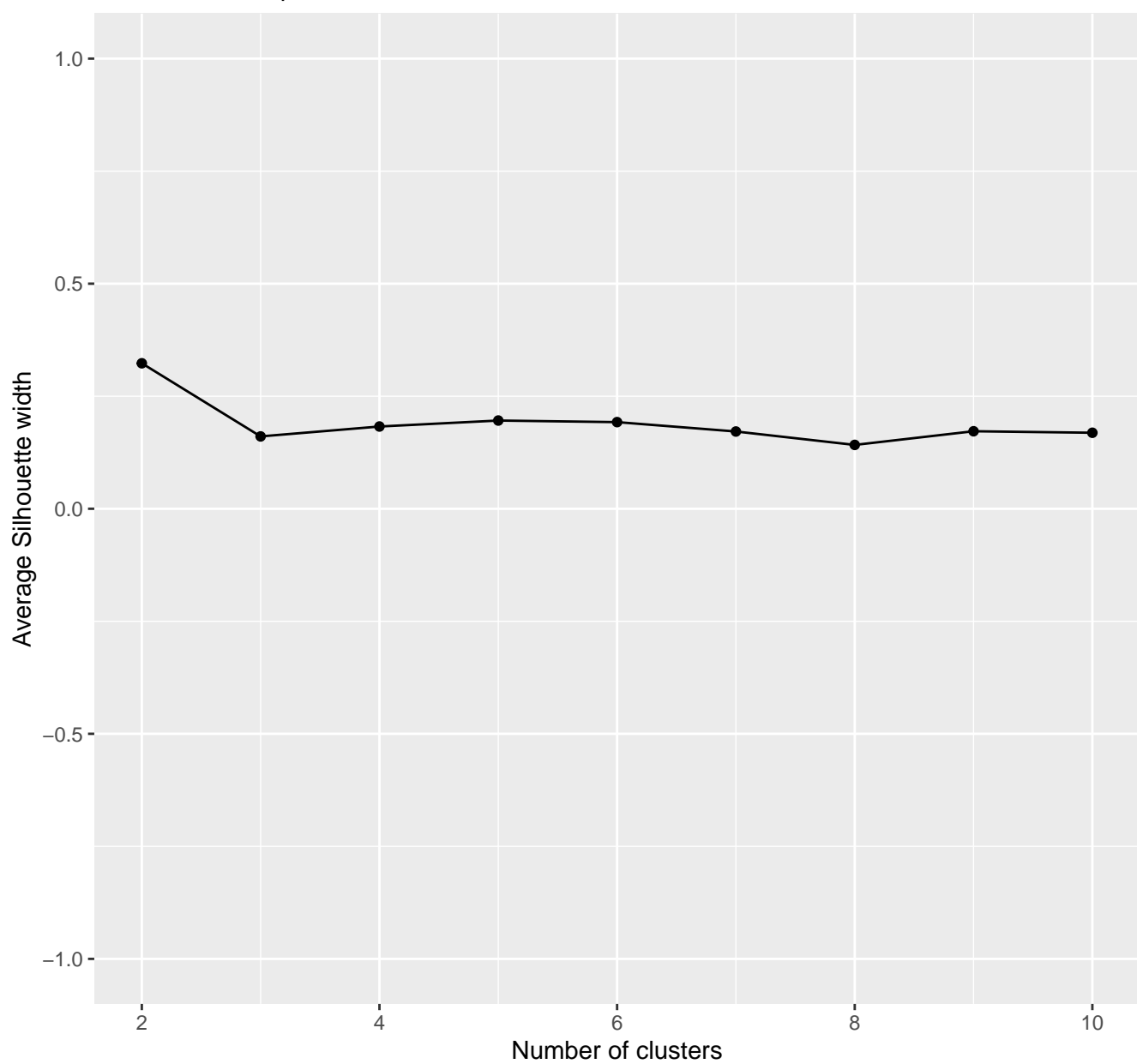


K-Means elbow plot

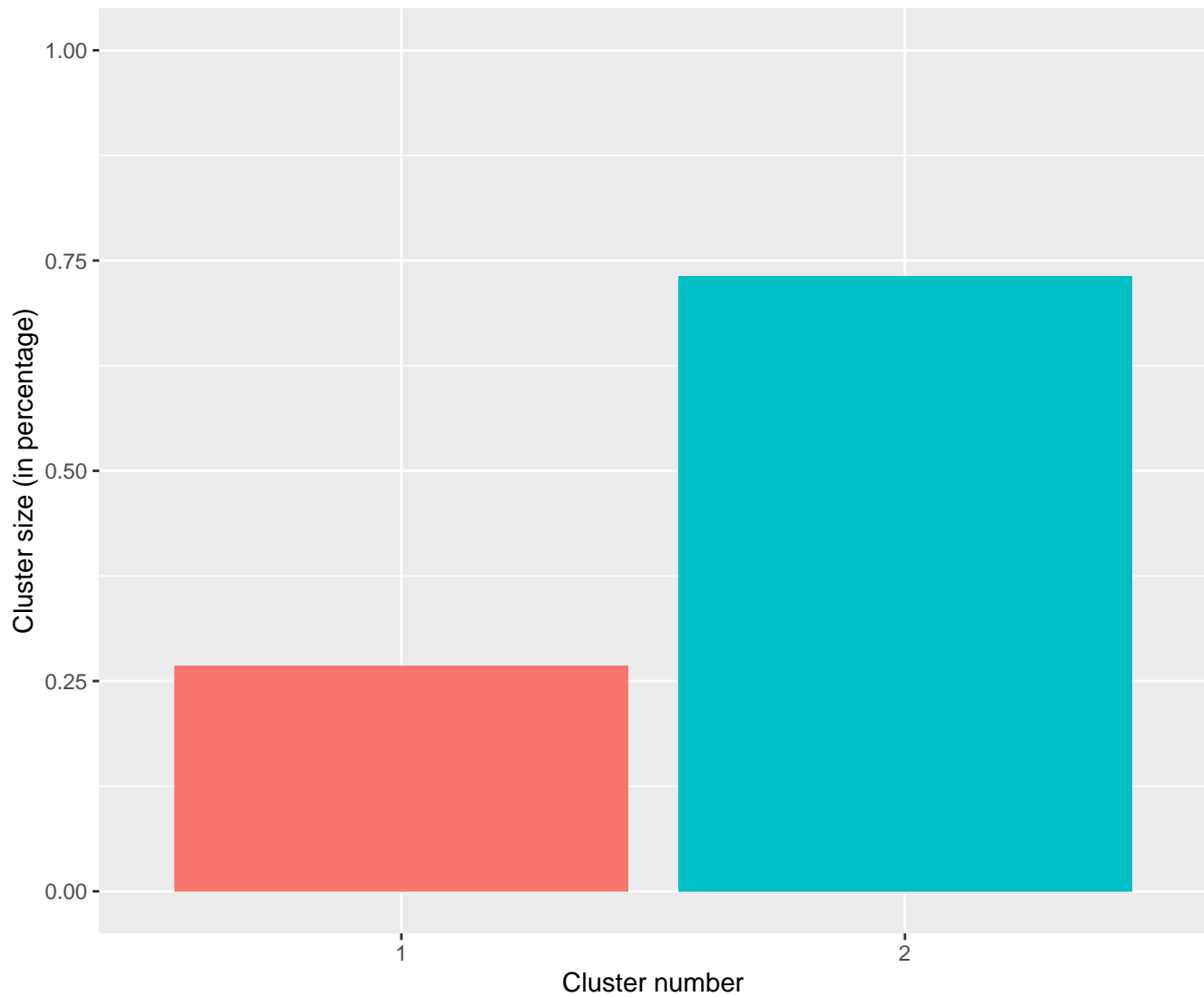


Dataset: journal.pone.0216416\_Takashi2019\_diabetes\_type1\_dataset\_preproces

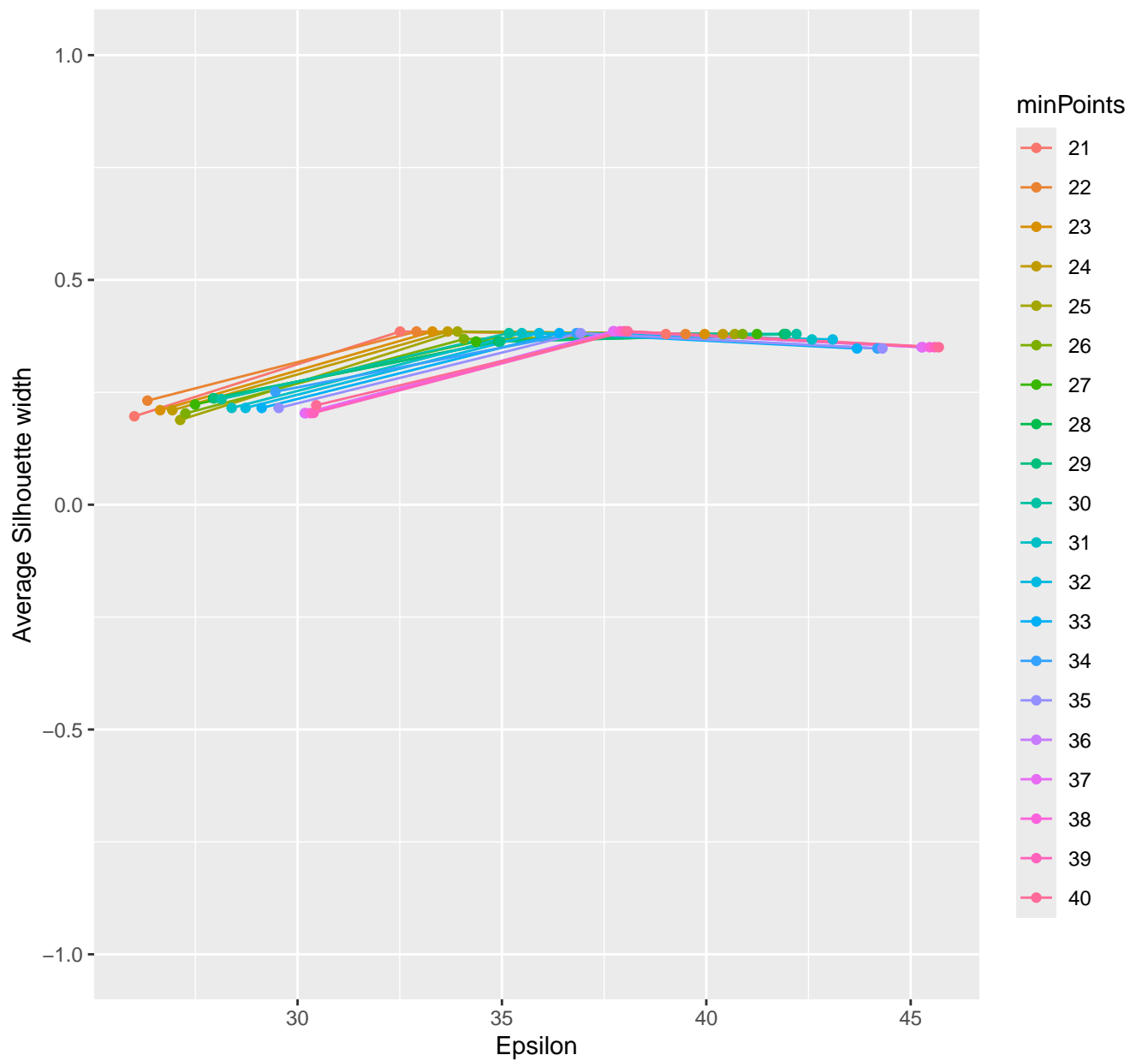
Clustering: KMEANS

Parameters used: k = 2

Cluster 1 2



## DBSCAN elbow plot

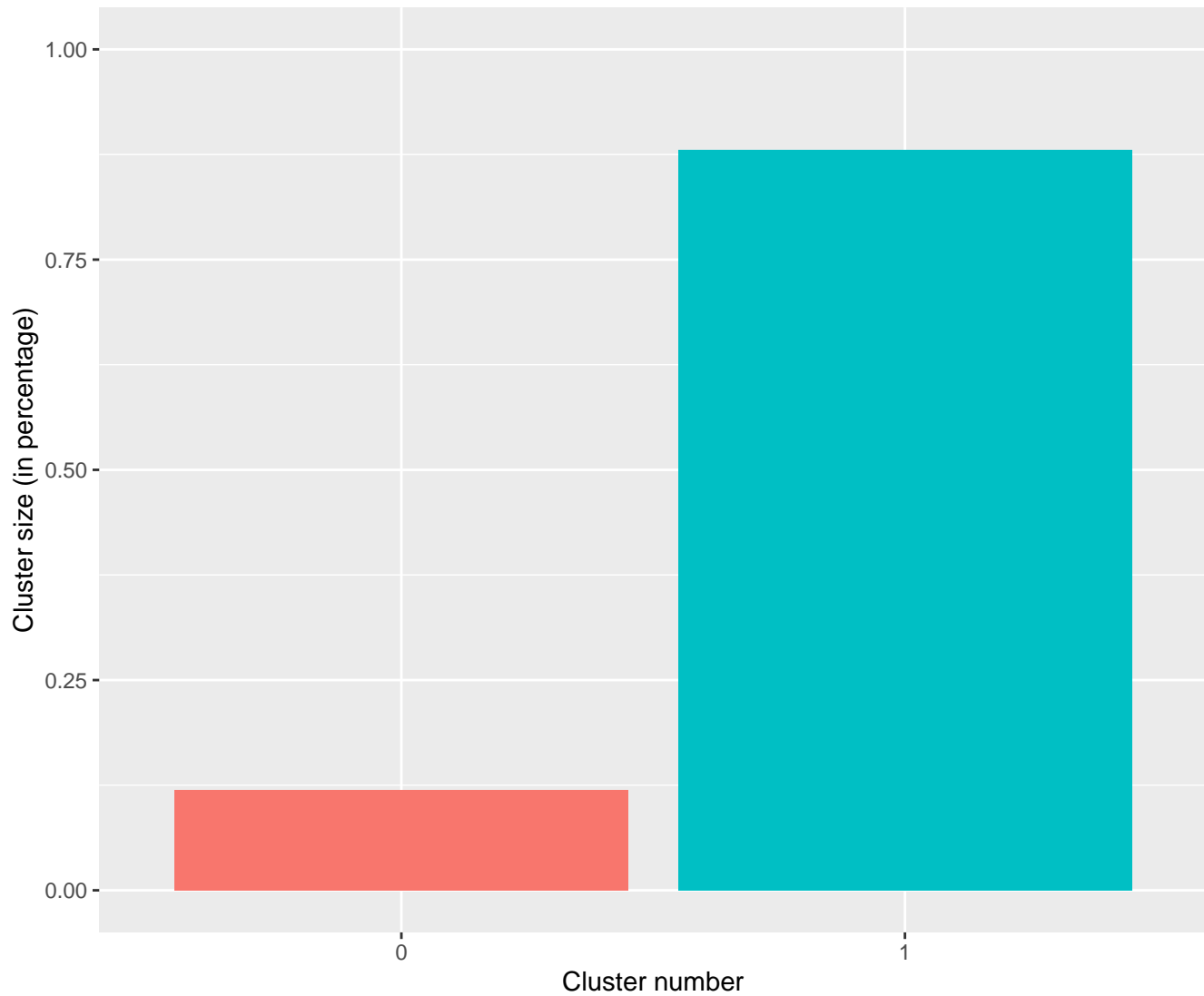


Dataset: journal.pone.0216416\_Takashi2019\_diabetes\_type1\_dataset\_preproces

Clustering: DBSCAN

Parameters used: minPoints = 36, epsilon = 37.72093

Cluster 0 1

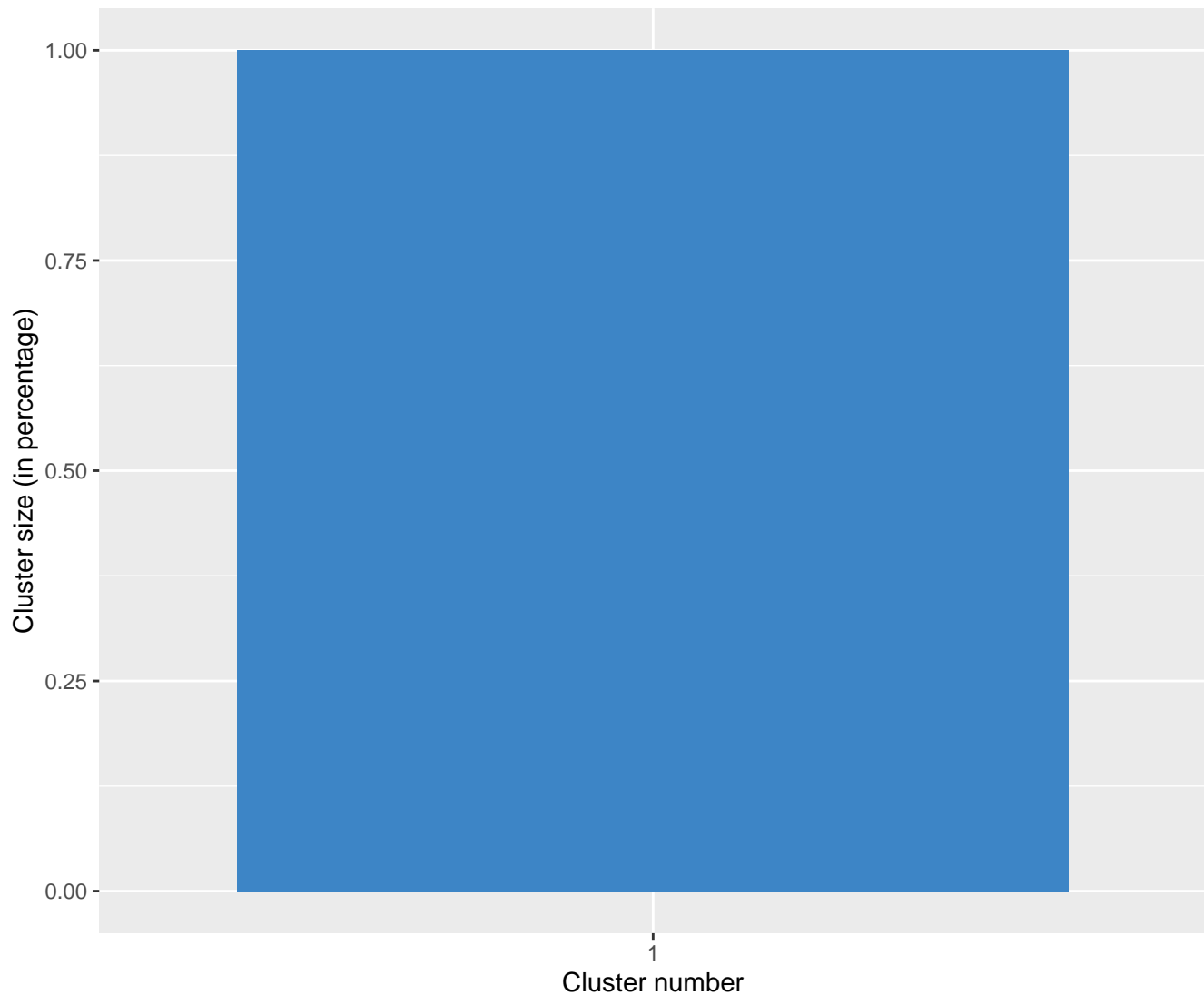


Dataset: journal.pone.0216416\_Takashi2019\_diabetes\_type1\_dataset\_preproces

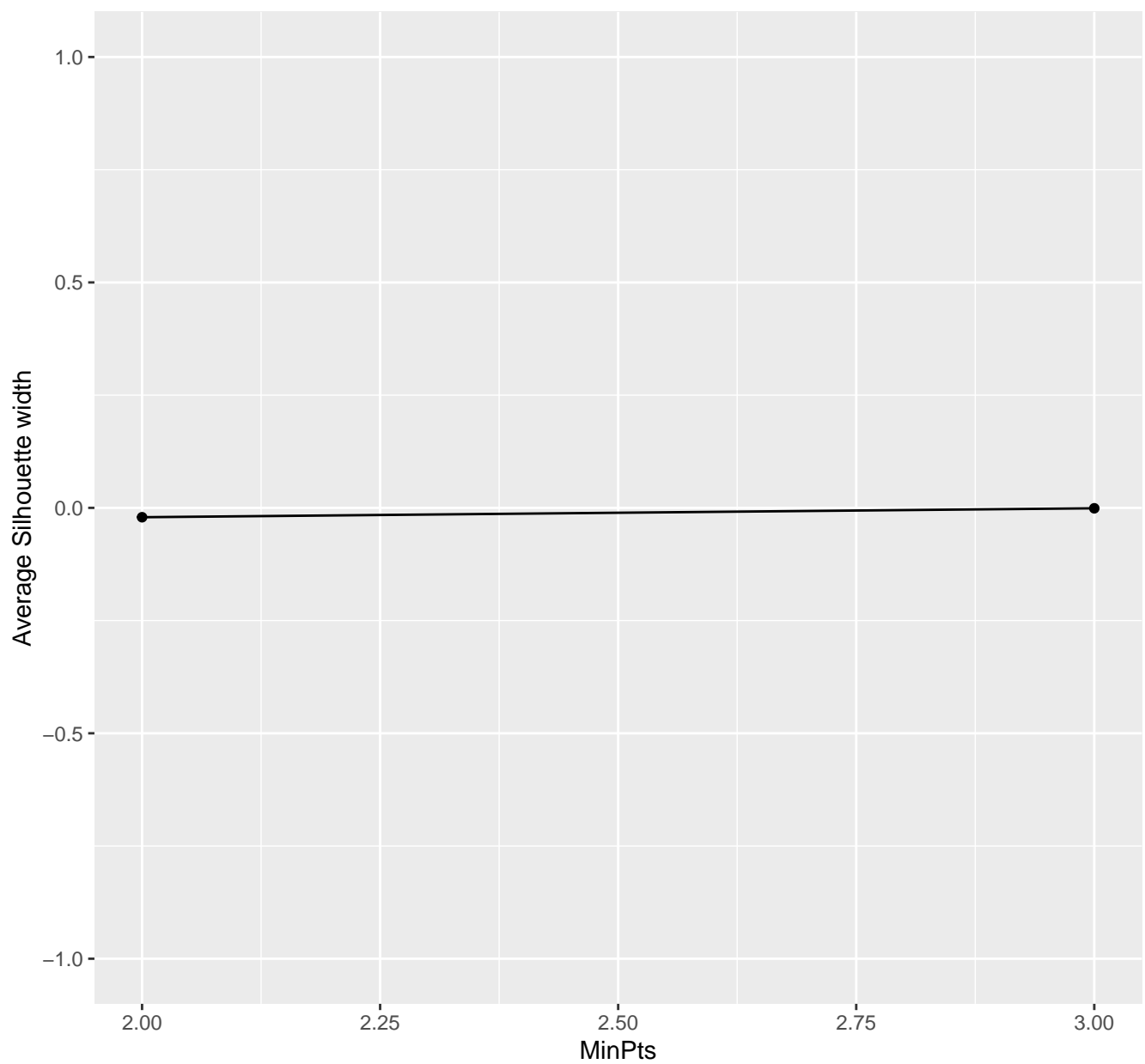
DBSCAN clustering with visual inspection

Parameters used: minPoints = 39, epsilon = 50

Cluster  1



HDBSCAN elbow plot

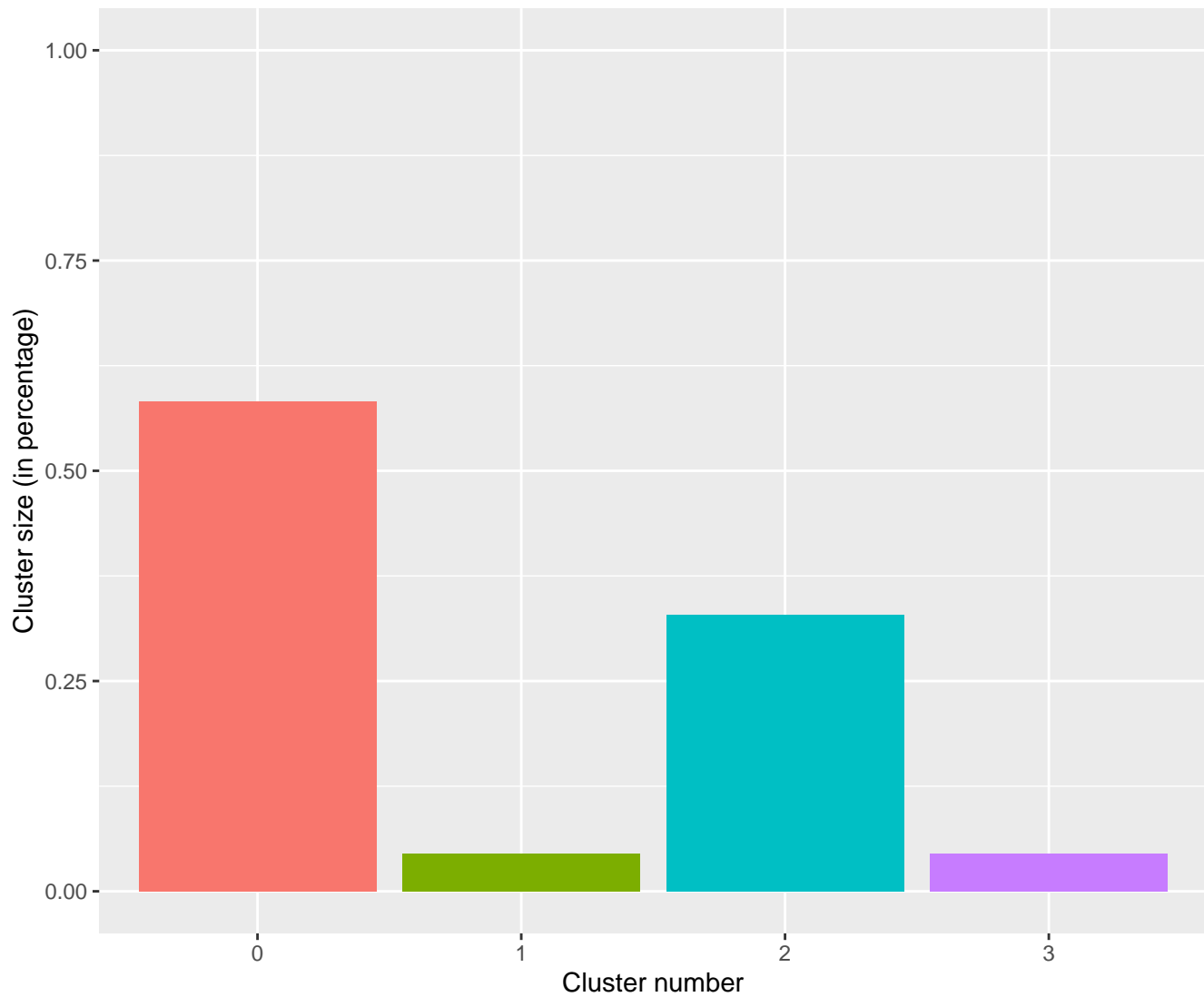


Dataset: journal.pone.0216416\_Takashi2019\_diabetes\_type1\_dataset\_preproces

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

