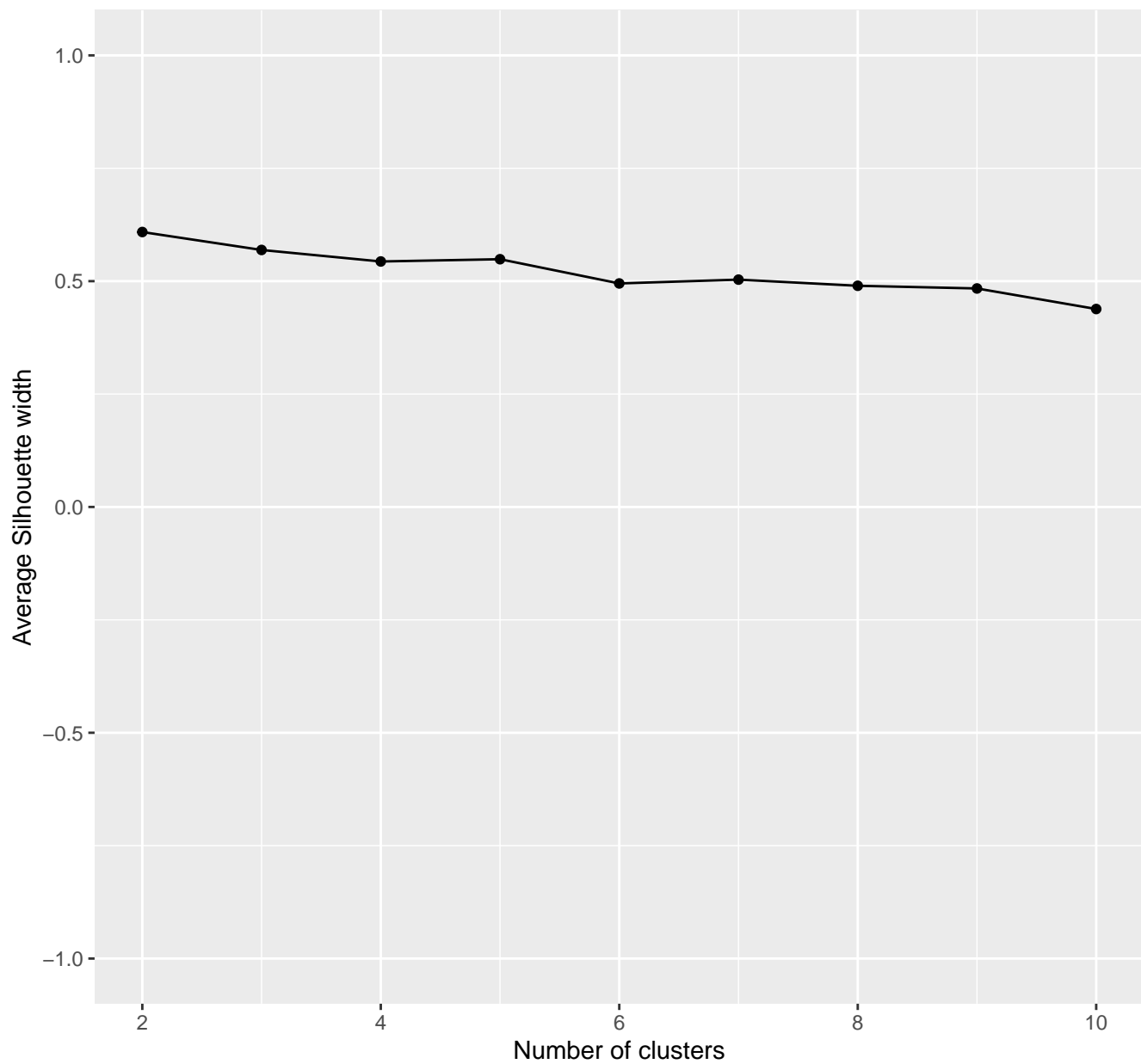


# K-Means elbow plot

Dataset: 10\_7717\_peerj\_5665\_dataYM2018\_neuroblastoma.csv

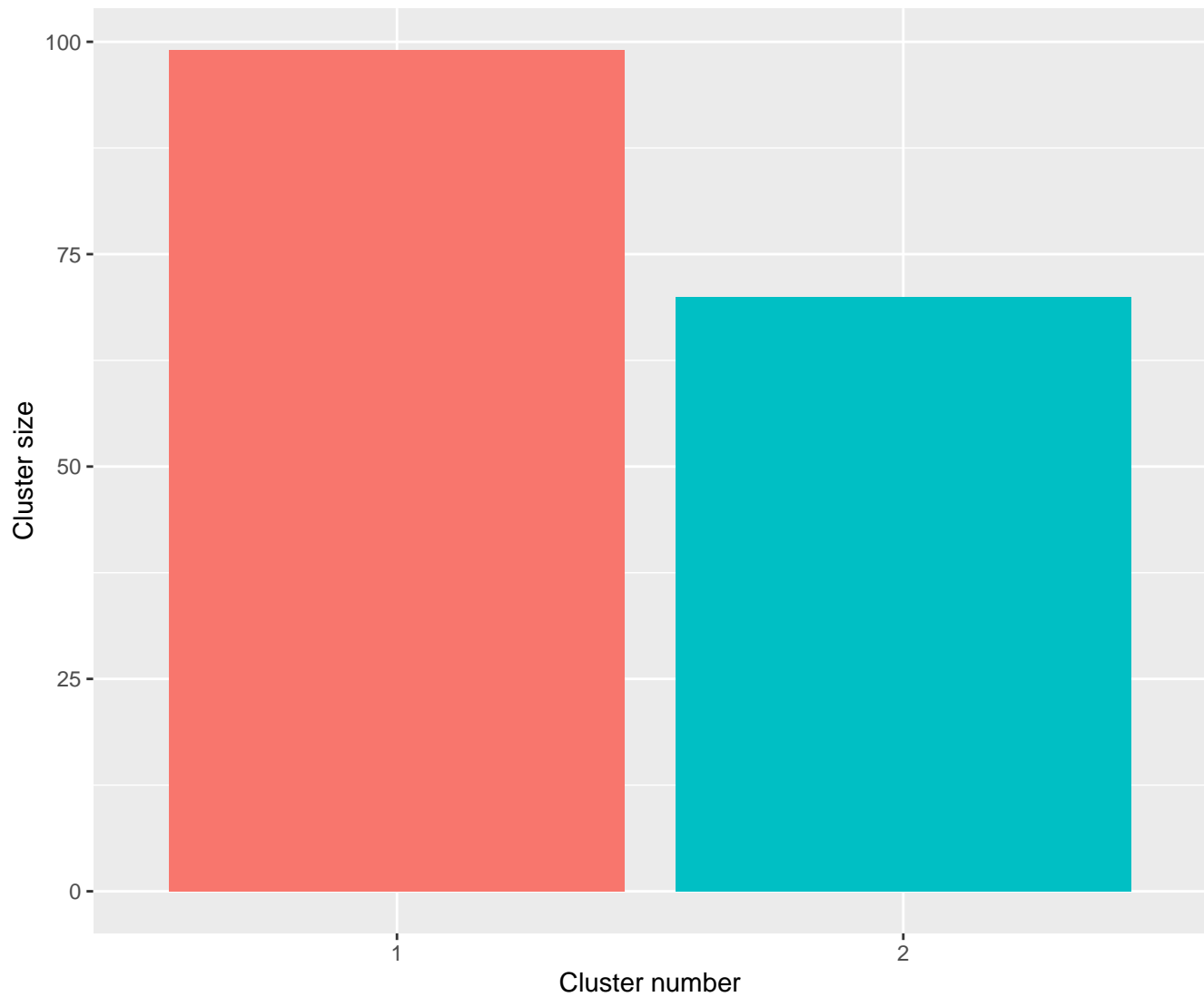


# Clustering: KMEANS

Dataset: 10\_7717\_peerj\_5665\_dataYM2018\_neuroblastoma.csv

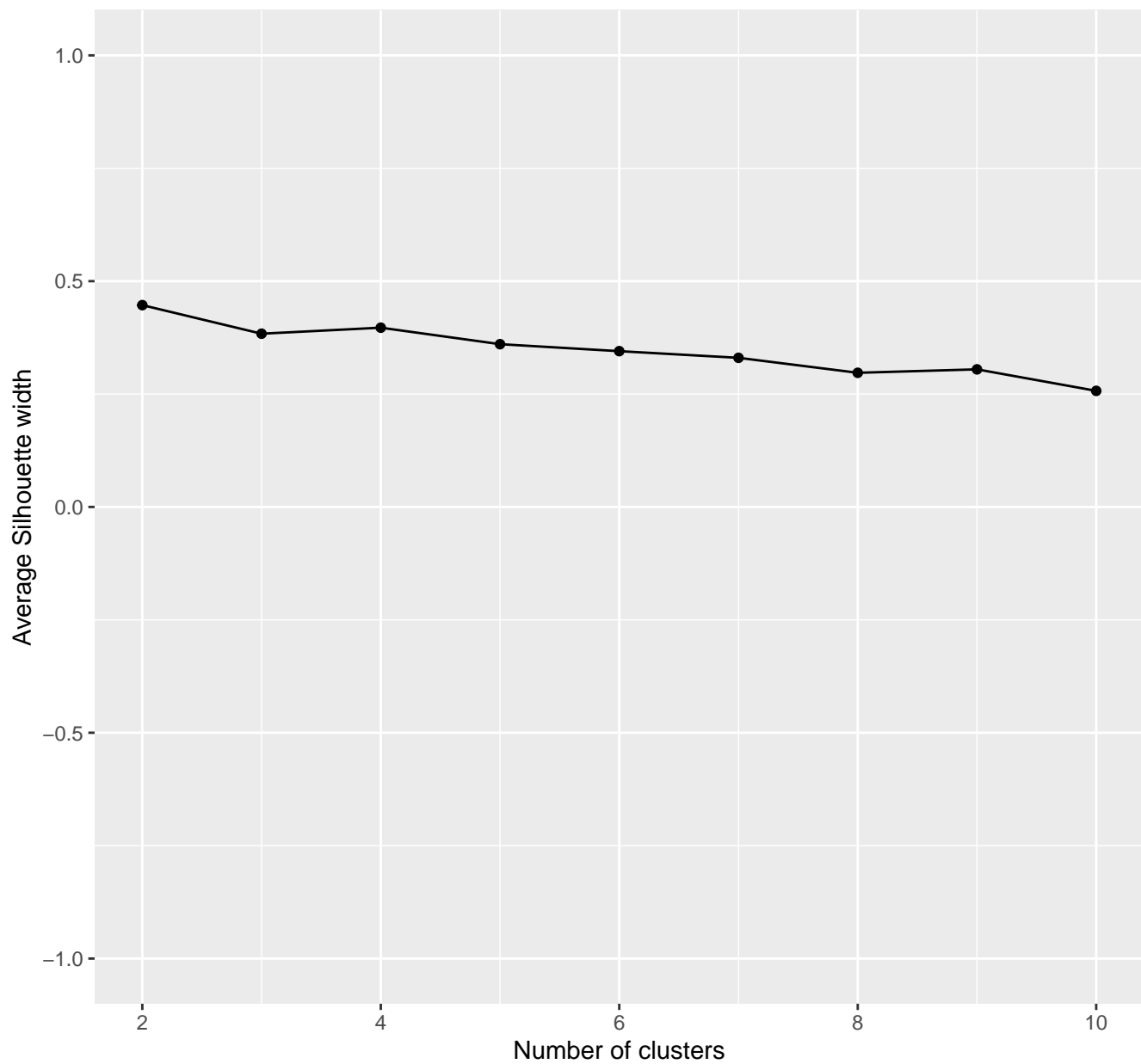
Parameters used: k = 2

Cluster 1 2



# K-Means elbow plot

Dataset: journal.pone.0148699\_S1\_Text\_Sepsis\_SIRS\_EDITED.csv

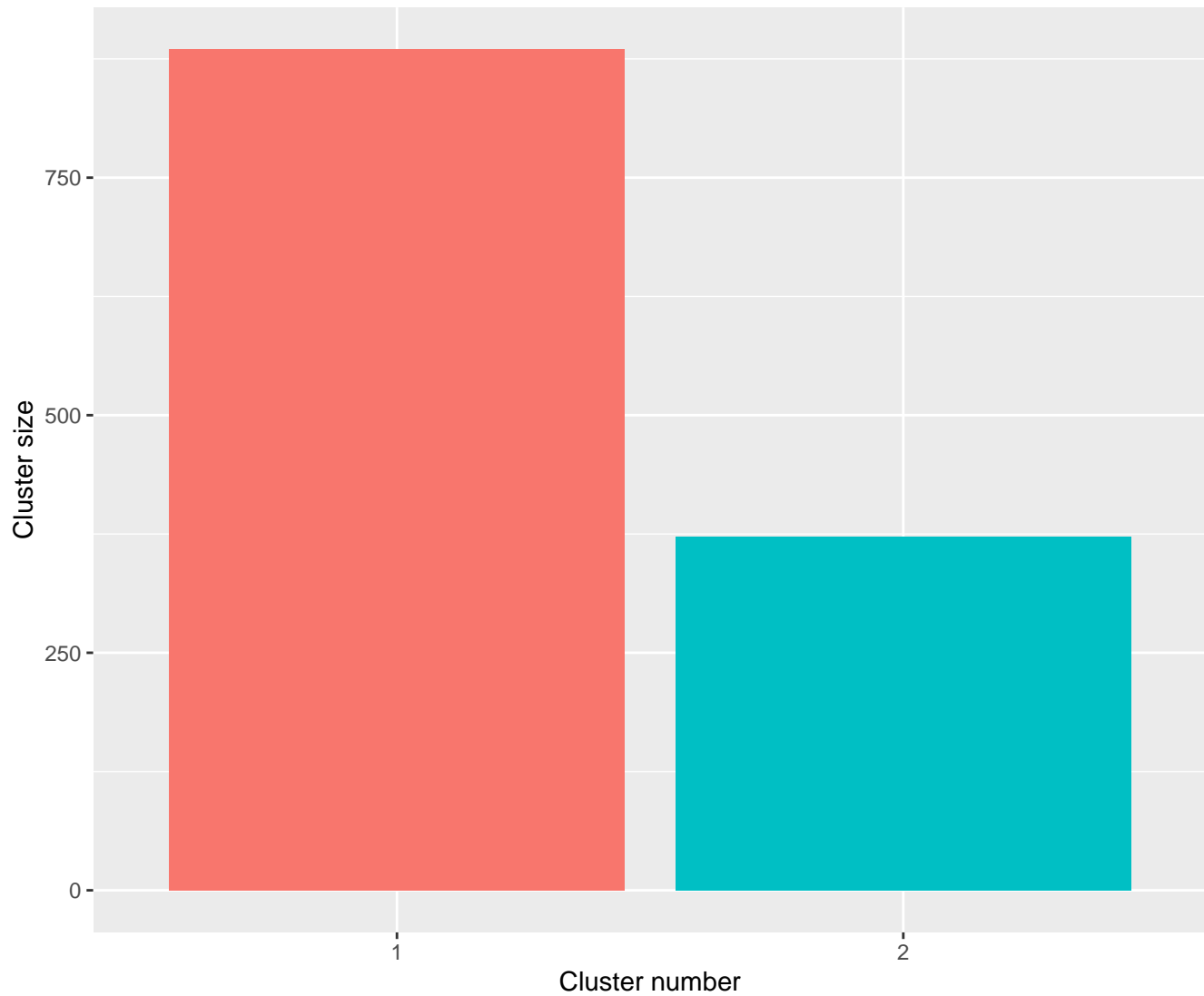


# Clustering: KMEANS

Dataset: journal.pone.0148699\_S1\_Text\_Sepsis\_SIRS\_EDITED.csv

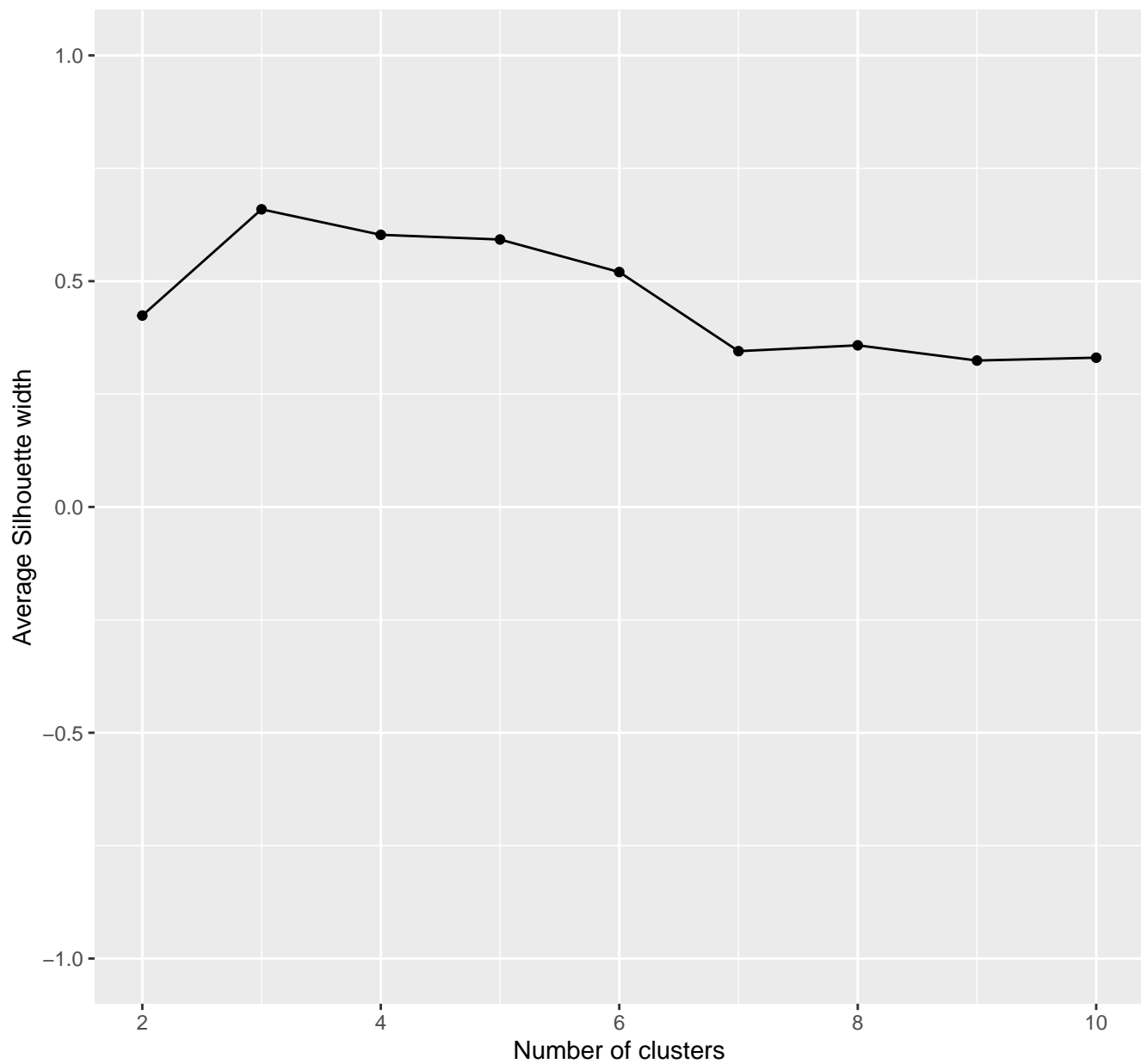
Parameters used: k = 2

Cluster 1 2



# K-Means elbow plot

Dataset: journal.pone.0158570\_S2File\_depression\_heart\_failure\_v2.csv

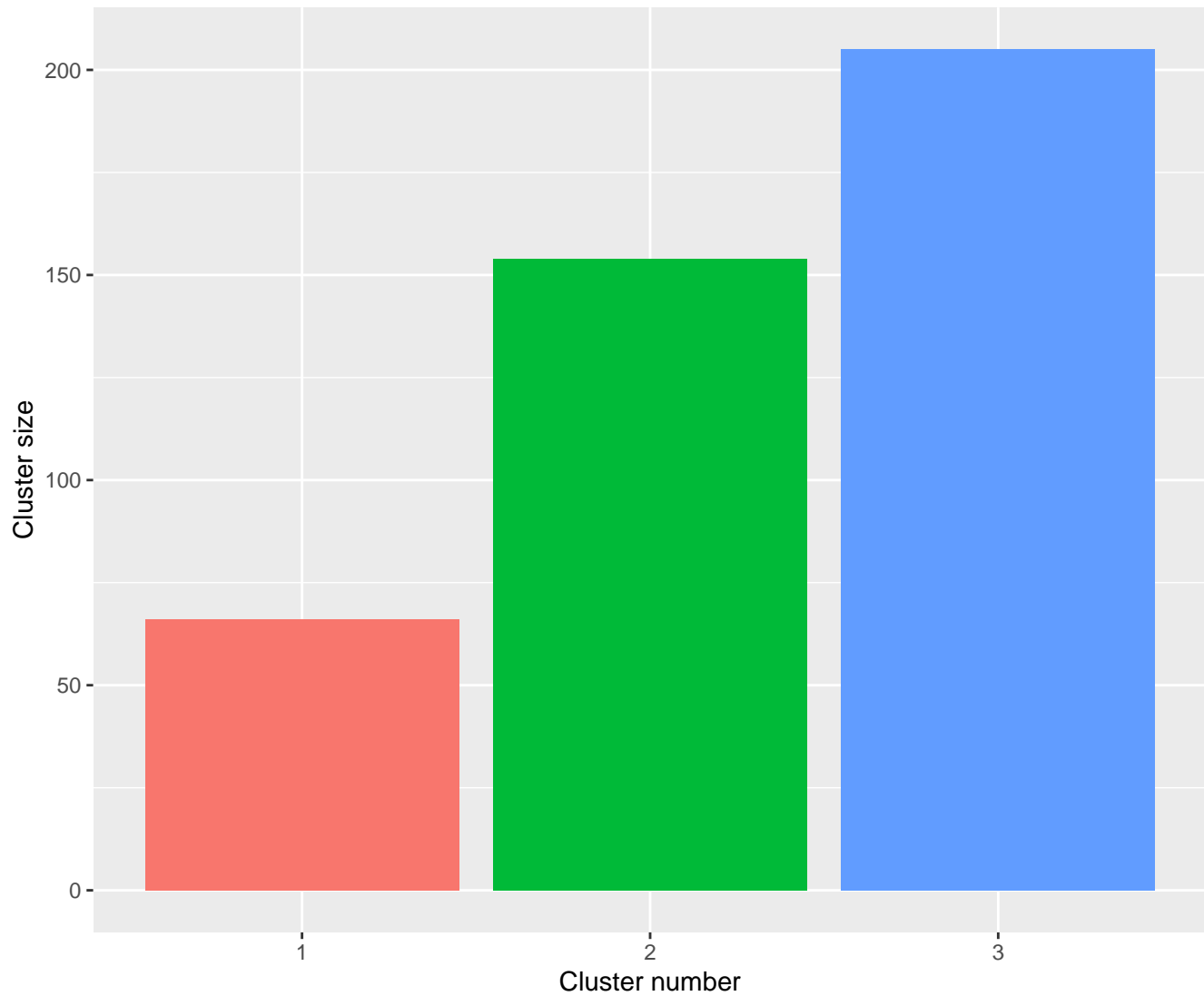


# Clustering: KMEANS

Dataset: journal.pone.0158570\_S2File\_depression\_heart\_failure\_v2.csv

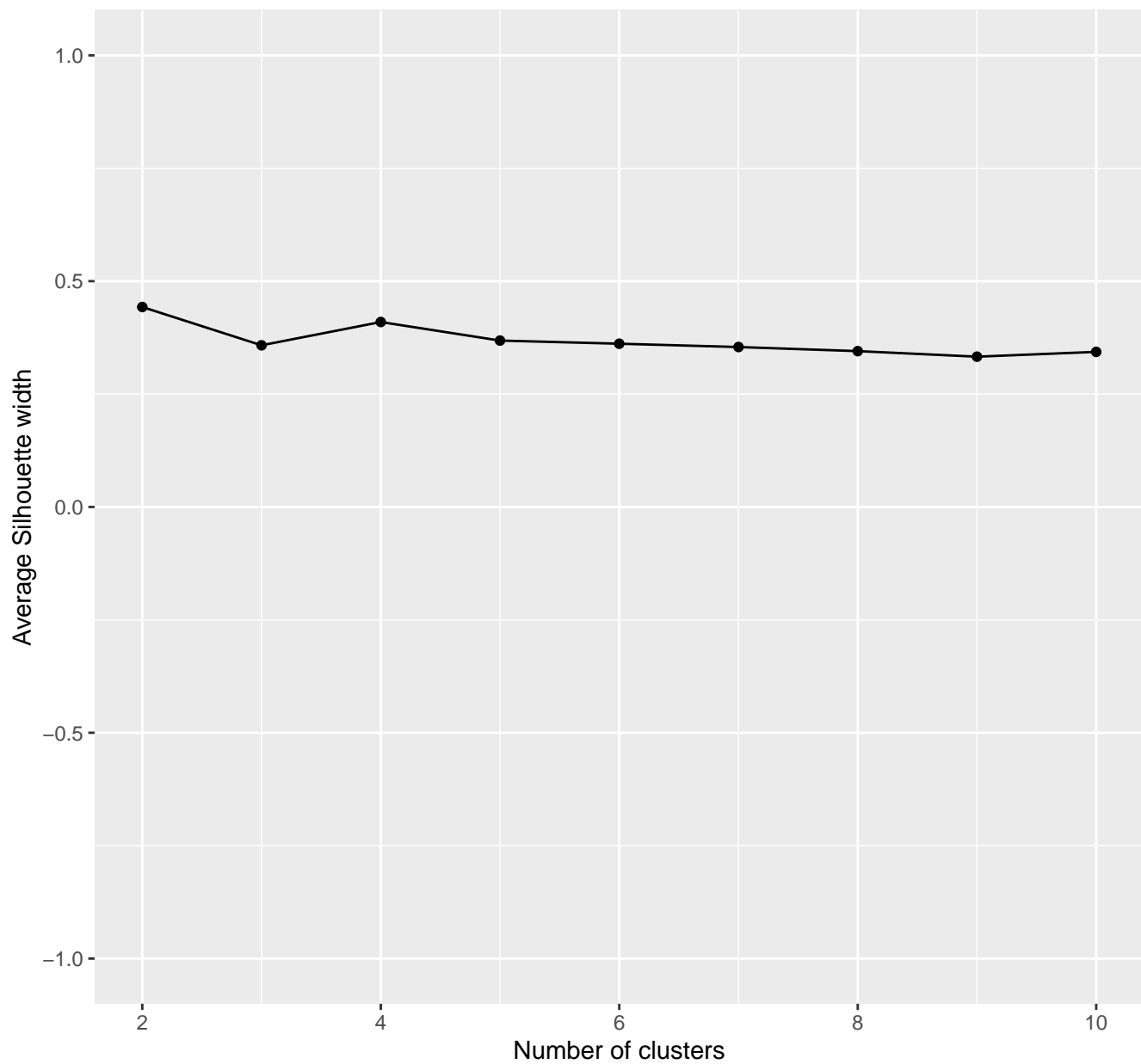
Parameters used: k = 3

Cluster 1 2 3



# K-Means elbow plot

Dataset: journal.pone.0175818\_S1Dataset\_Spain\_cardiac\_arrest\_EDITED..csv

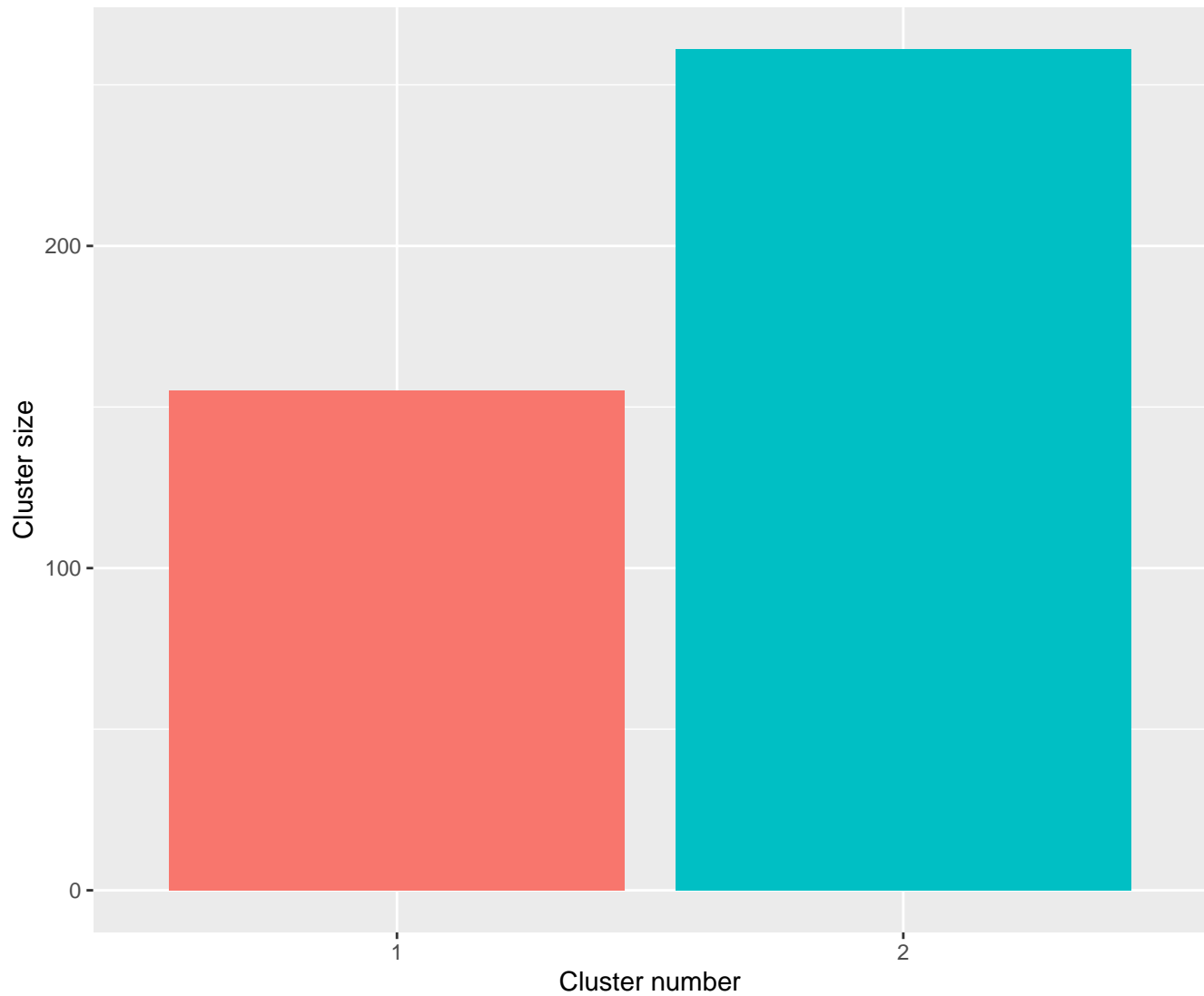


# Clustering: KMEANS

Dataset: journal.pone.0175818\_S1Dataset\_Spain\_cardiac\_arrest\_EDITED..csv

Parameters used: k = 2

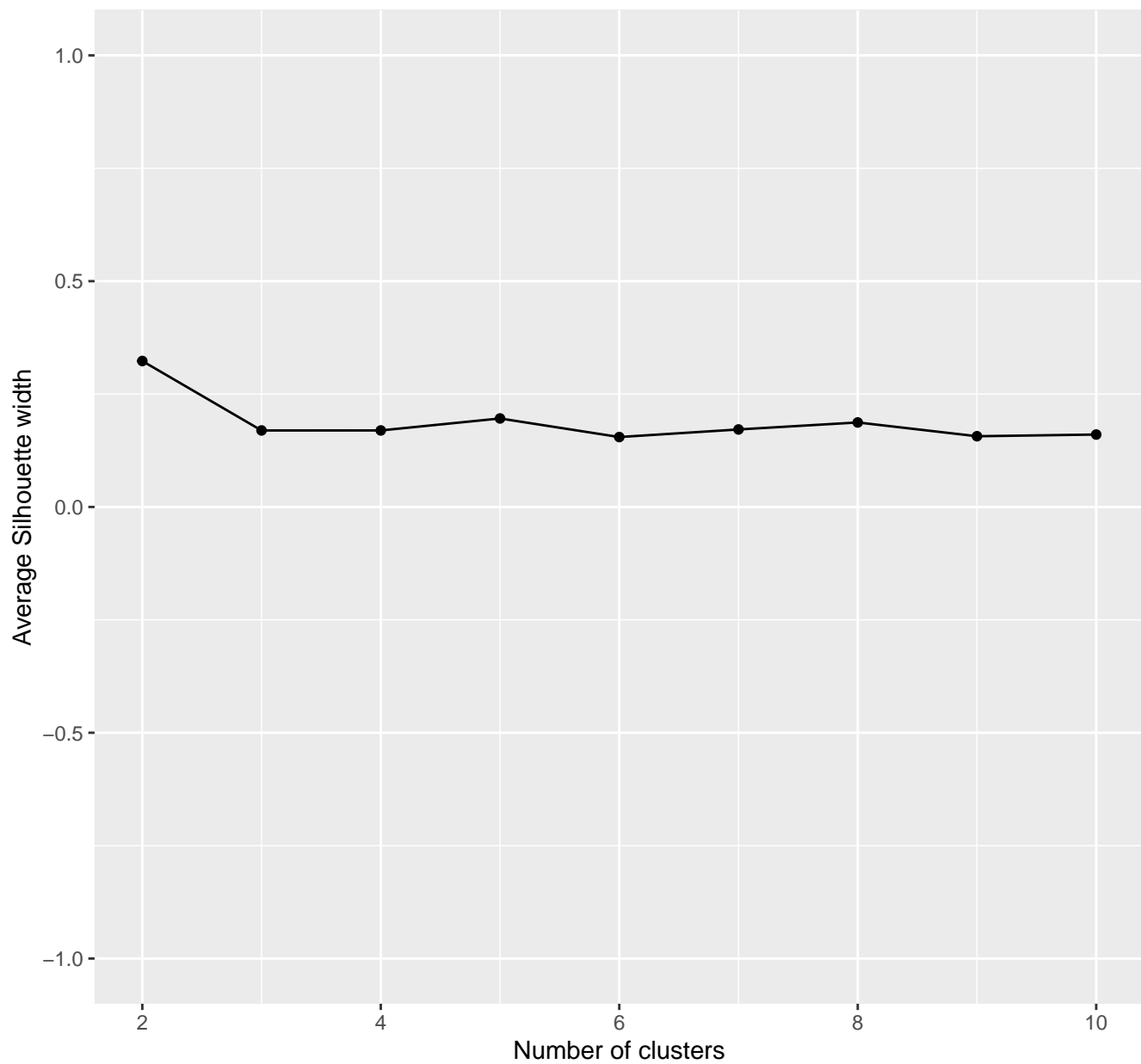
Cluster 1 2





# K-Means elbow plot

Dataset: journal.pone.0216416\_Takashi2019\_diabetes\_type1\_dataset\_preprocessed.csv



# Clustering: KMEANS

Dataset: journal.pone.0216416\_Takashi2019\_diabetes\_type1\_dataset\_preprocessed.csv

Parameters used: k = 2

Cluster ■ 1 ■ 2

