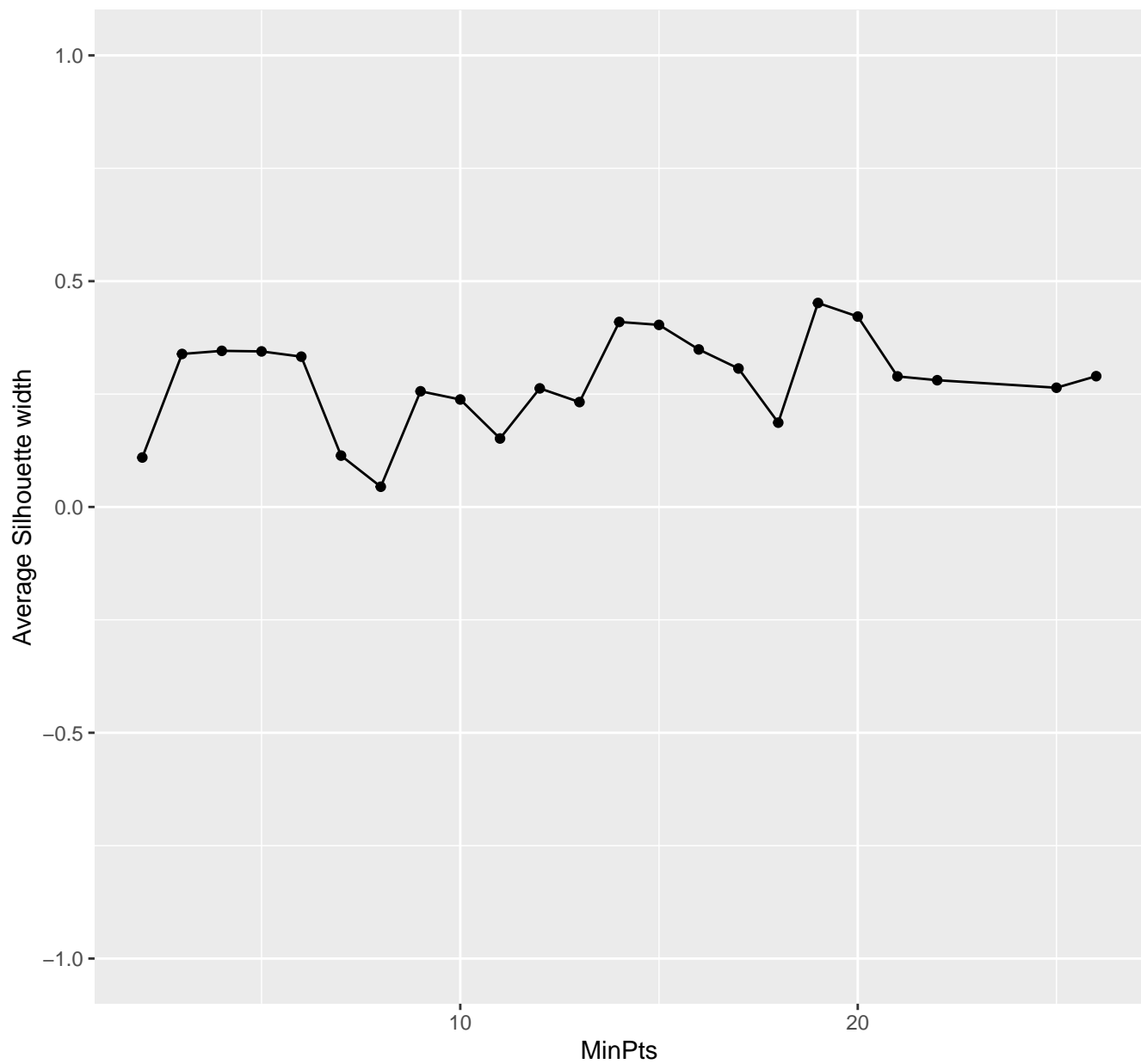


# HDBSCAN elbow plot

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

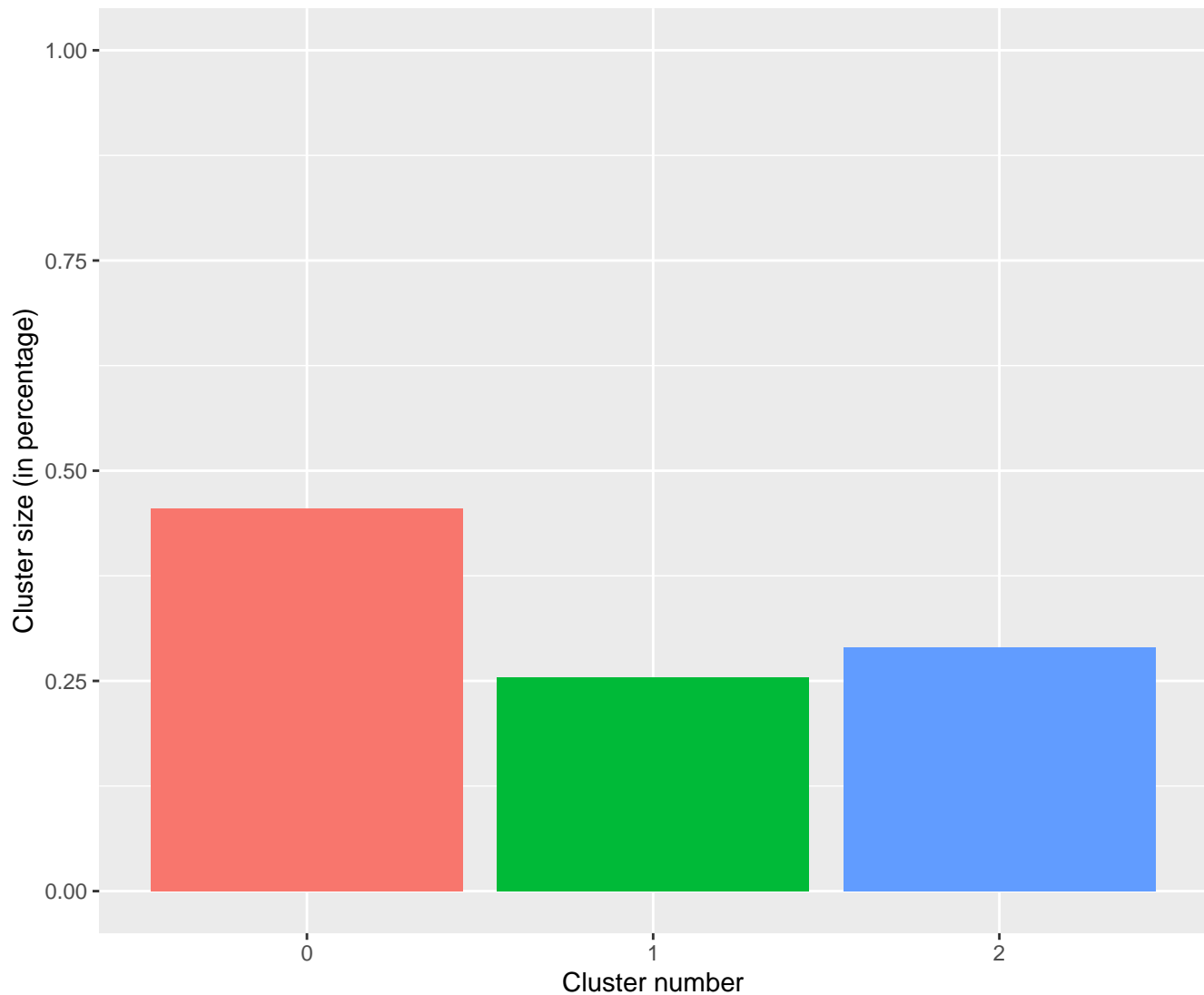


# Clustering: HDBSCAN

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

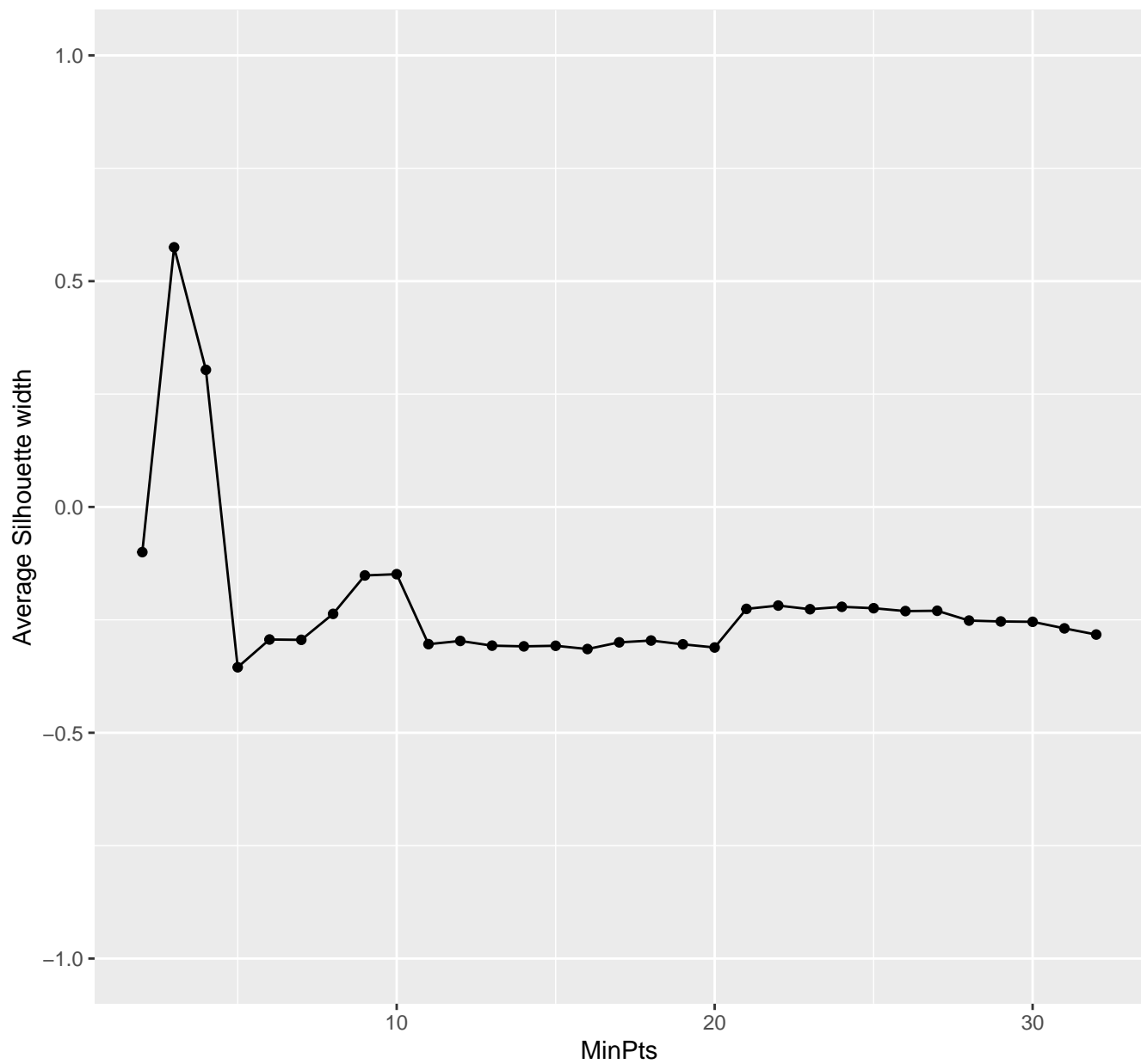
Parameters used: minPoints = 19

Cluster ■ 0 ■ 1 ■ 2



# HDBSCAN elbow plot

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

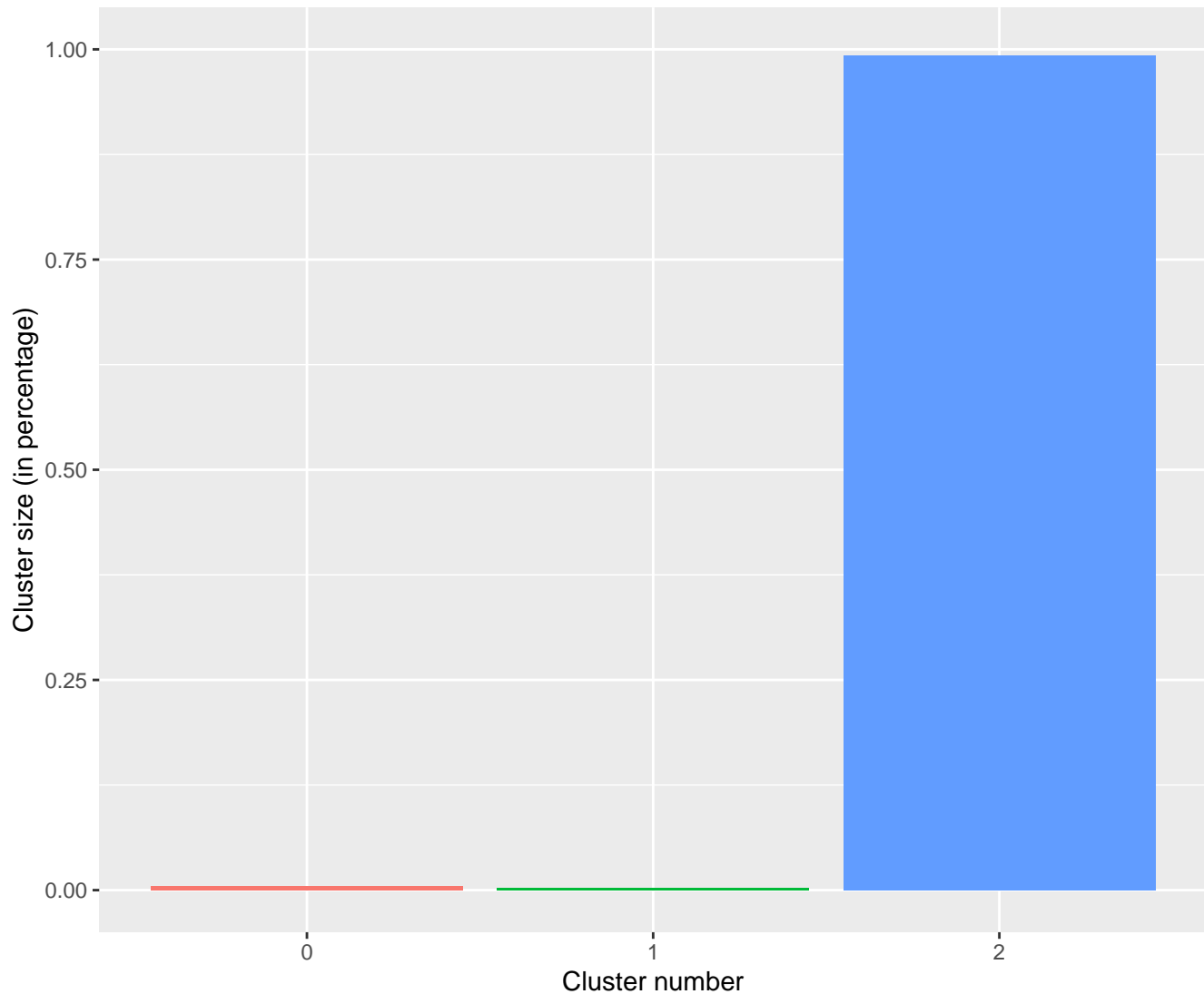


# Clustering: HDBSCAN

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

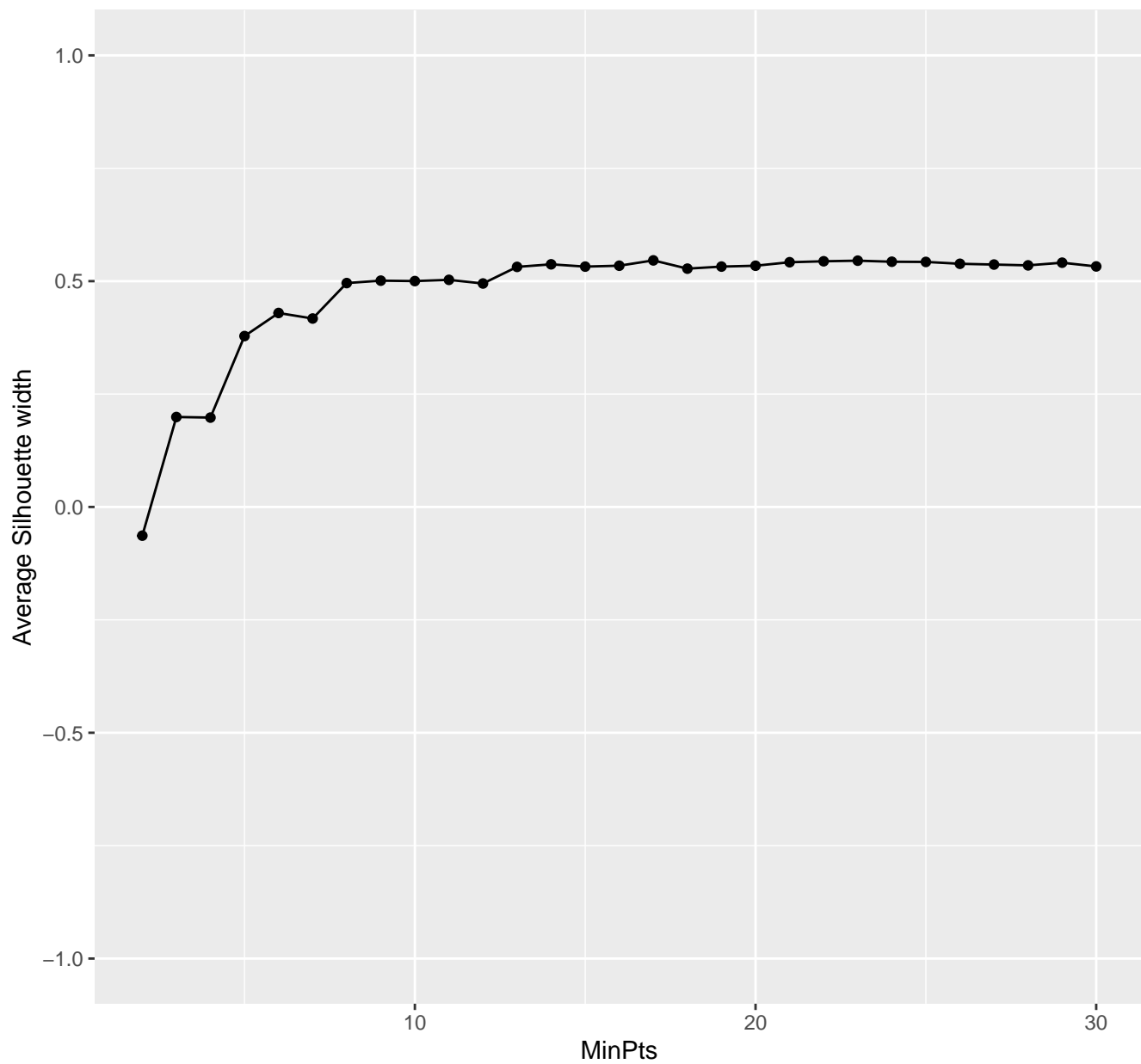
Parameters used: minPoints = 3

Cluster ■ 0 ■ 1 ■ 2



# HDBSCAN elbow plot

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

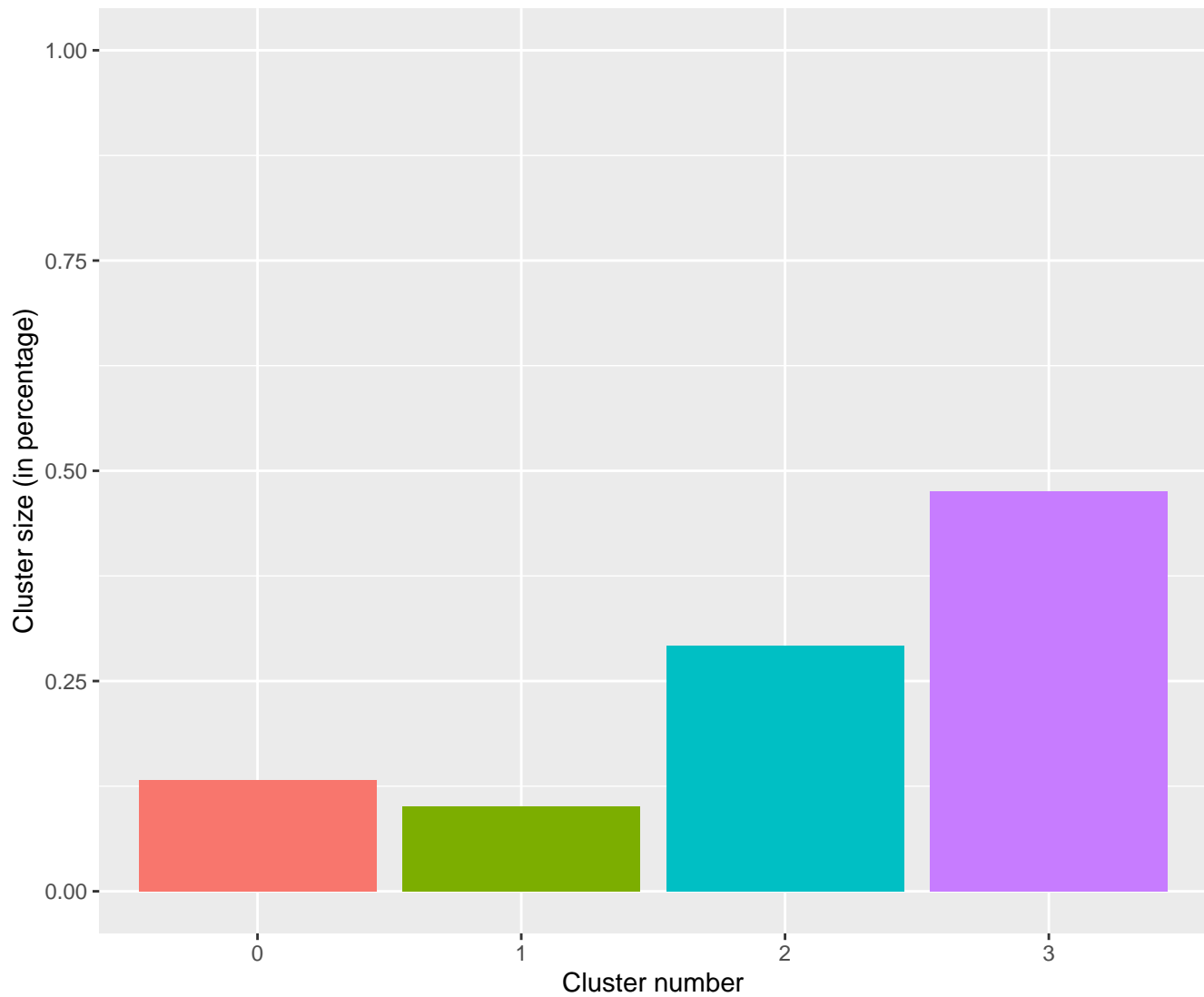


# Clustering: HDBSCAN

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

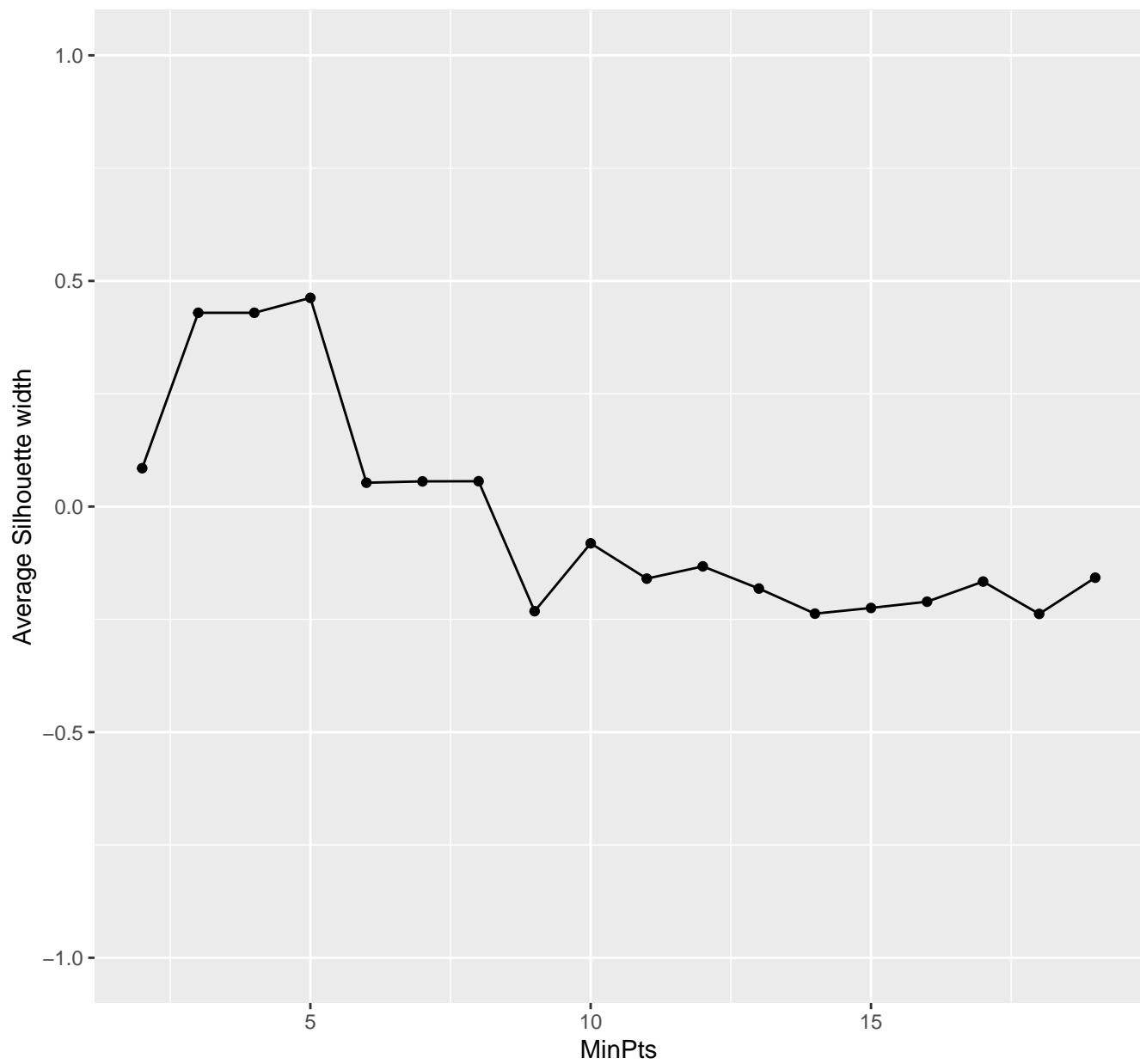
Parameters used: minPoints = 17

Cluster    0    1    2    3



# HDBSCAN elbow plot

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

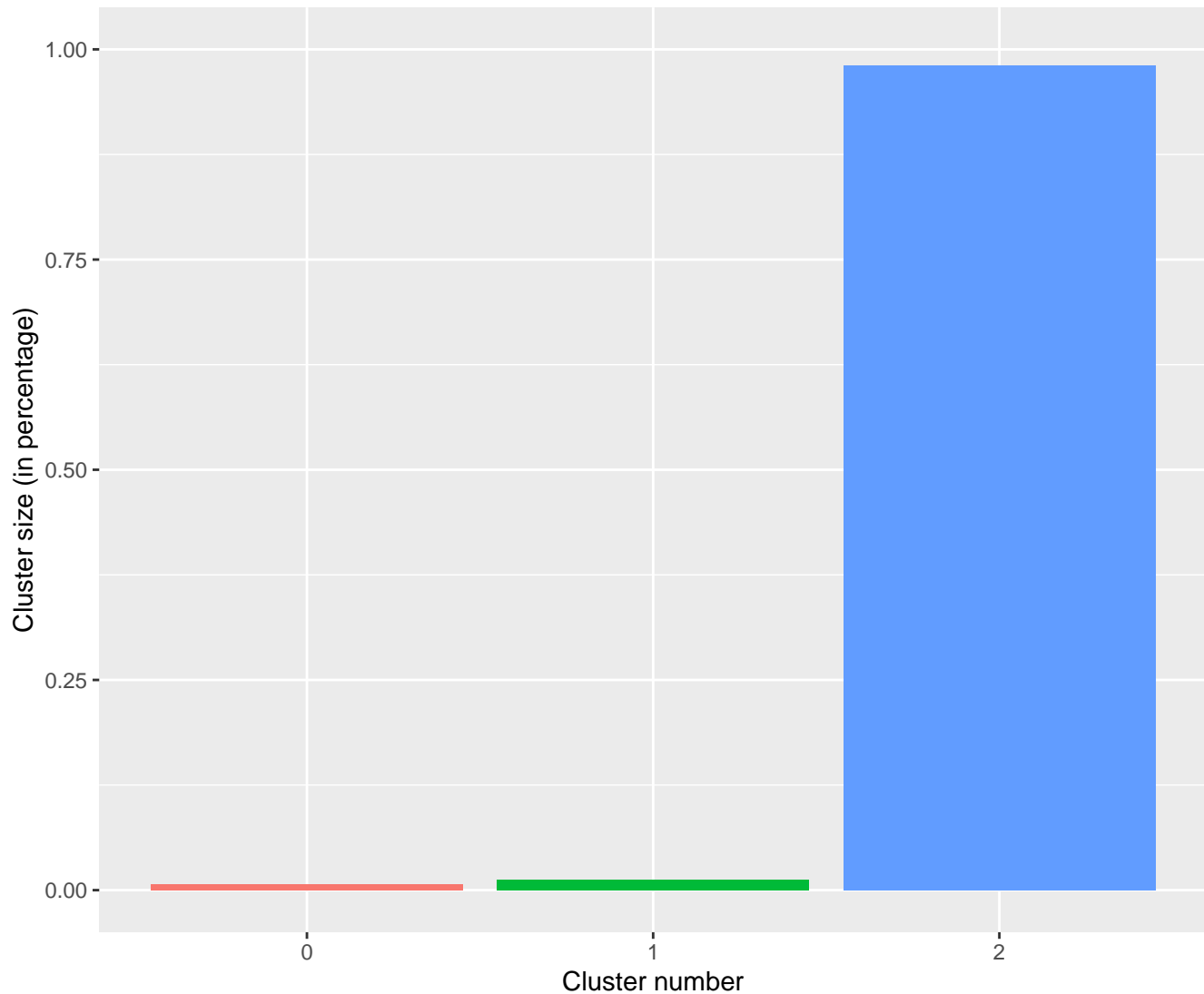


# Clustering: HDBSCAN

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

Parameters used: minPoints = 5

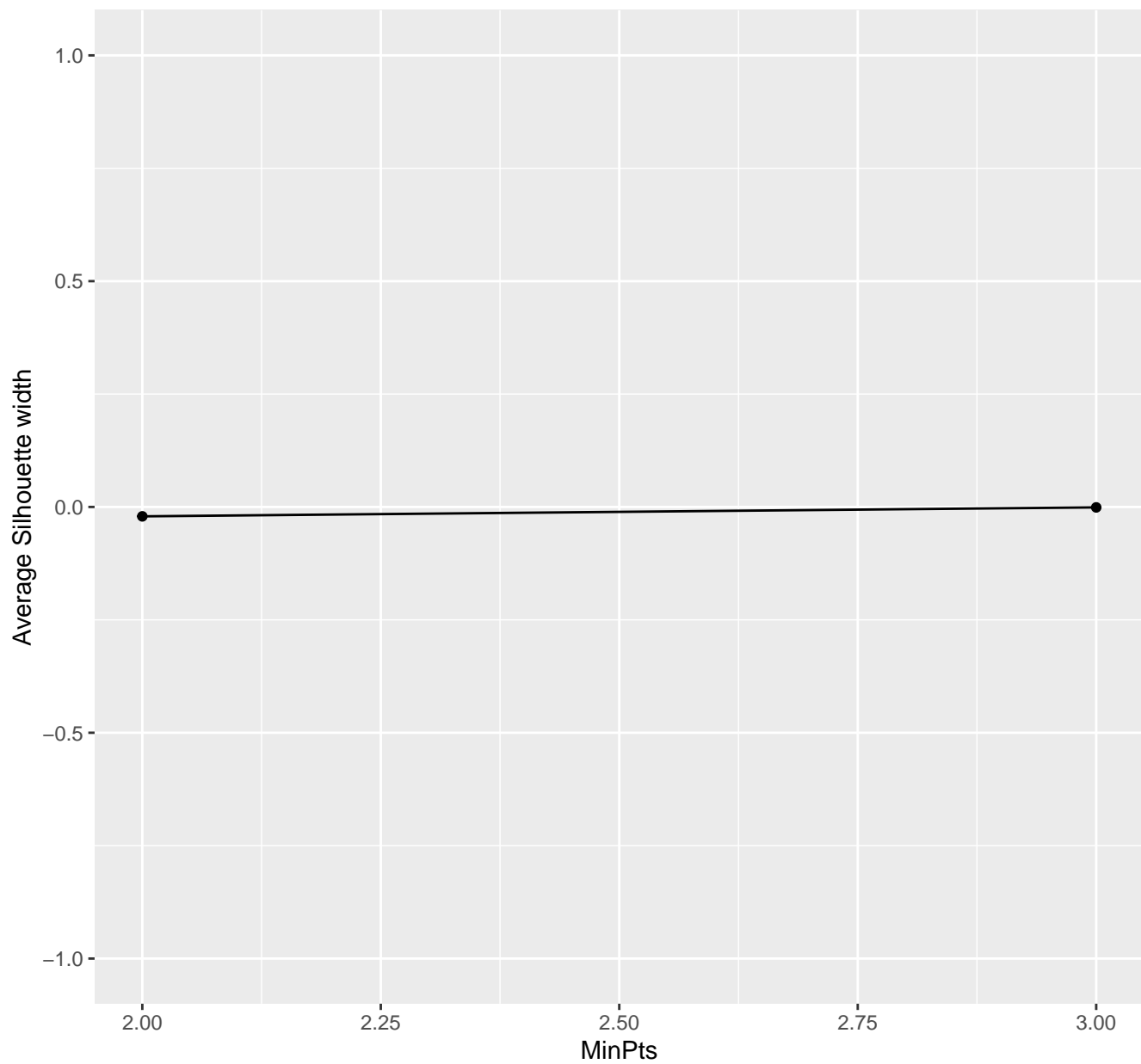
Cluster ■ 0 ■ 1 ■ 2





# HDBSCAN elbow plot

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

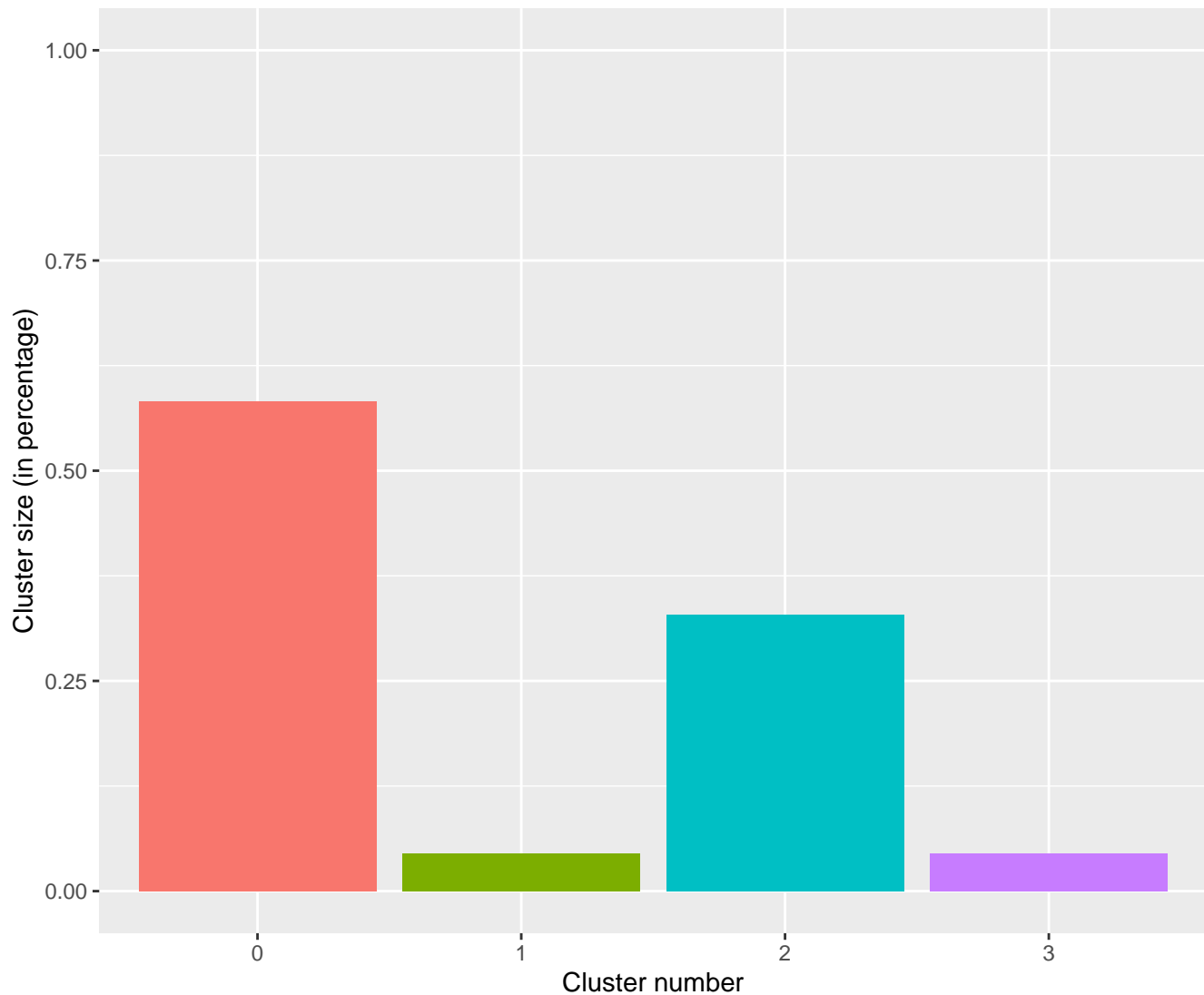


# Clustering: HDBSCAN

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

Parameters used: minPoints = 3

Cluster 0 1 2 3



# Clustering: HDBSCAN

Hyperparameter combinations ranked by its average Silhouette score

