

Silhouette plot of (x = cutree(hc5, k = 2), dist = dist(gene, method = "euclidean"

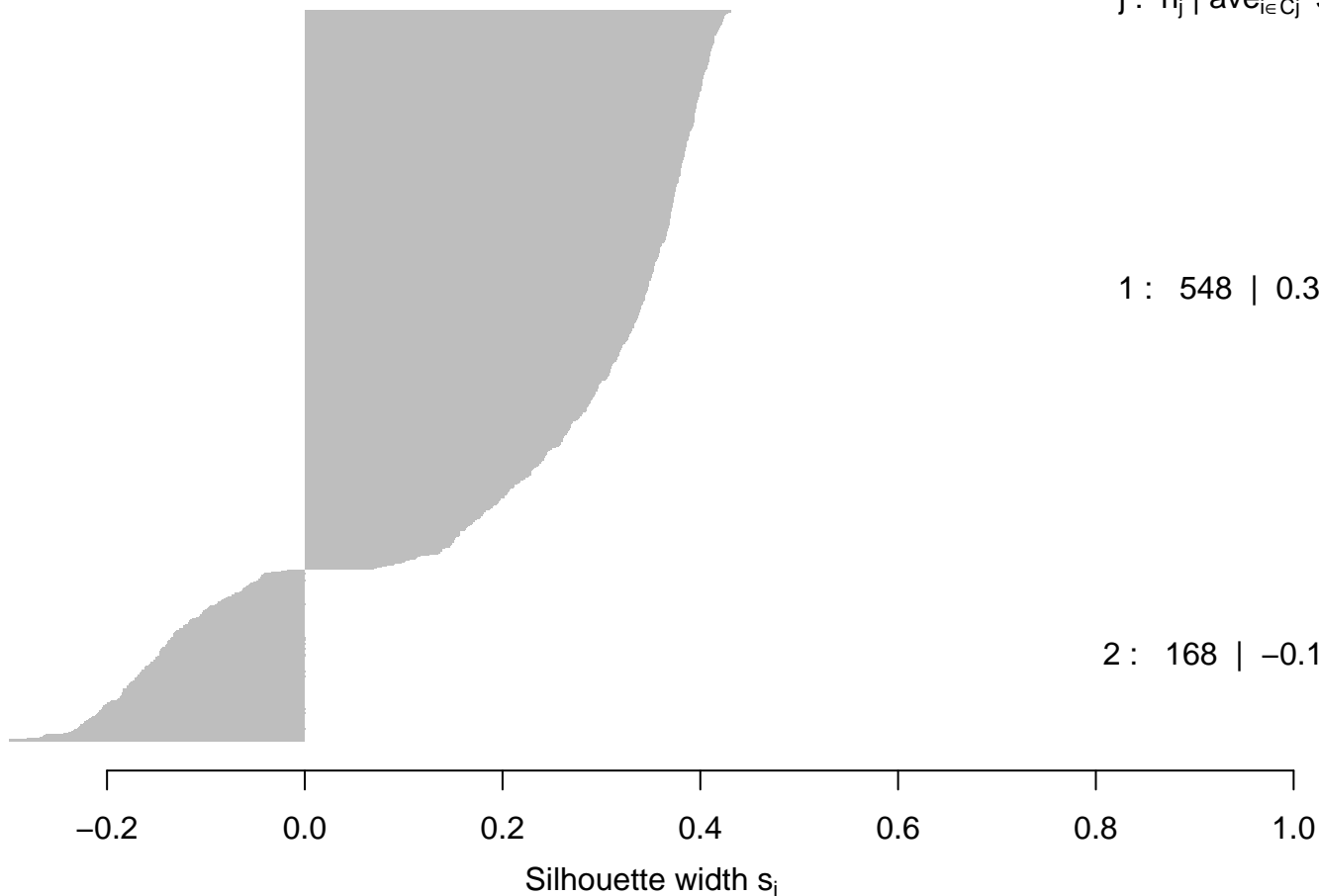
n = 716

2 clusters C_j

$j : n_j \mid \text{ave}_{i \in C_j} s_i$

1 : 548 | 0.32

2 : 168 | -0.15



Average silhouette width : 0.21