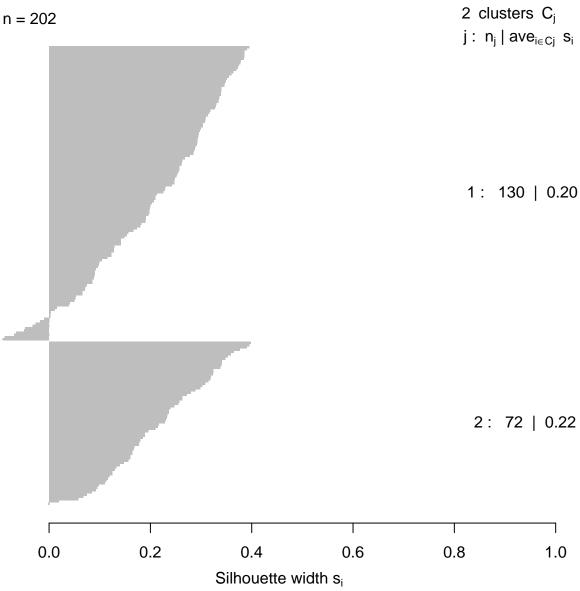
## Silhouette plot of pam(x = dist(distmatrix), k = k, diss = TRUE)



Average silhouette width: 0.21

PAM Metapatients/with\_entire\_count\_matrix/best\_k/LungCancer\_combat\_counts GSM214260&SM2142592 50 -GSM2142515 GSM2142449 0 -GSM21425 M2' Dim2 (10.3%) cluster GSM2142448SM2142522 2 M2142465 GSM2142618 -50 **-**GSM2142466 GSM2142568 GSM21/42602 -100 **-**A2620 -100-50 50 100 150 Dim1 (11.3%)

