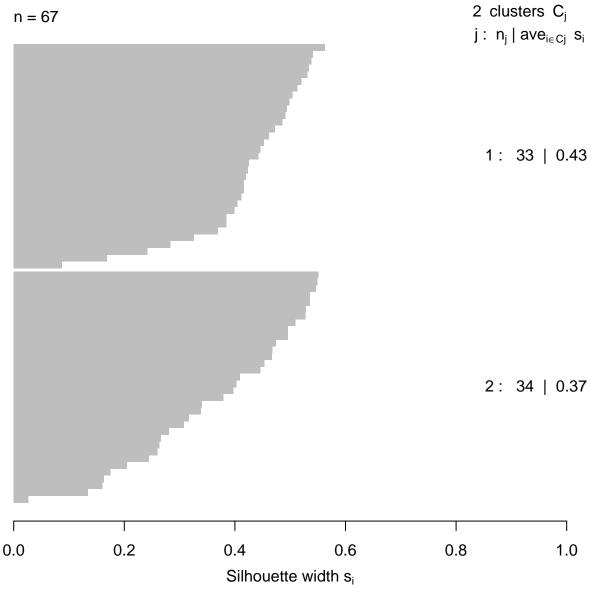
Silhouette plot of pam(x = dist(distmatrix), k = k, diss = TRUE)



Average silhouette width: 0.4

PAM Metapatients/with_entire_count_matrix/best_k/Ulcer_normalized_counts -GSM1598683 GSM1598458 GSM1598429 G\$M1598570 GSM1598540 **GSM15**2 GSM1598430 GSM1598670 GSM\$108698506 GSM1598546 0-M1\$98577 GSM1598434 GSM1598503 GSM159846 Dim2 (13.4%) GSM1598583 cluster GSM15985 GSM1598502 2 GSM1 598549 GSM159859 GSM1598545 G\$M1598455 -100 **-**3M1**5**98661 GSM15984 100 -100 -50 50 Dim1 (22.2%)

