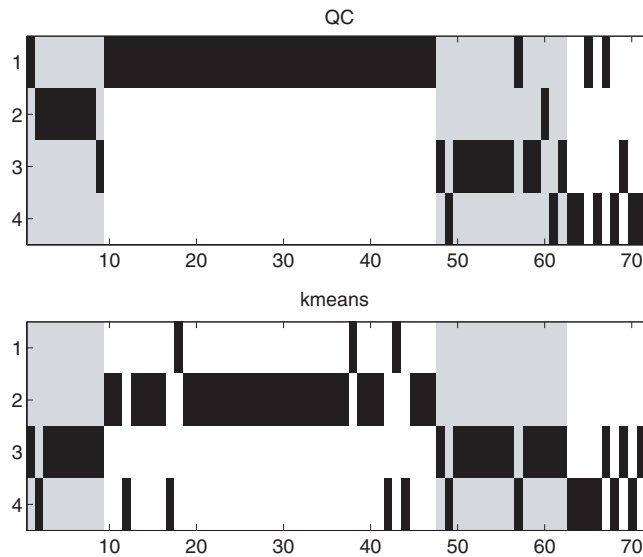


# Erratum

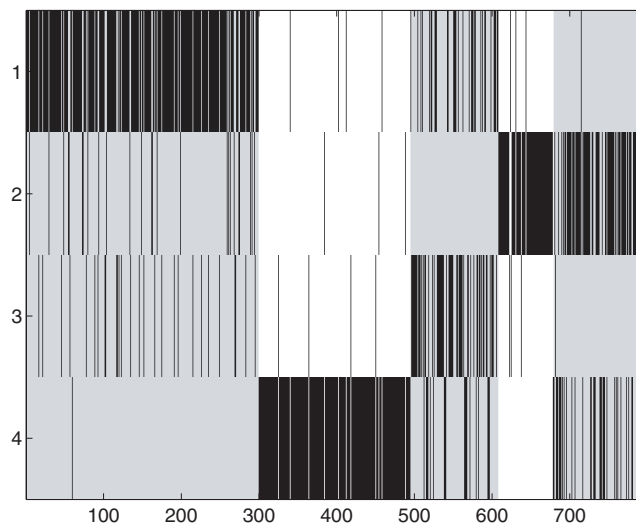
## Novel clustering algorithm for microarray expression data in a truncated SVD space

David Horn and Inon Axel  
*Bioinformatics* (2003) **19**, 1110–1115

The publisher wishes to apologize for accidentally printing the wrong versions of Figures 5 and 7 due to technical problems. The correct version of these figures is printed below.



**Fig. 5.** Clustering solutions for the AML/ALI, problem using QC with  $n = 0.54$  (upper frame) and  $k$ -means with  $k = 4$  (lower frame). The samples are ordered on the  $x$ -axis according to the true classification into four groups, indicated by alternative gray and white areas.



**Fig. 7.** Cluster assignments of genes for QC with  $\sigma = 0.46$  as compared to the classification by Spellman *et al.* (1998) shown as alternating gray and white areas.