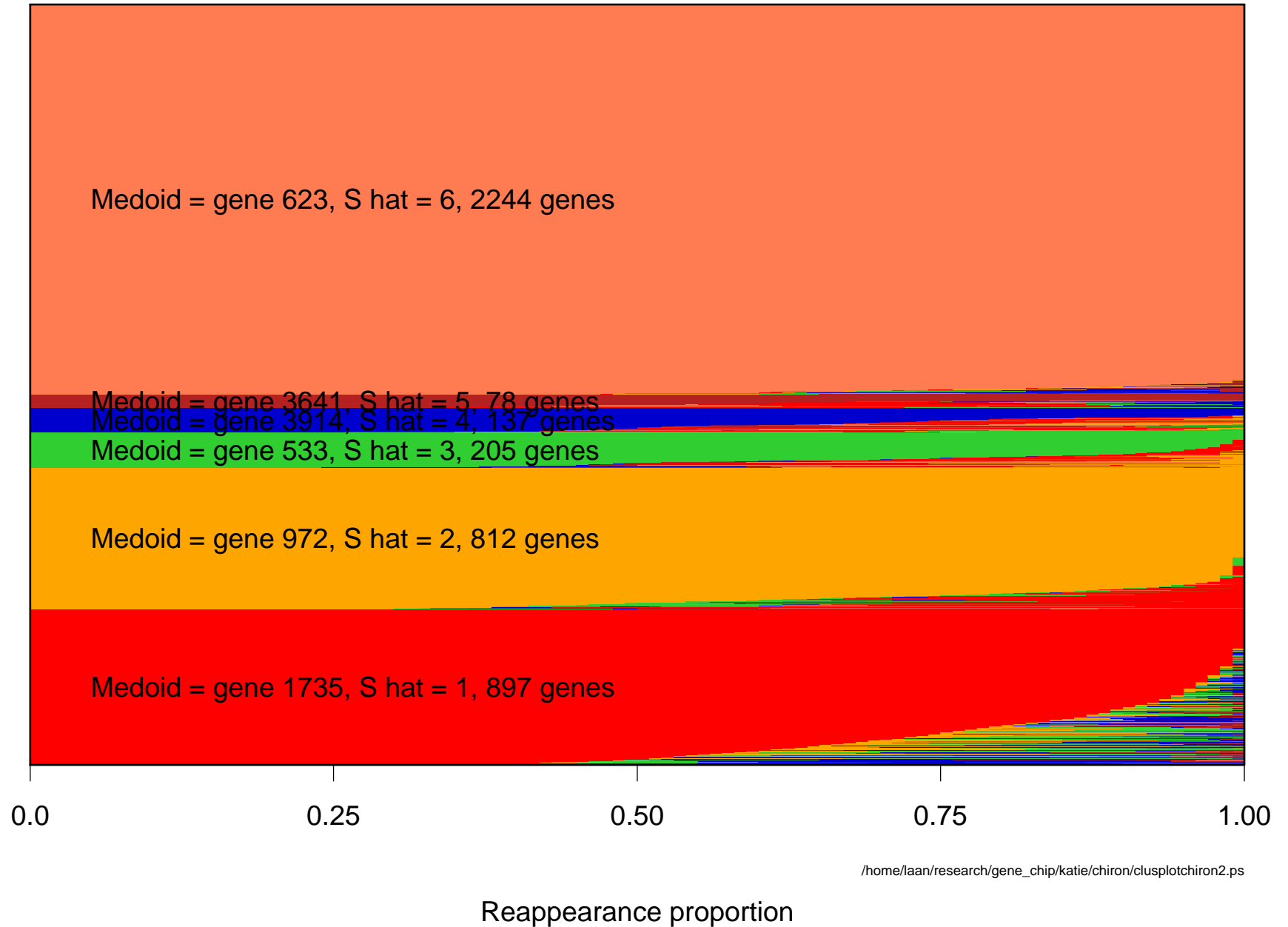


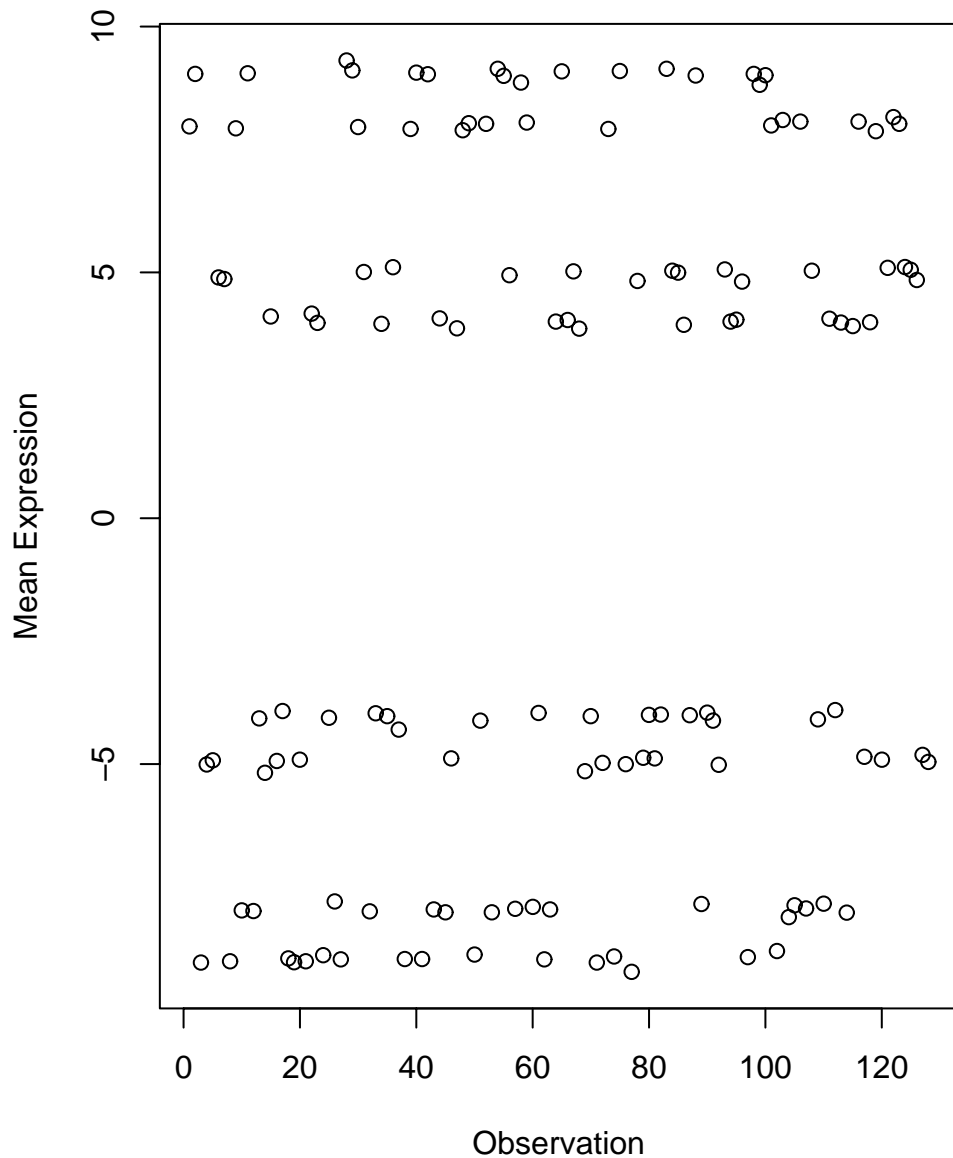
Reappearance proportions and cluster reproducibility

Subset contains 4373 genes.

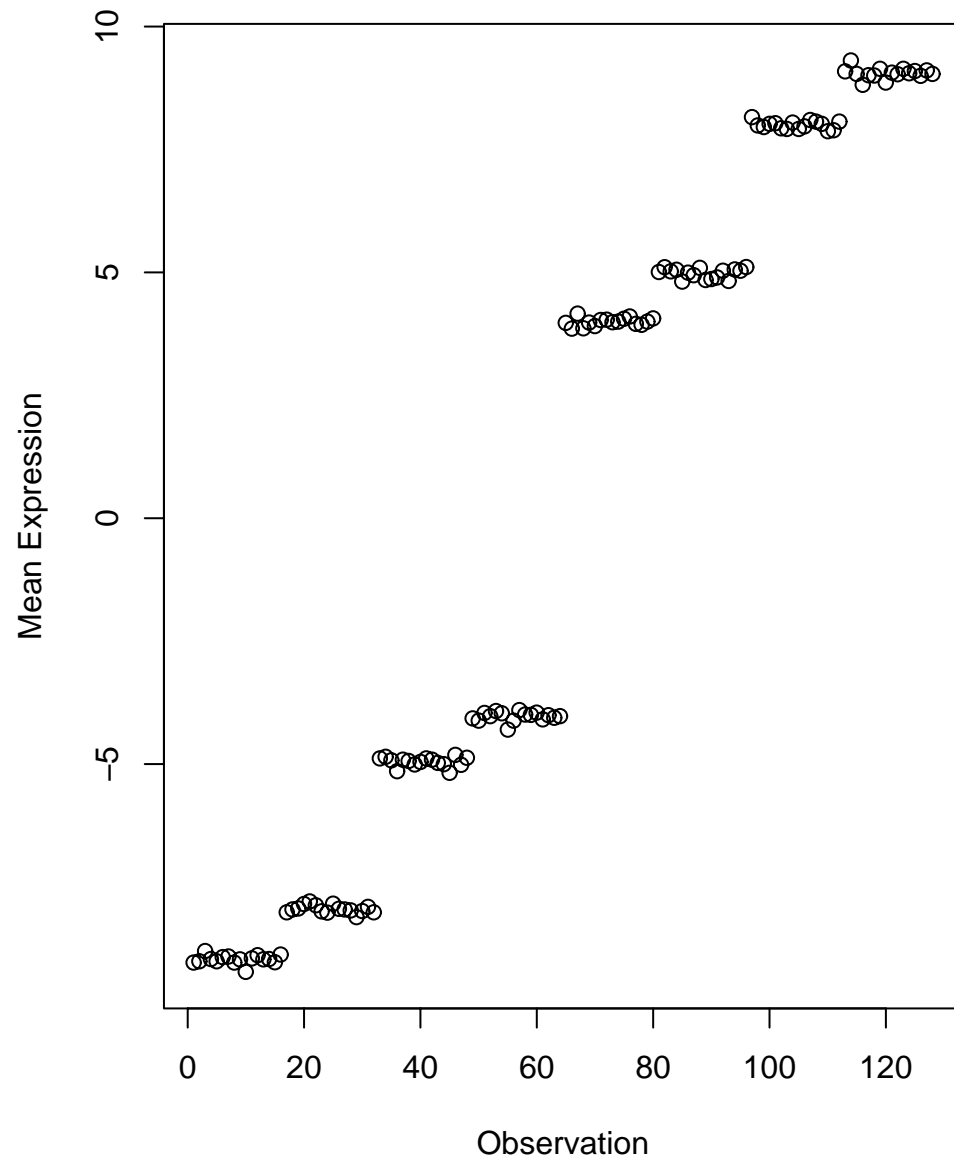


Data from 8 clusters of size 16: $\mu=(-9,-8,-5,-4,4,5,8,9)$, $\sigma=0.1$

Randomized Data



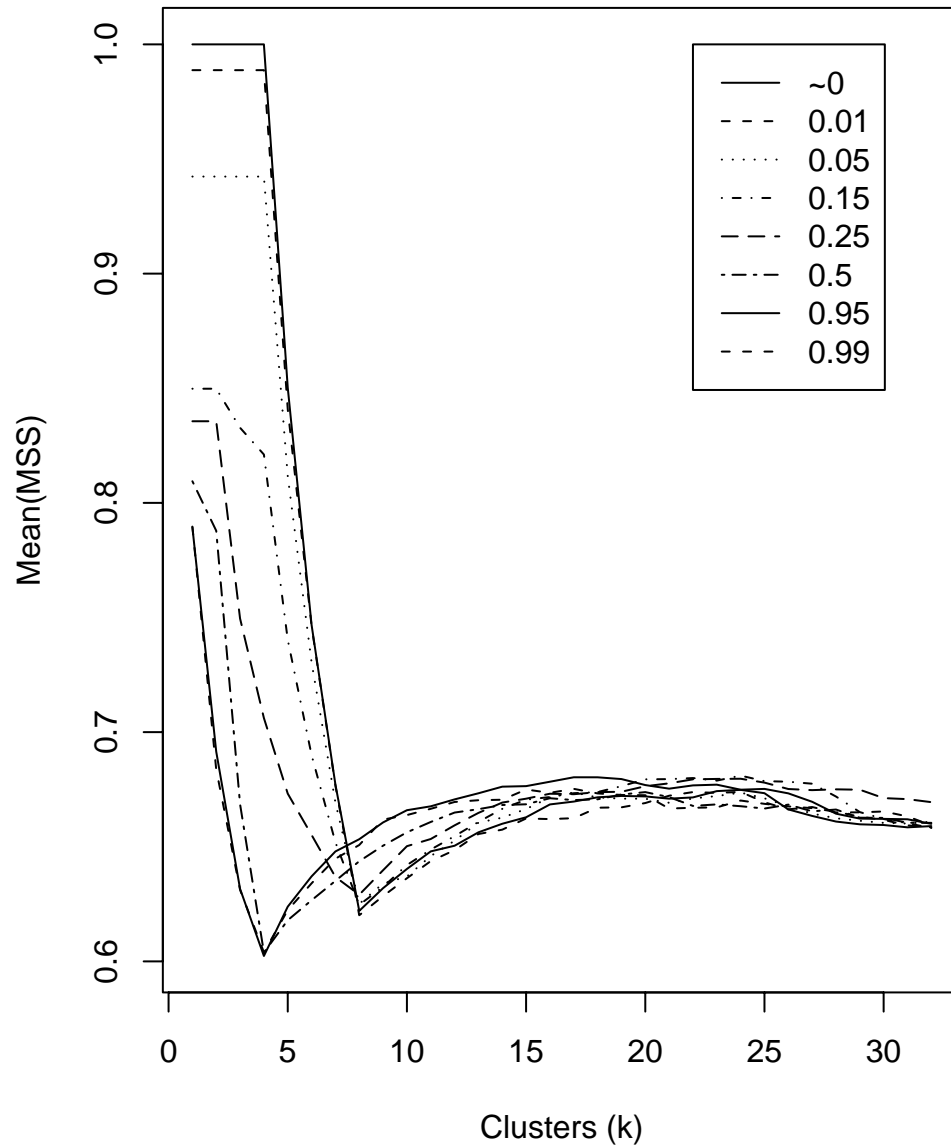
Ordered Data



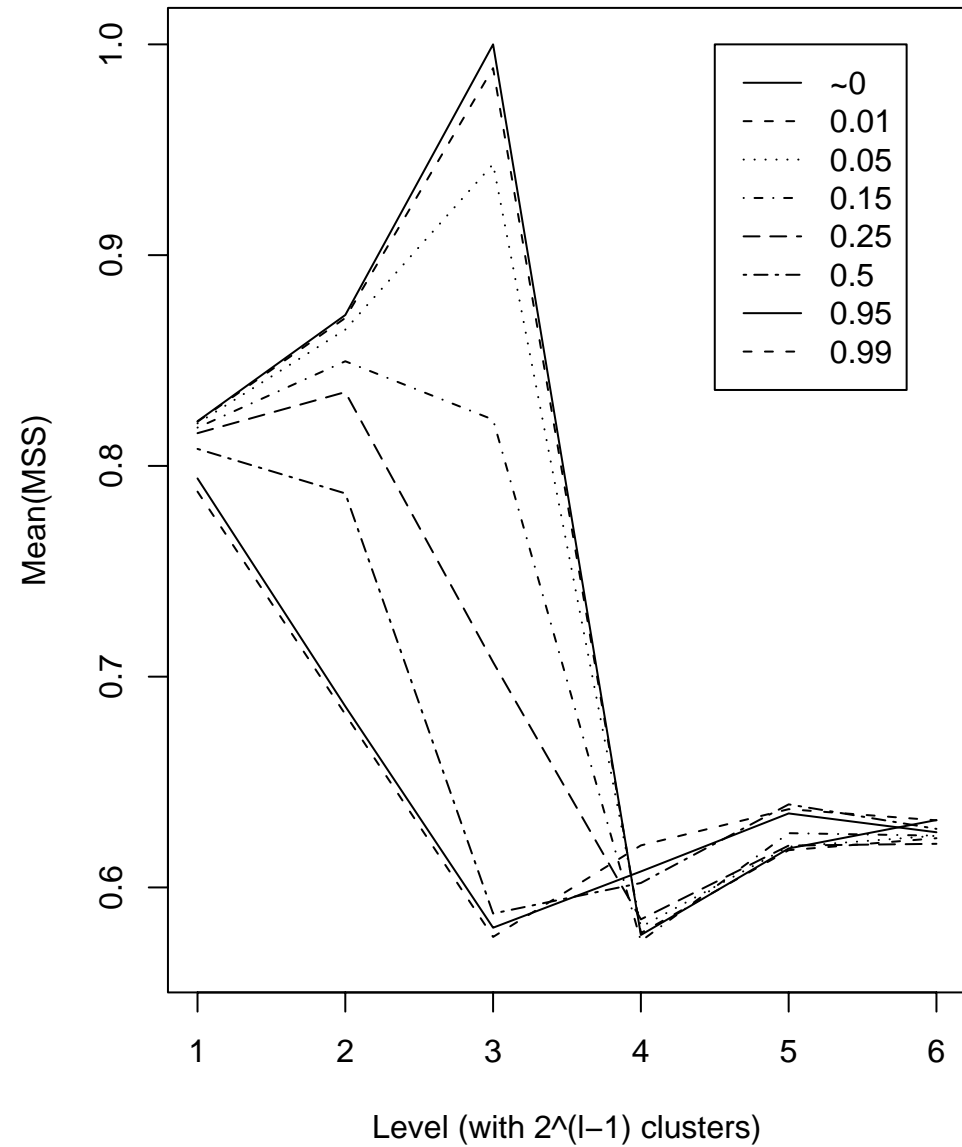
MSS versus number of clusters for eight values of sigma

$n=360, \mu=(1,2,5,6,14,15,18,19)$

PAM (MSS for k clusters)

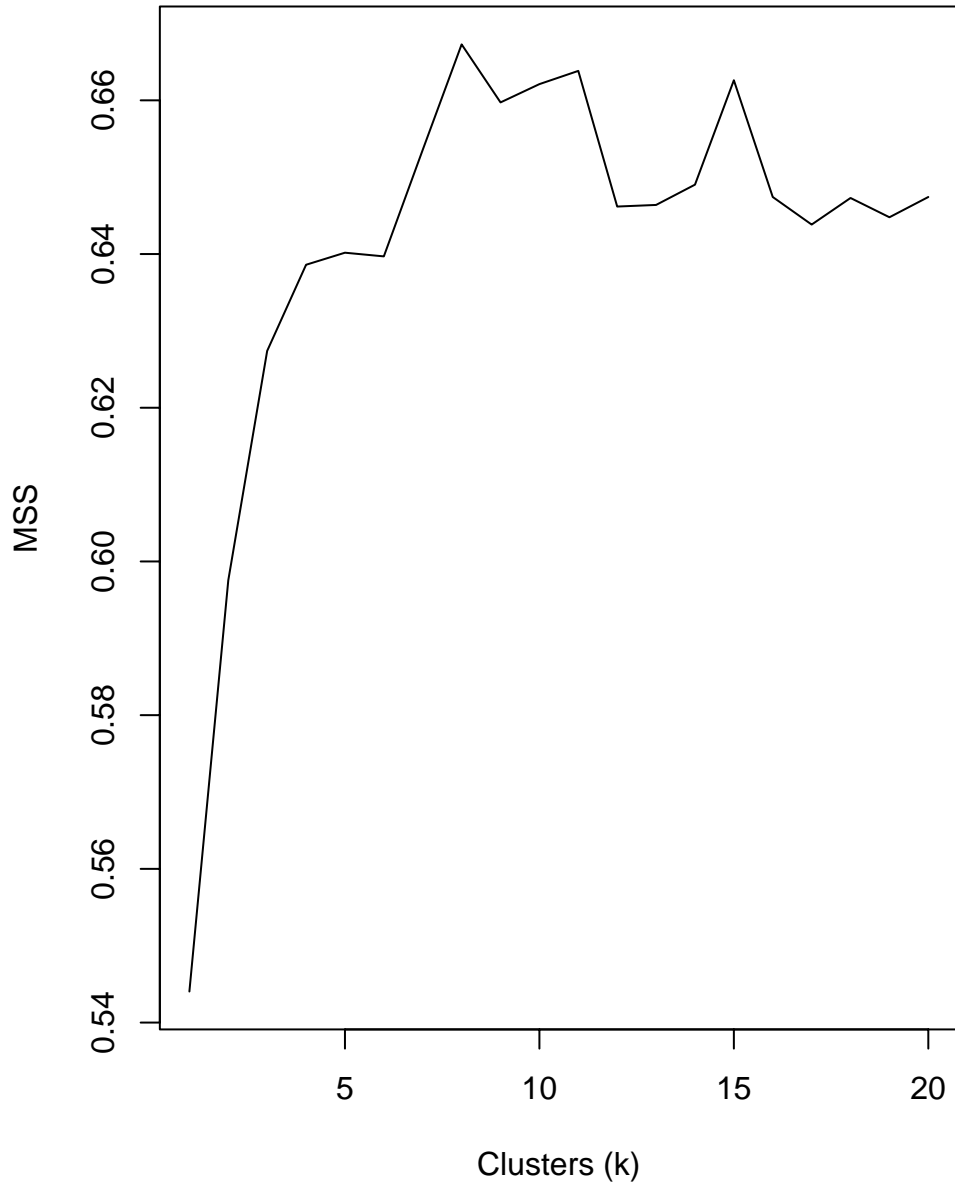


HOPACH (MSS for l levels)



Univariate normal data: n=360, p=1, mu=0, sigma=0.05

PAM



HOPACH

