Pearson dissimilarity, complete linkage

Height

0.5

0.4

0.3

0.2

0.1

0.0
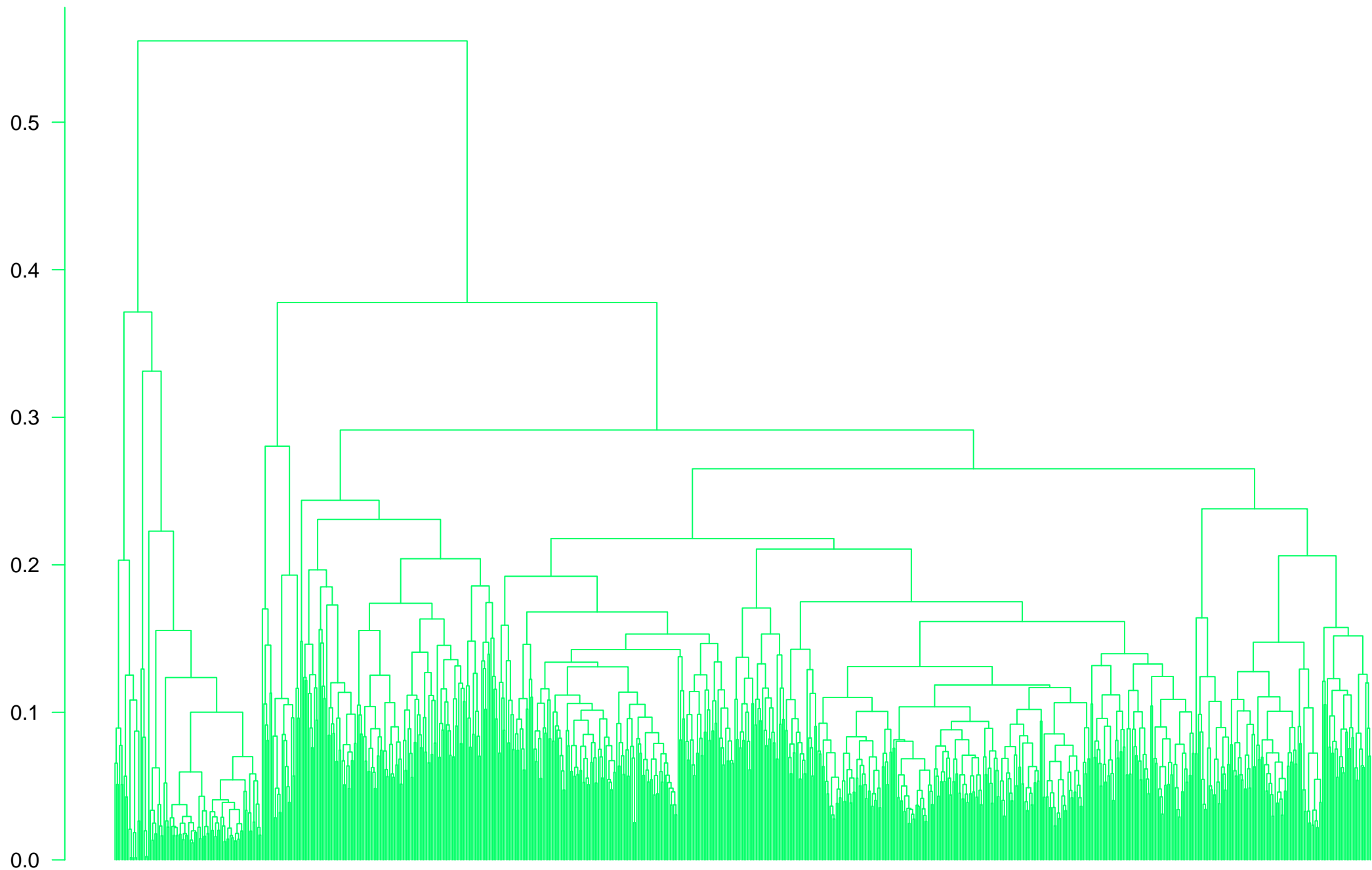
BIC.dist
hclust (*, "complete")