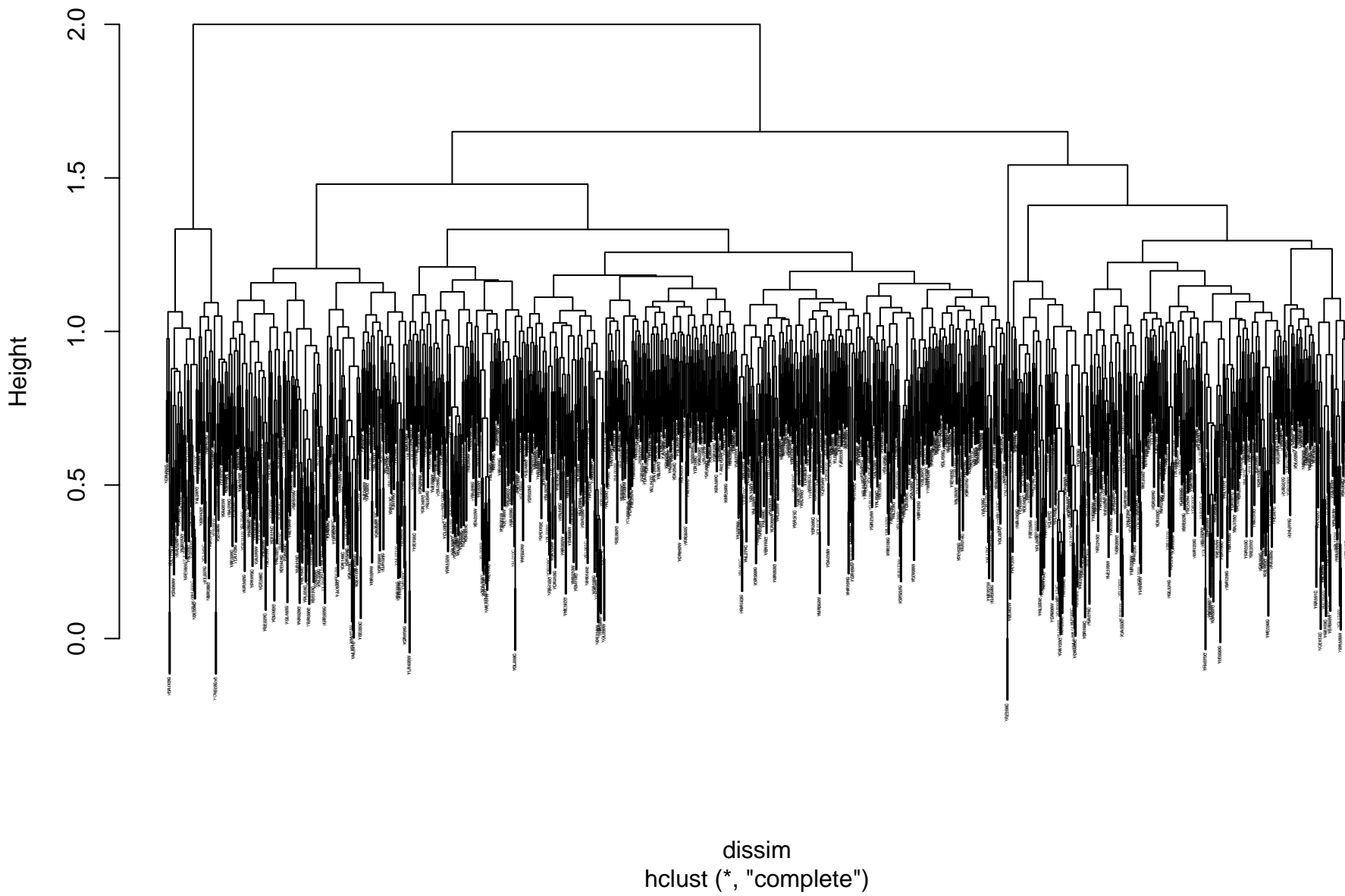
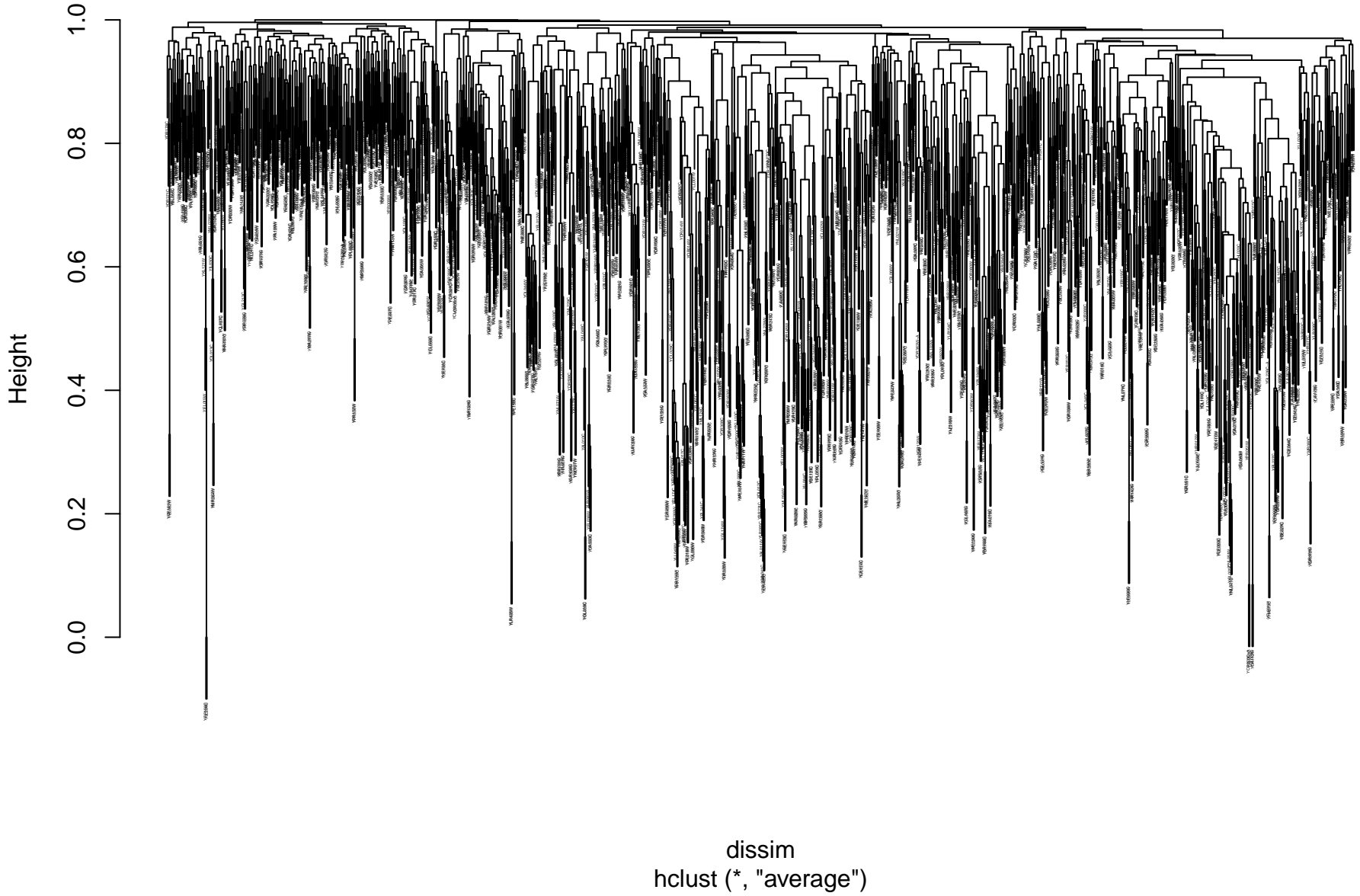


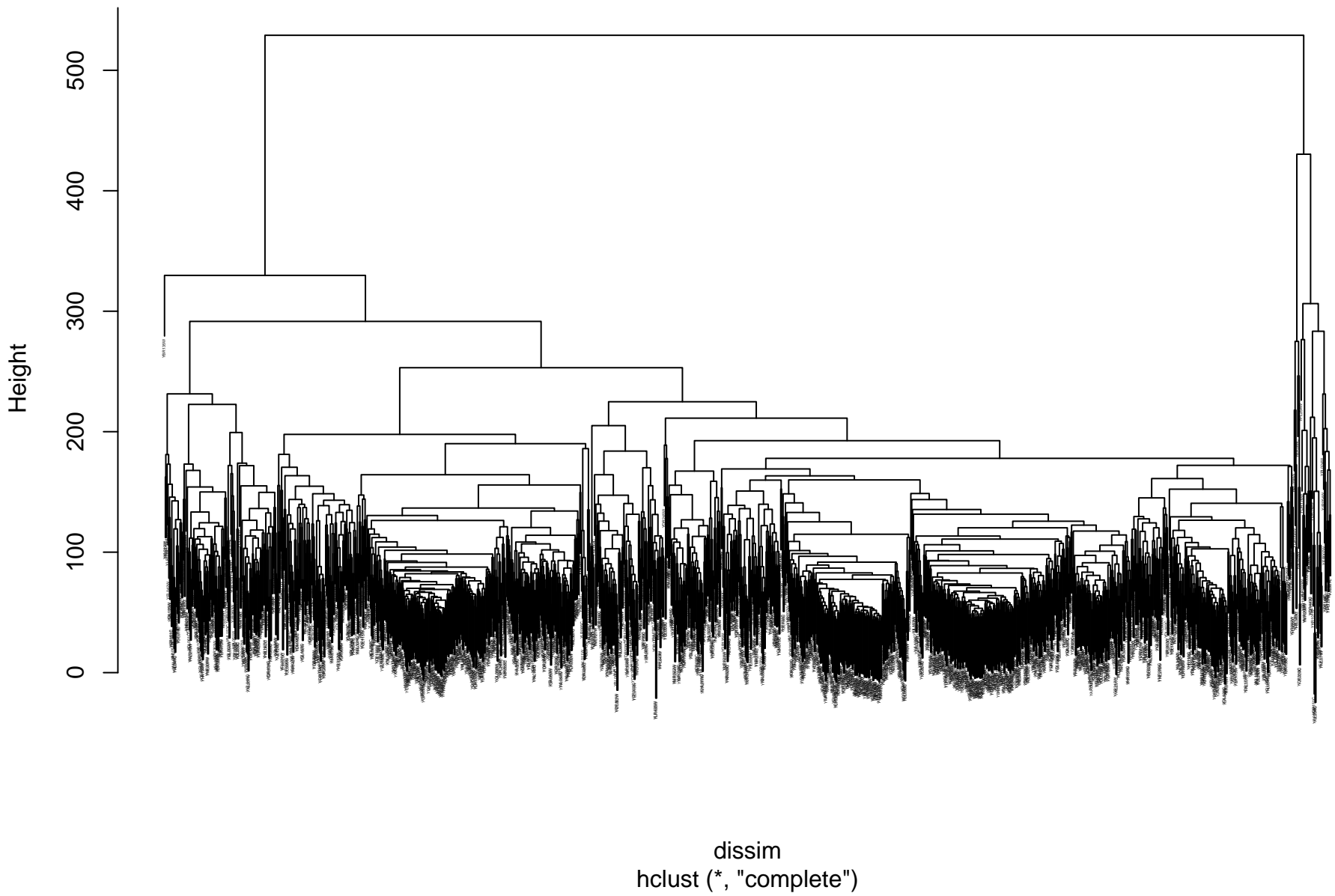
# library\_pearson\_complete

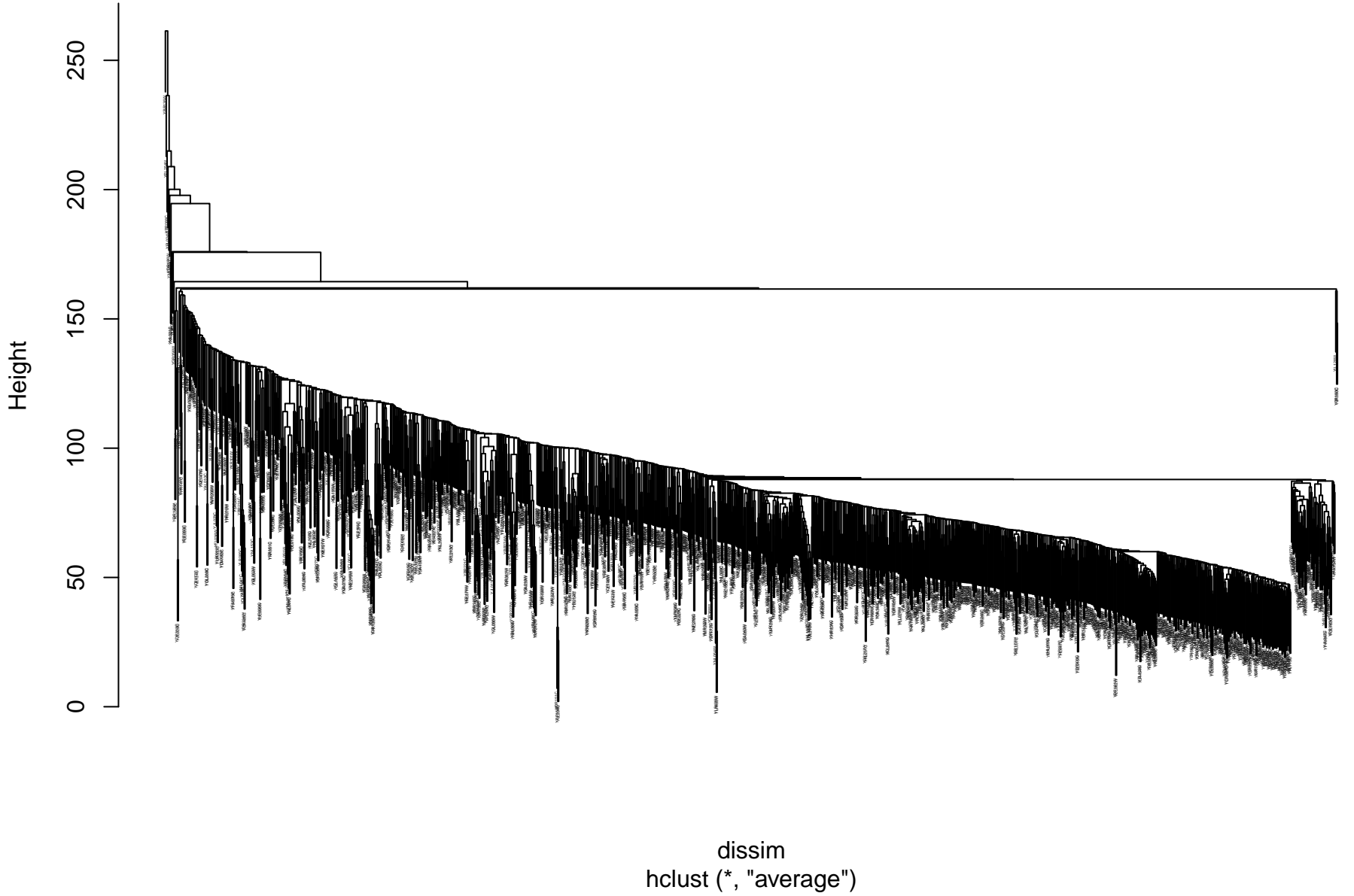


**library\_pearson\_average**

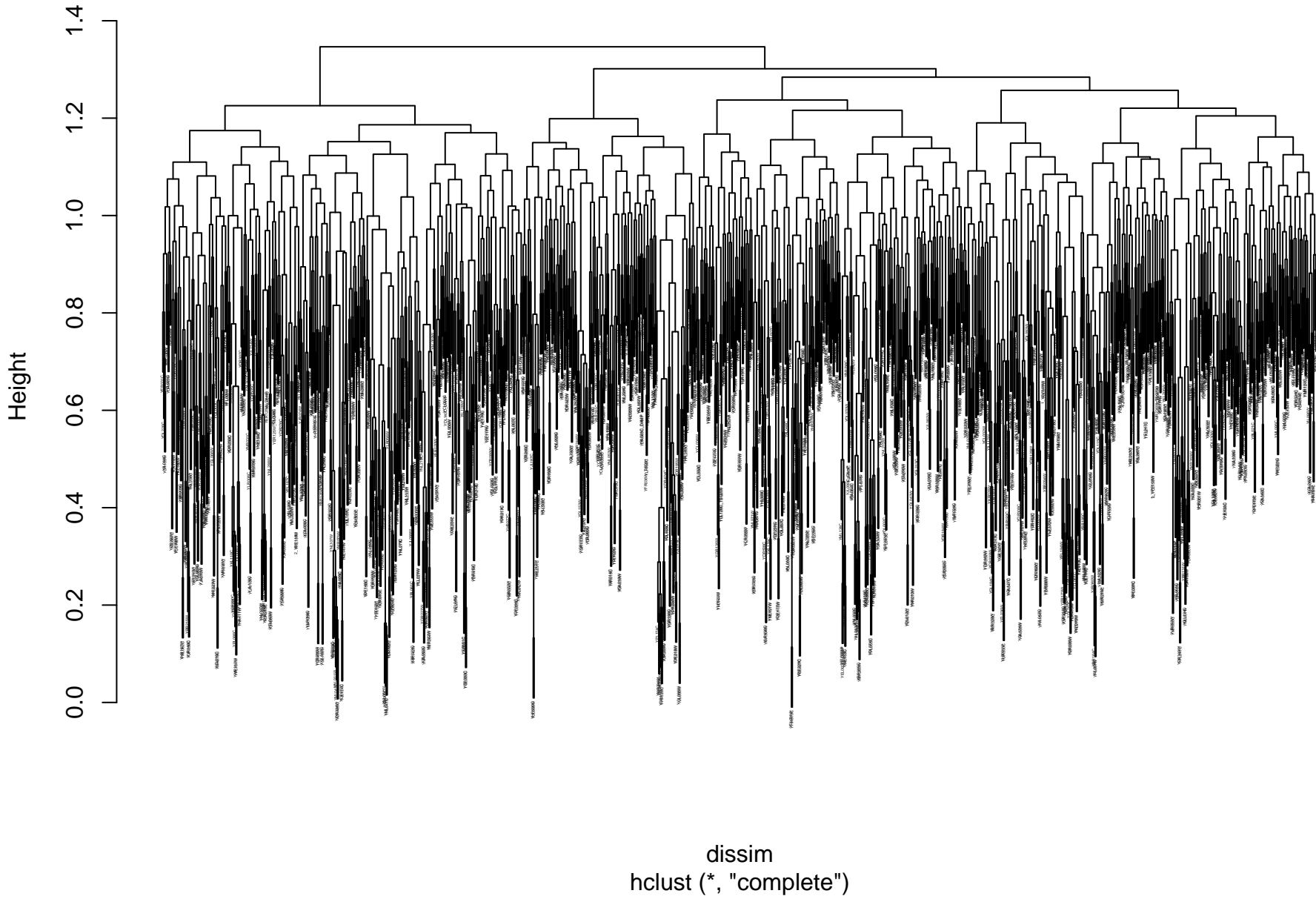




**library\_euclidean\_complete**

**library\_euclidean\_average**

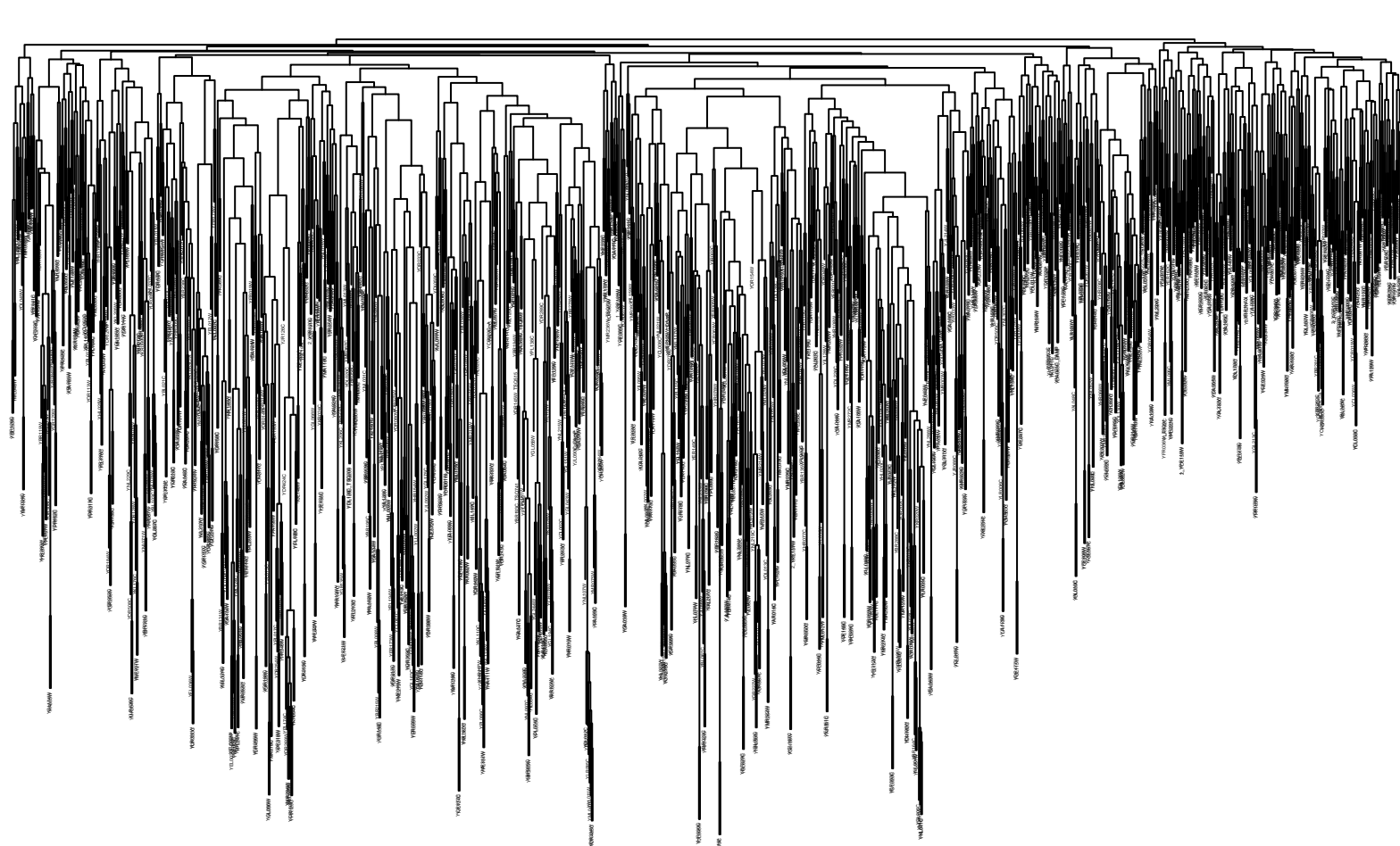
gene\_pearson\_complete



Height

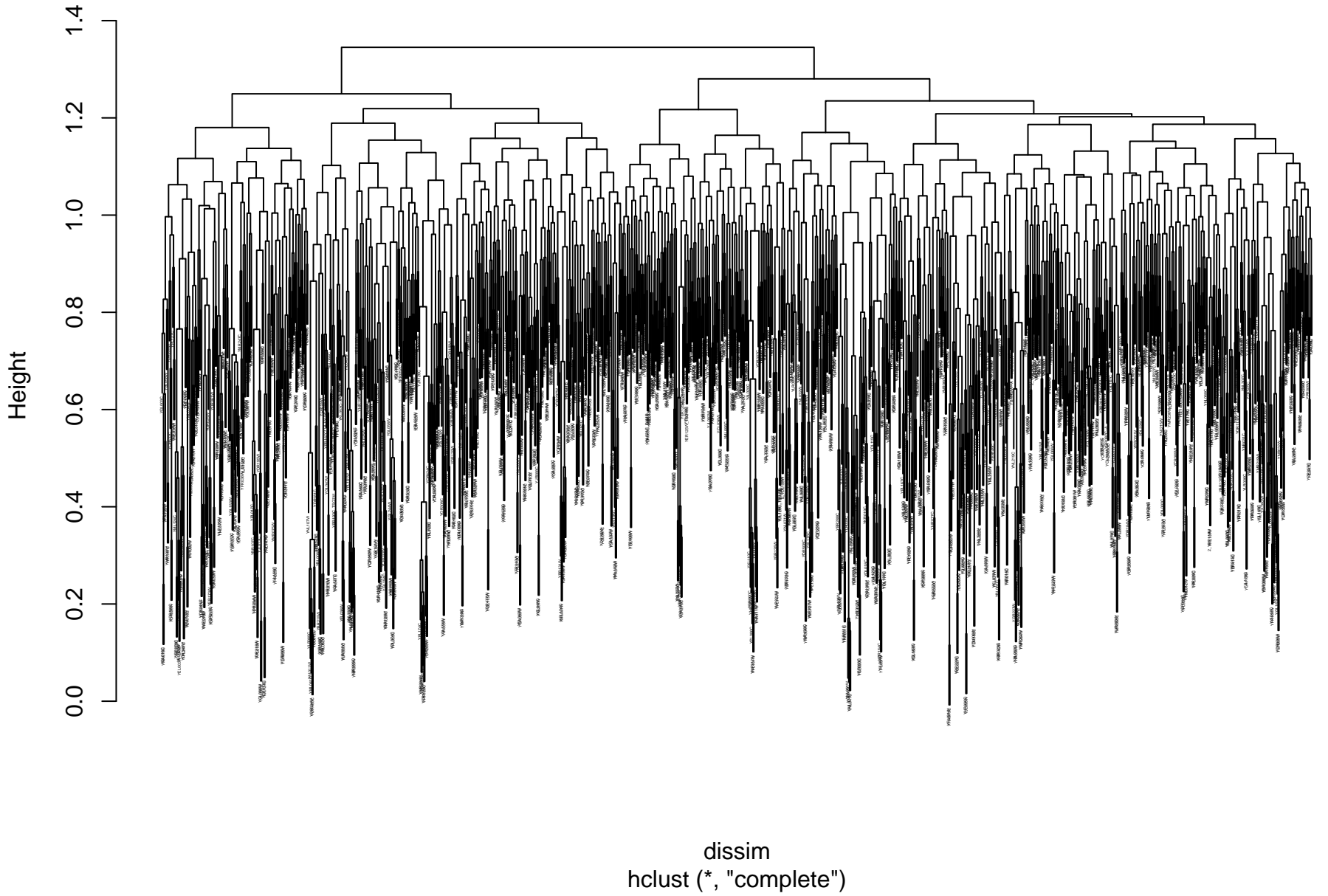
0.0 0.2 0.4 0.6 0.8 1.0

gene\_pearson\_average



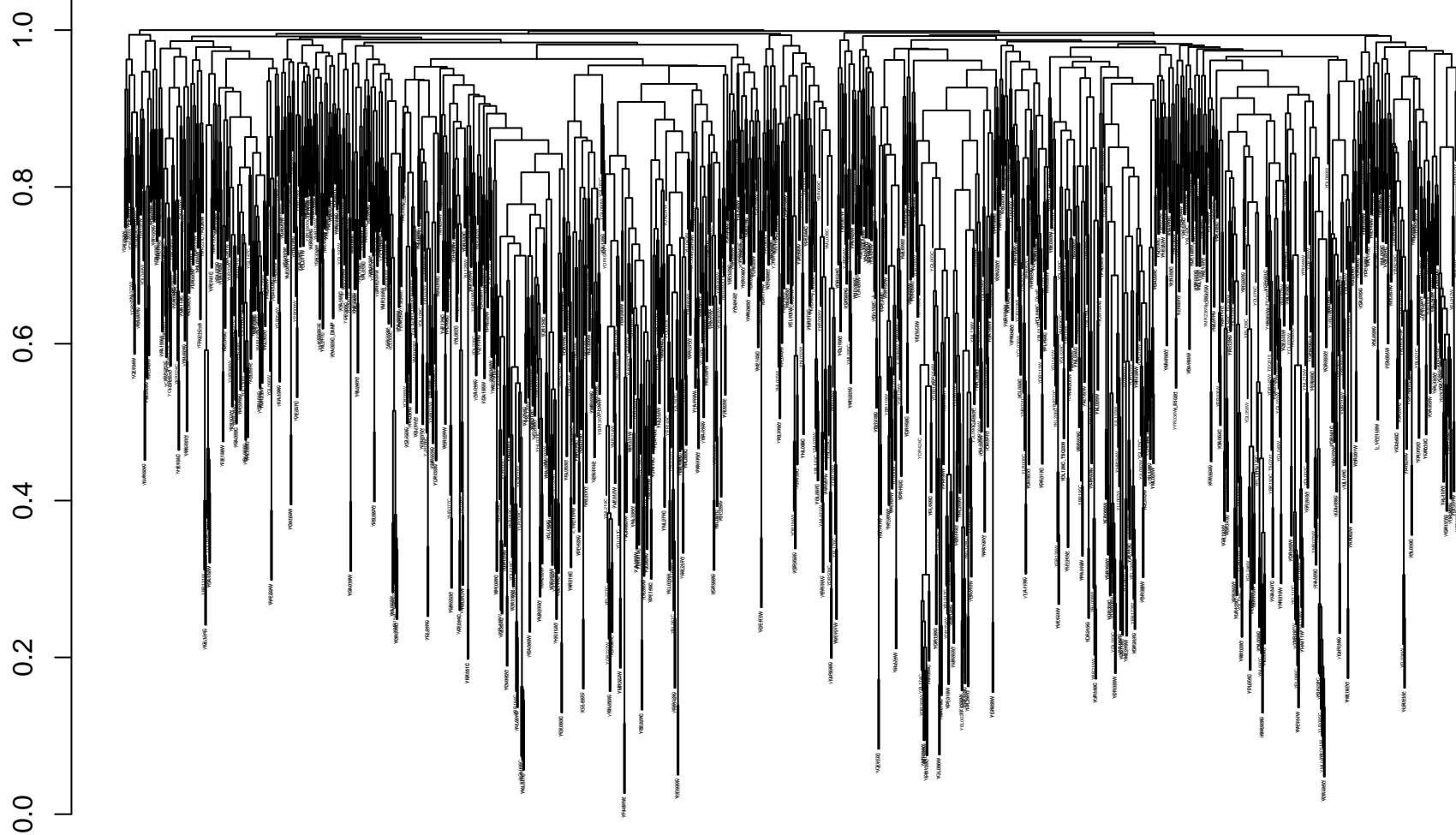


**gene\_correlation\_complete**



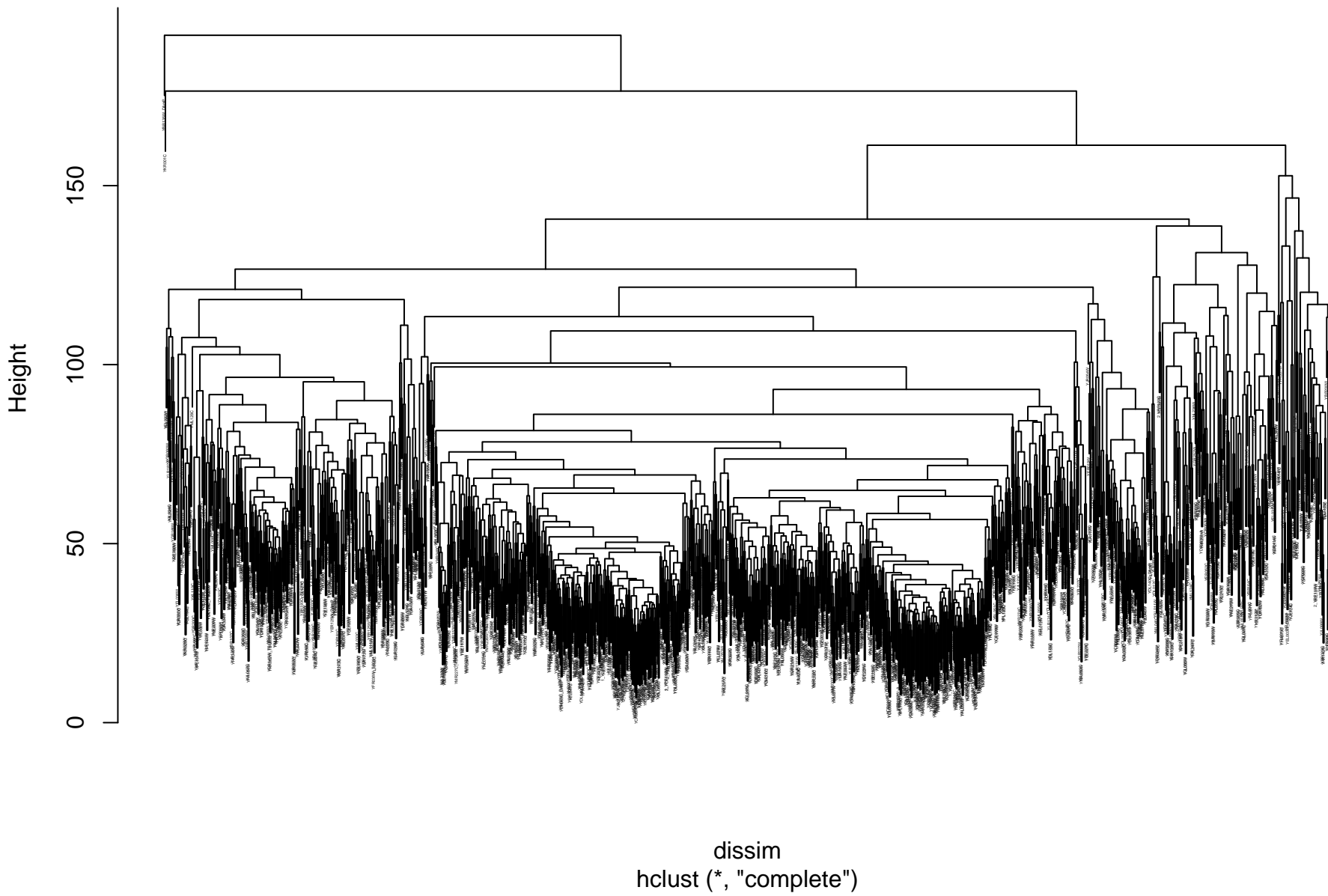
Height

gene\_correlation\_average



dissim  
hclust (\*, "average")

# gene\_euclidean\_complete



gene\_euclidean\_average

