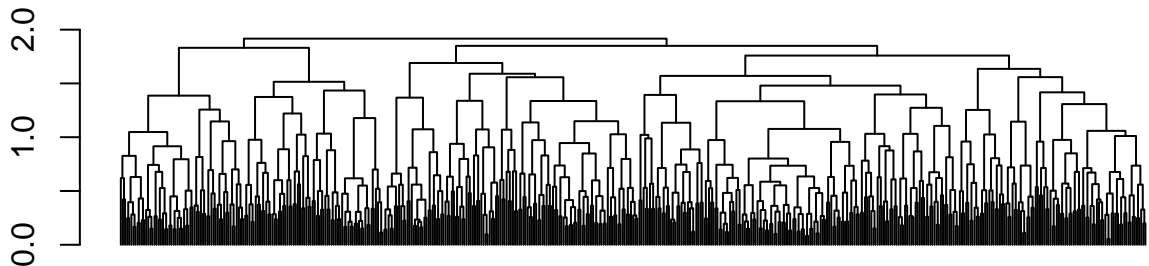


gènes

Height



cor_pearson_gene
hclust (*, "complete")