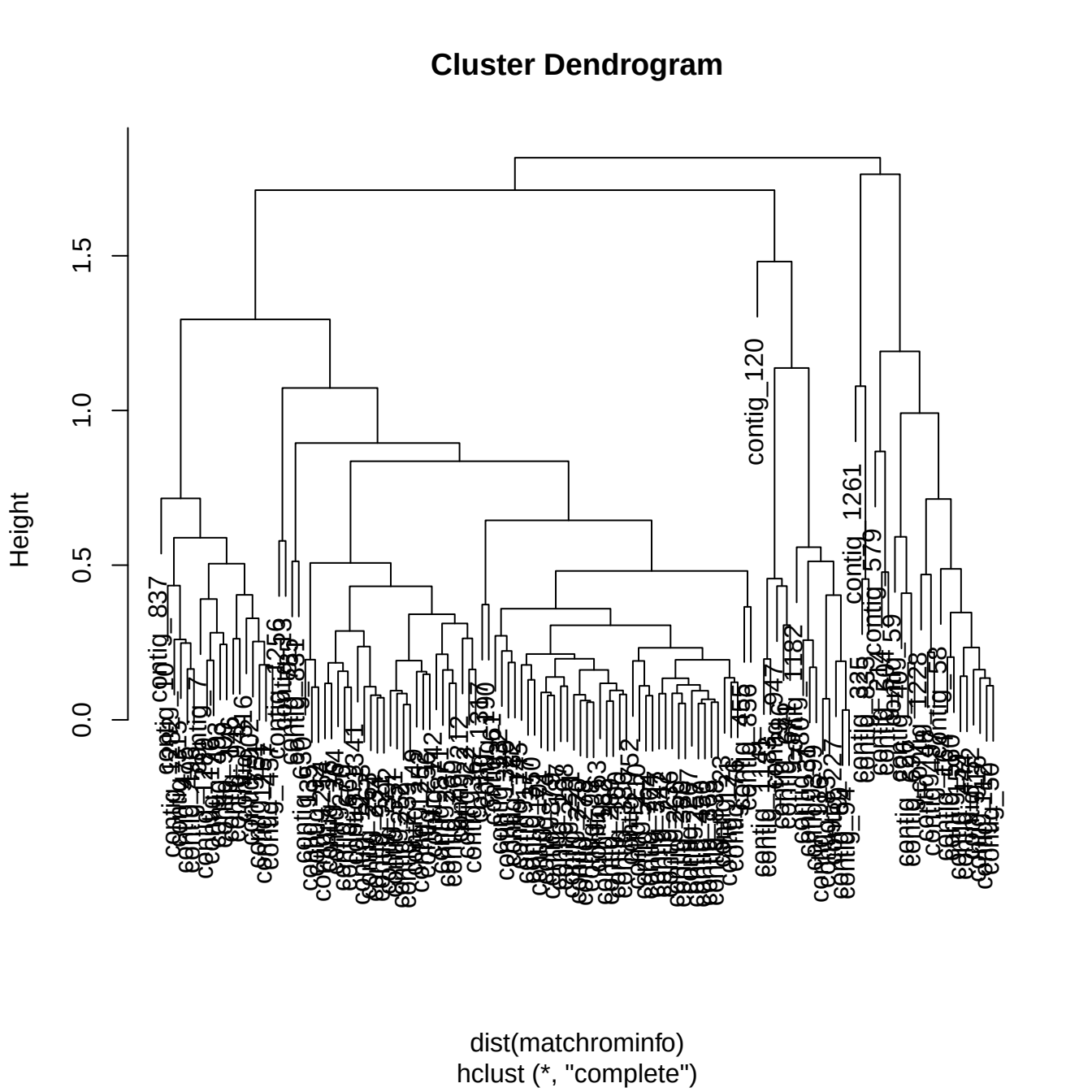
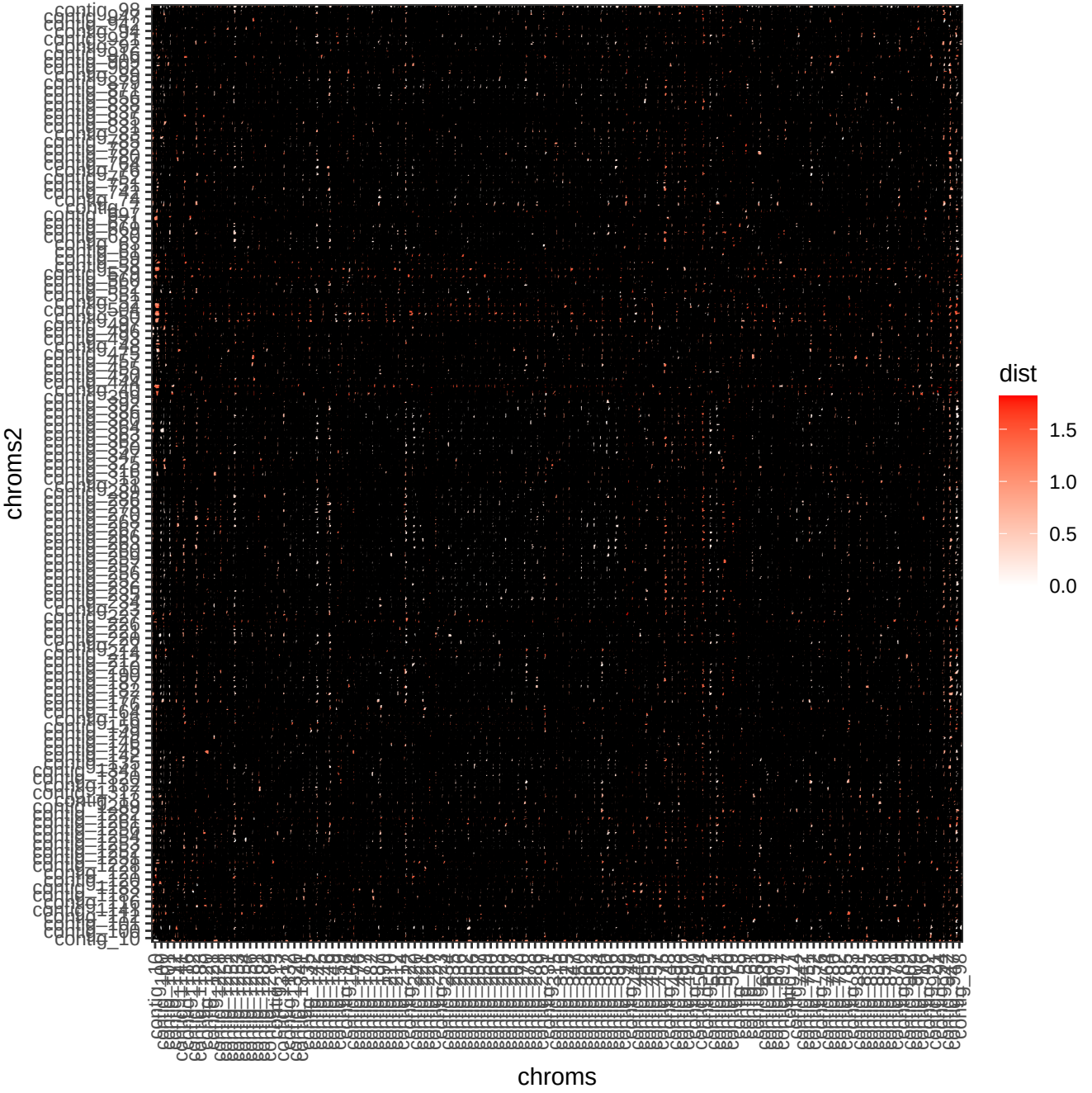

Cluster Dendrogram
Height
0.0
0.5
1.0
1.5
contig_120
contig_1261
contig_579
contig_837
dist(matchrominfo)
hclust (*, "complete")


chroms2
chroms
dist
1.5
1.0
0.5
0.0