cp decomposition glioma 73 test rank5

VIVI TANG

17/11/2021

1. Kmeans

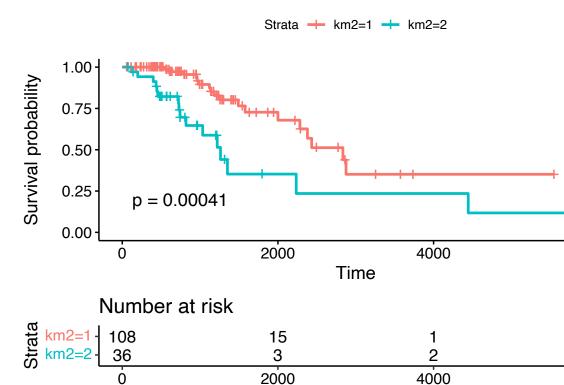
Warning: package 'lattice' was built under R version 3.6.2

 $\mbox{\tt \#\#}$ Warning: package 'survival' was built under R version 3.6.2

Warning: package 'survminer' was built under R version 3.6.2

Warning: package 'ggplot2' was built under R version 3.6.2

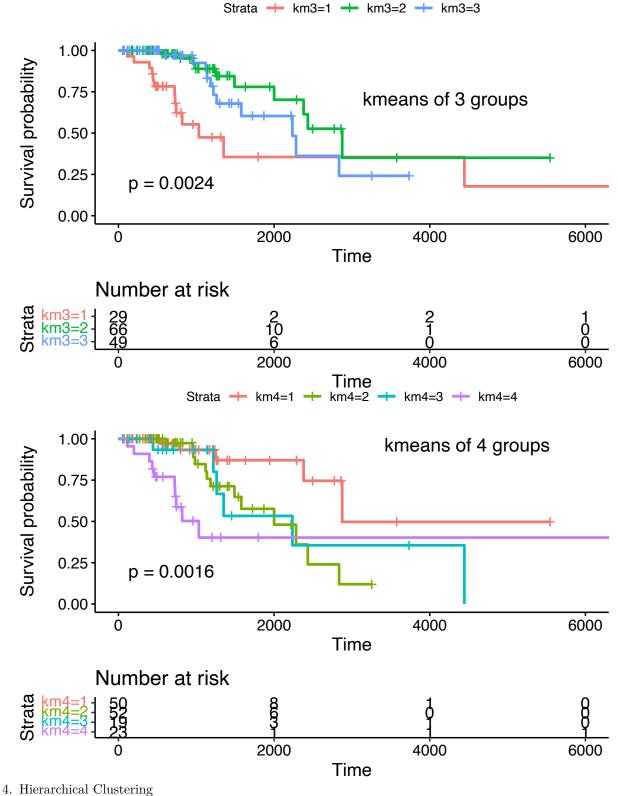
Warning: package 'ggpubr' was built under R version 3.6.2



Time

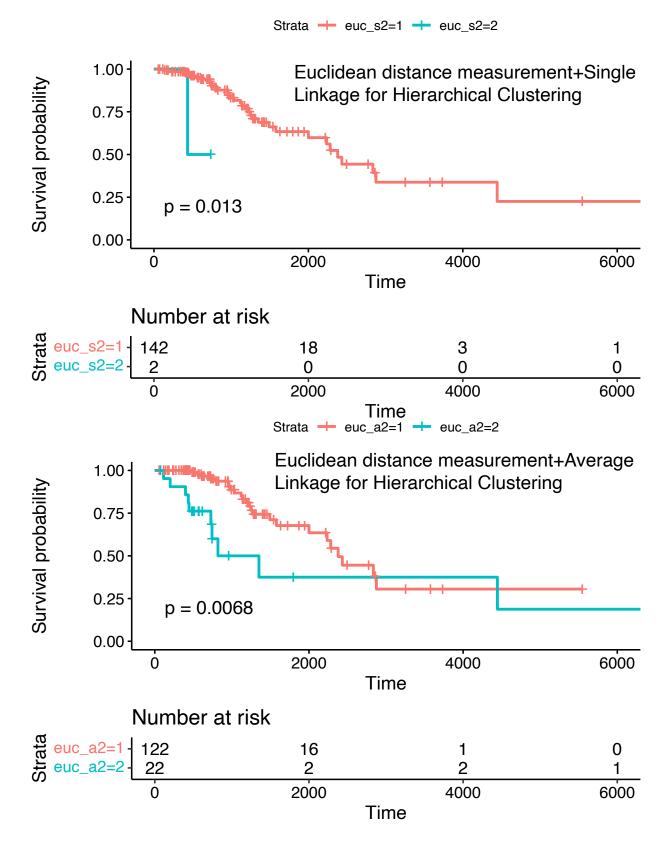
3. Fit the results of Kmeans

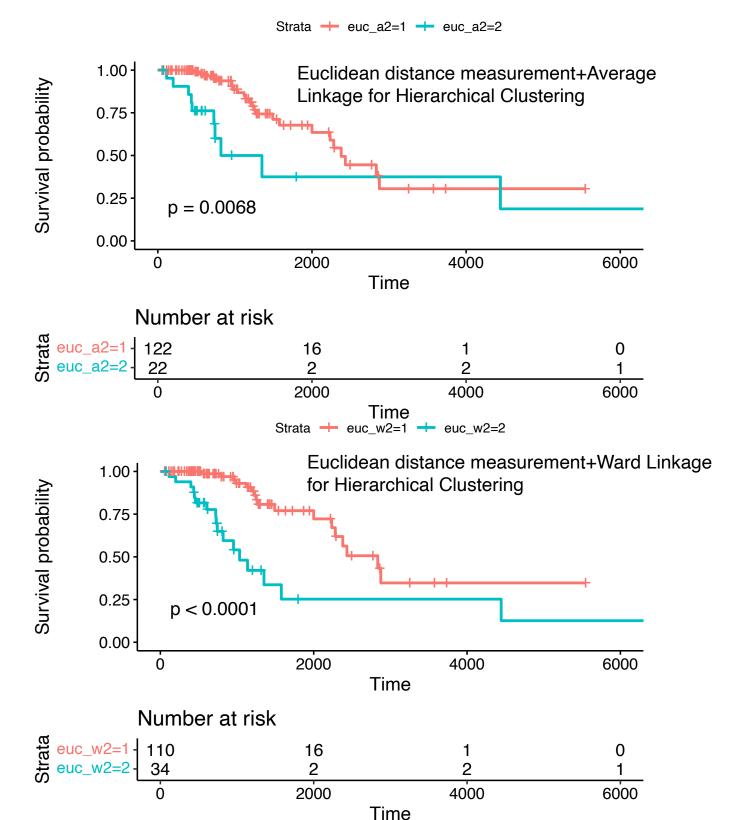
Kmeans of 2 groups

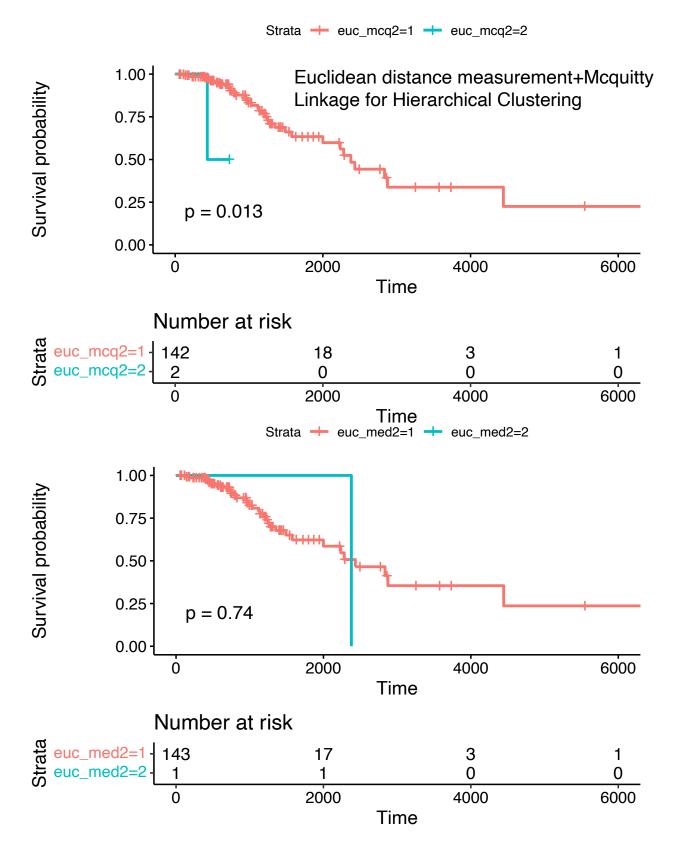


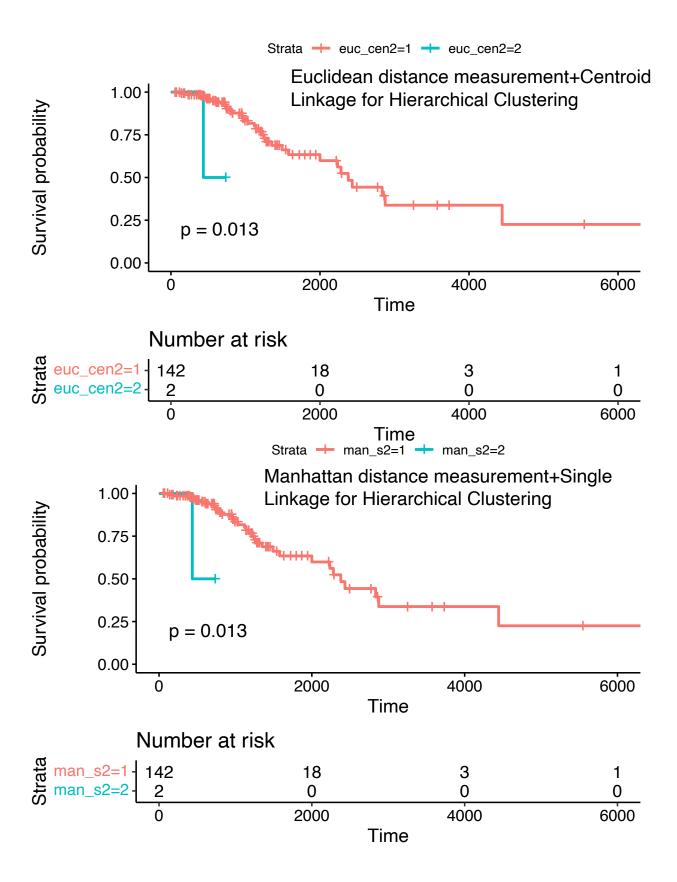
4. Theraremear Clustering

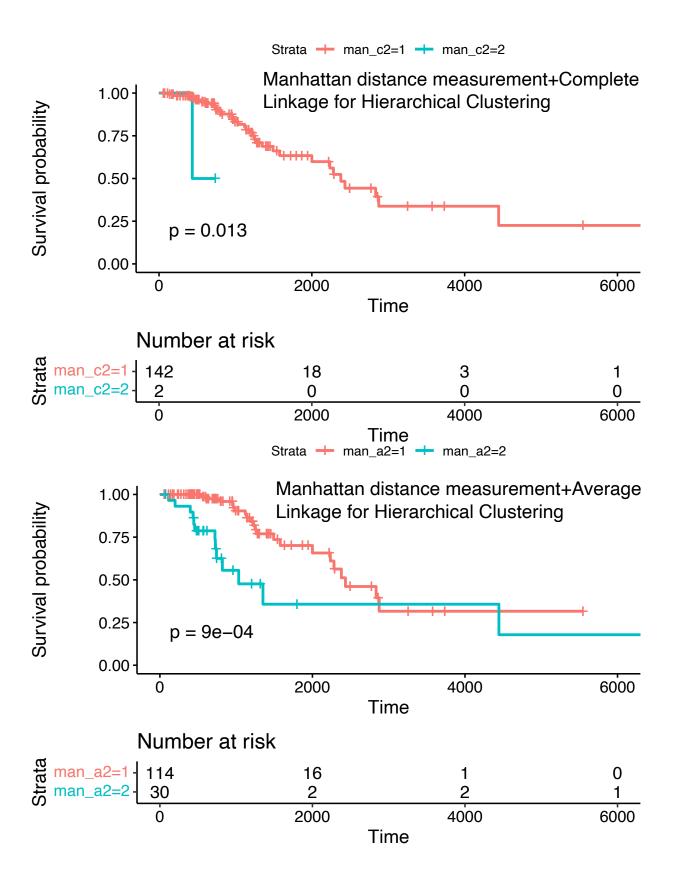
Warning: package 'proxy' was built under R version 3.6.2

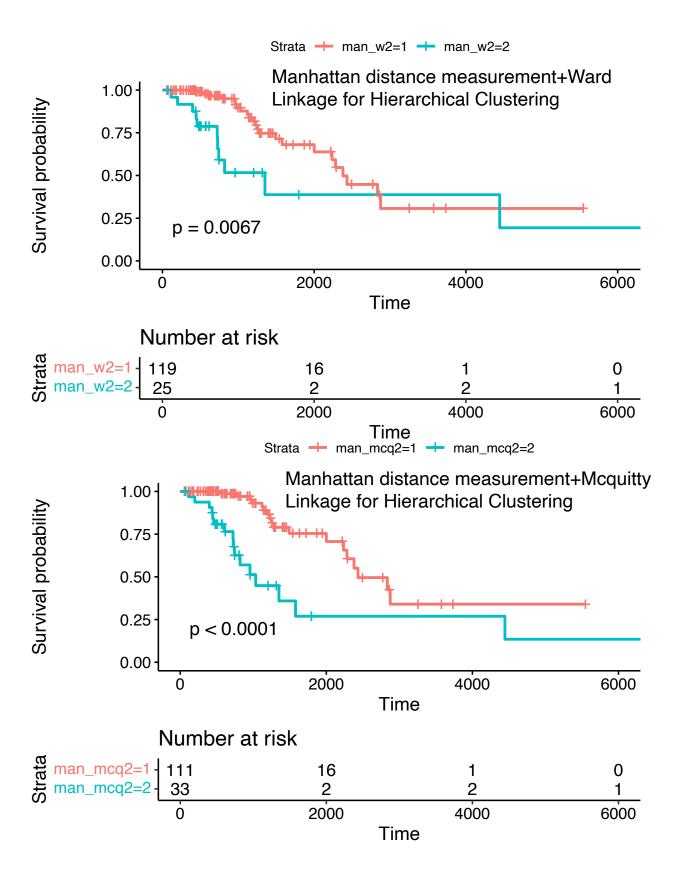


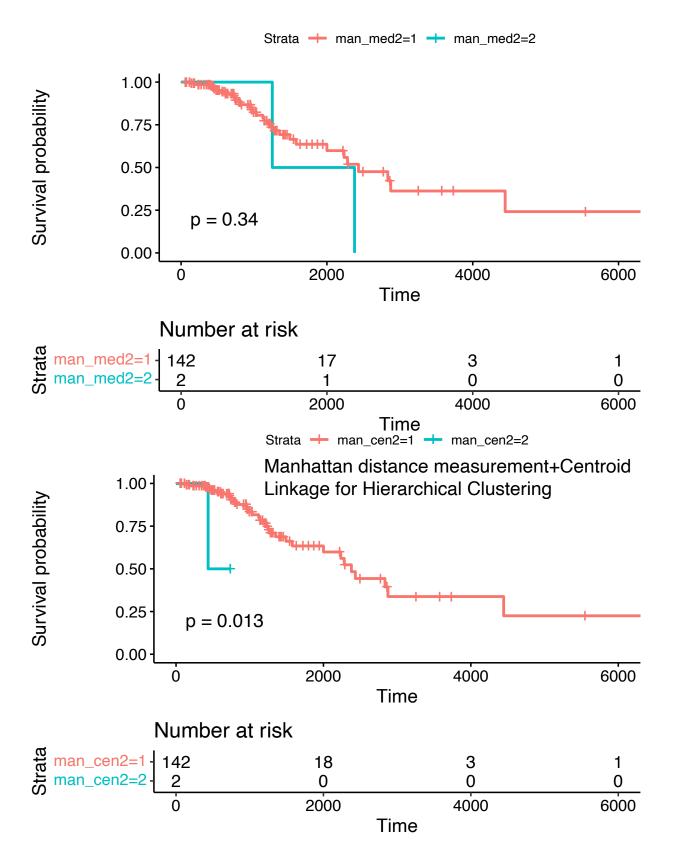


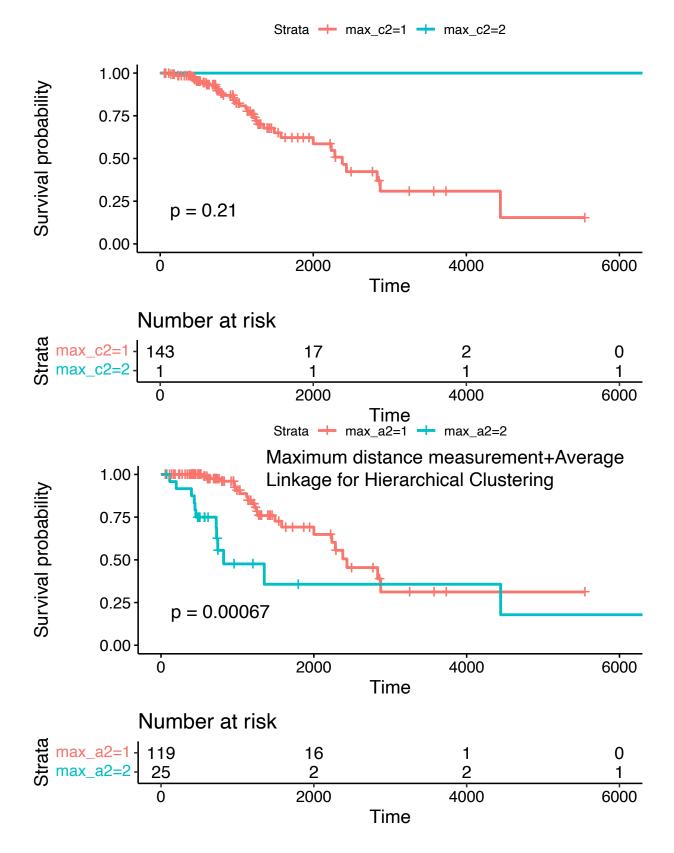


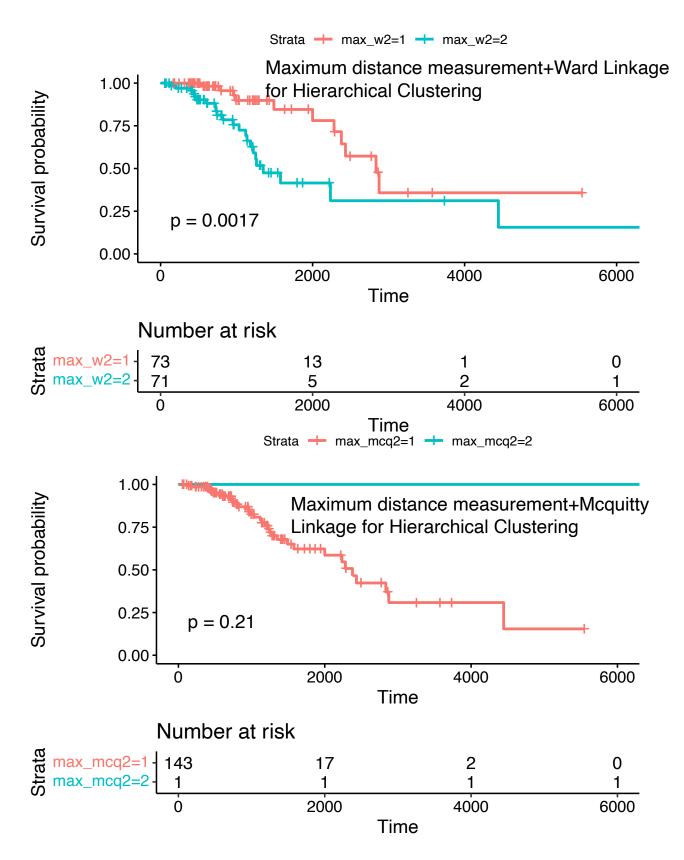




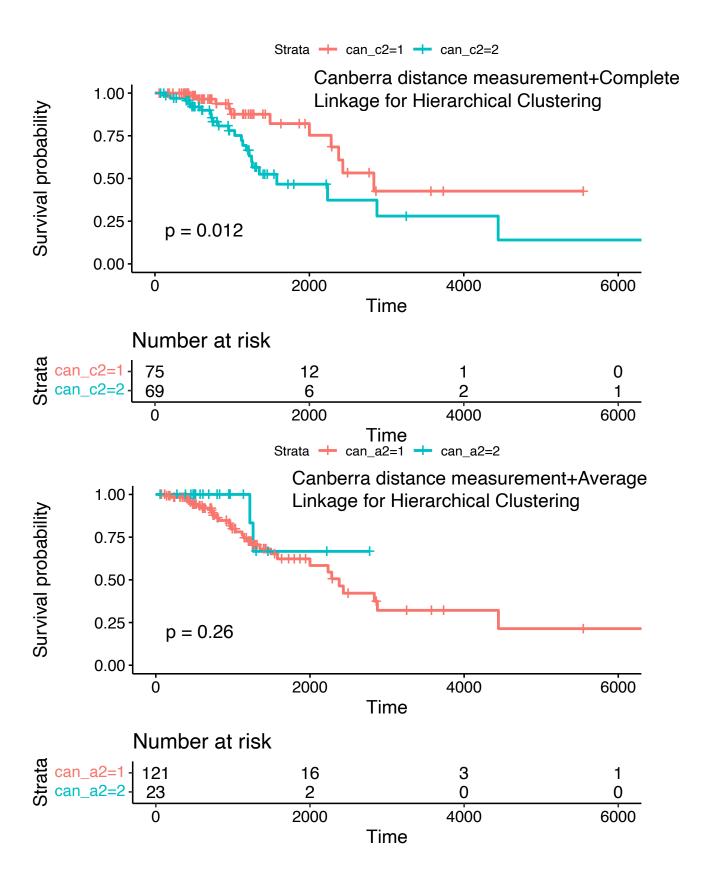


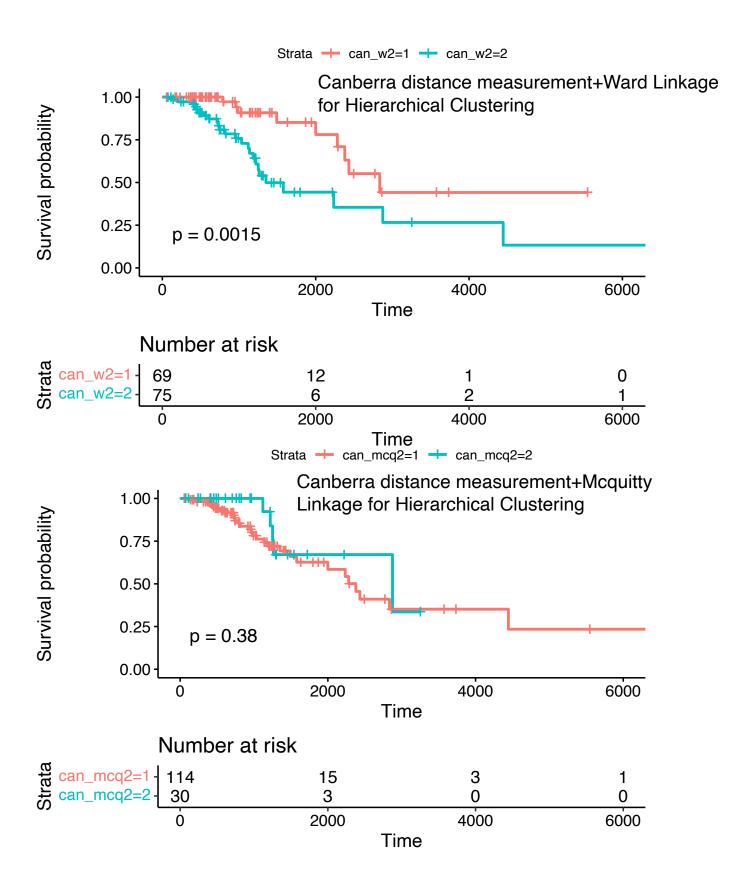


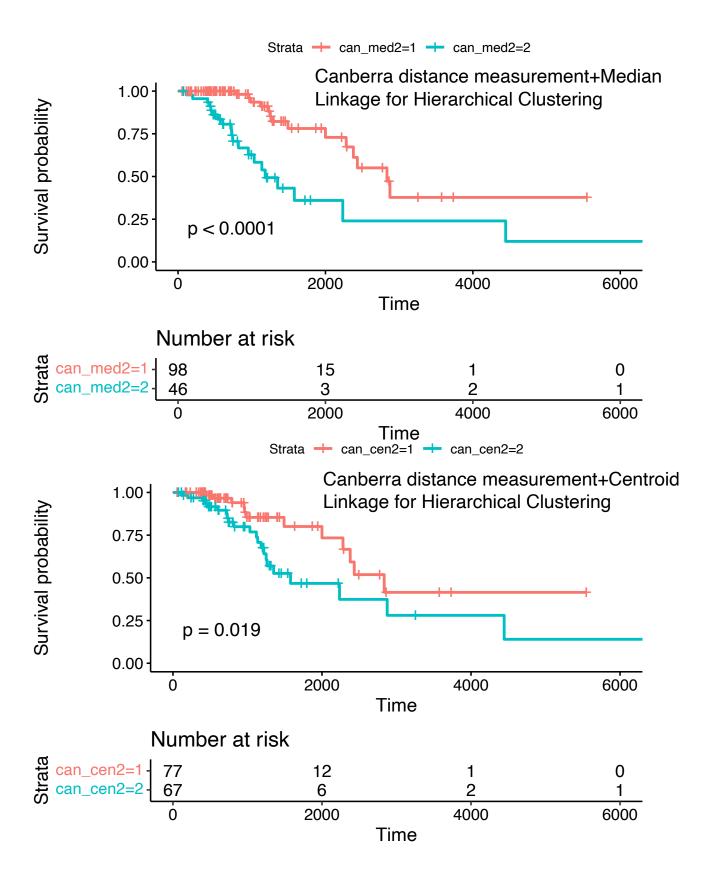


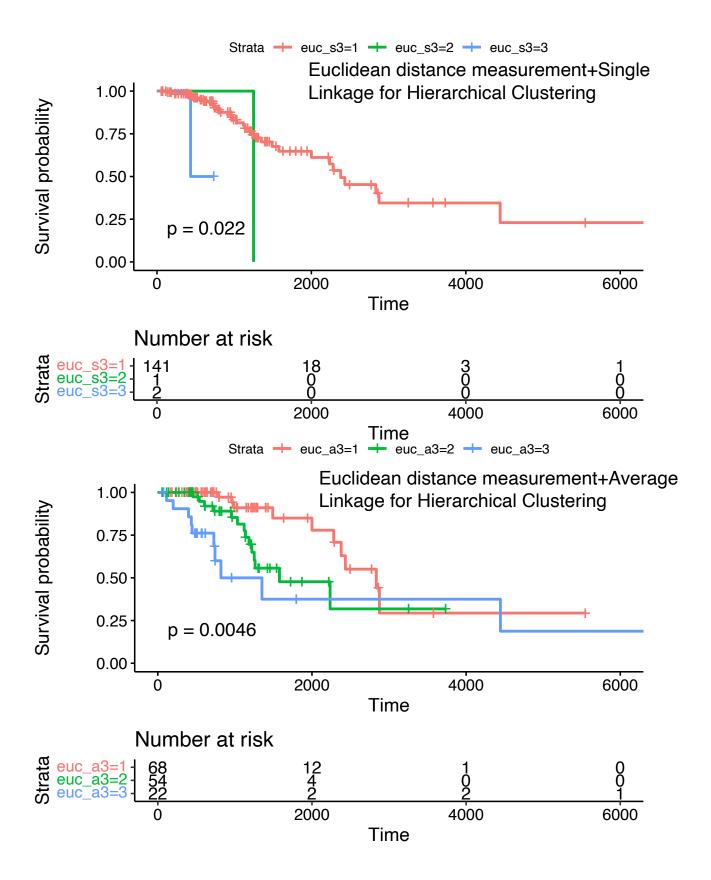


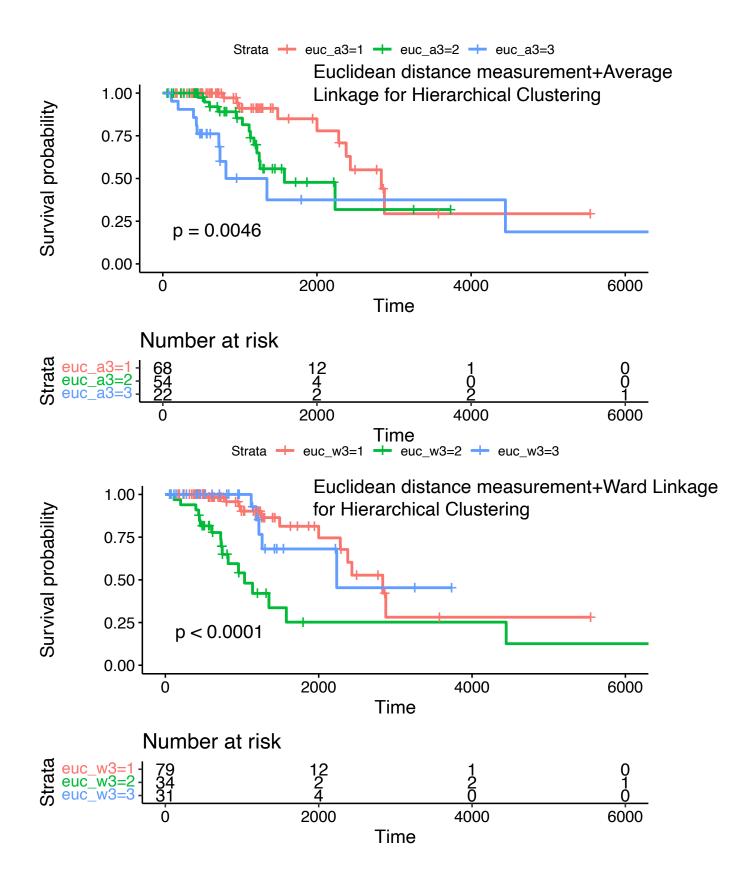
Warning in .pvalue(fit, data = data, method = method, pval = pval, pval.coord = pval.coord, : There
This is a null model.

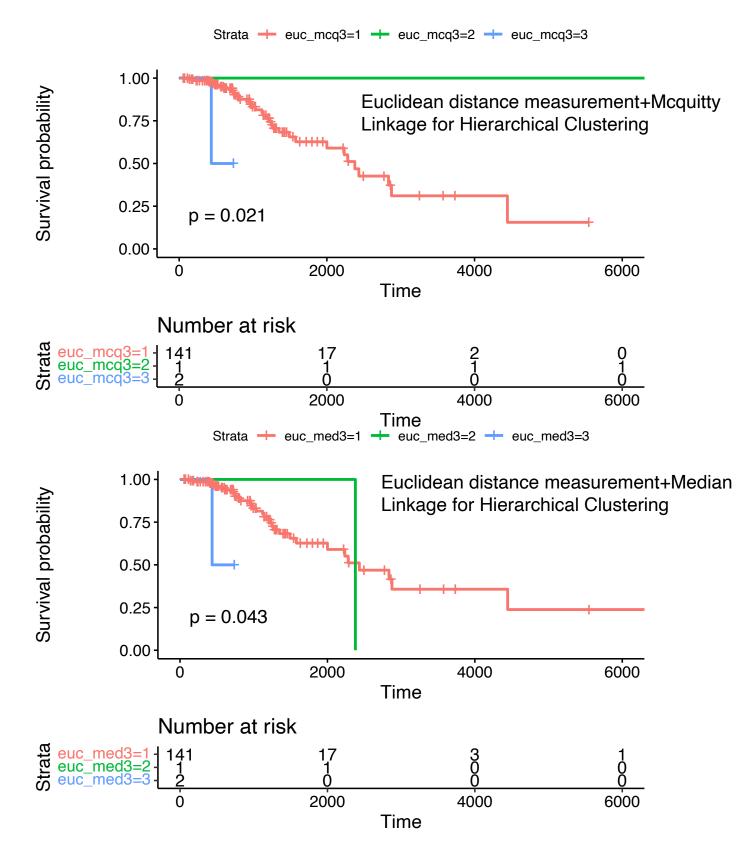


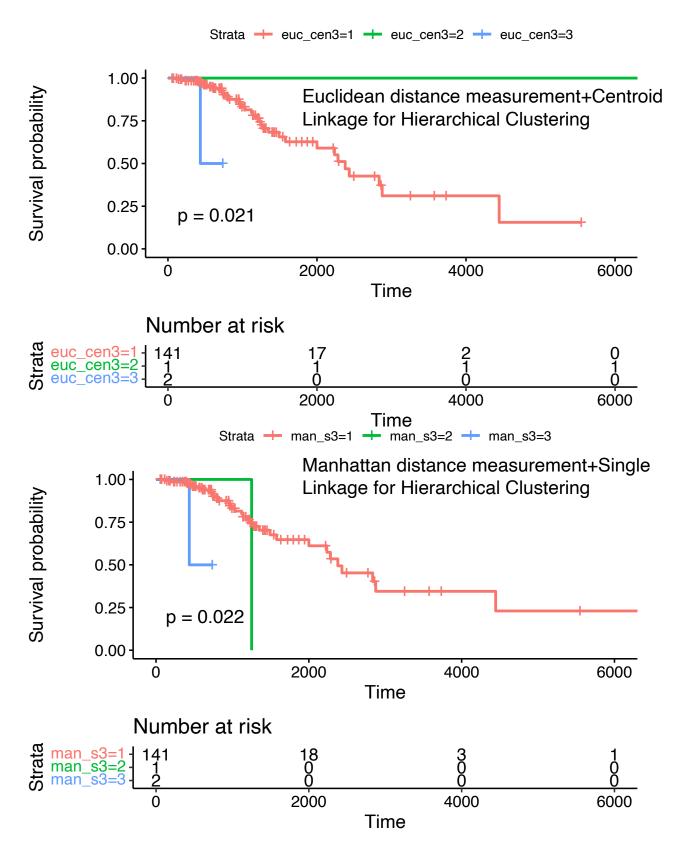


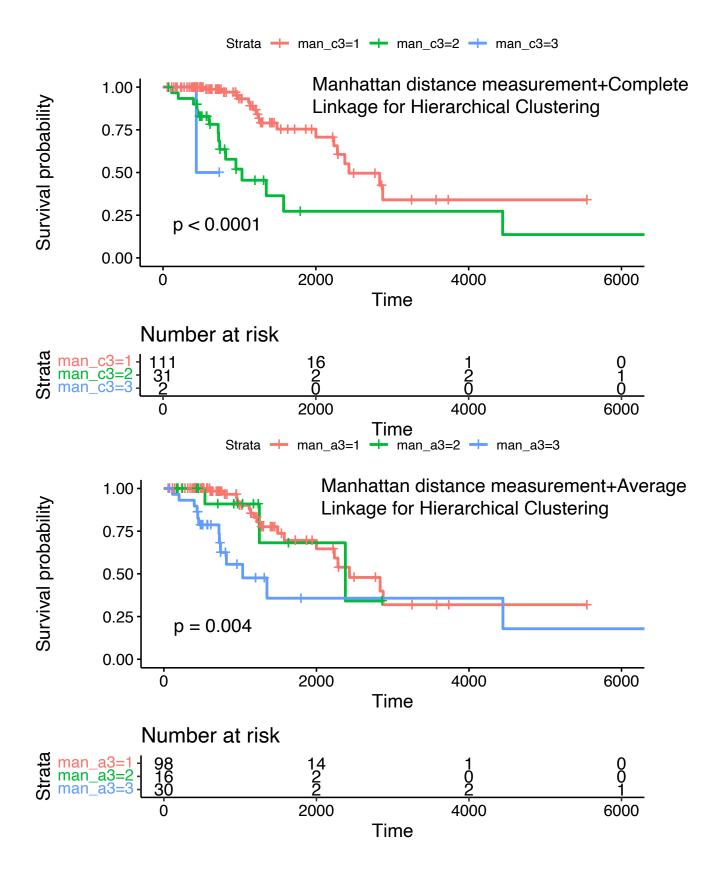


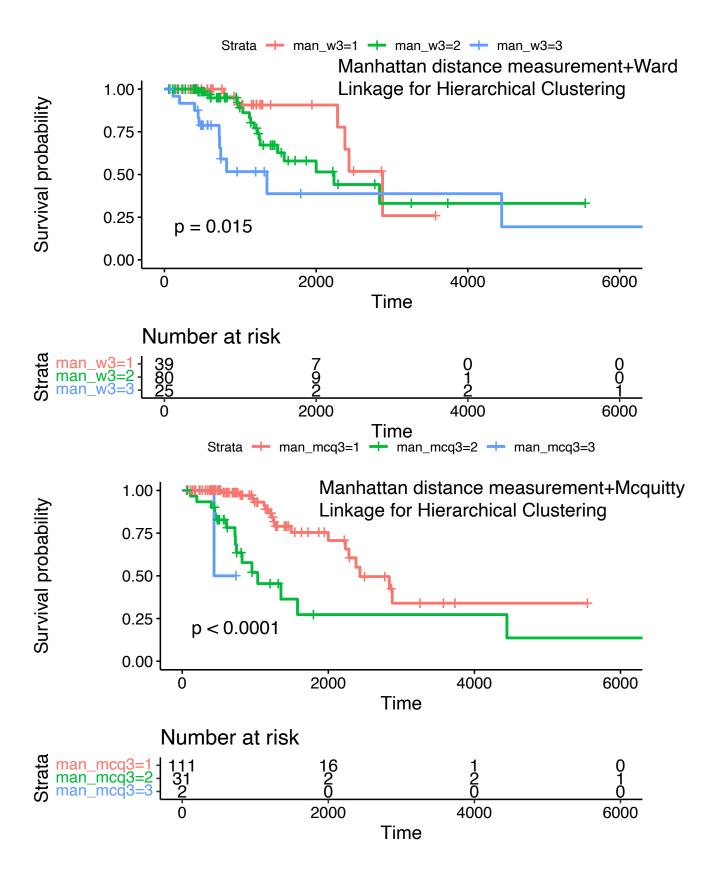


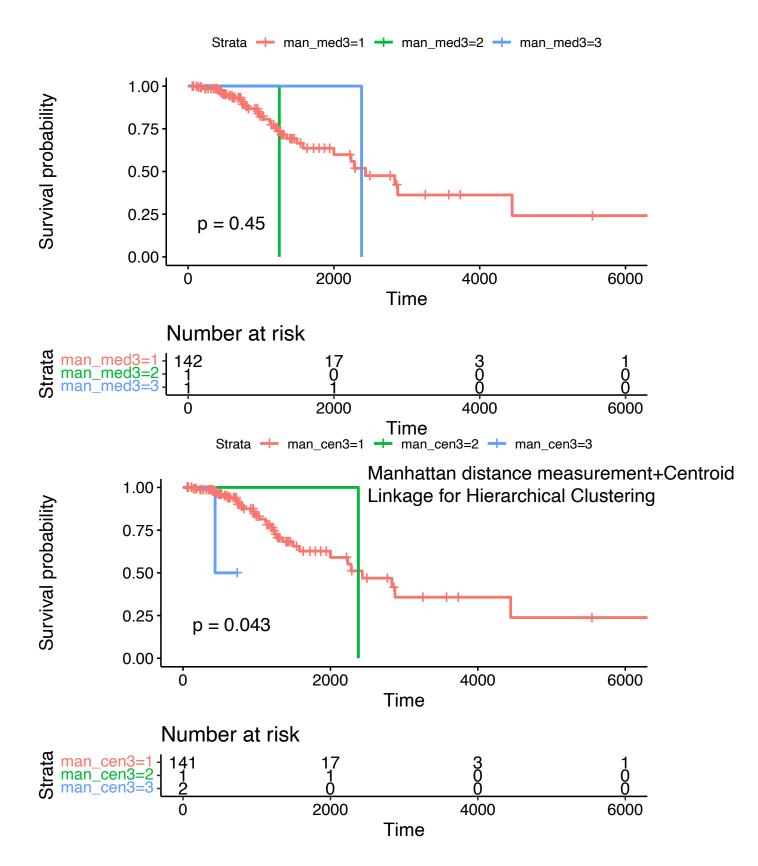


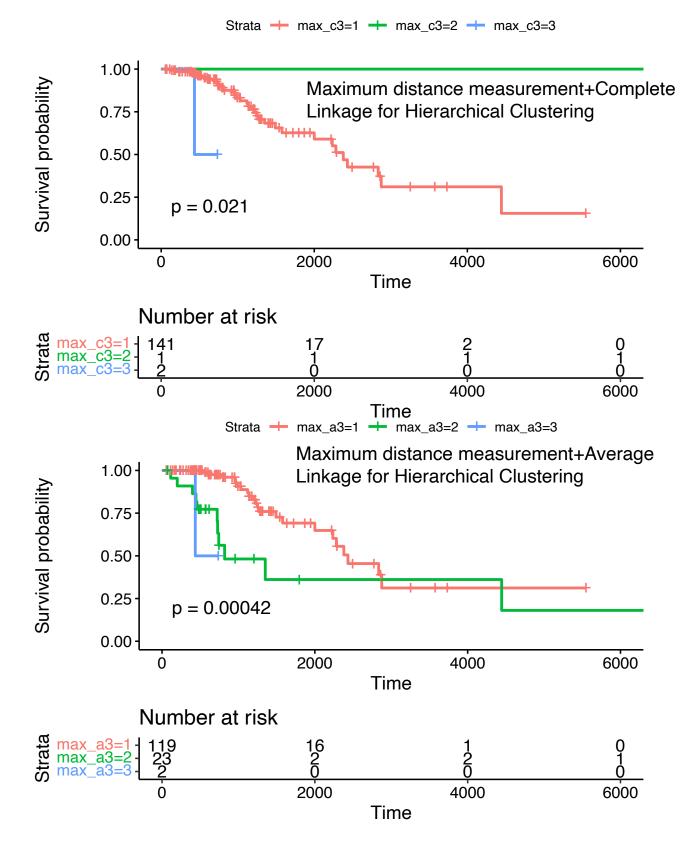


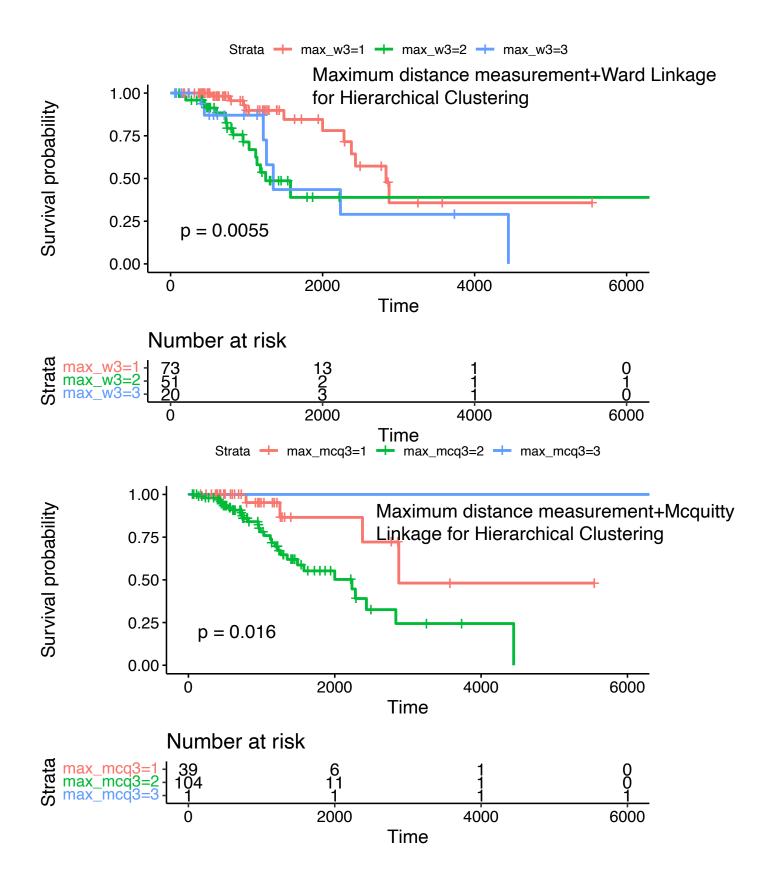


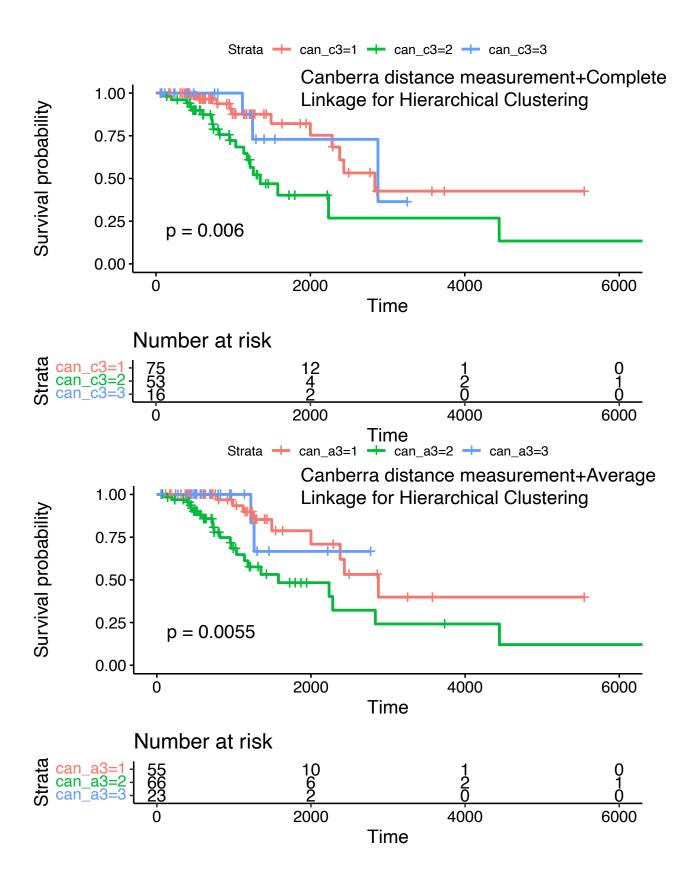


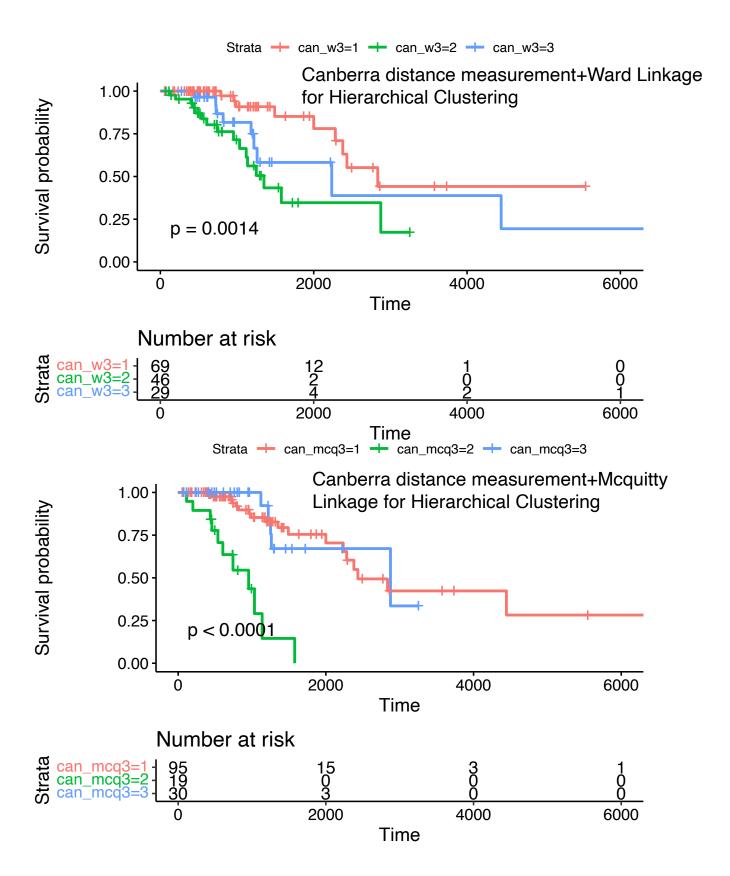


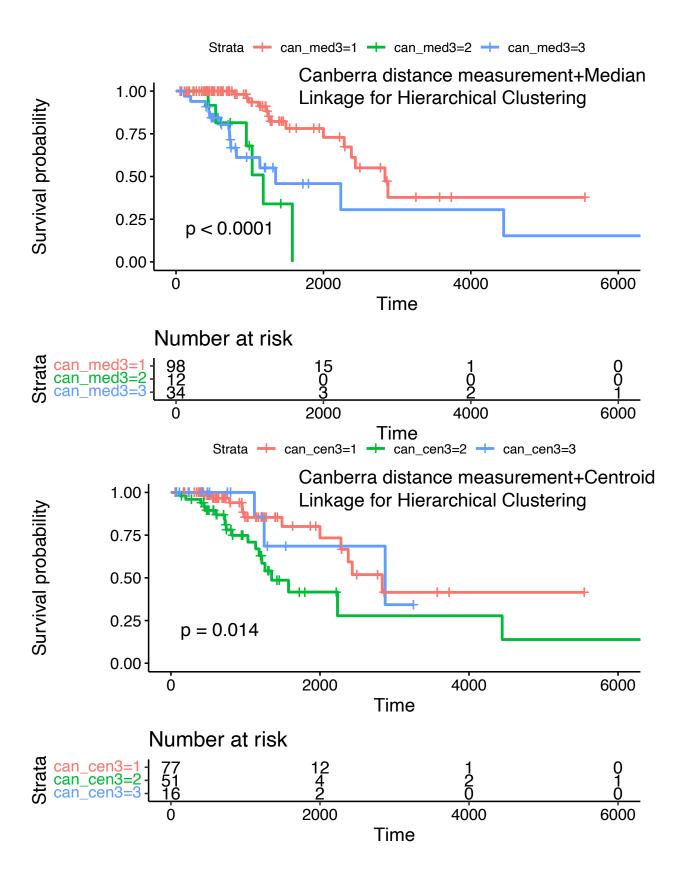


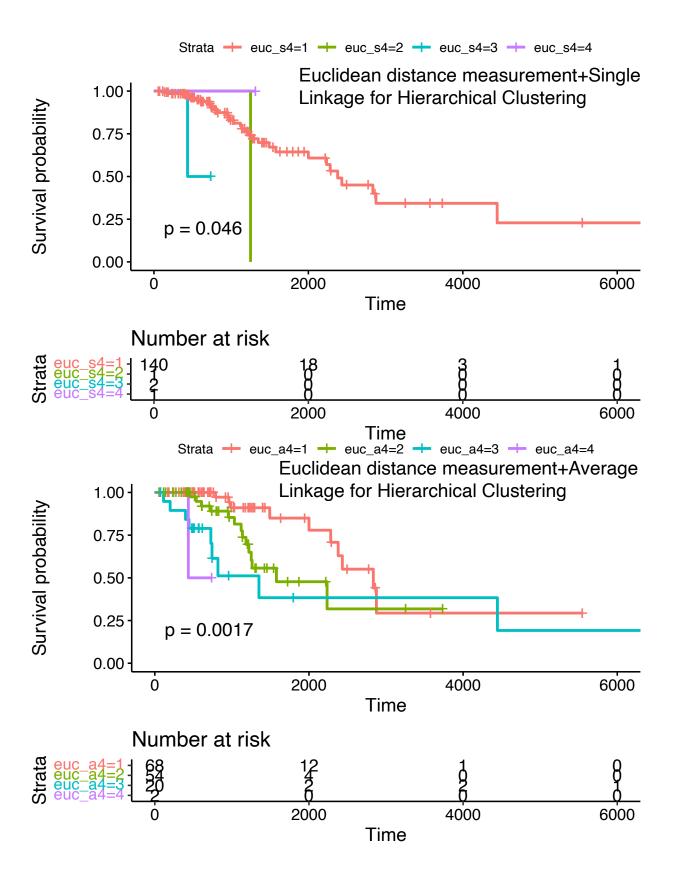


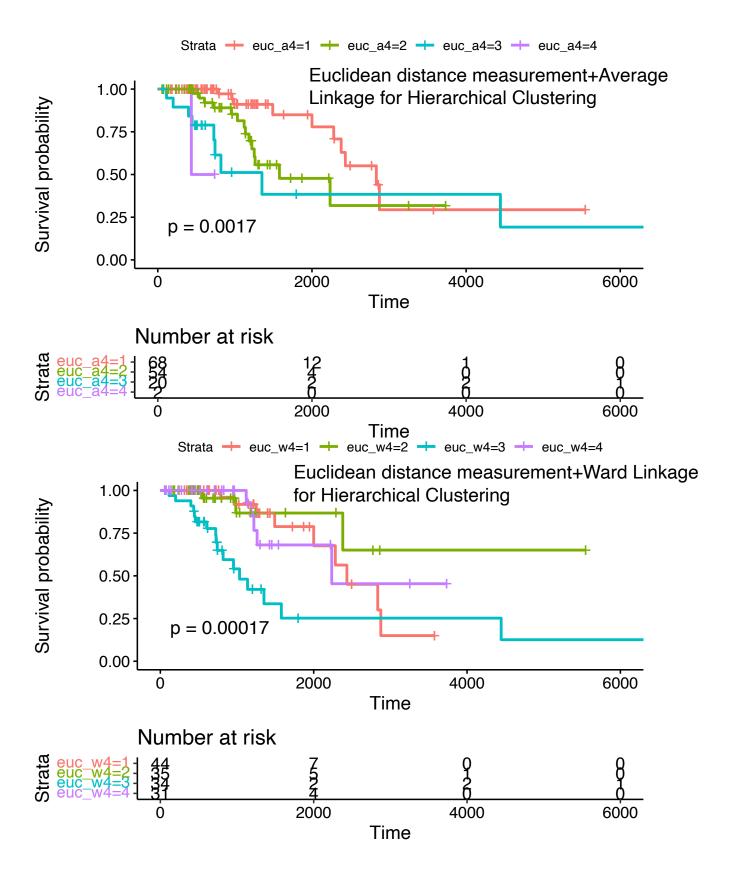


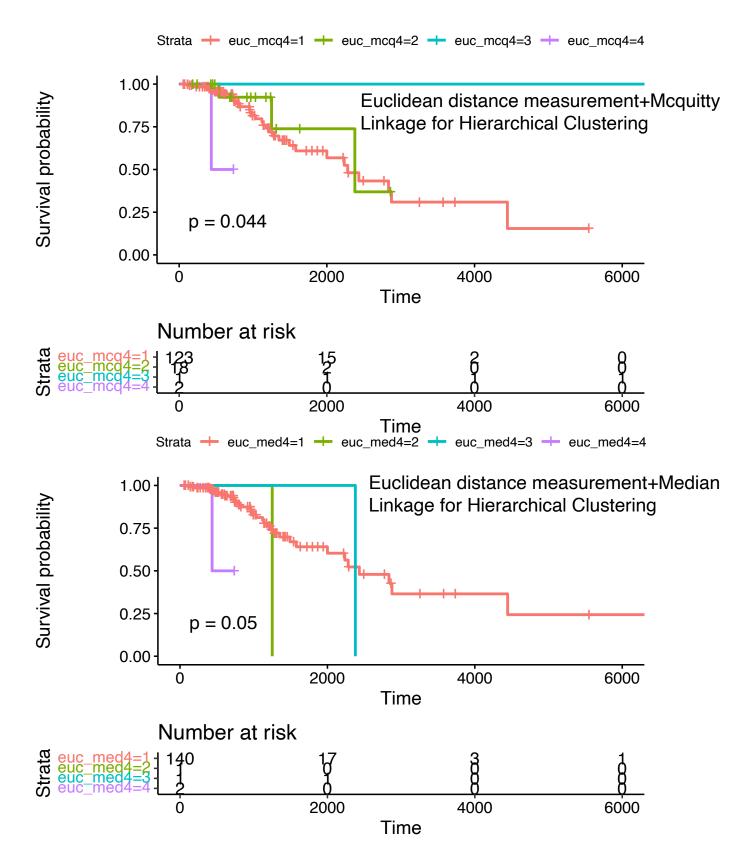


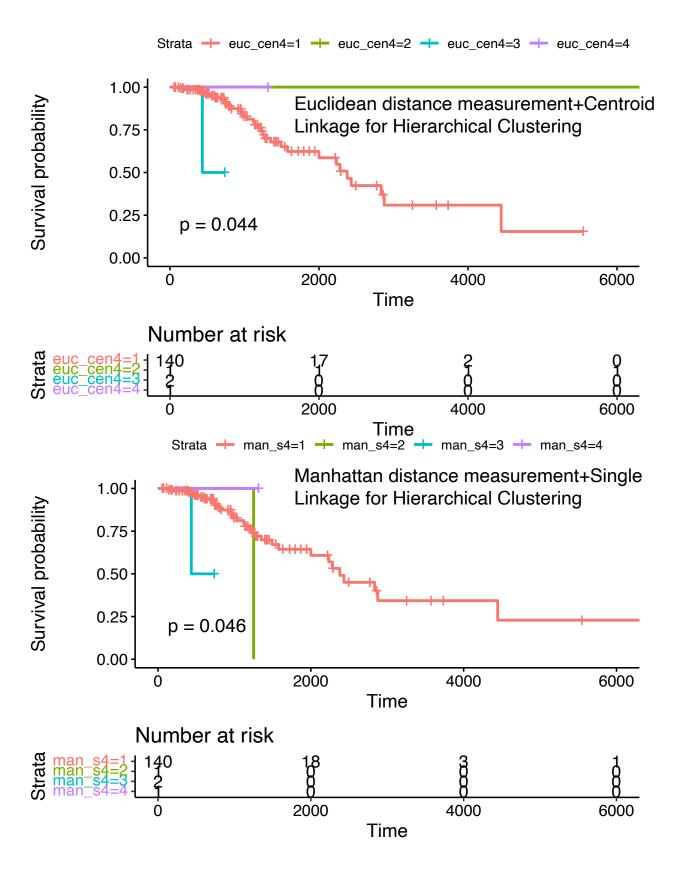


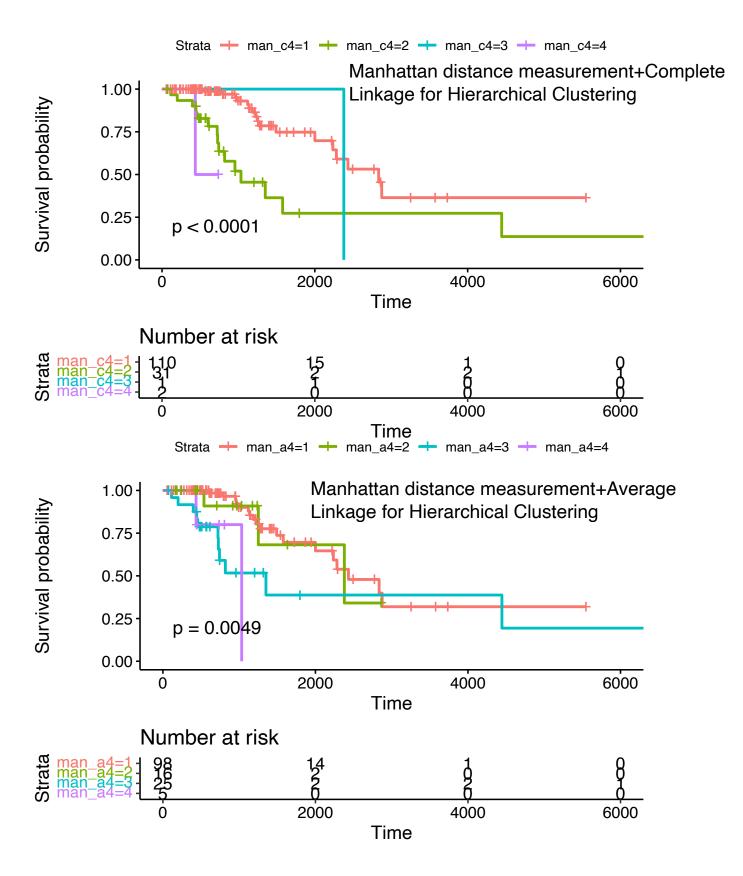


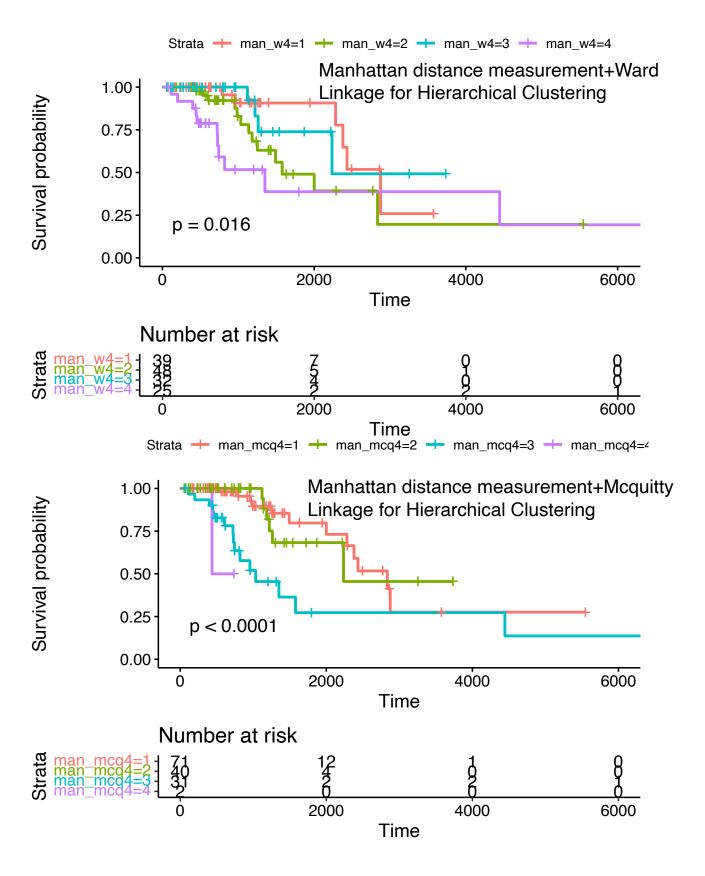


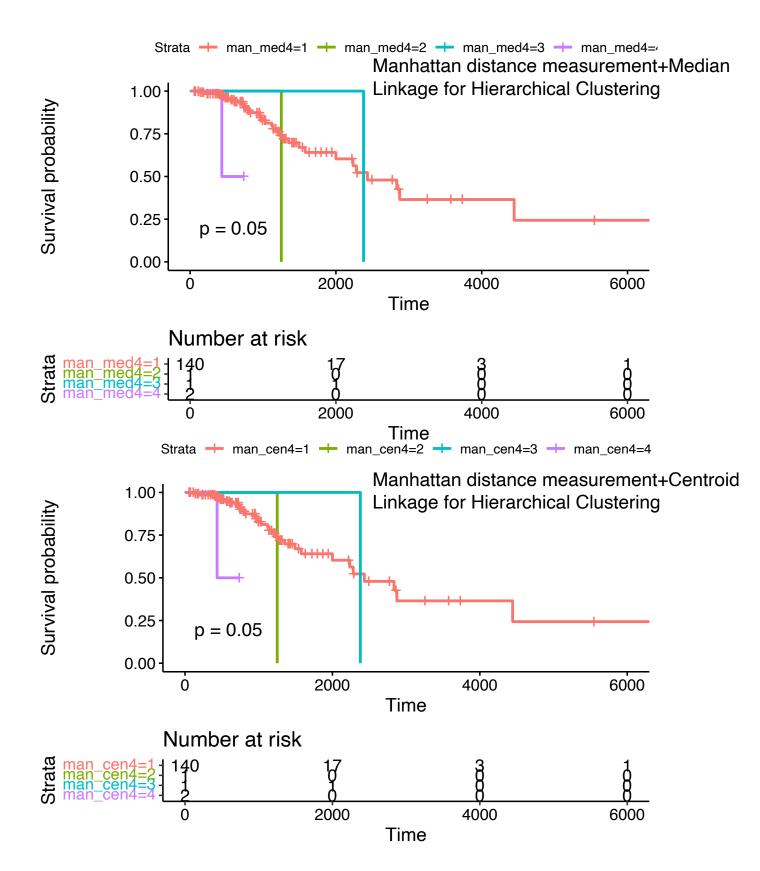


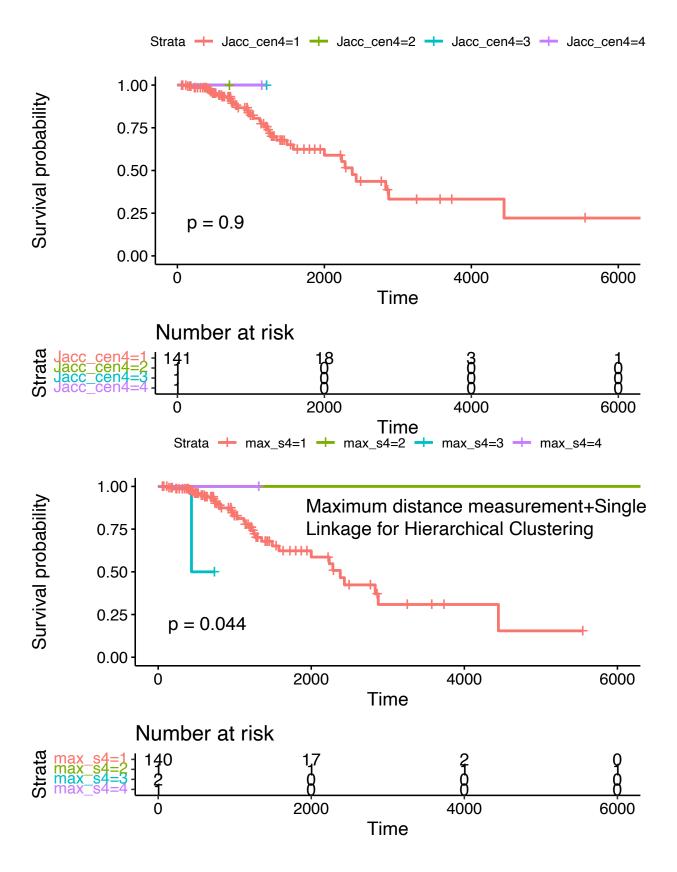


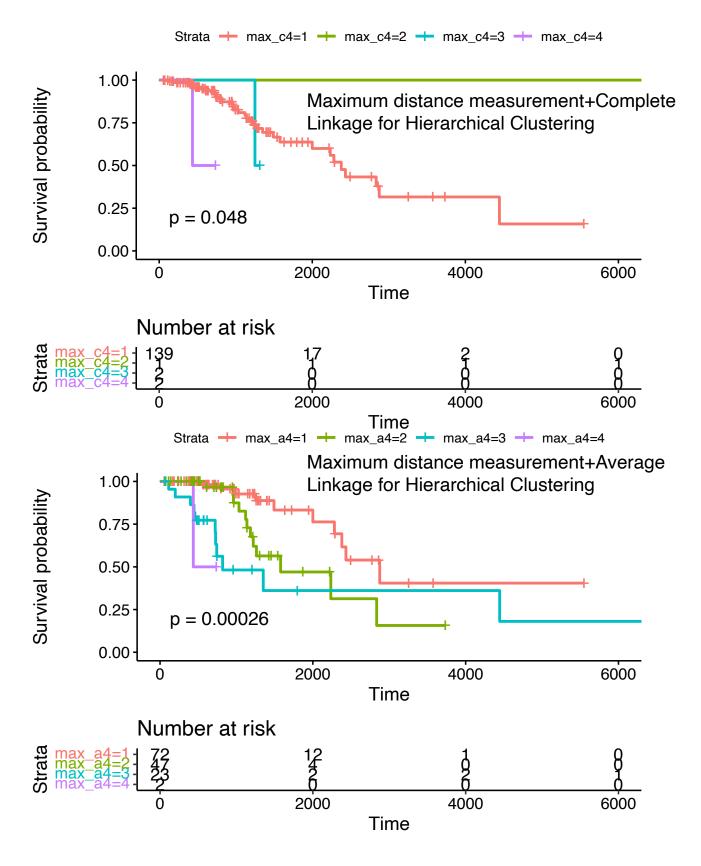


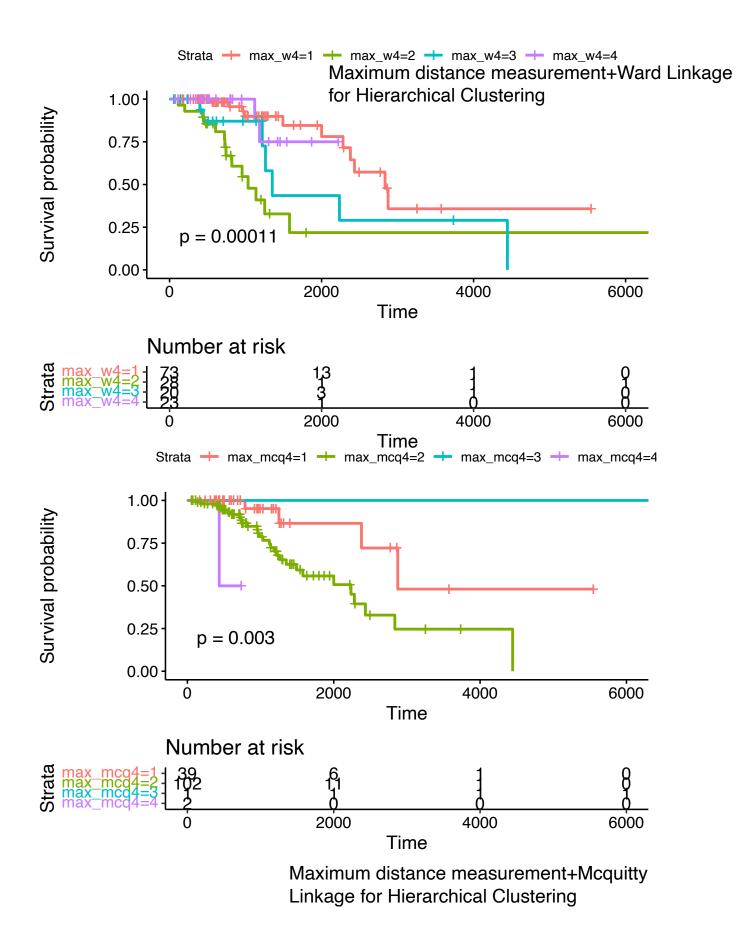


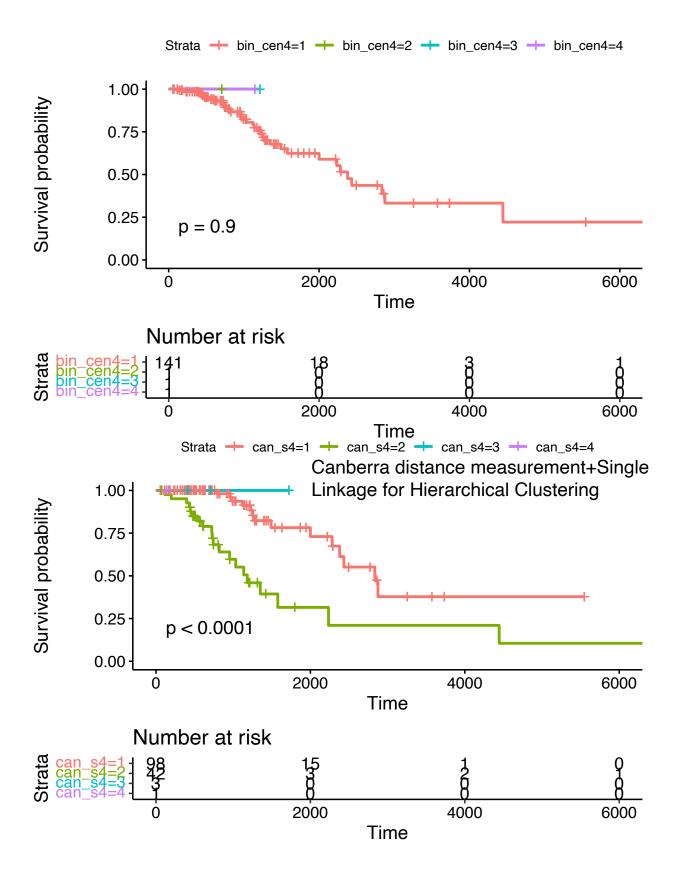


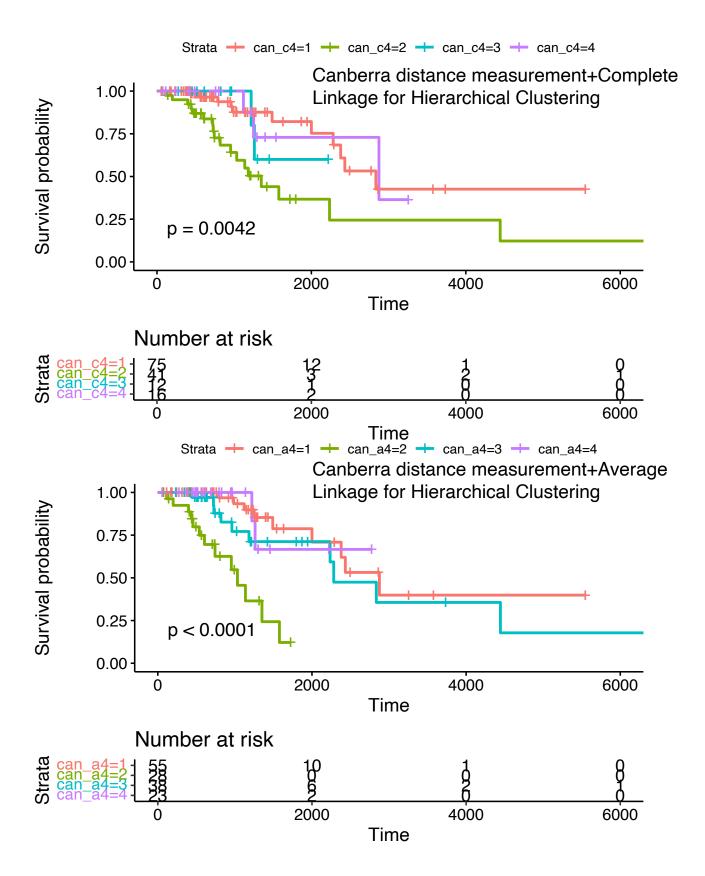


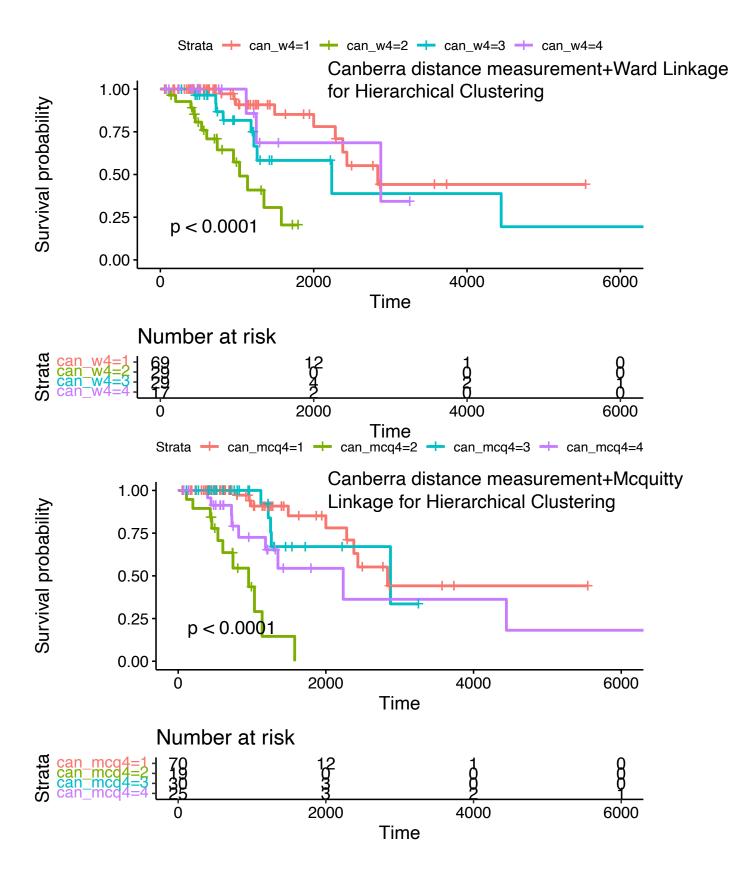


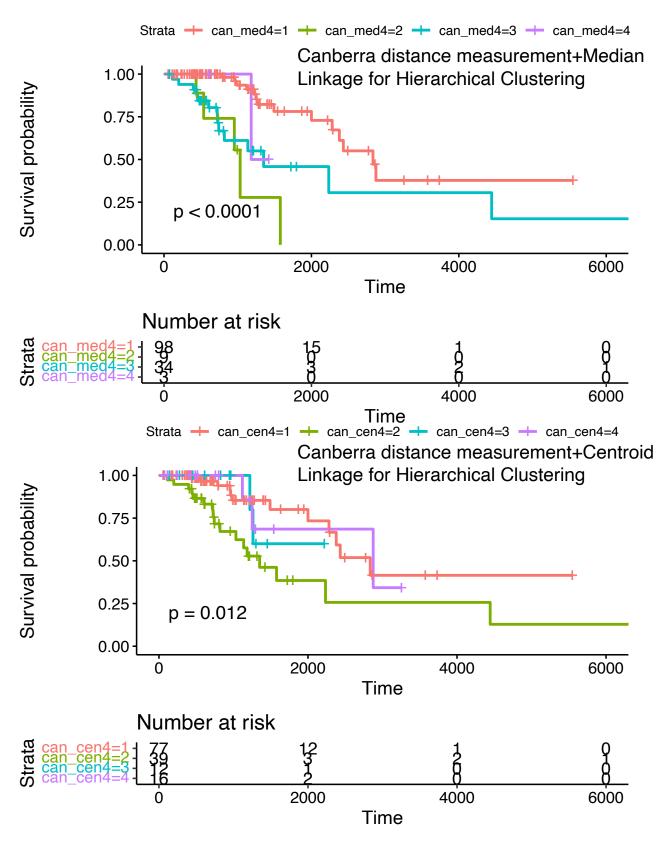












3. DBSCAN

Warning: package 'dbscan' was built under R version 3.6.2

