cp decomposition glioma 73 train rank5 test

VIVI TANG

17/11/2021

1. Kmeans

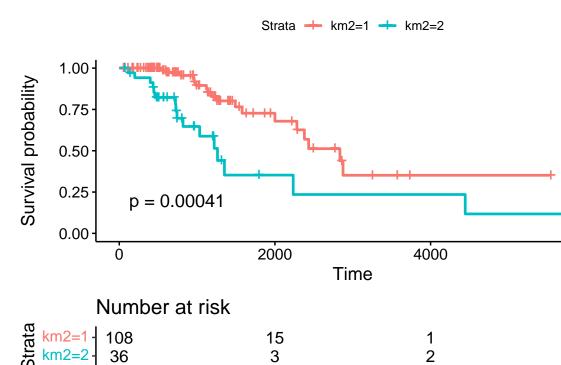
Warning: package 'lattice' was built under R version 3.6.2

 $\mbox{\tt \#\#}$ 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

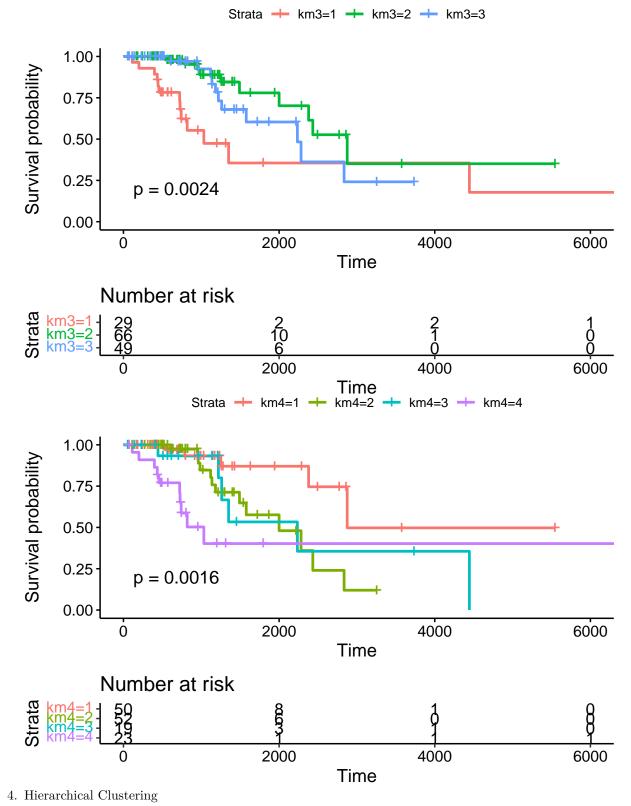


2000

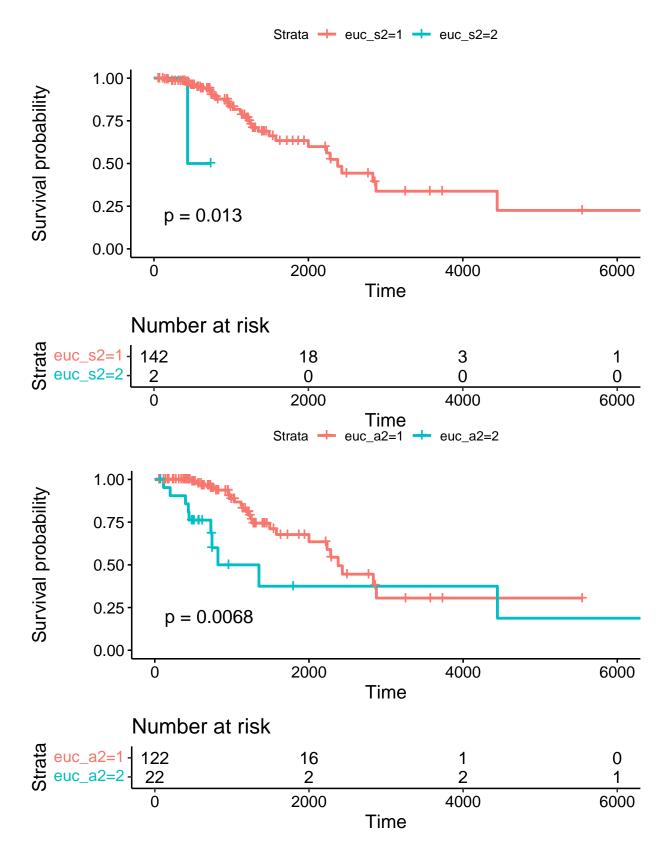
4000

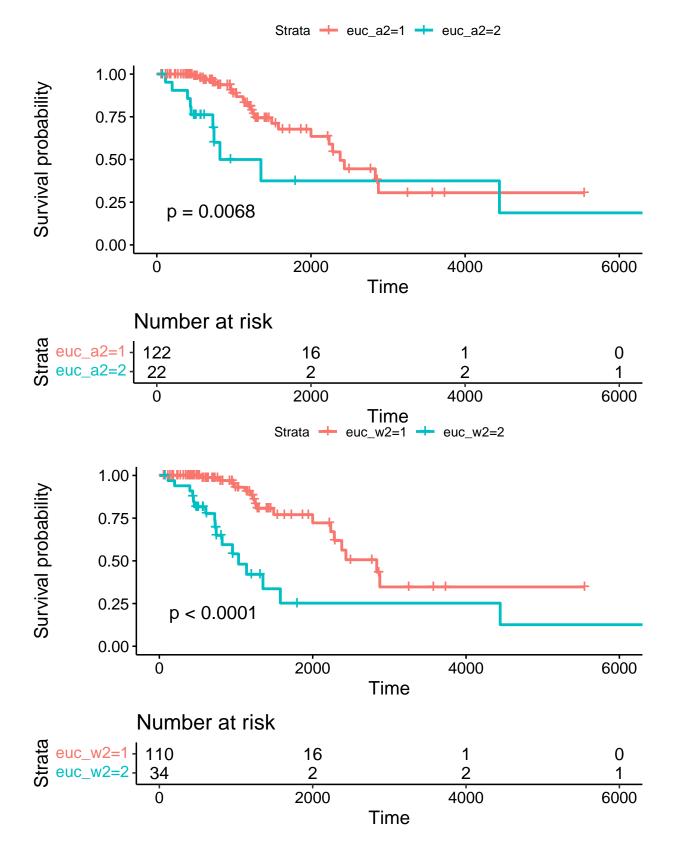
Time

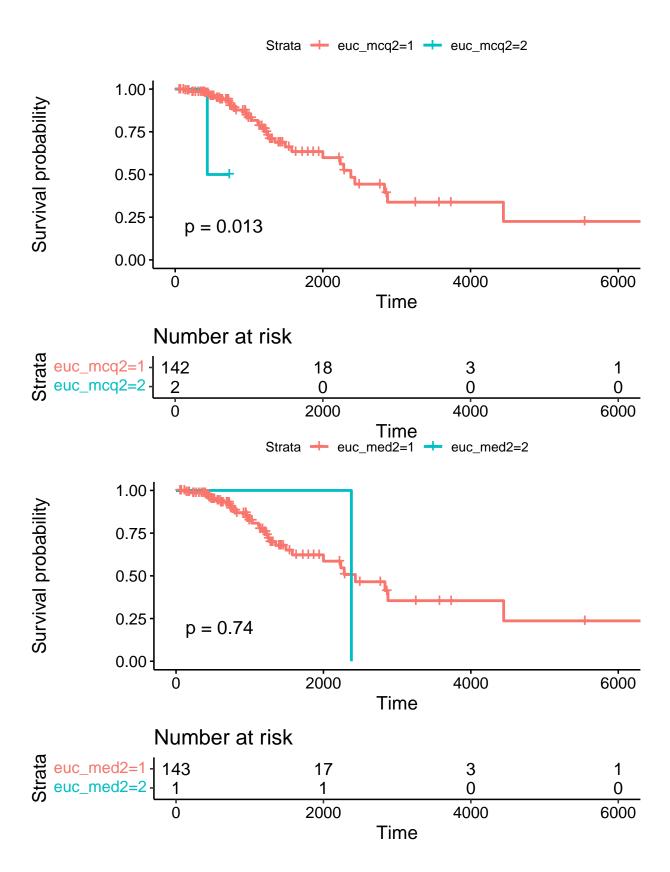
3. Fit the results of Kmeans

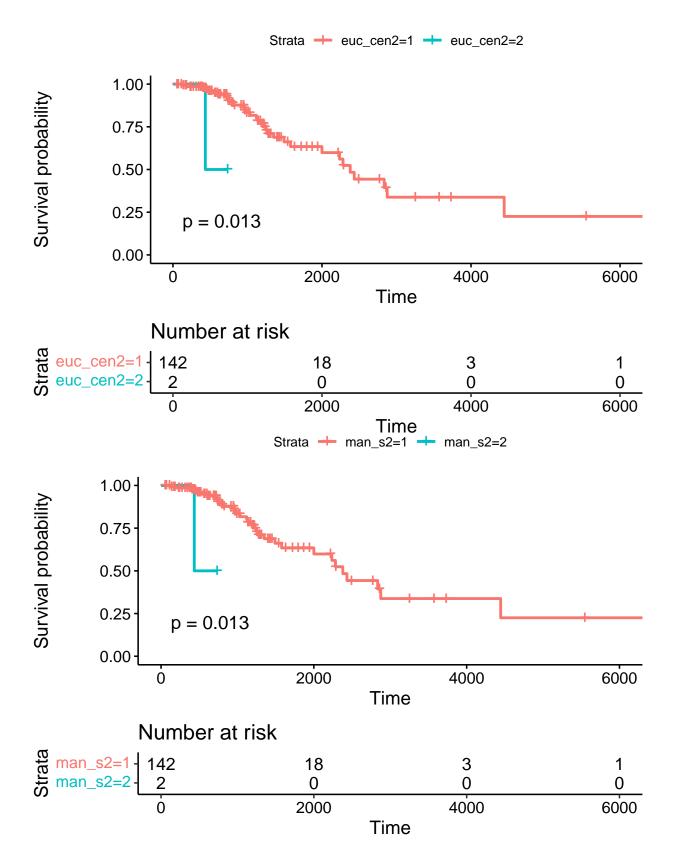


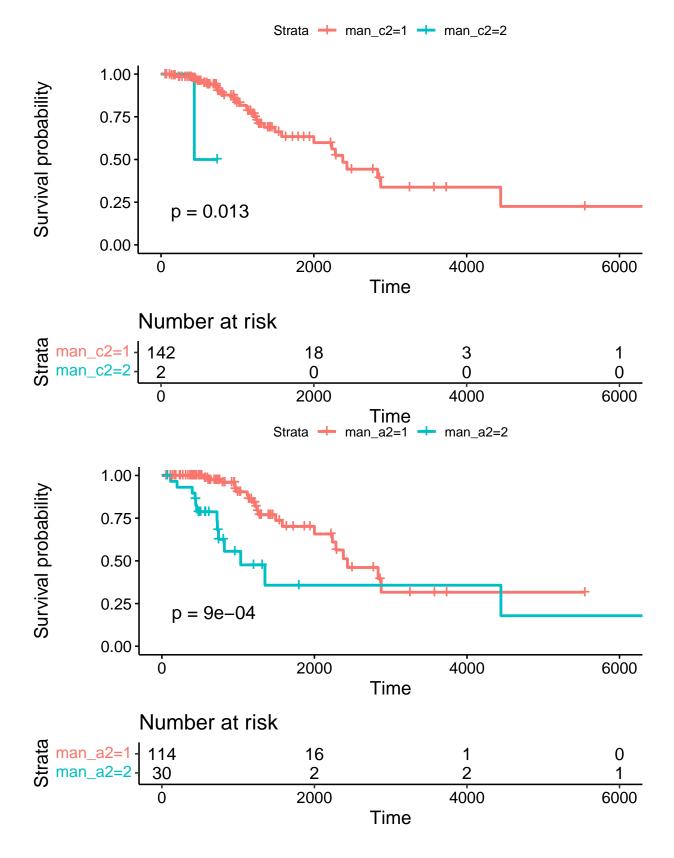
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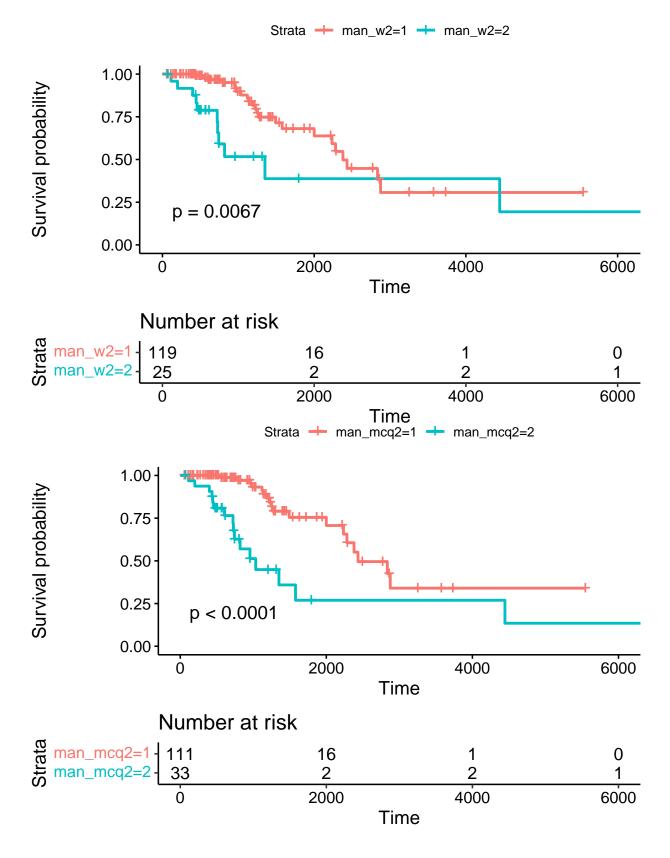


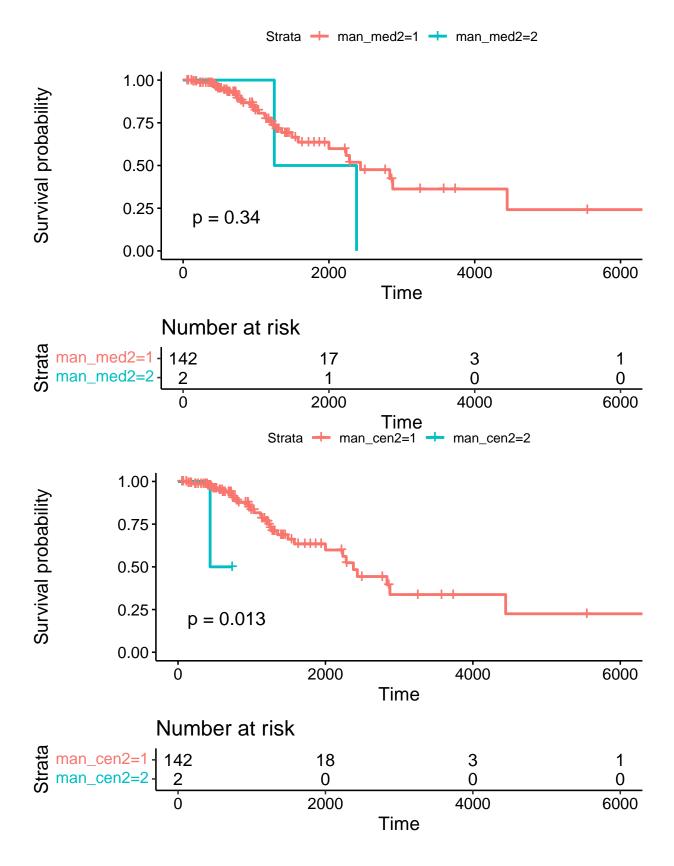


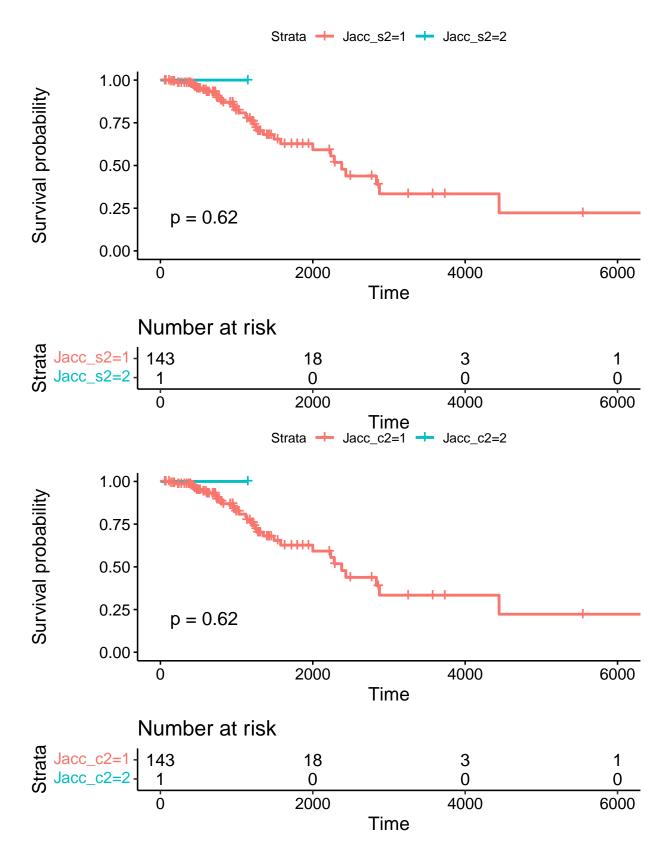


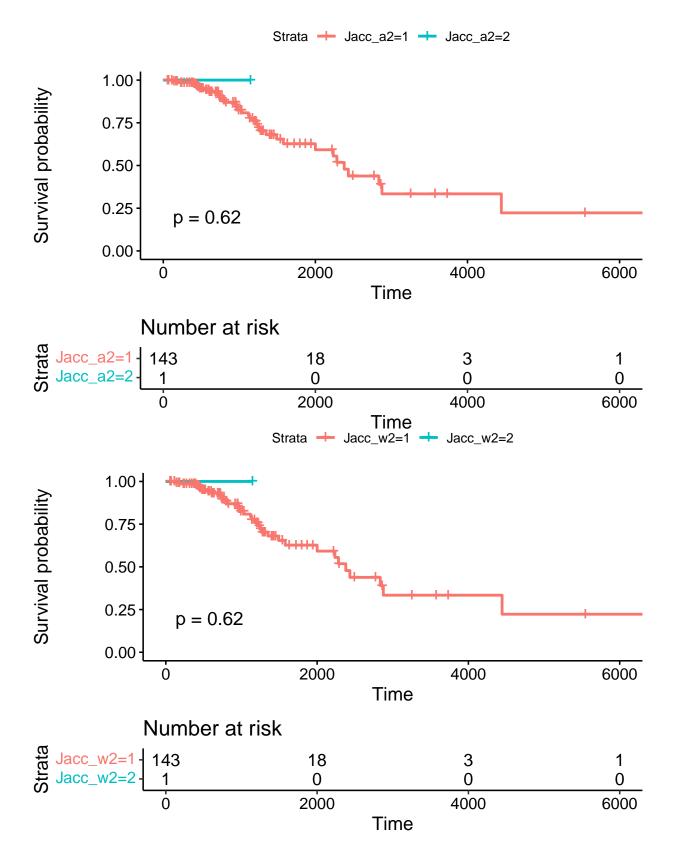


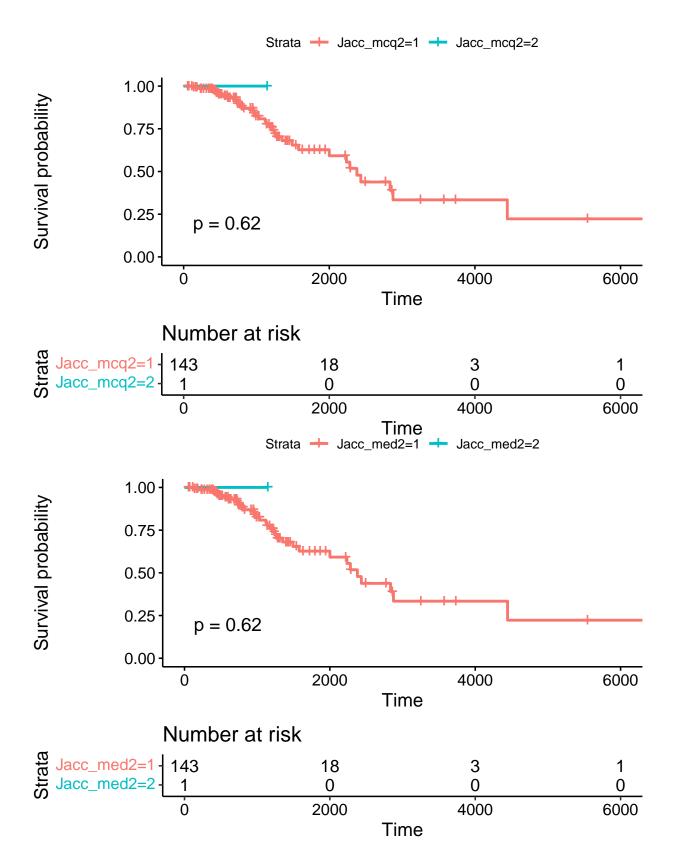


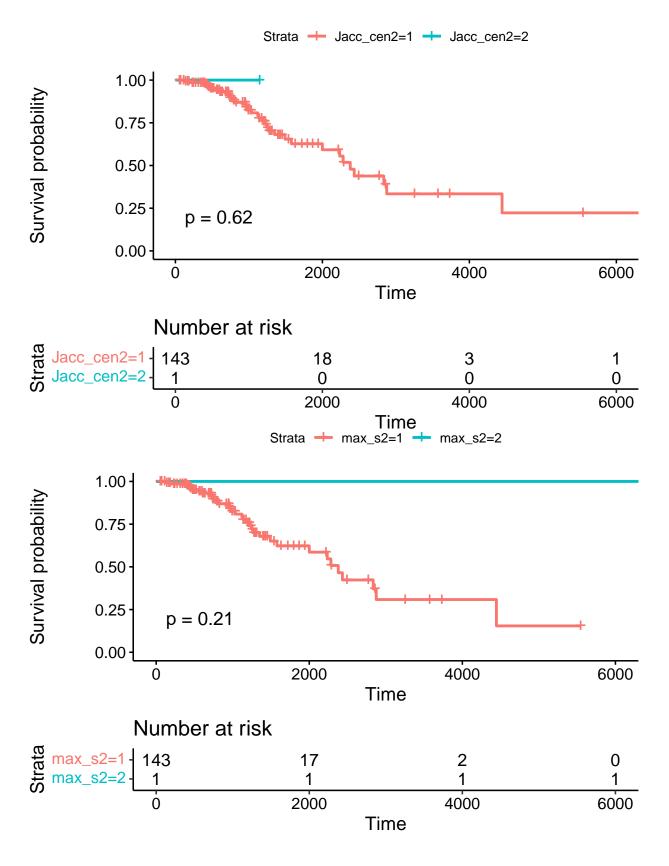


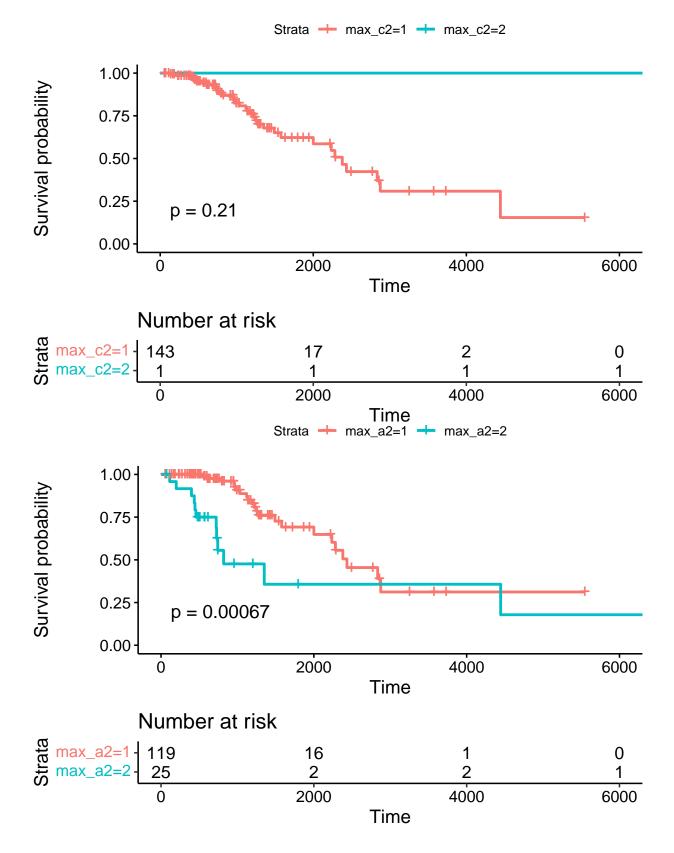


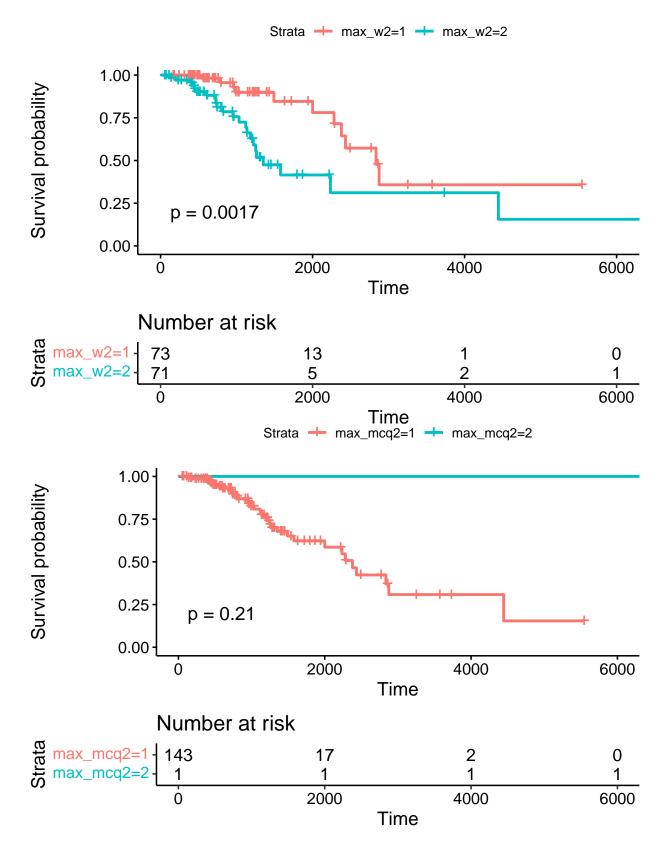




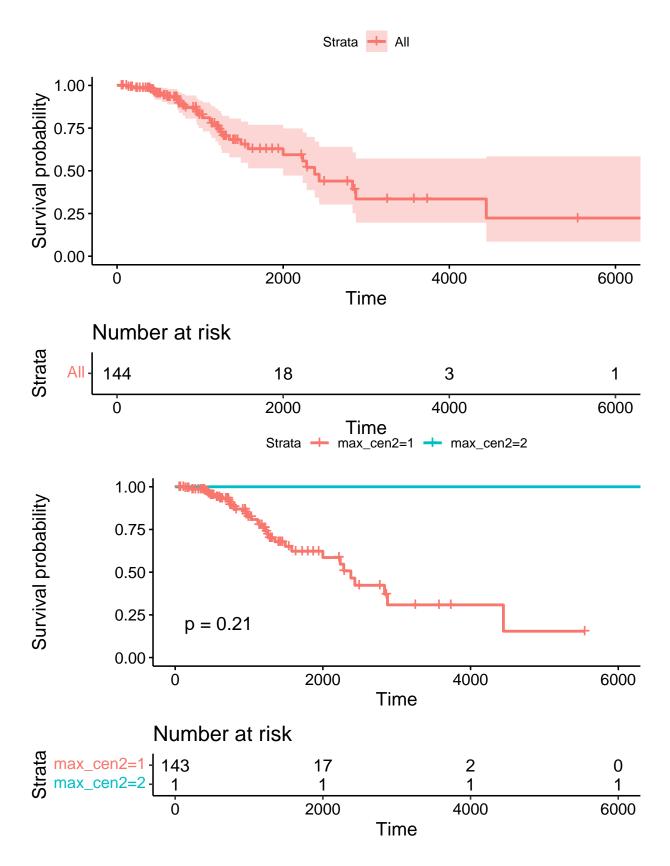


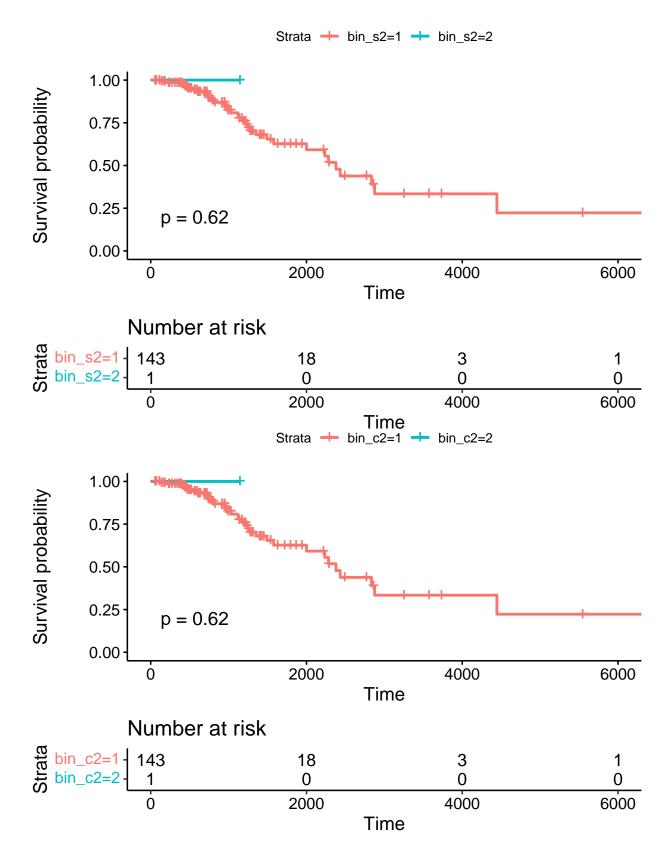


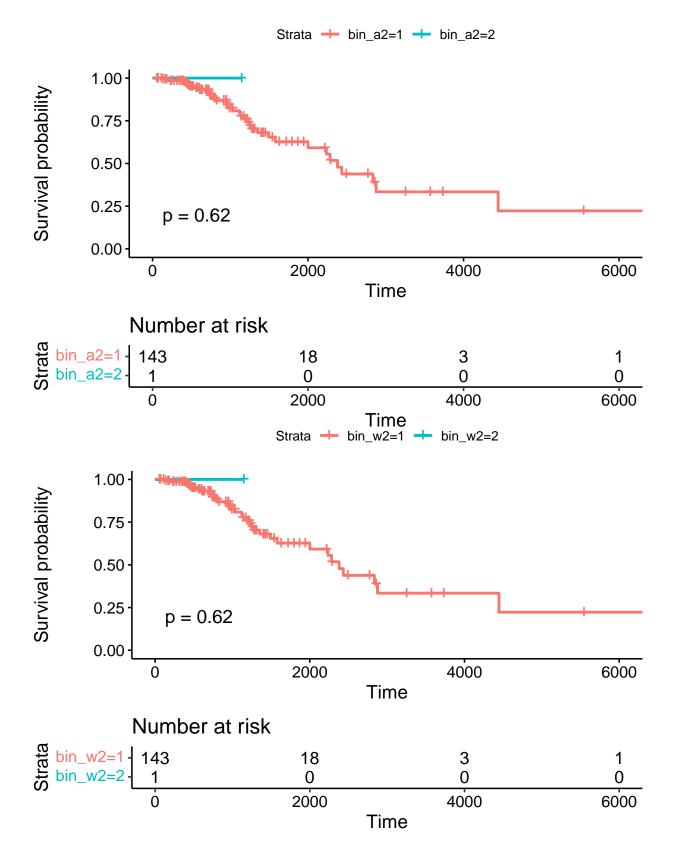


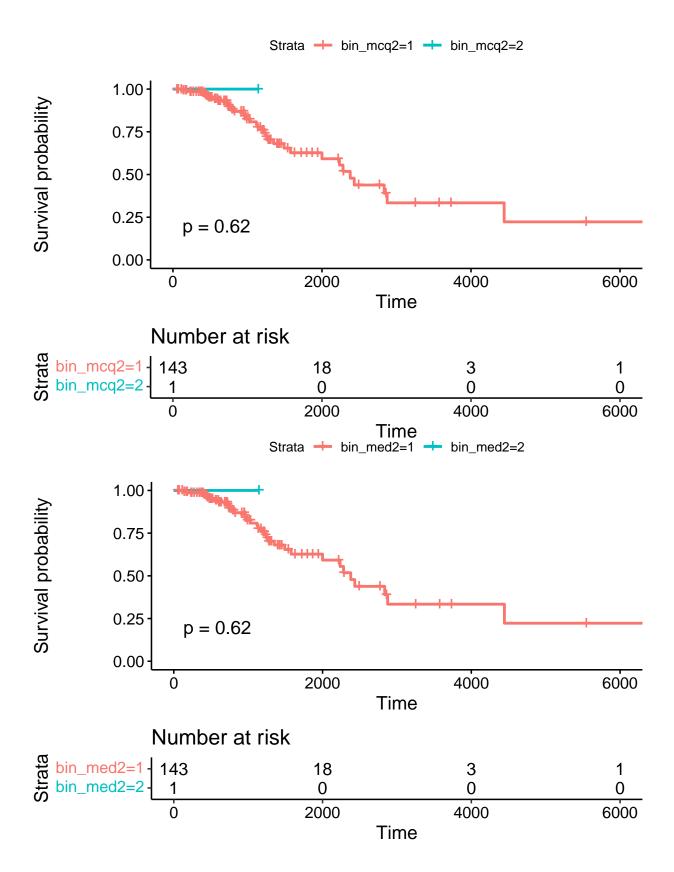


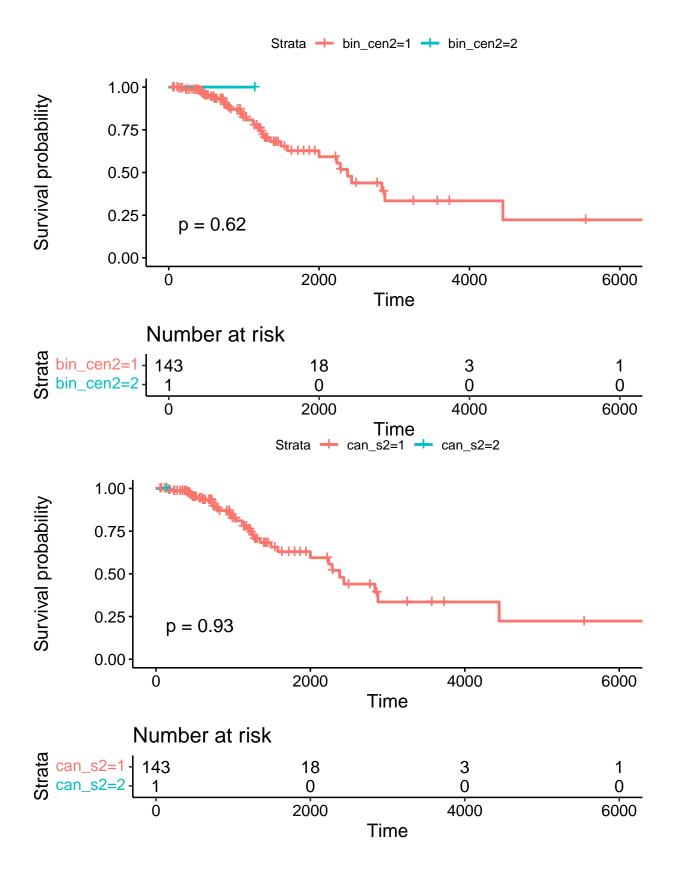
Warning in .pvalue(fit, data = data, method = method, pval = pval, pval.coord = pval.coord, : There
This is a null model.

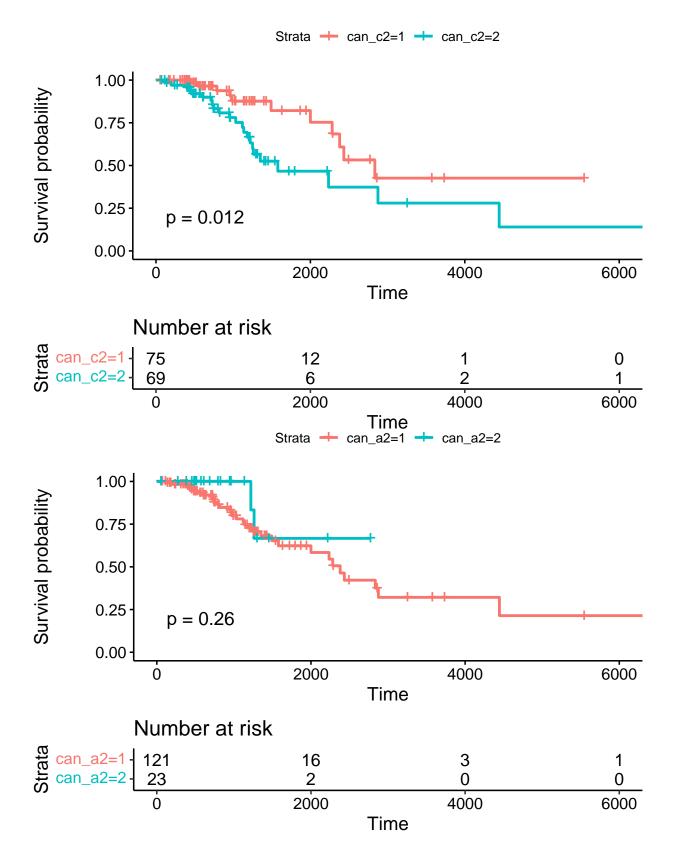


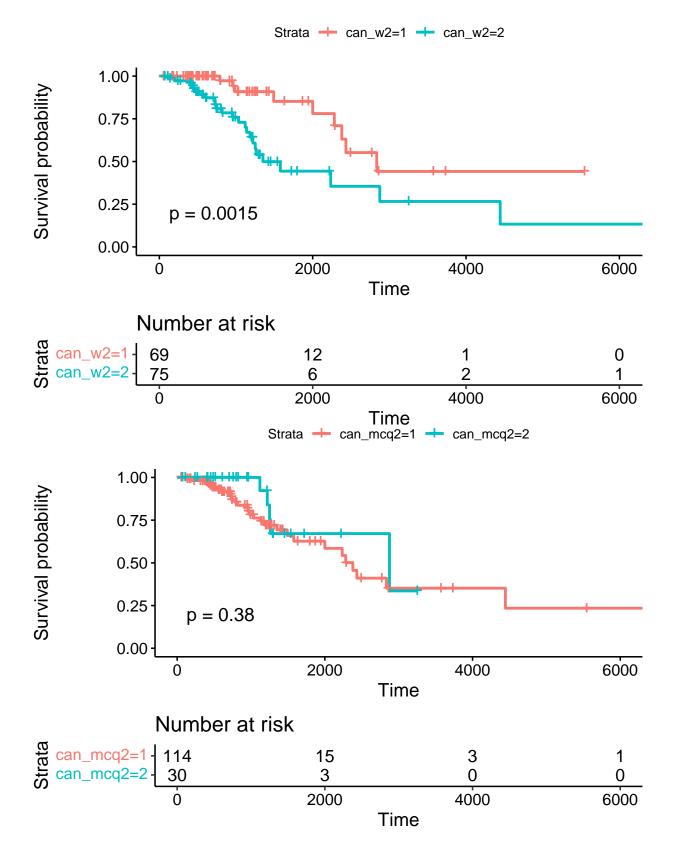


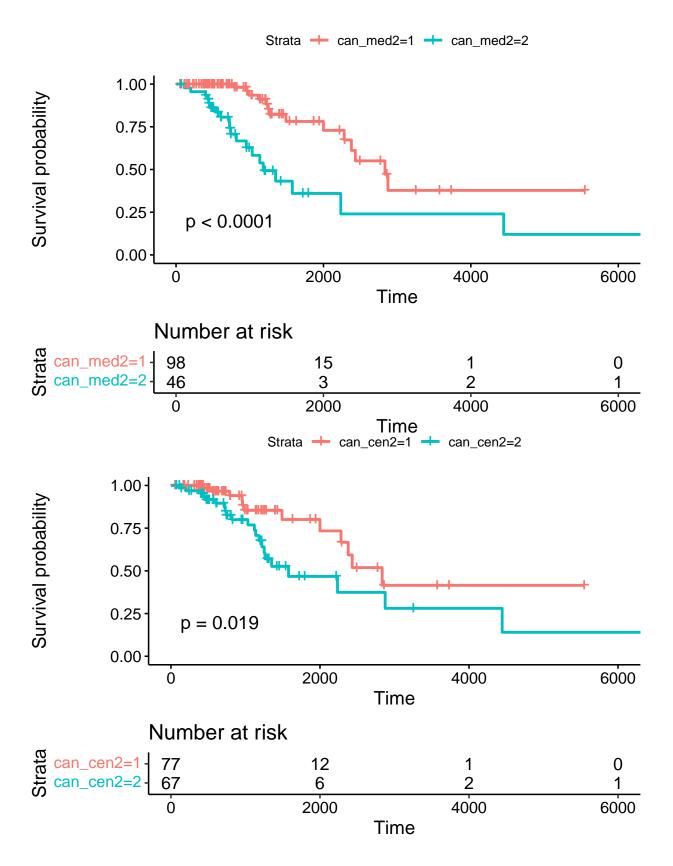


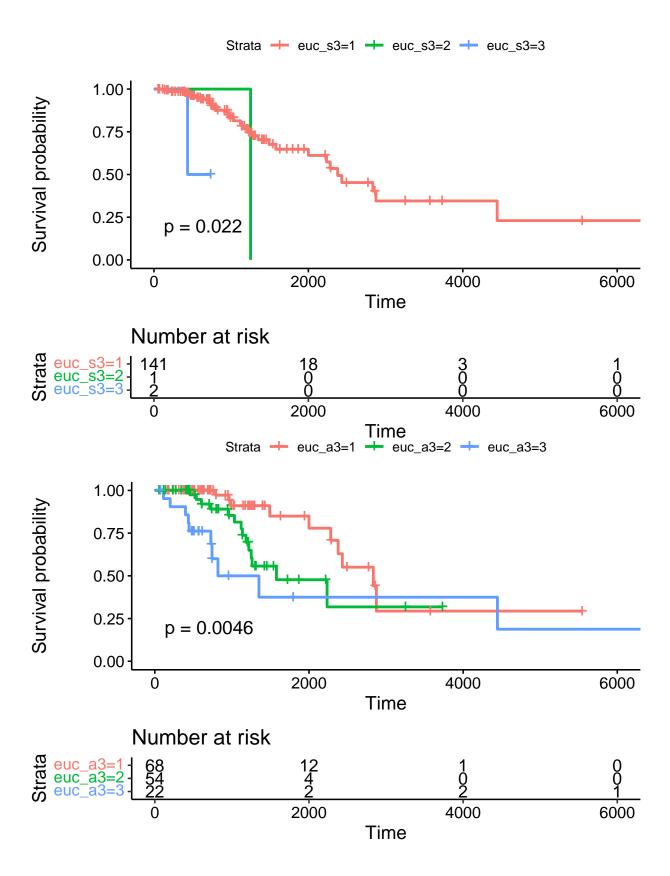


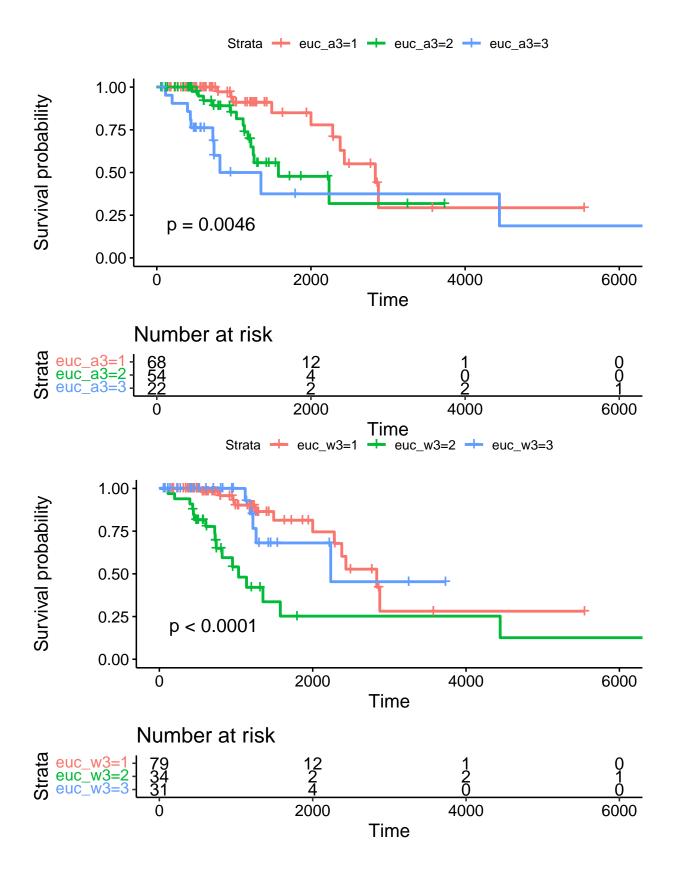


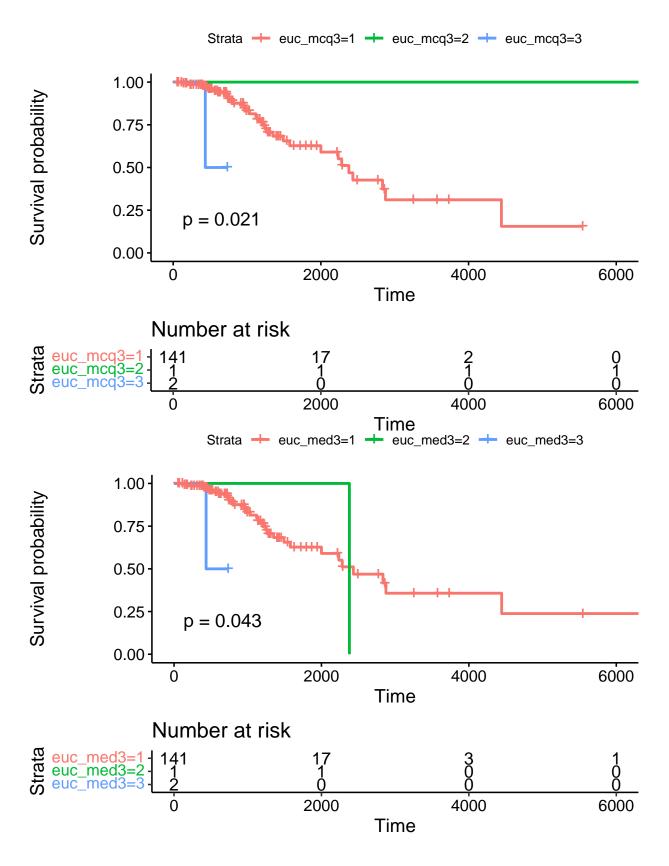


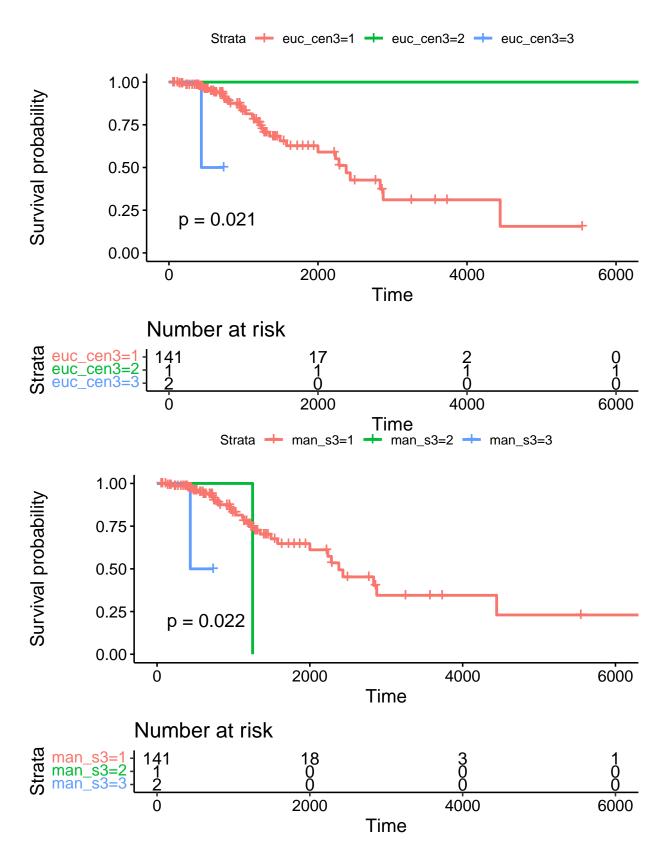


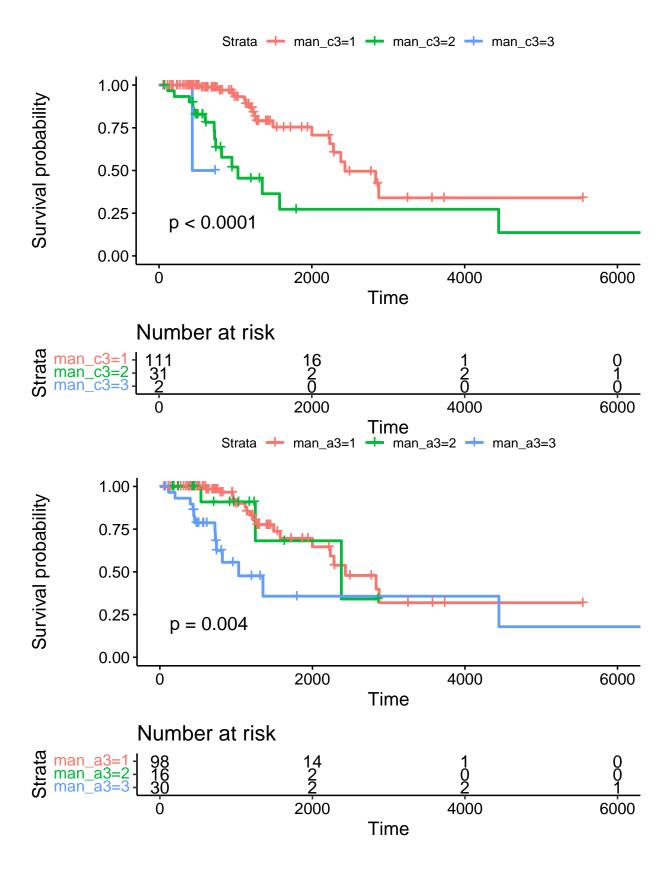


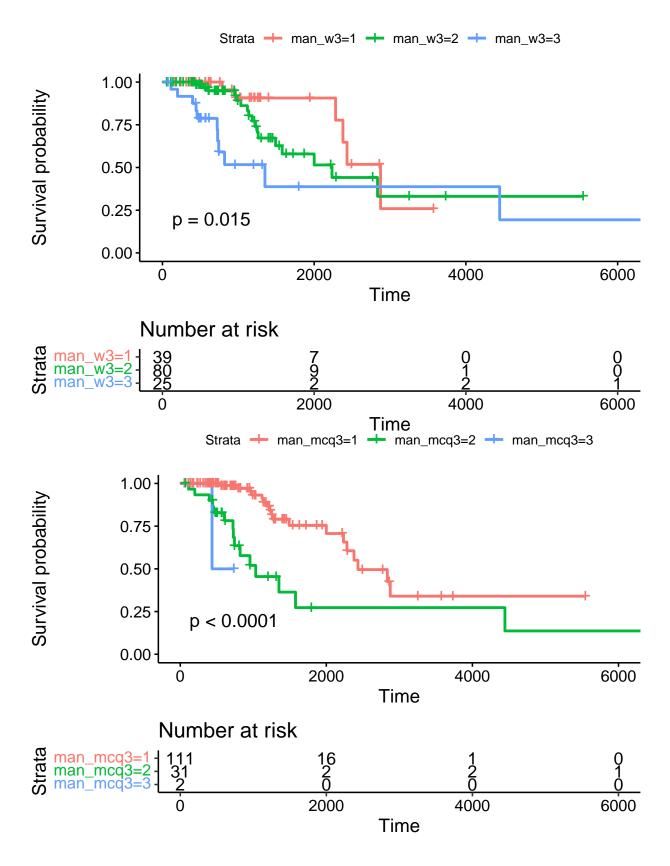


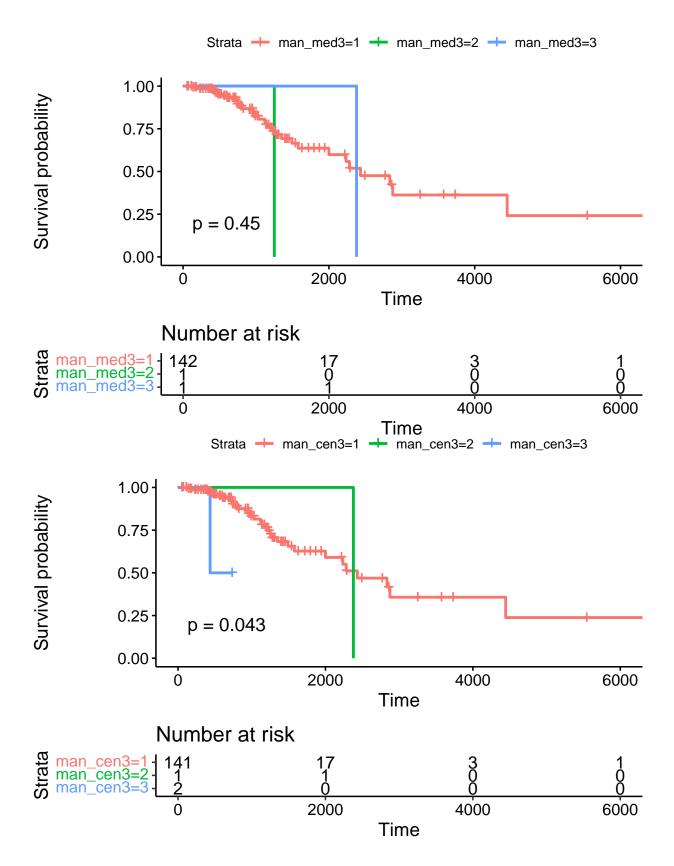


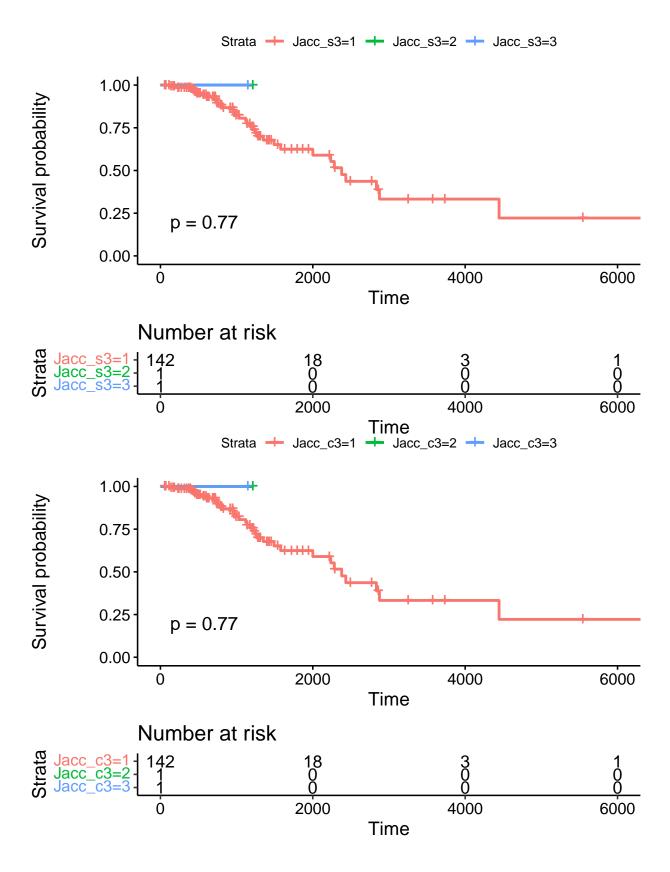


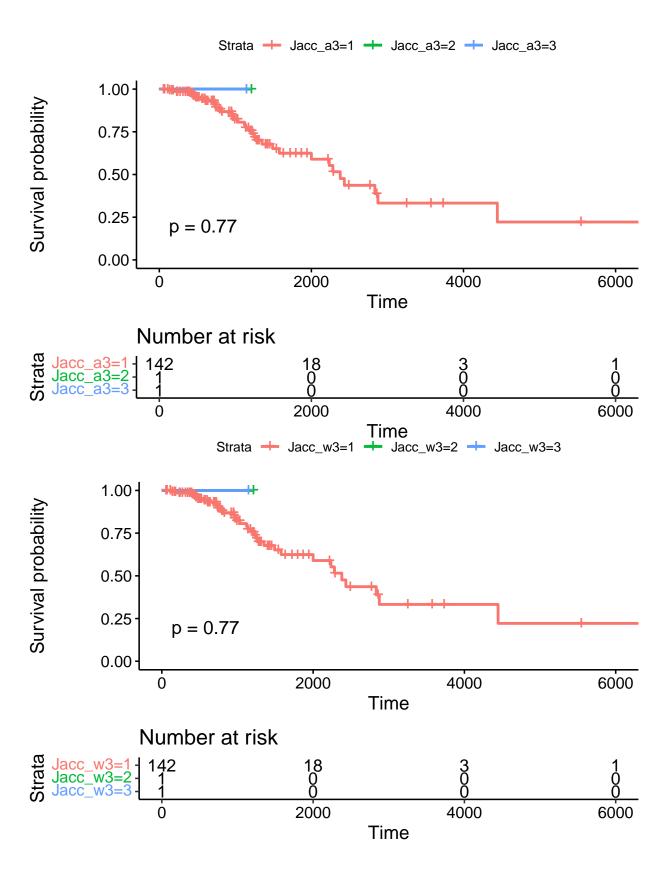


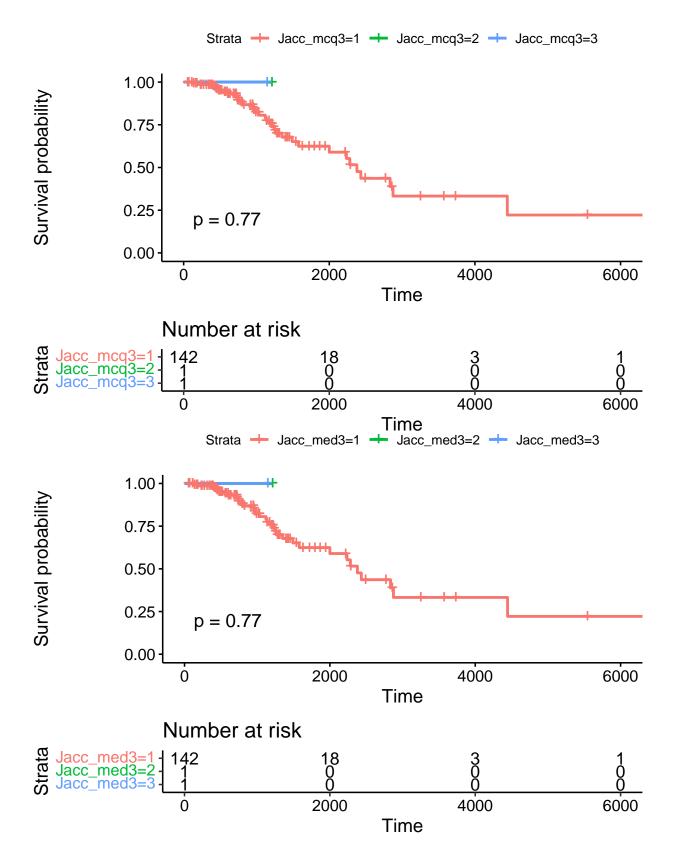


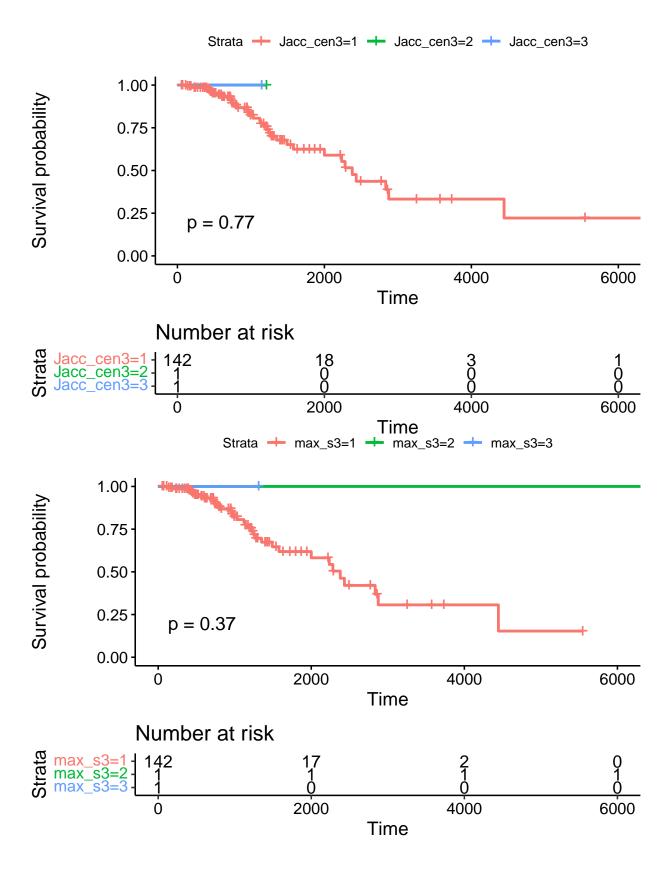


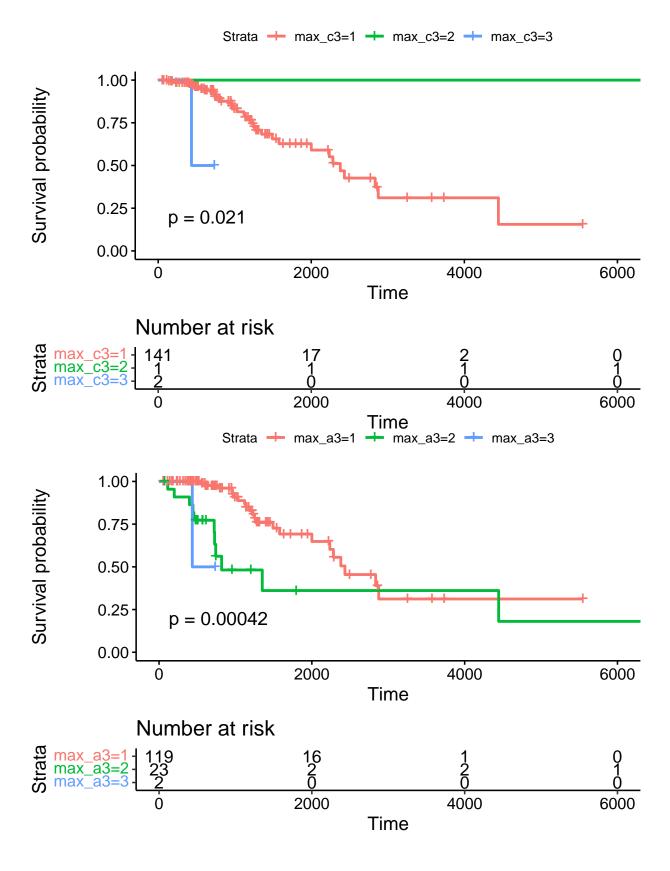


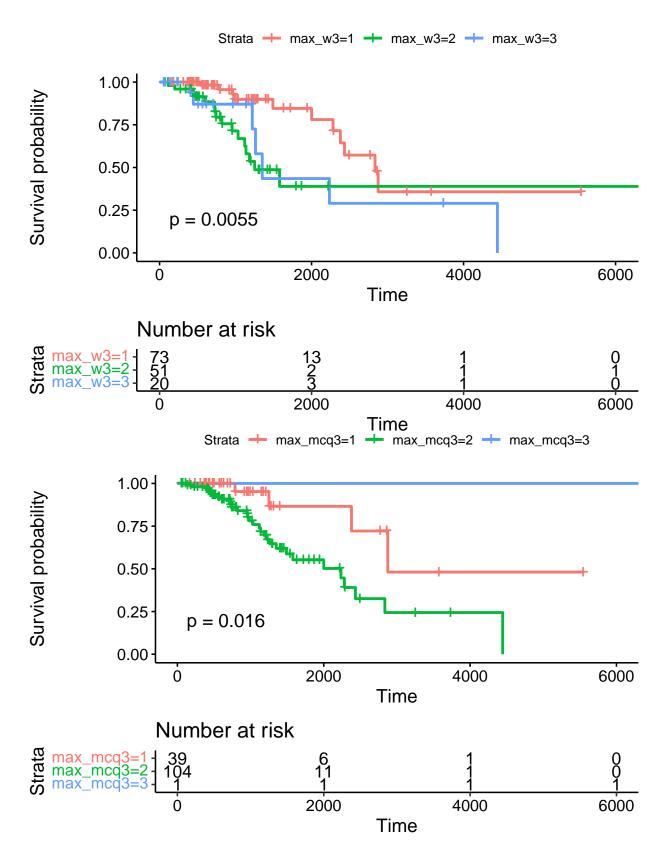


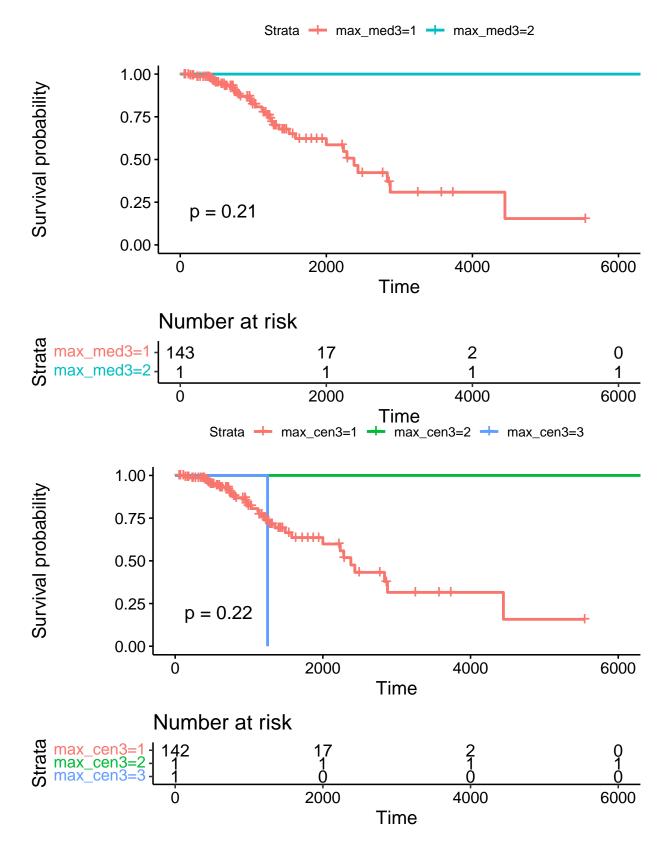


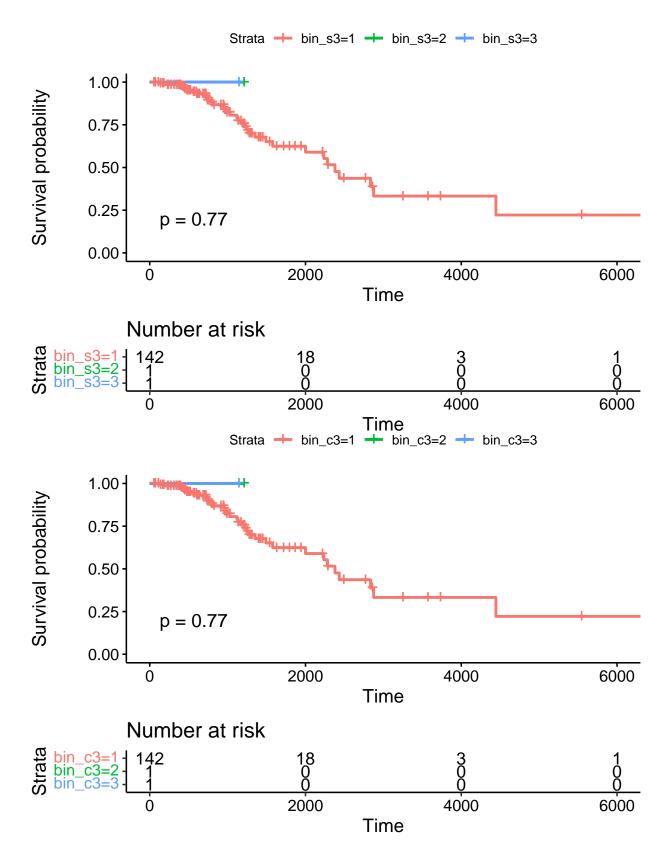


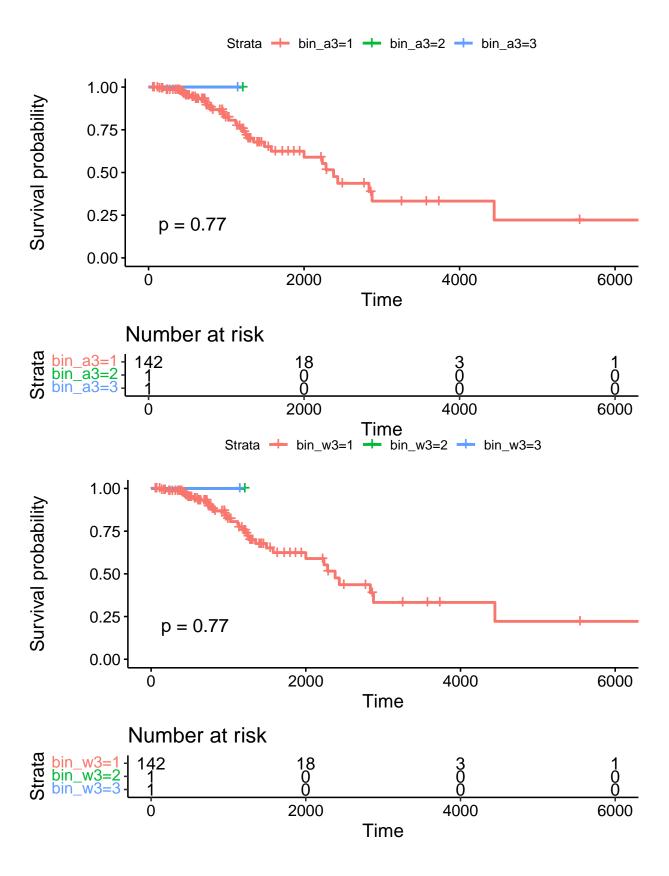


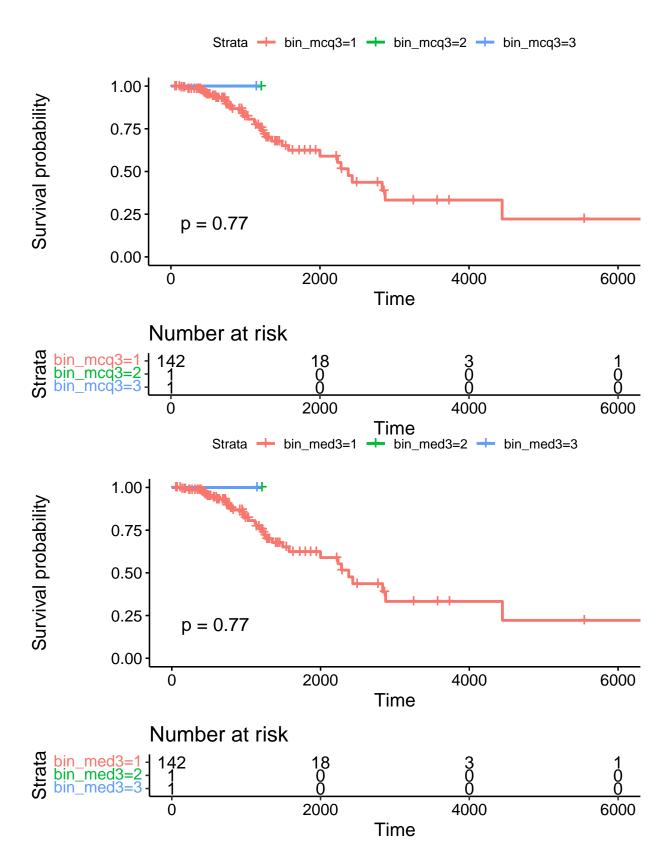


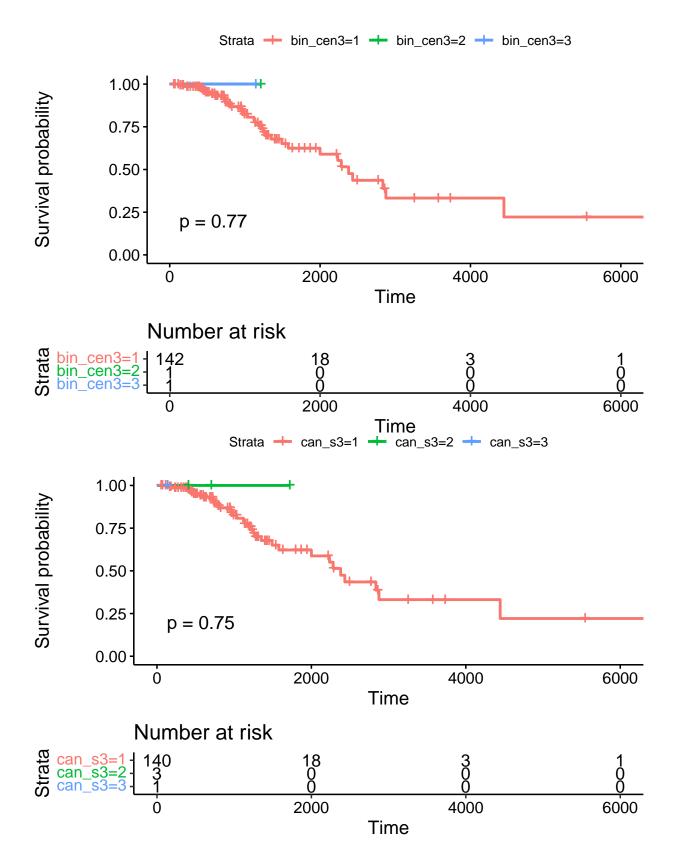


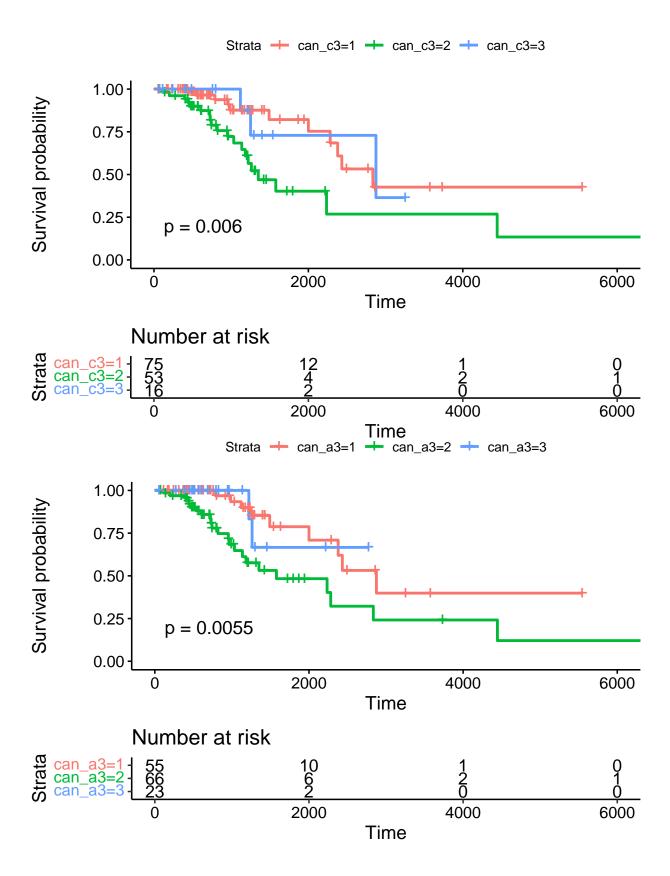


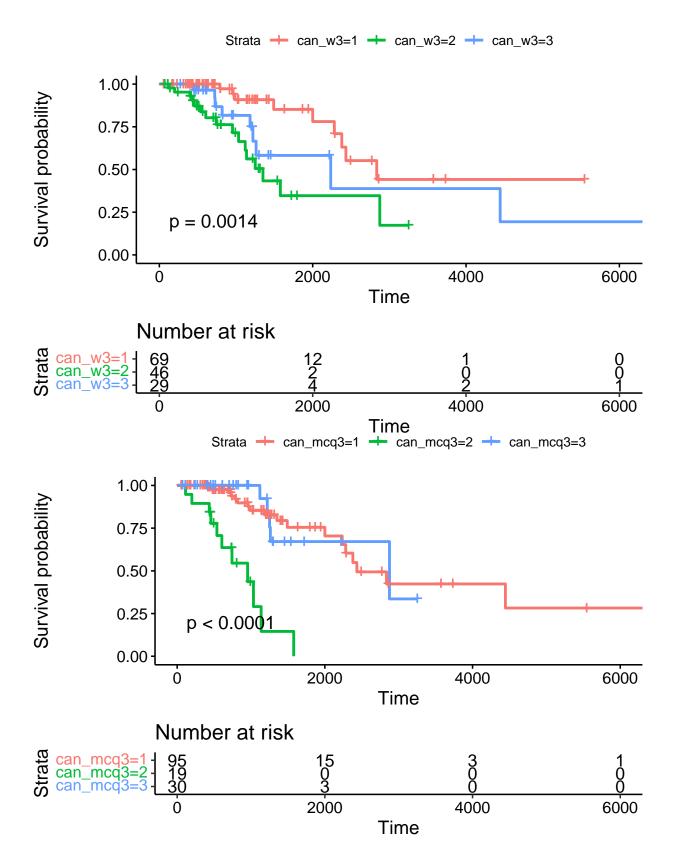


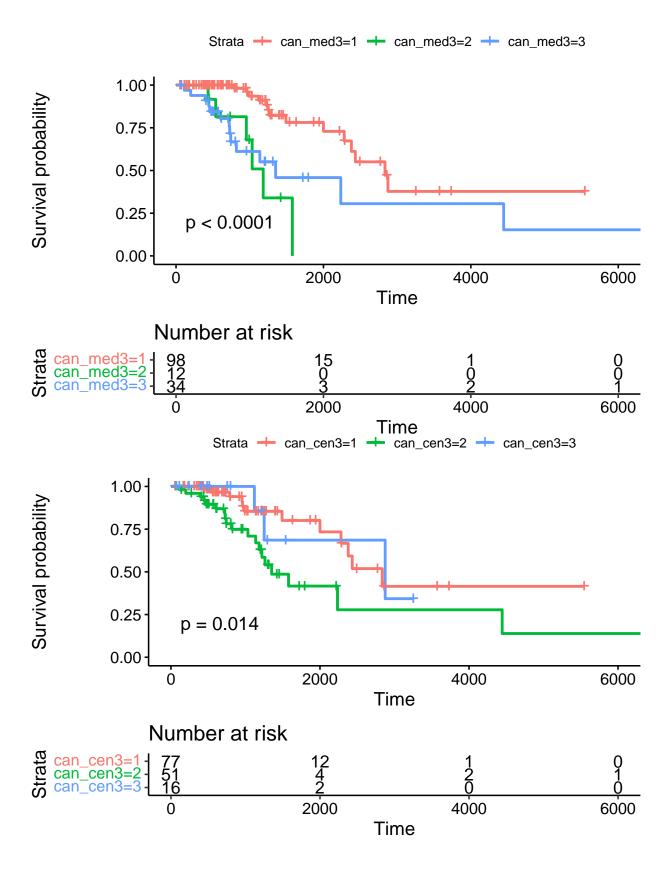


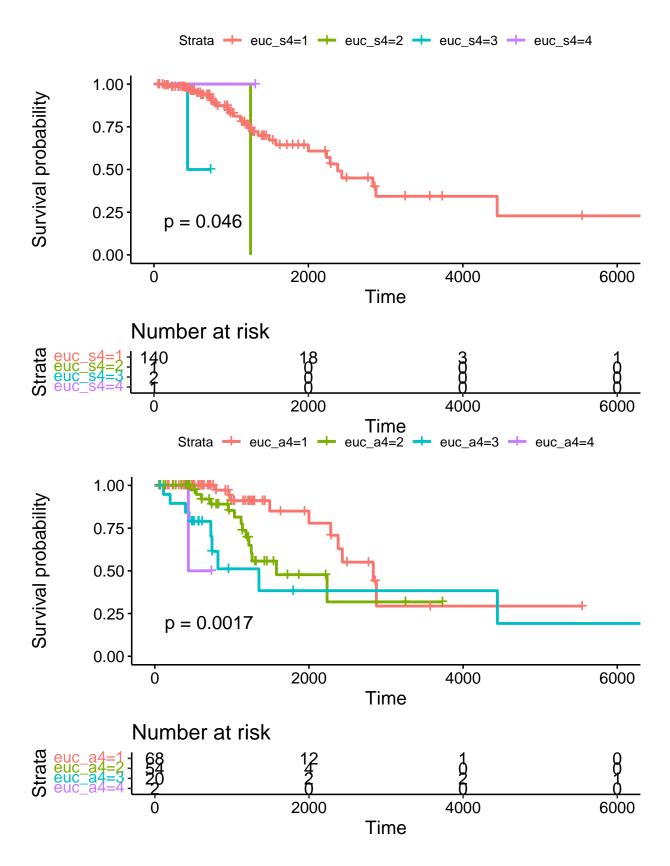


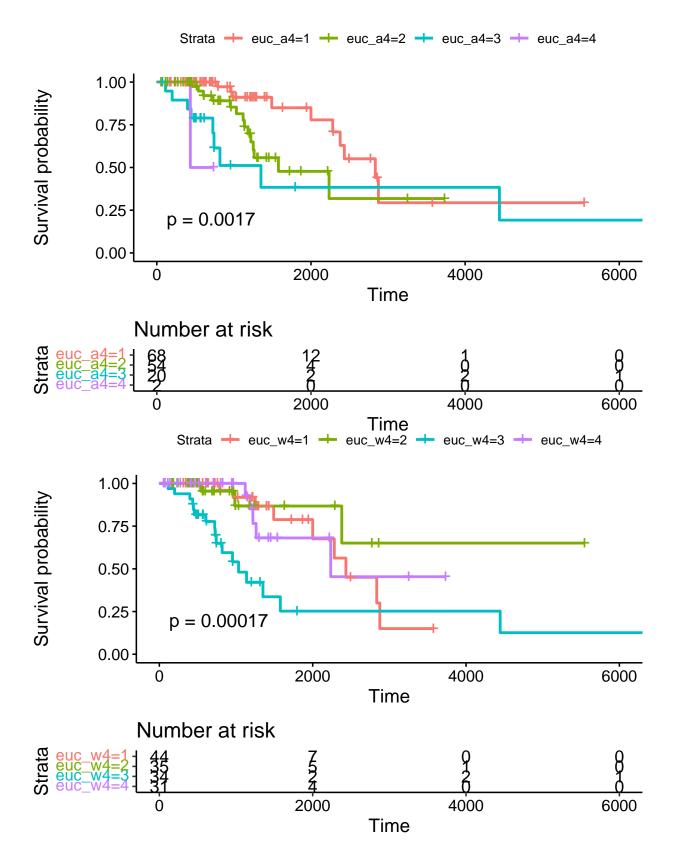


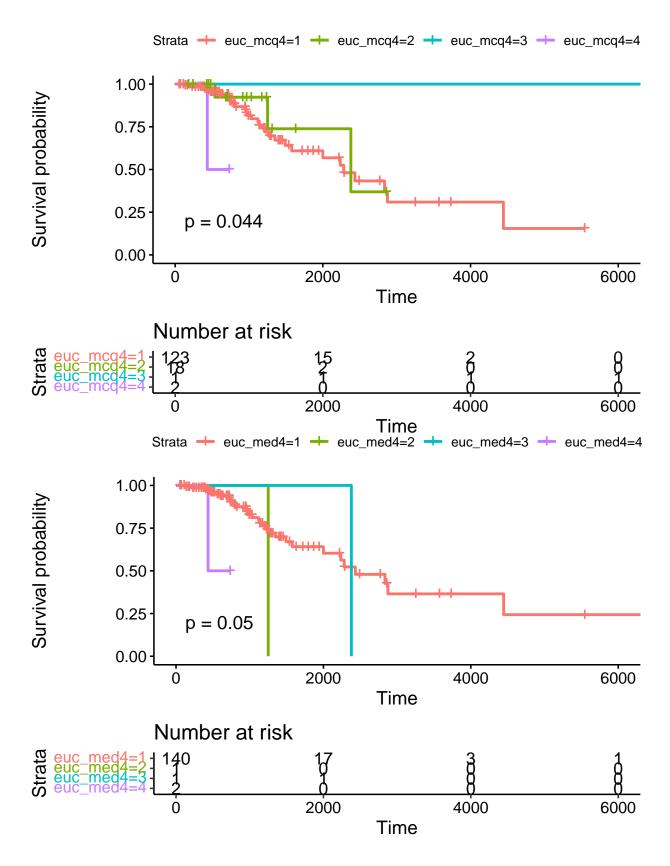


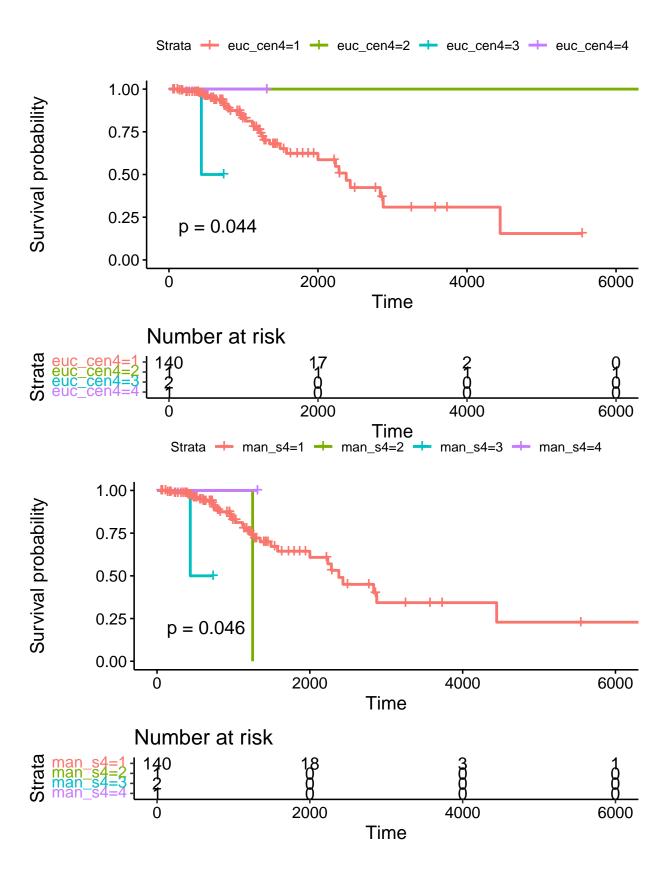


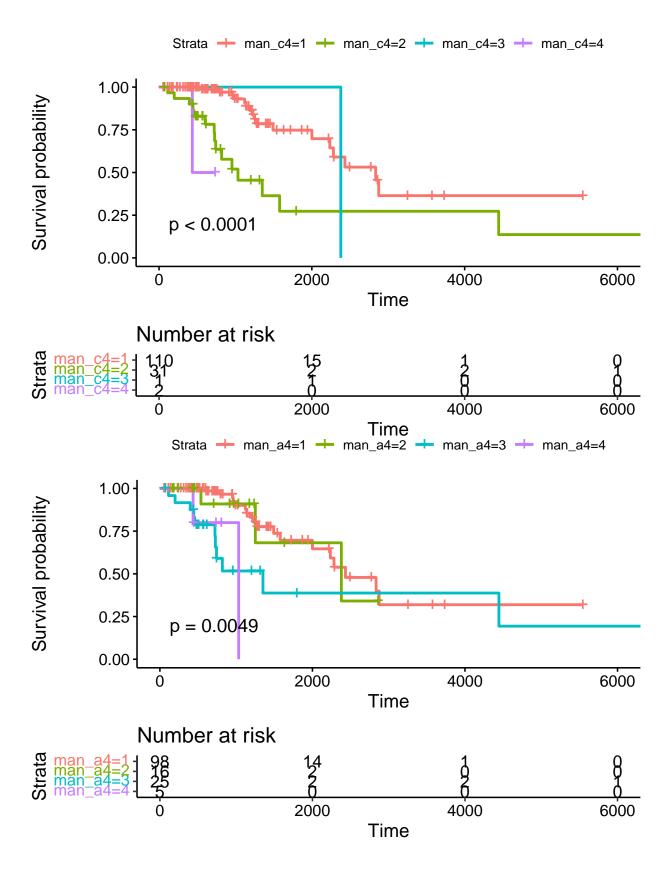


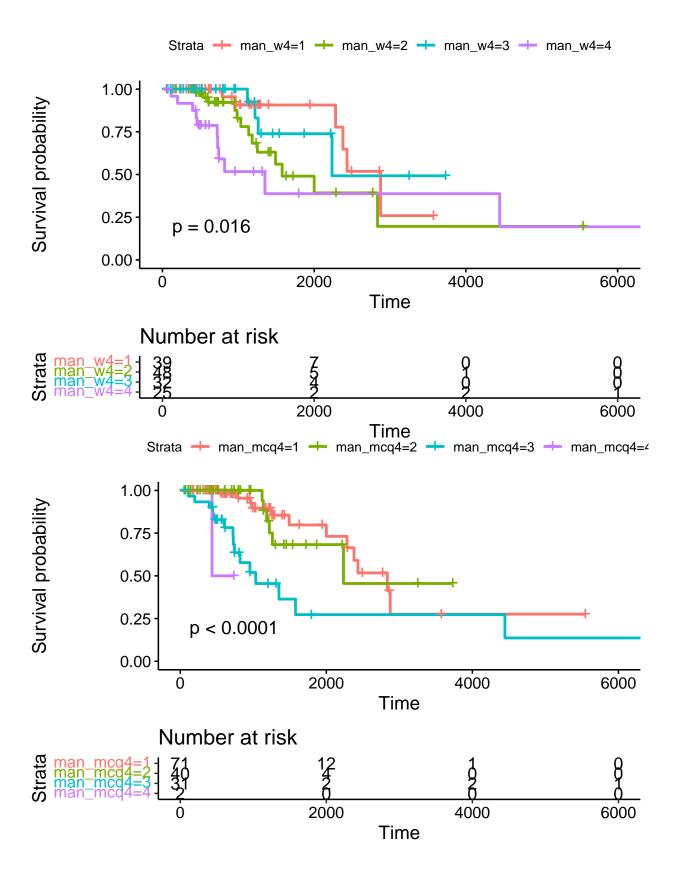


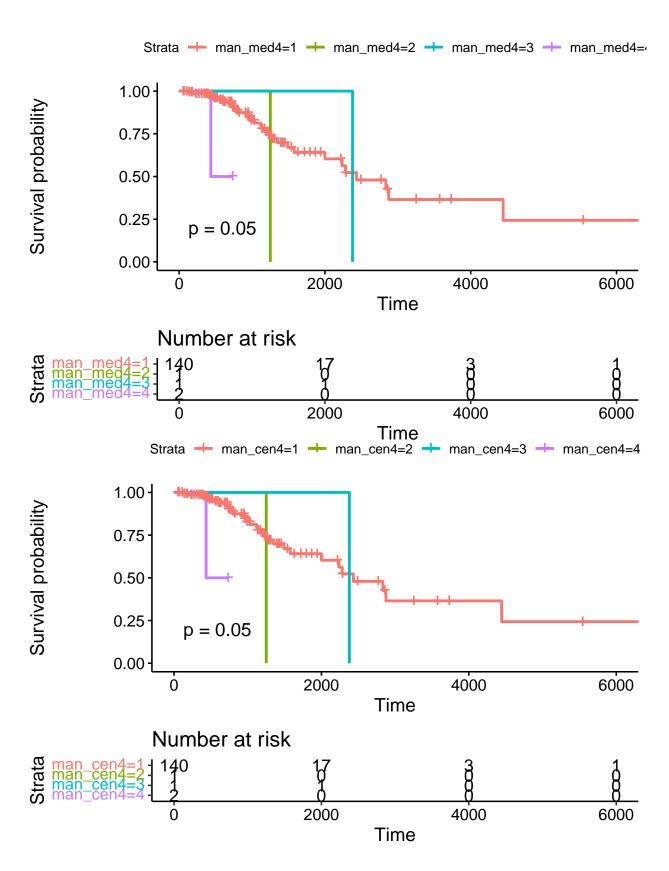


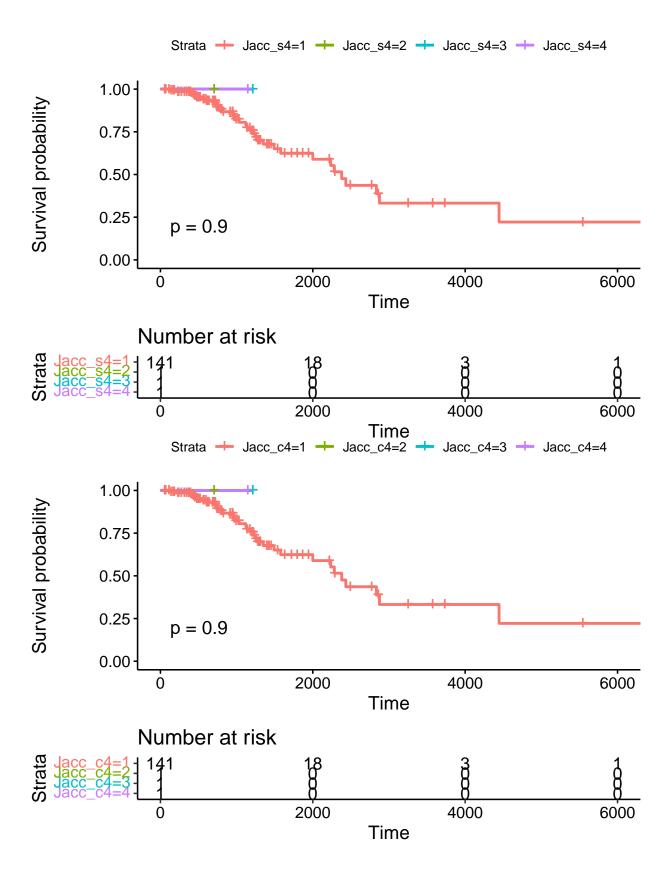


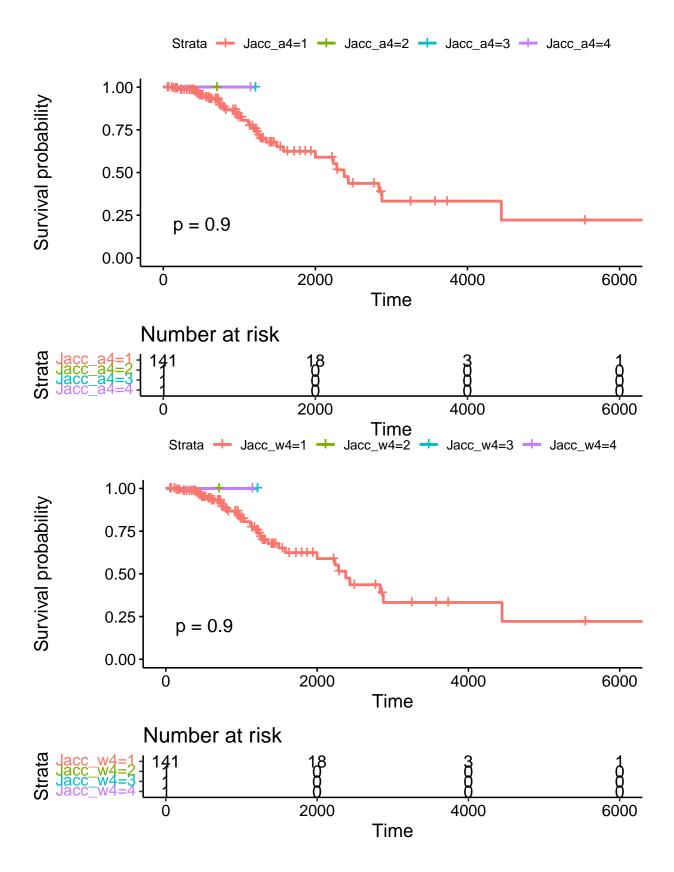


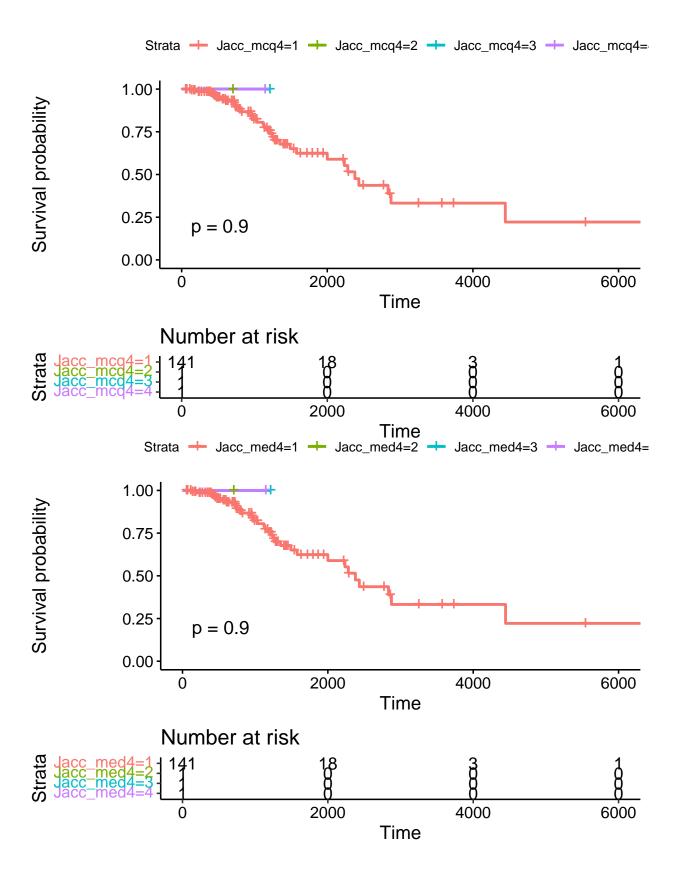


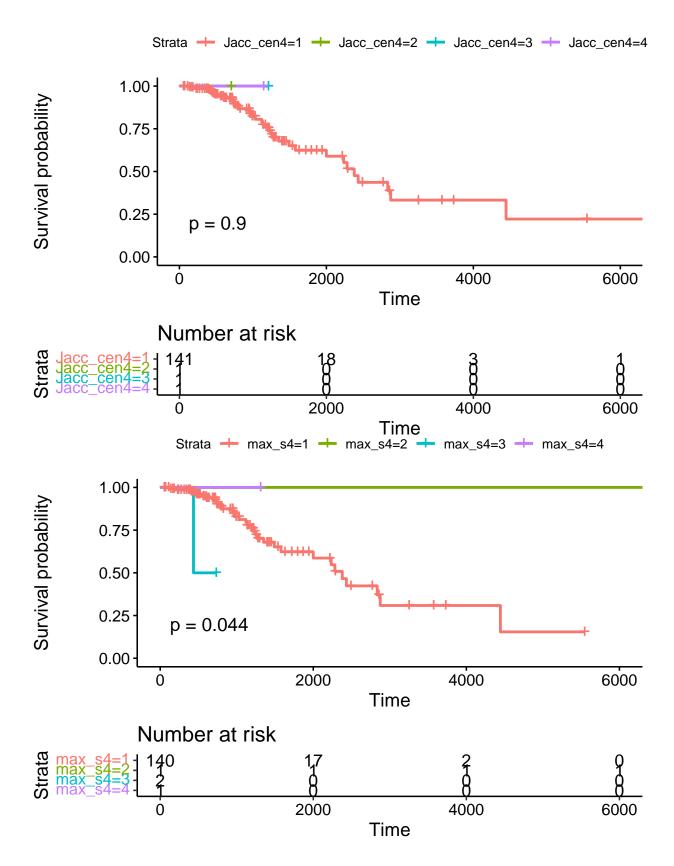


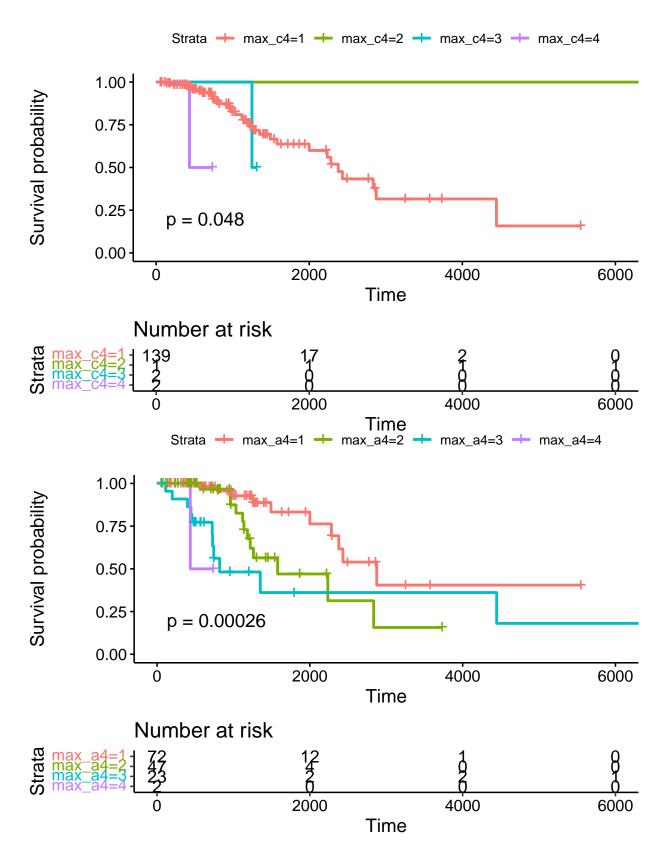


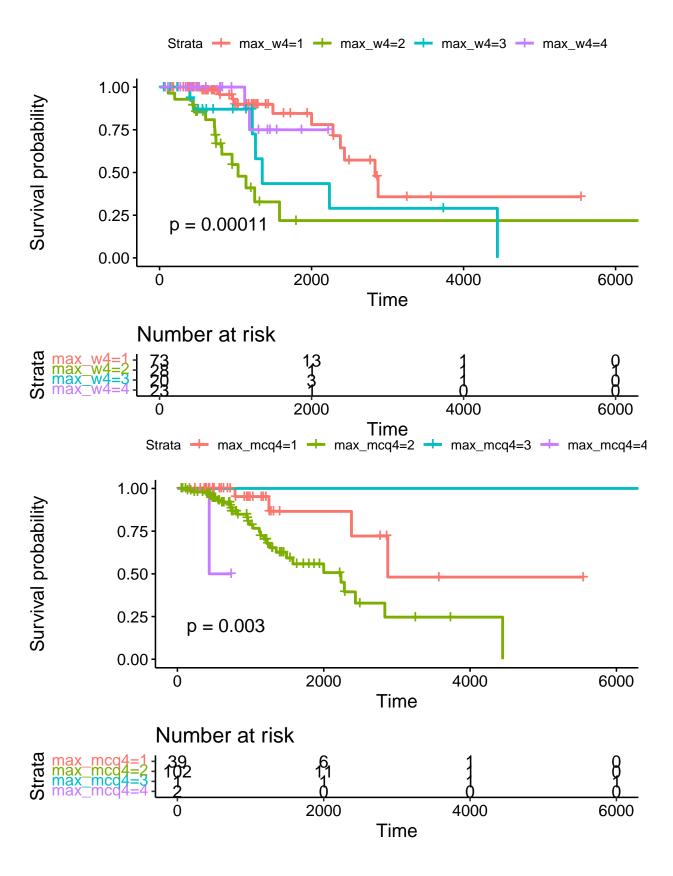


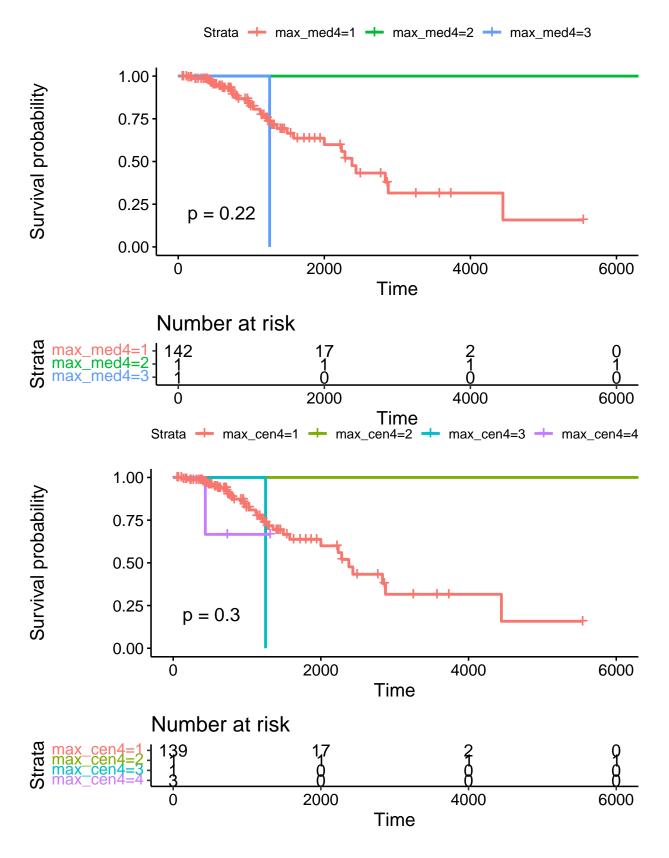


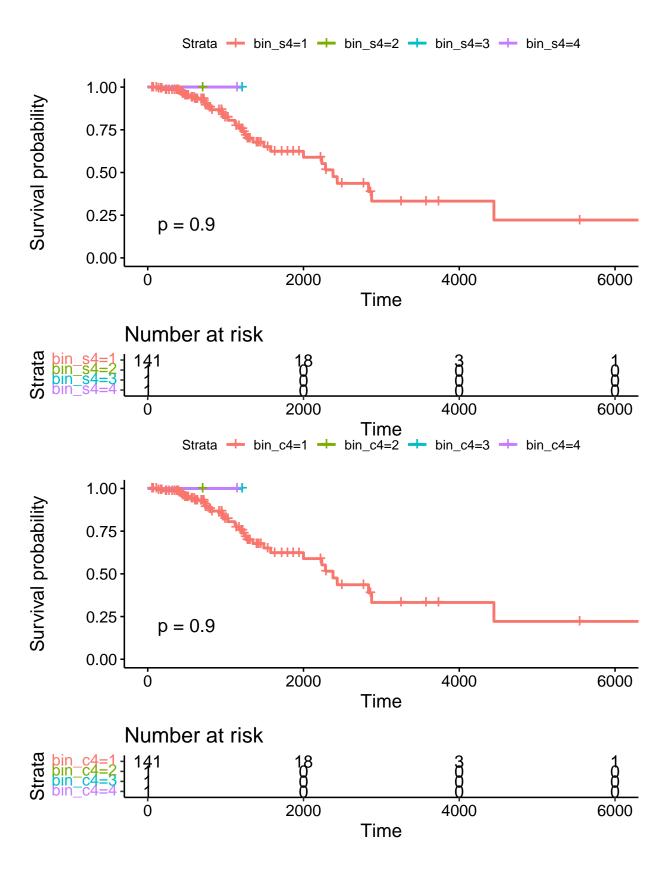


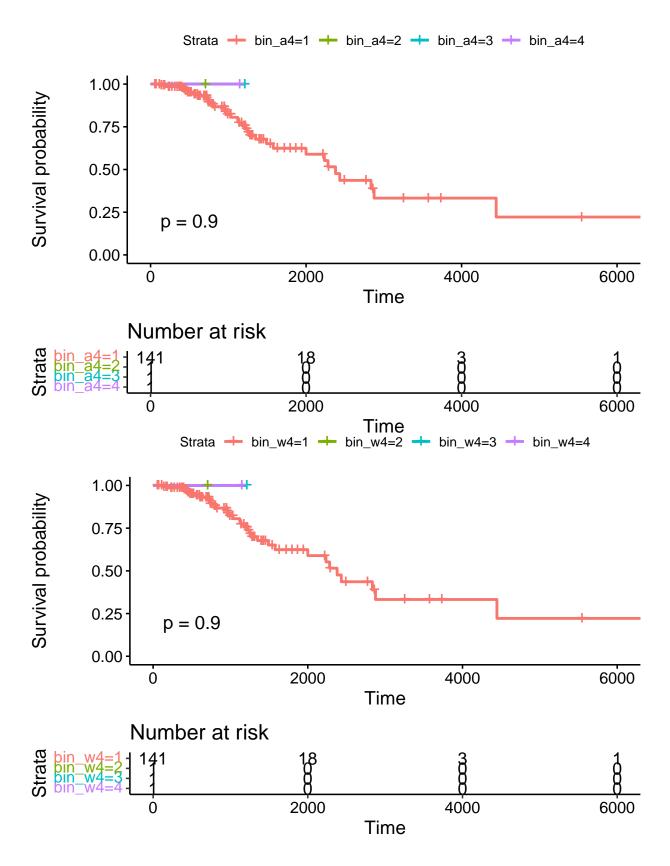


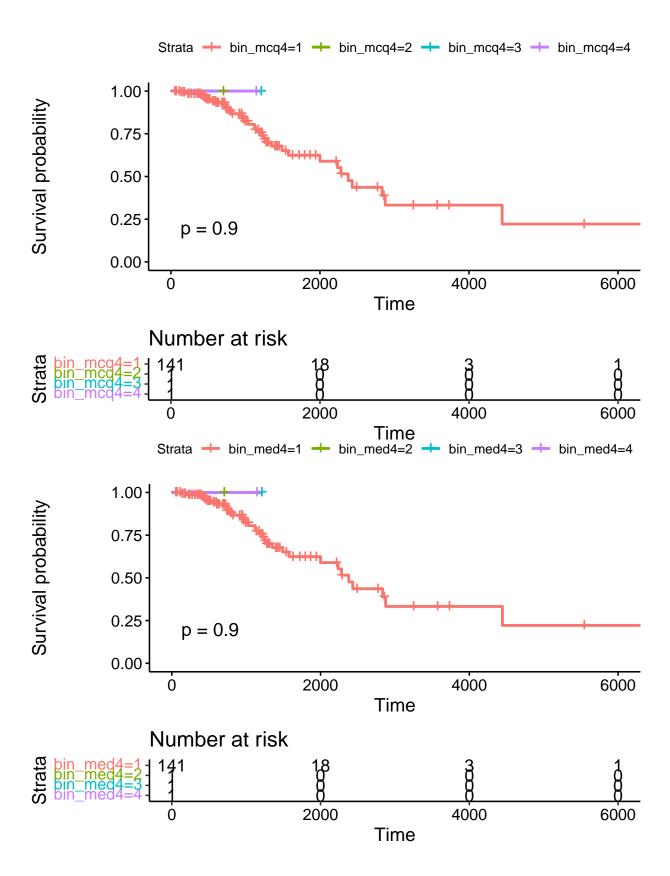


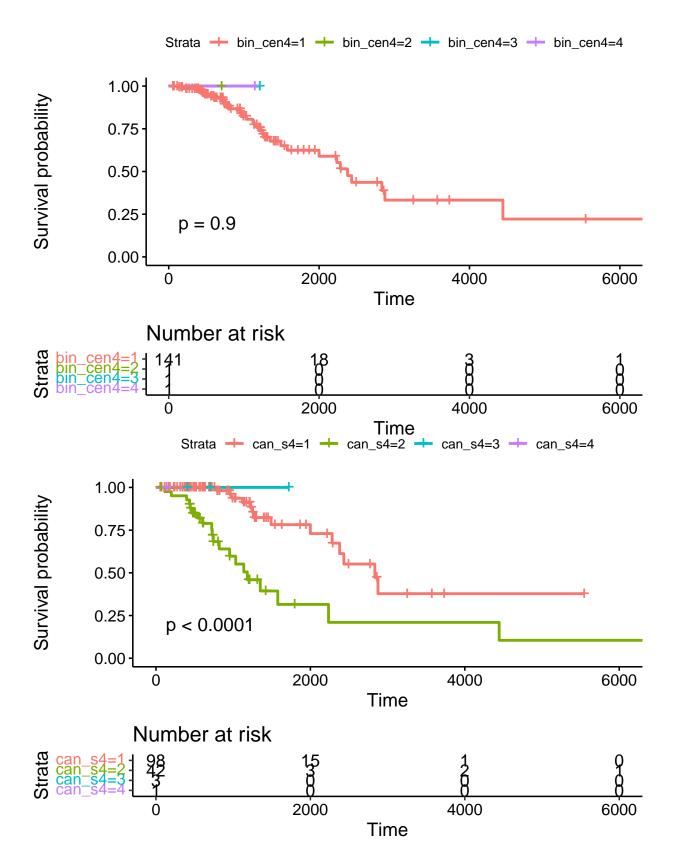


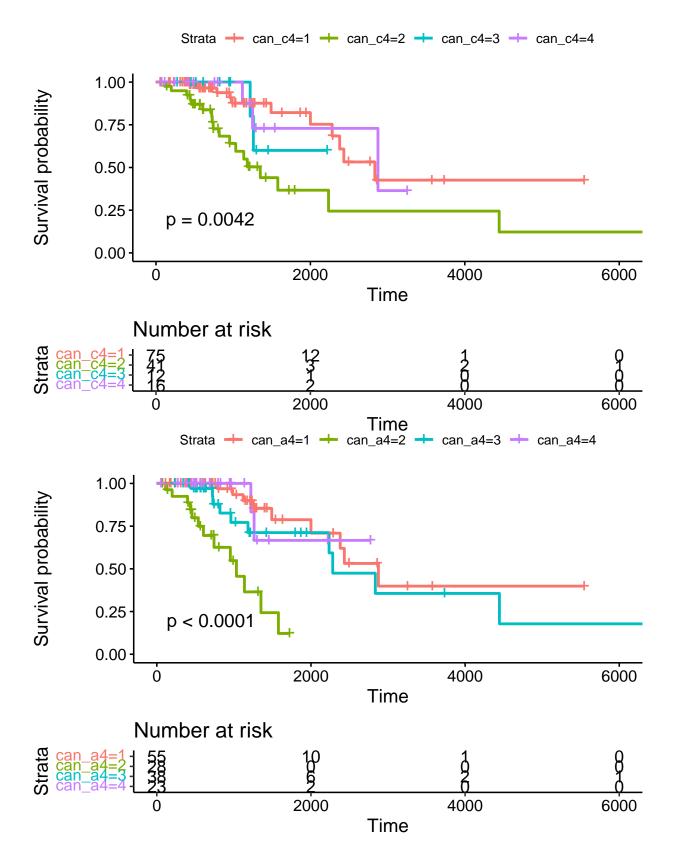


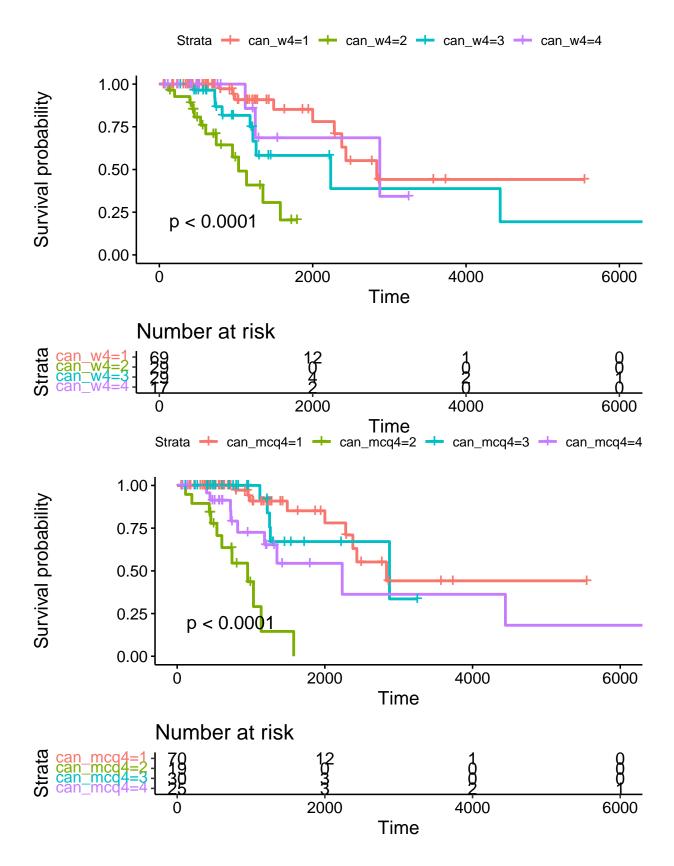


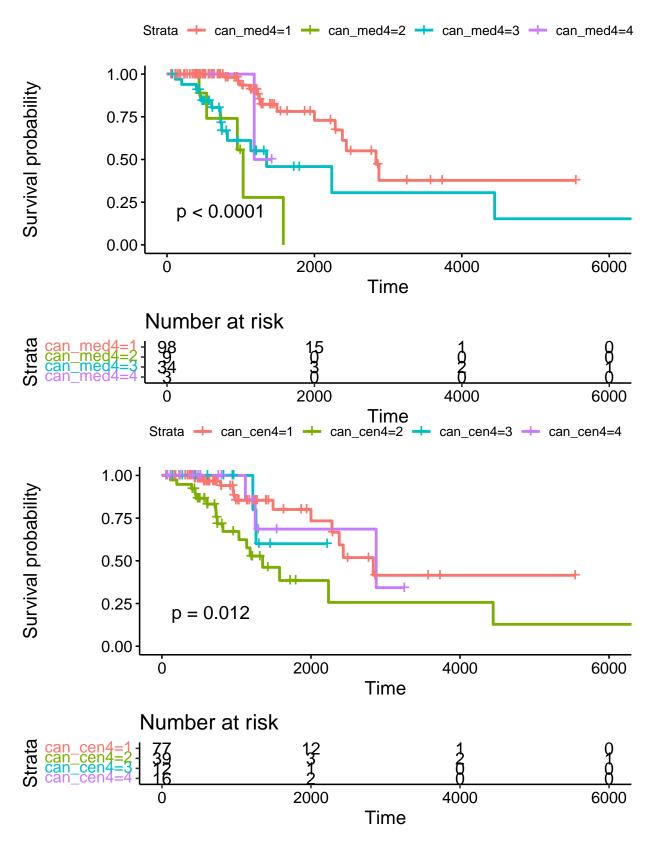






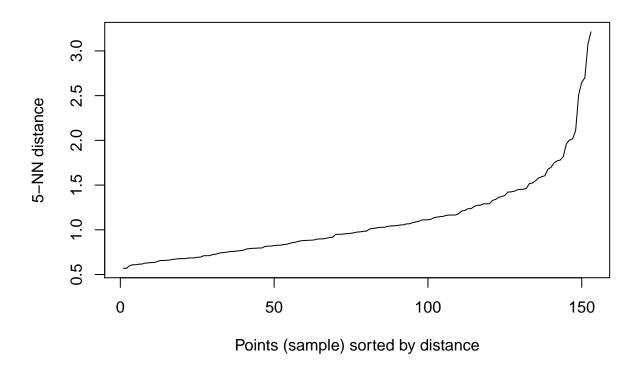






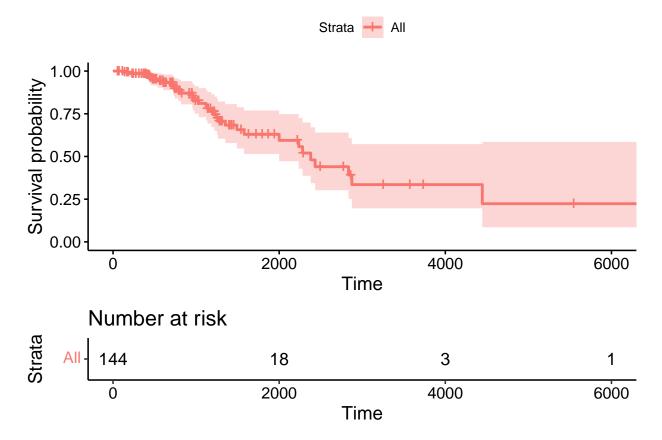
3. DBSCAN

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

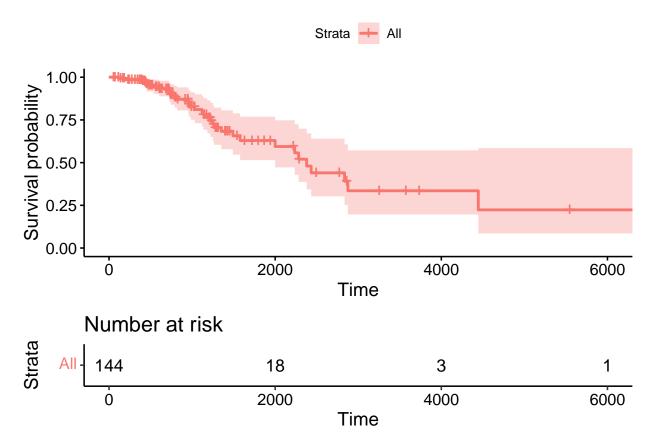


Warning in dbscan(tucker1, eps = 16, MinPts = 20): converting argument MinPts
(fpc) to minPts (dbscan)!

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



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



4. EM algorithm based on GMM

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

