

# cp decomposition glioma 73 train rank5

VIVI TANG

17/11/2021

## 1. Kmeans

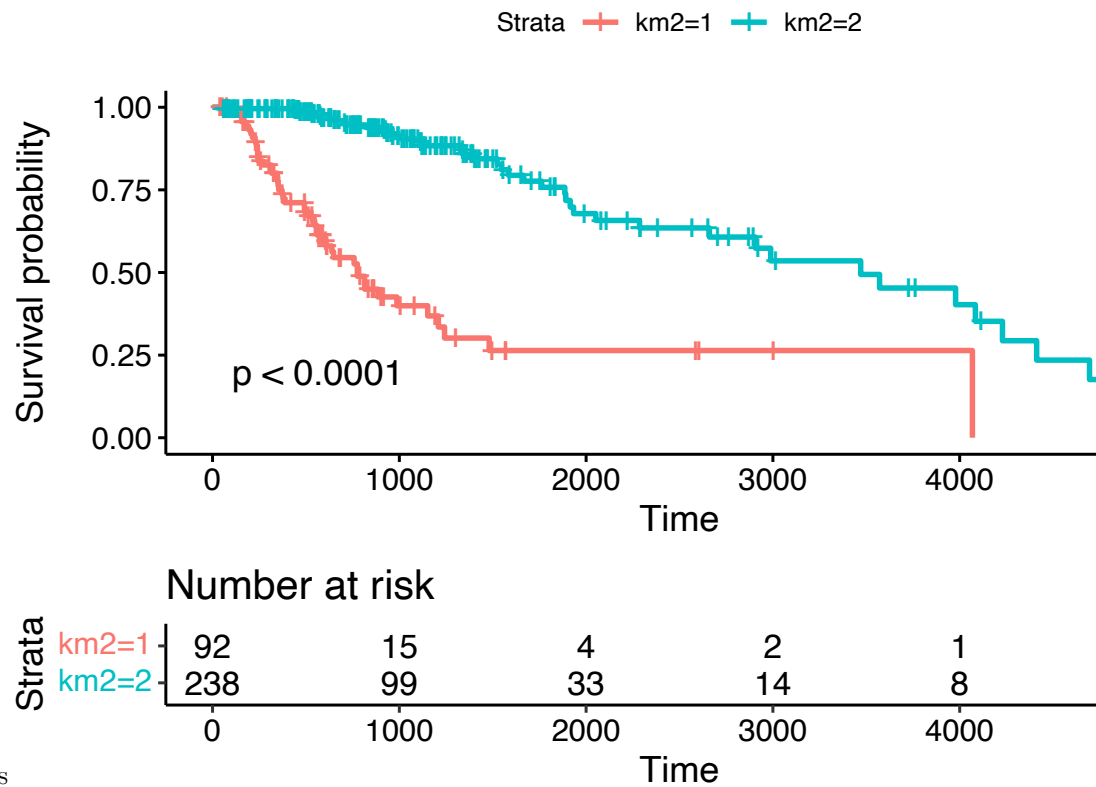
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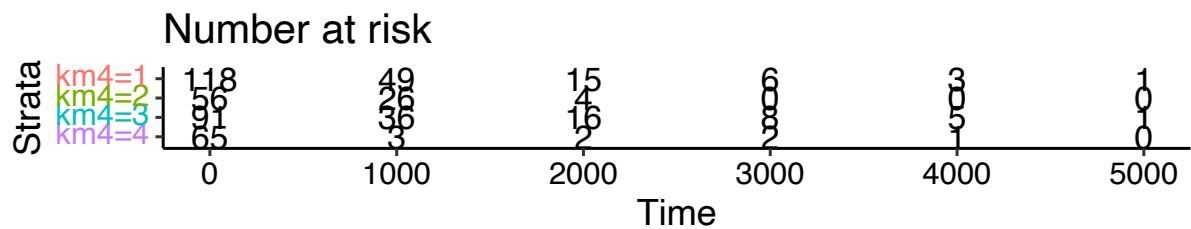
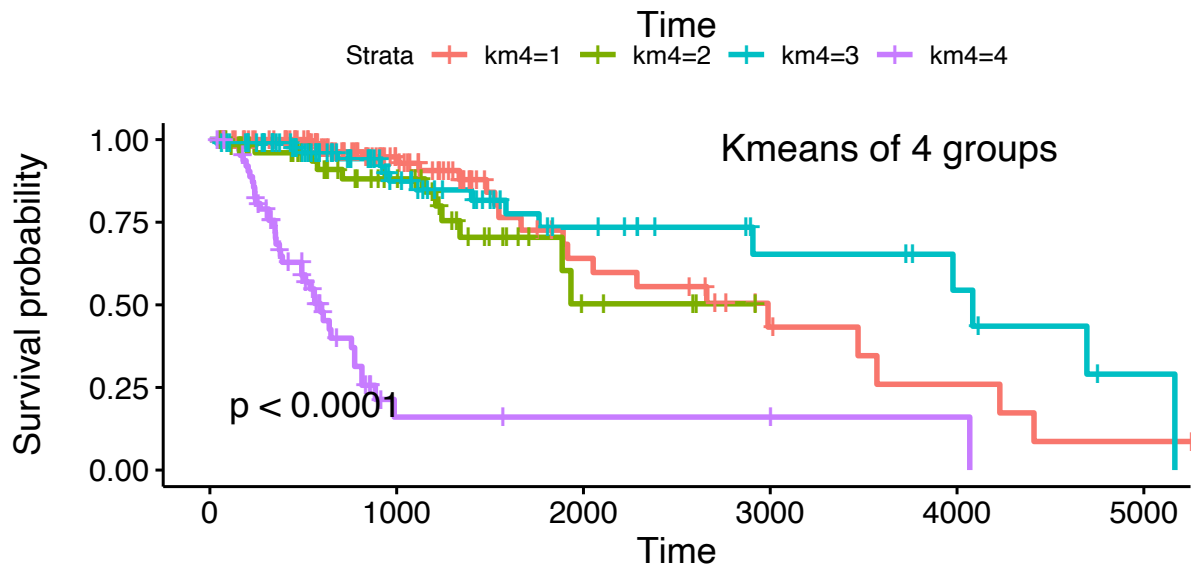
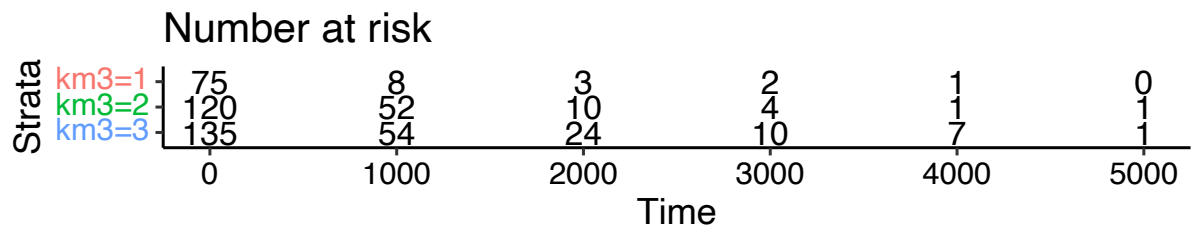
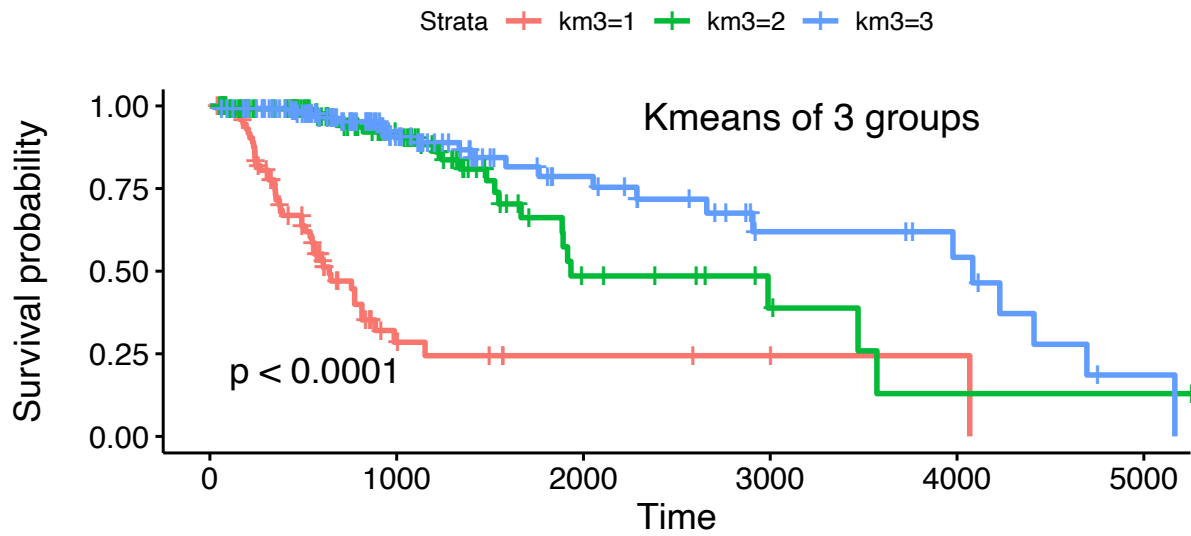
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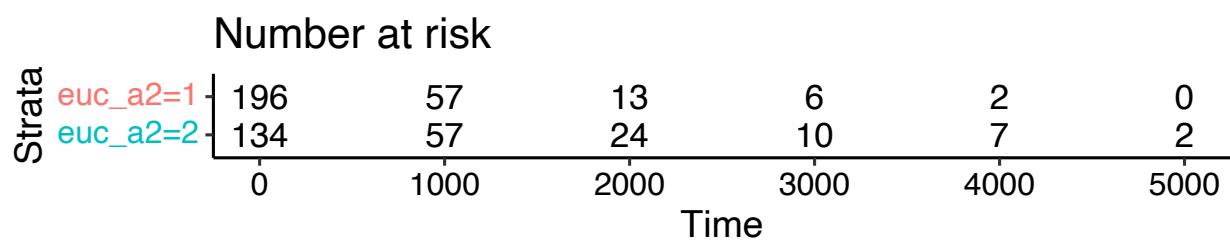
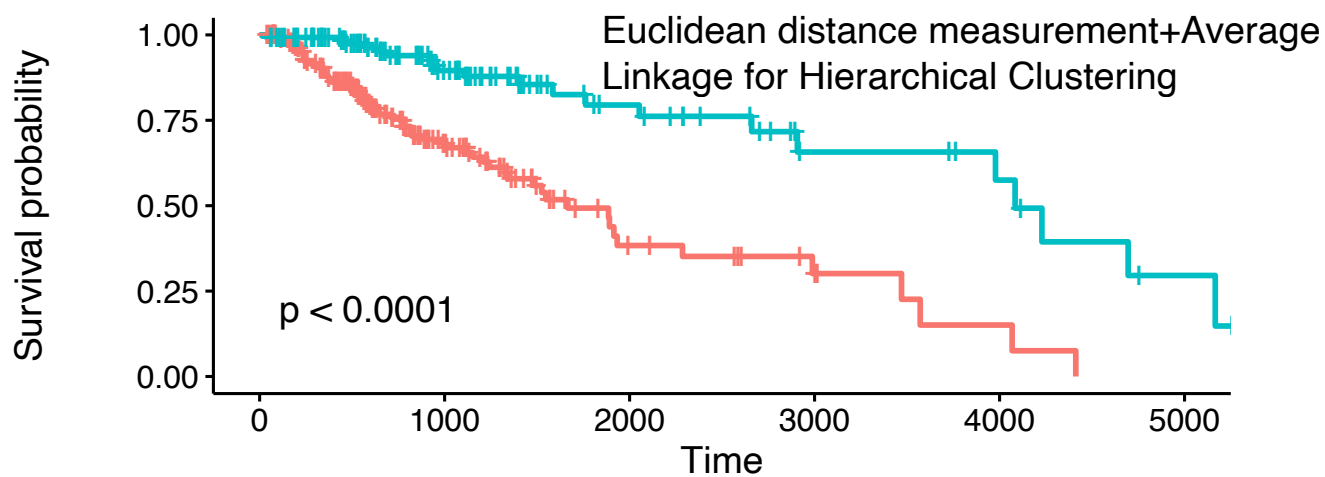
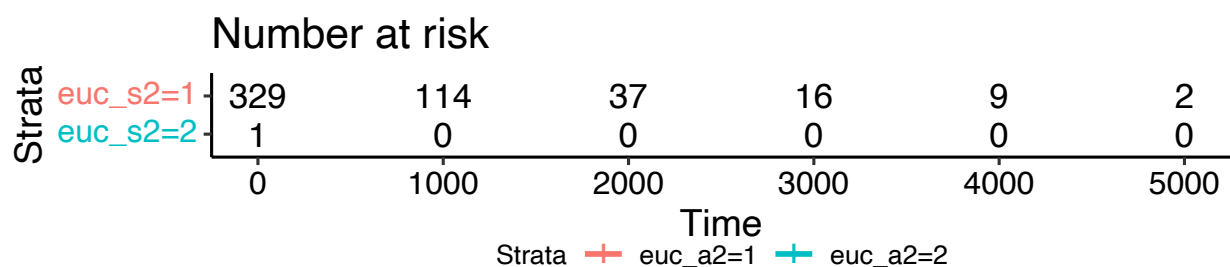
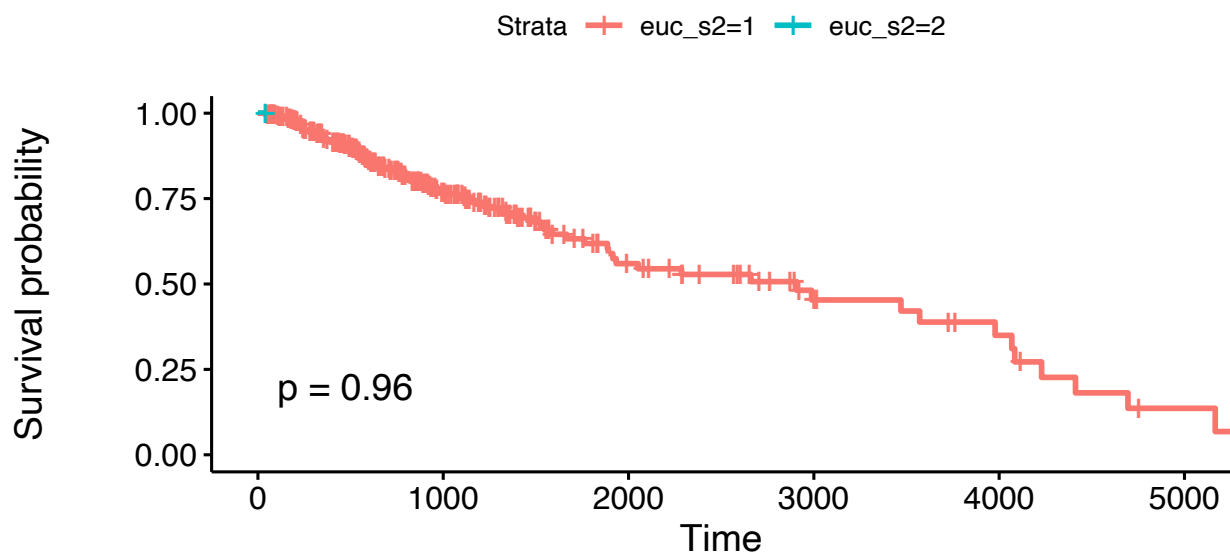
## 3. Fit the results of Kmeans

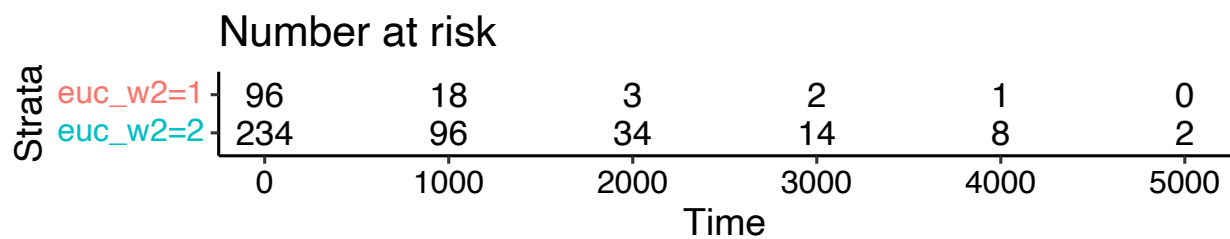
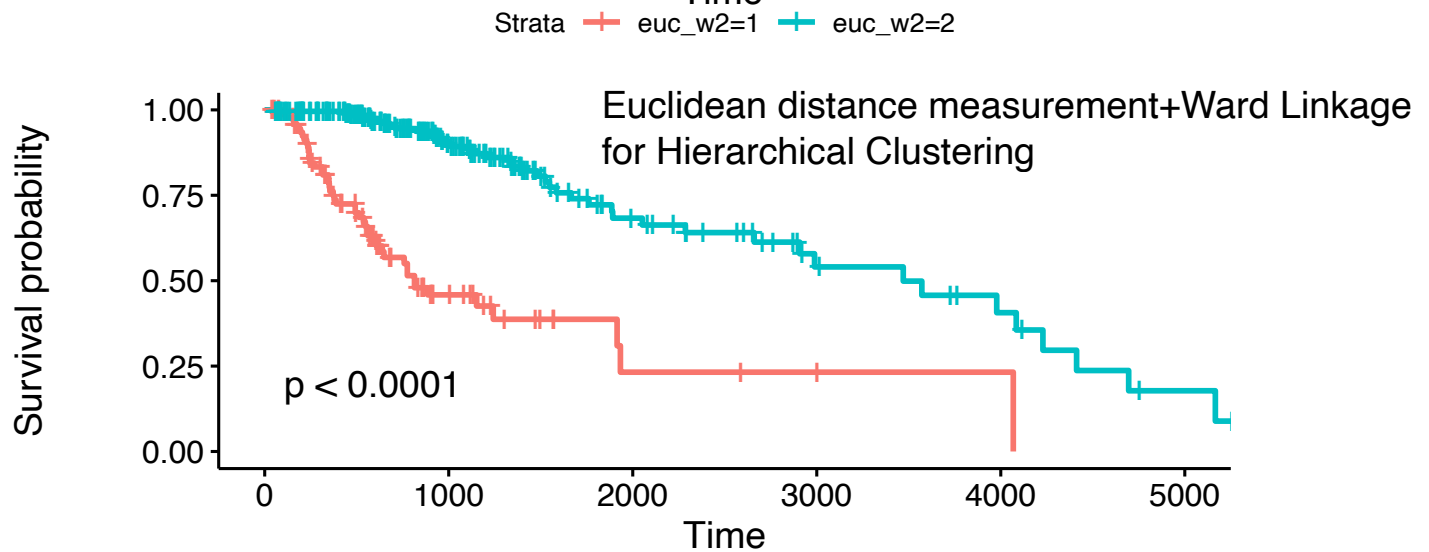
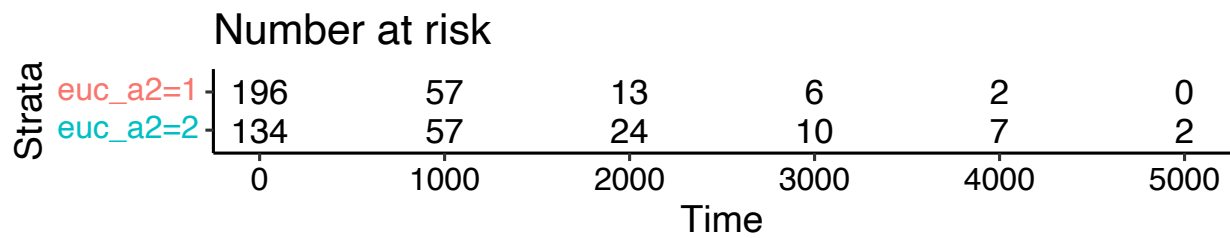
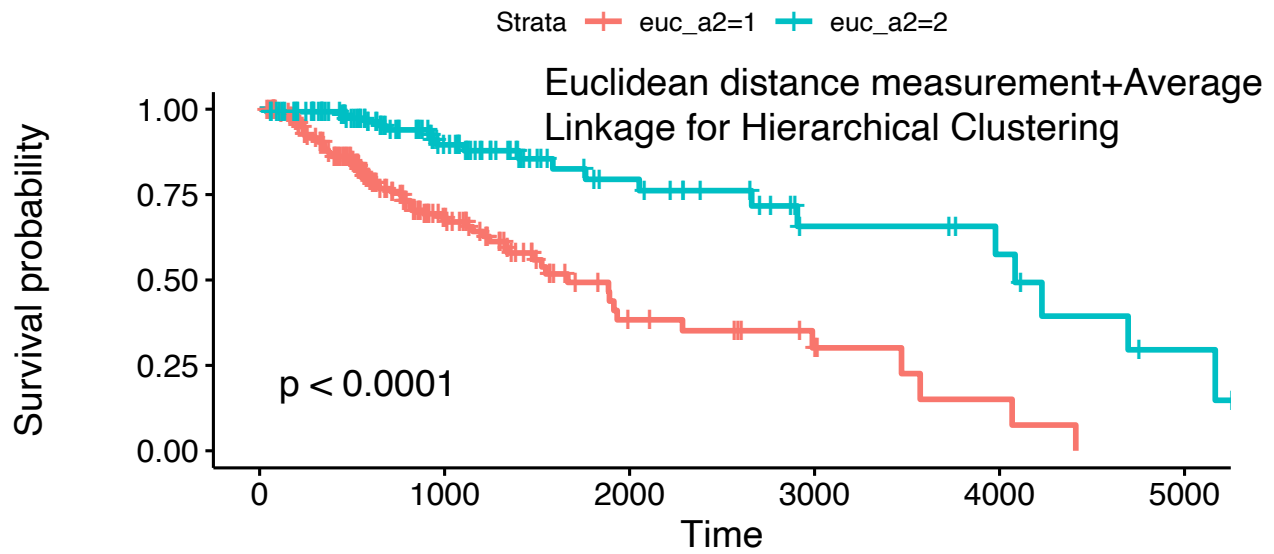
Kmeans of 2 groups

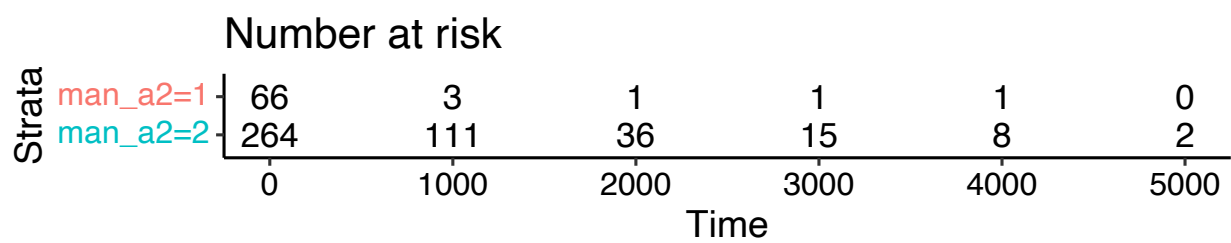
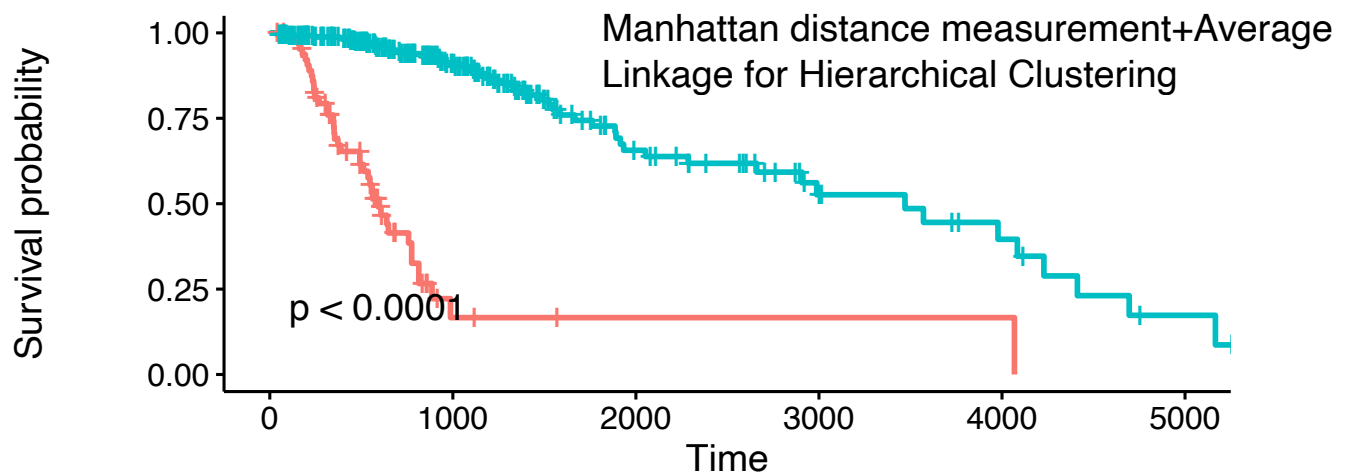
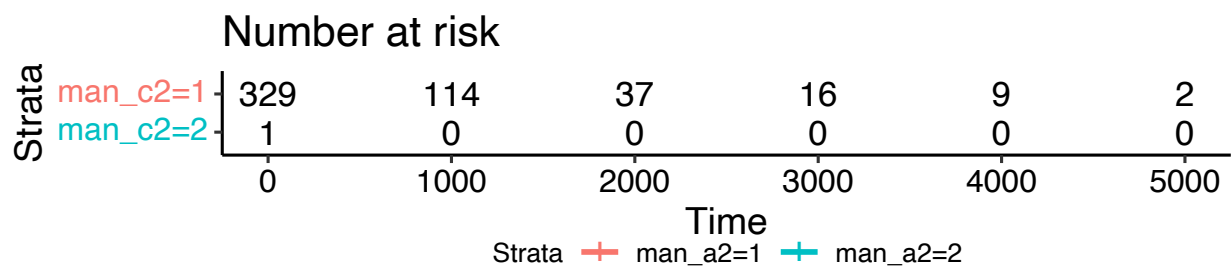
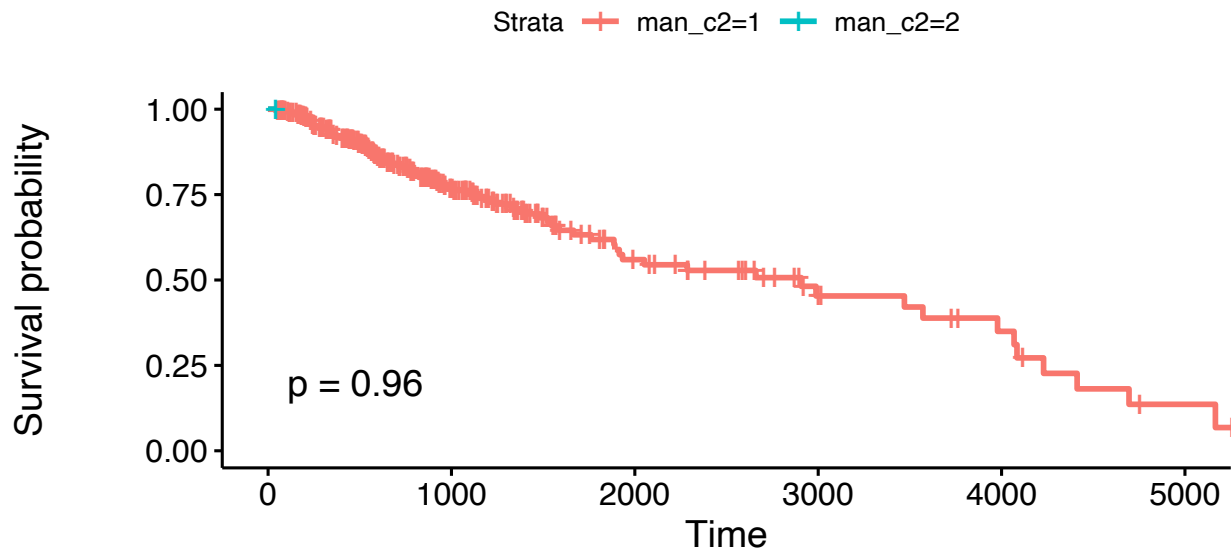


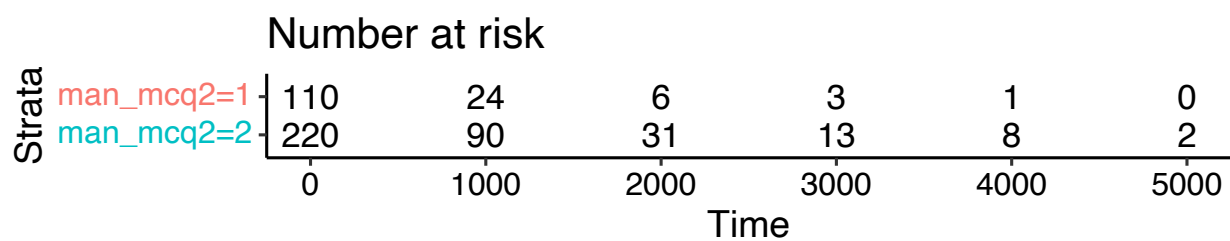
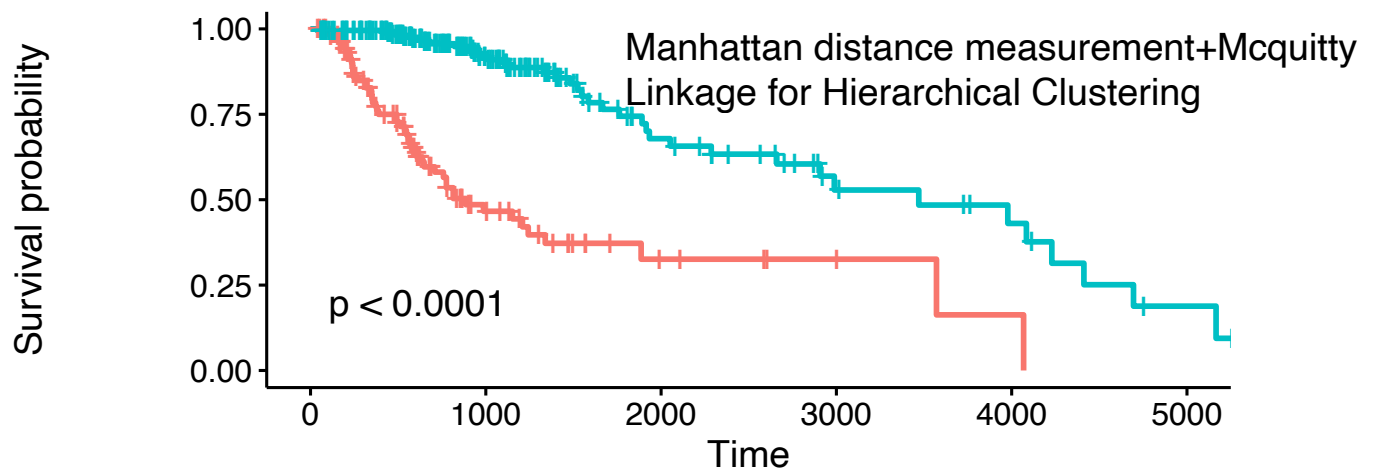
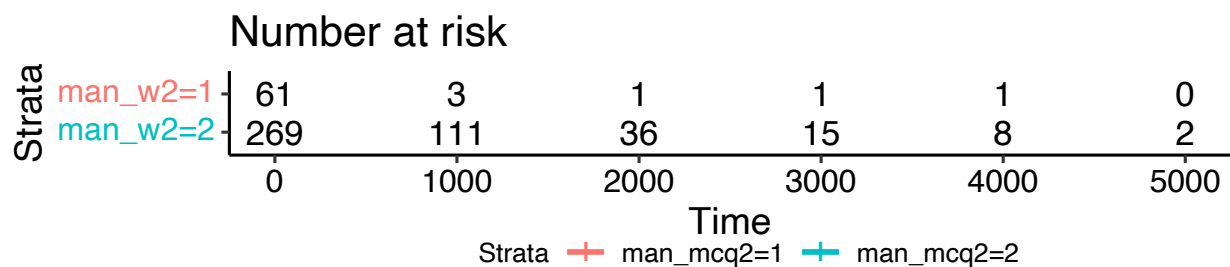
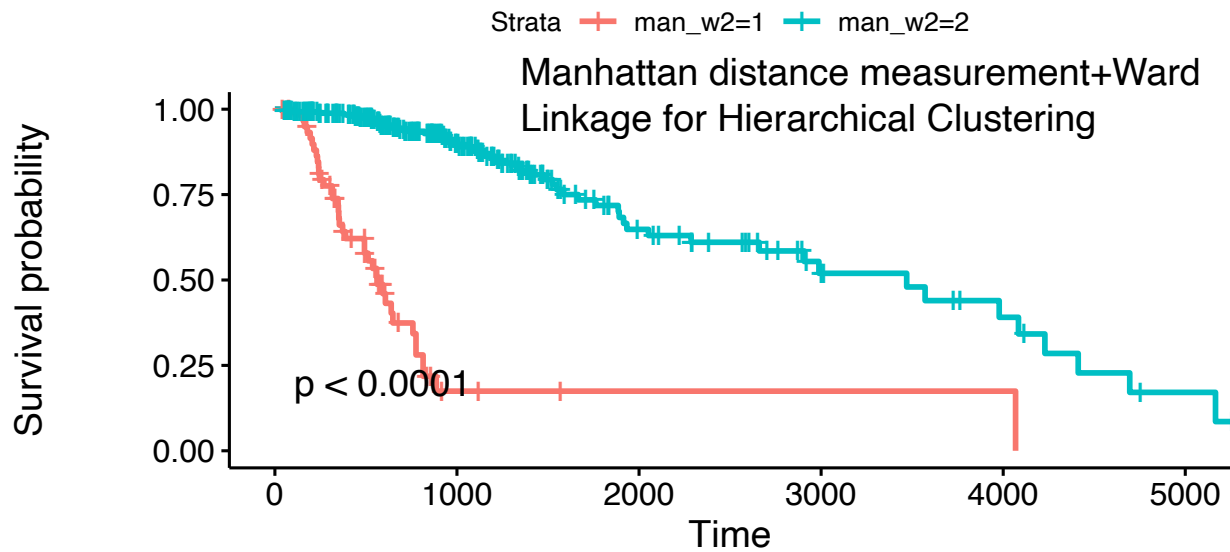
#### 4. Hierarchical Clustering

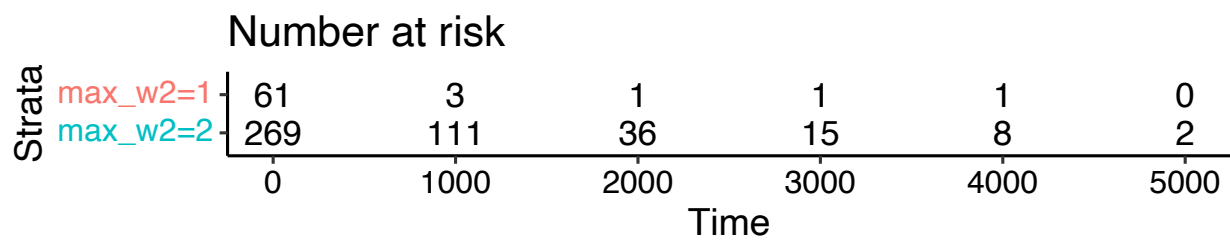
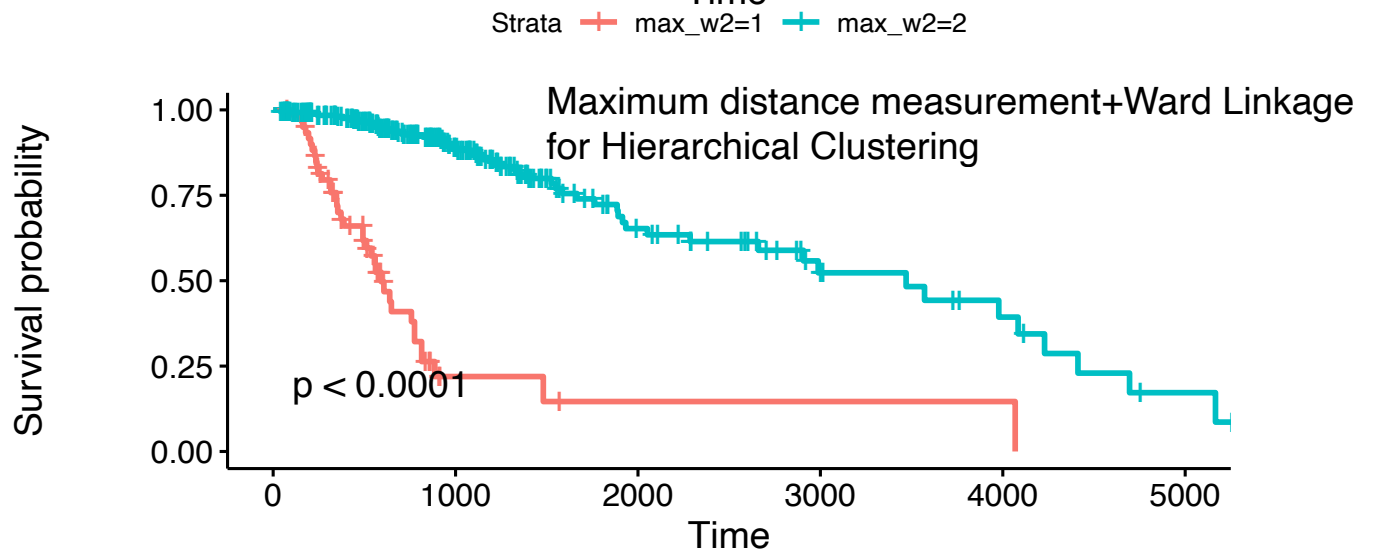
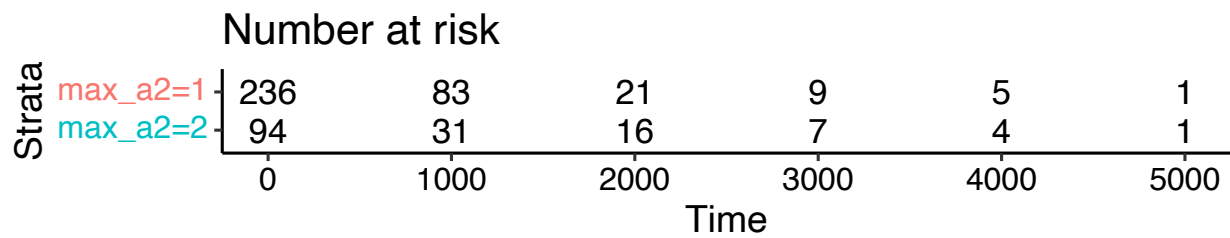
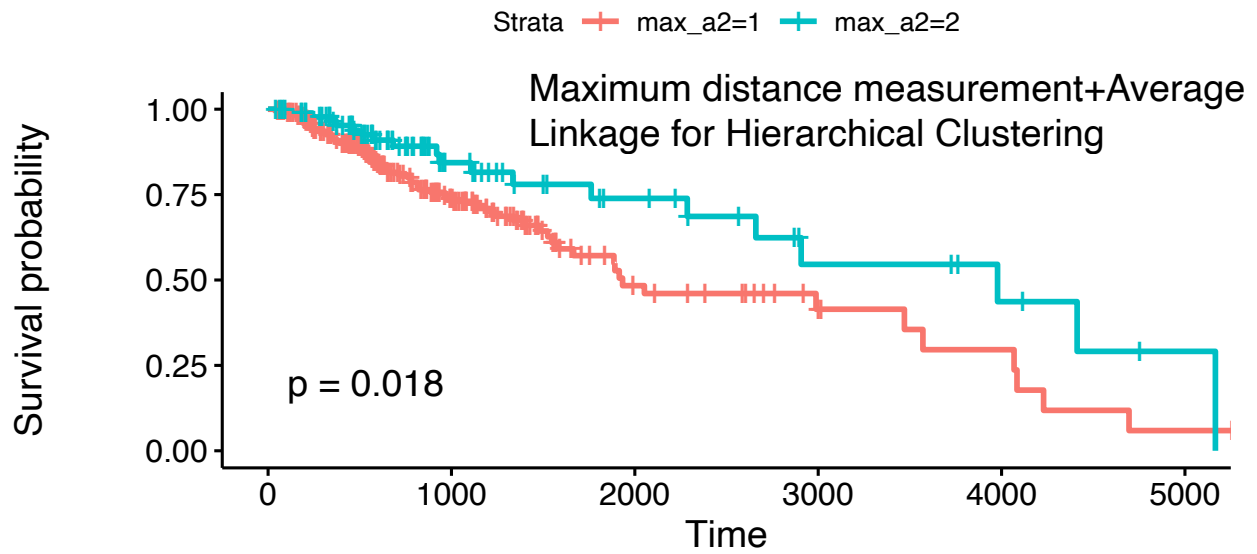
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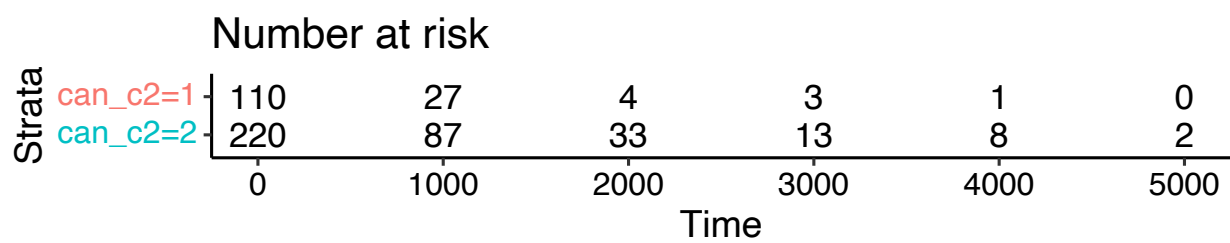
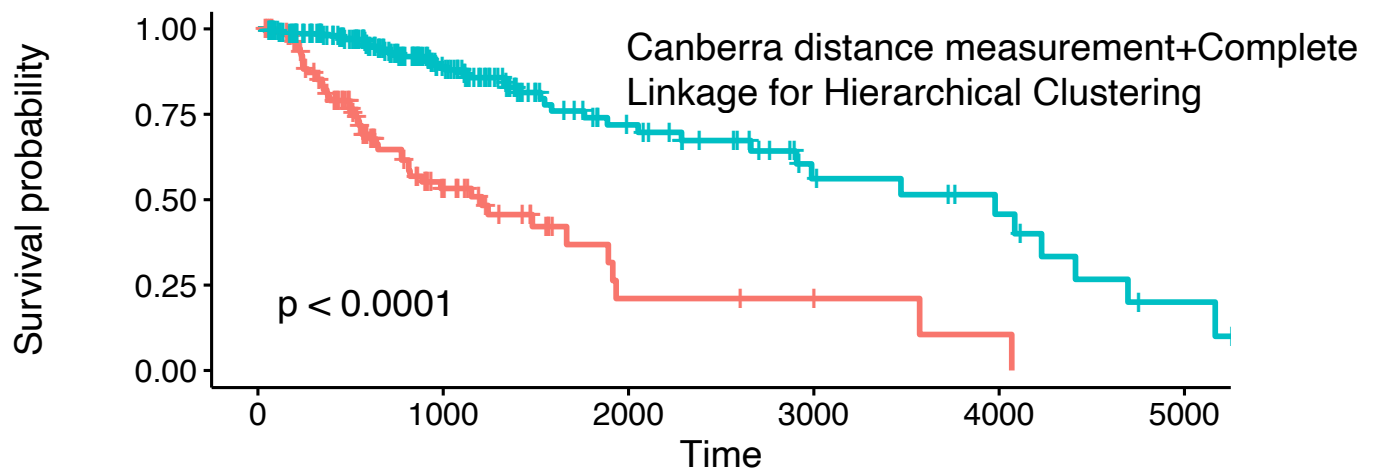
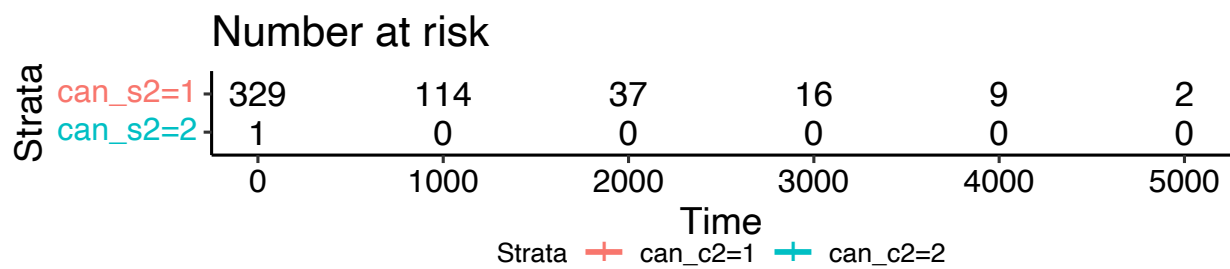
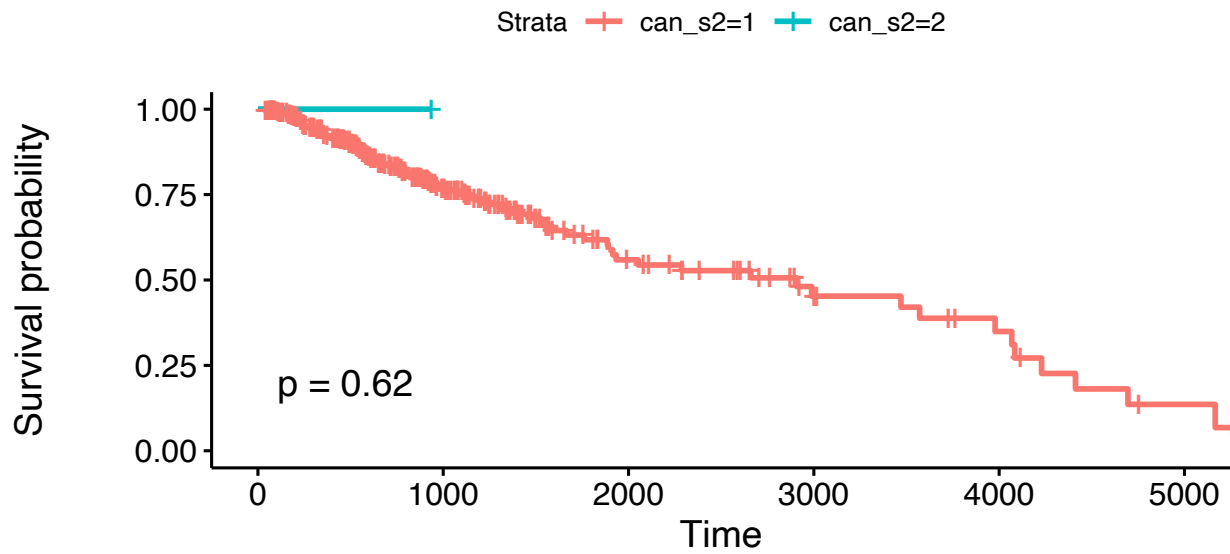




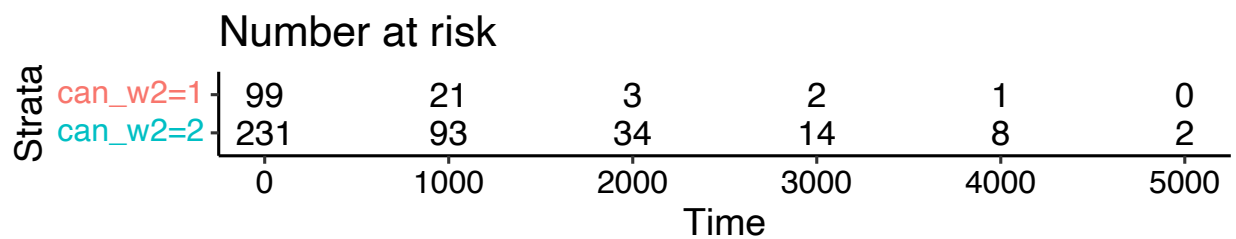
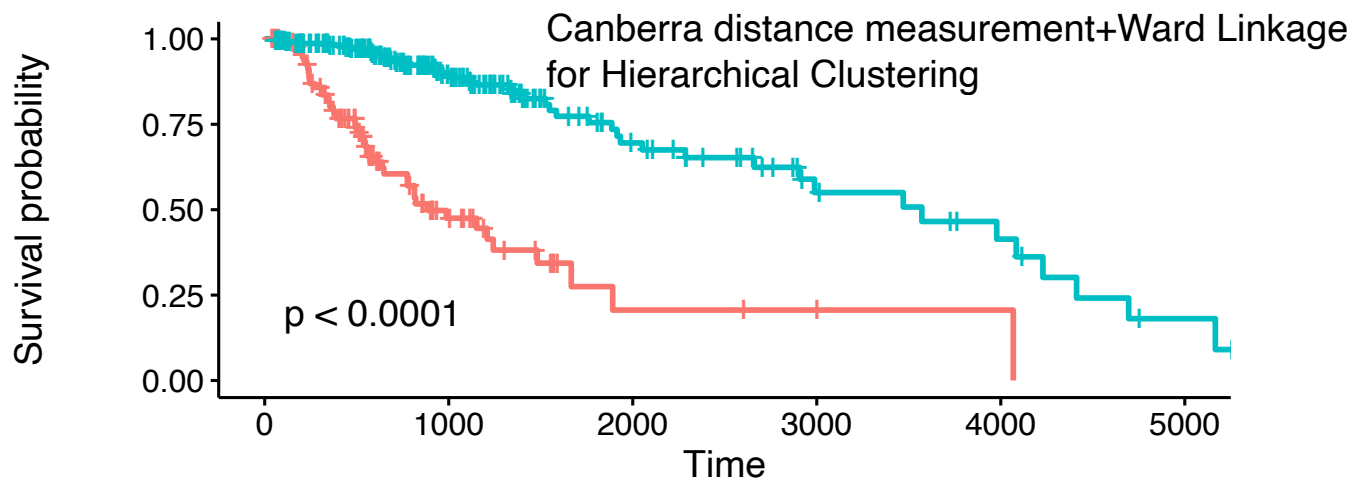
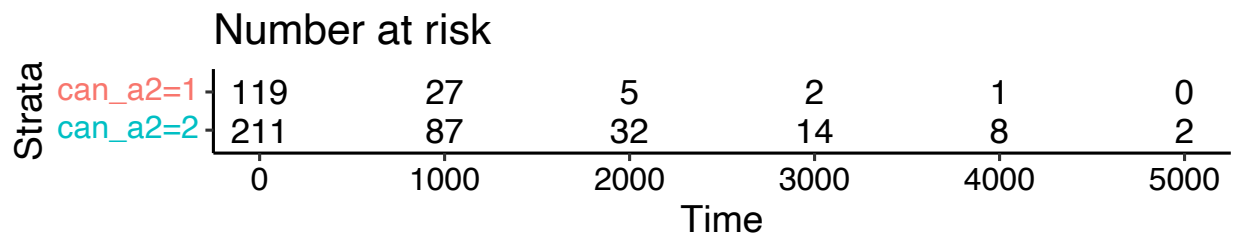
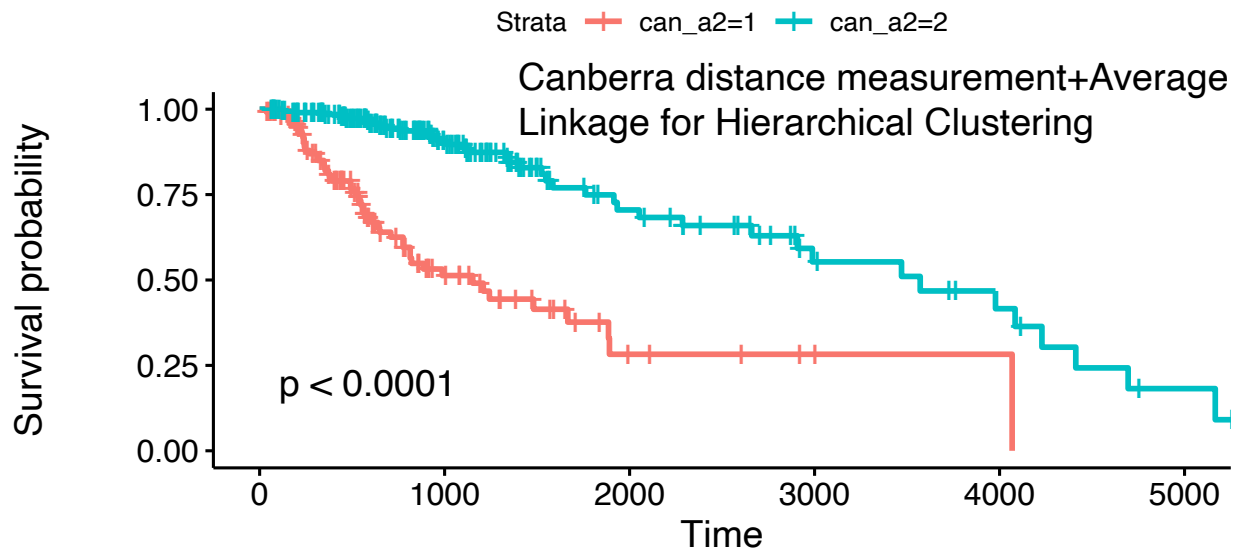


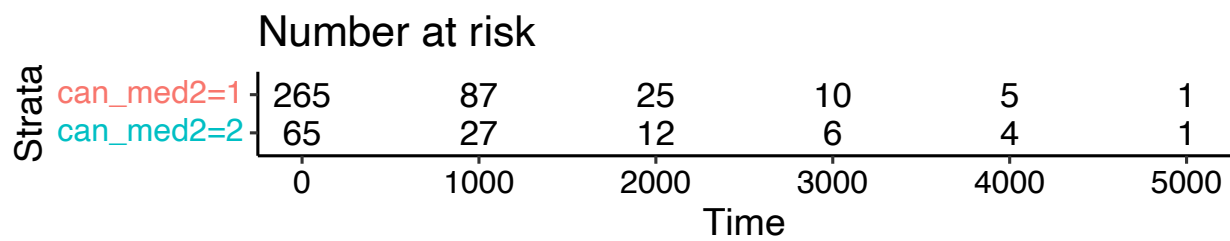
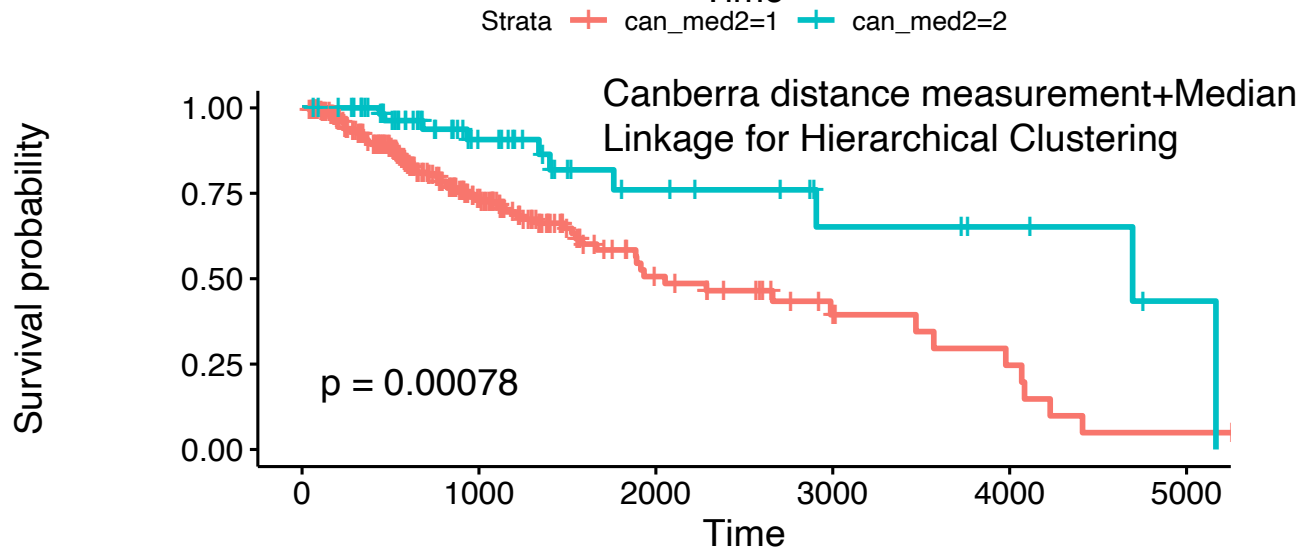
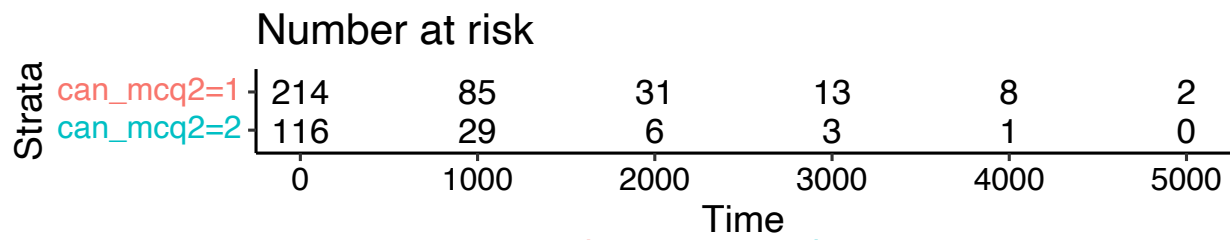
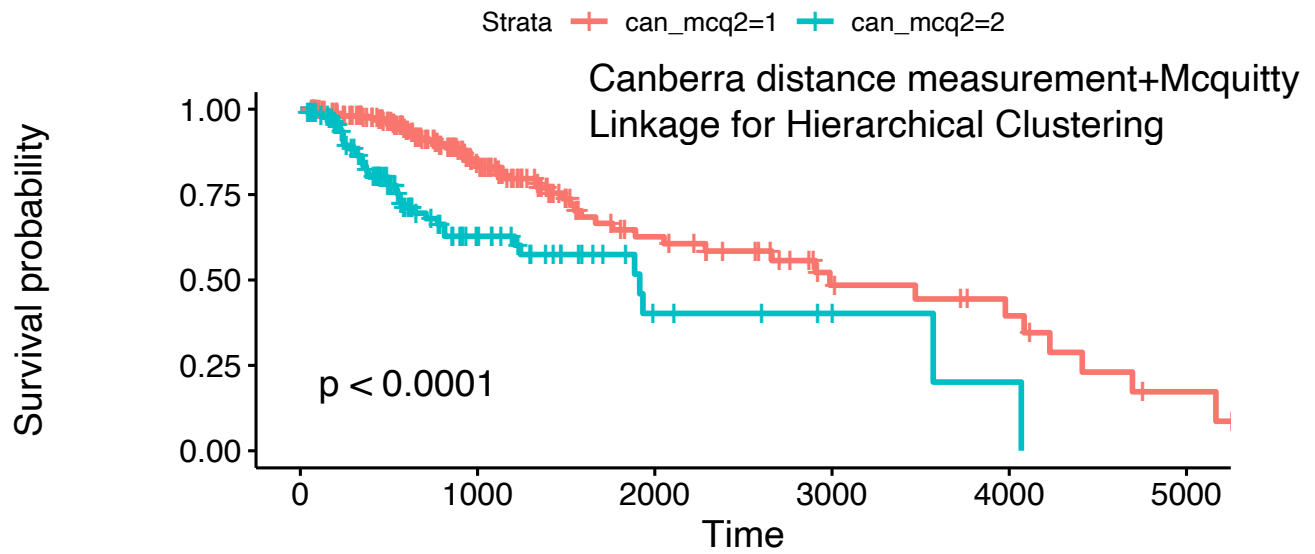


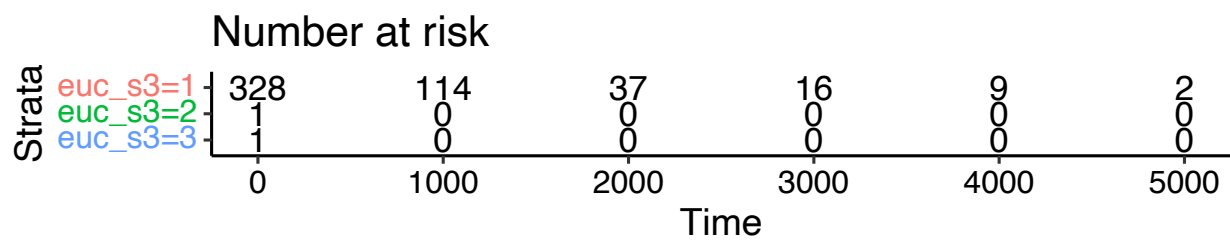
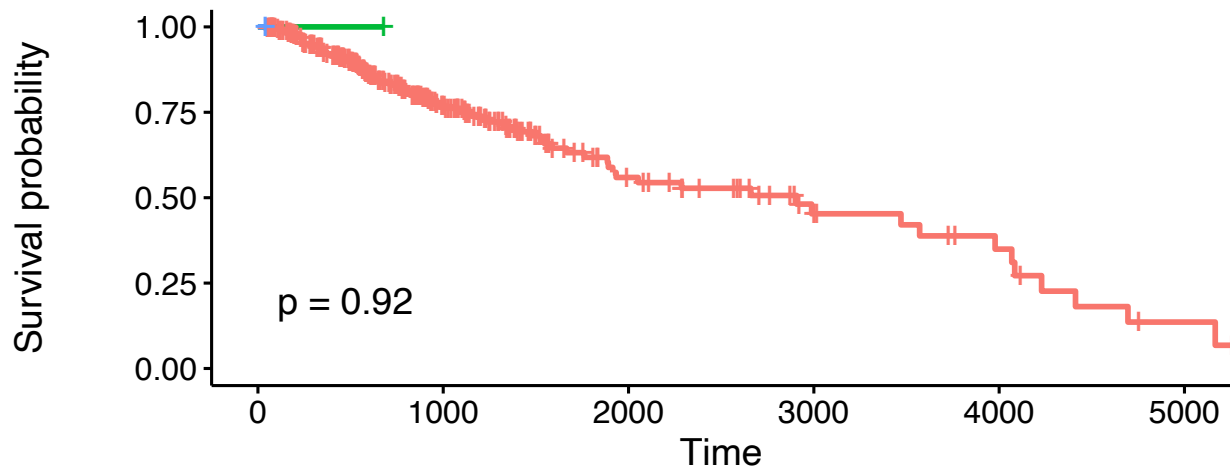
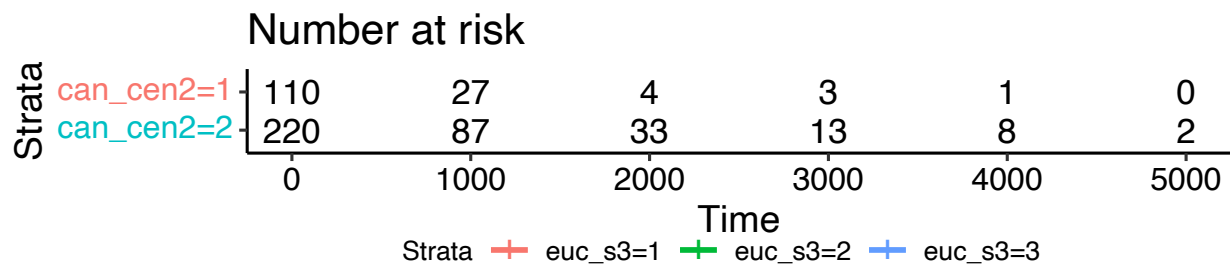
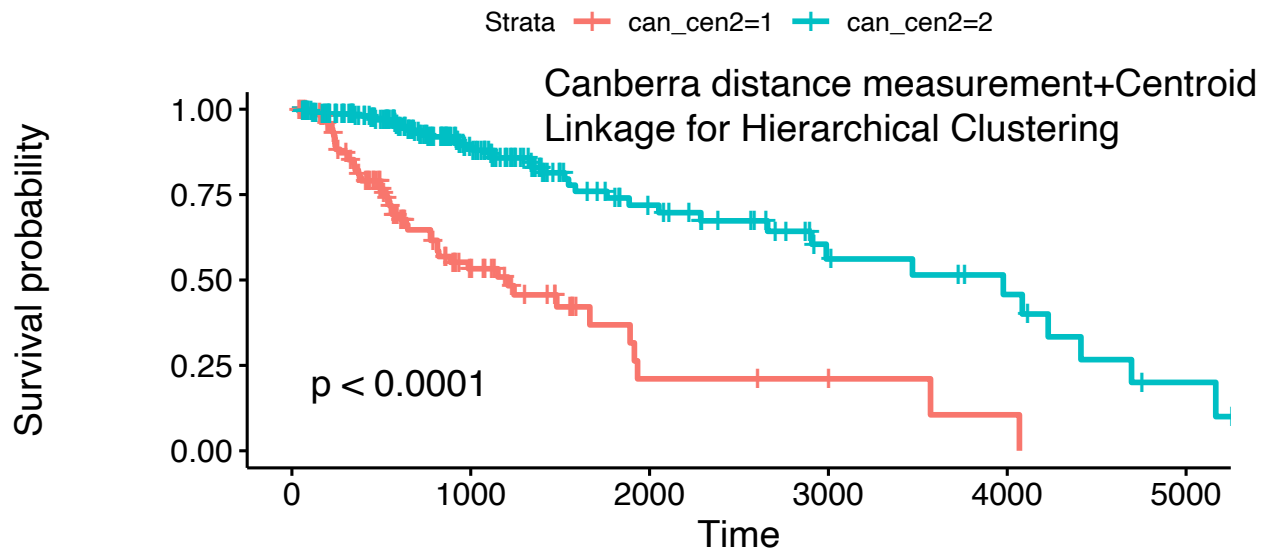


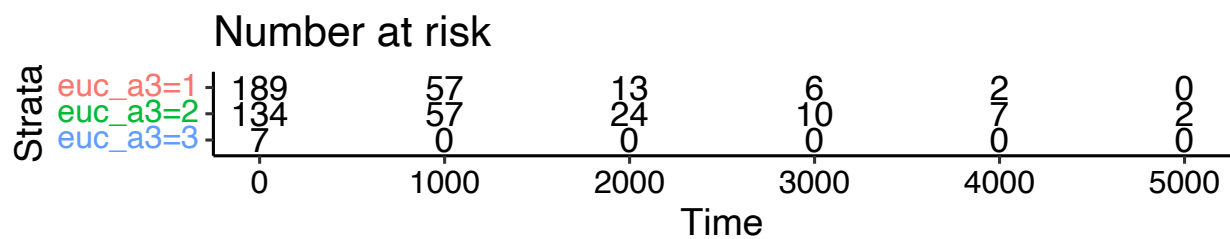
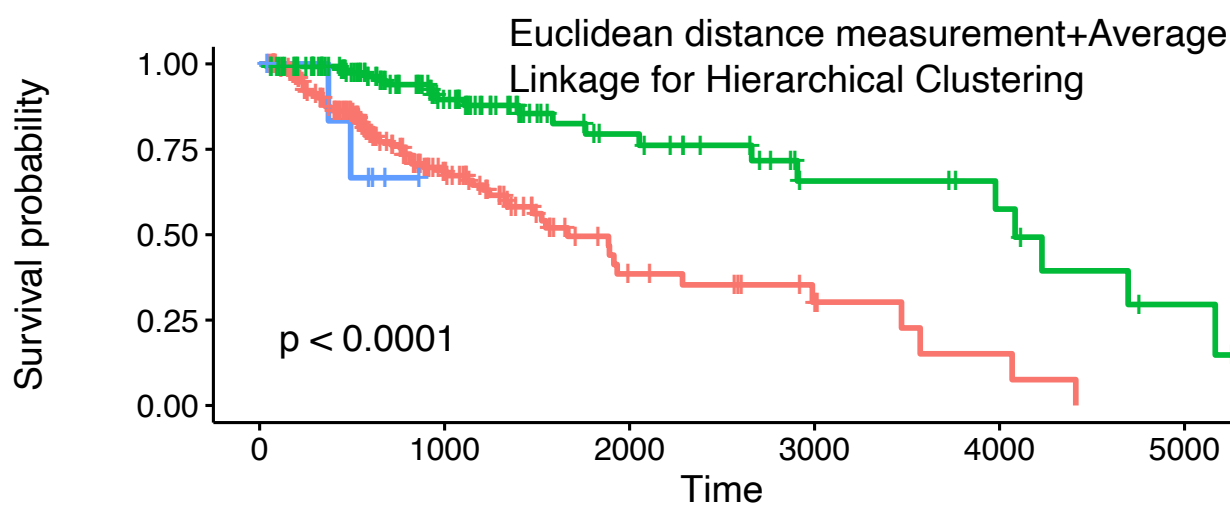
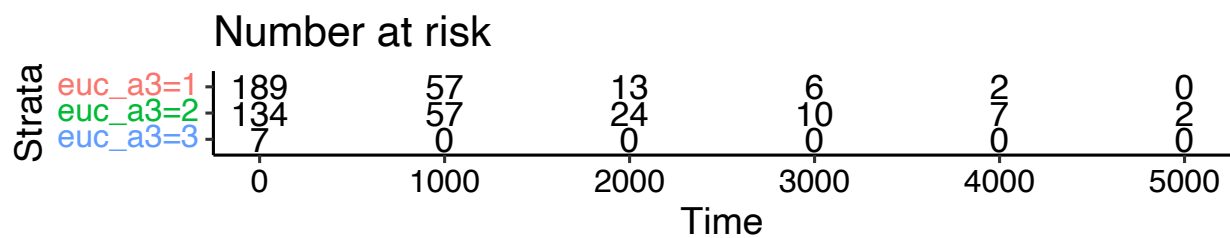
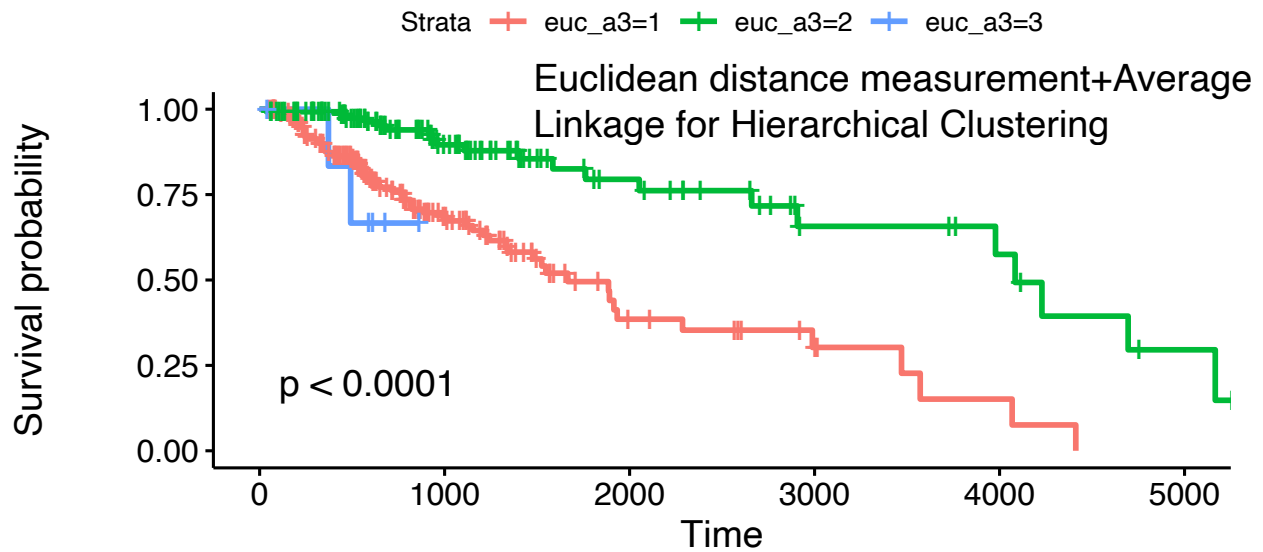


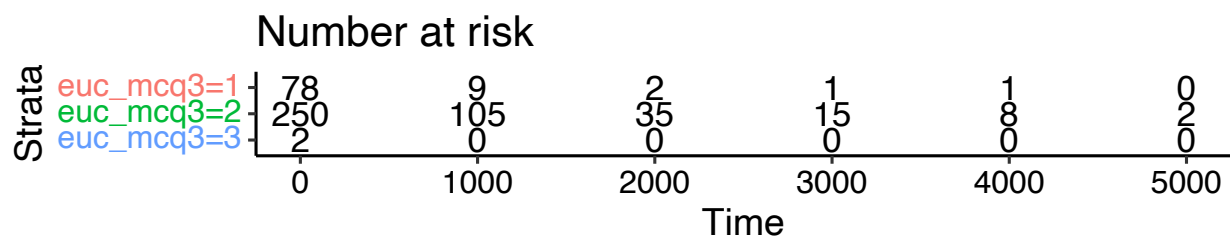
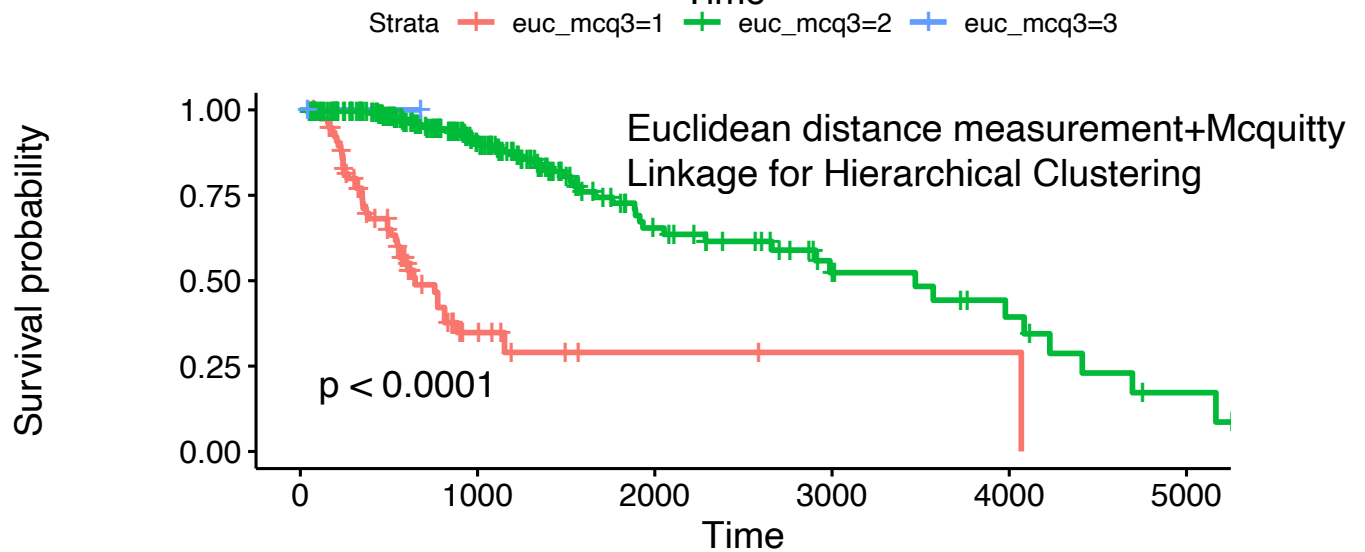
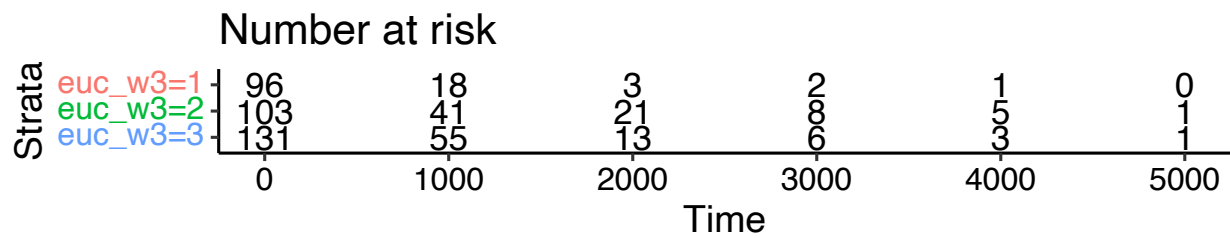
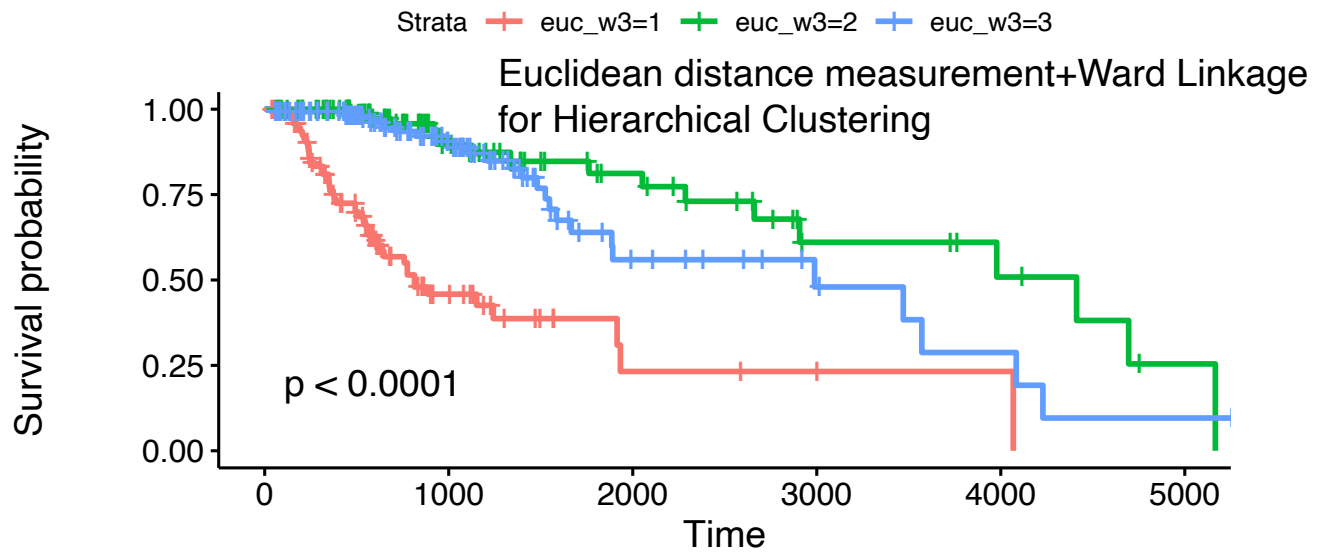


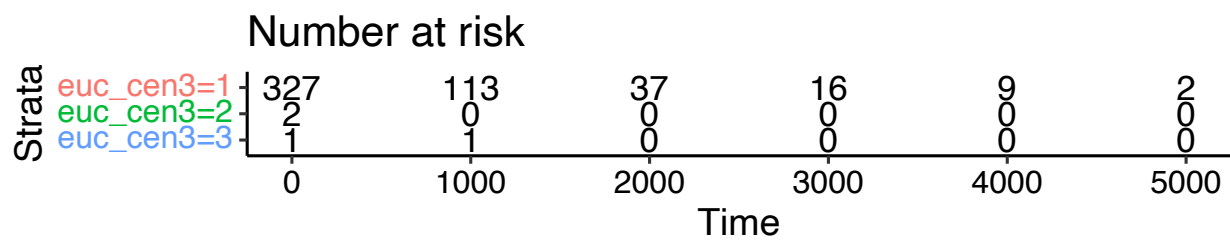
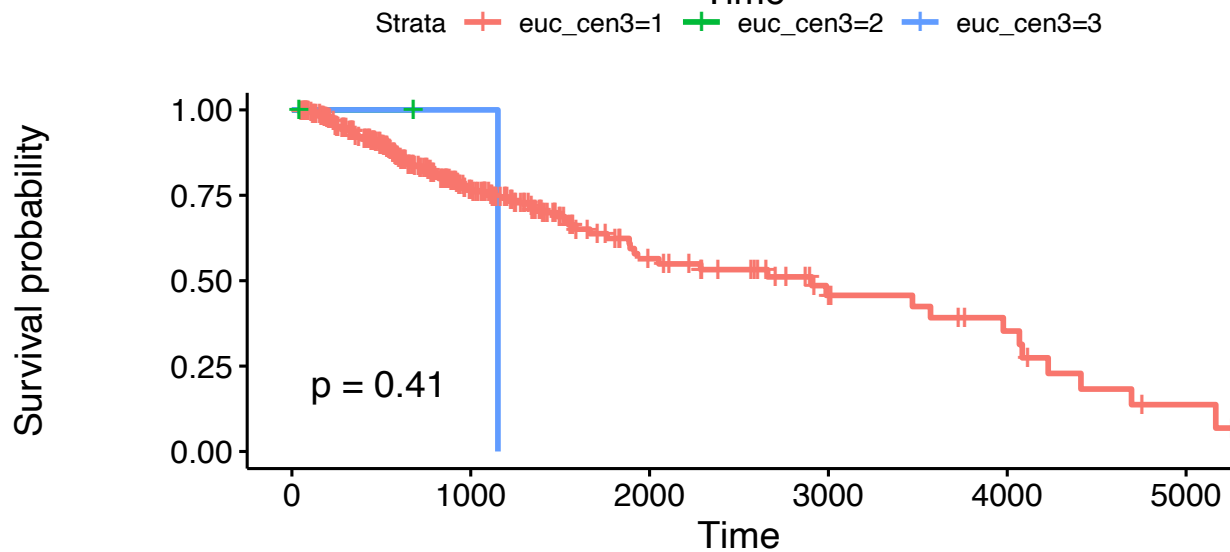
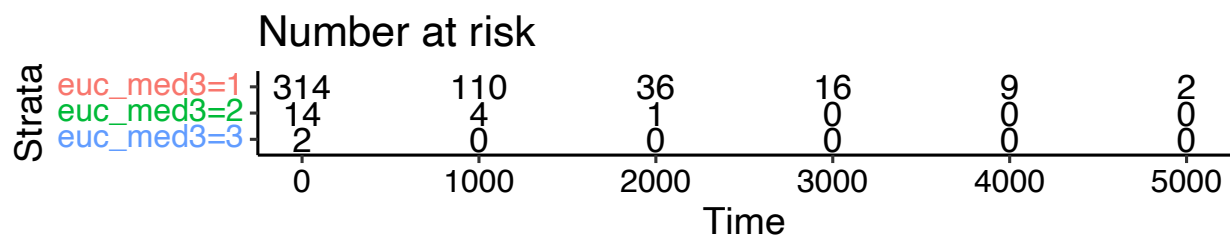
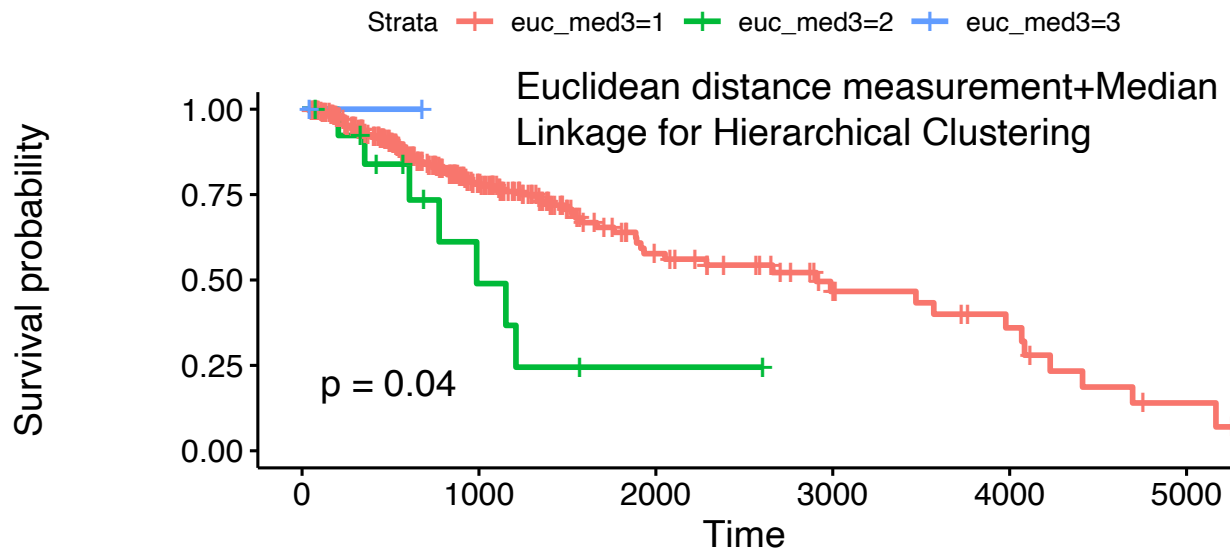


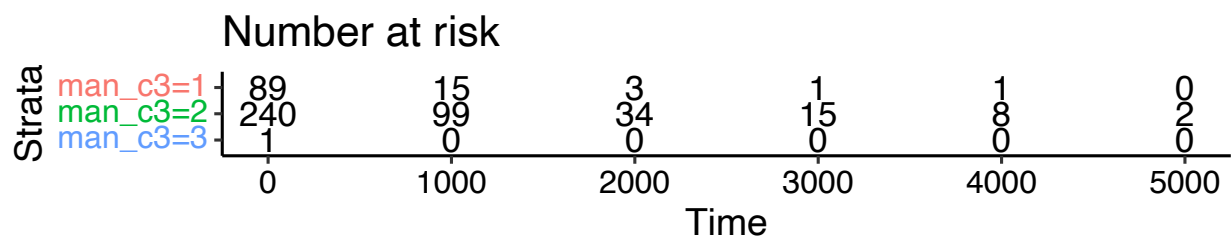
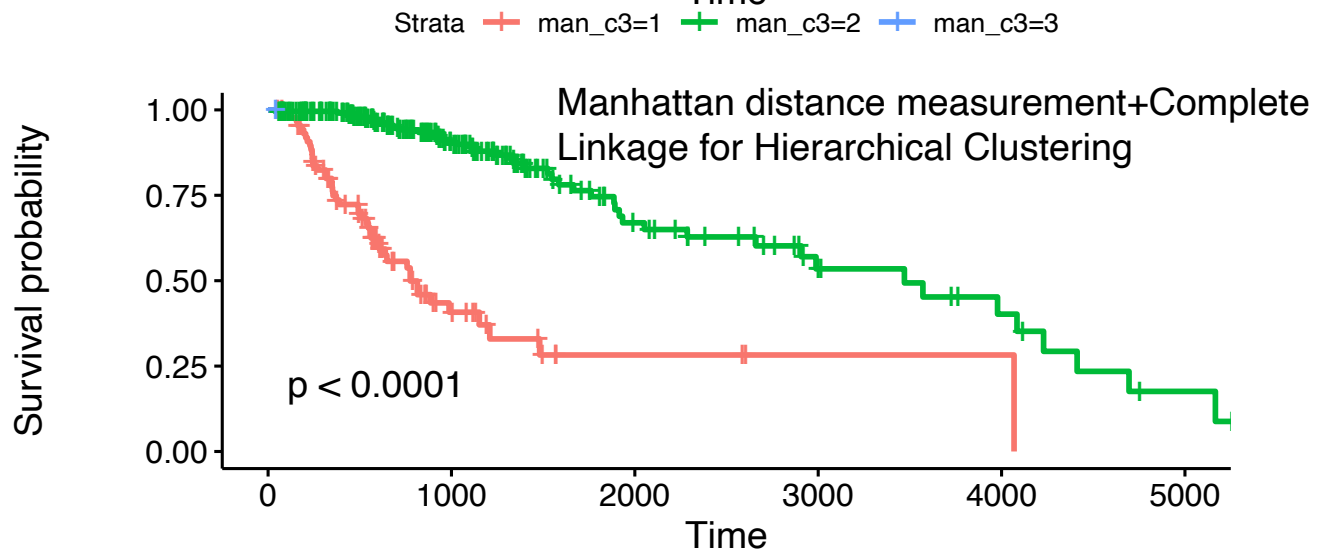
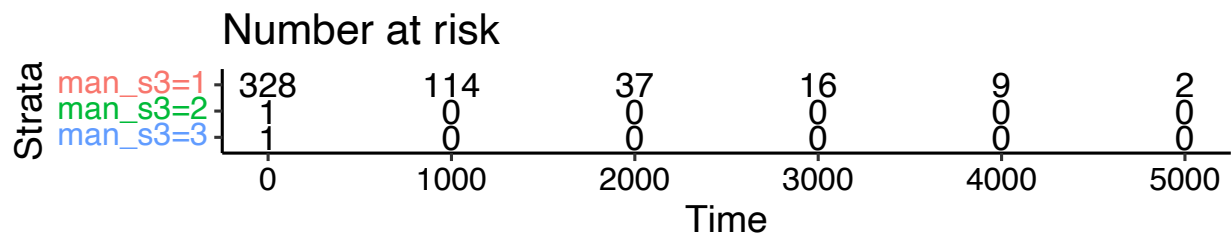
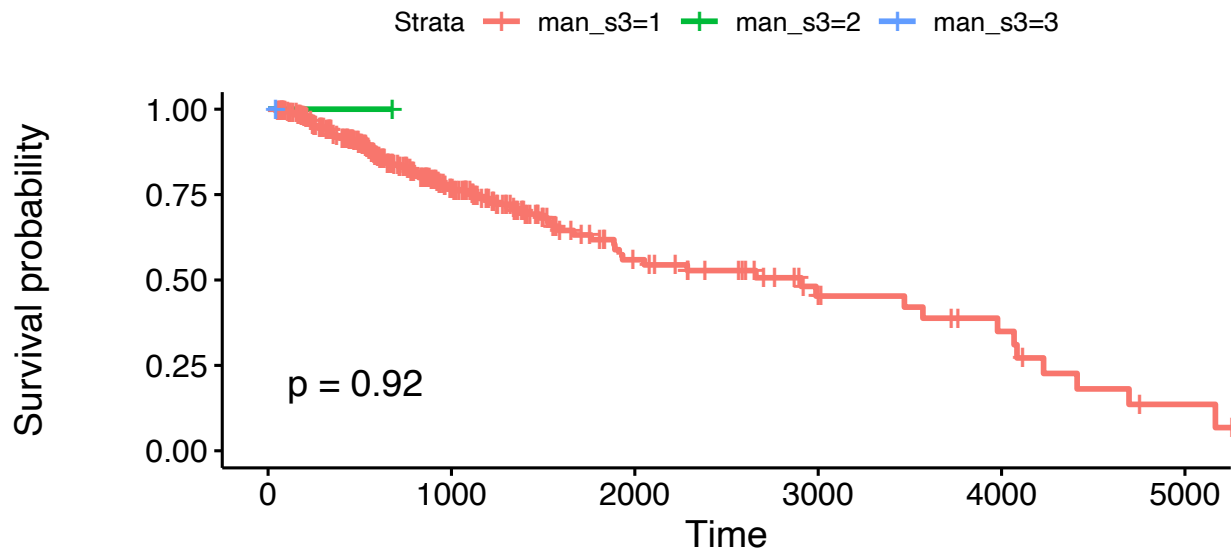


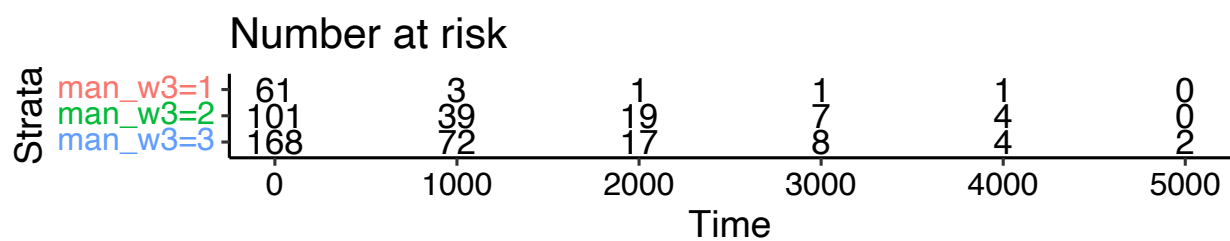
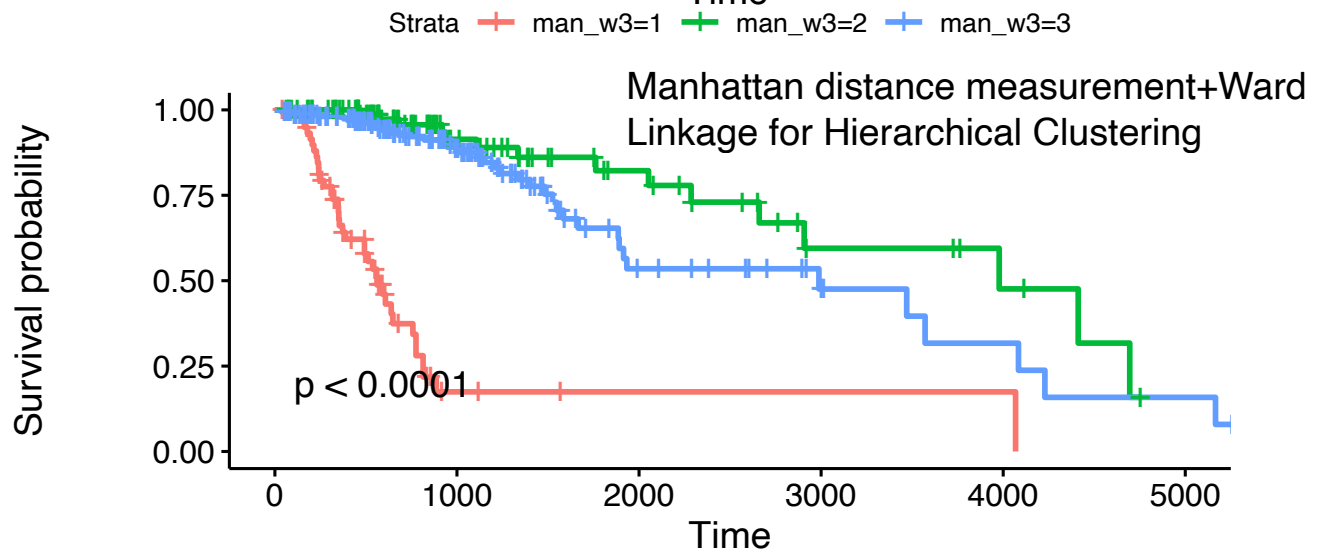
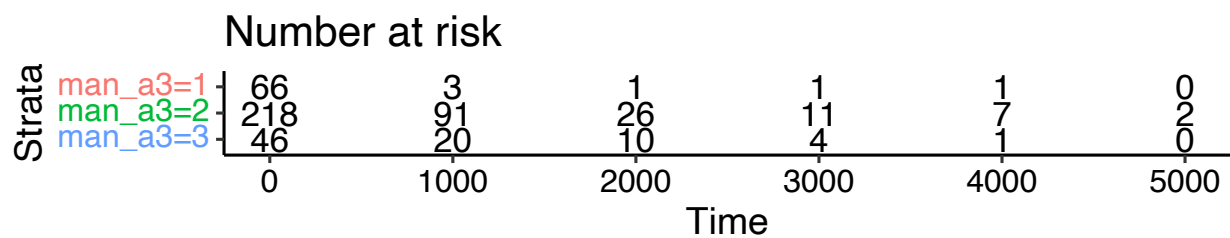
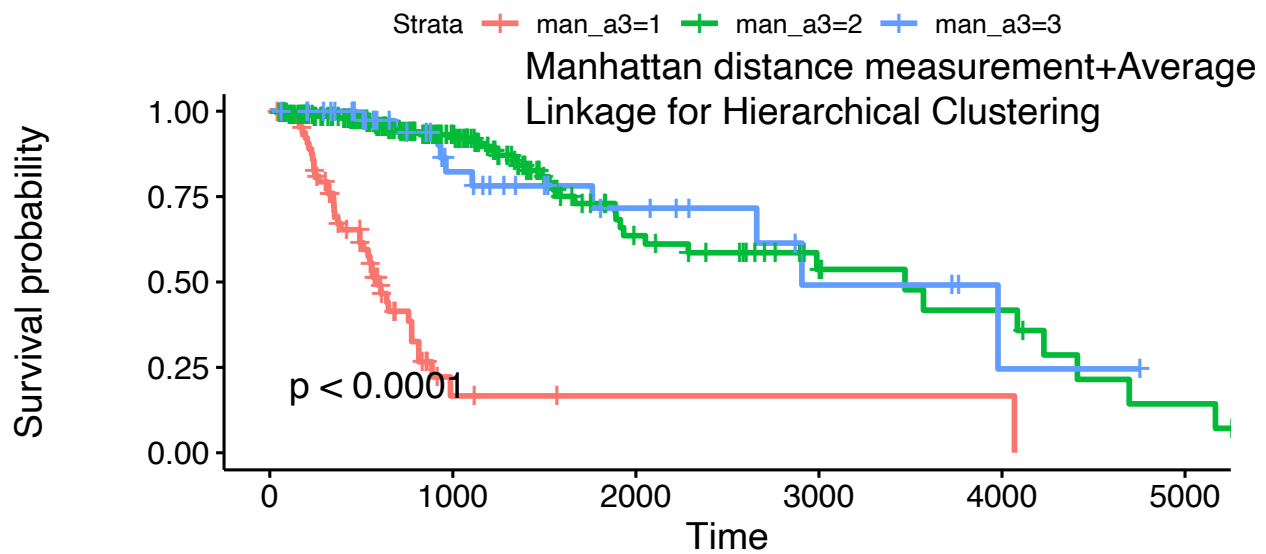




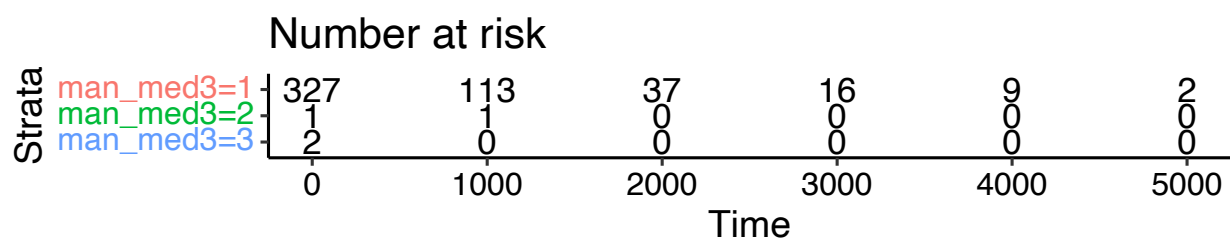
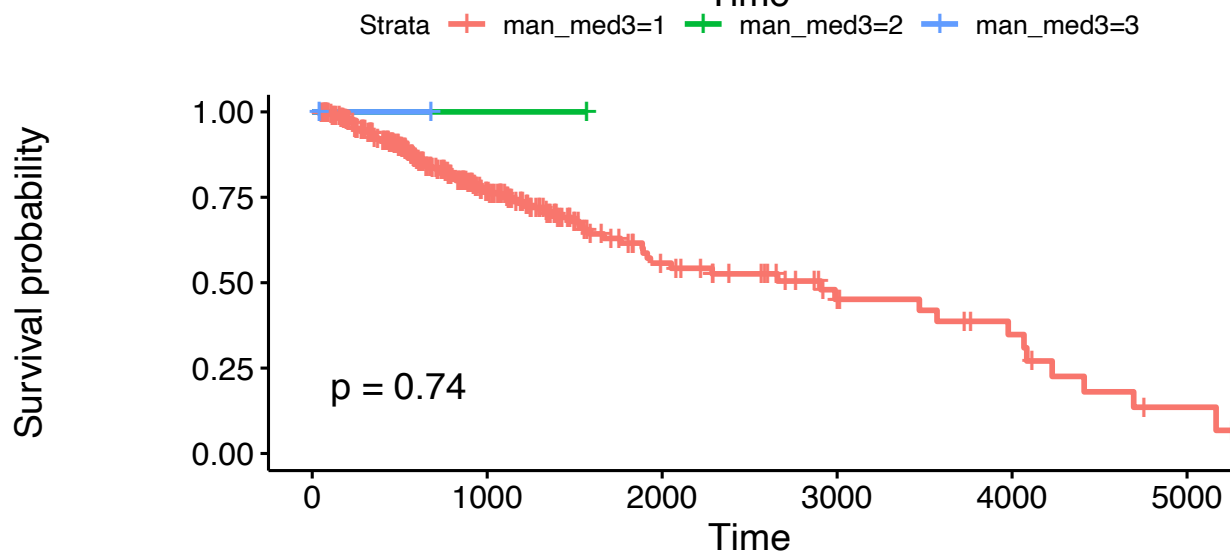
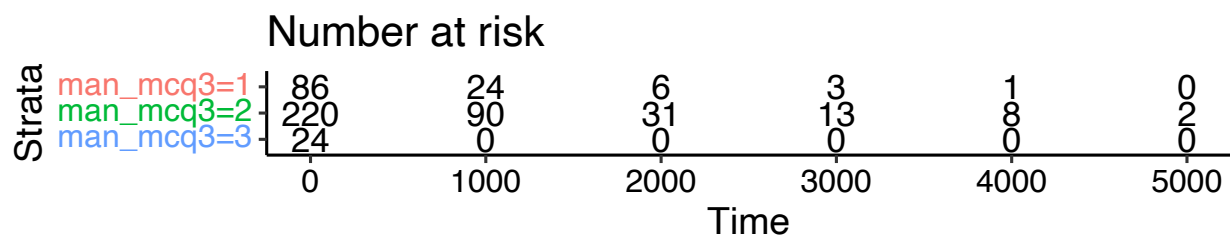
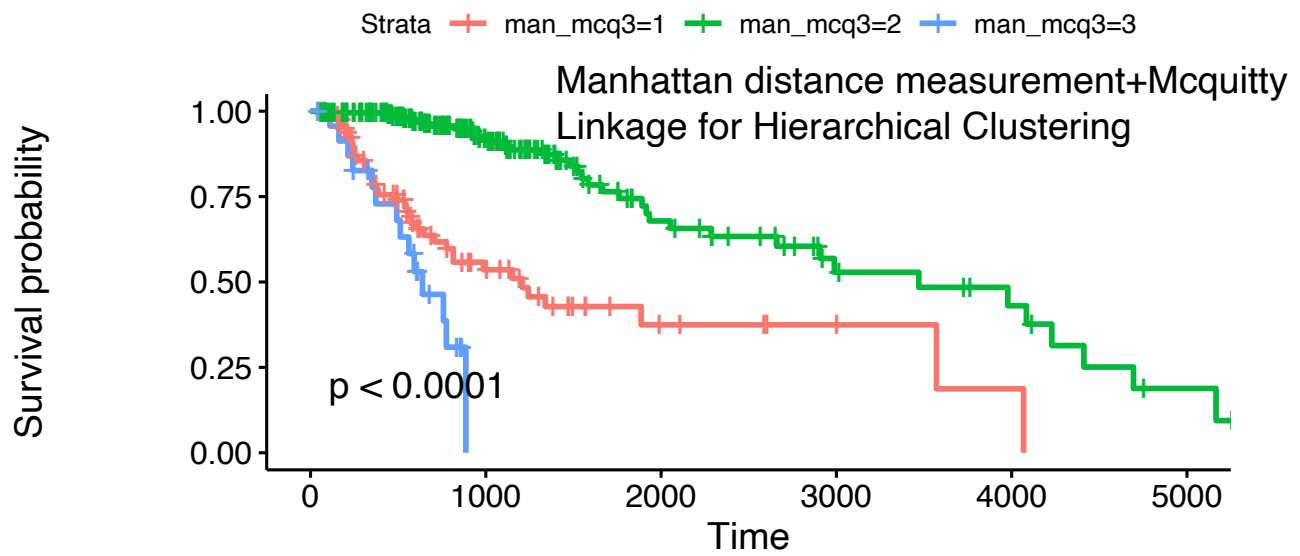


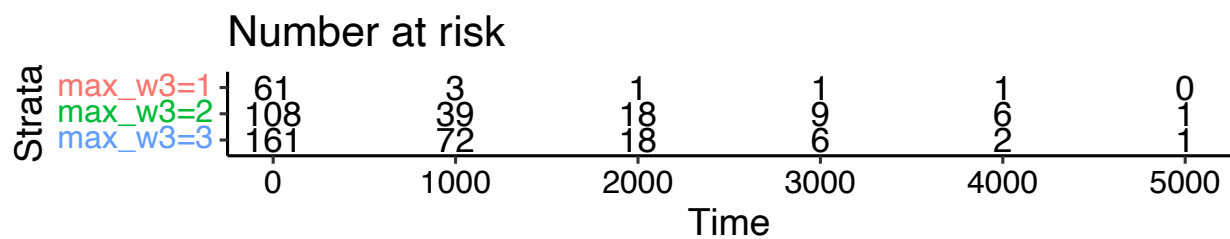
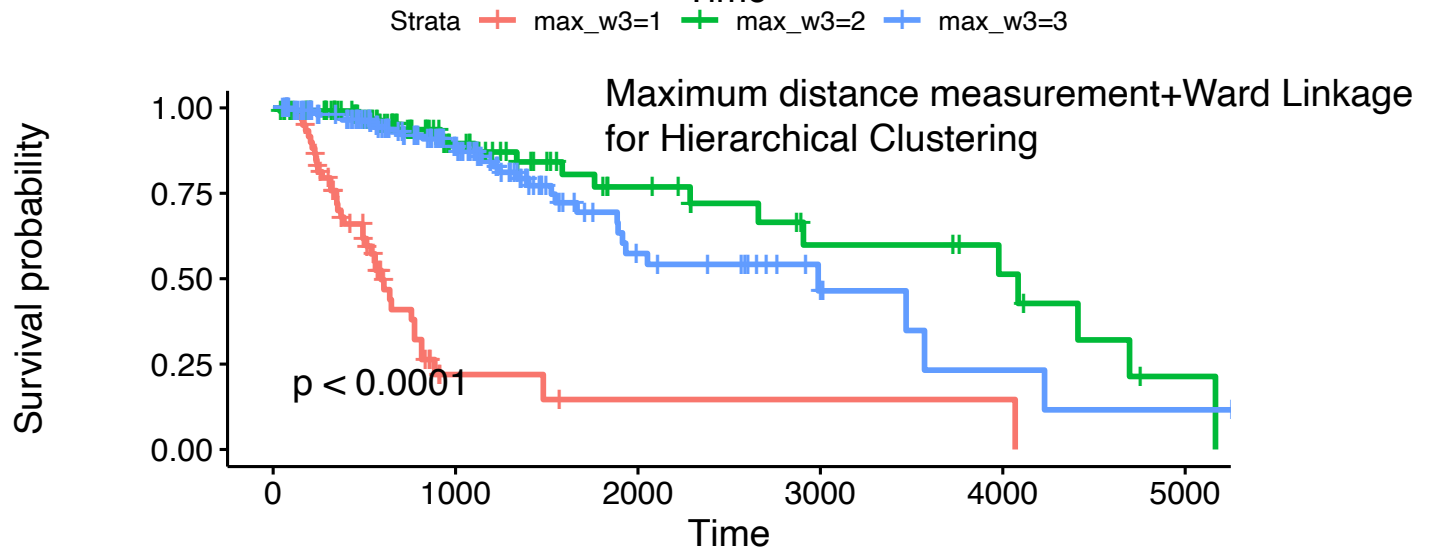
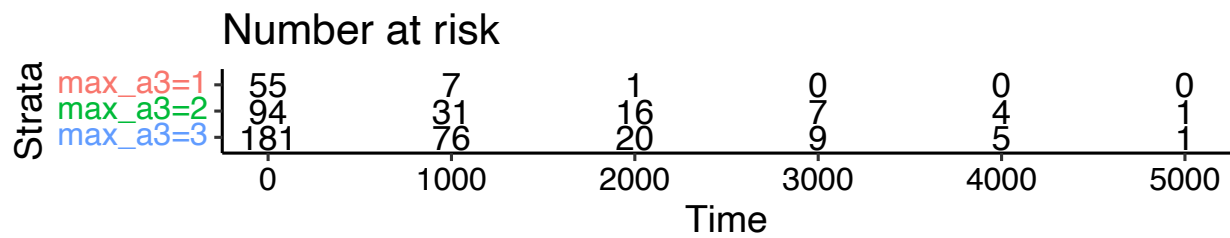
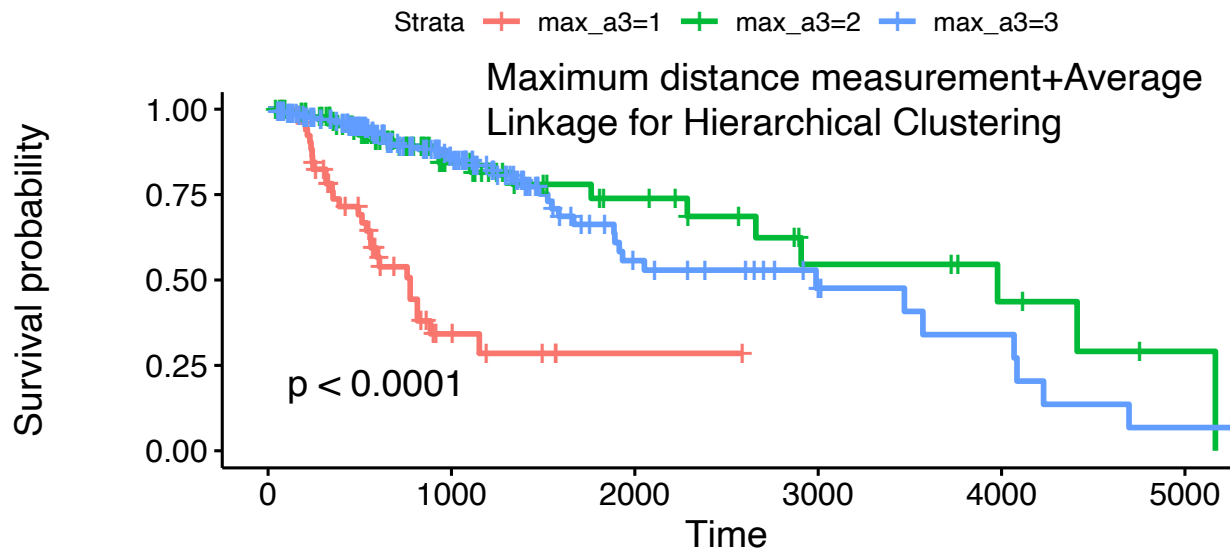


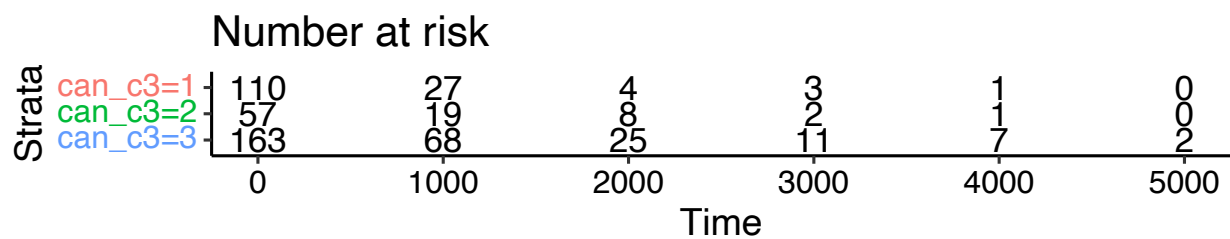
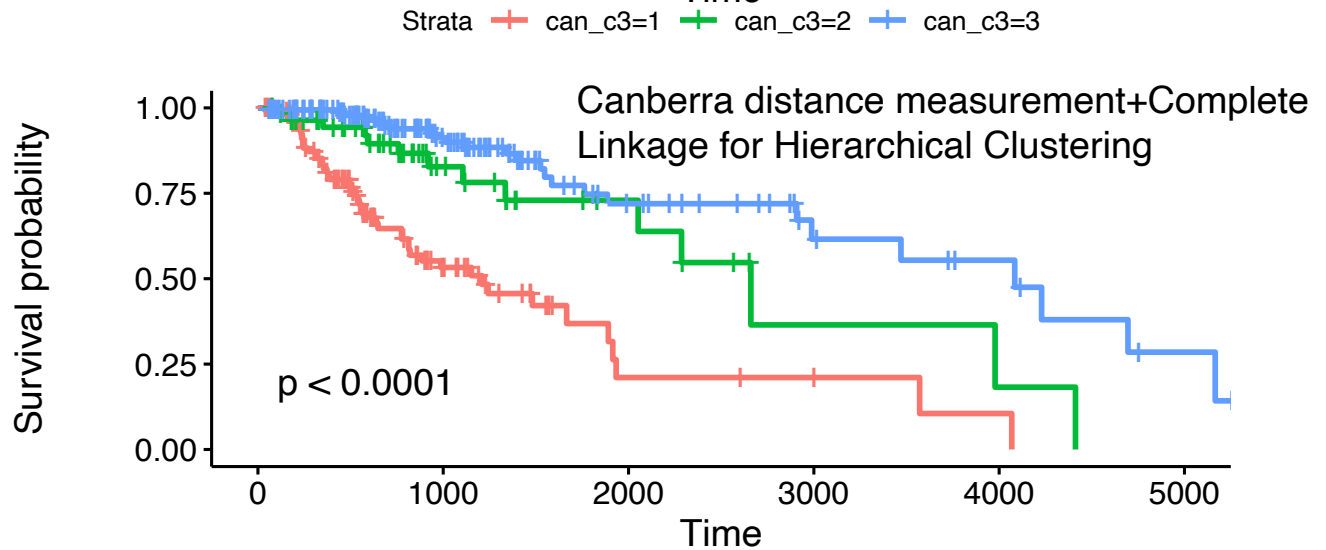
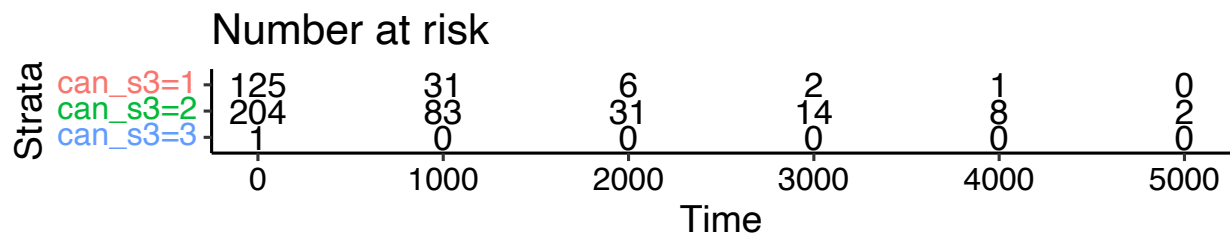
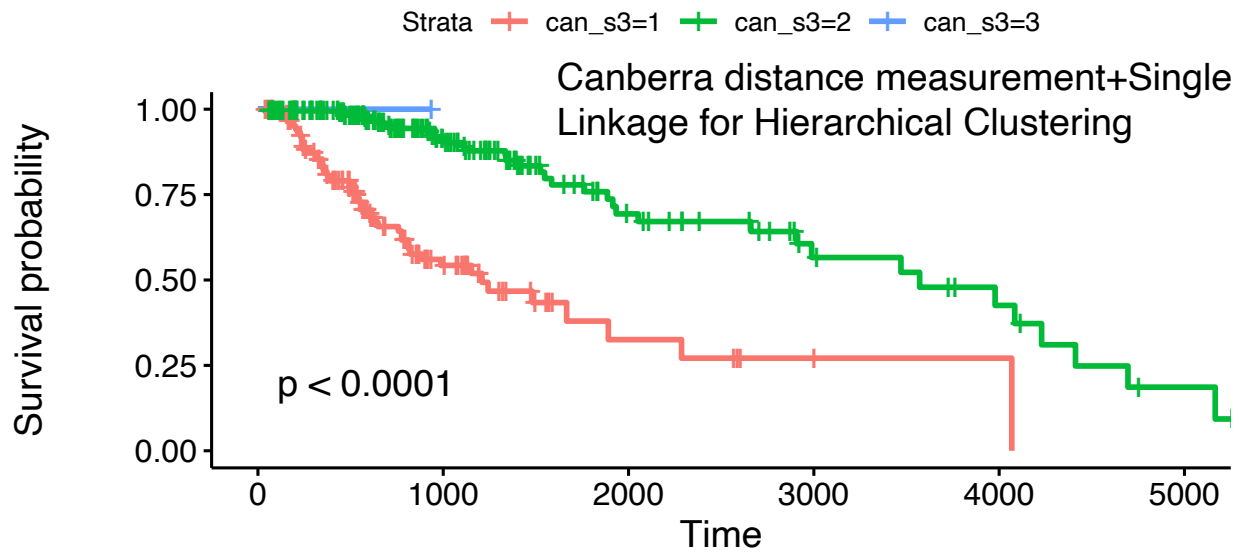


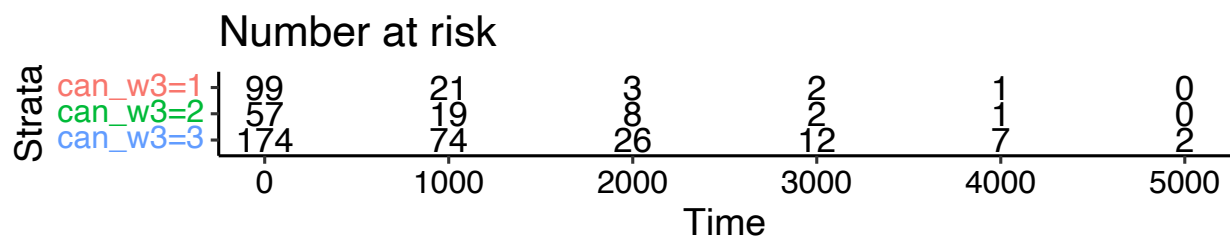
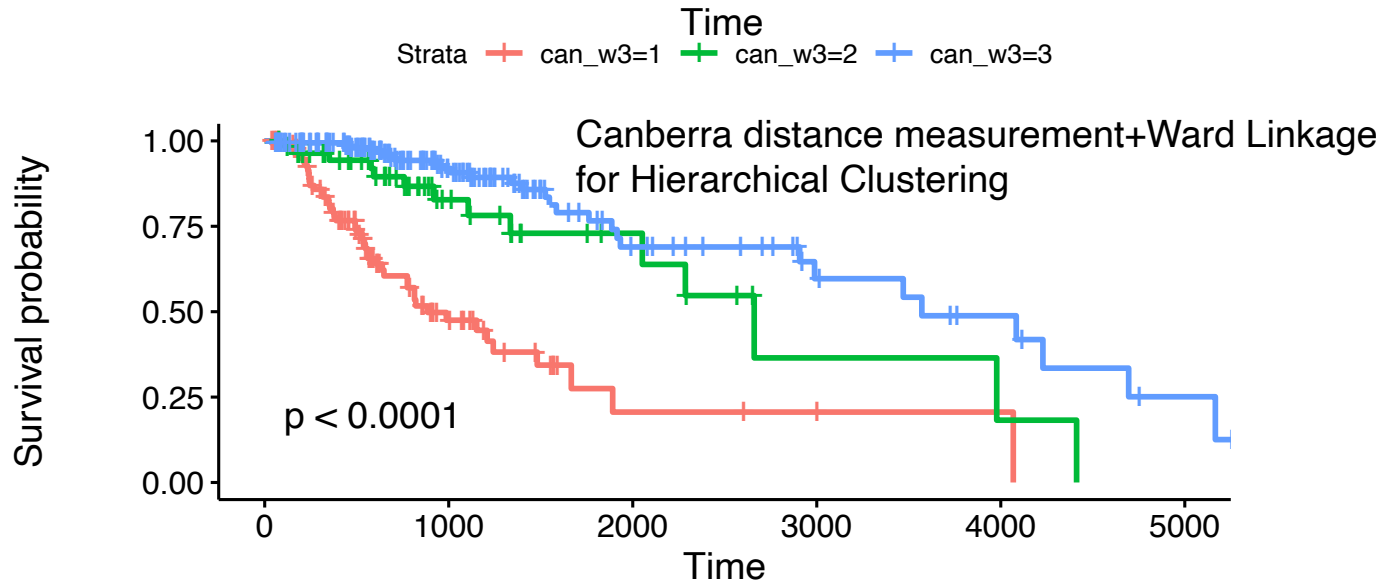
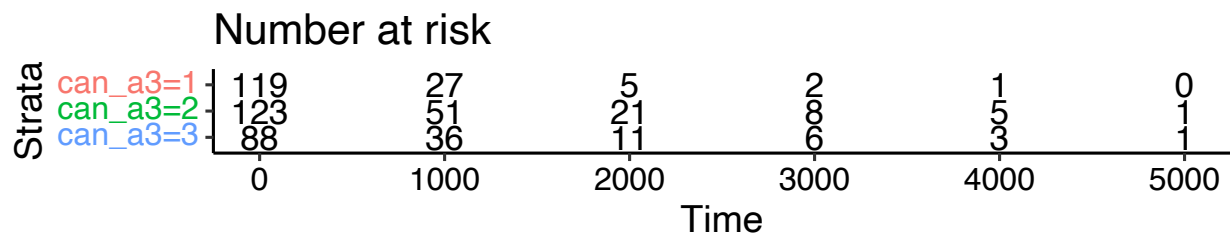
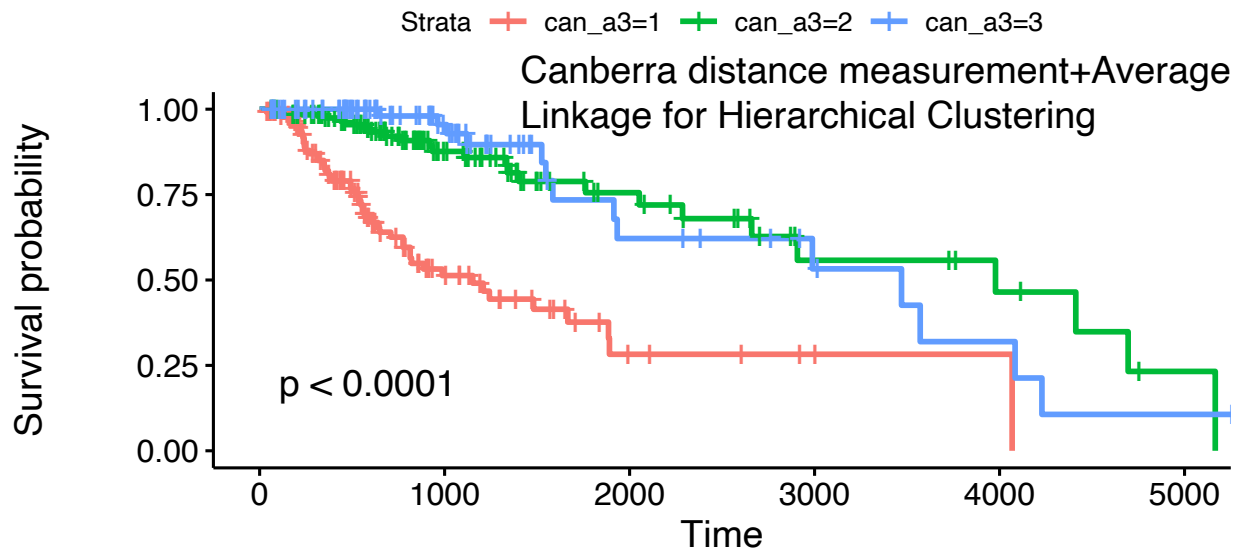


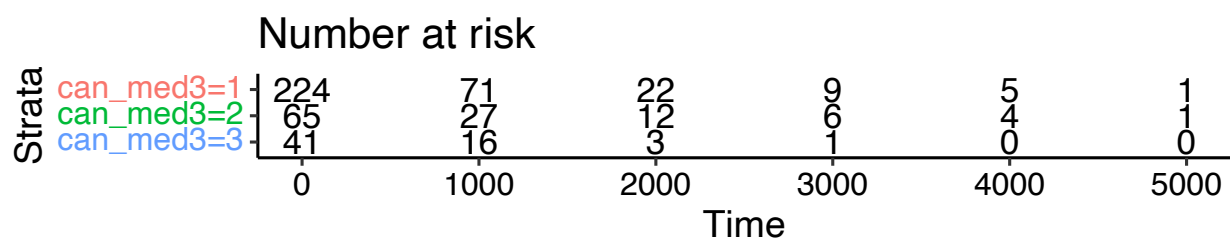
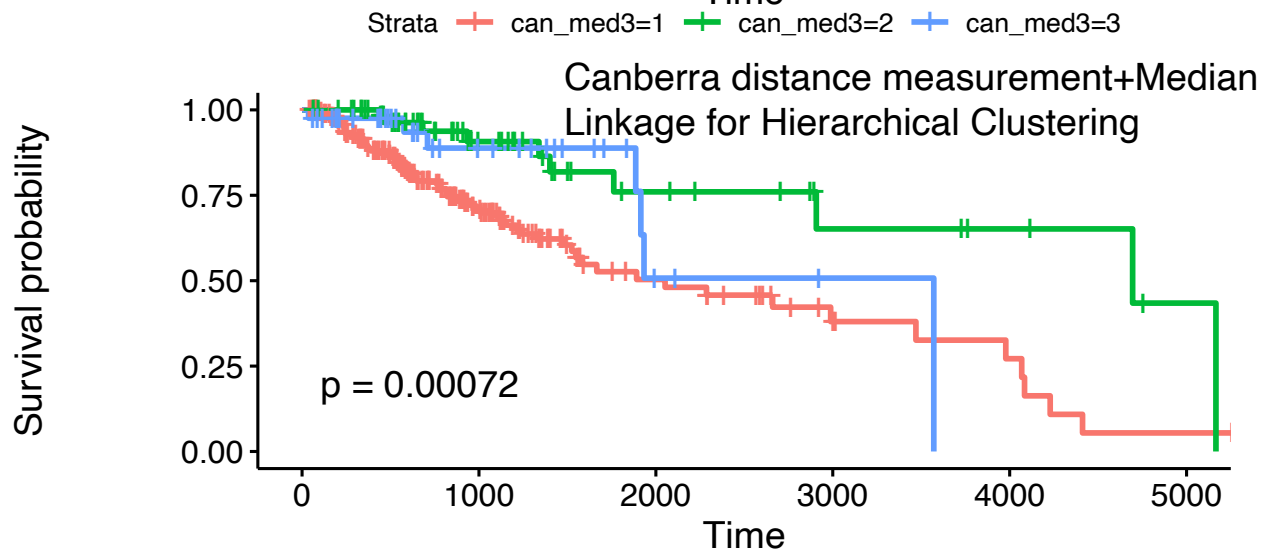
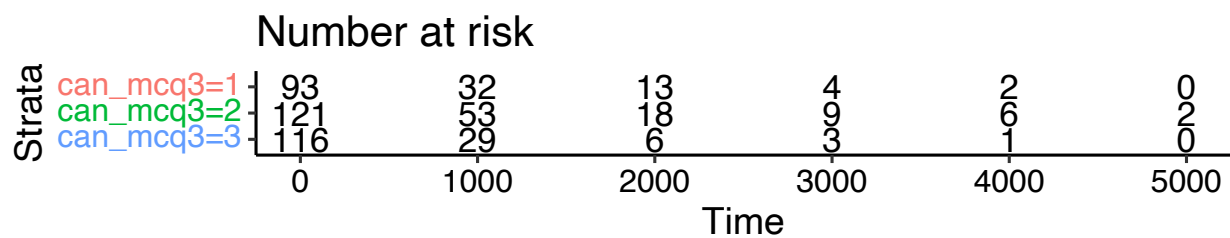
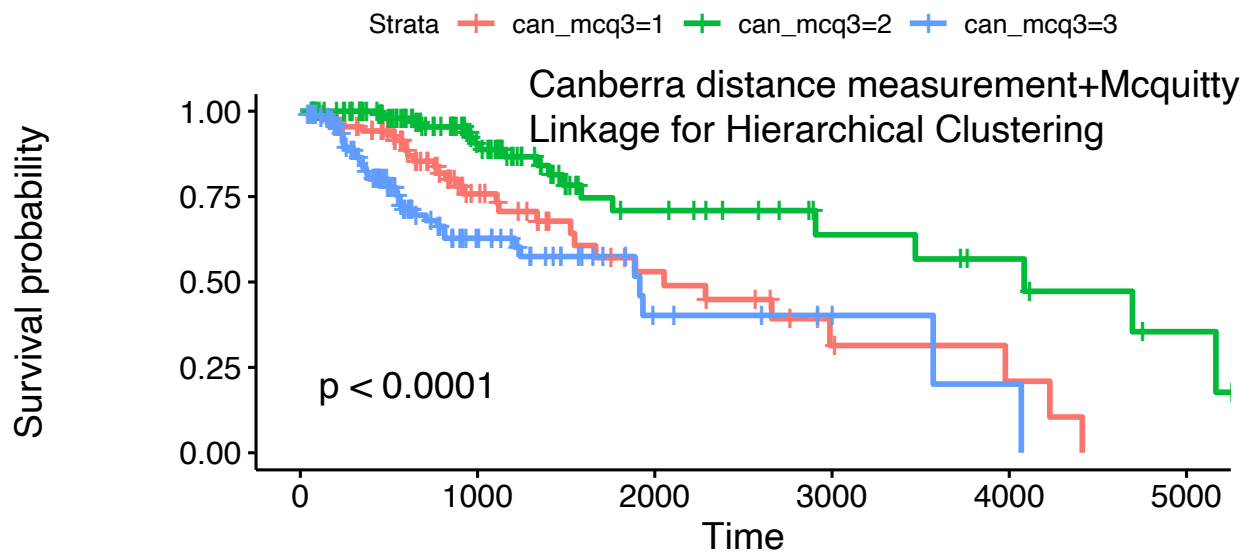


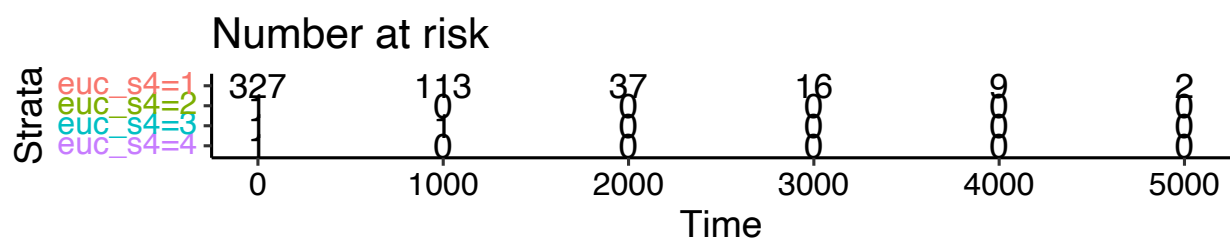
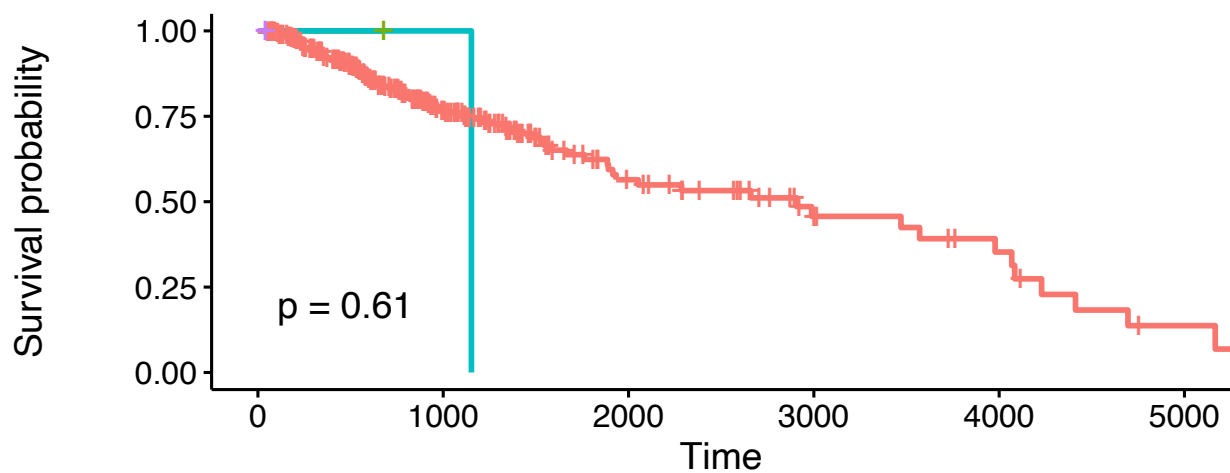
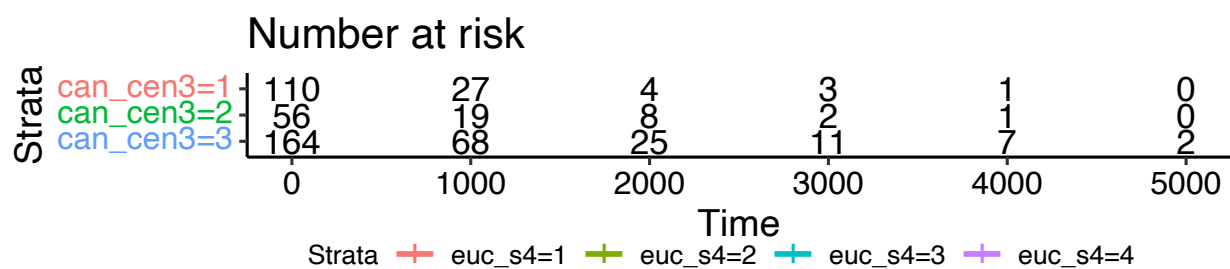
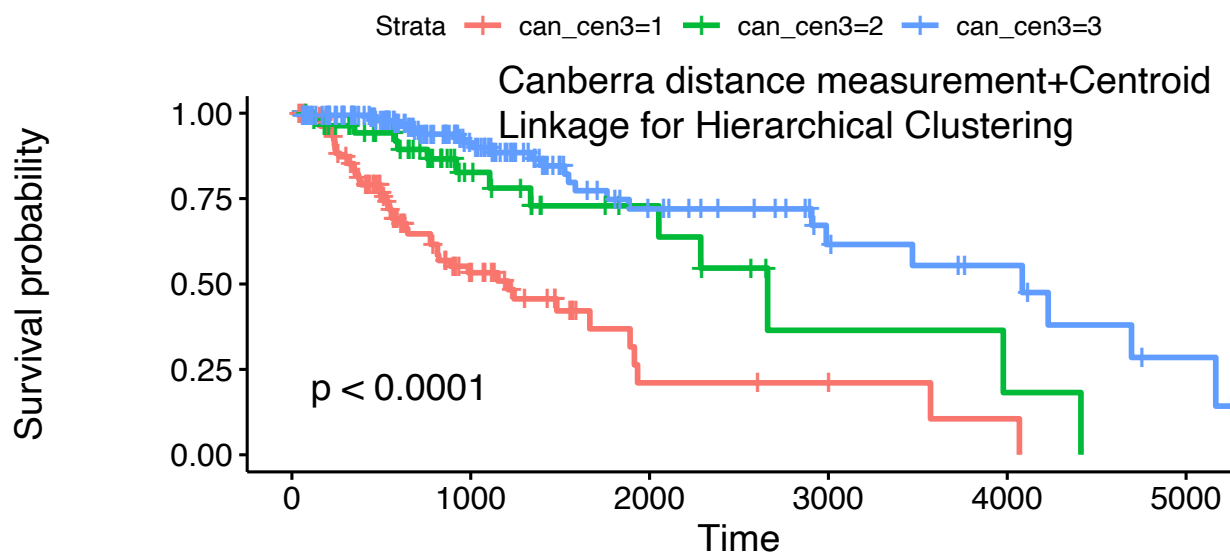


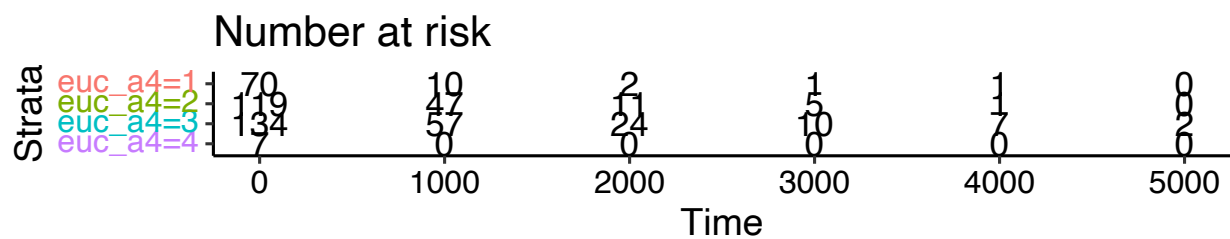
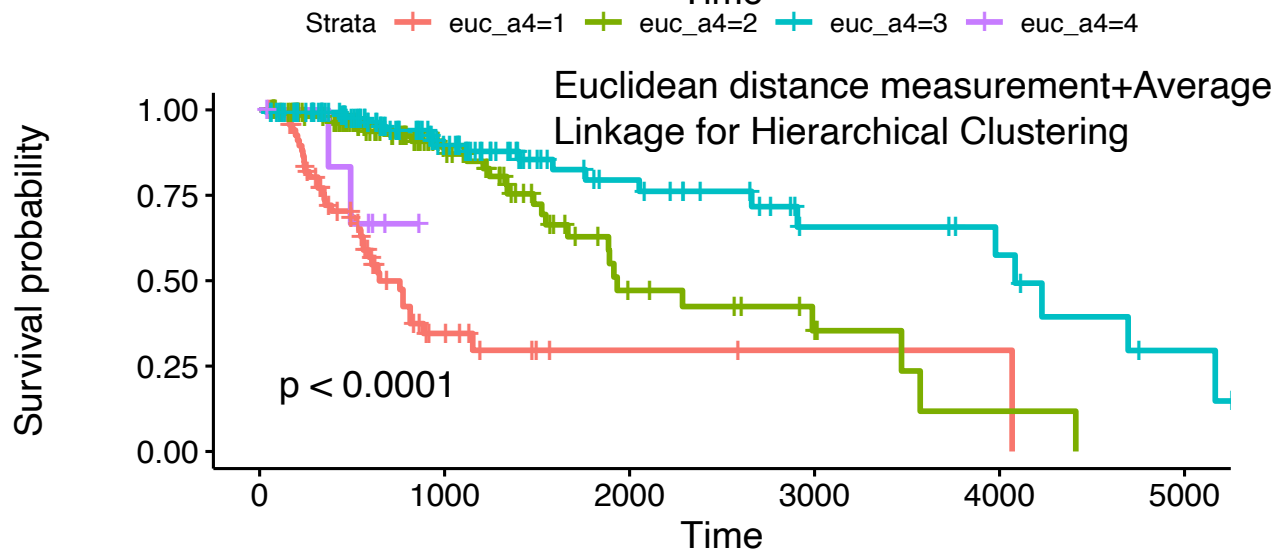
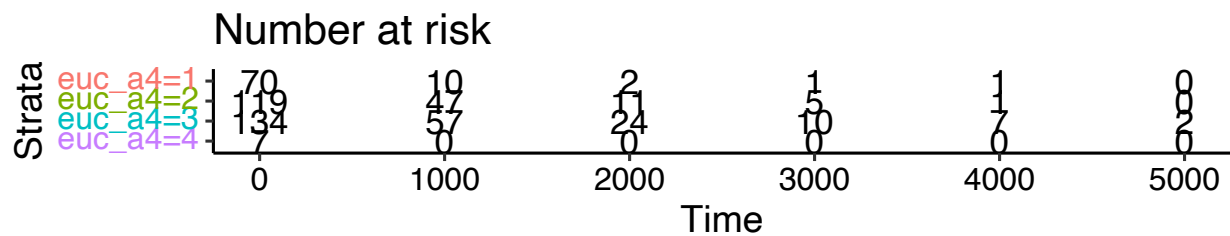
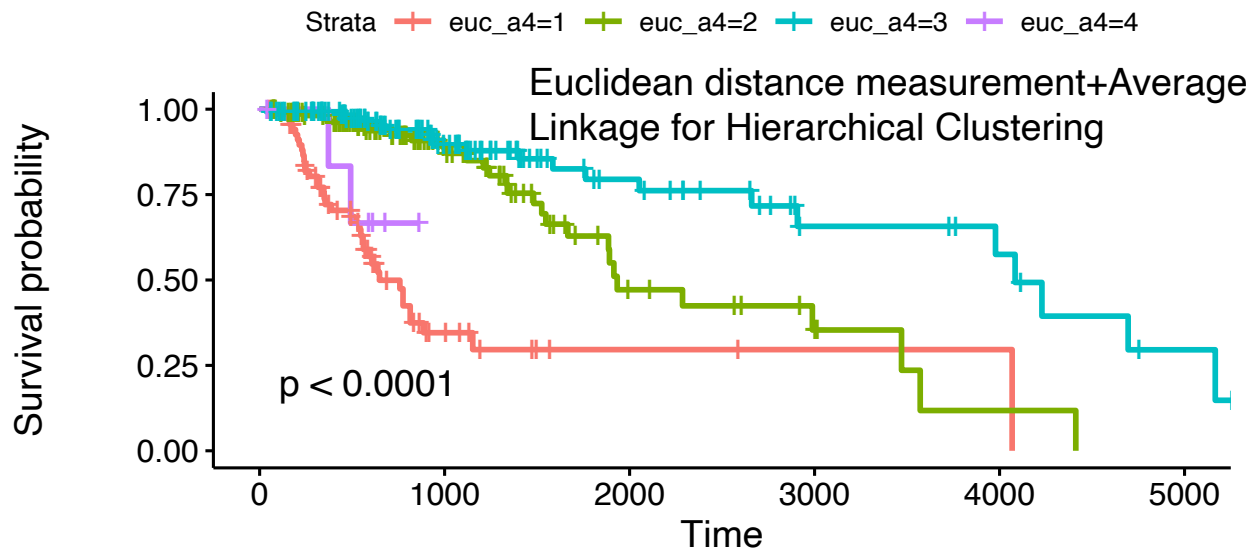


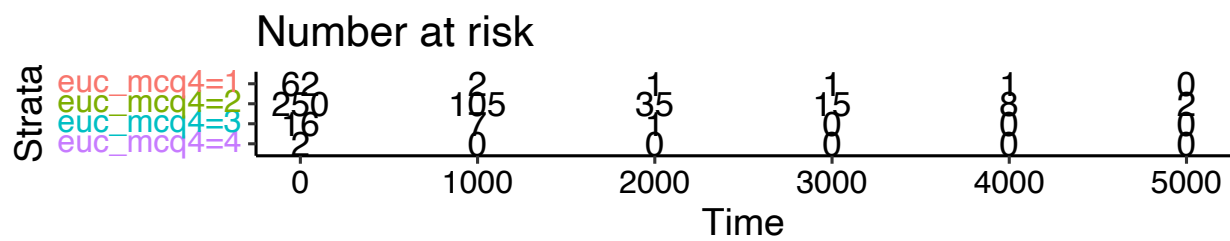
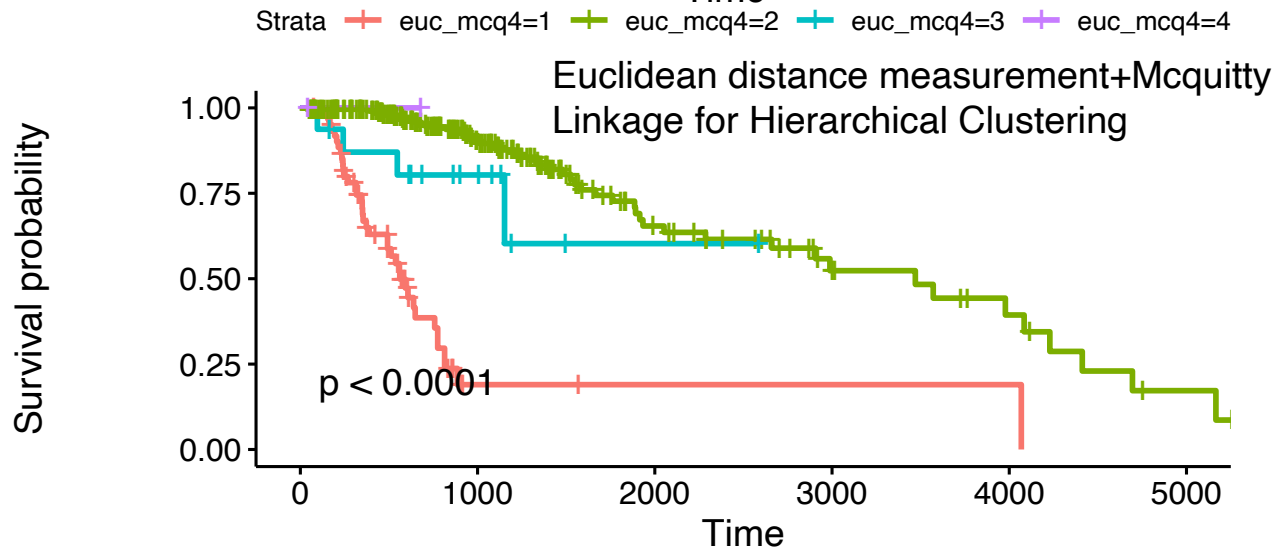
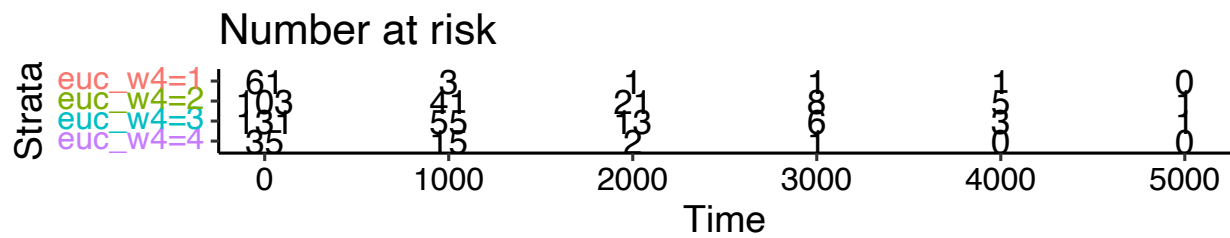
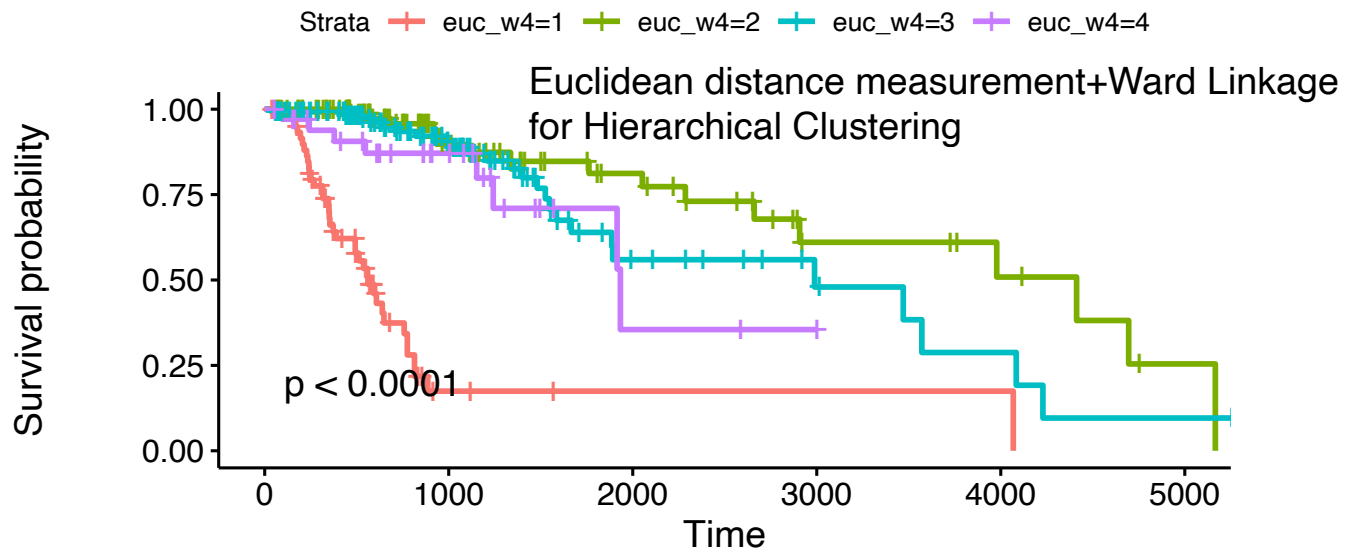




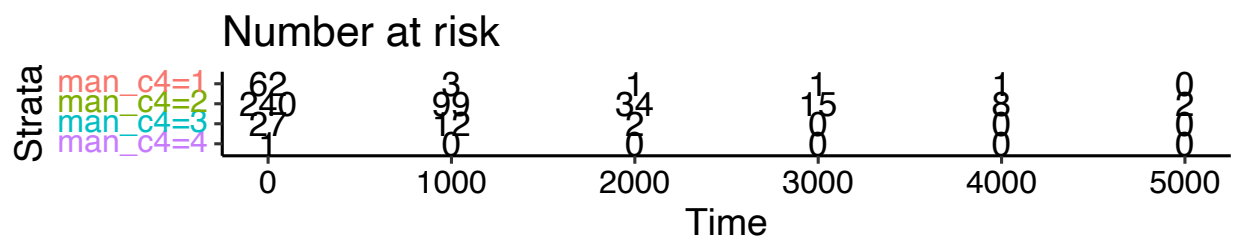
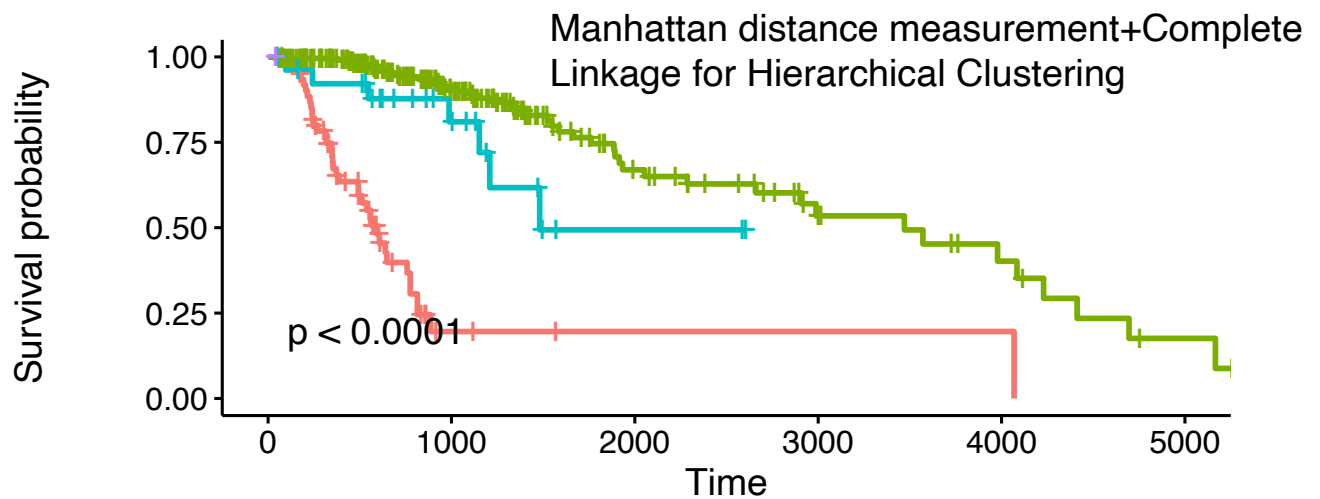
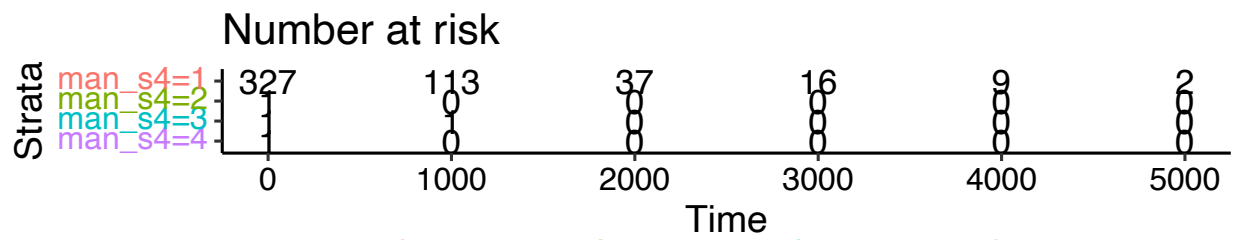
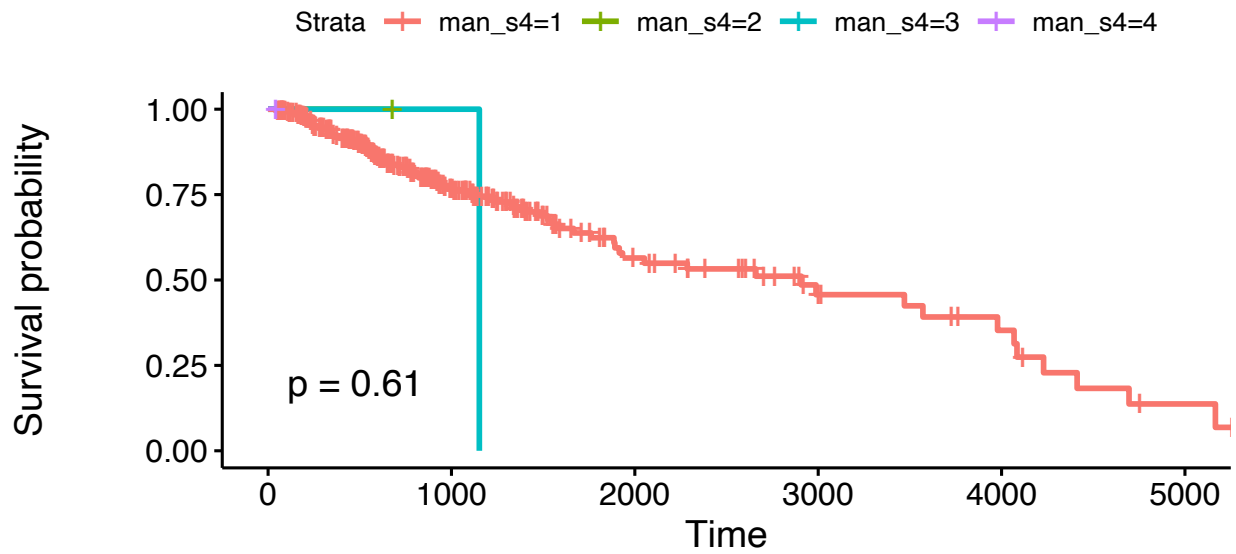


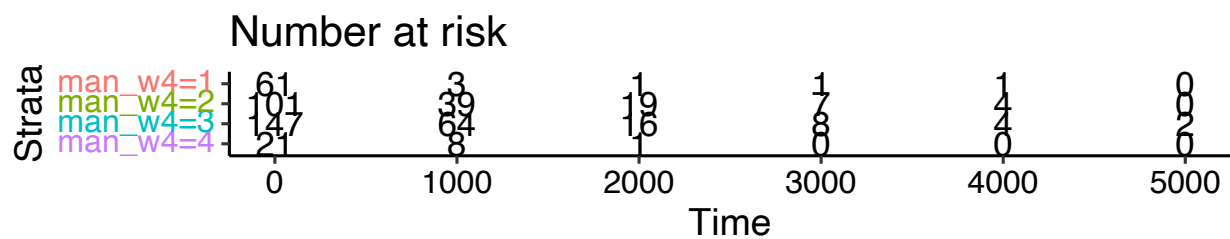
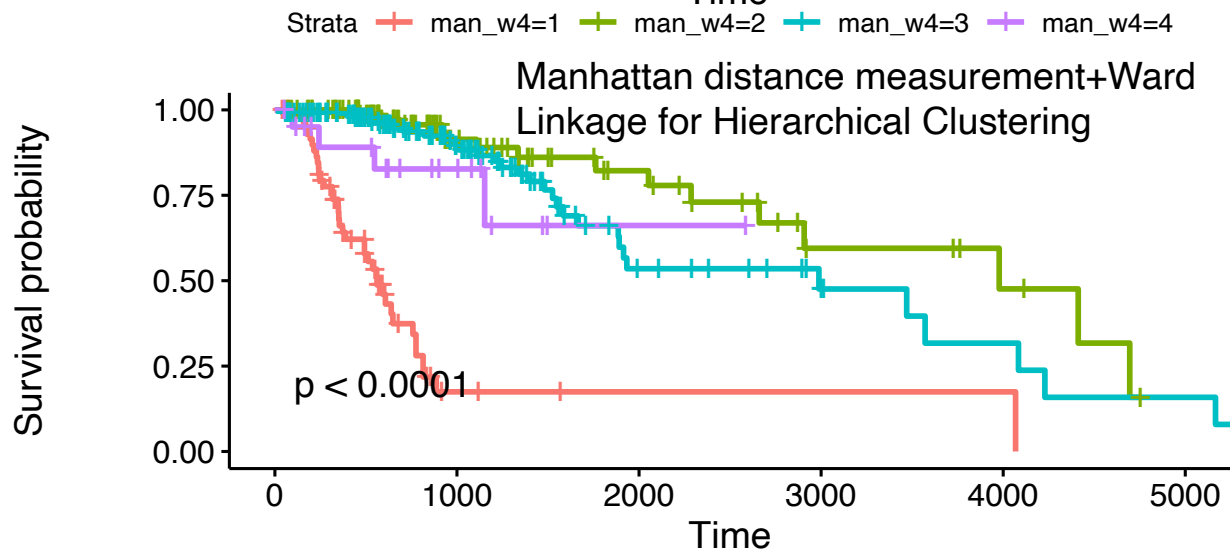
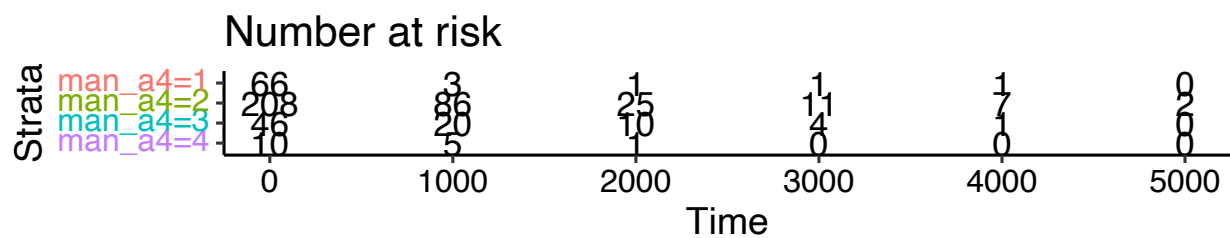
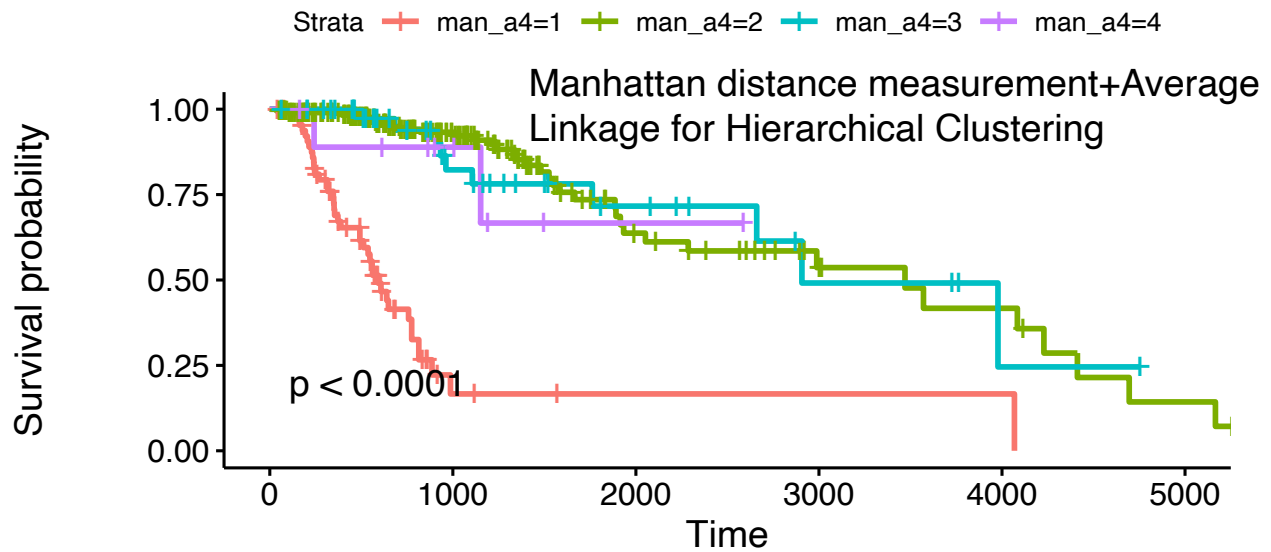


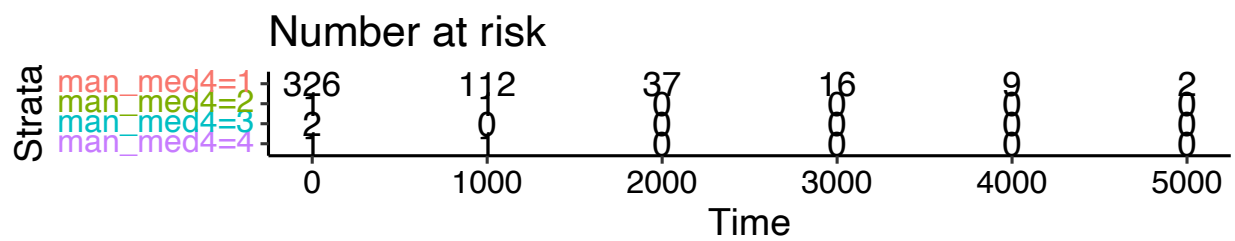
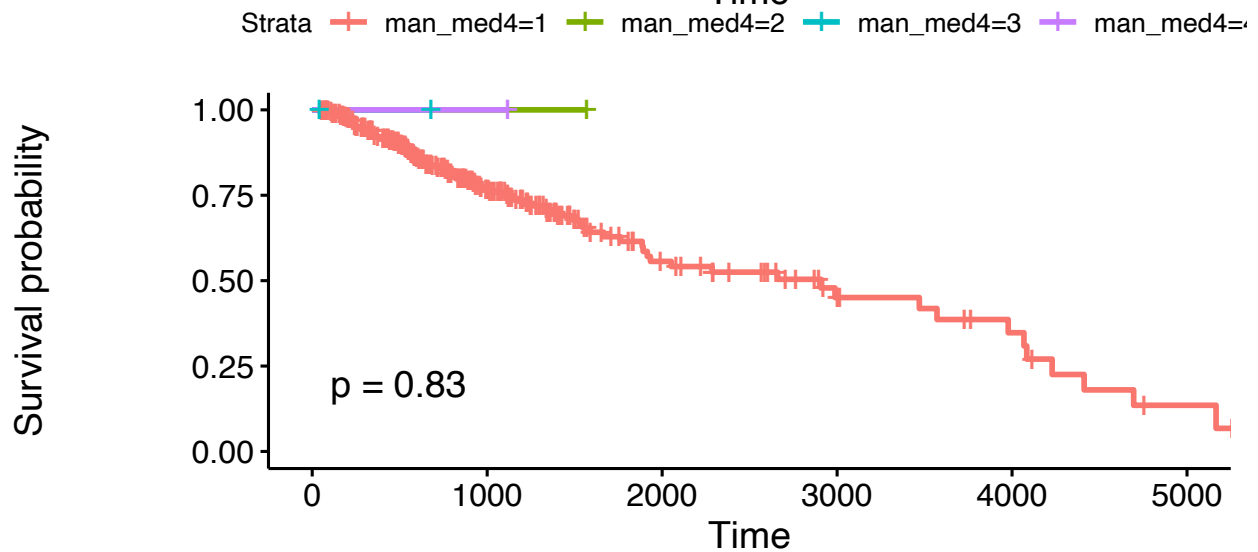
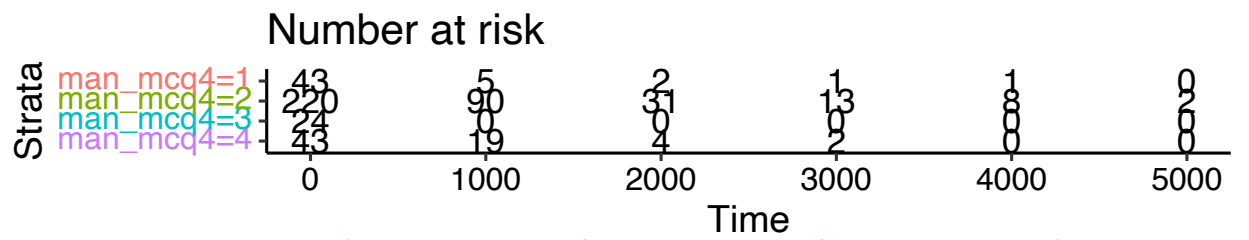
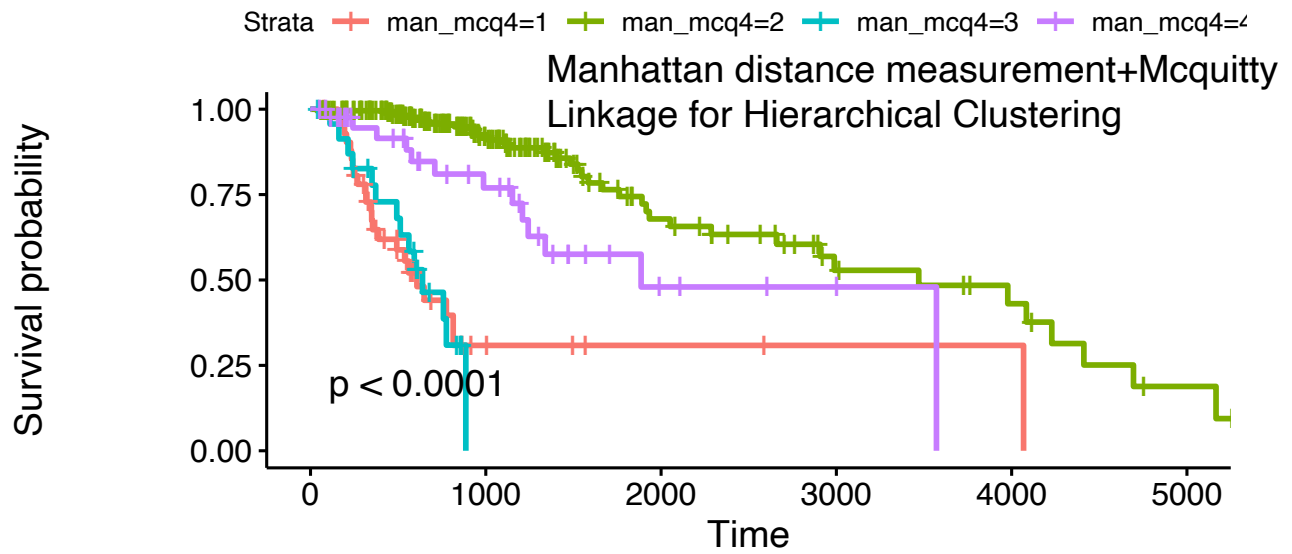


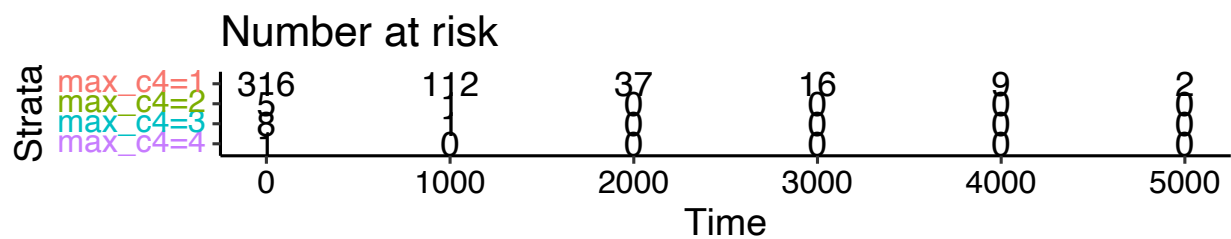
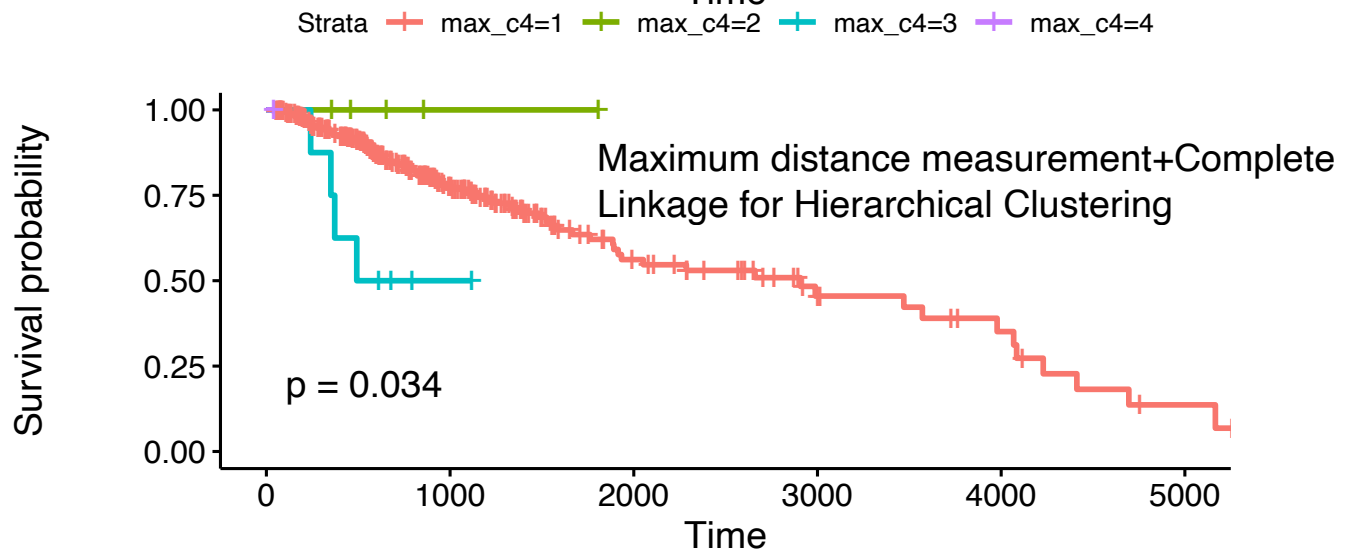
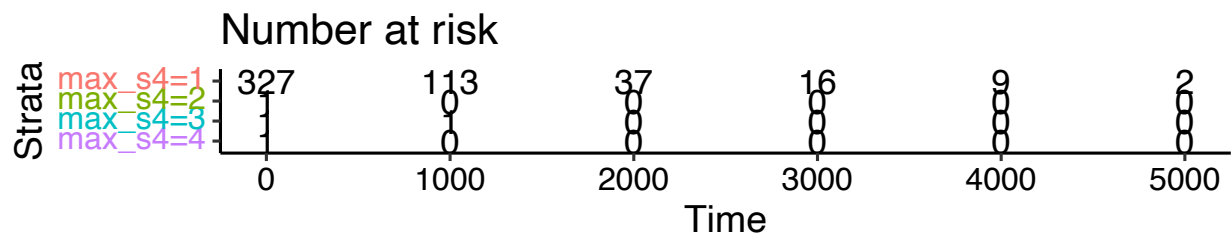
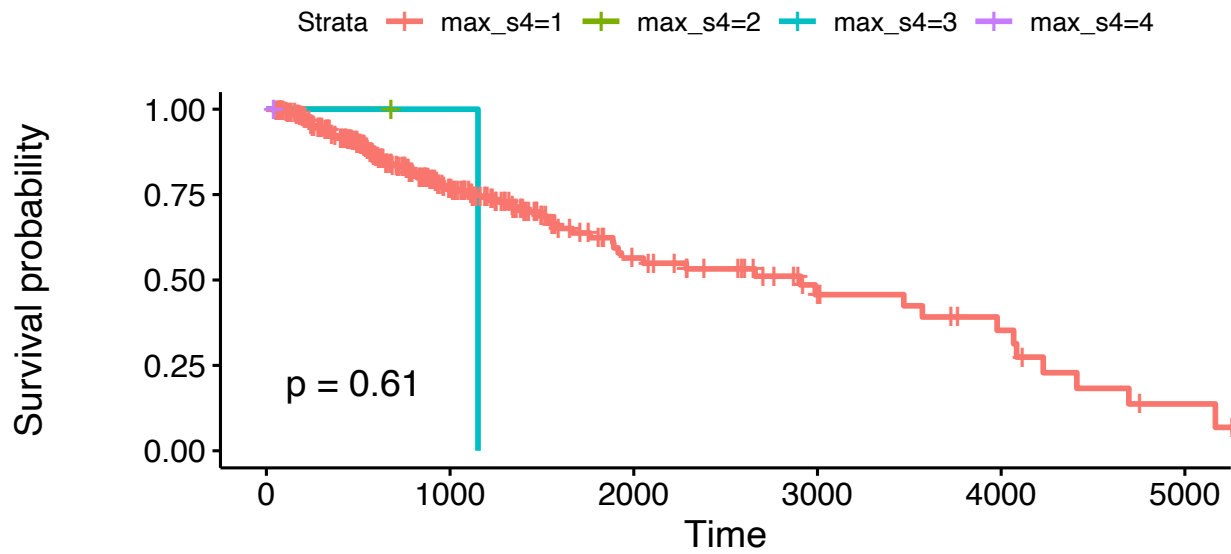


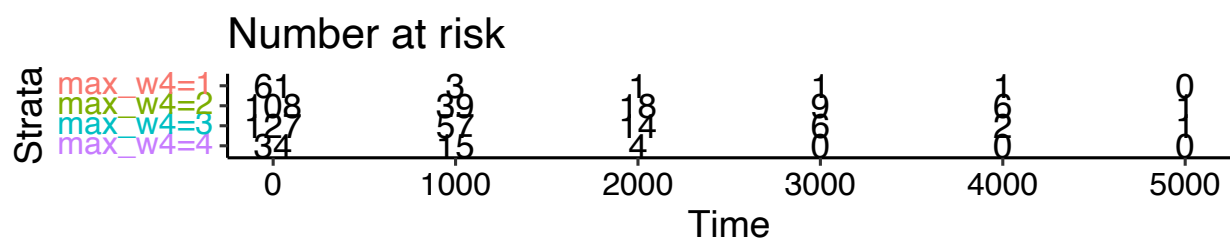
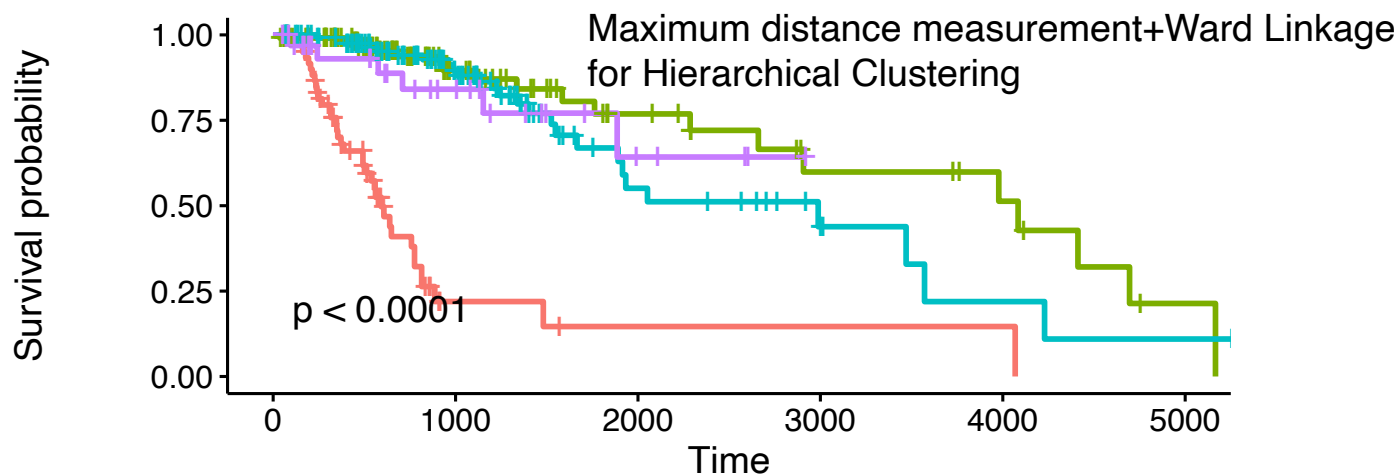
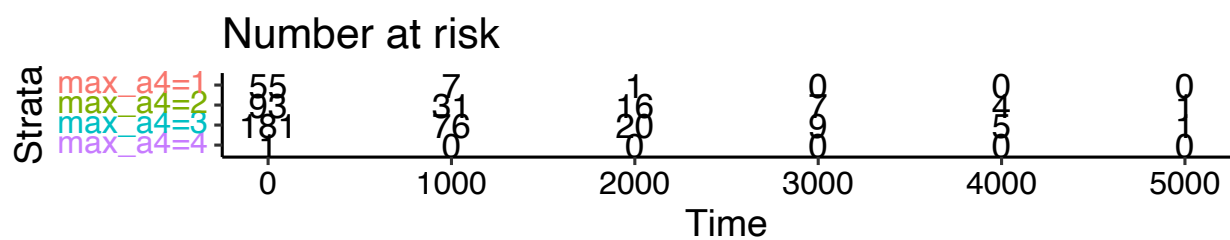
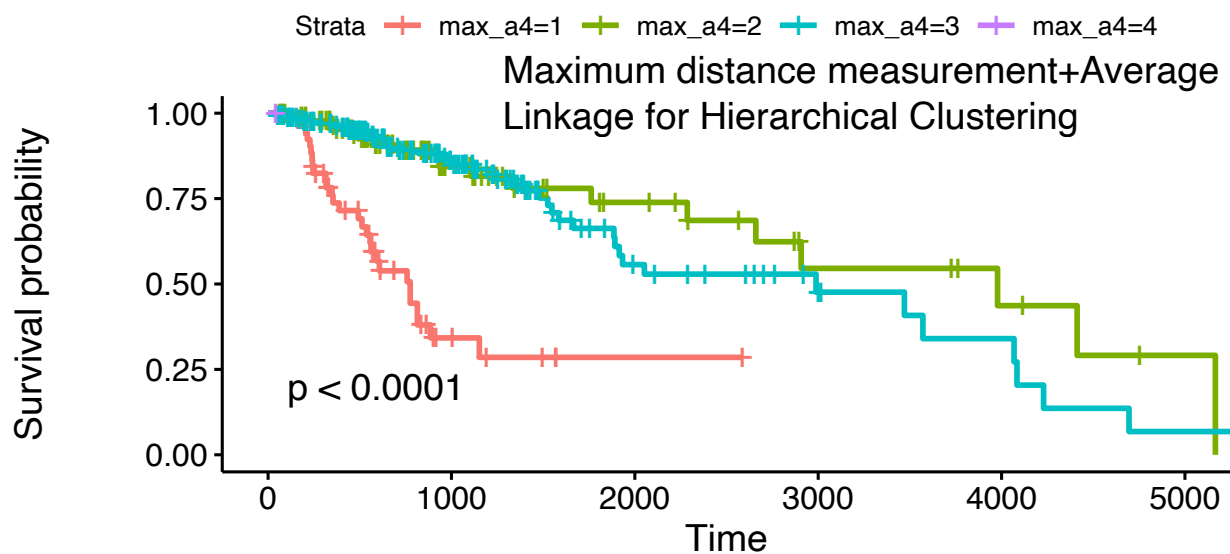


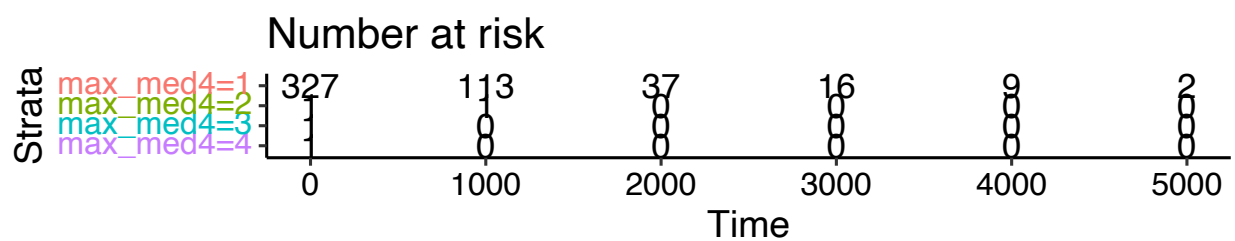
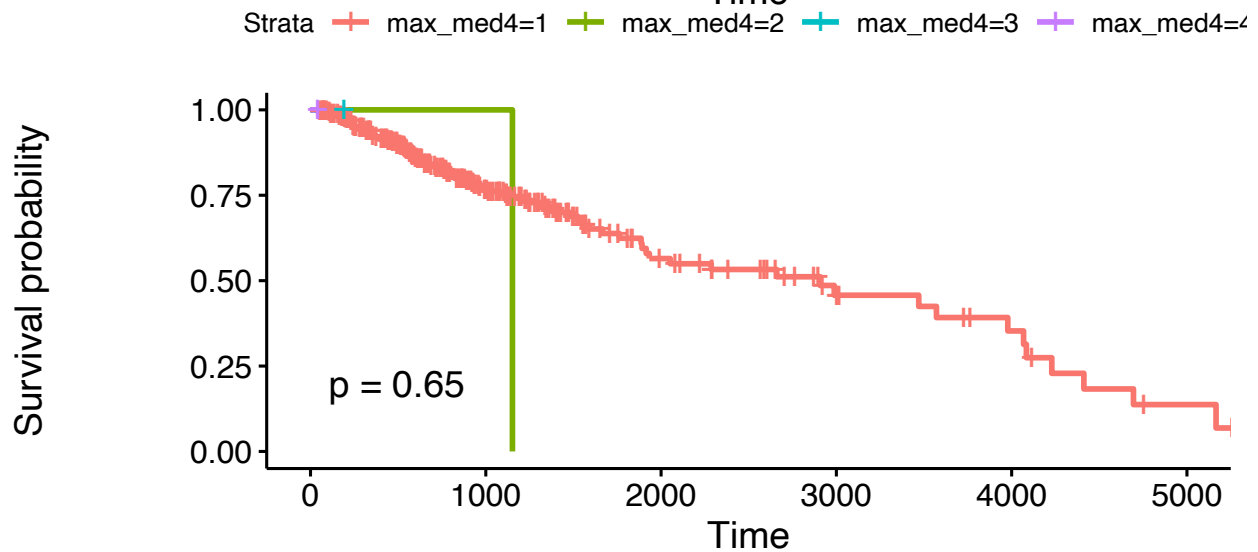
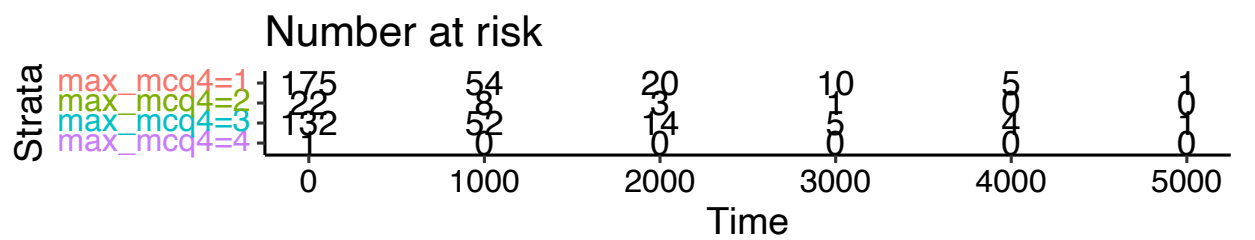
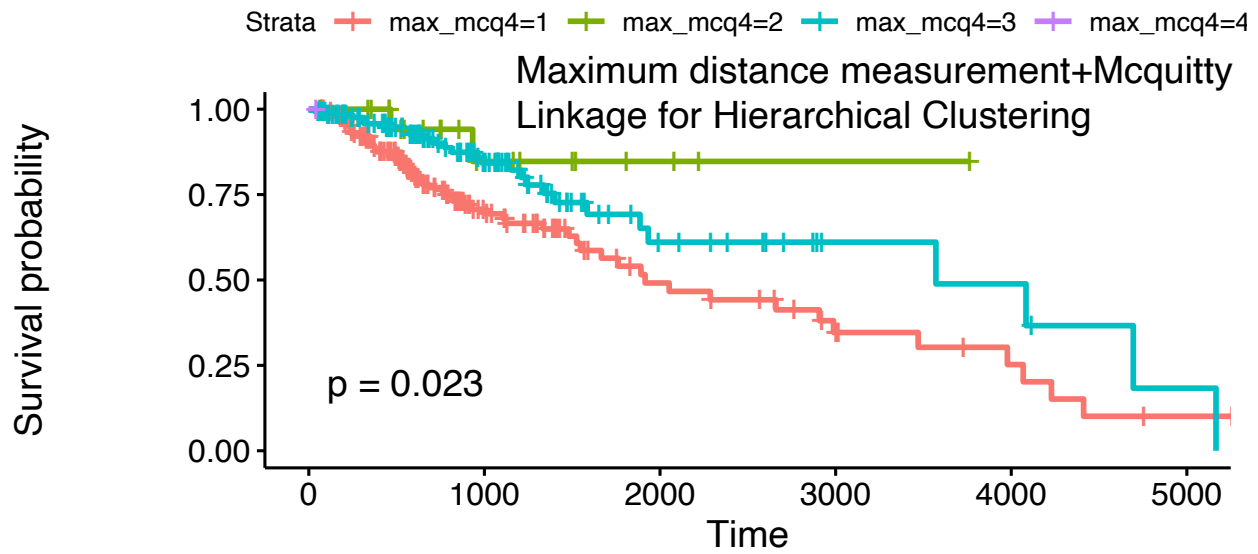


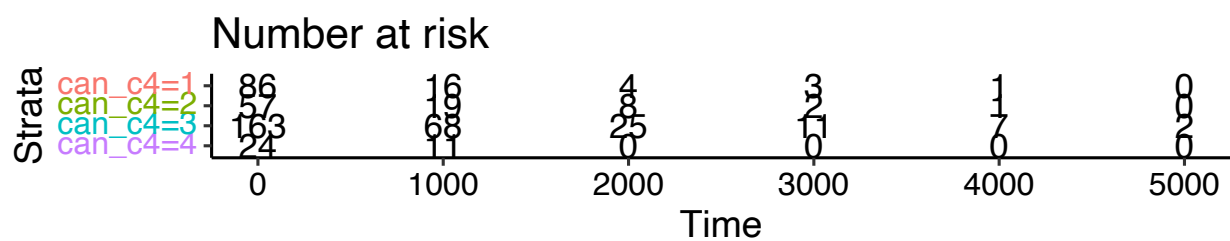
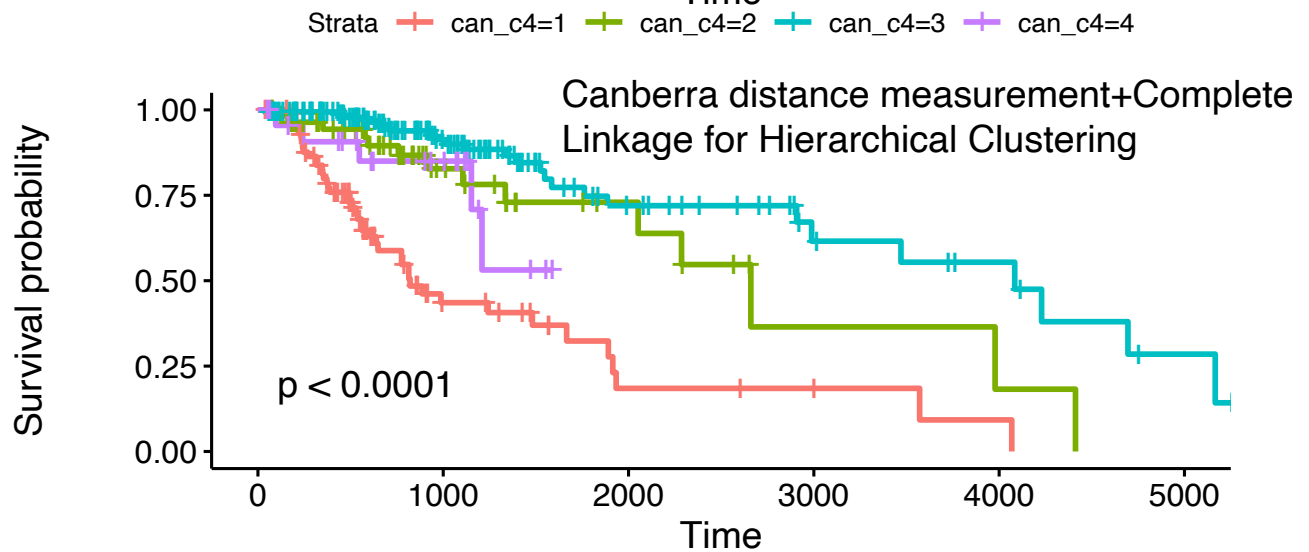
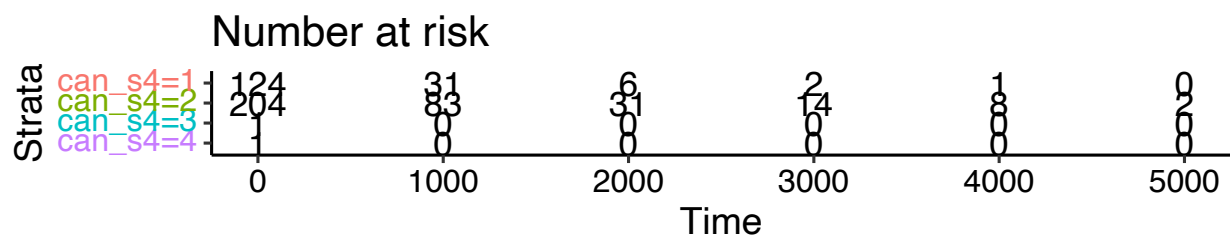
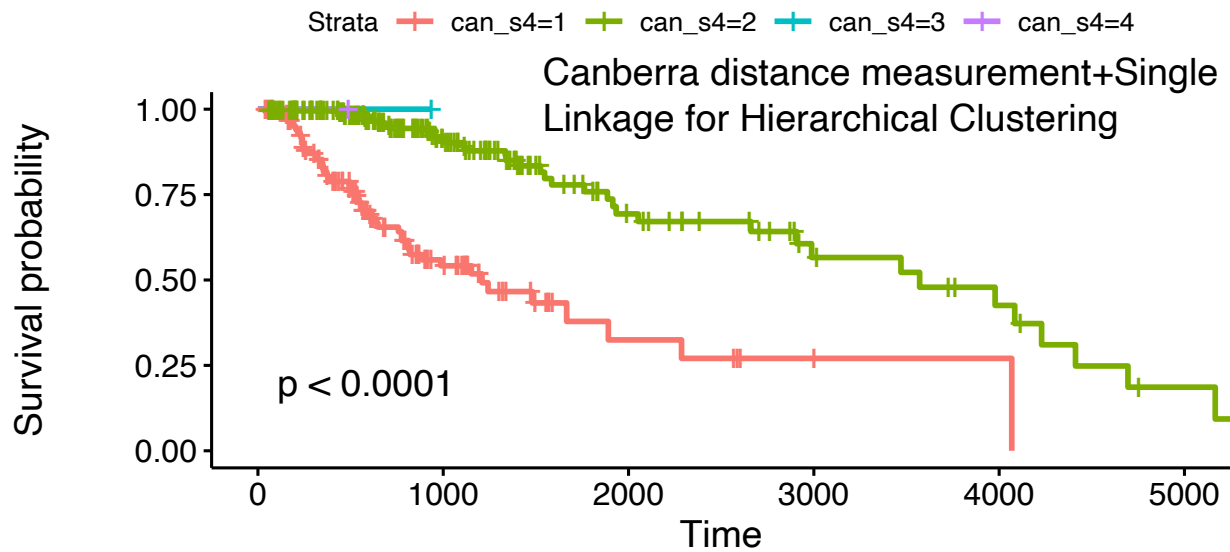


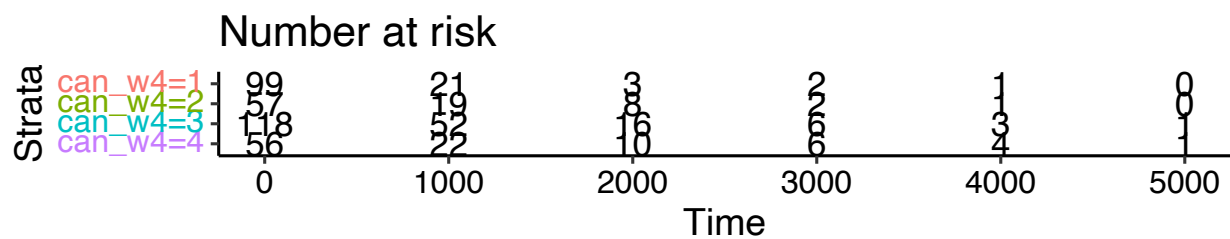
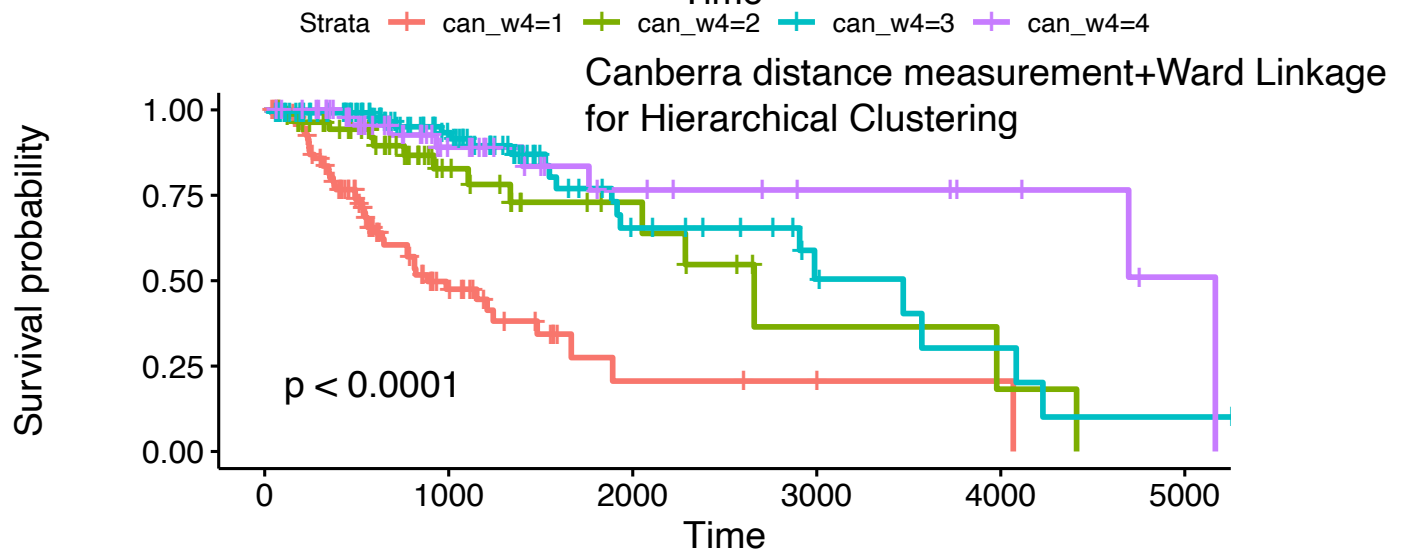
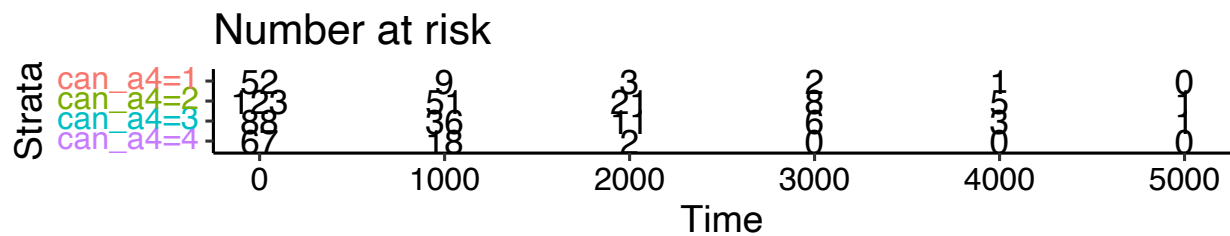
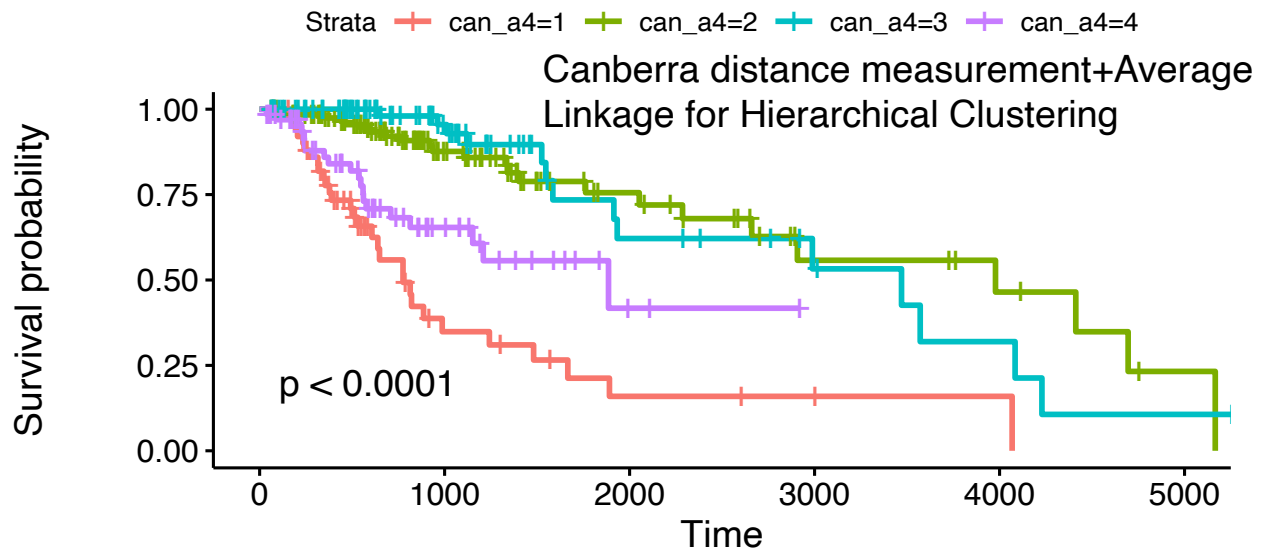




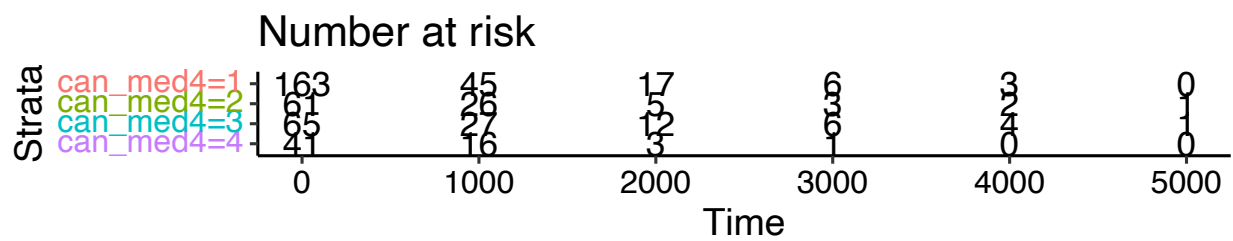
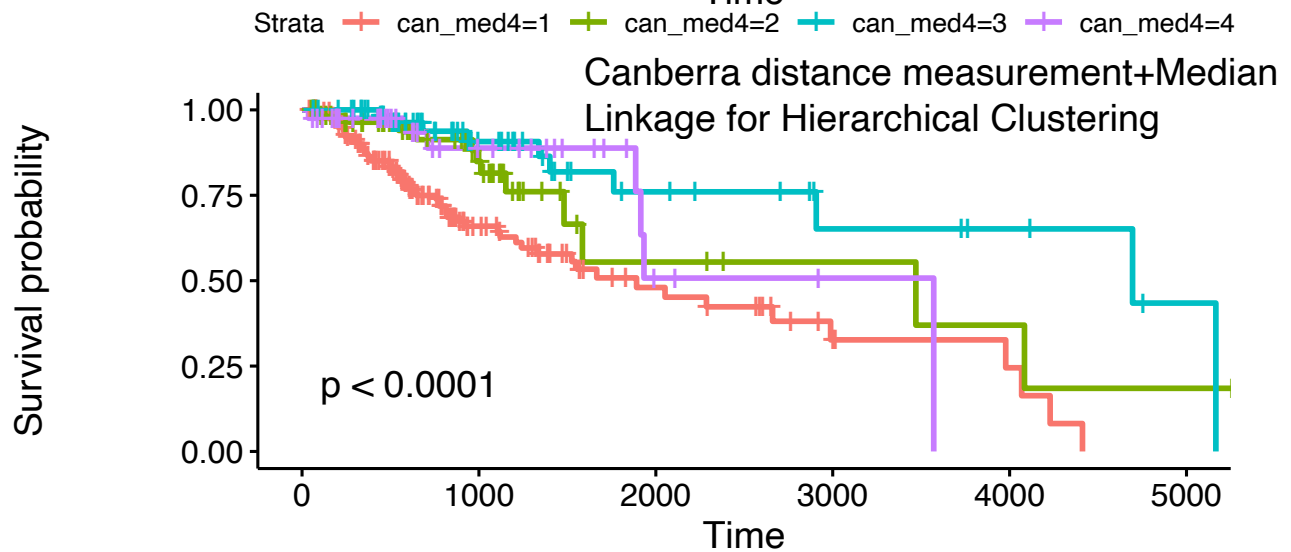
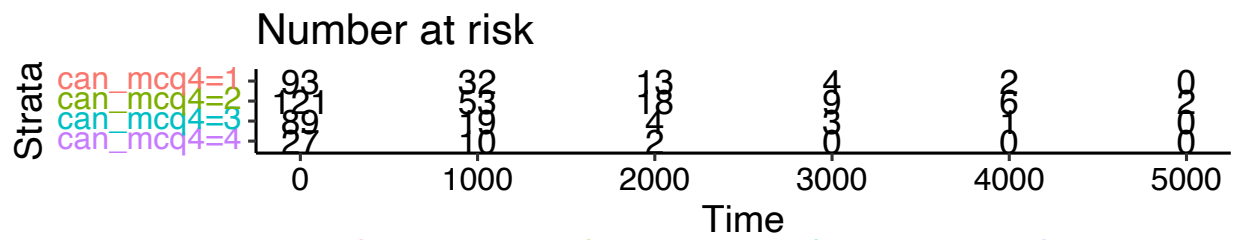
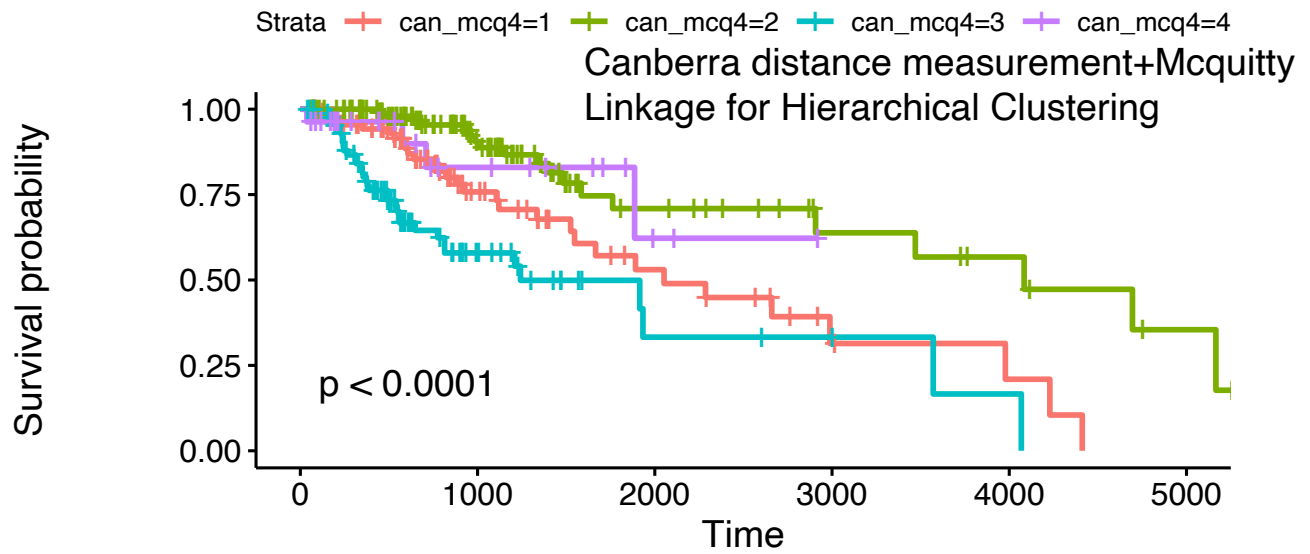


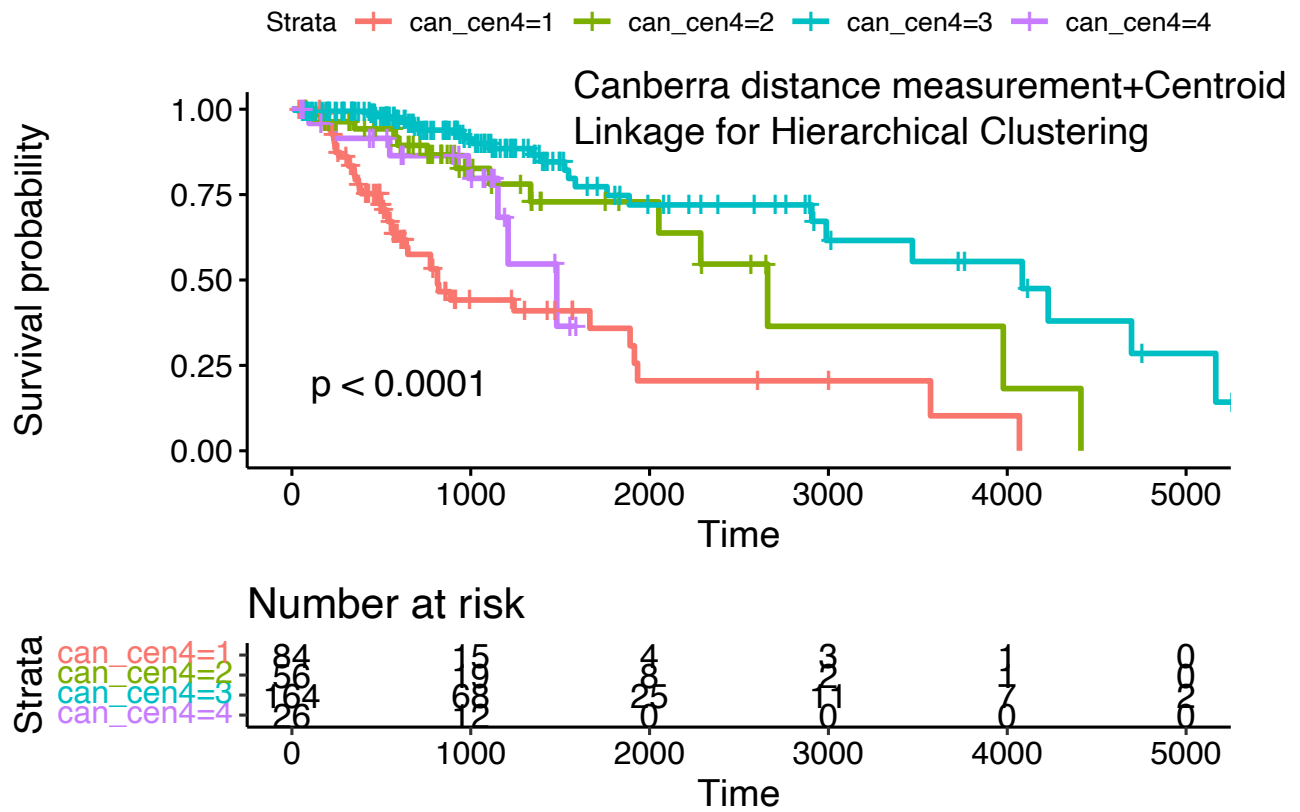






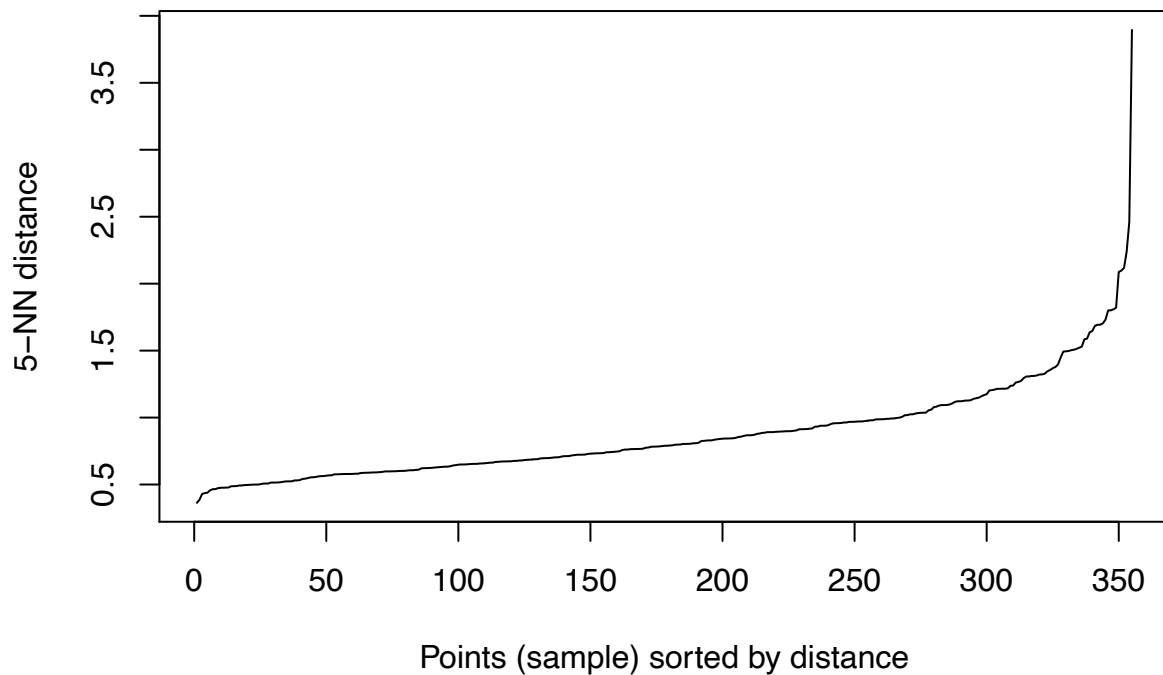






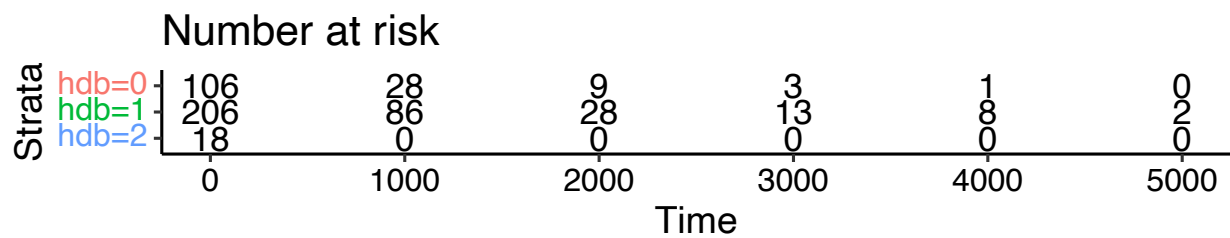
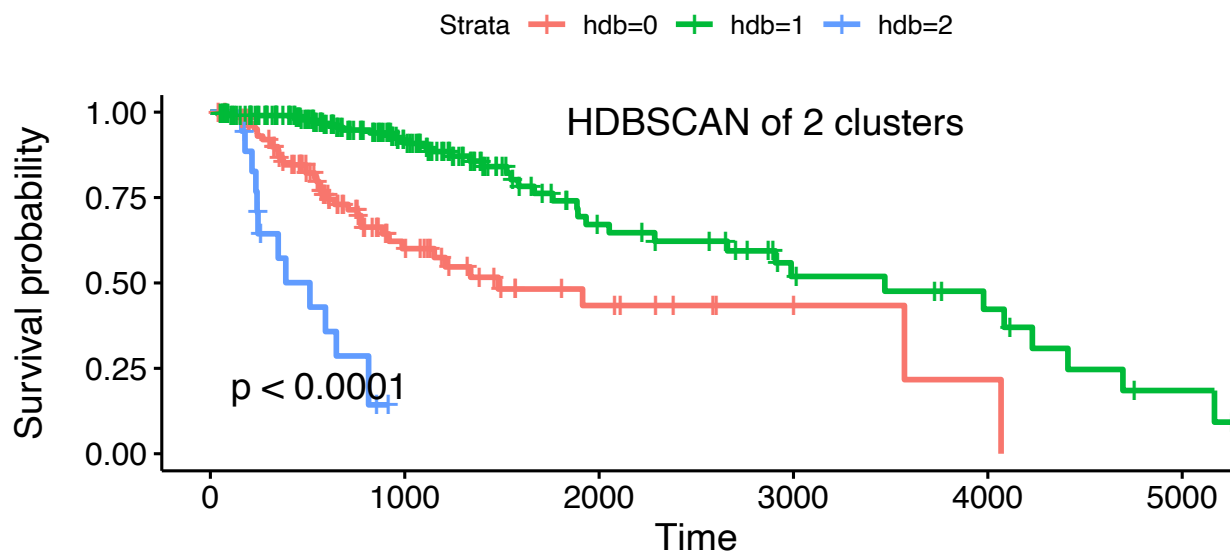
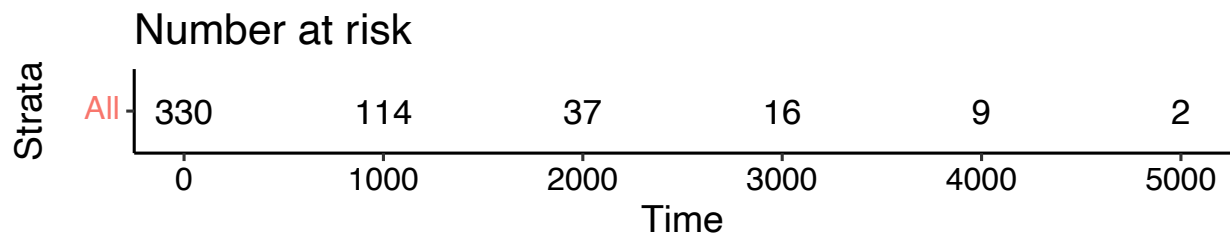
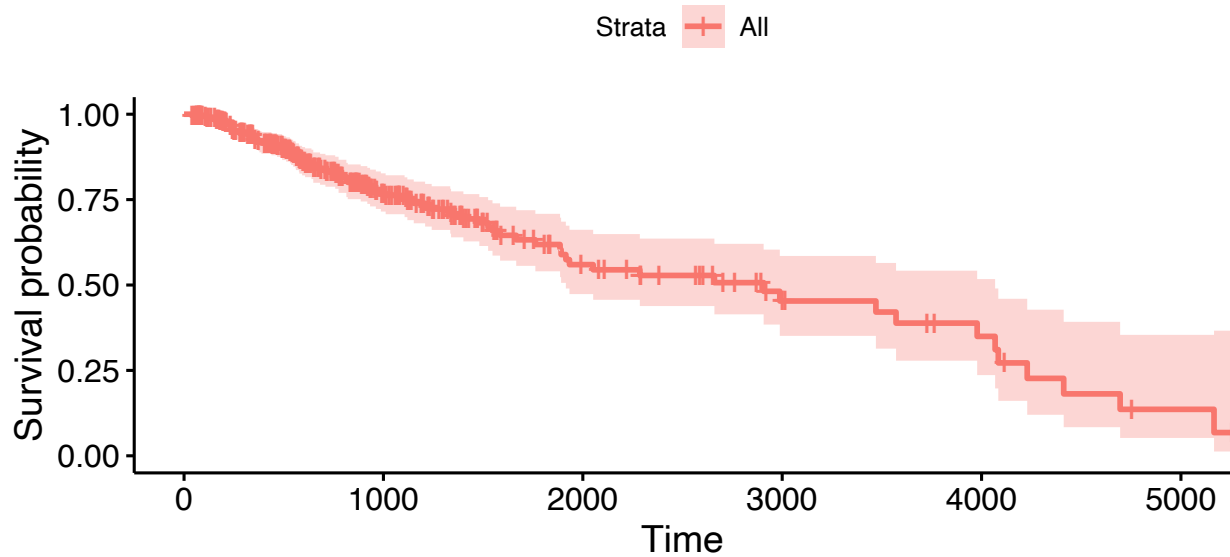
### 3. DBSCAN

## 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 a
## This is a null model.
```



#### 4. EM algorithm based on GMM

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

