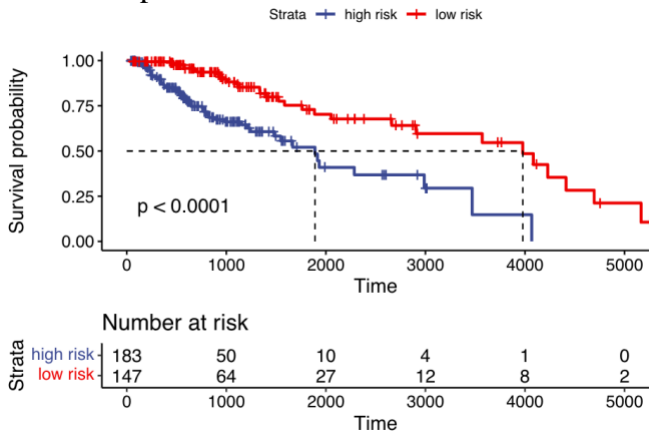
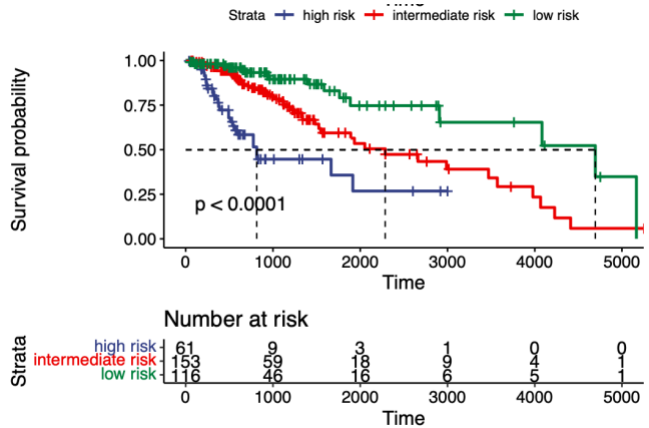


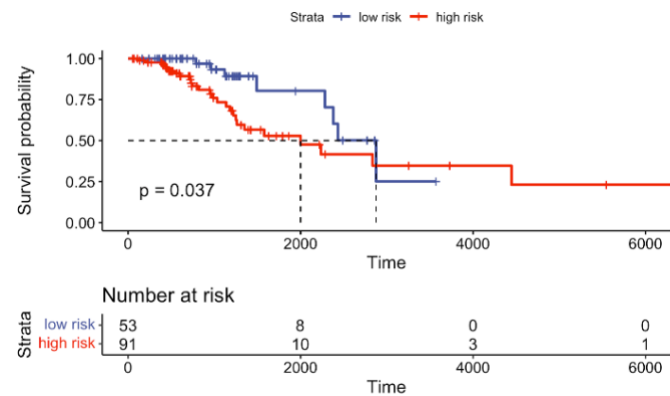
Since we change the legends of figures to present in the final report, we re-run all the codes of survival analysis again and might got different results, that is mainly due to the uncertainty brought by the clustering algorithms, but the overall results are unchanged. The results shown in the final reports are listed below:



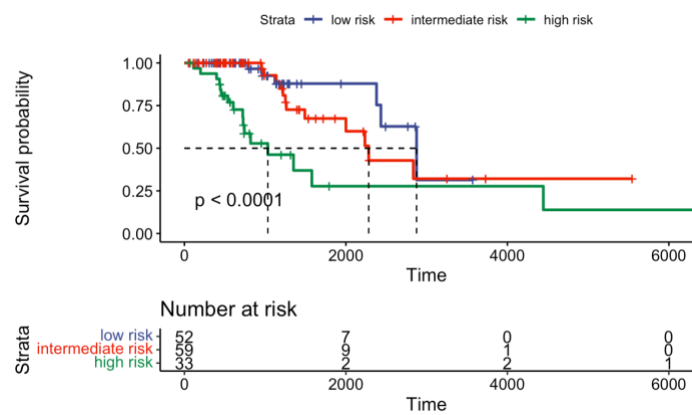
Glioma training set of 2 groups: Manhattan distance and Centroid linkage for the Hierarchical Clustering algorithm



Glioma training set of 3 groups: Canberra distance and Ward linkage for Hierarchical Clustering algorithm



Glioma test set of 2 groups: Euclidean distance and Mcquitty linkage for the Hierarchical Clustering algorithm



Glioma test set of 3 groups: Maximum distance and Ward linkage for Hierarchical Clustering algorithm