

Supplementary data 9: The R session info for the statistical and data analysis.

R version 4.1.0 (2021-05-18)
Platform: x86_64-w64-mingw32/x64 (64-bit)
Running under: Windows 8.1 x64 (build 9600)

Matrix products: default

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[3] LC_MONETARY=Icelandic_Iceland.1252 LC_NUMERIC=C
[5] LC_TIME=Icelandic_Iceland.1252

attached base packages:
[1] parallel stats4 stats graphics grDevices utils
datasets methods base

other attached packages:
[1] EnvStats_2.4.0 flexsurv_2.1
[3] muhaz_1.2.6.4 LogisticDx_0.2
[5] MKmisc_1.8 forcats_0.5.1
[7] purrr_0.3.4 tibble_3.1.2
[9] tidyverse_1.3.1 ggplotify_0.0.7
[11] DescTools_0.99.42 tidyr_1.1.3
[13] kableExtra_1.3.4 tableone_0.12.0
[15] pheatmap_1.0.12 survminer_0.4.9
[17] ggpubr_0.4.0 survival_3.2-11
[19] ggplot2_3.3.4 stringr_1.4.0
[21] dplyr_1.0.7 stringi_1.6.2
[23] readr_1.4.0 XML_3.99-0.6
[25] reshape2_1.4.4 Homo.sapiens_1.3.1
[27] TxDb.Hsapiens.UCSC.hg19.knownGene_3.2.2 org.Hs.eg.db_3.13.0
[29] GO.db_3.13.0 OrganismDbi_1.34.0
[31] GenomicFeatures_1.44.0 TCGAbiolinks_2.20.0
[33] clusterProfiler_4.0.0 enrichplot_1.12.1
[35] goseq_1.44.0 geneLenDataBase_1.28.0
[37] BiasedUrn_1.07 qusage_2.26.0
[39] limma_3.48.0 DO.db_2.9
[41] biomaRt_2.48.1 AnnotationDbi_1.54.1
[43] DESeq2_1.32.0
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[45] Biobase_2.52.0 MatrixGenerics_1.4.0
[47] matrixStats_0.59.0 GenomicRanges_1.44.0
[49] GenomeInfoDb_1.28.0 IRanges_2.26.0
[51] S4Vectors_0.30.0 BiocGenerics_0.38.0

loaded via a namespace (and not attached):
[1] estimability_1.3 rappdirs_0.3.3

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