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| Packages used in the analysis  [1] mixOmics\_6.3.2 MASS\_7.3-50 gdtools\_0.1.7  [4] bindrcpp\_0.2.2 rJava\_0.9-10 xlsx\_0.6.1  [7] officer\_0.3.2 rvg\_0.1.9 scales\_0.5.0  [10] viridis\_0.5.1 viridisLite\_0.3.0 broom\_0.4.5  [13] reshape2\_1.4.3 knitr\_1.20 tableone\_0.9.3  [16] ISOweek\_0.6-2 summarytools\_0.8.5 corrplot\_0.84  [19] Hmisc\_4.1-1 Formula\_1.2-3 survival\_2.42-3  [22] lattice\_0.20-35 forcats\_0.3.0 stringr\_1.3.1  [25] purrr\_0.2.5 readr\_1.1.1 tidyr\_0.8.1  [28] tibble\_1.4.2 tidyverse\_1.2.1 lubridate\_1.7.4  [31] readxl\_1.1.0 RColorBrewer\_1.1-2 dplyr\_0.7.6  [34] data.table\_1.11.4 rgdal\_1.3-3 sp\_1.3-1  [37] plyr\_1.8.4 ggplot2\_3.0.0 |