Normalizing

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```
library(ggplot2)
library(dplyr)
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
       filter, lag
##
## The following objects are masked from 'package:base':
##
       intersect, setdiff, setequal, union
library(tidyr)
library(gridExtra)
## Attaching package: 'gridExtra'
## The following object is masked from 'package:dplyr':
##
       combine
library(GGally)
## Registered S3 method overwritten by 'GGally':
    method from
           ggplot2
    +.gg
library(knitr)
library(corrplot)
## corrplot 0.95 loaded
load('combined_assay_data.RData')
```

```
X <- assay.data[,3:length(assay.data)]
assay.data$Toxicity <- factor(assay.data$Toxicity, levels = unique(assay.data$Toxicity))
y <- assay.data$Toxicity

# before logarithmizing
X_normalized <- apply(X[,2:length(X)], 2, function(x) x - X$Cmax)</pre>
```

Univariate analysis

Summary statistics

```
# Function to calculate the required statistics for each column
summary stats <- function(x) {</pre>
  c(
    minimum = min(x),
    Q1 = quantile(x, 0.25),
    median = median(x),
    mean = mean(x, na.rm = TRUE),
    Q3 = quantile(x, 0.75),
    maximum = max(x),
    variance = var(x),
    sd = sd(x),
    range = max(x) - min(x)
  )
}
# Apply the function to each column of the selected data
normalized_summary_df <- as.data.frame(t(apply(X_normalized, 2, summary_stats)))</pre>
# Print the result
print(normalized_summary_df)
```

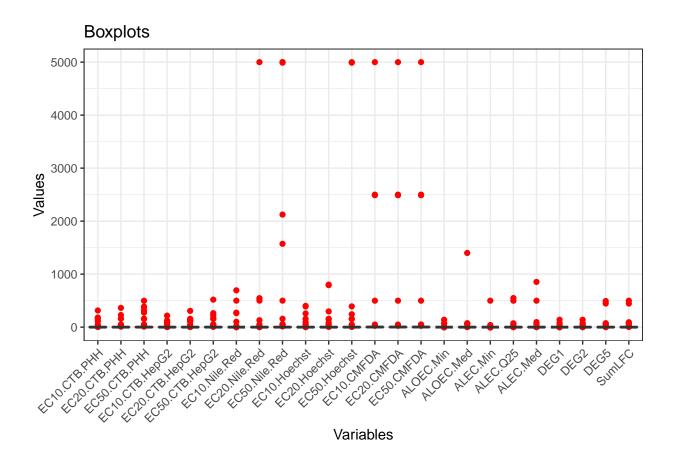
```
##
                                    Q1.25%
                                                                      Q3.75%
                      minimum
                                                 median
                                                              mean
## EC10.CTB.PHH -0.007762404 3.309113e-02 2.044170e-01 9.138898 1.57484713
## EC20.CTB.PHH
                -0.006875404 4.240748e-02 3.395719e-01 11.448512 2.29765775
                -0.005319404 8.436536e-02 7.082732e-01 25.204183 3.80918376
## EC50.CTB.PHH
## EC10.CTB.HepG2 -0.083363000 1.899247e-02 1.040546e-01 7.595209 0.90374870
## EC20.CTB.HepG2 -0.001636305 3.239827e-02 1.858485e-01 10.899380 1.29379018
## EC50.CTB.HepG2 -0.000991298 6.538990e-02 3.375496e-01 17.871516 2.47637046
## EC10.Nile.Red -0.649868351 5.998640e-03 3.674132e-02 19.108779 0.23612388
## EC20.Nile.Red -0.046552279 1.443276e-02 8.578657e-02 68.964544 0.49203619
## EC50.Nile.Red -0.023375293 6.170916e-02 4.427895e-01 198.186811 3.37338534
## EC10.Hoechst
                 -0.107019285 1.952632e-02 8.185485e-02 14.721234 0.73064420
## EC20.Hoechst -0.024270697 3.682121e-02 1.180755e-01 24.113290 1.85069702
## EC50.Hoechst -0.001494856 7.026302e-02 3.287699e-01 111.441390 2.49595471
## EC10.CMFDA
                 -0.015711105 4.990961e-01 2.483185e+00 132.116748 2.49981925
## EC20.CMFDA
                 -0.013411678 4.999040e-01 2.483185e+00 132.132939 2.49981925
## EC50.CMFDA
                -0.005927245 4.999963e-01 2.492753e+00 132.739345 2.49989849
## ALOEC.Min
                -6.148000000 -3.834621e-03 5.858060e-04 2.947964 0.01981741
                -0.049844507 7.315891e-04 4.107450e-02 16.754342 0.49853001
## ALOEC.Med
```

```
## ALEC.Min
                 -9.962211999 -4.338442e-03 5.181465e-05 5.426345 0.01590561
## ALEC.Q25
                 -0.170062880 1.053011e-04 4.226874e-02 12.821395 0.45686087
## ALEC.Med
                 -0.154089591 5.116237e-03 8.045704e-02 17.010121 0.55539543
## DEG1
                 -3.988000000 -4.639000e-04 3.431257e-03 3.752479 0.05436765
## DEG2
                 -0.179622098 2.567885e-05 1.243010e-02 4.439505 0.31567660
## DEG5
                 -0.179622098 1.893438e-04 4.524440e-02 12.650031 0.97898936
## SumLFC
                 -0.179622098 6.335976e-04 8.105122e-02 13.300521 1.66126441
##
                   maximum
                            variance
                                             sd
                                                   range
## EC10.CTB.PHH
                  314.6036
                            1659.3283 40.73485
                                                314.6114
## EC20.CTB.PHH
                  363.5386 2314.1995 48.10613 363.5455
## EC50.CTB.PHH
                  498.2800 7197.3277 84.83707 498.2854
## EC10.CTB.HepG2
                 217.5594
                            897.5322 29.95884 217.6427
## EC20.CTB.HepG2
                 306.9219
                            1733.6636 41.63729
                                                306.9236
## EC50.CTB.HepG2 519.1888
                            4481.5361 66.94428 519.1898
## EC10.Nile.Red
                  694.4848
                            8596.1514 92.71543 695.1347
## EC20.Nile.Red
                 4999.8920 256391.9112 506.35157 4999.9386
## EC50.Nile.Red 4999.9942 790395.6977 889.04201 5000.0176
## EC10.Hoechst
               401.1805
                            4020.9579 63.41102 401.2876
## EC20.Hoechst
                802.3669 13699.9147 117.04663 802.3911
## EC50.Hoechst 4999.9942 494158.0284 702.96375 4999.9957
## EC10.CMFDA
                 4999.8920 425957.9696 652.65456 4999.9077
## EC20.CMFDA
                 4999.8920 425953.6678 652.65126 4999.9054
## EC50.CMFDA
               4999.8920 425816.8599 652.54644 4999.8979
## ALOEC.Min
                             290.2402 17.03644 146.0400
                139.8920
## ALOEC.Med
                1399.8920 19651.5080 140.18384 1399.9418
## ALEC.Min
                 499.1267
                           2504.6202 50.04618 509.0889
## ALEC.Q25
                 548.6841
                            5508.1717
                                      74.21706 548.8542
## ALEC.Med
                  853.6160
                            9807.2029 99.03132 853.7701
## DEG1
                            300.2056 17.32644 143.8800
                 139.8920
## DEG2
                 139.8920
                            328.5098 18.12484 140.0716
                  492.8520
## DEG5
                            4443.3733 66.65863 493.0316
## SumLFC
                  499.1267
                            4543.6748 67.40679 499.3063
```

Boxplots

```
# Convert to long format
long_data <- as.data.frame(X_normalized) %>%
    pivot_longer(cols = everything(), names_to = "Variable", values_to = "Value")
# Ensure 'Variable' is a factor and ordered based on its appearance in the dataset
long_data$Variable <- factor(long_data$Variable, levels = colnames(X))

# Create boxplots for all numeric columns in one chart
ggplot(long_data, aes(x = Variable, y = Value)) +
    geom_boxplot(fill = "lightblue", outlier.color = "red") +
    theme_bw() +
    theme(axis.text.x = element_text(angle = 45, hjust = 1)) +
    labs(title = "Boxplots", x = "Variables", y = "Values")</pre>
```

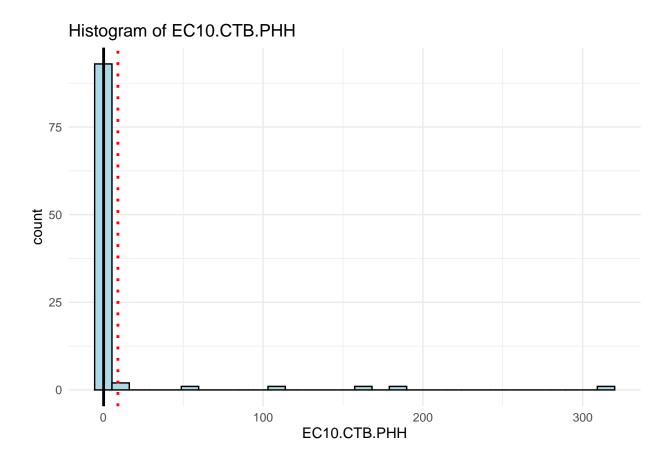


Histograms

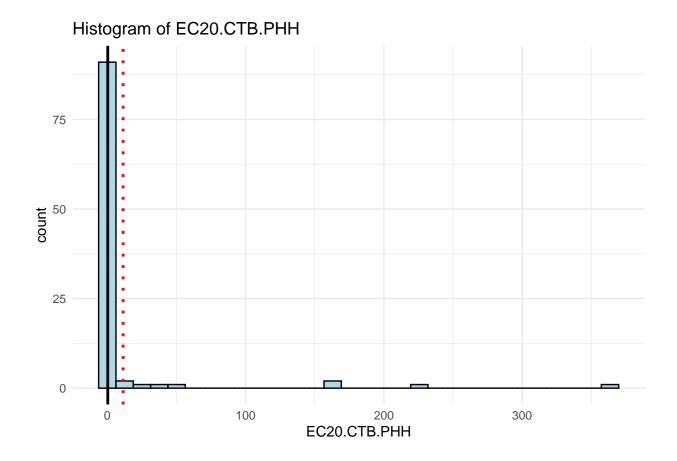
```
# Loop through each column of the dataset 'X'
for (i in 1:ncol(X_normalized)) {
  # Get the column name
  col_name <- colnames(X_normalized)[i]</pre>
  # Extract the mean and median for the current column from summary_df
  mean_val <- normalized_summary_df[col_name, "mean"]</pre>
  median_val <- normalized_summary_df[col_name, "median"]</pre>
  # Create the histogram and add vertical lines for mean and median
  gg <- ggplot(X_normalized, aes(x = get(col_name))) +</pre>
    geom_histogram( fill = "lightblue", color = "black") +
    geom_vline(aes(xintercept = mean_val), linetype = "dotted", color = "red", size = 1) + # Dotted red
    geom_vline(aes(xintercept = median_val), linetype = "solid", color = "black", size = 1) + # Solid b
    xlab(col_name) +
    theme_minimal() +
    labs(title = paste("Histogram of", col_name))
  # Print the plot
  print(gg)
```

```
## Warning: Using 'size' aesthetic for lines was deprecated in ggplot2 3.4.0.
## i Please use 'linewidth' instead.
## This warning is displayed once every 8 hours.
## Call 'lifecycle::last_lifecycle_warnings()' to see where this warning was
## generated.
```

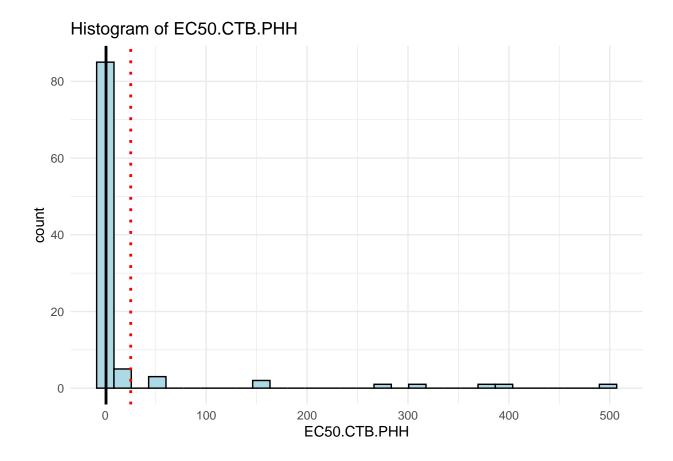
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



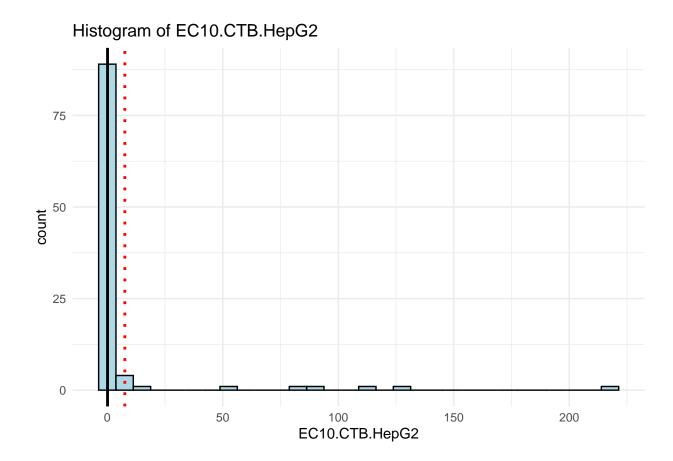
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



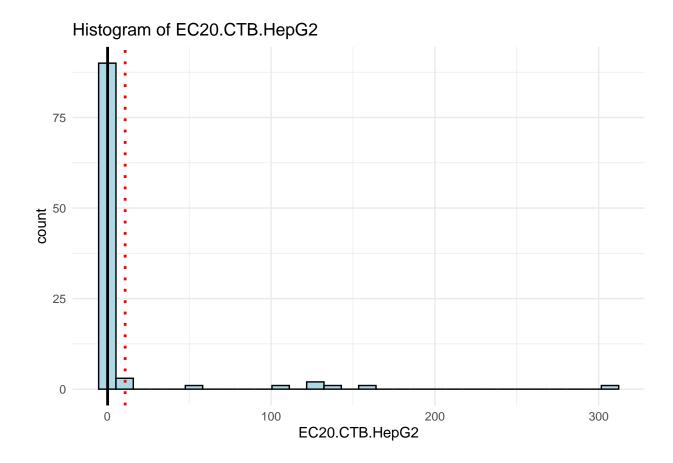
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



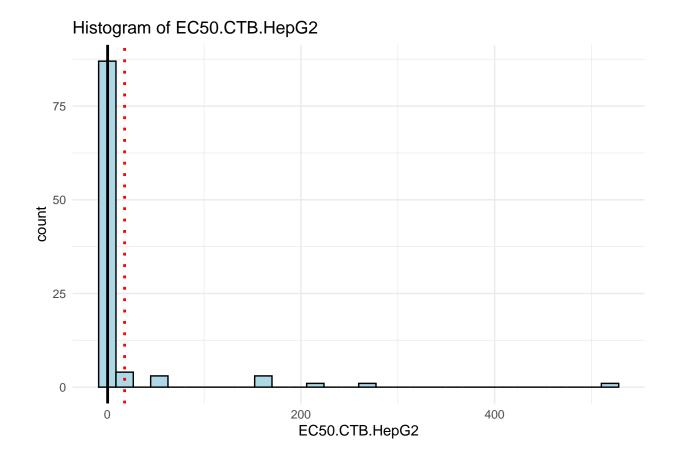
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



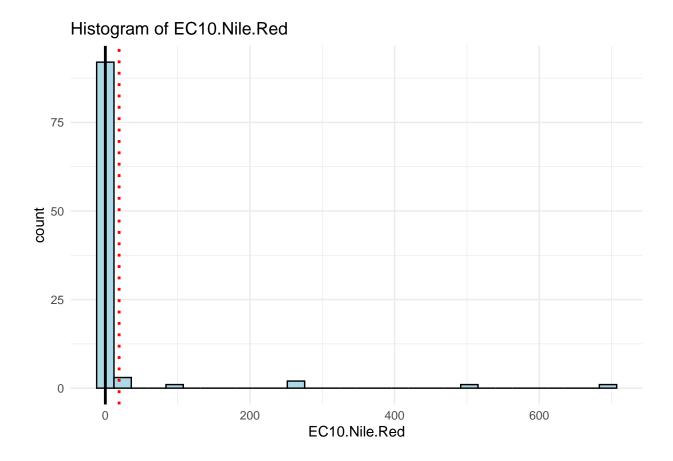
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



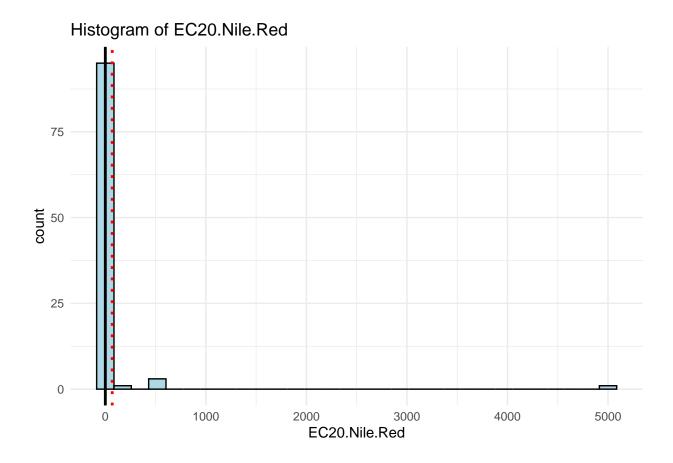
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



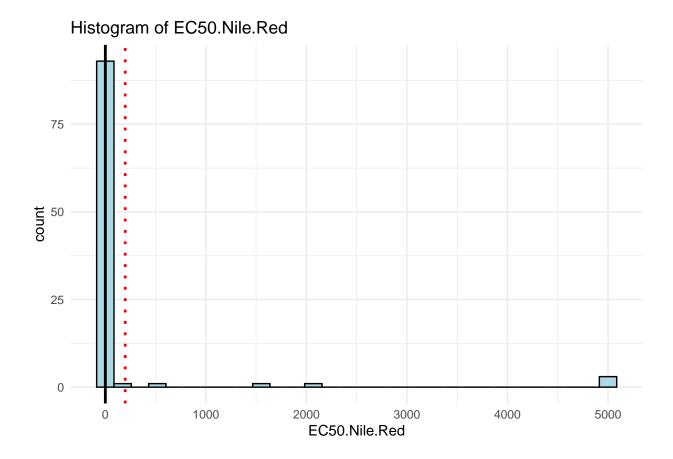
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



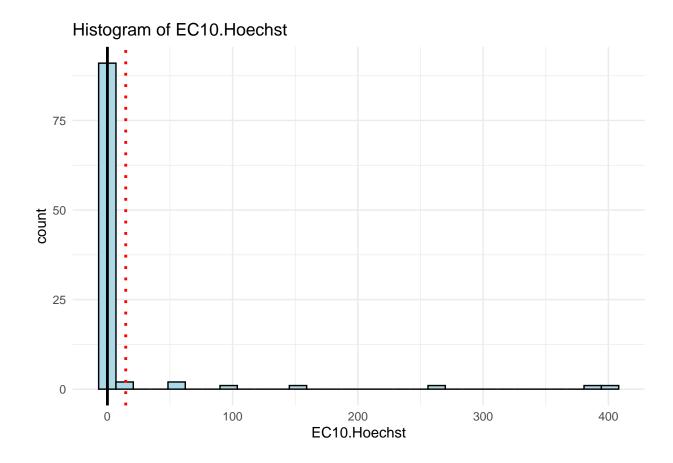
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



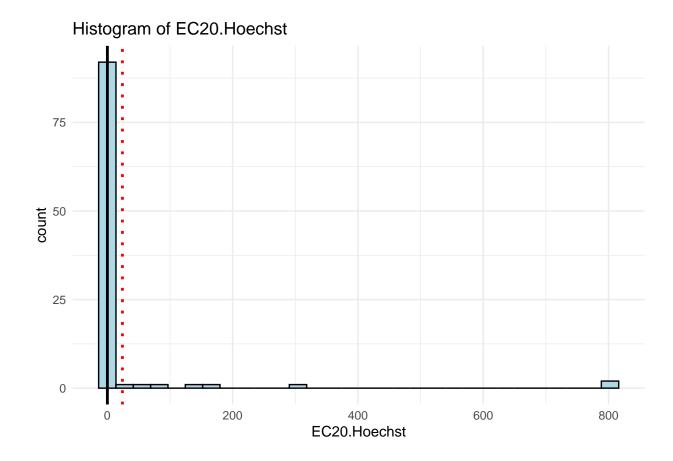
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



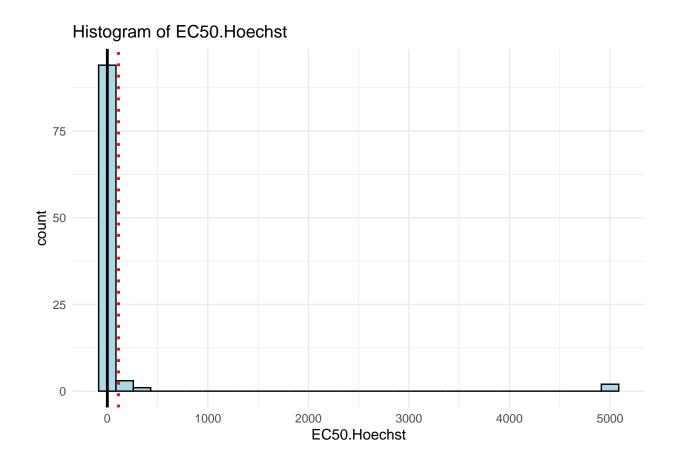
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



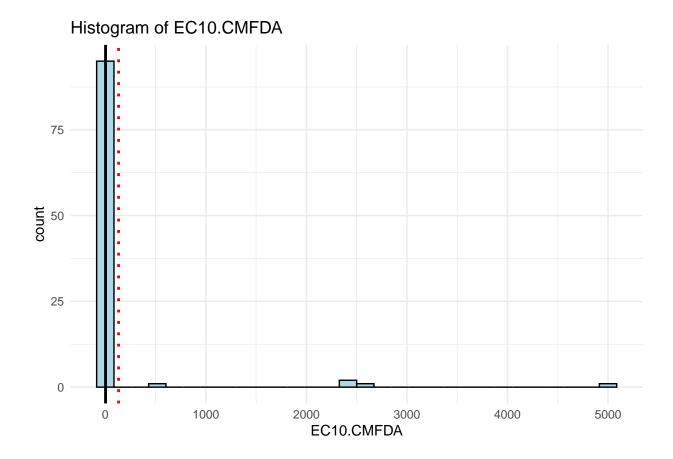
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



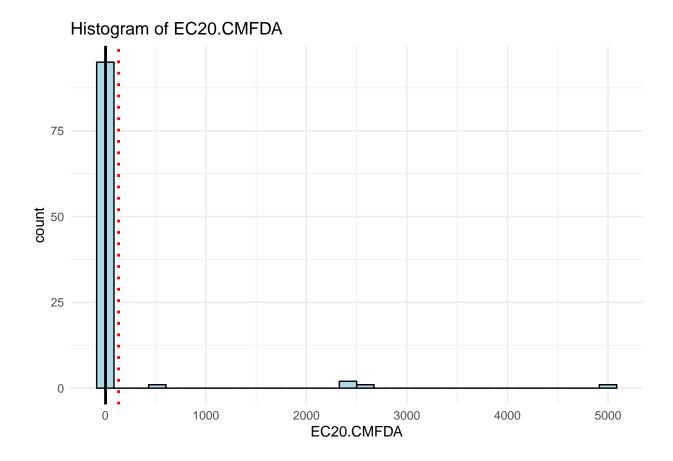
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



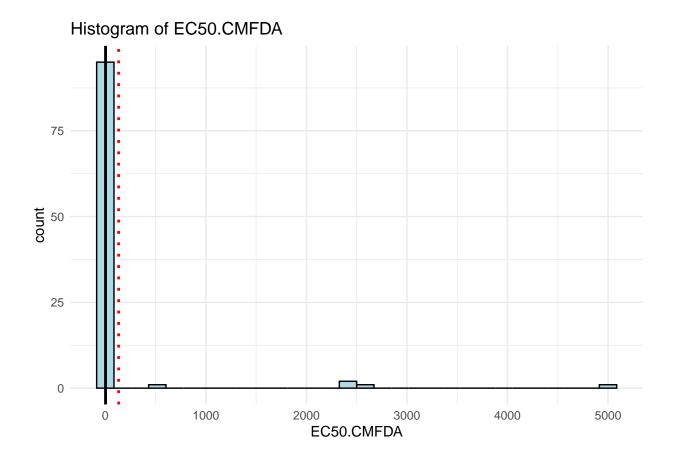
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



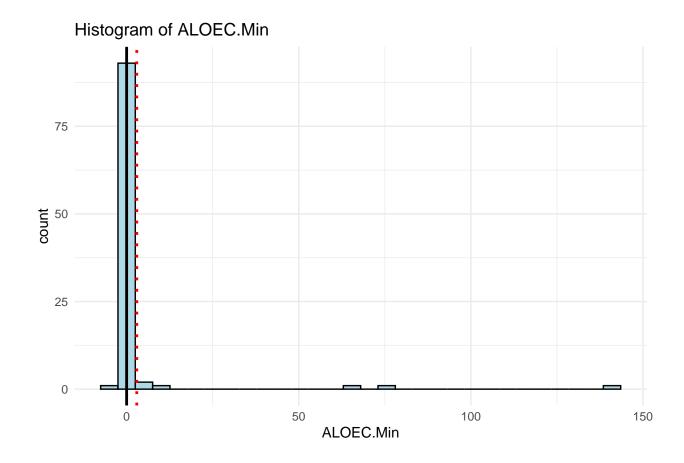
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



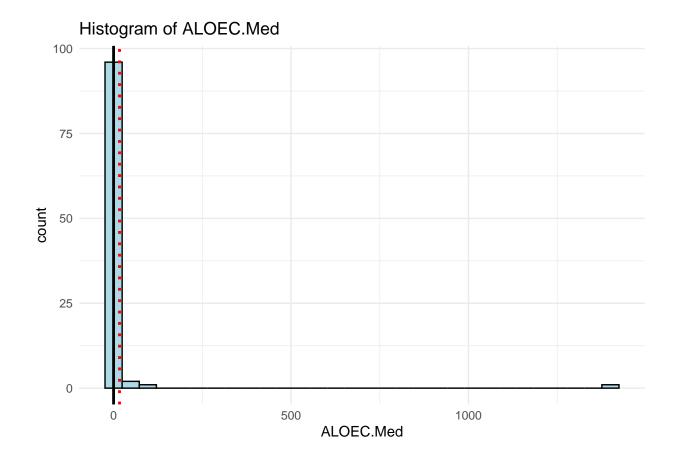
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



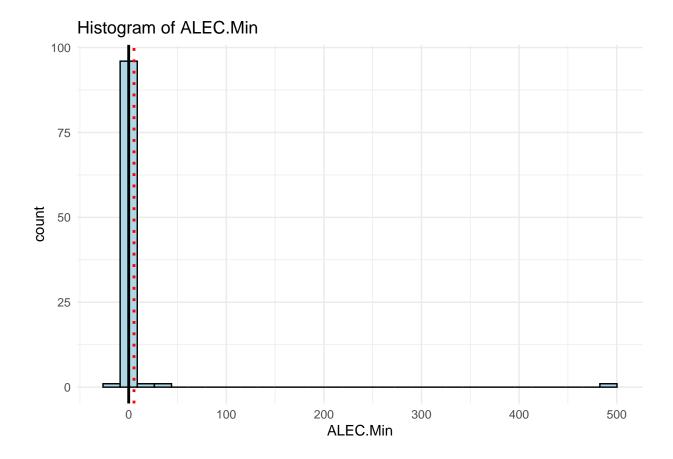
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



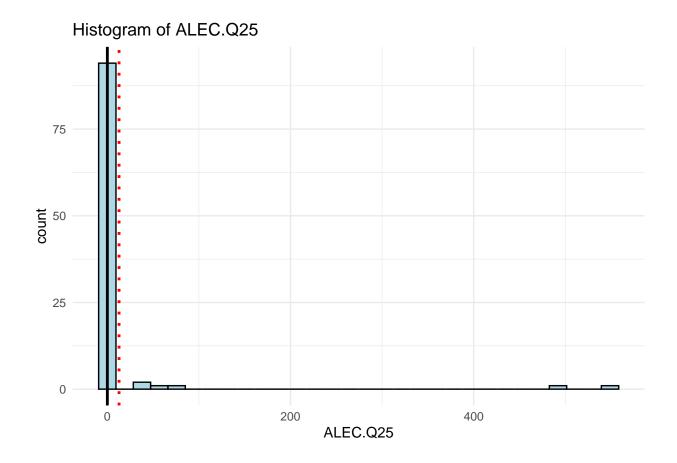
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



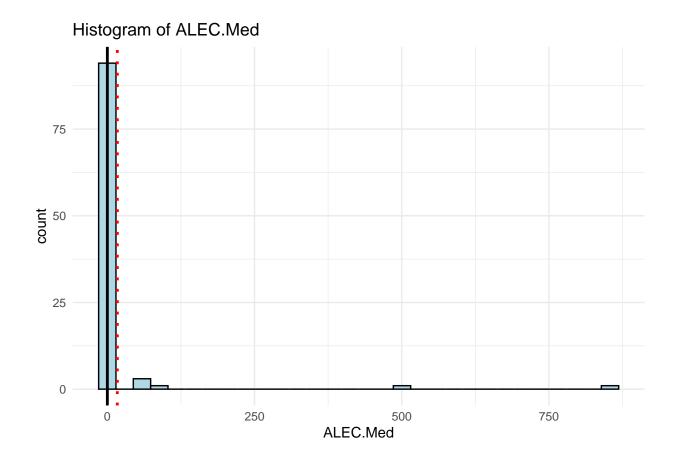
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



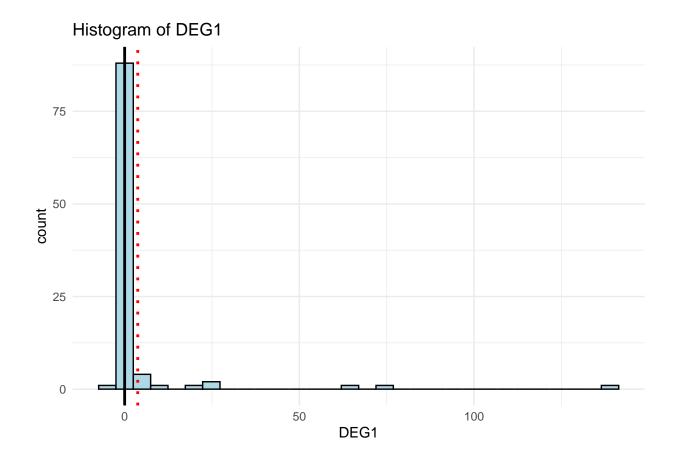
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



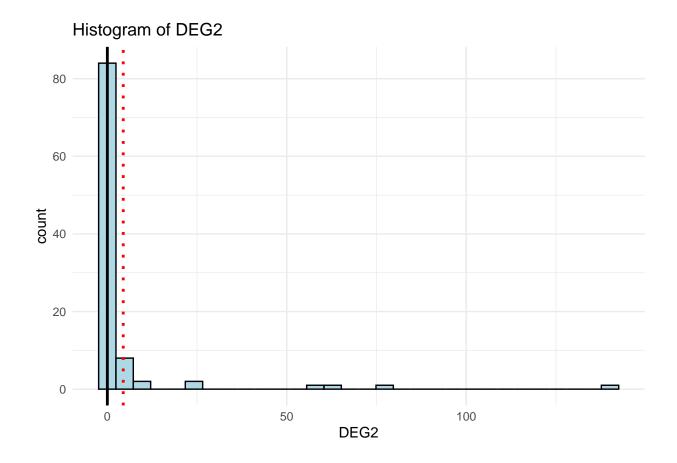
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



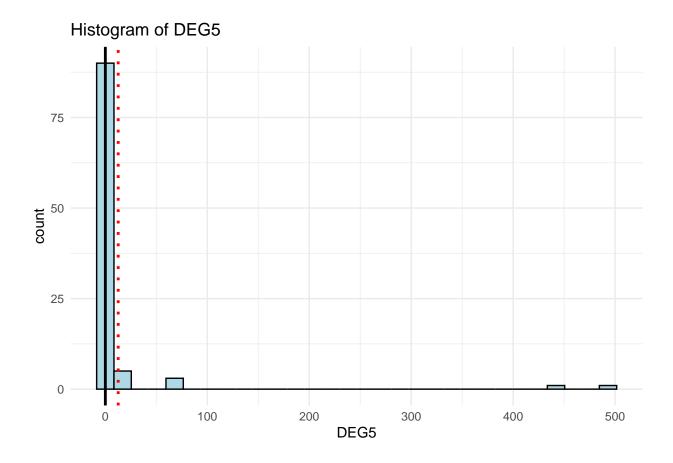
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



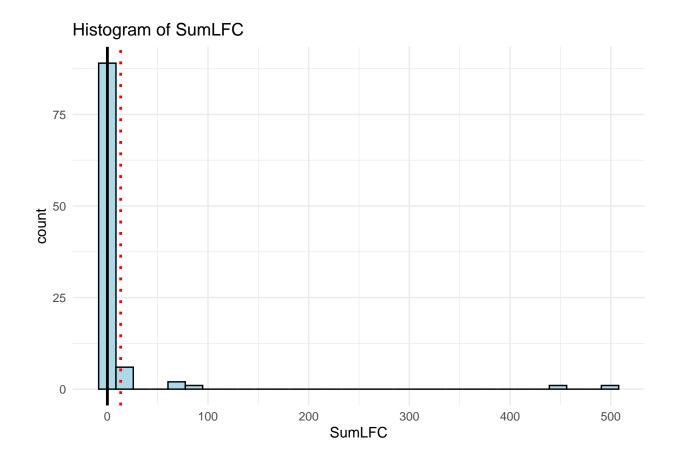
'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



'stat_bin()' using 'bins = 30'. Pick better value with 'binwidth'.



Bivariate analysis

```
cor_matrix4n <- cor(X_normalized)
# Find pairs of variables with correlation > 0.9
high_cor4n <- which(abs(cor_matrix4n) > 0.9, arr.ind = TRUE)
high_cor4n <- high_cor4n[high_cor4n[,1]!=high_cor4n[,2],]

# Get the variable names and their correlation values
high_cor_pairs4n <- data.frame(
    var1 = rownames(cor_matrix4n)[high_cor4n[, 1]],
    var2 = colnames(cor_matrix4n)[high_cor4n[, 2]],
    correlation = cor_matrix4n[high_cor4n]
)

# Remove duplicate pairs (since correlation matrix is symmetric)
high_cor_pairs4n <- high_cor_pairs4n[!duplicated(t(apply(high_cor_pairs4n, 1, sort))), ]

# Sort the pairs by correlation in descending order
high_cor_pairs4n <- high_cor_pairs4n[order(-abs(high_cor_pairs4n$correlation)), ]

# Display the sorted pairs of variables with high correlation
print(high_cor_pairs4n)</pre>
```

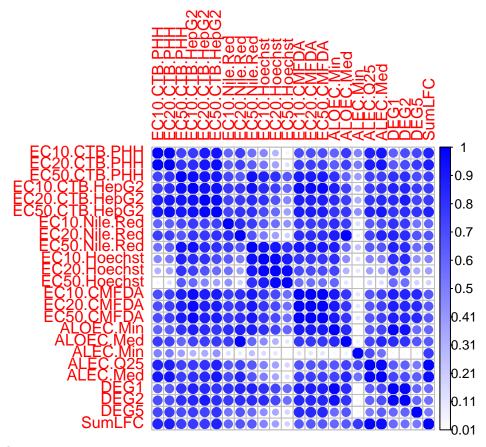
```
##
                                var2 correlation
                var1
## 45
                                        1.0000000
          EC20.CMFDA
                          EC10.CMFDA
                                        0.9999723
## 51
          EC50.CMFDA
                          EC20.CMFDA
## 46
          EC50.CMFDA
                          EC10.CMFDA
                                        0.9999713
## 1
        EC20.CTB.PHH
                       EC10.CTB.PHH
                                       0.9918579
## 29
                      EC20.Nile.Red
           ALOEC.Med
                                       0.9914217
## 65
                            ALEC.Q25
              SumLFC
                                        0.9893673
## 38
        EC20.Hoechst
                        EC10. Hoechst
                                        0.9774411
## 61
                DEG1
                           ALOEC.Min
                                        0.9763288
## 64
            ALEC.Med
                            ALEC.Q25
                                        0.9763105
## 72
                DEG2
                                DEG1
                                        0.9718249
## 22
      EC50.CTB.HepG2 EC20.CTB.HepG2
                                       0.9697559
## 41
        EC50.Hoechst
                        EC20.Hoechst
                                        0.9683496
                                        0.9603678
## 18
          EC50.CMFDA EC10.CTB.HepG2
## 17
          EC20.CMFDA EC10.CTB.HepG2
                                        0.9603066
## 16
          EC10.CMFDA EC10.CTB.HepG2
                                        0.9603056
                        EC20.CTB.PHH
## 6
      EC50.CTB.HepG2
                                        0.9559083
## 8
      EC10.CTB.HepG2
                        EC50.CTB.PHH
                                        0.9482806
              SumLFC
## 70
                            ALEC.Med
                                        0.9477058
## 32
        EC10.Hoechst
                      EC50.Nile.Red
                                        0.9430167
## 14 EC50.CTB.HepG2 EC10.CTB.HepG2
                                        0.9393316
       EC50.Nile.Red EC10.CTB.HepG2
                                        0.9366937
## 13 EC20.CTB.HepG2 EC10.CTB.HepG2
                                        0.9365660
       EC50.Nile.Red
                        EC50.CTB.PHH
## 11
                                        0.9337509
## 62
                DEG2
                           ALOEC.Min
                                        0.9334396
      EC50.CTB.HepG2
## 2
                        EC10.CTB.PHH
                                        0.9248035
## 9
      EC20.CTB.HepG2
                        EC50.CTB.PHH
                                        0.9246008
## 7
            ALEC.Med
                        EC20.CTB.PHH
                                        0.9222193
## 33
        EC20.Hoechst
                       EC50.Nile.Red
                                        0.9154158
## 5
      EC20.CTB.HepG2
                        EC20.CTB.PHH
                                        0.9149542
## 3
            ALEC.Med
                        EC10.CTB.PHH
                                        0.9136802
## 10 EC50.CTB.HepG2
                        EC50.CTB.PHH
                                        0.9130623
## 28
            ALEC.Med EC50.CTB.HepG2
                                        0.9112898
## 36
          EC50.CMFDA
                      EC50.Nile.Red
                                        0.9106312
## 35
          EC20.CMFDA
                      EC50.Nile.Red
                                        0.9102698
## 34
                      EC50.Nile.Red
          EC10.CMFDA
                                        0.9102636
## 52
           ALOEC.Min
                          EC20.CMFDA
                                        0.9032864
## 47
           ALOEC.Min
                          EC10.CMFDA
                                        0.9032862
## 57
           ALOEC.Min
                          EC50.CMFDA
                                        0.9032669
```

```
print(length(high_cor_pairs4n[,1]))
```

[1] 38

corrplot

```
corrplot(cor matrix4n, is.corr=FALSE, col=colorRampPalette(c("white", "blue"))(200) )
```



scatterplots

Scatterplot between tartet and explanatory variables

```
# Create pairwise scatter plots colored by 'Toxicity'
#ggpairs(cbind(X_normalized, Toxicity = y, aes(fill = Toxicity, alpha = 0.7), progress = FALSE) +
# theme_bw() +
#labs(title = "Pairwise Scatter Plots for Predictor Variables (Colored by Toxicity)")
```

Scatterplots between Toxicity and other variables

```
cols =colnames(X_normalized)
#cols = df_aucs$variable # in the order of sorted auc
plot_list = list()
for (i in 2:ncol(X_normalized)) {
   colname <- cols[i]
   p <- ggplot(X_normalized, aes(x = .data[[colname]], y)) +
        geom_point() +
        labs(x = colname, y = "Toxicity") +
        theme_minimal()
#print(p)
# Add each plot to the list
plot_list[[i-1]] <- p</pre>
```

```
# Arrange the plots in a grid (choose the number of rows/columns as needed)
grid.arrange(grobs = plot_list, ncol = 2)
```

