

## 13 ExpoStatLink module

### 13.1 Application domain

ExpoStatLink module is designed to find the statistical relationships between exposure factors and health outcome.

### 13.2 Theory

### 13.3 Work pipeline

Users can easily get the modeling results and their visualization plots with high quality by following the detailed instructions in each step. The statistical interpretations from the perspectives of subjects and whole dataset are both provided.

```
# The following two packages should be installed in advance
# devtools::install_github("ExposomeX/exstat", force = TRUE)
# devtools::install_github("ExposomeX/extidy", force = TRUE)

# library(exstat)
# library(extidy)
library(tidyverse)

# devtools::install_github("ExposomeX/exposomex", force = TRUE)
library(exposomex)
```

```
res = InitStatLink()
```

```
res1 = LoadStatLink(PID=res$PID,
                    UseExample="example#1")
res1$Expo$Data
```

```
## # A tibble: 20 x 14
##   SampleID Subjec~1 Y1 Y2 X1 X2 X3 X4 X5 X6 X7 X8
##   <chr> <chr> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
## 1 Tr1 S1 1 -101 7.76 10.2 24.2 1.47 1.06 1.43 2.00 1.87
## 2 Tr2 S2 0 -51 10.1 11.6 3.39 1.03 1.46 1.32 1.69 1.18
## 3 Tr3 S3 0 -37 8.54 9.52 15.8 1.23 1.42 1.64 1.52 1.95
## 4 Tr4 S4 1 -61 14.2 14.9 11.3 NA 1.75 1.23 1.91 1.29
## 5 Tr5 S5 0 -28 11.0 16.4 1.43 1.31 1.70 1.69 1.04 1.68
## 6 Tr6 S6 0 -8 11.3 12.7 8.37 1.29 1.25 1.38 1.26 1.17
## 7 Tr7 S7 1 -63 7.66 7.73 17.1 1.49 1.19 1.83 1.46 1.08
## 8 Tr8 S8 0 -35 11.3 8.25 13.3 1.48 1.91 1.07 1.30 1.09
## 9 Tr9 S9 0 -14 14.5 11.1 11.4 1.29 NA 1.30 1.46 1.23
## 10 Tr10 S10 1 -99 6.26 6.04 8.03 1.92 1.53 1.08 1.92 1.85
## 11 Tr11 S11 0 -60 3.43 10.3 8.45 1.76 1.79 1.13 1.73 1.07
## 12 Tr12 S12 0 -32 6.75 6.86 28.0 1.03 1.10 1.58 1.98 1.06
## 13 Tr13 S13 0 -73 7.71 9.47 22.3 NA 1.20 1.34 1.45 1.53
## 14 Tr14 S14 0 -18 12.6 8.97 9.23 1.23 1.13 NA 1.62 1.58
## 15 Tr15 S15 0 -48 11.7 8.60 9.31 1.20 1.33 1.99 1.85 1.94
## 16 Tr16 S16 0 -20 7.10 12.5 2.52 1.83 1.19 1.65 1.47 1.48
## 17 Tr17 S17 0 -9 6.32 49.1 8.52 1.26 1.94 1.53 1.95 1.18
## 18 Tr18 S18 1 -98 4.66 8.49 24.2 1.07 1.25 1.35 1.70 1.98
## 19 Tr19 S19 0 -70 2.44 -2.27 16.0 1.43 1.77 1.21 1.87 1.48
## 20 Tr20 S20 0 -36 10.6 7.07 10.0 1.99 1.61 1.65 1.84 1.94
## # ... with 2 more variables: X9 <dbl>, X10 <dbl>, and abbreviated variable name
```

```
## # 1: SubjectID
```

```
res1$Expo$Voca
```

```
## # A tibble: 12 x 5
##   SerialNo SerialNo_Raw FullName GroupName Lod
##   <chr>      <chr>      <chr>    <chr>    <dbl>
## 1 Y1        Y1          Y_disc Outcome    NA
## 2 Y2        Y2          Y_cont Outcome    NA
## 3 X1        X1          TE_1     Chemical  0.5
## 4 X2        X2          TE_2     Chemical  0.5
## 5 X3        X3          TE_3     Chemical  0.5
## 6 X4        X4          TE_4     Chemical  0.5
## 7 X5        X5          TE_5     Chemical  0.5
## 8 X6        X6          TE_6     Chemical  0.5
## 9 X7        X7          TE_7     Chemical  0.5
## 10 X8       X8          TE_8     Chemical  0.5
## 11 X9       X9          CH1      Chemical  5
## 12 X10     X10          CH2      Chemical  5
```

```
res2 = StatLinkCros(PID=res$PID,
                    OutPath = "default",
                    VarsY = "Y1" ,
                    VarsX = "all.x" ,
                    LinkModel = "ranger",
                    ObsrPartType = "raw" ,
                    ObsrPartNum = "50",
                    ObsrProfType = "partial" ,
                    ObsrProfNum = "100",
                    ObsrProfVars = "all.x" ,
                    ObsrProfGeom = "profiles",
                    SubjPredSeq = "S1,S2,S3",
                    SubjPartType = "break_down")
```

```
res2$CrosVipPlot
```

```
## $Y1_importance_ranger_raw
## [[1]]
## TableGrob (1 x 1) "arrange": 1 grobs
##   z      cells   name      grob
## 1 1 (1-1,1-1) arrange gtable[layout]
##
## attr("class")
## [1] "arrangelist" "list"
```

```
res2$CrosProfPlot
```

```
## $Y1_ranger_partial_profiles_by_X1
## [[1]]
## TableGrob (1 x 1) "arrange": 1 grobs
##   z      cells   name      grob
## 1 1 (1-1,1-1) arrange gtable[layout]
##
## attr("class")
## [1] "arrangelist" "list"
##
## $Y1_ranger_partial_profiles_by_X2
```

```

## [[1]]
## TableGrob (1 x 1) "arrange": 1 grobs
##   z      cells      name      grob
## 1 1 (1-1,1-1) arrange gtable[layout]
##
## attr("class")
## [1] "arrangelist" "list"
##
## $Y1_ranger_partial_profiles_by_X3
## [[1]]
## TableGrob (1 x 1) "arrange": 1 grobs
##   z      cells      name      grob
## 1 1 (1-1,1-1) arrange gtable[layout]
##
## attr("class")
## [1] "arrangelist" "list"
##
## $Y1_ranger_partial_profiles_by_X4
## [[1]]
## TableGrob (1 x 1) "arrange": 1 grobs
##   z      cells      name      grob
## 1 1 (1-1,1-1) arrange gtable[layout]
##
## attr("class")
## [1] "arrangelist" "list"
##
## $Y1_ranger_partial_profiles_by_X5
## [[1]]
## TableGrob (1 x 1) "arrange": 1 grobs
##   z      cells      name      grob
## 1 1 (1-1,1-1) arrange gtable[layout]
##
## attr("class")
## [1] "arrangelist" "list"
##
## $Y1_ranger_partial_profiles_by_X6
## [[1]]
## TableGrob (1 x 1) "arrange": 1 grobs
##   z      cells      name      grob
## 1 1 (1-1,1-1) arrange gtable[layout]
##
## attr("class")
## [1] "arrangelist" "list"
##
## $Y1_ranger_partial_profiles_by_X7
## [[1]]
## TableGrob (1 x 1) "arrange": 1 grobs
##   z      cells      name      grob
## 1 1 (1-1,1-1) arrange gtable[layout]
##
## attr("class")
## [1] "arrangelist" "list"
##
## $Y1_ranger_partial_profiles_by_X8

```

```

## [[1]]
## TableGrob (1 x 1) "arrange": 1 grobs
##   z      cells   name      grob
## 1 1 (1-1,1-1) arrange gtable[layout]
##
## attr("class")
## [1] "arrangelist" "list"
##
## $Y1_ranger_partial_profiles_by_X9
## [[1]]
## TableGrob (1 x 1) "arrange": 1 grobs
##   z      cells   name      grob
## 1 1 (1-1,1-1) arrange gtable[layout]
##
## attr("class")
## [1] "arrangelist" "list"
##
## $Y1_ranger_partial_profiles_by_X10
## [[1]]
## TableGrob (1 x 1) "arrange": 1 grobs
##   z      cells   name      grob
## 1 1 (1-1,1-1) arrange gtable[layout]
##
## attr("class")
## [1] "arrangelist" "list"
res2$CrosSubjPlot$Y1_ranger_break_down_S1

## [[1]]
## TableGrob (1 x 1) "arrange": 1 grobs
##   z      cells   name      grob
## 1 1 (1-1,1-1) arrange gtable[layout]
##
## attr("class")
## [1] "arrangelist" "list"
FuncExit(PID = res$PID)

## [1] "Success to exit. Thanks for using ExposomeX platform!"

```