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Package

HTSanalyzeR2 0.99.11

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1 Introduction

In this short tutorial, we give a detailed illustration for the shiny report part of **HTSanalyzeR2** to visualize the results and modify figures in different aspects.

2 Interactive Shiny report visualization

2.1 Visualize single GSCA object for individual data set

For single data set analysis, after analyzed by HTSanalyzeR2, we can get a *GSCA* object and use the function *report* to launch the shiny report.

```
data(d7_gsca)
report(d7_gsca)
```

2.1.1 The hypergeometric test result table[Figure 1]

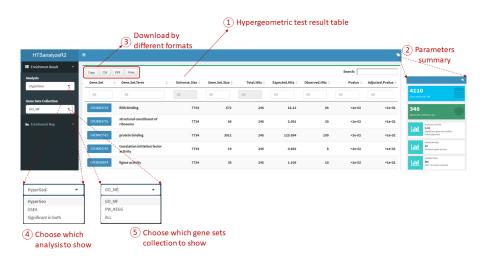


Figure 1: The hypergeometric test result table

2.1.2 Parameters for modifying the figure[Figure2]

There are five main parts for users to modify their figure by interactive operations including: Layout, Label, Node, Edge and Color Scheme. Following is a detailed explanation for each part.

- Layout: A widge to control the whole pattern
 - Mode: There are two layout modes inside this Shiny report, users can choose any of them to get better visualization based on their data.
 - Parameters: Two more parameters Egde Repel and Adjust Sizes could be used to make the figure looser.

- Gravity: Ranging from -50 to 50, users can adjust it step by step. The larger it is, the more even the pattern would be.
- **Label**: A widge to modify the label of nodes with three easy-to-understand parameters: *Text, Label Scale* and *Label color*.
- **Node**: A widge to modify the nodes with four easy-to-understand parameters: *Scale*, *Opacity*, *Border Width* and *Border Color*.
- **Edge**: A widge to modify the edges with two easy-to-understand parameters: *Edge Scale* and *Edge Color*.
- Color Scheme: A widge to change the default color.

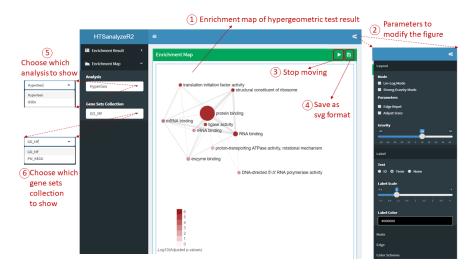


Figure 2: The enrichment map of hypergeometric test result

2.2 Visualize single *NWA* object for individual data set

```
data(d7_nwa)
report(d7_nwa)
```

2.2.1 The identified subnetwork[Figure3]

2.3 Visualize a list of *GSCA* objects for time series/comparative analysis

```
data(gscaTS)
## To make the figure more compact, we set a cutoff
## to move any other edges with low Jaccard coefficient.
reportAll(gscaTS, cutoff = 0.035)
```

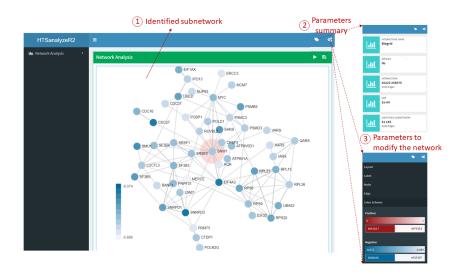


Figure 3: The identified subnetwork

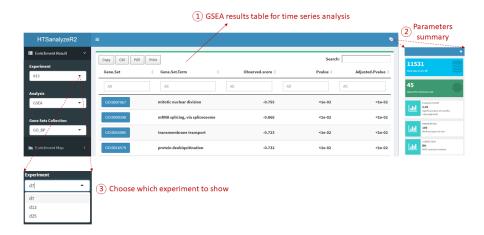


Figure 4: GSEA results table for time series/comparative analysis

- 2.3.1 GSEA results table for time series/comparative analysis[Figure4]
- 2.3.2 Union enrichment map for time series/comparative analysis[Figure5]
- 2.4 Visualize a list of *NWA* objects for time series/comparative analysis

```
data(nwaTS)
reportAll(nwaTS)
```

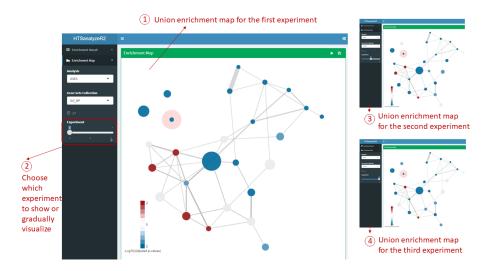


Figure 5: Union enrichment map for time series/comparative analysis

2.4.1 Union subnetwork for time series/comparative analysis[Figure6]

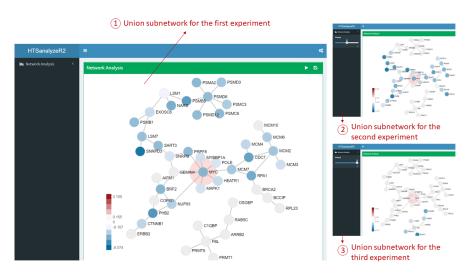


Figure 6: Union subnetwork for time series/comparative analysis

2.5 Visualize both GSCA and NWA objects simultaneously

reportAll(gsca = gscaTS, nwa = nwaTS)

2.5.1 Visualize both GSEA and netwrok result in the same report[Figure7]

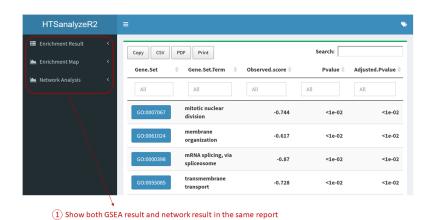


Figure 7: Visualize both GSEA and netwrok result in the same report

3 Session Info

```
## R version 3.4.2 (2017-09-28)
## Platform: x86_64-pc-linux-gnu (64-bit)
## Running under: Debian GNU/Linux 9 (stretch)
## Matrix products: default
## BLAS/LAPACK: /usr/lib/libopenblasp-r0.2.19.so
##
## locale:
## [1] LC_CTYPE=en_US.UTF-8
                                  LC_NUMERIC=C
## [3] LC_TIME=en_US.UTF-8
                                  LC_COLLATE=en_US.UTF-8
## [5] LC_MONETARY=en_US.UTF-8
                                  LC_MESSAGES=C
## [7] LC_PAPER=en_US.UTF-8
                                  LC_NAME=C
## [9] LC_ADDRESS=C
                                  LC_TELEPHONE=C
## [11] LC_MEASUREMENT=en_US.UTF-8 LC_IDENTIFICATION=C
## attached base packages:
## [1] stats
             graphics grDevices utils
                                              datasets methods
## other attached packages:
## [1] BiocStyle_2.6.0
##
## loaded via a namespace (and not attached):
## [1] compiler_3.4.2 backports_1.1.1 bookdown_0.5
                                                      magrittr_1.5
## [5] rprojroot_1.2 tools_3.4.2 htmltools_0.3.6 yaml_2.1.16
                                     rmarkdown_1.8
## [9] Rcpp_0.12.16 stringi_1.1.7
                                                      knitr_{-}1.17
## [13] stringr_1.3.0 digest_0.6.15
                                      evaluate_0.10.1
```