

# Package ‘toxSummary’

February 27, 2024

**Title** Visualize and Summarize Repeat-Dose Toxicology Study Results

**Version** 1.0.0

**Description** A 'shiny' app that generates plots and summary tables from repeat-dose toxicology study results to facilitate holistic evaluation of the drug safety of active pharmaceutical ingredients (API) prior to initiation of clinical trials.

**License** MIT + file LICENSE

**URL** <https://github.com/phuse-org/toxSummary>

**BugReports** <https://github.com/phuse-org/toxSummary/issues>

**Encoding** UTF-8

**RoxygenNote** 7.2.1

**Imports** cicerone, data.table, DBI, dplyr, DT, flextable, fs, ggiraph, ggplot2, htmltools, magrittr, patchwork, rlang, RSQLite, scales, shiny, shinycssloaders, shinydisconnect, stringr

**Depends** R (>= 2.10)

**Suggests** knitr, officer, rmarkdown

**VignetteBuilder** knitr

**NeedsCompilation** no

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**Repository** CRAN

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toxSummary_app	<i>Run toxSummary app</i>
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**Description**

Run toxSummary app

**Usage**

```
toxSummary_app(  
  database_path = NULL,  
  studyid_file = NULL,  
  save_file_path = NULL,  
  where_to_run = "local"  
)
```

**Arguments**

database_path	Optional, character file path for database
studyid_file	Optional, character file path for studyid
save_file_path	optional, character directory where files will be saved when app create any files. if NULL, all the files will be created in temp directory.
where_to_run	Optional, character where app will be running. Default is "local". Should use "rsconnect" when deploy to rsconnect.

**Value**

function run the app.

**Examples**

```
if(interactive()) {  
  toxSummary_app()  
}
```

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