

# Package: EpiLinx (via r-universe)

January 30, 2025

**Title** Interactive Visualization Tool for Nosocomial Outbreaks

**Version** 0.2.0-1

**Date** 2025-01-30

**Description** What the package does (one paragraph).

**License** GPL (>= 3)

**Encoding** UTF-8

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.3.2

**Depends** R (>= 4.1.0)

**Imports** stats, utils, lubridate, dplyr, tidyverse, ggraph, igraph,  
reshape2, shiny, ggnetwork, sna, shinydashboard,  
shinycssloaders, DT, shinyWidgets, shinyjs, shinyBS

**Suggests** knitr, rmarkdown, spelling, testthat (>= 3.0.0)

**Config/testthat/edition** 3

**VignetteBuilder** knitr

**Language** en-GB

**LazyData** true

**Config/pak/sysreqs** libfontconfig1-dev libfreetype6-dev libfribidi-dev  
libglpk-dev make libharfbuzz-dev libicu-dev libjpeg-dev  
libpng-dev libtiff-dev libxml2-dev libssl-dev libx11-dev  
zlib1g-dev

**Repository** <https://ku-awdc.r-universe.dev>

**RemoteUrl** <https://github.com/ku-awdc/EpiLinx>

**RemoteRef** v0.2.0

**RemoteSha** 8c4bbb4dd9cd82f0065c2119b207333e55befd3d

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launch_app	<i>Launch the EpiLinx app</i>
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**Description**

Launch the EpiLinx app

**Usage**

```
launch_app()
```

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test_long_2021	<i>Example dataset</i>
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**Description**

Some fake data to play around with

**Usage**

```
test_long_2021
```

**Format**

A data frame with:

**SSI\_id** Anonymised ID ...

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## \* datasets

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