

# Package ‘rTRMui’

May 16, 2025

**Type** Package

**Title** A shiny user interface for rTRM

**Version** 1.47.0

**Date** 2015-09-17

**Author** Diego Diez

**Imports** shiny (>= 0.9), rTRM, MotifDb, org.Hs.eg.db, org.Mm.eg.db

**Maintainer** Diego Diez <diego10ruiz@gmail.com>

**Description** This package provides a web interface to compute transcriptional regulatory modules with rTRM.

**License** GPL-3

**LazyLoad** yes

**ByteCompile** yes

**biocViews** Transcription, Network, GeneRegulation, GraphAndNetwork, GUI

**URL** <https://github.com/ddiez/rTRMui>

**BugReports** <https://github.com/ddiez/rTRMui/issues>

**git\_url** <https://git.bioconductor.org/packages/rTRMui>

**git\_branch** devel

**git\_last\_commit** 041cef4

**git\_last\_commit\_date** 2025-04-15

**Repository** Bioconductor 3.22

**Date/Publication** 2025-05-15

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rTRMui-package

*rTRM shiny UI*

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### Description

A web-based GUI for the rTRM package implemented with the package shiny.

### Details

Package: rTRMui  
Type: Package  
Version: 0.1  
Date: 2013-04-07  
License: GPL-3

### Author(s)

Diego Diez

Maintainer: Diego Diez <diego10ruiz@gmail.com>

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runTRM

*Runs an rTRM (shiny) web server.*

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### Description

This command runs a local web server (constructed with the shiny package), which implements a GUI for the rTRM package.

### Usage

```
runTRM()
```

### Author(s)

Diego Diez

### See Also

[runApp](#)

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