

A vignette for redirecting to the meshr package

Koki Tsuyuzaki^{1,4}, Gota Morota², Takeru Nakazato³ and Itoshi Nikaido⁴.

October 27, 2020

¹Department of Medical and Life Science, Tokyo University of Science.

²Department of Animal Sciences, University of Wisconsin-Madison.

³Database Center for Life Science, Research Organization of Information and Systems.

⁴Bioinformatics Research Unit, RIKEN Advanced Center for Computing and Communication.

`k.t.the-answer@hotmail.co.jp`, `dritoshi@gmail.com`

Description for any MeSH-related packages is written in the vignette of *meshr* package. Please follow the link below

<http://www.bioconductor.org/packages/release/bioc/html/meshr.html>

or just type

```
> if(!requireNamespace("BiocManager", quietly=TRUE)) > install.packages("BiocManager")
> BiocManager::install("meshr")
> library("meshr")
> vignette("MeSH")
```

in R console window.