

# A vignette for redirecting to the meshr package

Koki Tsuyuzaki<sup>1,4</sup>, Gota Morota<sup>2</sup>, Takeru Nakazato<sup>3</sup> and Itoshi Nikaido<sup>4</sup>.

October 30, 2018

<sup>1</sup>Department of Medical and Life Science, Tokyo University of Science.

<sup>2</sup>Department of Animal Sciences, University of Wisconsin-Madison.

<sup>3</sup>Database Center for Life Science, Research Organization of Information and Systems.

<sup>4</sup>Bioinformatics 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.