

# A vignette for redirecting to the meshr package

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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

```
> source("http://bioconductor.org/biocLite.R")
> biocLite("meshr")
> library("meshr")
> vignette("MeSH")
```

in R console window.