

# R

Note: R client library available from version 4 onwards

R library package is implemented and maintained for the latest R release and distributed through Bioconductor (<https://bioconductor.org/packages/release/bioc/html/cellbaseR.html>). For a quick install, please enter the R terminal and type:

```
source("https://bioconductor.org/biocLite.R")
biocLite("cellbaseR")
```

The R code is also distributed together with the rest of the CellBase code. R code can be found at: `cellbase/clients/R`

The R code provides a library for programmatic access to CellBase data. No CLI is implemented in R.

Comprehensive description of the library is provided by the Bioconductor documentation:

<https://bioconductor.org/packages/release/bioc/html/cellbaseR.html>

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