

# A vignette for redirecting to the scTensor package

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Description for any LRBase-related packages is written in the vignette of *scTensor* package.

Please follow the link below

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

or just type

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

in R console window.