

# Package ‘RSeqAn’

November 1, 2025

**Type** Package

**Title** R SeqAn

**Version** 1.30.0

**Description** Headers and some wrapper functions from the SeqAn C++ library for ease of usage in R.

**License** BSD\_3\_clause + file LICENSE

**BugReports** <https://github.com/compbiocore/RSeqAn/issues>

**LinkingTo** Rcpp

**Imports** Rcpp

**Suggests** knitr, rmarkdown, testthat

**VignetteBuilder** knitr

**biocViews** Infrastructure, Software

**RoxygenNote** 6.1.0

**git\_url** <https://git.bioconductor.org/packages/RSeqAn>

**git\_branch** RELEASE\_3\_22

**git\_last\_commit** 67ea6d9

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**Repository** Bioconductor 3.22

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|                  |                         |
|------------------|-------------------------|
| containment_test | <i>containment test</i> |
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**Description**

containment test

**Usage**

containment\_test(s1)

**Arguments**

|    |                              |
|----|------------------------------|
| s1 | Should be a character string |
|----|------------------------------|

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