

# Package ‘nanotubes’

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**Type** Package

**Title** Mouse nanotube CAGE data

**Version** 1.25.0

**Author** Malte Thodberg

**Maintainer** Malte Thodberg <maltethodberg@gmail.com>

**Description** Cap Analysis of Gene Expression (CAGE) data from ``Identification of Gene Transcription Start Sites and Enhancers Responding to Pulmonary Carbon Nanotube Exposure in Vivo" by Bornholdt et al. supplied as CAGE Transcription Start Sites (CTSSs).

**URL** <https://github.com/MalteThodberg/nanotubes>

**BugReports** <https://github.com/MalteThodberg/nanotubes/issues>

**License** GPL-3

**Encoding** UTF-8

**LazyData** FALSE

**Depends** R (>= 3.6)

**Suggests** knitr, rmarkdown, BiocStyle, GenomicRanges, rtracklayer, CAGEfightR

**RoxygenNote** 6.1.1

**VignetteBuilder** knitr

**biocViews** ExperimentData, SequencingData, ExpressionData

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

**git\_branch** devel

**git\_last\_commit** bf1b39c

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