Package 'gUtils'

January 20, 2018

Title R Package Providing Additional Capabilities and Speed for
GenomicRanges Operations

Version 0.2.0

Description R package providing additional capabilities and speed for GenomicRanges operations.

Depends R (>= 3.1.0), GenomicRanges (>= 1.18), data.table (>= 1.9)

Imports IRanges (>= 2.0), S4Vectors (>= 0.4), GenomeInfoDb (>= 1.2),
parallel, BiocGenerics(>= 0.12), methods, Matrix, stringr

Suggests BSgenome.Hsapiens.UCSC.hg19, testthat, rtracklayer

License GPL-2

BugReports <http://github.com/mskilab/gUtils/issues>

LazyData true

RoxygenNote 6.0.1.9000

NeedsCompilation no

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R topics documented: