Package ‘qcmetrics’

May 4, 2024

Type Package

Title A Framework for Quality Control

Version 1.42.0

Description The package provides a framework for generic quality control of data. It permits to create, manage and visualise individual or sets of quality control metrics and generate quality control reports in various formats.

Depends R (>= 3.3)

Imports Biobase, methods, knitr, tools, xtable, pander, S4Vectors

Suggests affy, MSnbase, ggplot2, lattice, mzR, BiocStyle, rmarkdown, markdown

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URL http://lgatto.github.io/qcmetrics/articles/qcmetrics.html

BugReports https://github.com/lgatto/qcmetrics/issues

biocViews ImmunoOncology, Software, QualityControl, Proteomics, Microarray, MassSpectrometry, Visualization, ReportWriting

VignetteBuilder knitr

RoxygenNote 6.1.0

git_url https://git.bioconductor.org/packages/qcmetrics

git_branch RELEASE_3_19

git_last_commit 8dae2f7

git_last_commit_date 2024-04-30

Repository Bioconductor 3.19

Date/Publication 2024-05-03

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Description

A simple wrapper for the QC of 15N labelling. The respective QC items are the distribution of PSM incorporation rates, distribution of log2 fold-changes and number of identified features. See the vignette for details.

Usage

n15qc(object,
  fcol = c("Protein_Accession", "Peptide_Sequence", "Number_Of_Unique_Peptides", "Variable_Modifications", "inc"),
  inctr = 97.5, lfctr = c(-0.5, 0.5), type, reportname)

Arguments

- **object**: An MSnSet to be quality controlled.
- **fcol**: The name of the feature variables for the protein identifiers (accession numbers for example), the peptide sequences, the number of unique peptides for each identified protein, the variable modifications identified on the peptides and the N15 incorporation rate. These must be provided in that order. Defaults are Protein_Accession, Peptide_Sequence, Number_Of_Unique_Peptides, Variable_Modifications, and inc.
- **inctr**: The minimum level of median incorporation rate to set the QC item status to TRUE. Default is 97.5.
- **lfctr**: The range of accepted median PSM log2 fold-change for the QC item status to be set to TRUE. Default is c(-0.5, 0.5).
- **type**: The type of report to be saved. If missing (default), no report is generated. See qcReport for details.
- **reportname**: The name of the report, in case a type is defined. If missing (default), the report will be names n15qcreport followed by the generation data and time.

Value

Invisibly returns the resulting QcMetrics instance.
Description

An example data for 15N metabolic labelling, distributed as an MSnSet to illustrate 15N QC.

Usage

data(n15psm)

References

See MSnSet and the MSnbase-demo vignette in the MSnbase package.

Examples

library("MSnbase")
data(n15psm)
psm

Qc2Tex 'QcMetric' sectioning functions

Description

These functions convert the $i$th QcMetric instance of the QcMetrics object into a section of the adequate format, i.e. TeX of R markdown.

Usage

Qc2Tex(object, i)
Qc2Tex2(object, i)
Qc2Tex3(object, i)
Qc2Rmd(object, i)

Arguments

object An instance of class QcMetrics with at least one QC item.
i A numeric of length 1 indicating the index of the item to be converted into text section.
Value

A character representing the QC item section.

Author(s)

Laurent Gatto

See Also

qcReport and the vignette.

QcMetadata-class  The "QcMetadata" class

Description

The QcMetadata class is a simple interface to metadata. The objects can be displayed with show for a summary and print for the content.

Objects from the Class

Objects can be created by calls of the form QcMetadata(...).

Slots

metadata: Object of class "list" that stores the metadata variables. The list must be named. NA and empty characters are not allowed.

Methods

[ signature(x = "QcMetadata"): subsets x as a new QcMetadata instance.
[[ signature(x = "QcMetadata"): extracts a single element of x.
 metadata signature(x = "QcMetadata"): return the object’s metadata list. Also available as mdata.
 metadata<- signature(x = "QcMetadata", value = "list"): sets the objects metadata. Also available as mdata.
 length signature(x = "QcMetadata"): returns the number of metadata variables.
 names signature(x = "QcMetadata"): returns the names of the metadata variables.

Author(s)

Laurent Gatto
Examples

QcMetadata(metadata =
  list(name = "John Doe",
       lab = "Big Lab in Big Uni"))
## less typing
qmd <- QcMetadata(list(name = "John Doe",
                        lab = "Big Lab in Big Uni"))
mdata(qmd)
show(qmd)
print(qmd)

QcMetric-class

The "QcMetric" class for QC items

Description

Data structure for individual QC metrics items.

Objects from the Class

Objects can be created using the constructor QcMetric(...), where slots are assigned individually. See example below.

Slots

name: Object of class "character" of length 1 naming the object.
description: Object of class "character" of arbitrary length describing the qc metric in more details.
qcdata: Object of class "environment" that stores the actual data.
plot: Object of class "function" to graphically represent the data and infer quality status.
show: Object of class "function" to print a short textual representation of the object. A reasonable default value is provided.
status: Object of class "logical" that indicates weather the data passes (TRUE) or fails (FALSE) the metric or has not yet been evaluated.

Methods

name signature(object = "QcMetric"): retrieves the name of the object.
name<- signature(object = "QcMetric", value = "character"): set the name of the object.
description signature(object = "QcMetric"): retrieves the description of the object.
description<- signature(object = "QcMetric", value = "character"): set the description of the object.
status signature(object = "QcMetric"): retrieves the status of the object.
status<- signature(object = "QcMetric", value = "logical"): sets the status of the objects.
**qcdada** signature(object = "QcMetric", x = "missing"): lists all the data objects that are associated with the objects.

**qcdada** signature(object = "QcMetric", x = "character"): retrieves the variable x for the object.

**qcdada<-** signature(object = "QcMetric", var): creates or overwrites (with a message) the data variable var by assigning the RHS value. If var is missing and the RHS expression is an environment, then qcdada is reset with all the variables in value.

**qceenv** signature(object = "QcMetric"): return the environment that stores the QC data.

**qceenv<-** signature(object = "QcMetric"): Set all variable in the RHS environment as qcdada variables. Equivalent to qcdada(object) <- x where x is an environment.

**show** signature(object = "QcMetric"): shows a textual summary of object. The default show implementation is available as the qcshow(object, qcdada) function. The second argument is a logical (default is TRUE) that specifies whether qcdada(object) should be displayed.

**show<-** signature(object = "QcMetric", value = "function"): sets a custom show method for object.

**plot** signature(x = "QcMetric", y = "missing"): plots the object using the provide show method.

**plot<-** signature(object = "QcMetric", value = "function"): sets a custom plot method for object.

**qcReport** signature(x = "QcMetric", ...): to generate quality reports. See qcReport for details.

**Author(s)**

Laurent Gatto

**See Also**

The QcMetrics class to bundle a set of QcMetric instances.

**Examples**

```r
(qc <- QcMetric())
qcdada(qc)
try(qcdada(qc, "x"))

x <- rnorm(10)
qcdada(qc, "qc1") <- x
qcdada(qc, "qc2") <- 1:10
qcdada(qc)
all.equal(qcdada(qc, "qc1"), x)
all.equal(qcdada(qc, "qc2"), 1:10)
name(qc) <- "My test QcMetric"
description(qc) <- "This qc metric describes bla bla bla, indicating possible issues in the third step of protocol bla bla bla."
status(qc) <- FALSE
cq
cq

# or
e <- new.env()
e$qc1 <- rnorm(100)
```
e$qc2 <- 1:100
cqevn(qc) <- e
length(qcdata(qc, "qc1"))
head(qcdata(qc, "qc2"))

show(qc)
show(qc) <- function(object) cat("Updated show method\n")
show(qc)
show(qc) <- qcshow
qc

plot(qc)
plot(qc) <-
  function(object, ...)
    plot(qcdata(object, "qc2"),
         qcdata(object, "qc1"),
         xlab = "qc1",
         ylab = "qc2",
         ...
    )
plot(qc)
plot(qc, col = "red", pch = 19)

---

The "QcMetrics" class for collections of QC items

Description

Data structure for storing lists of QcMetric items.

Objects from the Class

Objects can be created using the constructor QcMetrics(...), where slots are assigned individually. See example below.

In a standardised quality control pipeline, the QcMetrics and QcMetric object are not generated manually. Their creation is delegated to a wrapper function that reads a specific type of files, parses the data, produces the individual QcMetric instances and, eventually, the QcMetric object. See the package vignette for details and examples.

Slots

metadata: Object of class QcMetadata storing the metadata of the object. This list would typically contain the input file the data was read from, the date the object was generated, ... or fully fledged minimum information descriptions (see MIaxE), when available.

qcdata: Object of class "list" storing all the individual QcMetric instances.
Methods

signature(x = "QcMetrics"): subsets x as a new QcMetrics instance.
signature(x = "QcMetrics"): extracts a single QcMetric instance.
length signature(x = "QcMetrics"): returns the number of QcMetric instances populate x.
metadata signature(x = "QcMetrics"): return the object's metadata list. Also available as mdata.
metadata<- signature(x = "QcMetrics", value = "list"): sets the objects metadata. Also available as mdata.
metadata<- signature(x = "QcMetric", value = "QcMetadata"): sets the objects metadata. Also available as mdata.
name signature(object = "QcMetrics"): returns a character vector of length length(object) with the names of the QcMetric instances.
qcdata signature(object = "QcMetrics", x = "missing"): returns a list of all QcMetric instances.
qcdata<- signature(object = "QcMetrics", value = "list"): sets the qcdata of object.
show signature(object = "QcMetrics"): prints a short textual description of object.
status signature(object = "QcMetrics"): returns a vector of quality statuses (logicals).
status<- signature(object = "QcMetrics", value = "logical"): sets the quality statuses. Length of value and object must be identical.
as signature(object = "QcMetrics", "data.frame"): coerces object as a length(object) by 2 data frame with the respective QcMetric instances names and statuses.
qcReport signature(object = "QcMetrics"): ...

Author(s)
Laurent Gatto

See Also
The QcMetric class for individual QC items.

Examples

example(QcMetric)
show(qc)

qc2 <- QcMetric(name = "My other metric", status = TRUE)
qcdata(qc2, "x") <- rnorm(100)
qcdata(qc2, "k") <- rep(LETTERS[1:2], 50)

plot(qc2) <- function(object, ...) {
  require("lattice")
  d <- data.frame(x = qcdata(object, "x"),
                  k = qcdata(object, "k"))
  bwplot(x ~ k, data = d)
}
qcReport-methods

Automatic report generation

Description

The `qcReport` method generates report in various formats taking a `QcMetrics` instance as input. Each individual quality control item produces a section with description of the item and a assessment figure.

Details

The reporting functions take a `QcMetrics` instance as input, generate the source of the report and compile it into the final format that are currently available are `reporting_pdf`, `reporting_tex`, `reporting_rmd` and `reporting_html`. See `qcto` for details about the sectioning functions, that convert individual `QcMetric` objects into adequate report sections.

The package vignette documents the report generation in more details and describes possibilities for customisation.

Methods

```r
signature(object = "QcMetrics", reportname = "character", type = "character", author = "character", title = "character", ...)
```

generates a report for the `QcMetrics` object. The report will be named according the `reportname` (default is `qcreprt`) and `type`, the latter defining the output format and the extension. Possible types are `pdf` (default), `tex`, `Rmd`, `html` (all generated using the package `knitr`). A custom title can be provided; default is "Quality control report generated with qcmetrics". If no author is provided, the default value (`Sys.getenv("USER")`) is used. The addition of a table of contents (default is `FALSE`), a metadata section, a summary section and the session information can be controlled with the `toc`, `metadata`, `summary` and `sessioninformation` arguments. The metadata section is added to the report when present and the other have `TRUE` as default.
Intermediate files are deleted, unless clean is set to FALSE and verbose output can be turned on by setting quiet to FALSE.

The reporter and qcto arguments are used to convert QcMetric and QcMetrics objects into report source. See Details and the package vignette for details.

Addition parameters can be passed to inner functions. For the pdf report, passed to texi2pdf; for html, passed to markdown::markdownToHTML.

The method invisibly returns the name of the report that was generated.

Examples

```r
example(QcMetrics)
show(qcm)

destdir <- tempdir()
(report <- file.path(destdir, "testQCReport"))

## Not run:
## pdf report
qcReport(qcm, reportname = report)
## use pdflatex to generate the pdf file
qcReport(qcm, reportname = report, texi2dvi = "pdflatex")

## End(Not run)

## default html report
html <- qcReport(qcm, reportname = report, type = "html")
html
if (interactive())
  browseURL(html)
```
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