

# Package ‘MSstatsQCgui’

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**Version** 1.26.0

**Type** Package

**Title** A graphical user interface for MSstatsQC package

**Description** MSstatsQCgui is a Shiny app which provides longitudinal system suitability monitoring and quality control tools for proteomic experiments.

**License** Artistic License 2.0

**LazyData** TRUE

**URL** <http://msstats.org/msstatsqc>

**BugReports** <https://groups.google.com/forum/#!forum/msstatsqc>

**RoxygenNote** 7.1.0

**Imports** shiny, MSstatsQC, ggExtra, gridExtra, plotly, dplyr, grid

**Suggests** knitr

**VignetteBuilder** knitr

**biocViews** Software, QualityControl, Proteomics, MassSpectrometry, GUI

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

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`RunMSstatsQC`*Shiny connection for example datasets*

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**Description**

Shiny connection for example datasets

**Usage**

```
RunMSstatsQC()
```

**Value**

A connection for the shiny interface

**Examples**

```
## Not run: An example dataset can be found through MSstatsQC package
library(MSstatsQC)
data <- MSstatsQC::S9Site54
## Not run: RunMSstatsQC()
```

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`S9Site54`*CPTAC study 9.1 site 54 dataset*

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**Description**

system suitability testing results generated during CPTAC Study 9.1 for Site 54

**Usage**

```
data(S9Site54)
```

**Format**

csv

**Details**

CPTAC system suitability testing data for Site 54 from Study 9.1

**Value**

An example dataset generated from CPTAC study 9.1

**References**

<http://www.mcponline.org/content/early/2015/02/18/mcp.M114.047050>

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