Package: IDSL.UFAx (via r-universe)

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Type Package

Title Exhaustive Chemical Enumeration for United Formula Annotation

Version 1.9.1

Depends R (>= 4.0)

Imports IDSL.IPA (>= 2.7), IDSL.UFA , readxl, RcppAlgos

Author Sadjad Fakouri-Baygi [aut] (<https://orcid.org/0000-0002-6864-6911>), Dinesh Barupal [cre, aut] (<https://orcid.org/0000-0002-9954-8628>)

Maintainer Dinesh Barupal <dinesh.barupal@mssm.edu>

Description A pipeline to annotate a number of peaks from the 'IDSL.IPA' peaklists using an exhaustive chemical enumeration-based approach. This package can perform elemental composition calculations using the following 15 elements : C, B, Br, Cl, K, S, Si, N, H, As, F, I, Na, O, and P.

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URL https://github.com/idslme/idsl.ufax

BugReports https://github.com/idslme/idsl.ufax/issues

Encoding UTF-8

Archs i386, x64

Repository https://idslme.r-universe.dev

RemoteUrl https://github.com/idslme/idsl.ufax

RemoteRef HEAD

RemoteSha 7d587efc246b7593cb155b53933343541350584e

Contents

UFAx_workflow *UFAx Workflow*

Description

This function runs the exhaustive chemical enumeration part of the IDSL.UFAx pipeline.

Usage

```
UFAx_workflow(spreadsheet)
```

Arguments

spreadsheet	IDSL.UFAx parameter spreadsheet

Value

The MolecularFormulaAnnotationTable is saved in the assigned folder in the parameter spreadsheet.

Note

You should load the IDSL.UFA package to run the IDSL.UFAx functions.

Examples

```
library(IDSL.UFAx)
s_path <- system.file("extdata", package = "IDSL.UFAx")
SSh1 <- paste0(s_path,"/UFAx_parameters.xlsx")
temp_wd <- tempdir() # update this address
temp_wd_zip <- paste0(temp_wd,"/003.mzML_UFAx_testfiles.zip")
spreadsheet <- readxl::read_xlsx(SSh1)
download.file(
  paste0("https://github.com/idslme/IDSL.UFAx/blob/main/UFAx_educational_files/",
         "003.mzML_UFAx_testfiles.zip?raw=true"), destfile = temp_wd_zip, mode = "wb")
unzip(temp_wd_zip, exdir = temp_wd)
spreadsheet[1, 4] <- temp_wd
spreadsheet[3, 4] <- temp_wd
spreadsheet[6, 4] <- temp_wd
spreadsheet[5, 4] <- "seq(1, 100, 1)" # peak IDs to process
UFAx_results <- UFAx_workflow(spreadsheet)
```

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