

# Additional Functions for PFAS Suspect List

## Function Description

For the analysis and useful application of the suspect list, additional functions were generated for R to process the dataset.

**Set up R environment** To open Microsoft Excel files, you must install the 'xlsx' package.

```
install.packages("xlsx")
```

```
source("fn/get_suspectlist.R")
get_suspectlist("PFAS Suspect List.xlsx")
# You can designate the list be saved as a different file
# name if you want.
```

**get\_suspectlist:** To download the most recent version of the PFAS suspect list, use this function.

```
source("fn/open_suspectlist.R")
suspectlist <- open_suspectlist("PFAS Suspect List.xlsx")
## use the file location if this is different
```

**open\_suspectlist:** Open the suspect list into the R environment (you must do this before any other steps)

```
source("fn/add_ionization.R")
suspectlist_w_ionization <- add_ionization(suspectlist, ionstate = c("M-H",
  "M-H2O+H", "M+H", "M+K", "M+Na"))
# These are all the possible ion states
write.csv(suspectlist_w_ionization, "suspectlist_w_ionization.csv",
  row.names = FALSE)
```

**add\_ionization:** calculates the exact m/z value for indicated ionization states.

```
source("fn/calculate_residual.R")
exactmasses <- readRDS("fn/exactmasses.RDS")
suspectlist_w_residuals <- add_residual(suspectlist, rep_unit = "CF2",
  exactmasses, adduct = "neutral")
# adduct can be 'neutral', 'M-H', 'M-H2O+H', 'M+H', 'M+K',
# 'M+Na', 'M+', or 'M-'
write.csv(suspectlist_w_residuals, "suspectlist_w_residuals.csv",
  row.names = FALSE)
```

***add\_residual***: calculates the residual mass after removing the repeating units (default = CF2)

```
source("fn/expand_additional.R")
suspectlist_w_additional <- expand_additional(suspectlist)
write.csv(suspectlist_w_additional, "suspectlist_w_additional.csv",
  row.names = FALSE)
```

***expand\_additional***: all delimited information stored in the ADDITIONAL column is expanded as additional columns. **NOTE:** There will be a significant number of new columns with NA values for most compounds.

```
source("fn/get_properties.R")
suspectlist_w_properties <- get_properties(suspectlist)
write.csv(suspectlist_w_properties, "suspectlist_w_properties.csv",
  row.names = FALSE)
```

***get\_properties***: using RDKit (<https://www.rdkit.org/>), calculates multiple 2D descriptors from the suspect list compounds. Requires the installation of RDKit and the use of the *tidychem* R Package. Definitions of the descriptors can be found at: <https://www.rdkit.org/docs/GettingStartedInPython.html#list-of-available-descriptors>

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