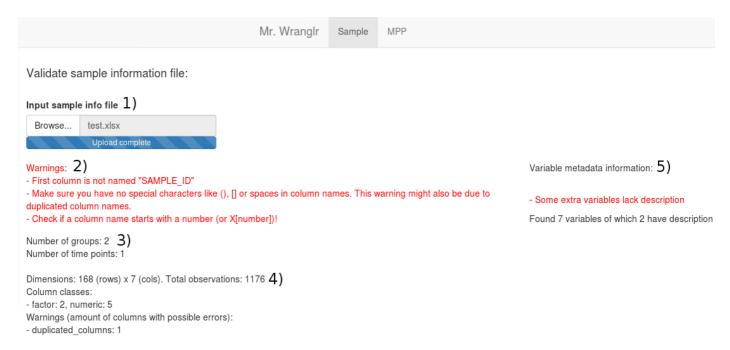
Wranglr instructions

Sample tab

Input sample information in a .xlsx file (see specification file). Wranglr will process the sample information file and produce files that are compatible with MPP and Worklist.

Wranglr provides useful information to check that your sample information file meets the requirements and that the file is read correctly.



- 1) Input Excel sheet as described in sample information form specification.
- 2) Wranglr displays warnings if your file does not meet the requirements. Any warnings here will prevent processing and must be dealt with.
- 3) If column(s) named GROUP and/or TIME are found, the number of groups and/or time points found will be displayed.
- 4) Some extra information on the sample information sheet. Warnings here do not prevent processing.
- 5) Information on the variable metadata. Warnings here do not prevent processing.

Randomize and add QC Project title: Demo Project home folder The complete path to project home folder. D:\MassHunter\Data\Demo\ Defaults to D:\MassHunter\Data\[project title]\ Sample run order Random Choose between original and randomized. Sample positioning type 96-well plate 54-vial plate Number of samples before every QC: Choose between multiple intervals, possible 12 to run without any QC. Modes ✓ HILIC neg ✓ HILIC pos ✓ RP neg ✓ RP pos Choose which modes will be run. Number of QC before beginning of mode: HILIC neg **HILIC** pos RP neg RP pos 10 10 10 Second column for MPP Choose the column used for grouping in MPP.

Note that the number of QC samples in the beginning of modes can change in course of analysis. In this case, instead of manually changing the worklist file, you can rerun the processing with corrected values. The randomization of run order calculates its seed number from the project title, so the run order will remain unchanged when the sample information file of any individual project is reprocessed.

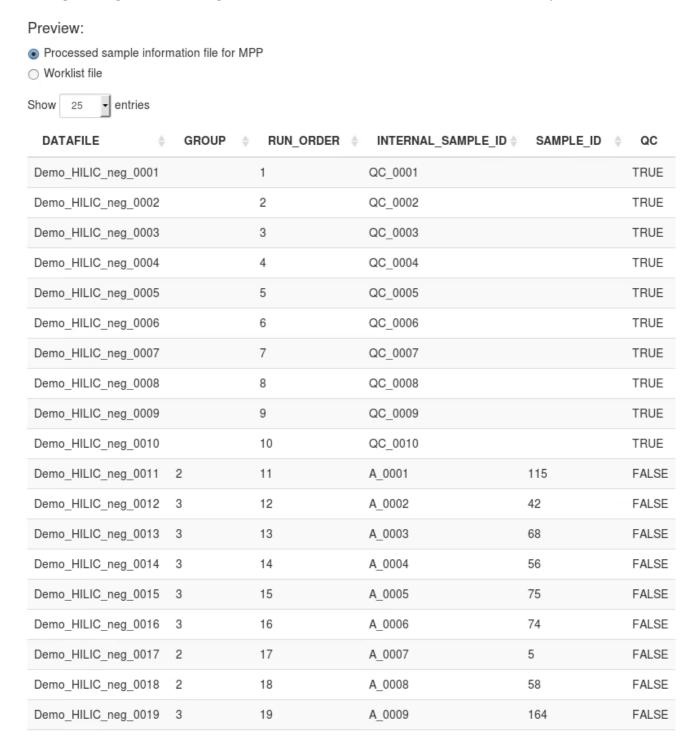
defaults to QC.

Defaults to GROUP column if found, else

QC

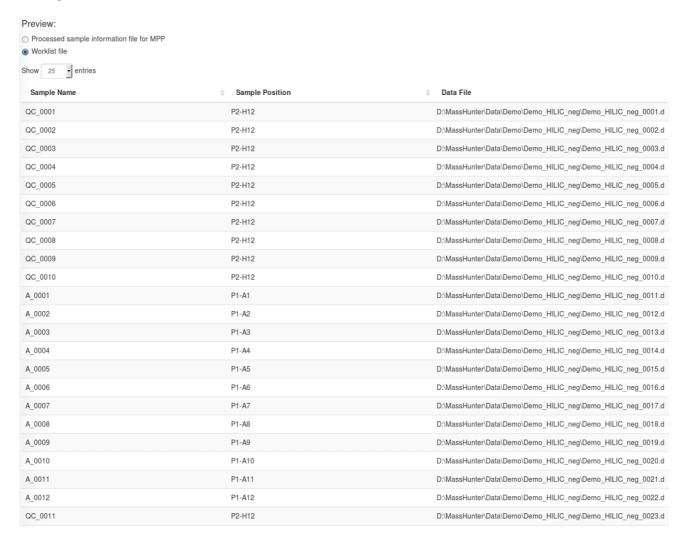
Submit & Process

Example of a processed sample information file used in MPP and further analysis:



Wranglr generates four new columns: DATAFILE, RUN_ORDER, INTERNAL_SAMPLE_ID and QC. The processed sample information file will also include all the columns of the original sample information file.

Example of a worklist file:



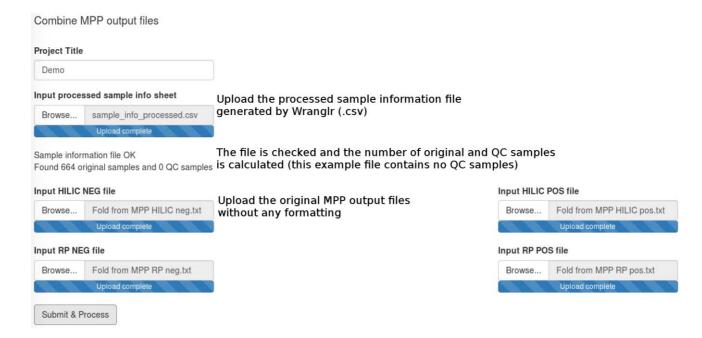
The Sample Name column is identical to INTERNAL_SAMPLE_ID column in the processed sample information file. Sample position is generated automatically. The last position on the second plate is reserved for QC samples. The Data File column includes the complete file paths. Wranglr will generate a subfolder for each mode.

An individual worklist file will be generated for each mode. All the files generated by Wranglr can be downloaded as .csv files. The original variable metadata sheet can also be downloaded as .csv file if needed.

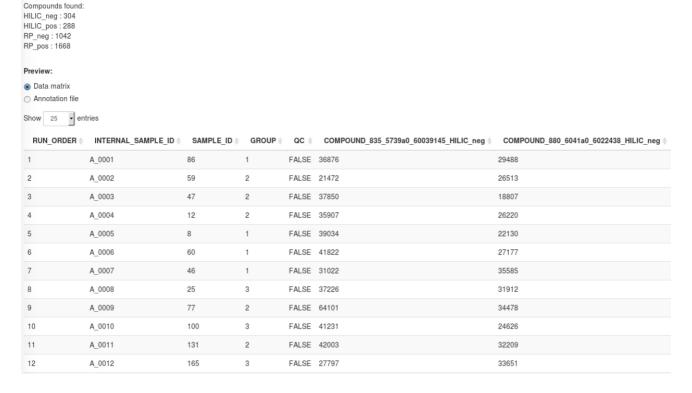
MPP tab

In the MPP tab you can combine MPP output files with sample information. The MPP output files should include the raw abundance values and the following columns: Compound, Mass, Retention Time, CompositeSpectrum, Frequency, CompoundAlgo and ionization mode.

The combined data matrix is ready for further data analysis. Other information of the compounds is stored in separate annotation file. Both files can be downloaded as .csv files.

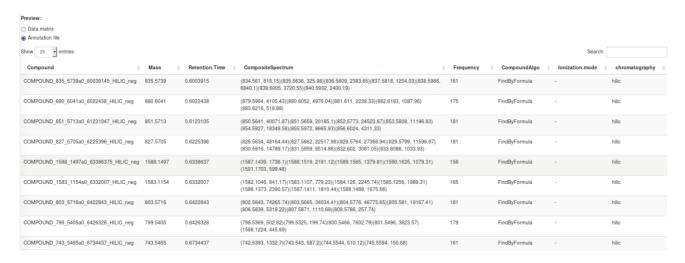


Example of the combined data matrix:



The data matrix has one row per sample. The columns from the processed sample sheet are combined with abundance values of all compounds found in all modes. The compound names are modified to better suit data analysis software.

Example of an annotation file:



The chromatography column is added. Compound names correspond to those in the data matrix.