
Subject: Feature Request: Reference List Filter
Posted by [richards99](#) on Sat, 30 May 2020 21:43:47 GMT

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Hi,
It would be really useful if with a dataset loaded, you can open up another dataset and use this dataset as a filter against the original data table (based on a specified column such as Structure). So besides the Append File.. option, maybe a Filter From File.. option.

Thanks,

Simon.

Subject: Re: Feature Request: Reference List Filter
Posted by [nbehrnd](#) on Tue, 02 Jun 2020 18:18:06 GMT

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Hi Simon,

might the merge/append and merge/replace function of DW an option for you?
<http://www.openmolecules.org/help/import.html#MergeData>

Norwid

Subject: Re: Feature Request: Reference List Filter
Posted by [richards99](#) on Tue, 02 Jun 2020 19:06:00 GMT

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I'm not sure this will do the function I want.
As an example, I have a data table with structures and data and I want to remove rows with specific structures which I have contained in a separate data file.

Simon.

Subject: Re: Feature Request: Reference List Filter
Posted by [nbehrnd](#) on Thu, 04 Jun 2020 12:54:12 GMT

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Hi Simon,

assuming both .dwar files contain a column of structures, DW's function «find similar compounds in file» possibly may perform the kind of sort you seek.

As a minimal example, I created a library of a few compounds around benzene and pyridine with some data computed by DW like molecular weight. This file ([larger_group.dwar](#), attached below) was closed.

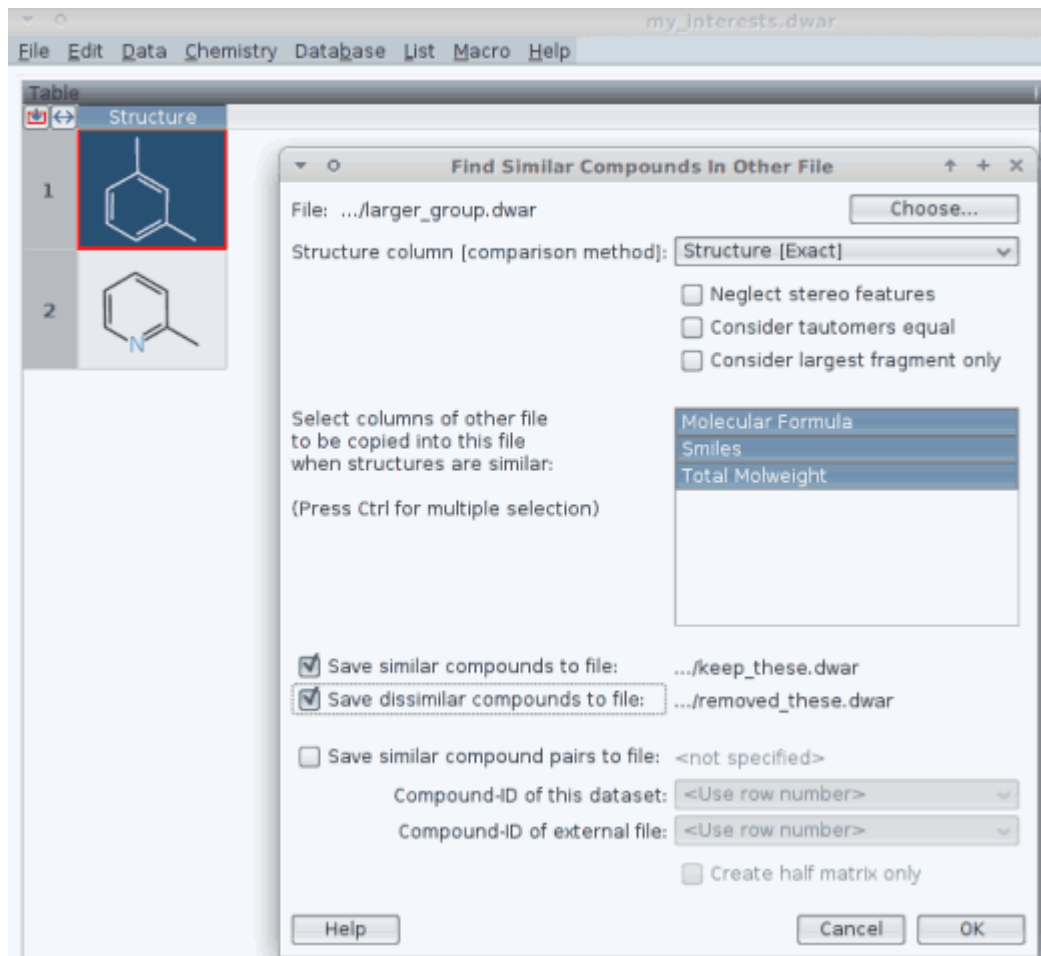
A second file was defined to contain only the structures of o-xylene, and 2-methyl pyridine ([my_interests.dwar](#)). From this file, I accessed chemistry -> Find Similar Compounds In File to encounter a new menu shown below. The file to scrutinize was indicated in the first field ([larger_group.dwar](#)), and the structure criterion was set to be exact. It is possible to use a less strict, and adjustable structure comparison, which is the one suggested by default, though.

Then, I opted to retain all of the data already computed in [larger_group.dwar](#) to be transmitted to either the new file about matching structures ([keep_these.dwar](#)), or to export the entries not needed now ([file_removed_these.dwar](#)).

The scope of structures of the two files newly written is complementary to each other, and adds up to the set of structures in the initial file, [larger_group.dwar](#).

File Attachments

- 1) [my_interests.dwar](#), downloaded 728 times
- 2) [larger_group.dwar](#), downloaded 707 times
- 3) [removed_these.dwar](#), downloaded 722 times
- 4) [keep_these.dwar](#), downloaded 693 times
- 5) [GUI_entries-or8.png](#), downloaded 1276 times



Subject: Re: Feature Request: Reference List Filter
Posted by [thomas](#) on Sun, 07 Jun 2020 18:45:14 GMT
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You also could add a structure list filter (substructure of similarity) and open the other file from within the filter structure list with a right mouse click:

File Attachments

1) [msf.png](#), downloaded 1153 times

