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**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.

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**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

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**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.

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**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.

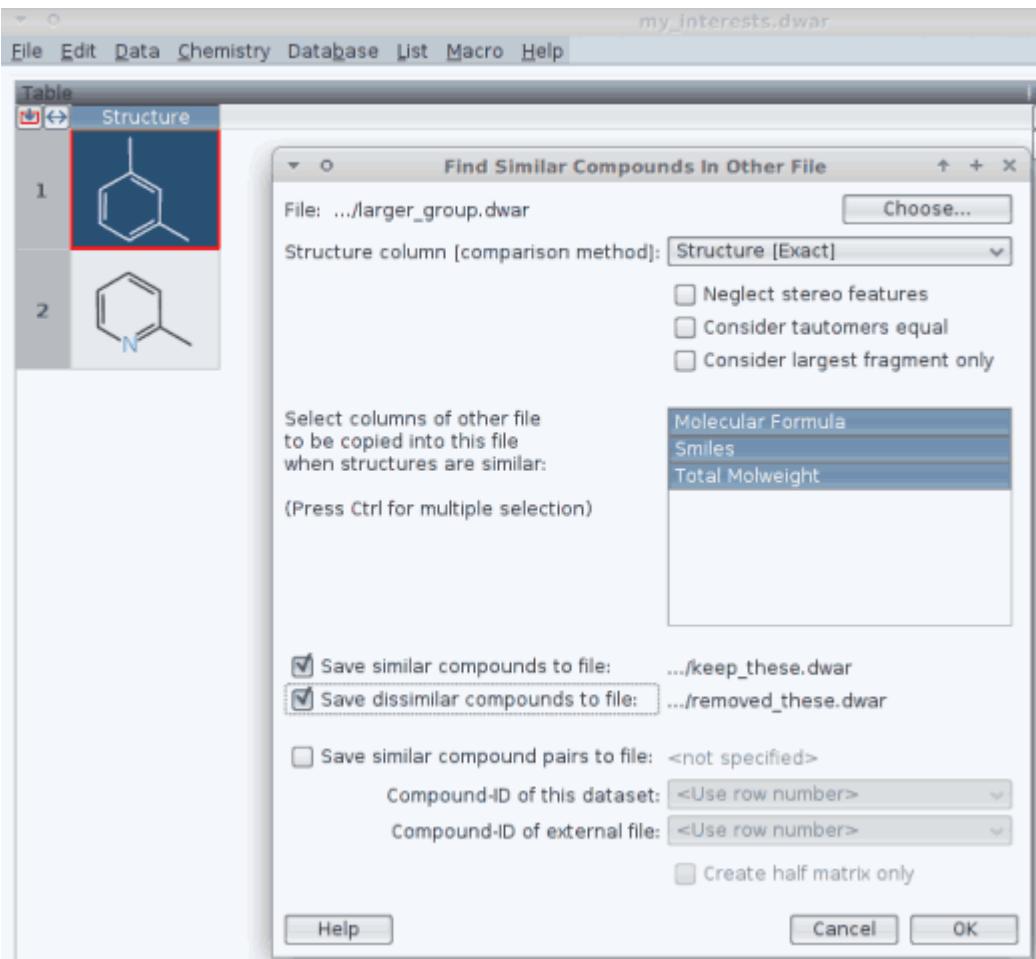
Then, I opted to retain all of the data already computed in larger\_group.dwar to be transmitted to either the new file about matching structures (keep\_these.dwar), or to export the entries not needed now (file removed\_these.dwar).

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

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- 1) [my\\_interests.dwar](#), downloaded 651 times
- 2) [larger\\_group.dwar](#), downloaded 621 times
- 3) [removed\\_these.dwar](#), downloaded 642 times
- 4) [keep\\_these.dwar](#), downloaded 598 times
- 5) [GUI\\_entries-or8.png](#), downloaded 1030 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 or similarity) and open the other file from within the filter structure list with a right mouse click:

#### File Attachments

1) [msf.png](#), downloaded 913 times

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## Structure



is similar to [FragFp]



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- [Add...](#)
- [Paste](#)
- [Add From File...](#)

