## Subject: Toggle-off absolute configuration while determining Murcko scaffold Posted by nbehrnd on Wed, 21 Aug 2019 12:00:58 GMT

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Dear Thomas.

using both DW as well as third-party python module rdkit to determine the Murcko scaffolds of molecules including those with a stereogenic center, I noticed DataWarrior (DW) retains the stereochemical information in the trimmed fragment. Reading the same SMILES string as DW (5.0.0), rdkit (version 2019.1 with Python2) however trims this information off.

Question: Is there an option to instruct DW equally to 'forget' about this piece of information when writing the SMILES string?

Pristine SMILES string in question used in both programs: C(=O)(C)O[C@H]1[C@H]([C@H](n2c3c(c(ncn3)N)nc2)O[C@@H]1CO)O

DW's output SMILES string about the Murcko scaffold: C(C1)CO[C@H]1n1c2ncncc2nc1

MWE for processing with rdkit:

from rdkit.Chem.Scaffolds import MurckoScaffold from rdkit.Chem import AllChem

source = 'C(=O)(C)O[C@H]1[C@H]([C@H](n2c3c(c(ncn3)N)nc2)O[C@@H]1CO)O' mol = Chem.MolFromSmiles(source) core = MurckoScaffold.GetScaffoldForMol(mol) print(Chem.MolToSmiles(core))

>>> c1ncc2ncn(C3CCCO3)c2n1

which constitutionally is about the same molecular structure.

This may be more than a cosmetic issue, because the search for the Murcko scaffold of

Brc1ccc([C@@]2(CC(=O)CCC2)CN(=O)=O)cc1

actually leads DW to invert the absolute configuration from initial R to now S (expressed by the lost of one @ in string O=C(CCC1)C[C@H]1c1cccc1).

## File Attachments

1) DWclip.png, downloaded 348 times

- 2) rdkit\_ACADOS.png, downloaded 310 times
- 3) ABEGUL\_DW\_inversion.png, downloaded 341 times

Subject: Re: Toggle-off absolute configuration while determining Murcko scaffold Posted by thomas on Fri, 23 Aug 2019 23:09:03 GMT

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Dear Norwid.

to me the current handling is correct: the change from R to S in your sample SMILES is correct, because after the removal of the methylene-nitro group, the stereo-chemistry is correctly retained in the Murcko-scaffold.

If you generate a SMILES from the scaffold, then you may use 'Find and replace' to remove all '@' symbols from them.

**Thomas** 

## File Attachments

1) t.png, downloaded 546 times

Subject: Re: Toggle-off absolute configuration while determining Murcko scaffold Posted by nbehrnd on Sun, 25 Aug 2019 16:56:16 GMT View Forum Message <> Reply to Message

Dear Thomas,

as probed, the «Find and Replace» (under the Edit tab) indeed offers exactly the tool I needed to simplify the SMILES as intended. This change both propagates well in subsequent stages of structure display

as well in the how to retrieve -- either by structure, or by string -- structures sharing a Murcko scaffold in common.

Thank you, Norwid

## File Attachments

- 1) example.png, downloaded 529 times
- 2) example.dwar, downloaded 298 times