Guideline For The Use Of Der/Xml Composer – Rat Metabolism v.5.2

Prepared for use with DER Composer, a product resulting from the joint cooperation between the U.S. Environmental Protection Agency (USEPA) and the Laboratory of Mathematical Chemistry (LMC-Bourgas, Bulgaria).

Richard Kolanczyk

Great Lakes Toxicology and Ecology Division

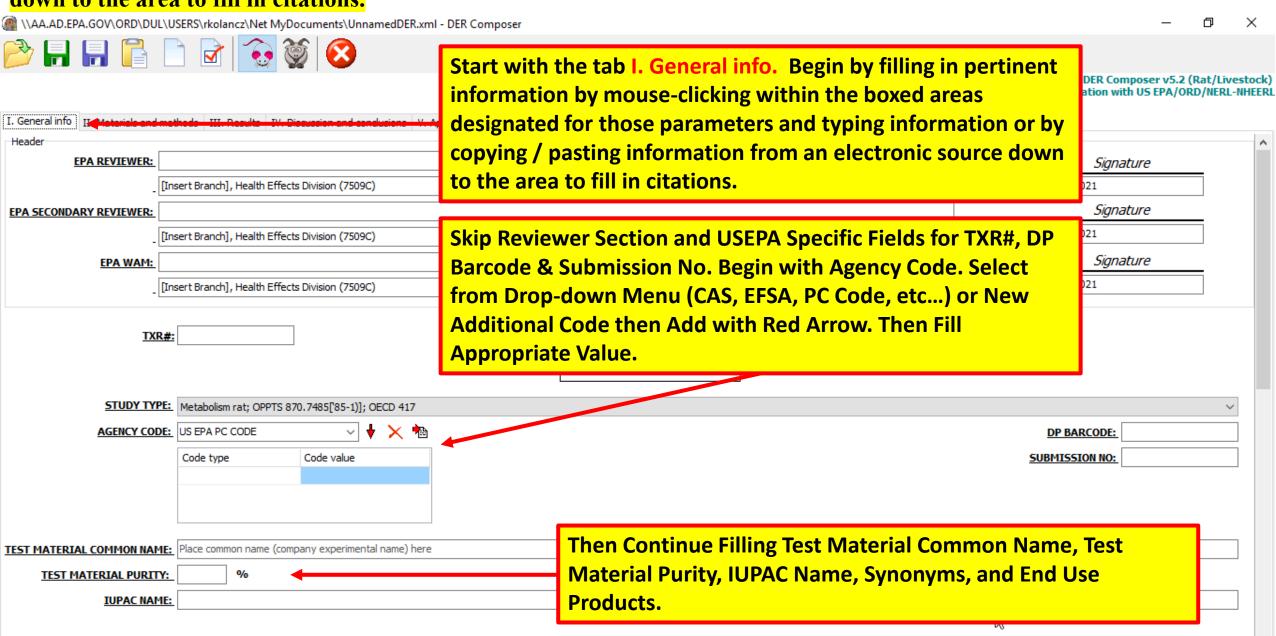
Center for Computational Toxicology and Exposure

Office of Research and Development, U.S. Environmental Protection Agency

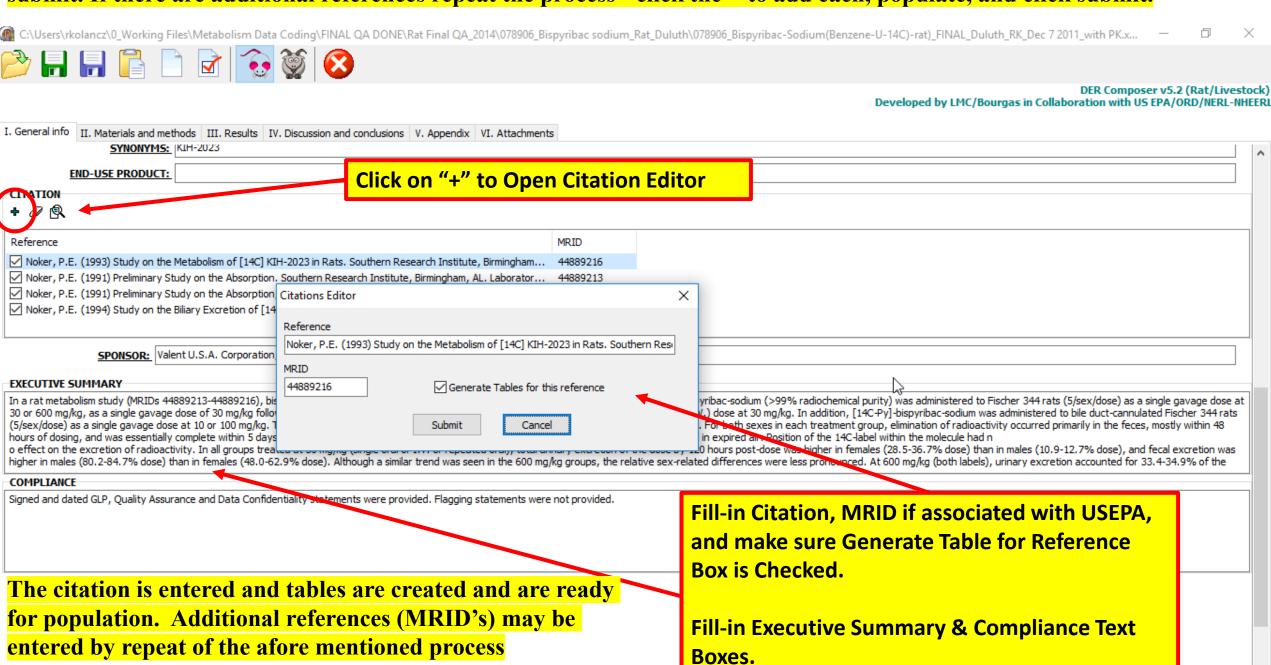
Duluth, MN, USA

- Opening DER
- General info
- Material and methods

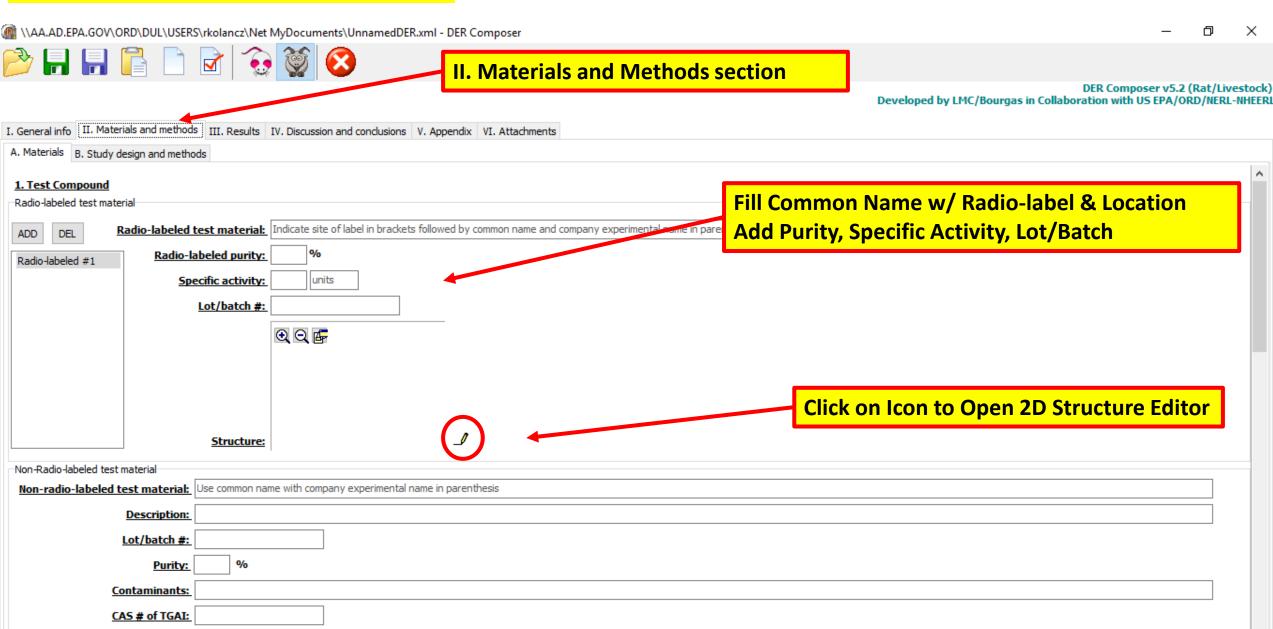
Open the program – start up will appear as follows: \\AA.AD.EPA.GOV\ORD\DUL\USERS\rkolancz\Net MyDocuments\UnnamedDER.xml - DER Composer **To Exit the Program** DER Composer v5.2 (Rat/Livestock) Collaboration with US EPA/ORD/NERL-NHEERL Icon – To Select Option for Livestock I. General info II. Materia's and methods III. Results IV. Discussion and conclusion. V. Appendix VI. Attachments Composer – Now Obsolete with MSS **Livestock Composer EPA REVIEWER** Signature (nsert Branch), Health Effects Division (7509C) EPA SECONDARY REVIEWER: Icon – To Select Option for Rat Signature 2021 [Insert Branch], Health Effects Division (7509C) **Metabolism Composer to Capture** EPA WAM: Signature **ADME Studies** [Insert Branck], Health Effects Division (75090 TXR # Icon – For Options Relating to System A EVALUATION RECOR Icon – To Reset **Warnings for Saving XML** the Composer, STUDY TYPE: Metabolism rat; OPPTS 870.7485['85-1)]; QECD 417 **To Start Over** AGENCY CODE: US ENA PC CODE Options Code value Code type Warnings Icon - Used to Generate ✓ Warn if saving with "RadioLabelled test material" left blank Icon - To Save a DER in the Form of a Warn if exiting with an unsaved DER File as XML ✓ Warn on exit **WORD Document** TEST MATERIAL COMPON Icon - To Save As Icon - To Open a Restore default Save changes X Cancel Command - Useful to **Previously Saved XML File for Editing** Make a Copy of a **Currently Saved XML** THE HET BROBLICT



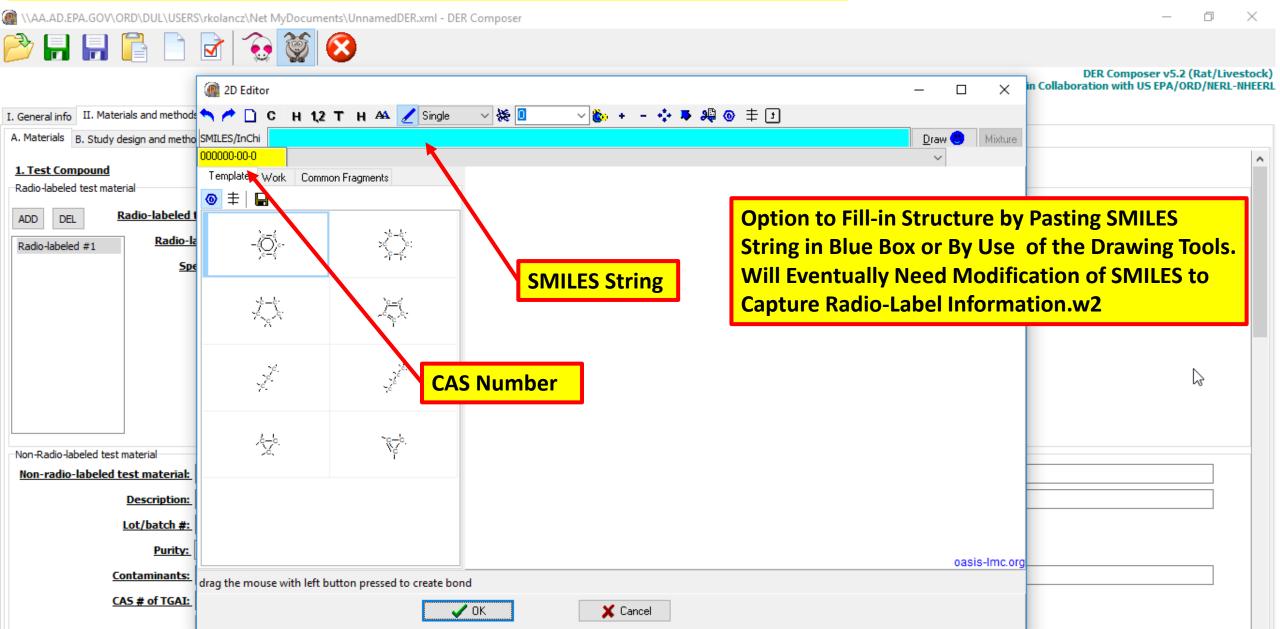
A CITATION EDITOR box pops up. Fill in reference, MRID number and click generate tables, followed by clicking on submit. If there are additional references repeat the process - click the + to add each, populate, and click submit.



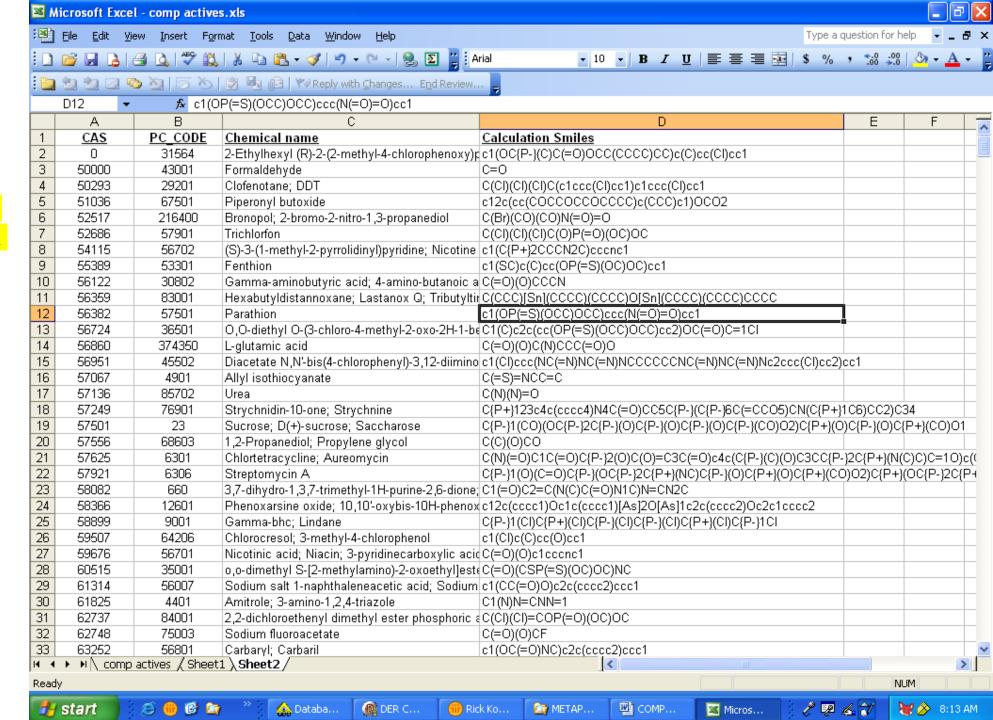
Next the tab II. Materials and methods and sub-tab A. Materials may be populated. Data is filled in via directly typing or copy/paste from electronic documents until reaching the structure entry. To enter the Radio-labeled Test Material and Non-radio-labeled Test Material structures:



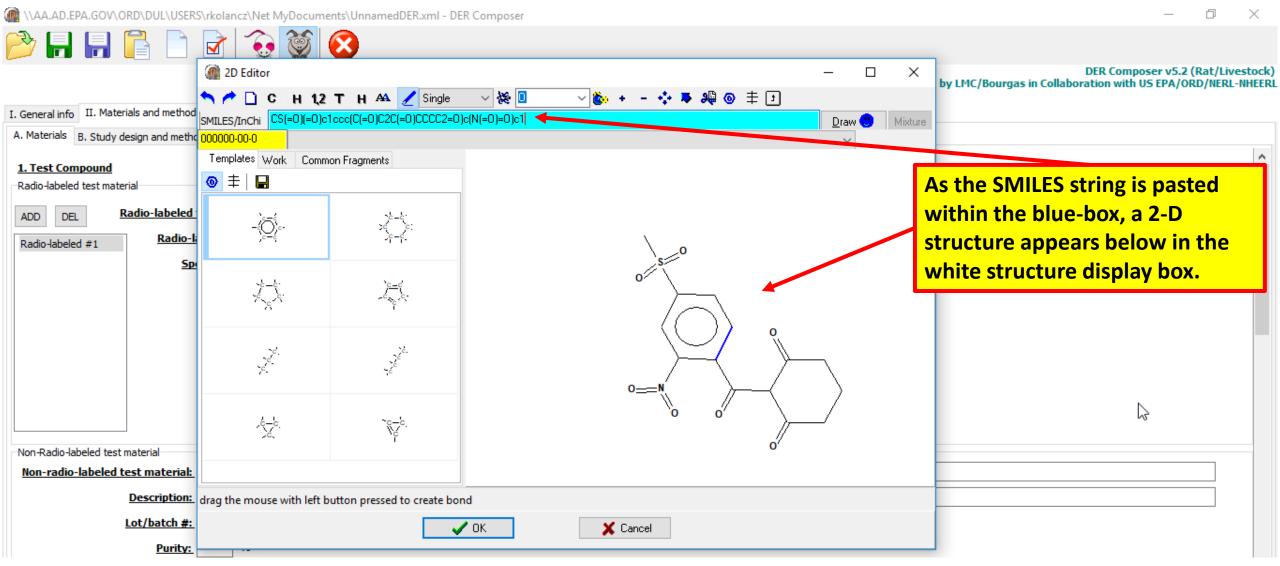
Below is a graphic of the STRUCTURE DRAWING editor pop-up box. The large white area is the drawing workspace, the light-blue box is where a SMILES string may be entered or displayed, and the yellow box is where a CAS number can be entered. Scrolling over the top of the icons will give some indication of their utility.



To get started a list of parent 2-D structures represented as linear "SMILES strings" will be made available. Locate the appropriate structure from the excel file, copy the SMILES string.



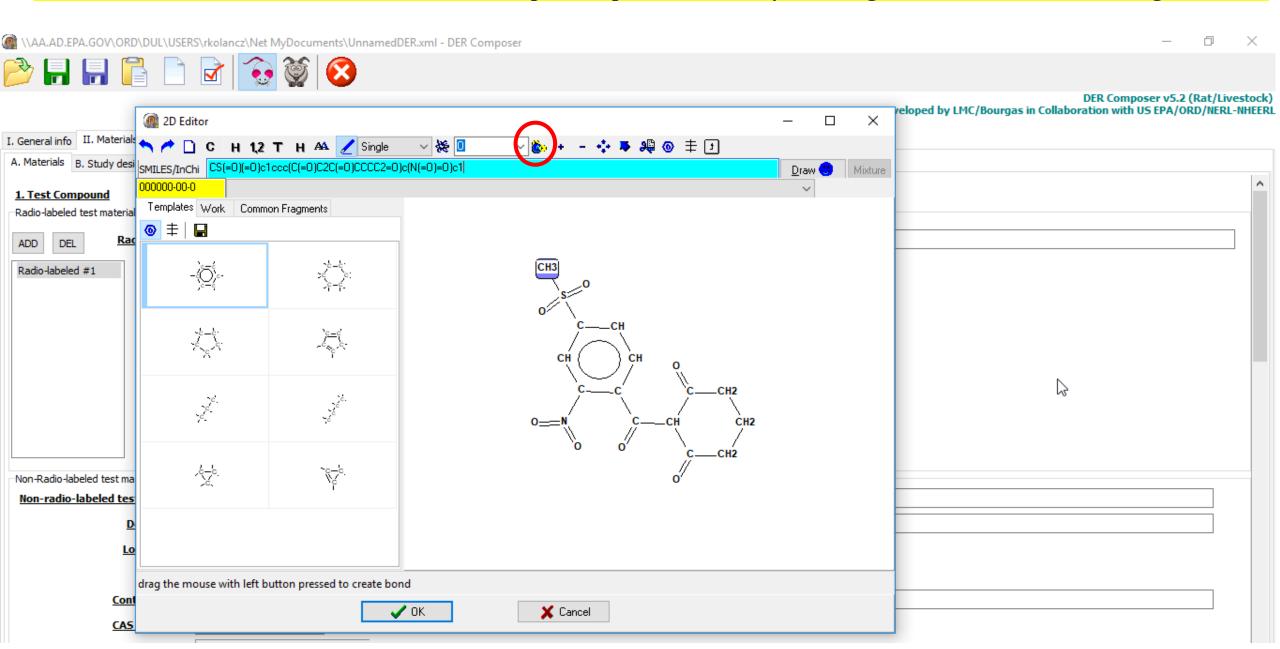
Perform a right-hand click of the mouse in the light-blue box of the STRUCTURE DRAWING package and select paste to enter the parent structure.



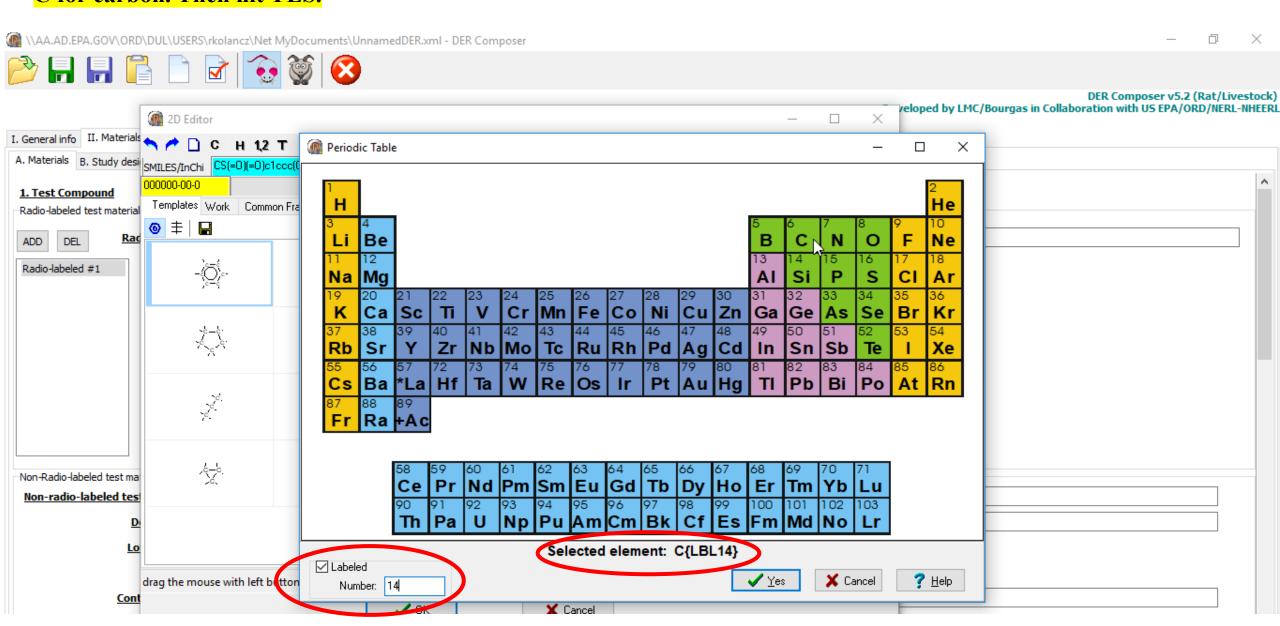
The parent structure (now present in the STRUCTURE DRAWING editor) may be modified utilizing tools within the editor. Specifically a label may be introduced in the structure of the radio-labeled parent.

Radio labeling of atoms

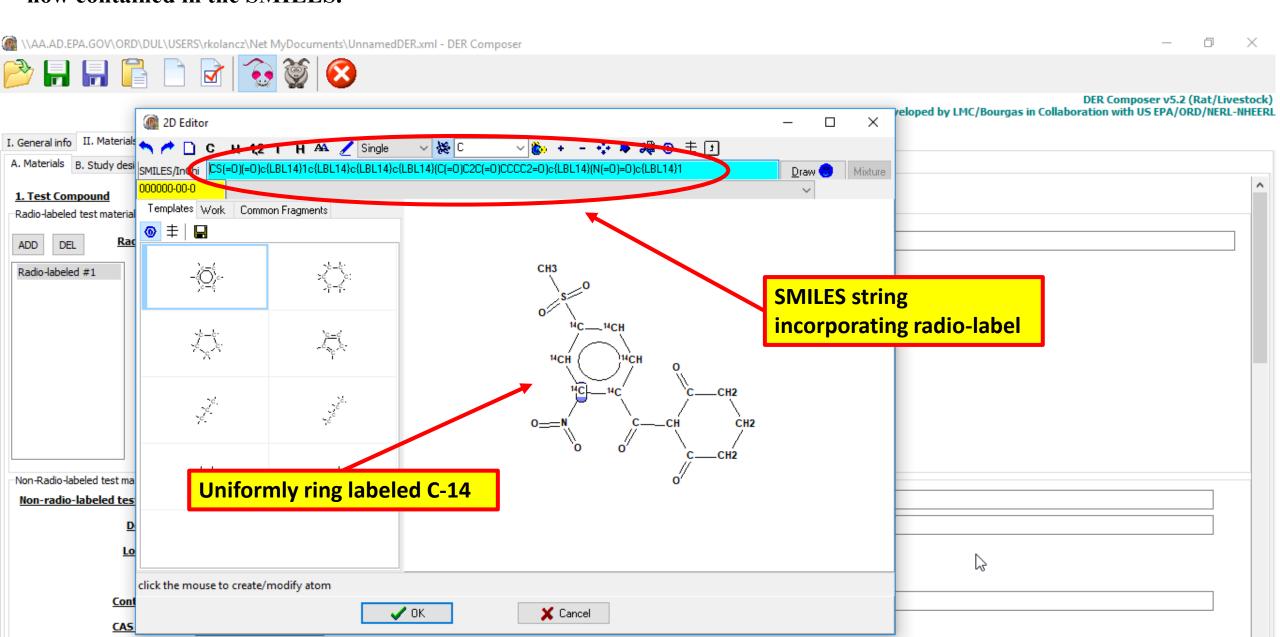
Within the STRUCTURE DRAWING window, open the periodic table by selecting the icon as circled in the figure below.

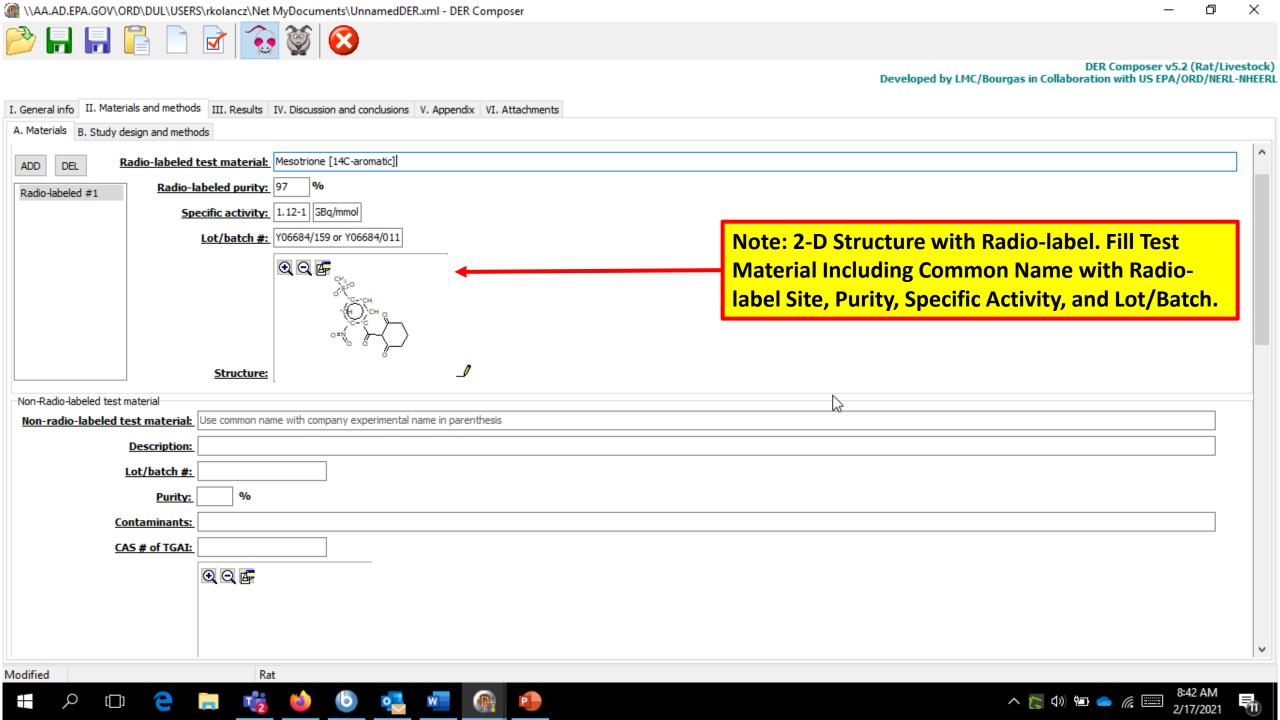


A periodic table screen comes up where you should check labeled, in this example add 14 in the number box and click on C for carbon. Then hit YES.

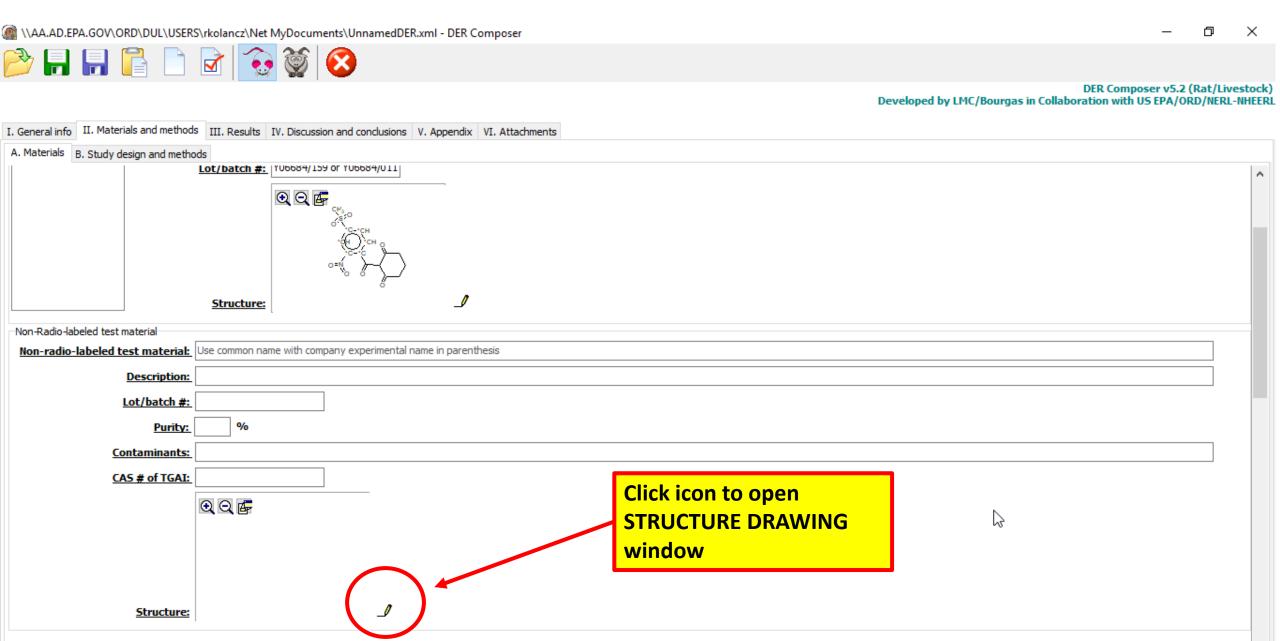


After clicking Yes on the previous screen, the periodic table closes, then you can add the C-14 label to each carbon in the example. The example below happens to be a uniformly labeled phenyl ring. Note that the information for the labeling is now contained in the SMILES.



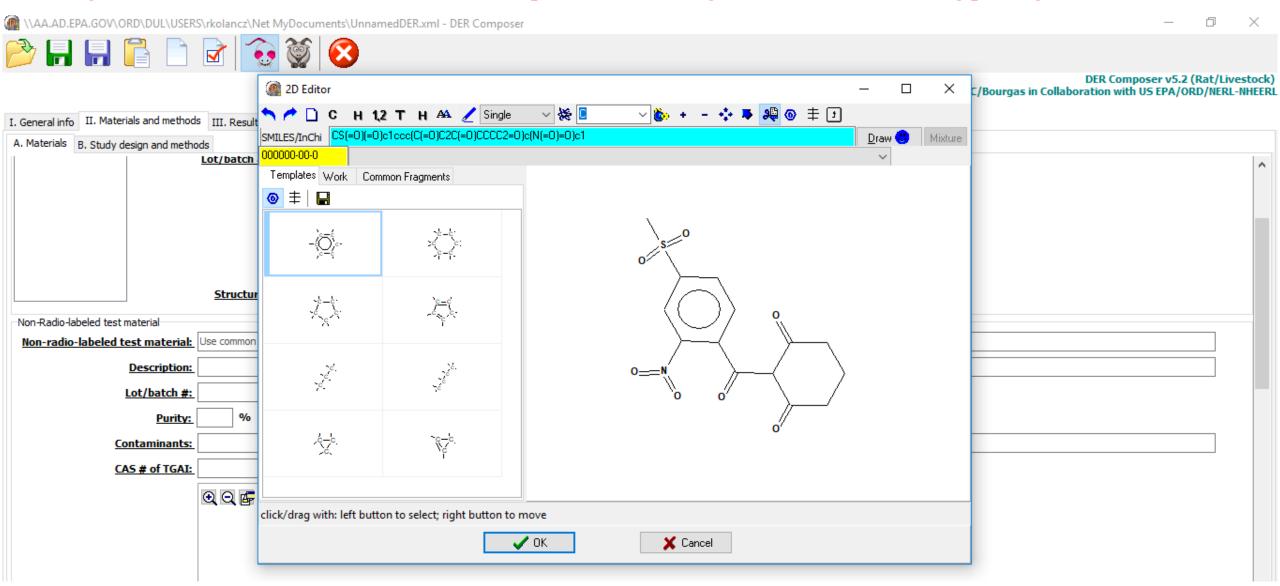


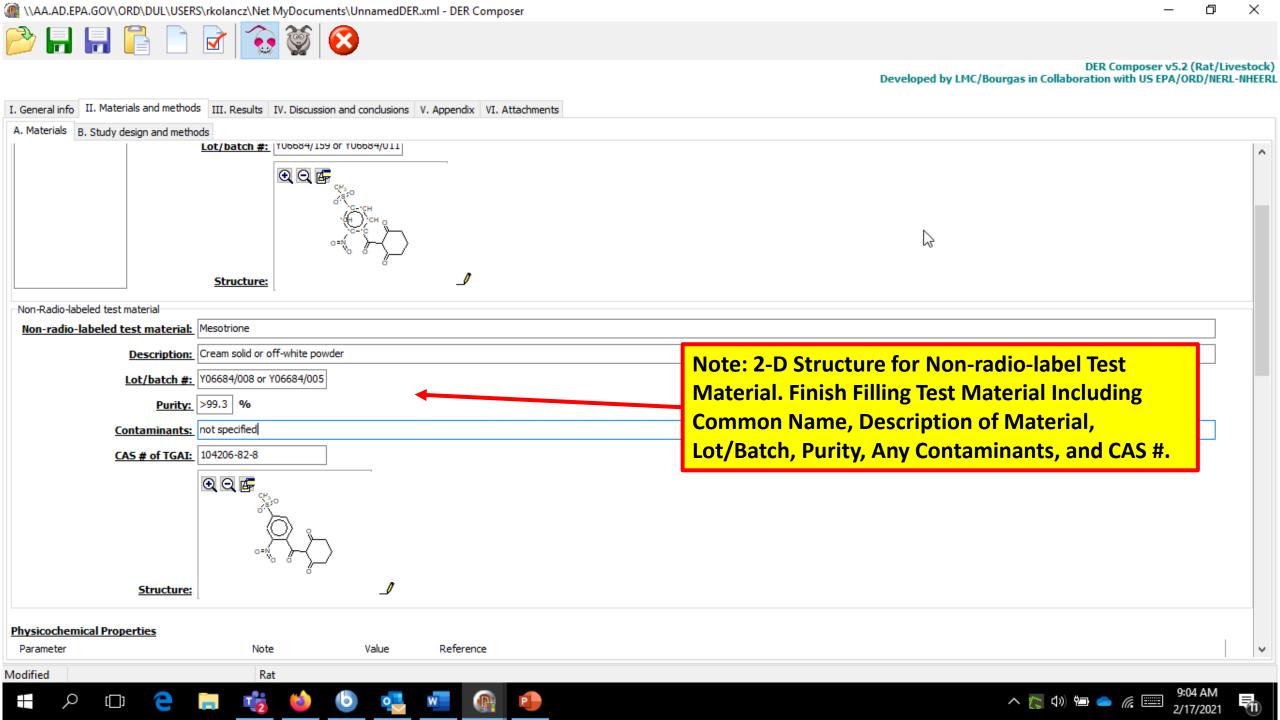
The Non-radiolabeled Test Material may be entered in the same fashion by opening the STRUCTURE DRAWING PACKAGE.



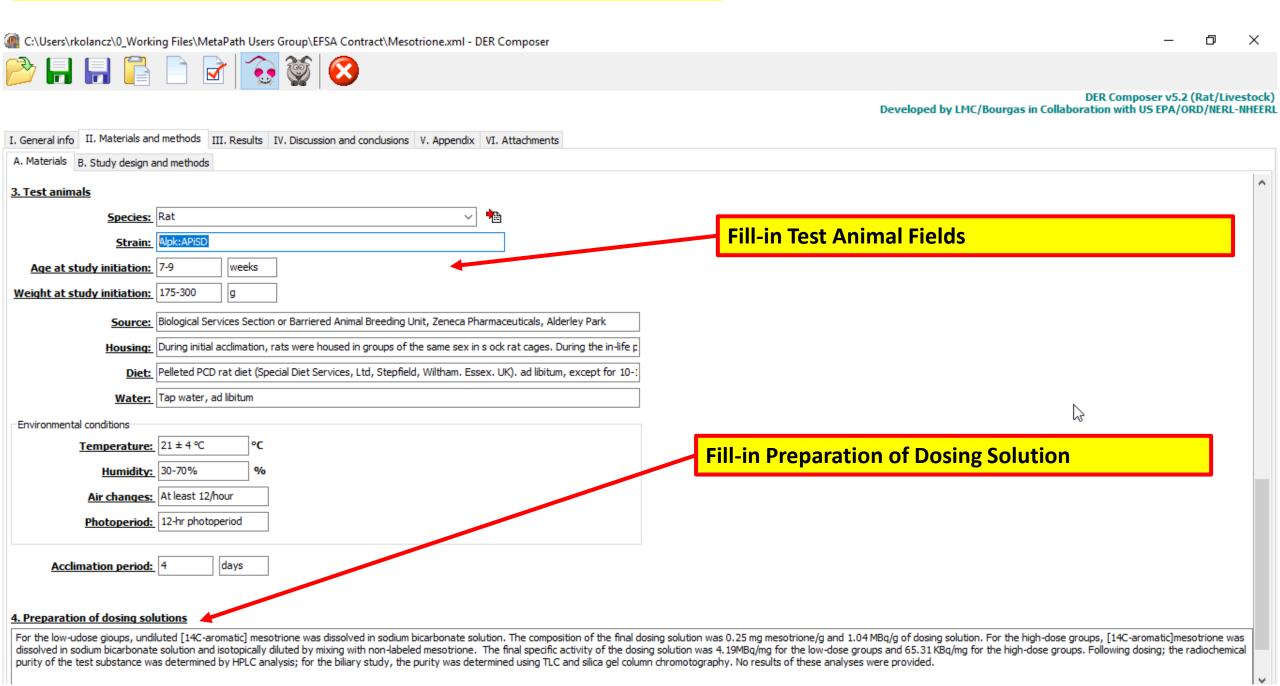
The SMILES string (from the excel list of parent structures) is entered in the light-blue box of the editor and the 2-D structure is immediately shown.

NOTE: The use of COPY/PASTE SMILES strings to generate the 2-D structures of parent chemicals serves to save time drawing structures, however the structures can be produced utilizing the tools of the drawing package.





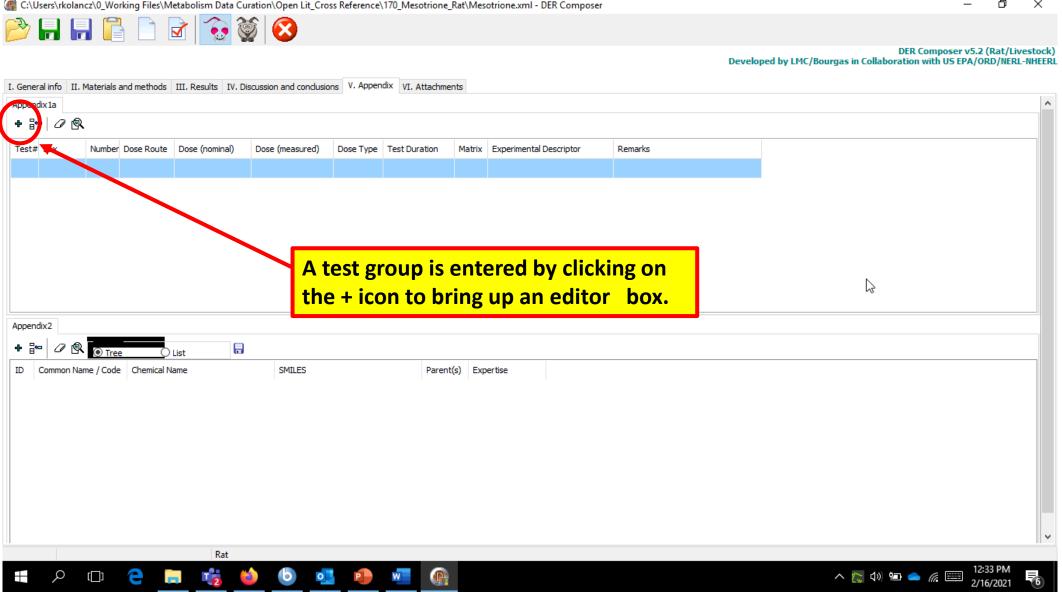
Continue filling out the rest of II. Material and methods A. Materials



- Appendix 1
- Appendix 2
- Material and methods
- Study design

Next go to tab V. Appendix – It is within this section that the various treatment groups are defined and listed as a TEST in the appendix 1 table below. A treatment group may be defined by gender, age, dose amount, dose route, sample matrix or other experimental descriptor (parameters that when varied may give rise to a different metabolic map for a particular chemical).

© C:\Users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\rho|users\



Cilck on submit to accept test — continue to add new tests via the same process until completed. Screen should appear as below: © C:\Users\rkolancz\0_\Working Files\Metabolism Data Curation\Open Lit_Cross Reference\170_Mesotrione_Rat\Mesotrione.xml - DER Composer — □ × DER Composer v5.2 (Rat/Livestock) Developed by LMC/Bourgas in Collaboration with US EPA/ORD/NERL-NHEERL

I. General info II. Materials and methods III. Results IV. Discussion and conclusions V. Appendix VI. Attachments + 2 0 0 Test# Sex Number Dose Route Dose (nominal) Dose (measured) Dose Type Test Duration Matrix | Experimental Descriptor Remarks pooled samples of feces, urine, and bil Female 2 Oral 50.0 mg/kg 48.5 mg/kg 48 hrs single Urine pooled samples of feces, urine, and bil Female Oral 50.0 mg/kg 48.5 mg/kg single 48 hrs Feces 50.0 mg/kg 48.5 mg/kg 48 hrs Bile pooled samples of feces, urine, and bil Oral single 104.8 mg/kg used solely as a source of fecal and uri Oral 100.0 mg/kg single 48 hrs Urine used solely as a source of fecal and uri 100.0 mg/kg 104.8 mg/kg Oral single 48 hrs Feces used solely as a source of fecal and uri Oral 100.0 mg/kg 110.7 mg/kg single 48 hrs Urine Female used solely as a source of fecal and un 100.0 mg/kg 110.7 mg/kg Female Oral single 48 hrs Feces Appendix2 Common Name / Code | Chemical Name SMILES Parent(s) Expertise

The completed Appendix 1 will automatically populate the group arrangements Table 1 of section/tab II. Materials and methods sub-tab B. Study design and methods. This may be observed by clicking on the appropriate tabs.

© C:\Users\rkolancz\0_Working Files\Metabolism Data Curation\Open Lit_Cross Reference\170_Mesotrione_Rat\Mesotrione.xml - DER Composer

| C:\Users\rkolancz\0_Working Files\Metabolism Data Curation\Open Lit_Cross Reference\170_Mesotrione_Rat\Mesotrione.xml - DER Composer

| DER Composer v5.2 (Rat/Livestock)

Developed by LMC/Bourgas in Collaboration with US EPA/ORD/NERL-NHEERL I. General info II. Materials and methods III. Results IV. Discussion and conclusions V. Appendix VI. Attachments A. Materials B. Study design and methods 1. Group arrangements Animals were assigned to the test groups noted in Table 1 Table 1a Treatment Group Sex Number Dose Route Dose (nominal) Dose (measured) Dose Type Remarks 100mg/kg Mal 5 Oral 100.11mg/kg single Feces, Tissue Fer 5 100mg/kg 98.79mg/kg Oral single Feces, Tissue Mal 5 Oral 1.0mg/kg 1.0mg/kg single Feces, Tissue Fer 5 Oral 1.0mg/kg 1.0mg/kg single Feces, Tissue Mal 5 Oral 1.0mg/kg 0.99mg/kg multiple Feces, Tissue Fer 5 1.02mg/kg multiple Oral 1.0mg/kg Feces, Tissue Mal 5 I.V. 1.0mg/kg 0.99mg/kg single Feces, Tissue 2. Dosing and sample collection Briefly describe dosing methods and sample collection Table2a

Analytical Detection

Remarks

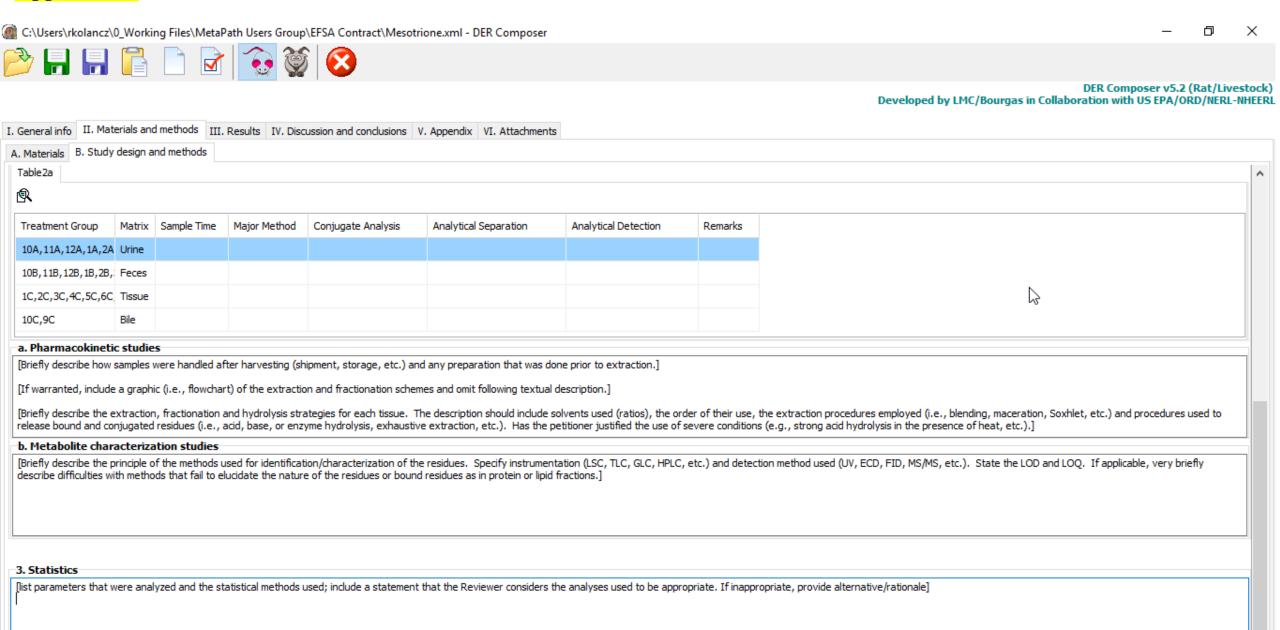
Matrix Sample Time Major Method

Conjugate Analysis

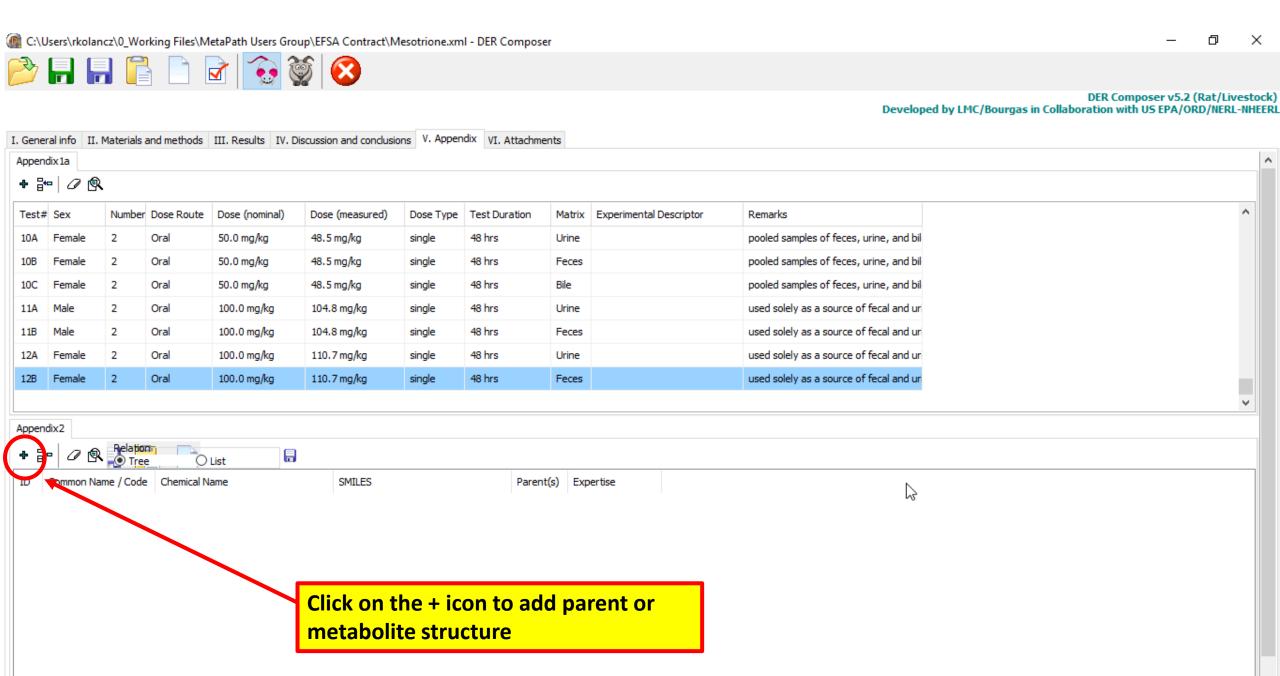
Analytical Separation

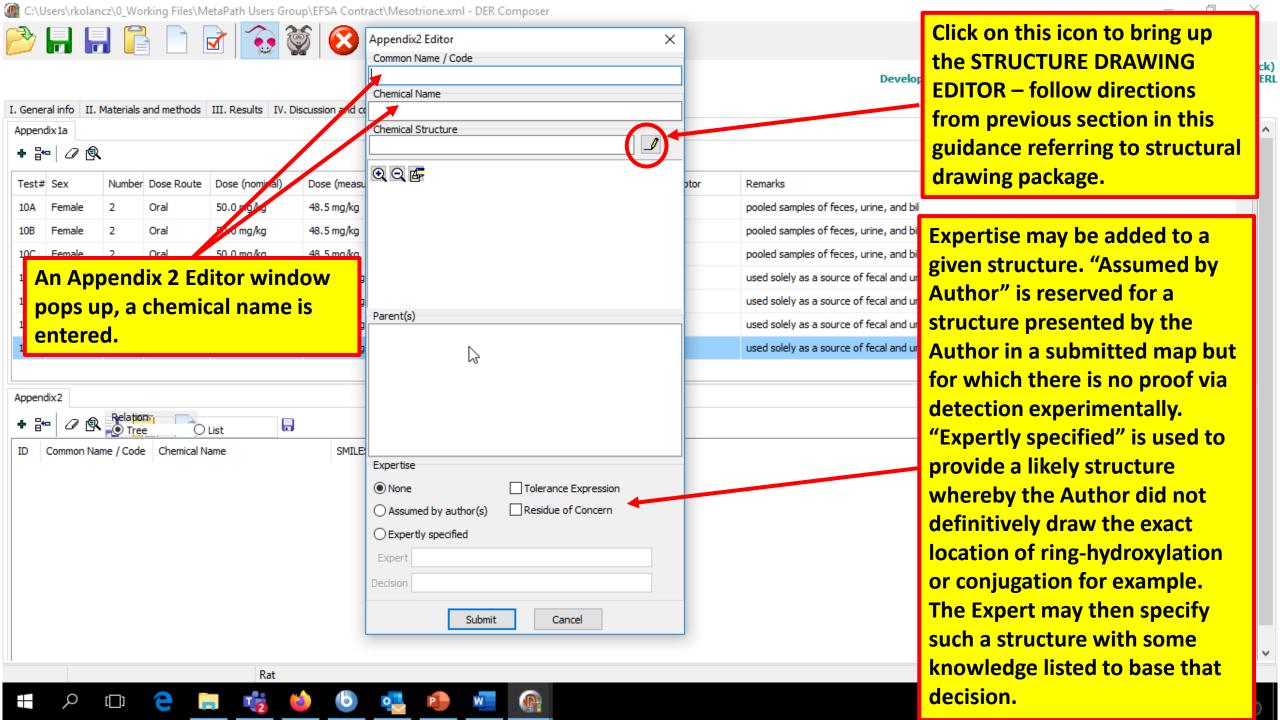
Treatment Group

In addition, an automatic partial entry of the dosing and sample collection Table 2 of section/tab II. Materials and methods sub-tab B. Study design and methods takes place. We will return to complete this table after completion of Appendix 2.

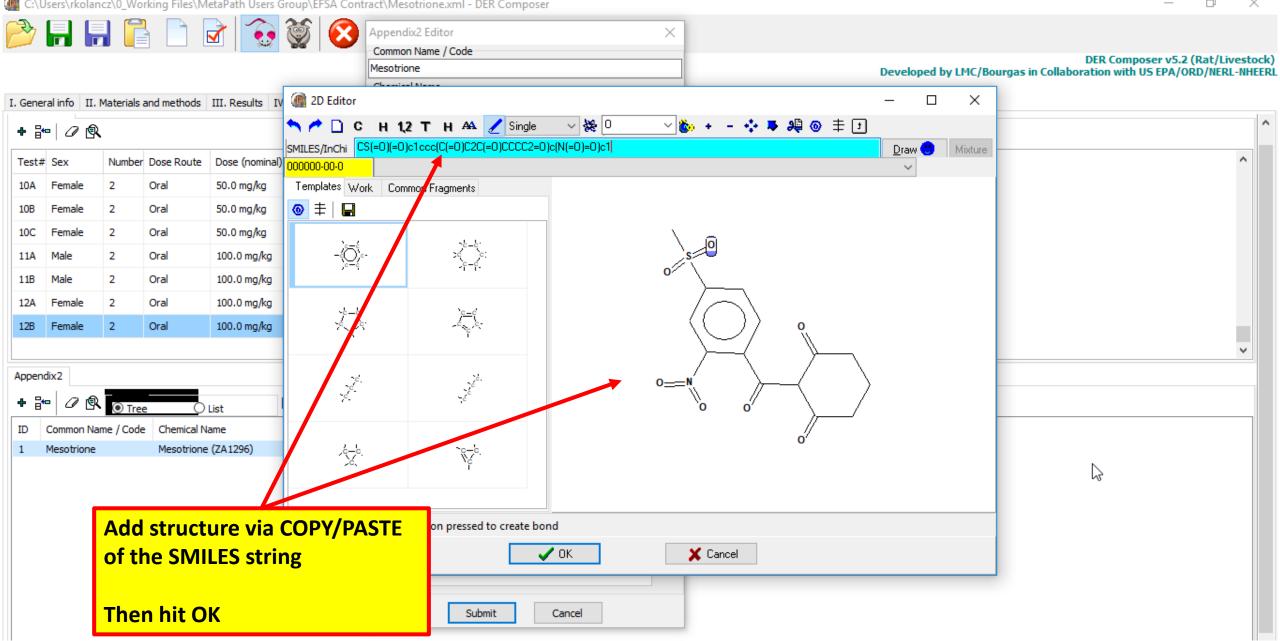


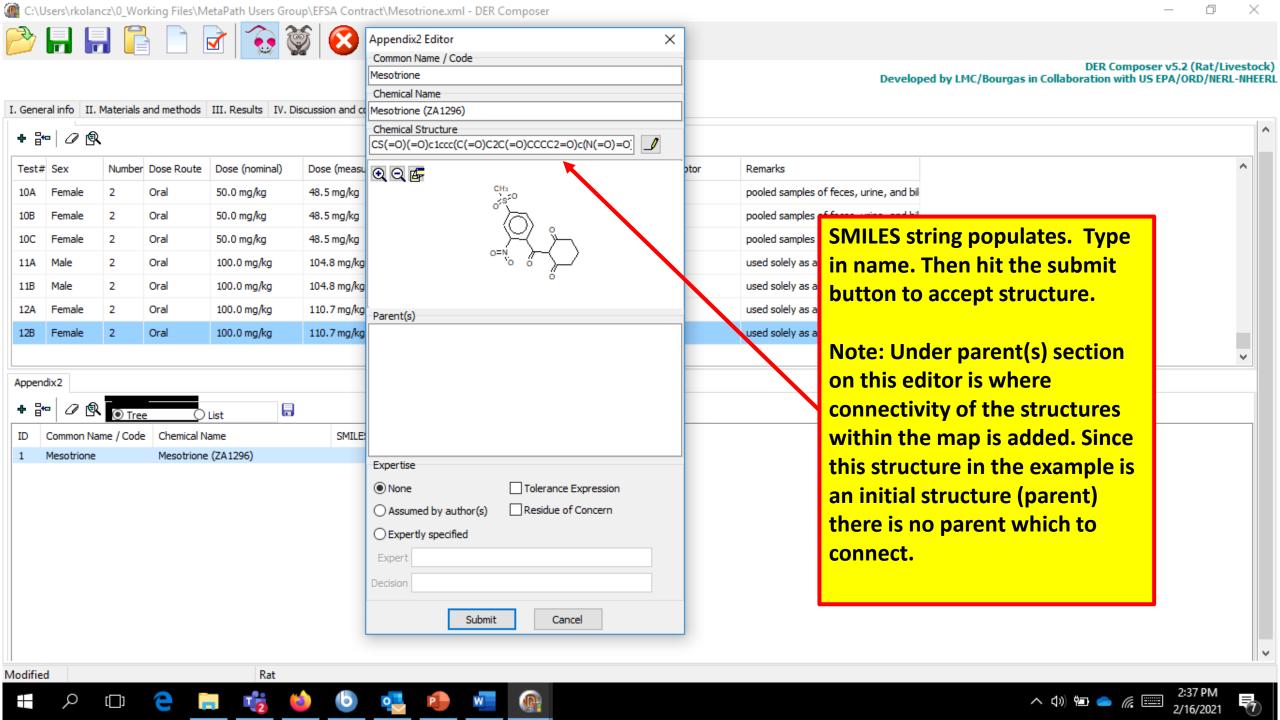
Next a Metabolite Inventory table should be completed as Appendix 2 of tab V. Appendix.

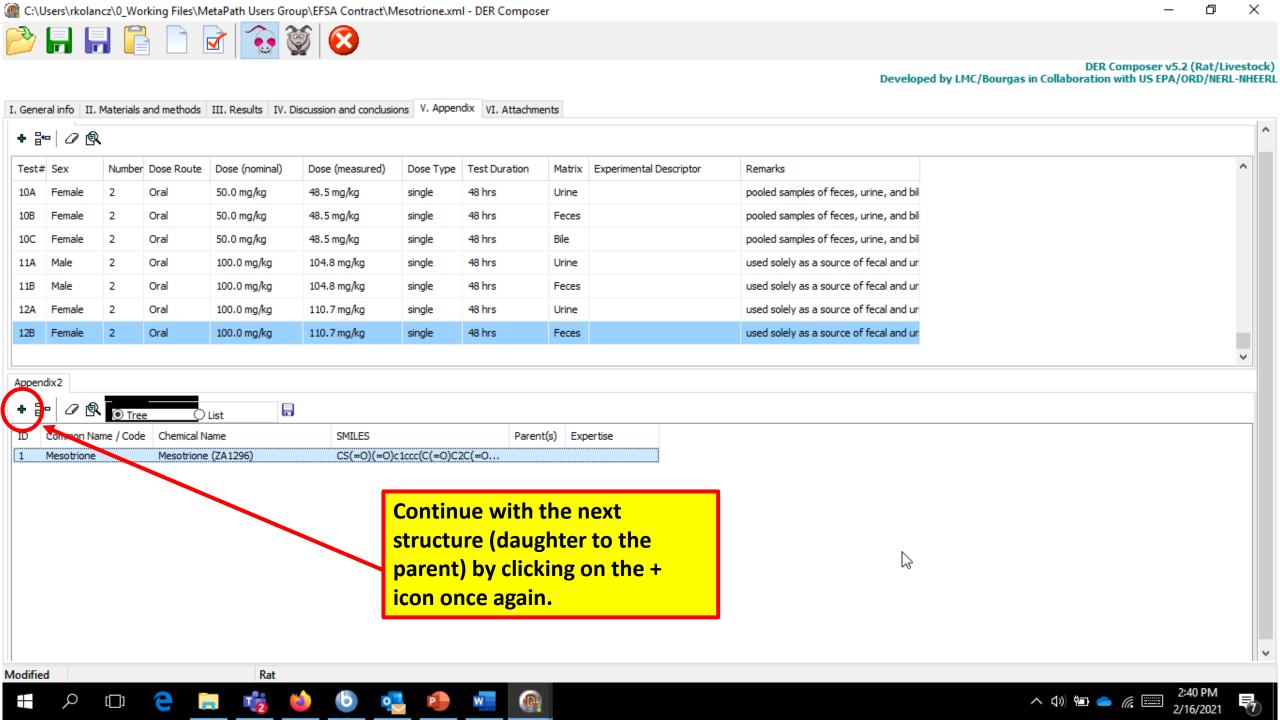


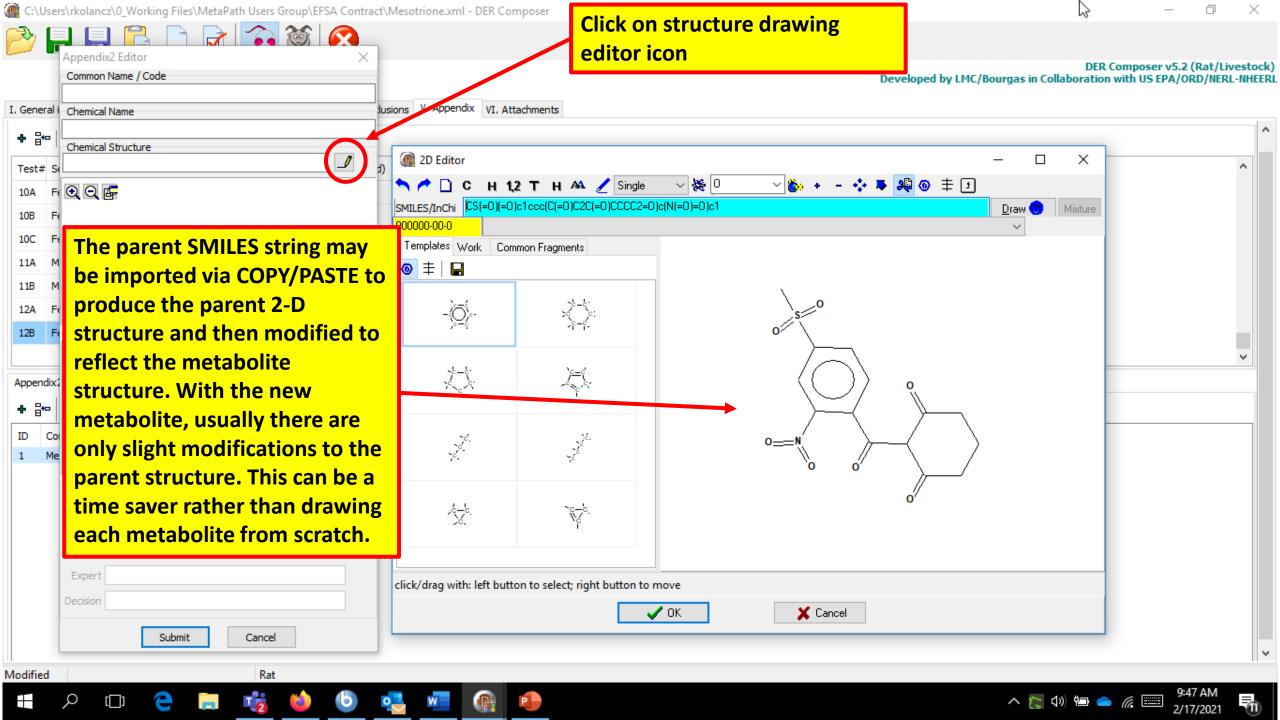


We will start by adding the parent structure – as was done in the materials & methods using COPY/PASTE of the SMILES string. (@ C:\Users\rkolancz\0_Working Files\MetaPath Users Group\EFSA Contract\Mesotrione.xml - DER Composer Appendix2 Editor Common Name / Code DER Composer v5.2 (Rat/Livestock) Mesotrione

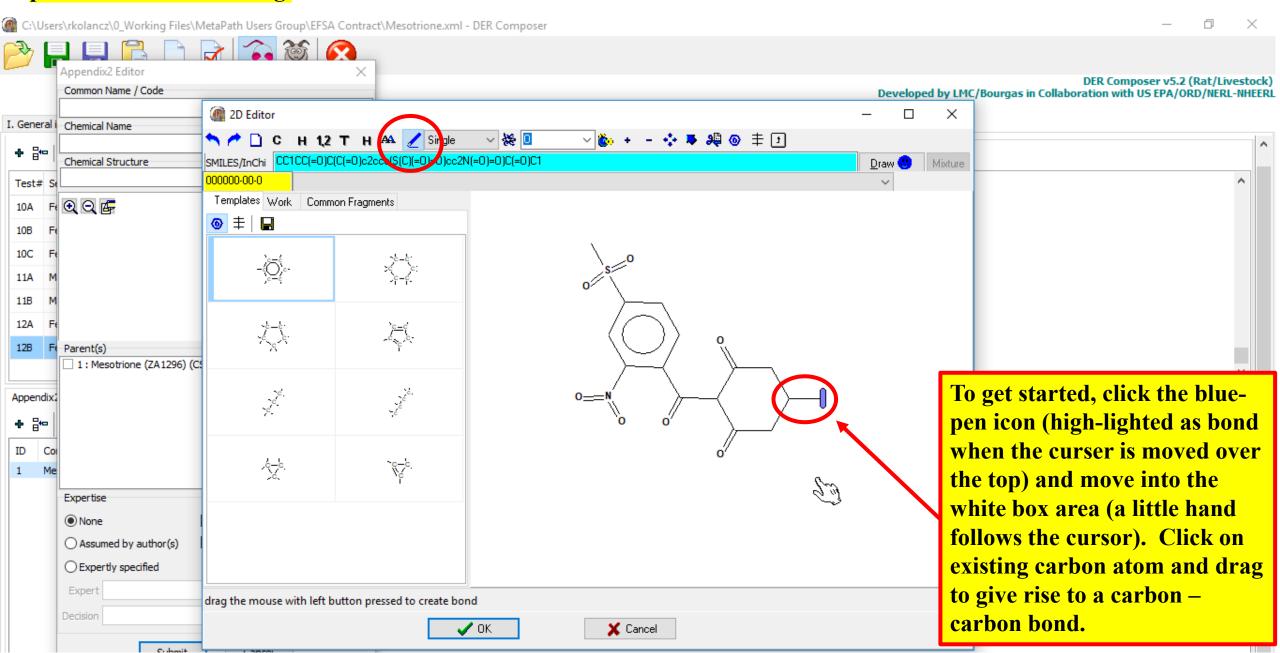




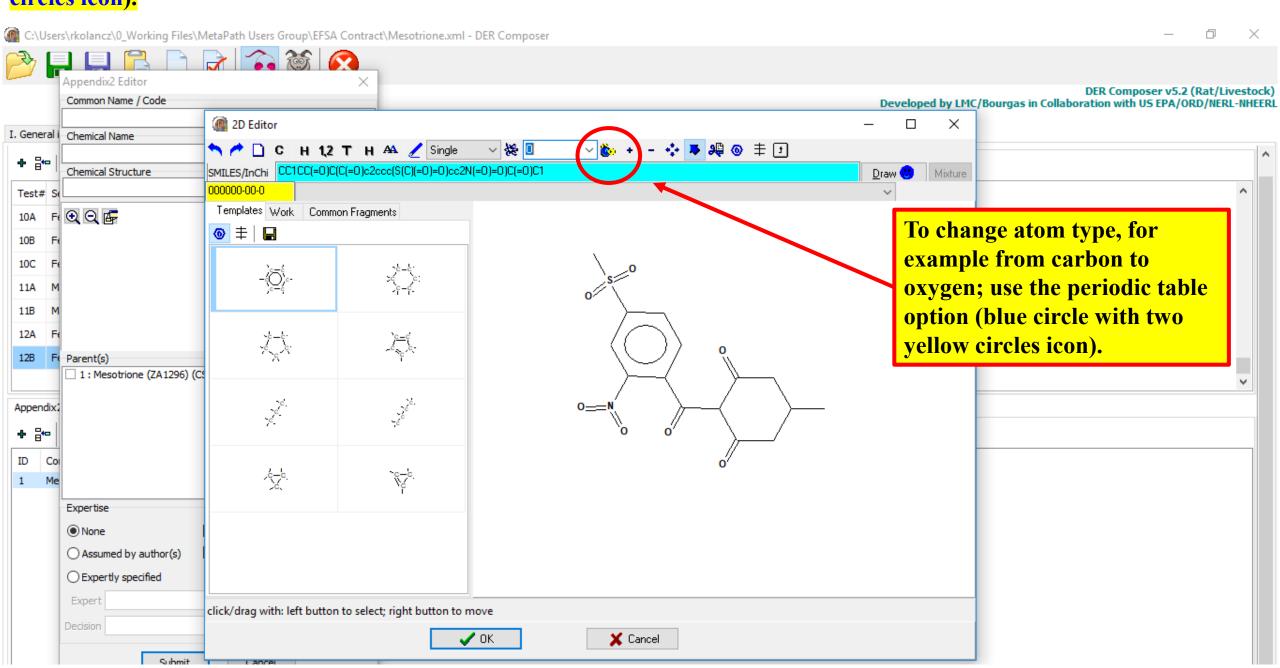




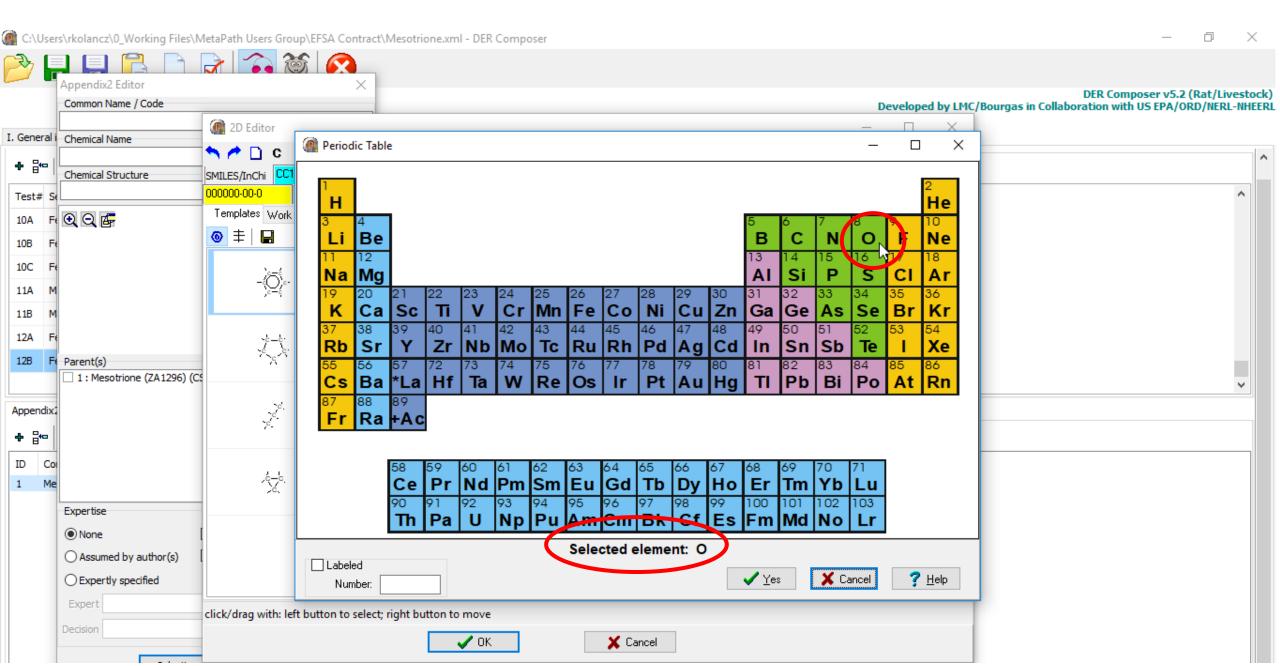
In this example the metabolite is 5-hydroxy-mesotrione. The following steps will introduce a hydroxy group in the 5-position of the dione ring.



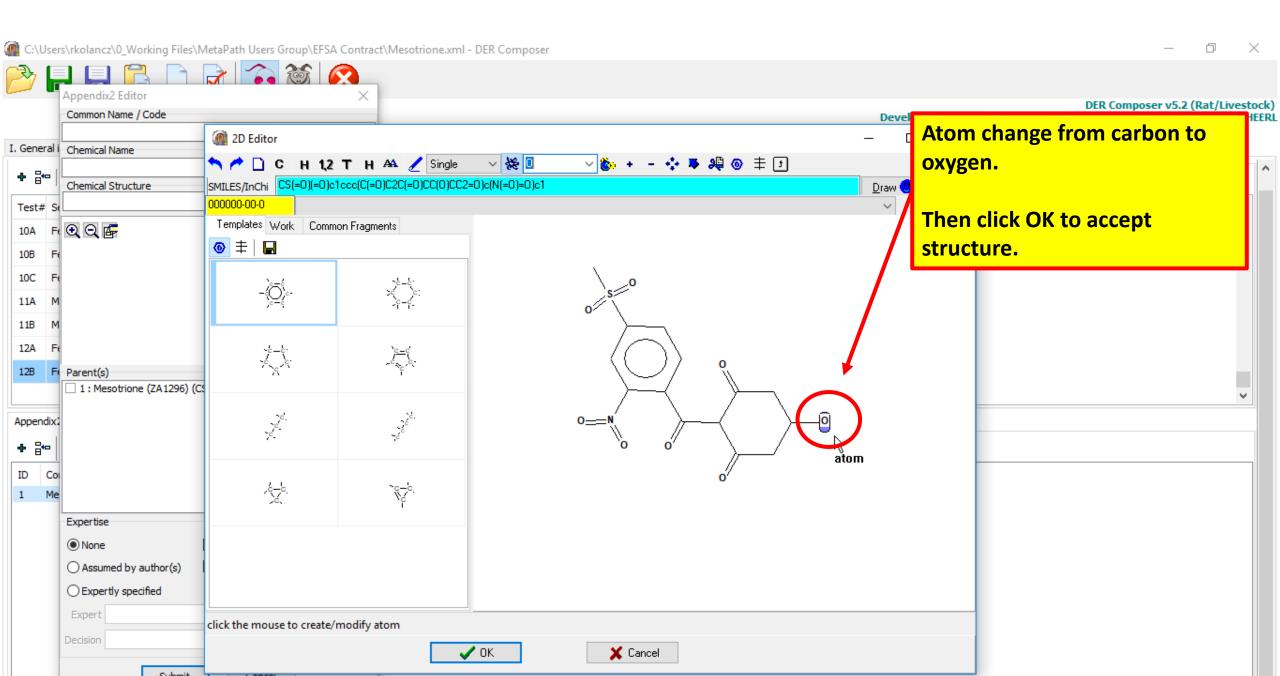
To change atom type, for example from carbon to oxygen; use the periodic table option (blue circle with two yellow circles icon).



The periodic table opens, click on atom choice, click Yes to accept choice and the table goes away.

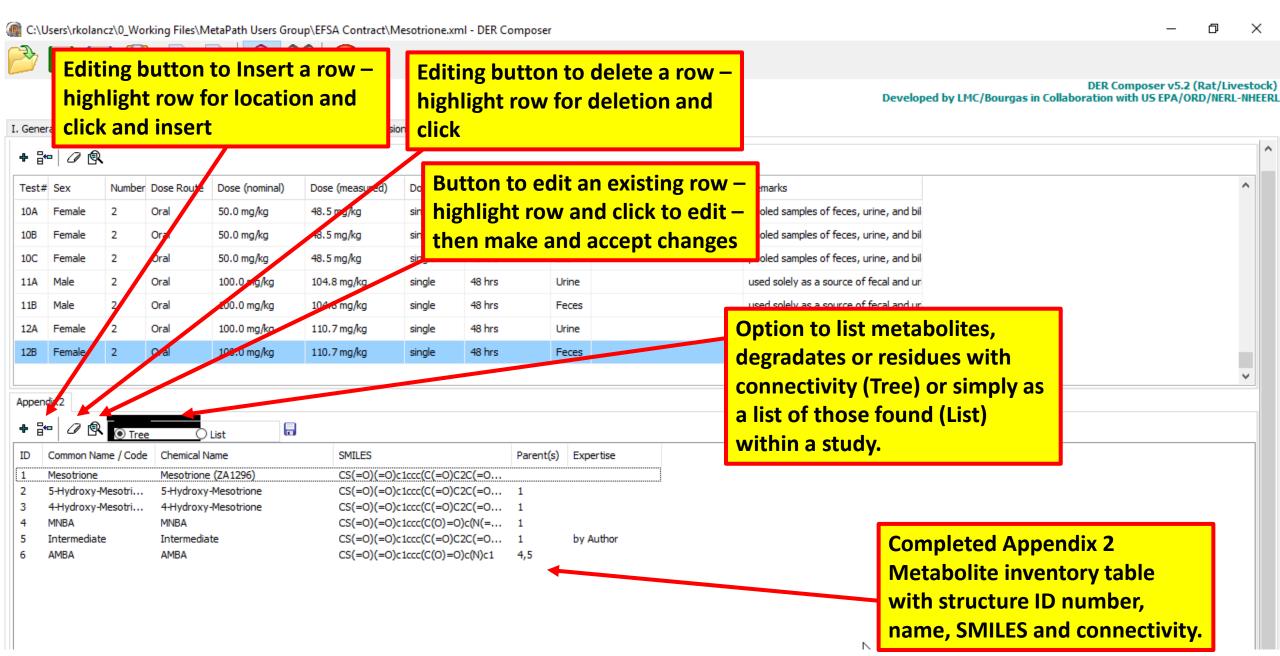


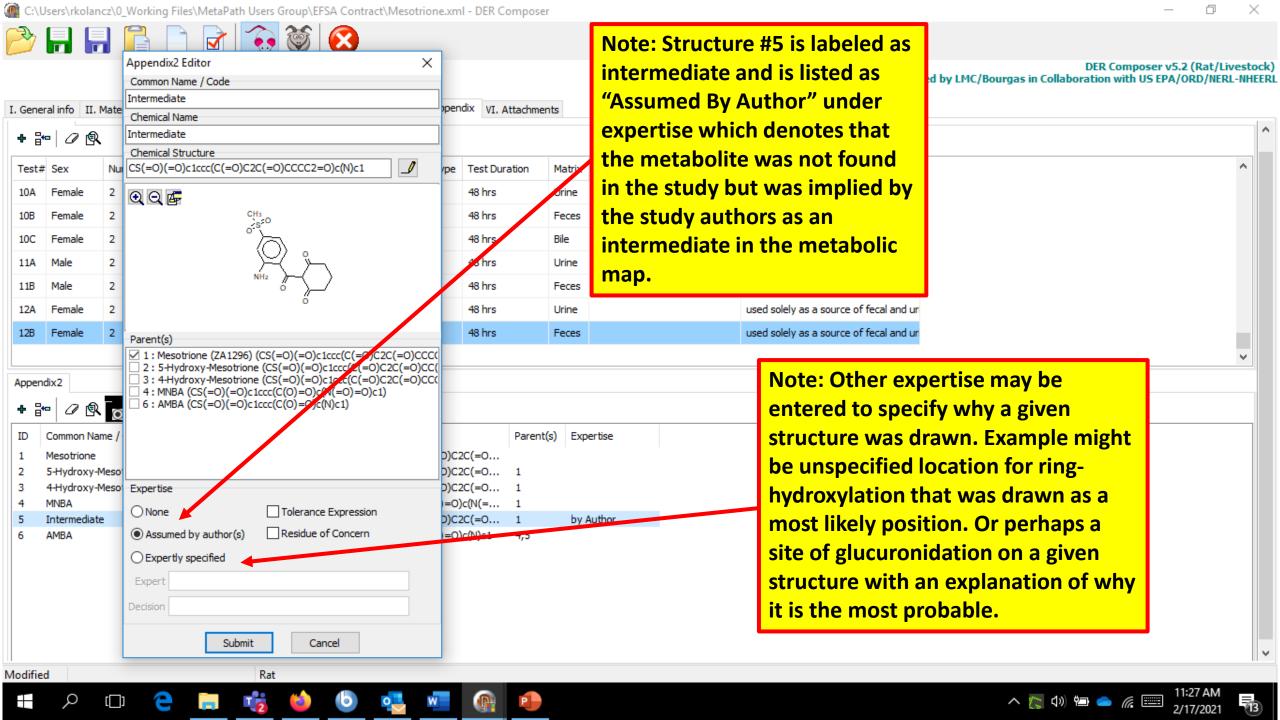
Simply click on the atom in the structure that you wish to replace and the substitution will be made.



SMILES for metabolite is entered into EDITOR. Add metabolite name to Chemical Name and check affiliation box of parent structure for this metabolite. Hit Submit to accept. C:\Users\rkolancz\0_Working Files\MetaPath Users Group\EFSA Contract\Mesotrione.xml - DER Composer Х Appendix2 Editor DER Composer v5.2 (Rat/Livestock) Developed by LMC/Bourgas in Collaboration with US EPA/ORD/NERL-NHEERI Common Name / Code 5-Hydroxy-Mesotrione I. General info II. Mate pendix VI. Attachments Chemical Name 5-Hydroxy-Mesotrione **Check box here to denote** Chemical Structure Nul CS(=0)(=0)c1ccc(C(=0)C2C(=0)CC(0)CC2=0)c(N(=0) Test# Sex Test Duration Matrix Experimental Descriptor connectivity of structure 2 (5-Female 48 hrs Urine QQ 👺 Hydroxy-Mesotrione), the 48 hrs Female Feces metabolite of the parent Female 48 hrs Bile structure 1 (Mesotrione). Urine Male 48 hrs 48 hrs Feces Female 48 hrs Urine Then hit submit. Female 48 hrs Parent(s) **Note: In more complex maps** Appendix2 the same metabolite may originate from more than one Parent(s) Expertise D)C2C(=0... Mesotrione source. Expertise None Tolerance Expression Assumed by author(s) Residue of Concern Expertly specified Expert Decision Submit Cancel

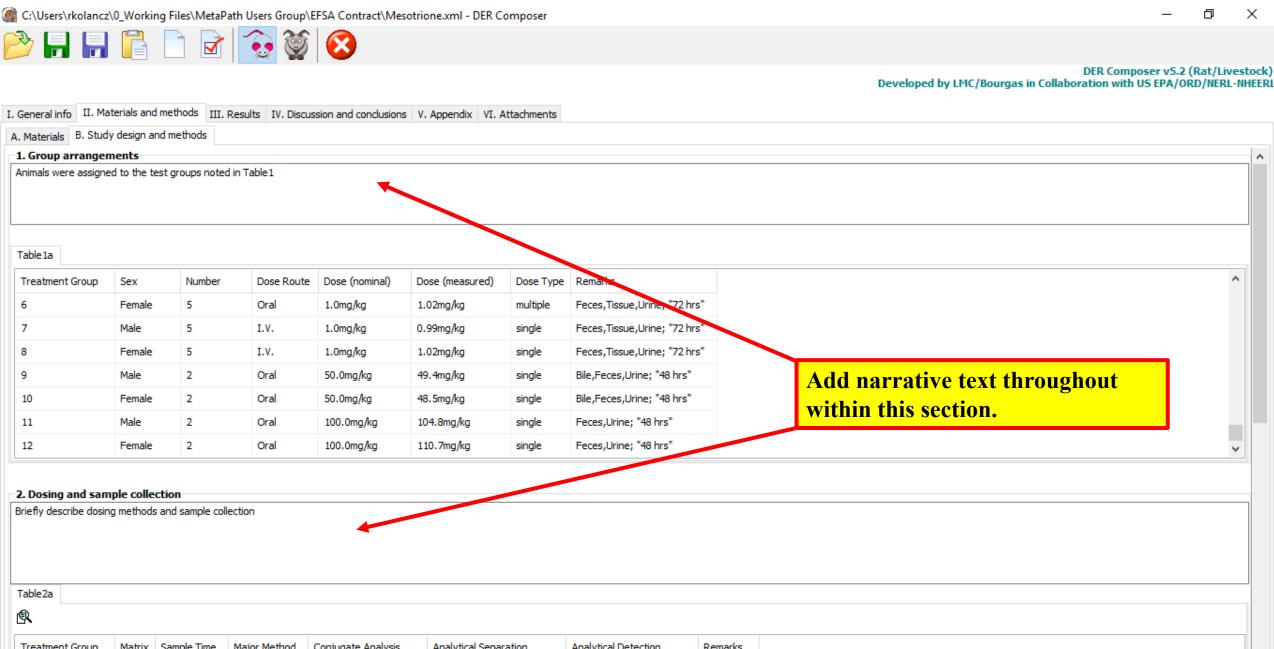
Continue filling in structures with connectivity information until the resulting table is sufficiently completed to represent the metabolic map.





Next go back to II. Materials and methods tab & sub-tab B. Study design and methods and fill in narrative text sections under 1. Group arrangements and 2. Dosing and sample collection. Tables 1a auto-populates from Appendix 1.

C:\Users\rkolancz\0 Working Files\MetaPath Users Group\EFSA Contract\Mesotrione.xml - DER Composer

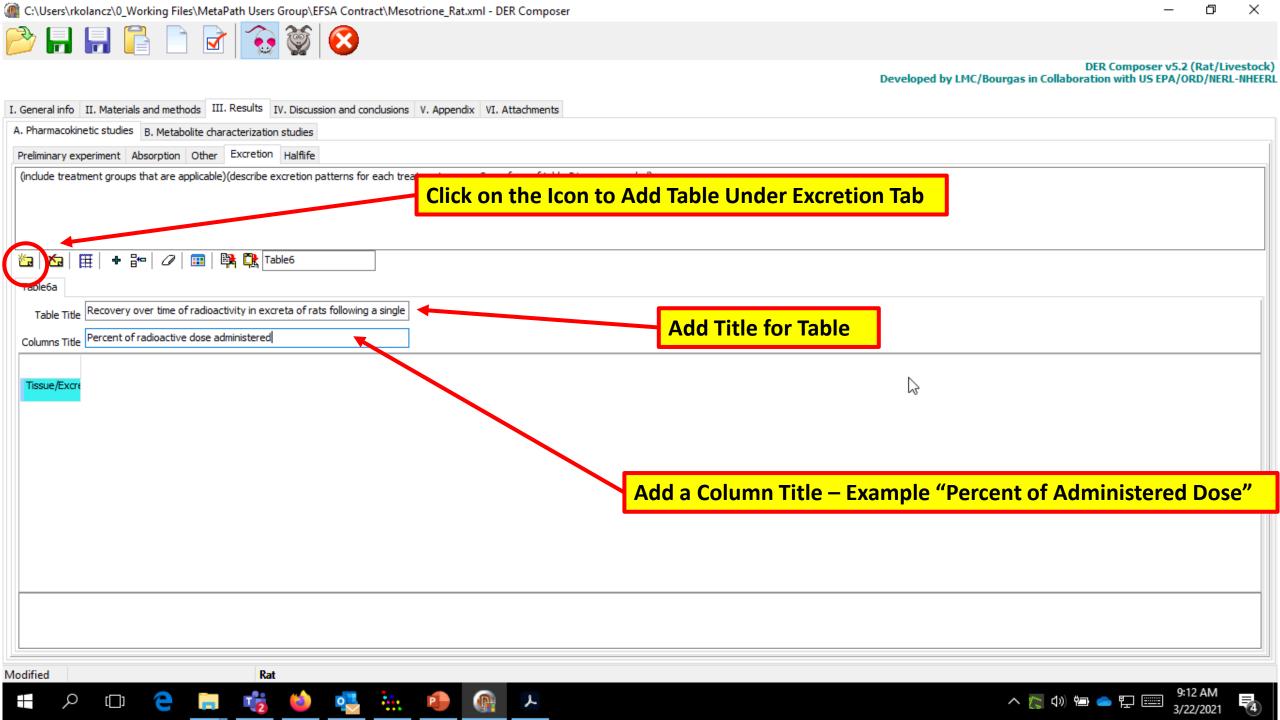


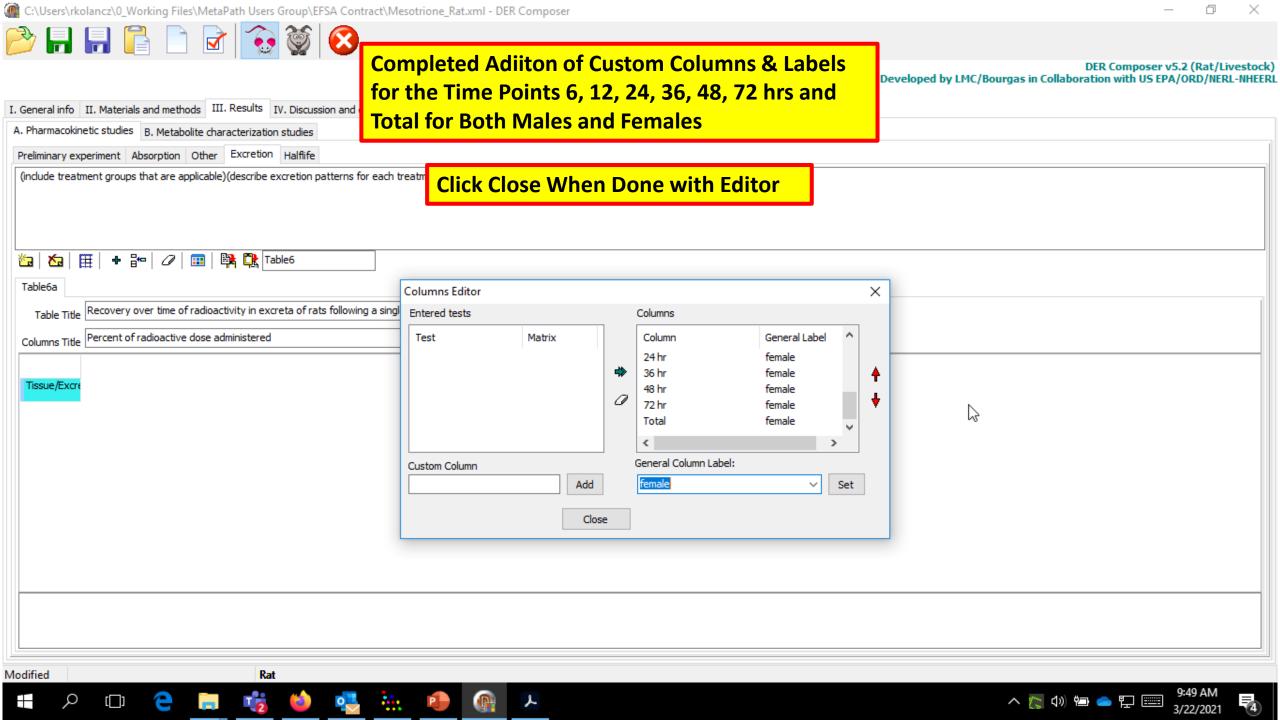
Down further in II. Materials and methods tab & sub-tab B. Study design and methods Tables 2a auto-populates part way from Appendix 1. 🤡 🖫 😂 DER Composer v5.2 (Rat/Livestock) Developed by LMC/Bourgas in Collaboration with US EPA/ORD/NERL-NHEERL A. Materials B. Study design and methods Table2 Editor Х Matrix Sample Time **®** Urine Sample Process Major Method Sample Process Conjugate Analysis Major N Treatment Group Matrix | Sample Time tection Remarks 10A, 1A, 12A, 1A, 2A Urine Analytical Separation Analytical Detection 10B, 1 B, 12B, 1B, 2B, Feces Remarks 1C,2C, C,4C,5C,6C Tissue 10C,9C Bile Submit Cancel a. Pharm cokinetic studies [Briefly desaribe how samples were handled after harvesting (shipment, storage, etc.) and any preparation that was done prior to extraction.] [If warranted, include a graphic (i.e., flowchart) of the extraction and fractionation schemes and omit following textual description.] [Briefly describe the extraction, fractionation and hydrolysis strategies for each tissue. The description should include solvents used (ratios), the order of their use, the extraction procedures employed (i.e., blending, maceration, Soxhlet, etc.) and procedures used to release bound and conjugated residues (i.e., acid, base, or enzyme hydrolysis, exhaustive extraction, etc.). Has the petitioner justified the use of severe conditions (e.g., strong acid hydrolysis in Finnish adding text b. Metabolite characterization studies [Briefly describe he principle of the methods used for identification/characterization of the residues. Specify instrumentation (LSC, TLC, GLC, HPLC, etc.) and detaction method used (UV, ECD) ound residues as in protein or lipid fractions. To finish filling out information in Table 2a, move cursor to line in table to be edited and click this EDIT button. The blue-box Table 2 nent that the Reviewer considers the analyses used to be appropriate. If inappropriate, provide alternative/rationale] **EDITOR** will come up. Make edits and click Submit. Modified Rat ^ [5] (1) 1 = - (6] (IIII

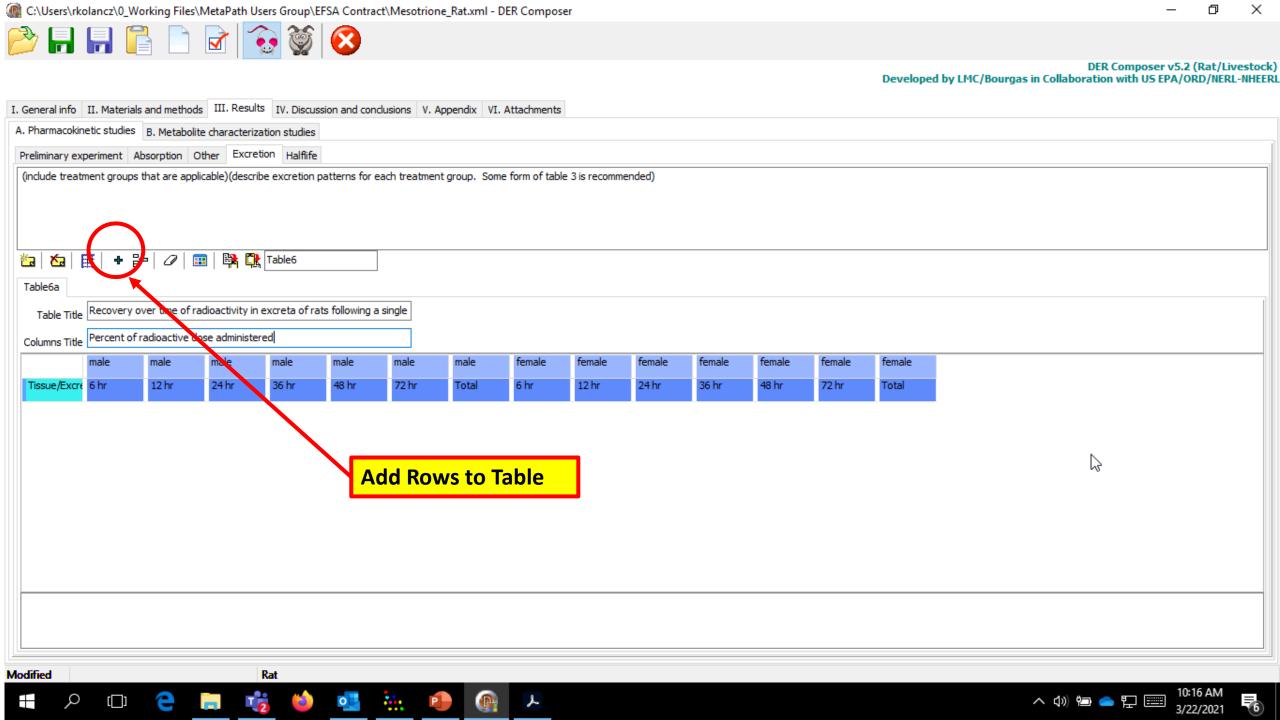
Results

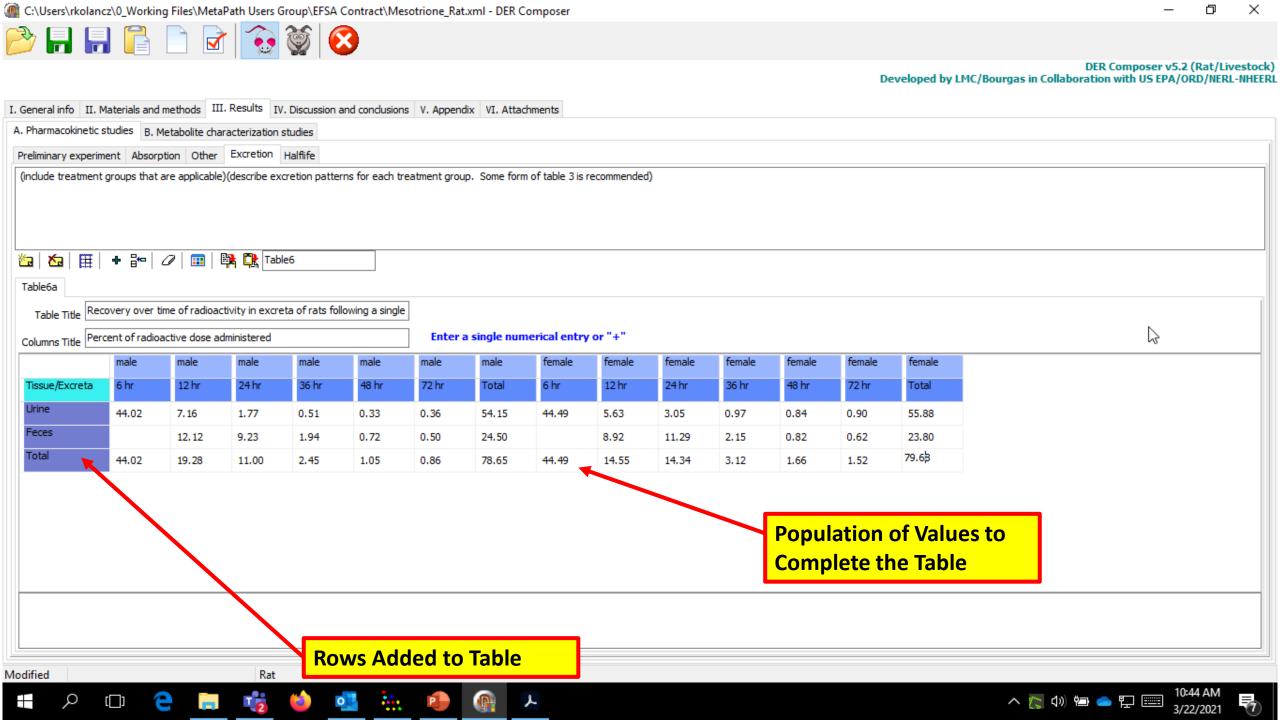
(Pharmacokinetic studies)

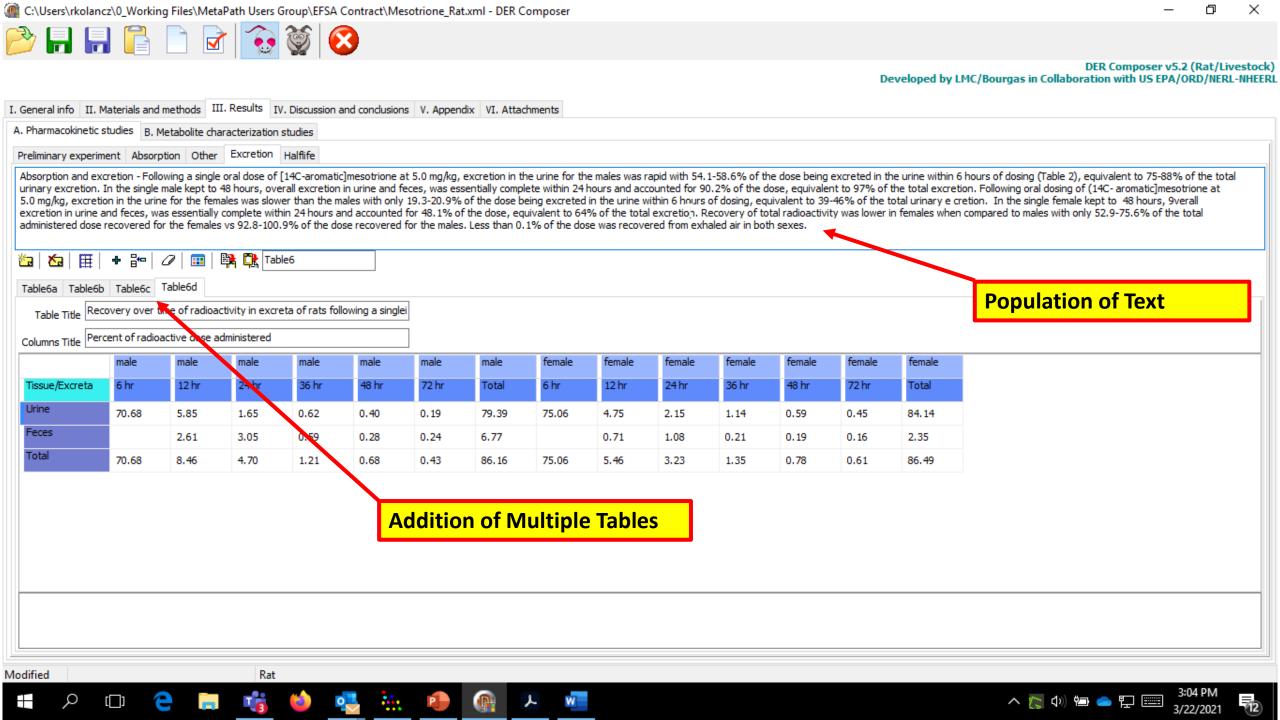
X III. Results tab & sub-tab A. Pharmacokinetic studies. There are sub-tabs for preliminary experiment, Absorption, Other, Excretion, and Half-life. Each will display a sample table and table construction will essentially follow the same process for each tab. Below are functions of button bar icons. Developed by LMC/Bourgas in Collaboration with US EPA/ORD/NERL-NHEERL I. General info II. Materials and methods III. Results IV. Discussion and conclusions V. Appendix VI. Attachments A. Pharmacokinetic studies B. Metabolite characterization studies Preliminary experiment Absorption Other Excretion Halflife Sample Table (if applicable) (Briefly describe results) Icon – To Hide or Show Sample Table immary of Storage Conditions - Example Matrix | Matrix | Actual Storage Duration | Interval of Demonstrated Storage Stability Icon – To Copy a Whole Table + 🔐 🕖 🔠 🖺 🟗 Table Title Icon – To Paste a Whole Table olumns Title Icon – Delete a Row Icon – Manage Columns Icon - Insert a Row Icon - Delete Table Icon - Add a Row Rat Icon – Add Table へ 燯 🗘 垣 🧒 🖫 📟





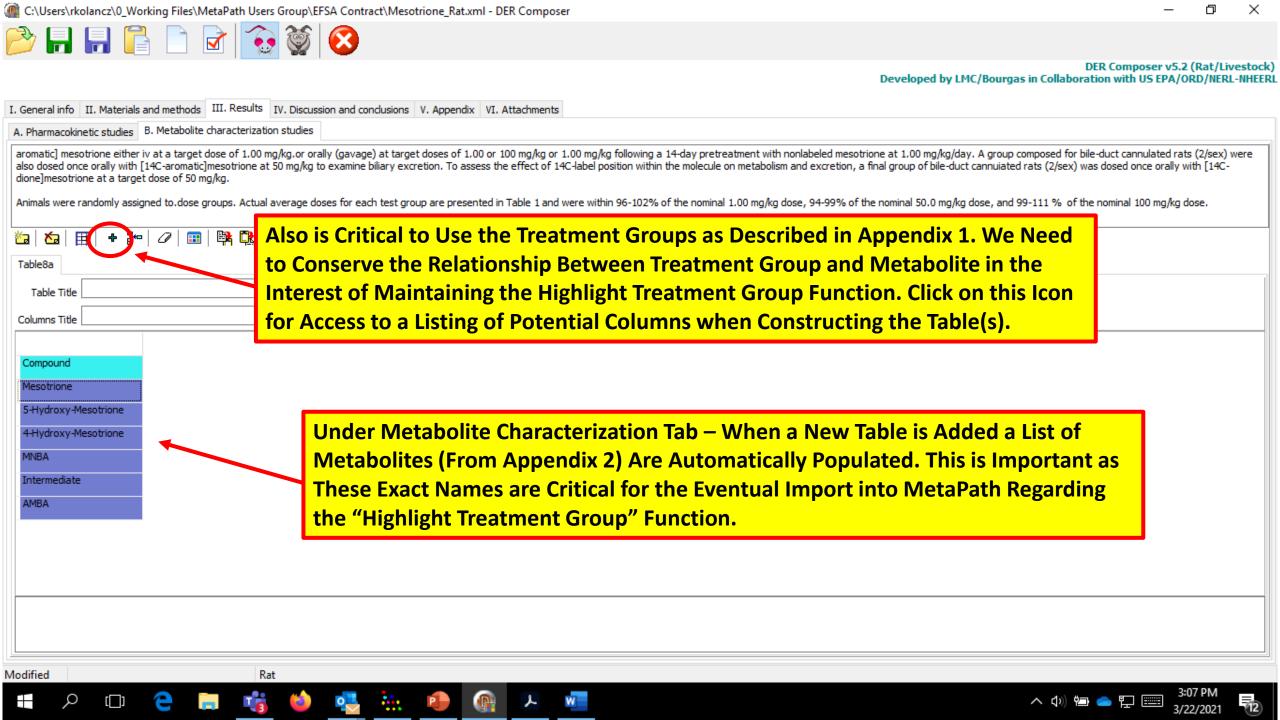


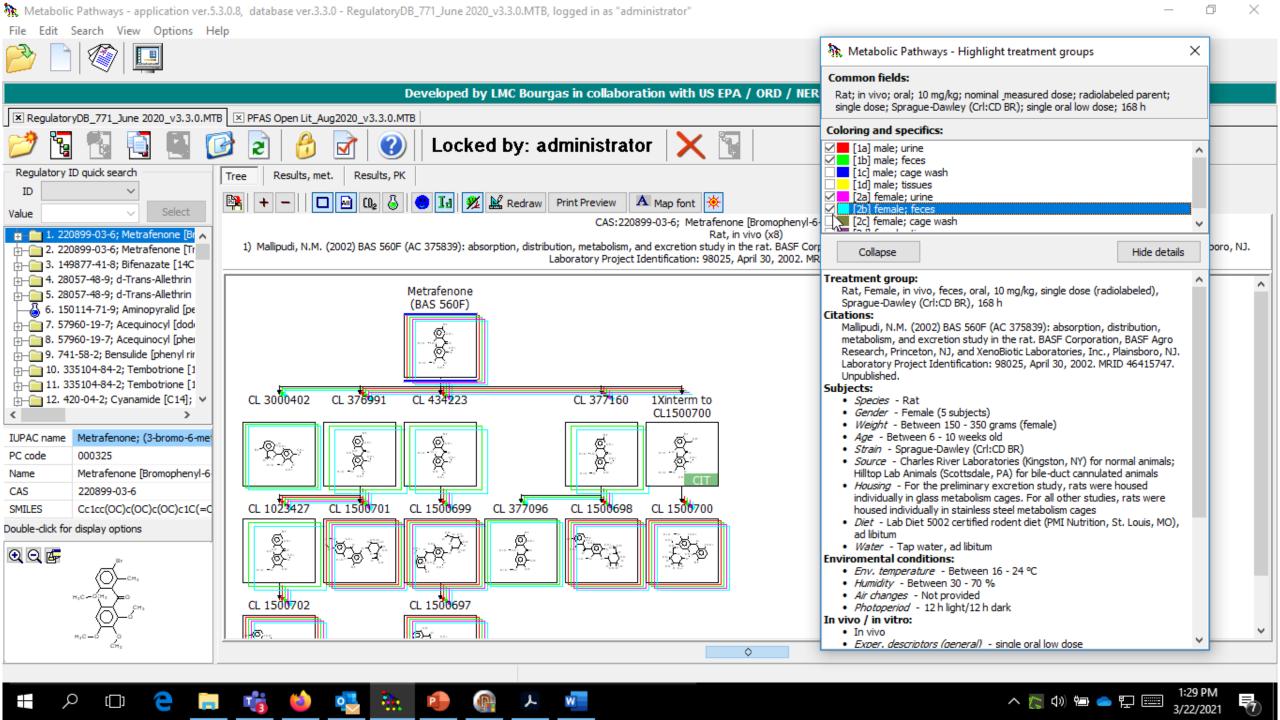


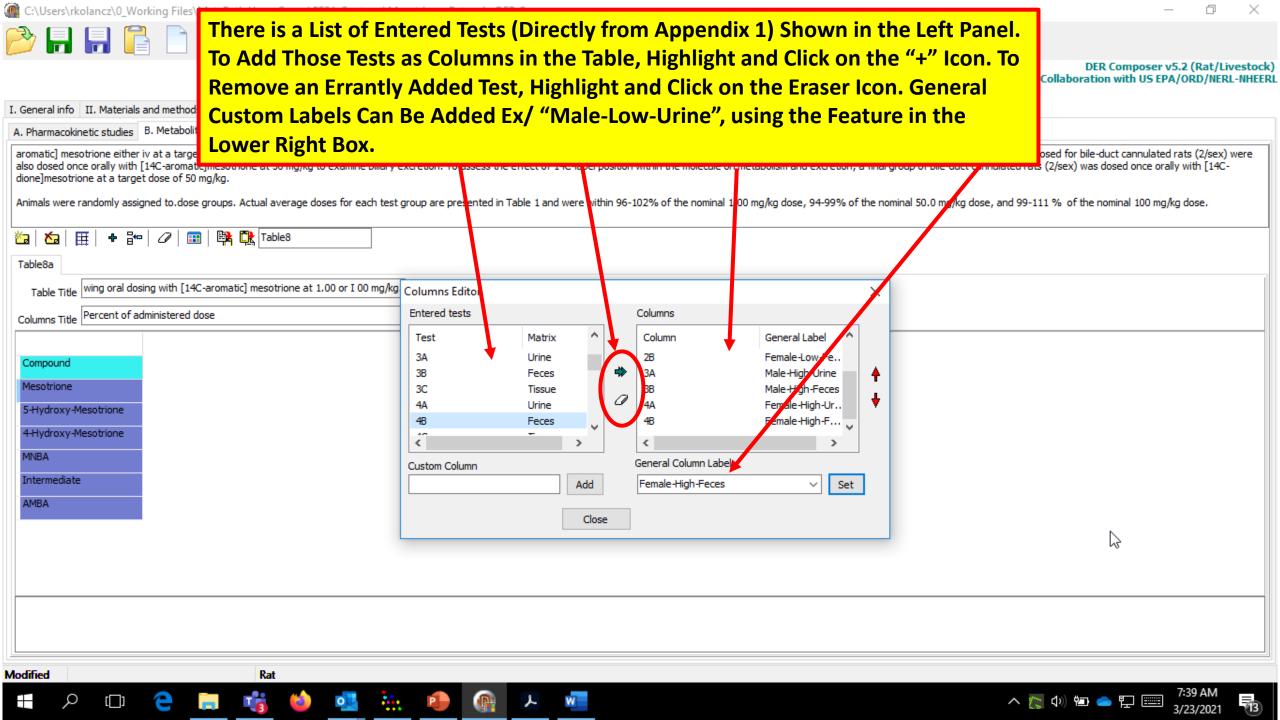


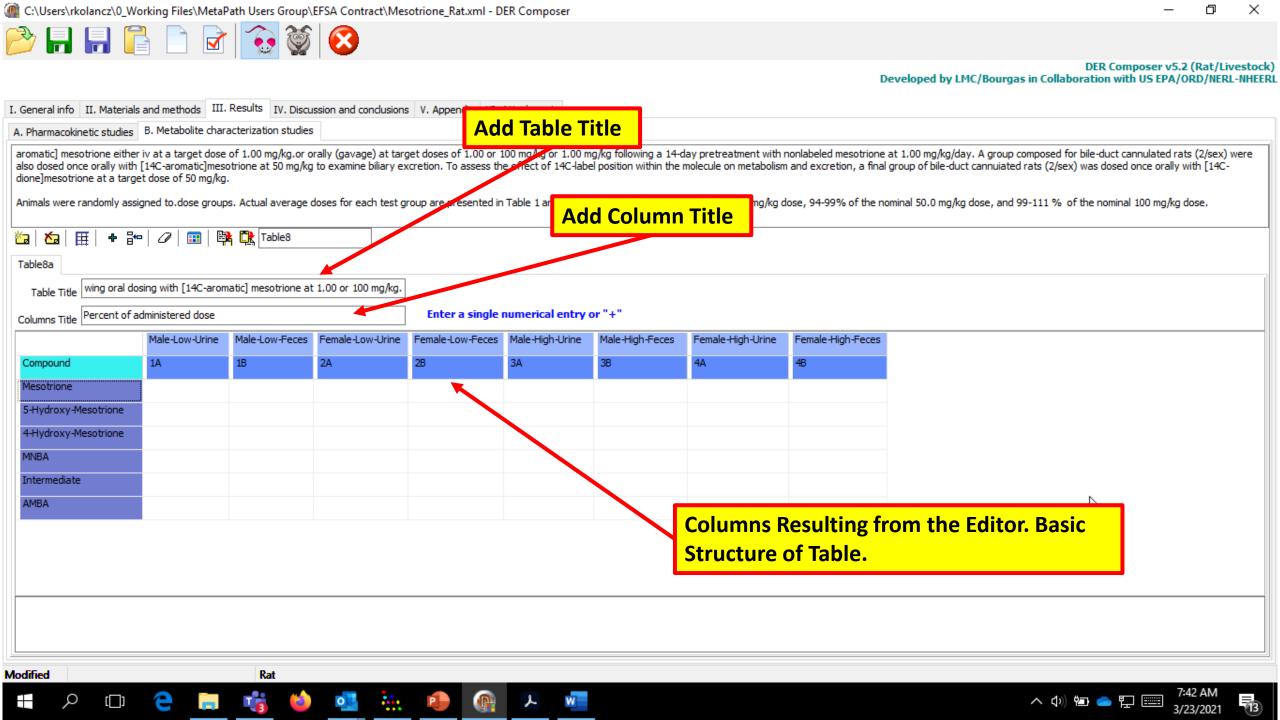
Results

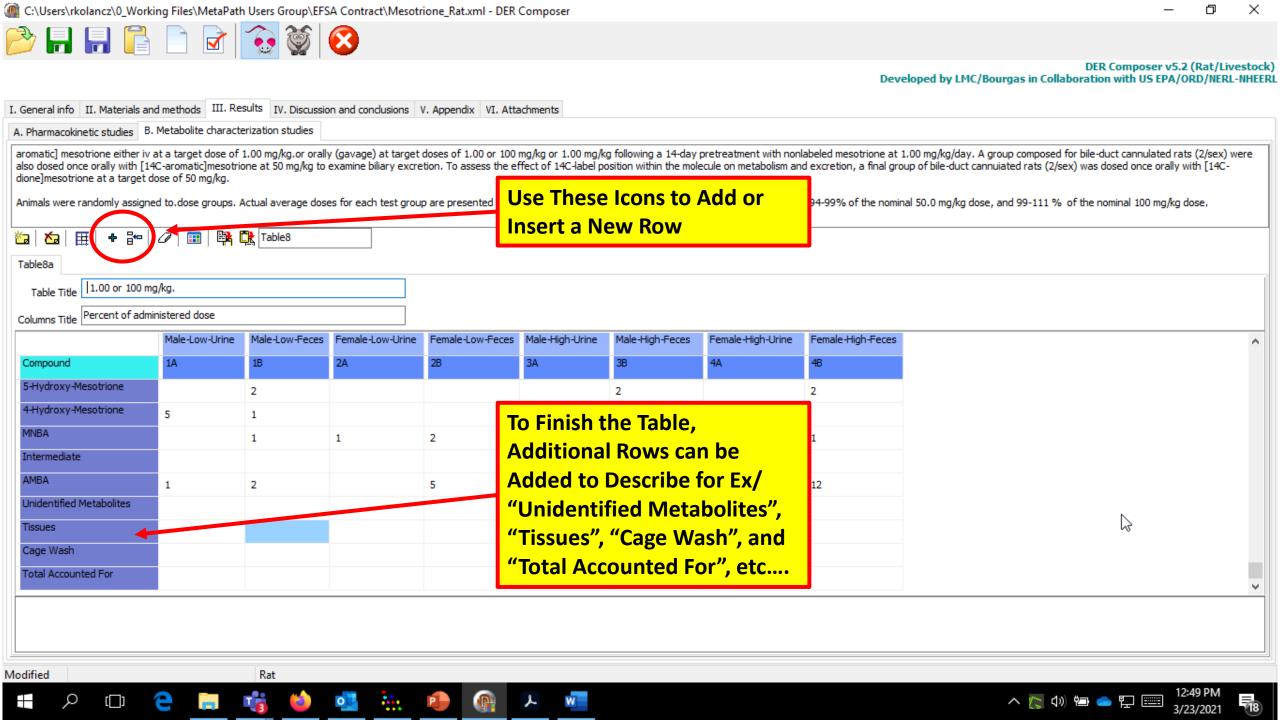
(Metabolite characterization studies)





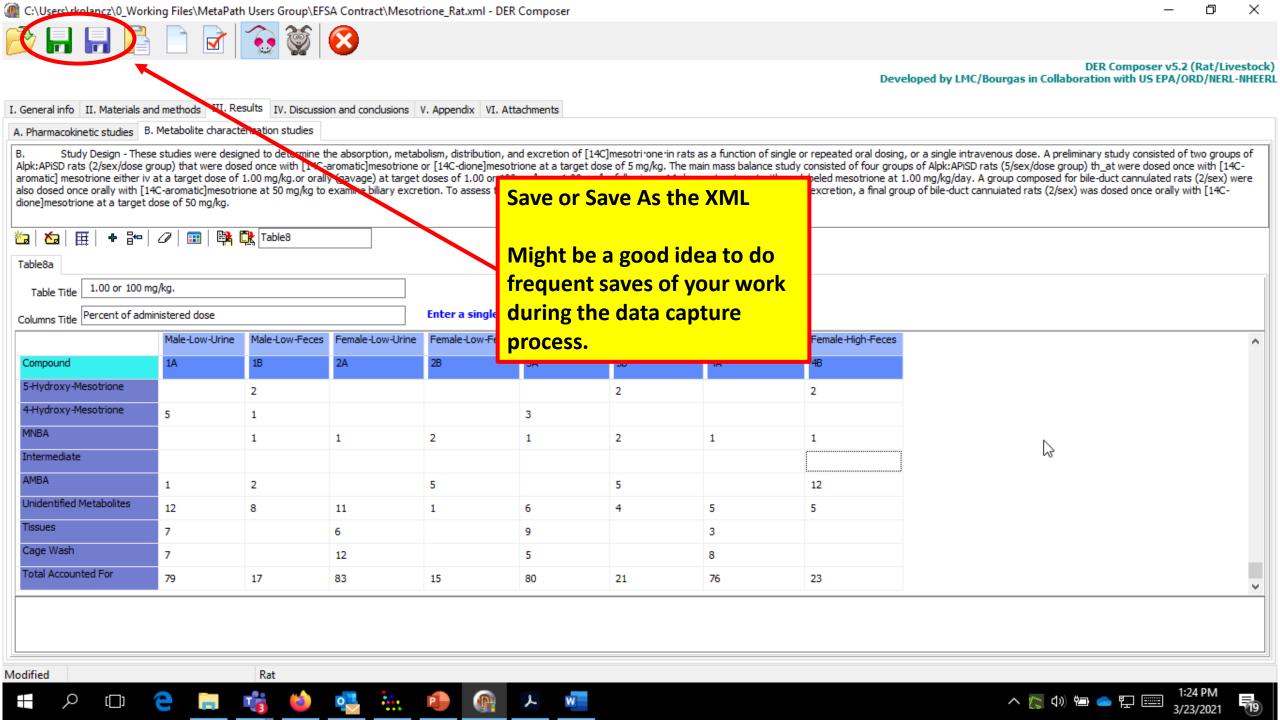






Conclusions

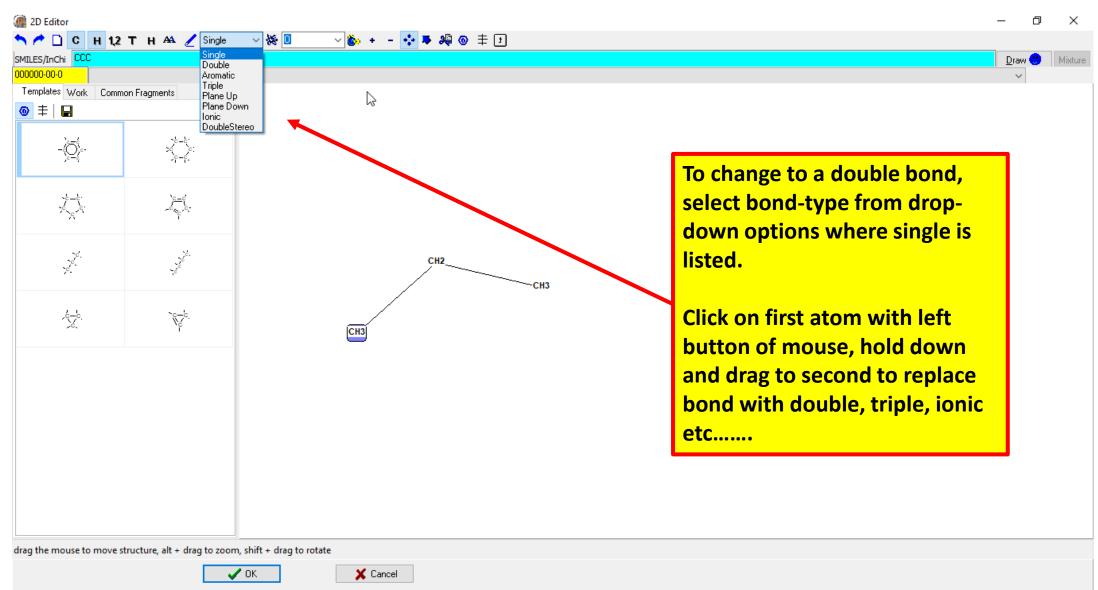
- XML
- .DOC File Generation of Report



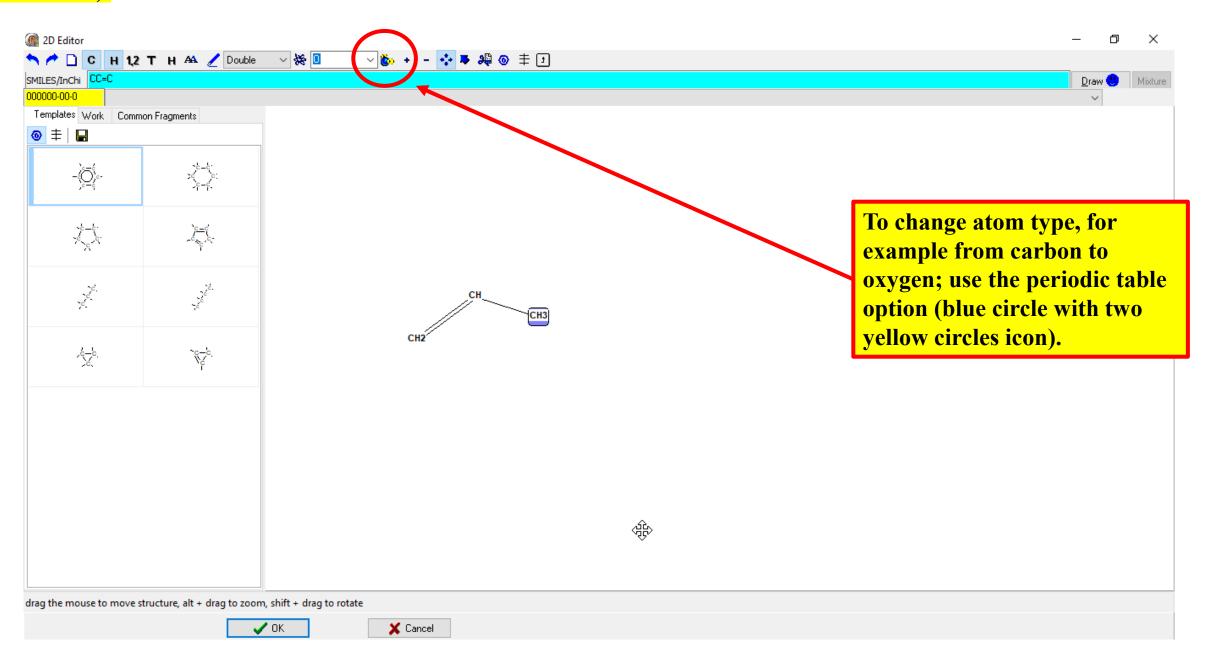
- Drawing Tools
- Structure Editor

STRUCTURE EDITING

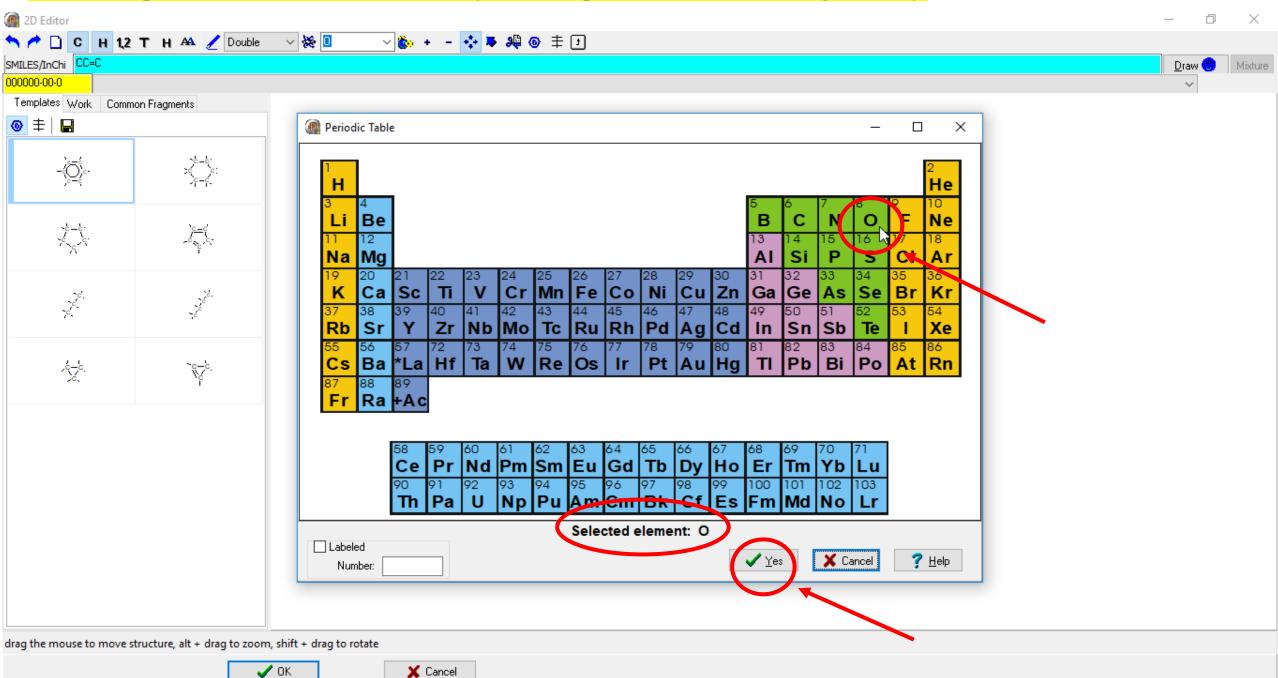
The following screen-shots illustrate some other functions of the STRUCTURE DRAWING package that may be used to modify/edit/draw 2-D structures of parent/metabolites.



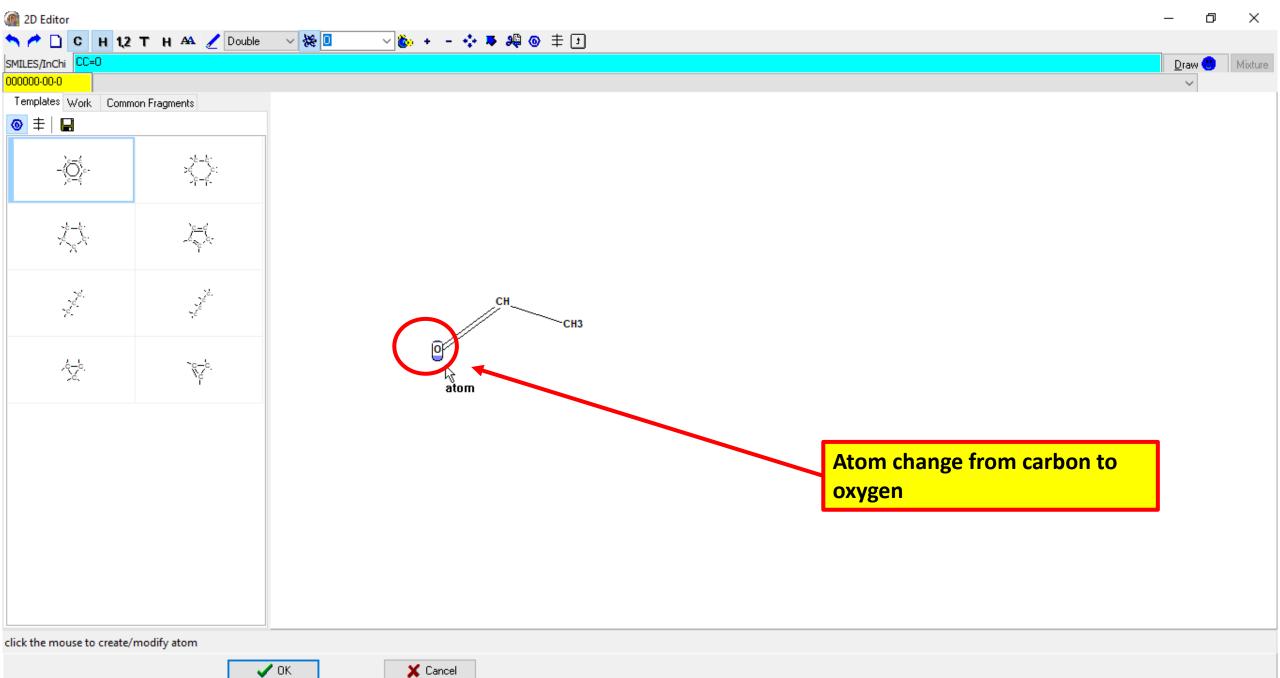
To change atom type, for example from carbon to oxygen; use the periodic table option (blue circle with two yellow circles icon).

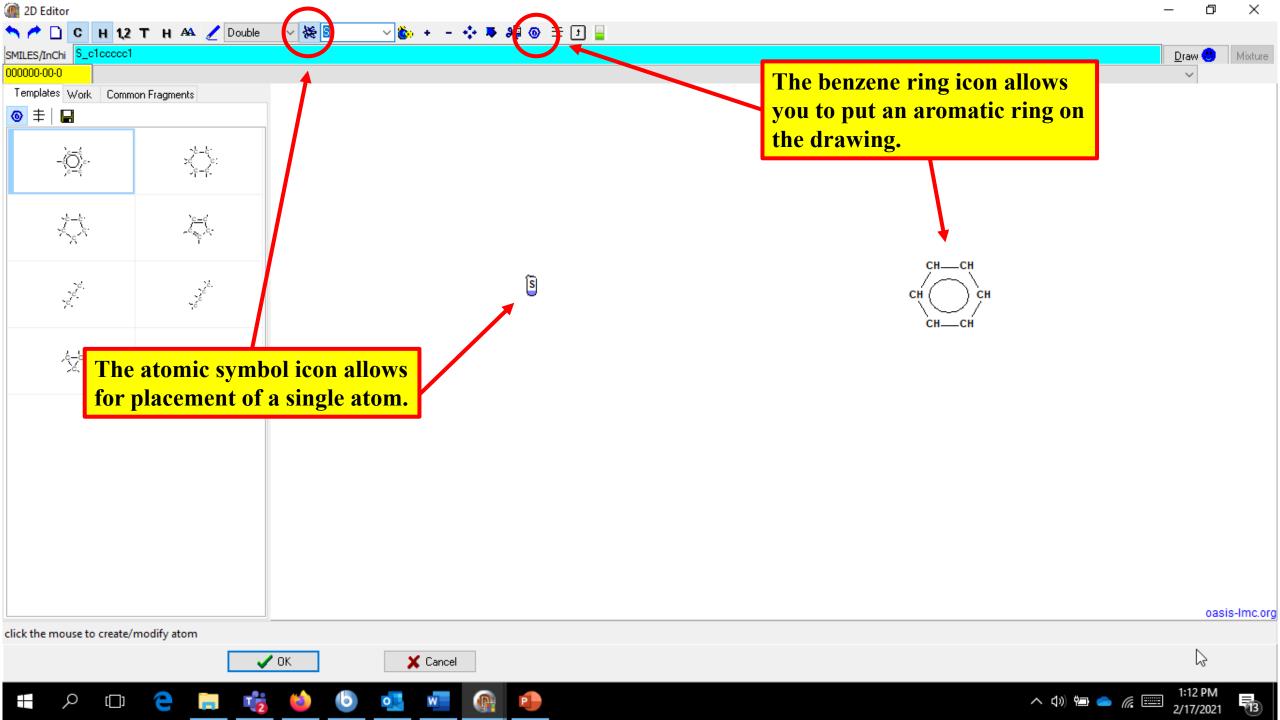


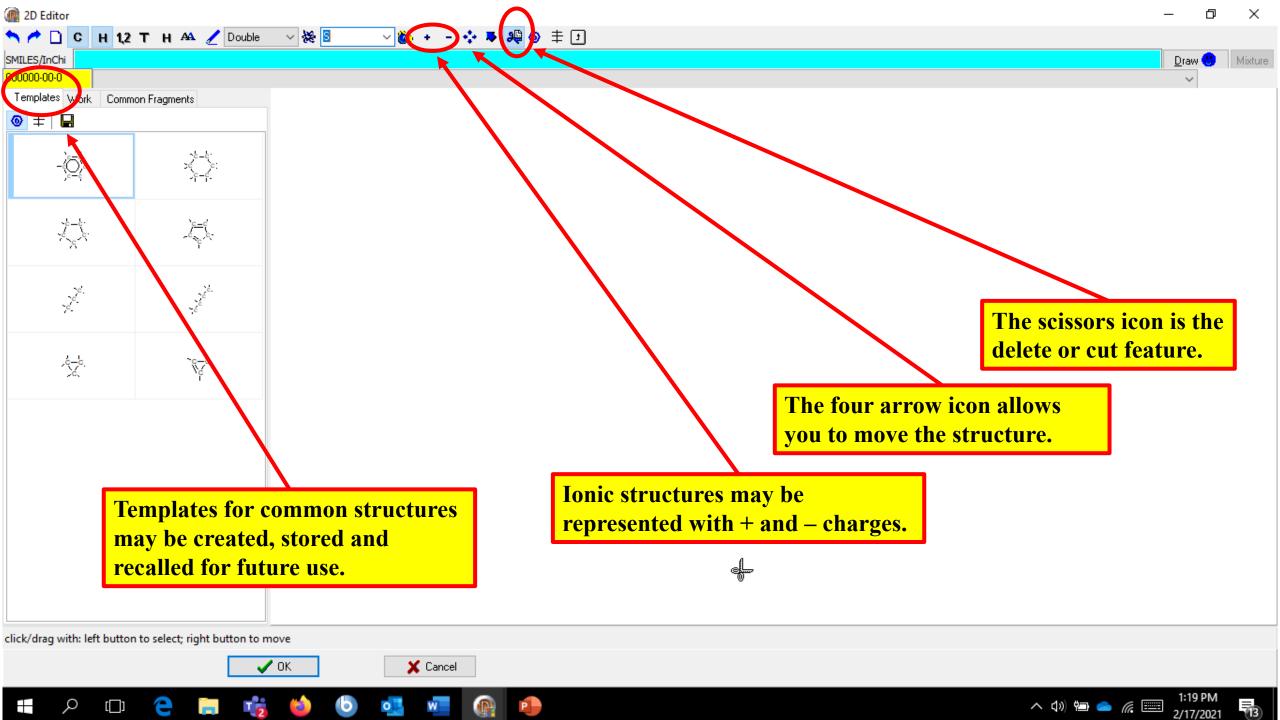
The table opens, click on atom choice, click yes to accept choice and the table goes away.



Simply click on the atom in the structure that you wish to replace and the substitution will be made.







Once a structure is drawn, the SMILES string will be auto-generated for that structure.

✓ OK

🗶 Cancel

