The <u>S</u>tochastic <u>H</u>uman <u>E</u>xposure and <u>D</u>ose <u>Si</u>mulation Model: High-Throughput

SHEDS-HT Beta Version 0.1.10

Quick Start Guide

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ACKNOWLEDGMENTS AND DISCLAIMER

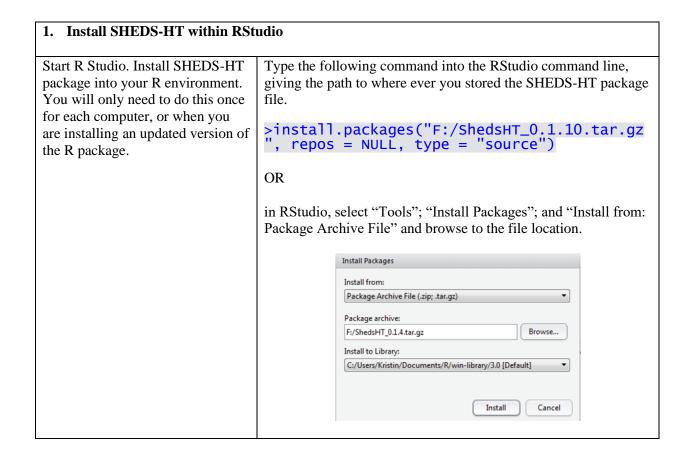
The United States Environmental Protection Agency through its Office of Research and Development funded and collaborated in the research and development of this software, in part under Contract EP-C-14-001 to ICF International. The model is publicly available in Beta version form. All input data used for a given application should be reviewed by the researcher so that the model results are based on appropriate data sources for the given application. This model, default input files, and R package are under continued development and testing. The model equations and approach are published in the peer-reviewed literature (Isaacs et al. Environ. Sci. Technol. 2014, 48, 12750-12759). The data included herein do not represent and should not be construed to represent any Agency determination or policy.

This tutorial will guide you through 1) Installing the SHEDS-HT R Package and corresponding data and 2) running an example run included in the R package (a run of chemicals identified via MSDS sheets as present in various categories of consumer products). This tutorial assumes that you have already:

- Installed R
- Installed RStudio
- Downloaded the most recent version of the SHEDS-HT R package. The most recent SHEDS-HT release can be obtained https://github.com/HumanExposure/SHEDSHTRPackage. Navigate to the folder "R Package", click on the current file name (e.g., "ShedsHT_0.1.X.tar.gz), and select the "Download" button:



Most of the below steps will only need to be done once. Also included with the SHEDS-HT distribution package is a script titled that, with editing for paths, will perform the steps in this tutorial.



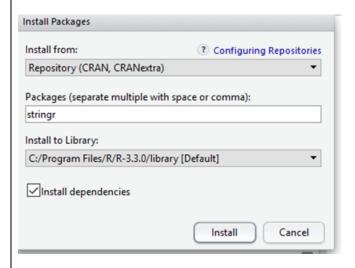
Load other R packages required by SHEDS-HT (**data.table**, **stringr**, **plyr**, and **ggplot**). You will only need to do this once for each computer.

Type the following commands into the RStudio command line

```
>install.packages("data.table")
>install.packages("stringr")
>install.packages("plyr")
>install.packages("ggplot2")
```

OR

in RStudio, select "Tools"; "Install Packages"; and "Install from: Repository" and type in a package name. Repeat for all 4 packages.



2. Set up a SHEDS-HT Run

Load the ShedsHT Package functions into the current R Studio session. (This must be done every time you start a new R session.)

Type the following command into the RStudio command line:

>library(ShedsHT)

Decide on a SHEDS home folder location for storing your input and output files for any runs. This is where SHEDS will create "inputs" and "output" subfolders. You can create as many SHEDS-HT home folders as you want (with any name), accessing them at different times via the setup() step below. Here, for example, is a SHEDS home folder called "C:/SHEDSFORTESTING". (Note that the inputs and output



folders will be created by SHEDS)

Run the "setup" function on the SHEDS home location folder that you created above, so SHEDS knows where to store materials. (This must be done every time you start a new R session or want to switch between Home locations.)

If when running "setup()" you receive the message

Error: cannot open file
'R/Sheds_HT.R': No such file or directory

Or on some systems:

Error in file(filename, "r", encoding = encoding) : cannot open the connection

then you have an old version of SHEDS-HT functions loaded in your R workspace. From the RStudio menu select" Session->Clear Workspace" and try again.

If this is the first time you are using SHEDS, or if you have created a new home location, call unpack(). The default SHEDS input files included in the R package are written into the "inputs" directory.

Note: If you run unpack() again in the same home location it will overwrite any of the ShedsHT package default input files you have altered. (It will not overwrite new files vou may have created - only those with the same names as files included with SHEDS).

Type the following command into the RStudio command line, using the name of your home directory that you created in Step 1:

> setup("C:/SHEDSFORTESTING")

The version information and EPA Disclaimer will display, for example:

ShedsHT Version 0.1.10 (04/23/2024)
Disclaimer
The United States Environmental Protection Agency through its Office of Research and Development
funded and collaborated in the research and development of this software, in part under Contract EP-C-14-001
to ICF International. The model is publicly available in Beta version form. All input data used for a given
application should be reviewed by the researcher so that the model results are based on appropriate data
sources for the given application. This model, default input files, and R package are under continued develop
and testing. The model equations and approach are published in the peer-reviewed literature
(Isaacs et al. Environ. Sci. Technol. 2014, 48, 12750-12759). The data included herein do not represent
and should not be construed to represent any Agency determination or policy.

Inputs folder was created in the working directory

>unpack()

This will produce the output:

```
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/activity_diaries.csv
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/chem_props.csv
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/dret_diaries.csv
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/dret_diaries.csv
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/fugacity.csv
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/mpusiology.csv
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/population.csv
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/run_artsandcrafts.tt
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/run_artsandcrafts.tt
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/run_other_sources.txt
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Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/run_copdats/max.txt
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/source_chem_cod.csv
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/source_chem_cod.csv
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/cource_chem_cod.csv
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/source_chem_cod.csv
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/cource_chem_cod.csv
Unpacking C:/Users/kisaacs1/AppData/Local/Programs/R/R-4.2.2/library/ShedsHT/extdata/cource_chem_cod.csv
```

Etc.

3. Perform a SHEDS-HT Run

Call the SHEDS run function with a SHEDS **Run file** as argument. SHEDS comes with several example run files described in the Technical Manual. They are located in the **input** directory of your SHEDS home location. Here, we run the

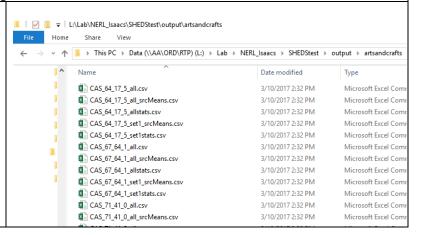
"run_artsandcrafts" example, which runs all the current SHEDS-HT default product composition data for arts and crafts products, which was developed from EPA's CPDat data. Alternatively, one could run all of the default SHEDS-HT data from EPA's CPDat V3 (all product types and chemicals) by calling the run "run_cpdatv3" run file. Note: the run_cpdatv3 run may take 2 or more hours to complete, depending on your computational resources.

>run("run_artsandcrafts.txt")

This will produce the output:

4. Locate the SHEDS-HT Output Files

Examine the SHEDS Output files in the Output folder of the SHEDS home folder you designated with **setup**(). The files will be in a subfolder under "Output" with the run name provided in the run file (E.g., either "artsandcrafts" or "CPDatV3max").



5. Combine SHEDS Output for All Chemicals into a Summary File

Combine the Percentile Data for all chemicals (i.e. the "AllStats" files created in the output folder) into a single file for other analyses, using the combine_output function, providing a run name and a name for the output file. The file is placed in the output folder for the run.

combine_output(run.name="artsandcrafts",
out.file="artsandcrafts.csv")

This will produce the output:

```
Processing chemical 1
                      of
                          21
Processing chemical 2
                      of
                          21
Processing chemical 3 of
                          21
Processing chemical 4 of
                          21
Processing chemical 5 of
                          21
Processing chemical 6 of
                          21
Processing chemical 7
                     of
                          21
Processing chemical 8 of
                          21
Processing chemical 9 of
                          21
Processing chemical 10 of
                           21
Processing chemical 11 of
                           21
Processing chemical 12
                           21
Processing chemical 13
                           21
Processing chemical 14 of
                           21
Processing chemical 15 of
                           21
Processing chemical 16 of
                           21
Processing chemical 17
                           21
                       of
Processing chemical 18 of
                           21
Processing chemical 19 of
                           21
Processing chemical 20 of
                           21
Processing chemical 21 of
                           21
Combining data...
```

5. Visualize Run Outputs

Due to the fact that the outputs are so rich, custom visualization of SHEDS results are usually desirable, but SHEDS does provide a two simple functions (see User Manual Section 5) for visualizing results. These include allstats.variable.rank.plot which provides a plot of select output variables and statistical metrics across chemicals.

```
allstats.variable.rank.plot("artsandcrafts"
, cohort.col = c("Total",
"Females"),output.variable =
"abs.tot.mgkg", metrics =
c("5%","50%","95%"),label_chem=T)
```

This will produce the output:

