

httk: High-Throughput Toxicokinetics Virtual Training

Breakout Group Worksheet: Beginner

This worksheet was developed for the Breakout Group session of the httk Virtual Training, hosted by the U.S. Environmental Protection Agency's Center for Computational Toxicology and Exposure on November 8-9, 2023.

For more information about httk:

- Visit the httk package manual: <http://r-project.org>
- Explore httk vignettes: <https://CRAN.R-project.org/package=httk>
- Learn more about EPA's [Rapid Chemical Exposure and Dose Research | US EPA](#)
- Read more in
 - Wambaugh et al. (2018): <https://doi.org/10.1093/toxsci/kfy020>
 - Pearce et al. (2017): [10.18637/jss.v079.i04](https://doi.org/10.18637/jss.v079.i04)
- For questions, contact: NAM@epa.gov

Goals

- Use R's package httk to explore toxicokinetics of a compound of interest.
- Identify physiological and compound-specific parameters.
- Simulate tissue concentrations of chemicals using different exposure scenarios.
- Extract experimental dosing and time course data from data provided in Wambaugh et al. (2018) and replicate the experiment in httk.

Tools

- R software environment (release \geq 2.10) and a graphical user interface (RStudio recommended)
- R packages:
 - httk (version 2.2.2)
 - readxl
- Excel file of experimental data: BeginnerFiles_toxsci.xlsx. (This file contains subsets=from the Supplementary data provided in Wambaugh et al. (2018): <https://doi.org/10.1093/toxsci/kfy020>).

Hints

- Navigate to the "help" tab in the pane in the bottom right corner of your Rstudio screen. Here, you can search for httk functions. You can also type "help([function])" into the

console. For example, type “help(solve_pbt)” into the console and you will see the help page for the pbt model.

- Chemical-specific data and in vitro data for a compound can be found in the table “chem.physical_and_invitro.data”. One way to subset this table and see data for a specific chemical is the following command:
- `subset(chem.physical_and_invitro.data, Compound==chem.name)`

```
# Load packages
```

```
# Make sure the following packages are installed:
```

```
#readxl, httk, ggplot2, car, openxlsx
```

```
library(readxl)
```

```
## Warning: package 'readxl' was built under R version 4.1.3
```

```
library(httk)
```

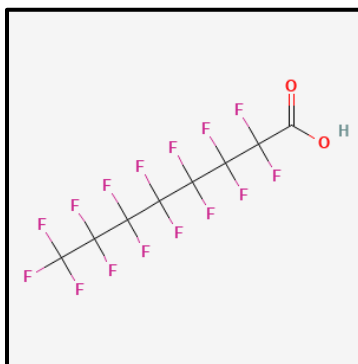
```
## Warning: package 'httk' was built under R version 4.1.3
```

```
rm(list=ls()) #To start with a clean environment, clear previous calculations.
```

Directions

A. Chemical of Interest

Perfluorooctanoic acid



Perfluorooctanoic acid (PFOA) is a perfluorinated carboxylic acid, belonging to the per- and polyfluorinated substances (PFAS) group of chemicals. It is a lab-made compound that began being used commercially in the mid-1950s as an industrial surfactant. Other applications include textiles and floor wax. Low levels of PFOA can be found in drinking water, some foods, and household dust.

Preferred IUPAC Name: Pentadecafluorooctanoic acid



Other names: Perfluorooctanoic acid, PFOA, C8, Perfluorooctanoate, PFO, Perfluorocaprylic acid, C₈-PFCA, FC-143, F-*n*-octanoic acid

Name in httk: Perfluorooctanoic acid

CASRN: 335-67-1

DSSTox substance identifier (DTXSID): DTXSID8031865

Molecular Formula: C₈HF₁₅O₂

SMILES: OC(=O)C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)C(F)(F)F

1. Confirm the DTXSID and CAS number for the chemical Perfluorooctanoic acid.
2. Say we wish to simulate the concentration of Perfluorooctanoic acid in the liver.
 - a. Which of the following models would be appropriate for this: 1comp, 3comp, pbtk? (**Hint:** Search solve_ 'modelname' in the help file to see a diagram of the compartment models for each model.)
 - b. Which model could we use if we needed concentration in the kidney?
 - c. What compartments does the pbtk model use? What compartments does 3comp use?
3. Find chemical and physical parameters within httk.
 - a. Find logHenry and logP for this chemical (**Hint:** Search chem.physical_and_invitro.data or get_physchem_param)
 - b. Find human and rat plasma fraction unbound. [**Hint:** Use chem.physical_and_invitro.data]
 - c. Run parameterize_pbtk or parameterize_3comp to find the fraction unbound (funbound.plasma) in the plasma for rabbits. Were you able to find a value or did you run into an error?
 - d. Say that you still want to generate parameters and simulate solutions for rabbits. Mitigate the issue from part (c) and find the funbound.plasma value (**Hint:** Use the "default.to.human" input.)

4. We are now going to simulate Perfluorooctanoic acid tissue concentrations to mirror the experimental data found in Wambaugh et al. (2018). The excel file `BeginnerFiles_toxsci.xlsx` contains multiple data points for multiple experimental chemicals, including Perfluorooctanoic acid.
 - Load the excel file.
 - Make sure the `readxl` package is installed [`install.packages("readxl")`] and loaded into R [`library("readxl")`]
 - Read the `.xlsx` file into R and name it: `my_data = read_excel("path_to_file/ BeginnerFiles_toxsci.xlsx ")`
 - Extract the data pertaining only to Pentadecafluorooctanoic acid. (**Hint:** *You can do this by creating a subset-- `subset(my_data, Compound == chem.name)`*)
- a. Notice that there are multiple routes of exposure and subjects tested for our chemical. Also note that each subject corresponds to one longitudinal experiment in which the subject was given a single dose at the start of the experiment. Our first goal is to compare oral dose `httk` simulations to the oral dose experimental concentrations obtained from subject 81, so we start by extracting experiment specifications for this case.
 - i. Find the oral dose and dose units for subject 81 (**Hint:** *Route="po" pertains to oral dosing.*)
 - ii. What is the output media and output units? What species was studied?
 - iii. How long did the experiment last?
- b. Using these specs, generate oral dose results using both the `3comp` and `pbtk` models and plot them against the experimental data.

Hints:

 - Use the `solve_3comp()` and `solve_pbtk()` functions with appropriate inputs.
 - Output units: `httk` has units in days. use `days = number-of-hours-in-experiment/24`. When plotting, make sure to convert `httk` output units into hours.
 - Note that `ug/mL = mg/L`
 - Make sure to use `iv.dose=FALSE`
 - When plotting overlaid items, you can use the following commands (and make sure units of time are the same):

```
plot(time.experimental, conc.experimental) # will plot points for experimental data
```

```
lines(time.simulation1, conc.simulation1) # will plot a line for simulation 1
```

```
lines(time.simulation2, conc.simulation2) # will plot a line for simulation 2
```

Explain the results. Which model produced results simulations that seem to fit the data well? Consider why this might be the case.

- c. Our second goal is to compare the IV dose experimental data pertaining to subject 85 with IV dose htk simulations. Extract the experiment specifications for IV.
 - i. Find the IV dose and IV dose units for subject 85.
 - ii. What is the output media and output units? What species was studied?
 - iii. How long did the experiment last?
- d. Using these specs, generate iv dose results using both the 3comp and pbtk models and plot them against the experimental data. (**Hint:** *Make sure to use `iv.dose=TRUE`*)
- e. Explain the results. Which model produced results simulations that seem to fit the data well? Consider why this might be the case.

- i. Compare the oral dose and IV dose time curves. Do they make sense?
 - ii. Compare the AUC for both cases and explain similarities or differences.

5. Some chemical exposure occurs frequently. In the case of PFOS, a subject might be exposed to multiple doses over a period of time rather than just one single initial exposure. Say you wish to investigate how multiple exposures affect the concentration of the substance in the plasma by splitting the 1 mg/kg oral dose into 3 even oral bolus doses: 33% of the chemical was given at time 0, 33% was given at 24 hours, and 33% was given at 36 hours. The experiment lasts for 3 days total. (**Hint:** Use a dosing matrix in the solve function and set dose = 0)
 - Example of how to create a dosing matrix. Use the following example of a simple arbitrary dosing matrix to create one for the above scenario.
 - Example: At time = 0 hours a dose of 10 mg/kg was given and at time = 6 hours a dose of 20 mg/kg was given.
dose.mat = matrix(c(0, 0.25, 10, 20), nrow=2)
colnames(dose.mat) = c("time", "dose")
 - a. Use the model you think gave the best fit (3comp or pbtk). Plot the concentration in plasma over time. How does the solution differ from that in [4b](#)?

6. Find conversion factor for chemical from ppmv to uM for PFOA. (**Hint:** Use `convert_units()`)

7. htk contains physiological/biochemical parameters for some or all of the following species: humans, rats, rabbits, and mice. For which species are there parameters for Perfluorooctanoic acid?

8. Find the following parameters using the pbtk model (**Hint:** Use `parameterize_pbtk()`):
 - a. Liver to plasma partition coefficient for PFOA. (**Hint:** Partition coefficients have the suffix "2pu".)

- b. Fractional volume of the kidney and fractional blood flow rate to kidney for an average human.