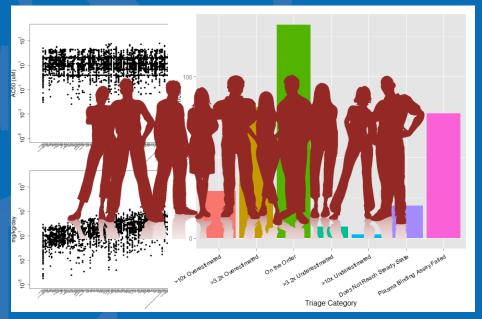


# R Package "httk"

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*Figure includes image from Thinkstock* 

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### Introduction

- Toxicokinetics (TK) provides a bridge between HTS and HTE by predicting tissue concentrations due to exposure
  - Traditional TK methods are resource intensive
- Relatively high throughput TK (HTTK) methods have been used by the pharmaceutical industry to determine range of efficacious doses and to prospectively evaluate success of planned clinical trials (Jamei, et al., 2009; Wang, 2010)
  - A key application of HTTK has been "reverse dosimetry" (also called Reverse TK or RTK)
  - RTK can approximately convert *in vitro* HTS results to daily doses needed to produce similar levels in a human for comparison to exposure data (Wetmore, *et al.*, 2012)

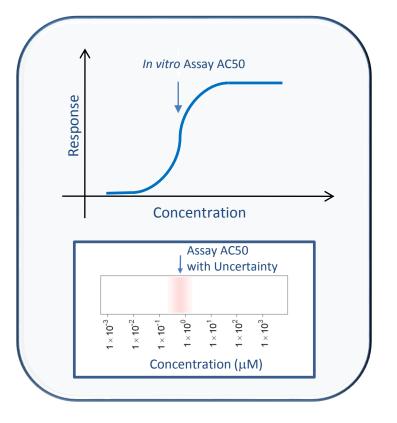


### **High-Throughput Bioactivity**

 Tox21: Examining >10,000 chemicals using ~50 assays intended to identify interactions with biological pathways (Schmidt, 2009)

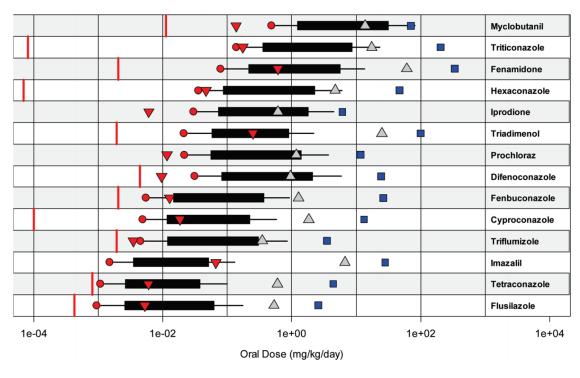


- ToxCast: For a subset (>1000) of Tox21 chemicals ran >500 additional assays (Judson et al., 2010)
- Most assays conducted in doseresponse format (identify 50% activity concentration – AC50 – and efficacy if data described by a Hill function)
- All data is public: http://actor.epa.gov/





### *In vitro* Bioactivity, HTTK, and *in Vivo* Toxic Doses



Comparison of HTTK predicted oral equivalent doses (box and whisker plots in mg/kg/day) with doses for no effect and low effect groups in animal studies

Lowest Observed Effect Level

<sup>A</sup> No Observed Effect Level (NEL)

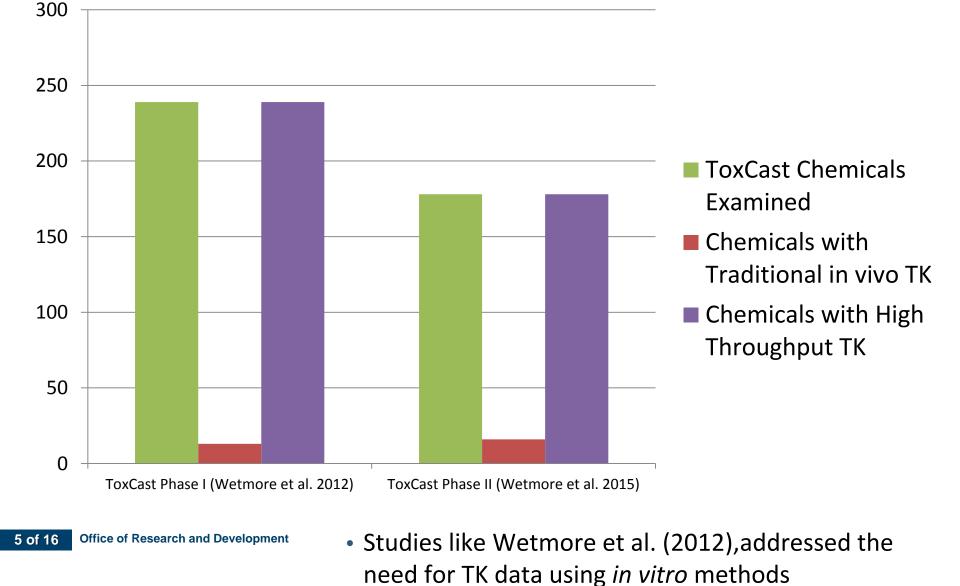
**v** NEL/100

Estimated chronic exposure levels from food residues are indicated by vertical red lines. All values are in mg/kg/day.

Judson et al. (2011)

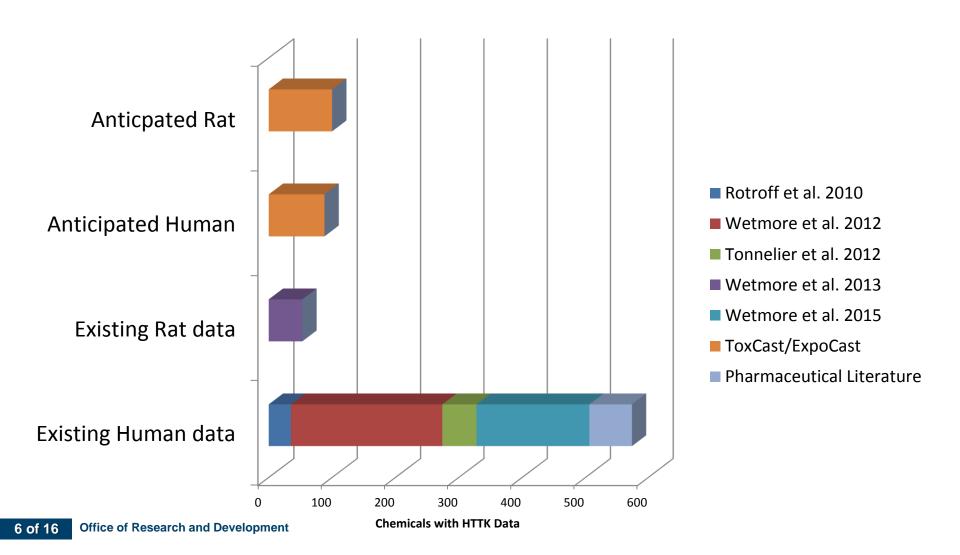


### The Need for *In Vitro* Toxicokinetics





### **Chemicals with HTTK Data**



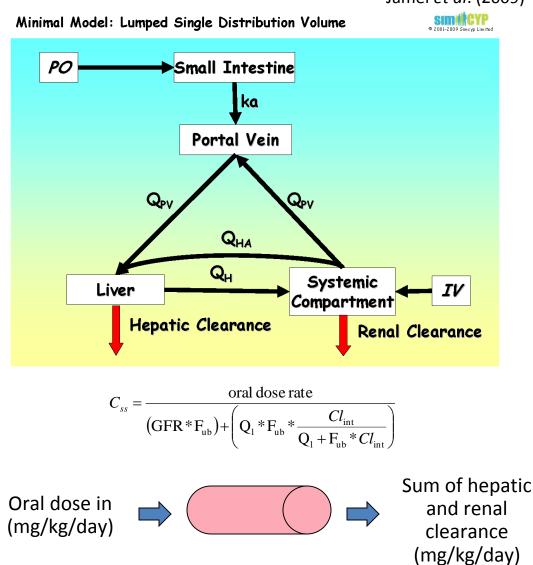


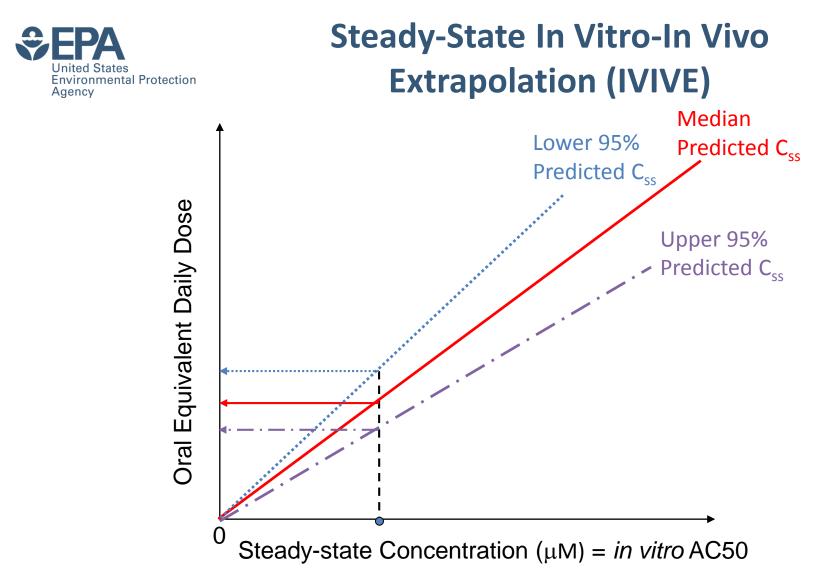
## High Throughput Toxicokinetics (HTTK)

 In vitro plasma protein binding and metabolic clearance assays allow approximate hepatic and renal clearances to be calculated

 At steady state this allows conversion from concentration to administered dose

 100% bioavailability assumed



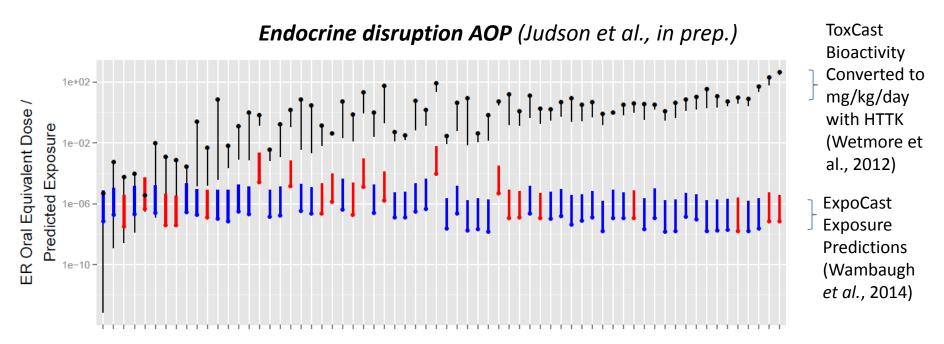


The higher the predicted  $C_{ss}$ , the lower the oral equivalent dose, so the upper 95% predicted  $C_{ss}$ from the MC has a lower oral equivalent dose

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### **Dosimetry and Exposure Provides Context for HTS**



#### **ToxCast Chemicals**

December, 2015 Panel:

"Scientific Issues Associated with Integrated Endocrine Bioactivity and Exposure-Based Prioritization and Screening"

DOCKET NUMBER: EPA-HQ-OPP-2014-0614



## Steady State Concentrations with httk R Package

<ul> <li>C f f https://cran.r-project.org/web/packages/httlk/index.html</li> <li>Q 2 x ≡</li> <li>httk: High-Throughput Toxicokinetics</li> <li>Functions and data tables for simulation and statistical analysis of chemical toxicokinetics ("TK") using data obtained from relatively high throughput, in vitro studies. Both physiologically-based ("PBTK") and empirical (e.g., one compartment) "TK" models can be parameterized for several hundred chemicals and multiple species. These models are solved efficiently, often using compiled (C-based) code. A Monte Carlo sampler is included for simulating biological variability and measurement limitations. Functions are also provided for exporting "PBTK" models to "SBML" and "JARNAC" for use with other simulation software. These functions and data provide a set of tools for in vitro-in vivo extrapolation ("IVIVE") of high throughput screening data (e.g., ToxCast) to real-world exposures via reverse dosimetry (also known as "RTK").</li> <li>Version: 1.2</li> <li>Imports: deSolve, msm.</li> <li>Suggests: ggplot2</li> <li>Published: 2015-05-11</li> <li>Author: John Wambaugh and Robert Pearce, Schmitt method implementation by Jimena Davis, dynamic model adapted from code by R. Woodrow Setzer, Rabbit parameters from Nisha Sipes</li> <li>Maintainer: John Wambaugh <wambaugh.john at="" epa.gov=""></wambaugh.john></li> <li>License: GPL-3</li> <li>NeedsCompilation: yes</li> <li>CP AM shoelva:</li> </ul>	RAN - Package httk ×		John 👝 🗆 🗙	
Functions and data tables for simulation and statistical analysis of chemical toxicokinetics ("TK") using data obtained from relatively high throughput, in vitro studies. Both physiologically-based ("PBTK") and empirical (e.g., one compartment) "TK" models can be parameterized for several hundred chemicals and multiple species. These models are solved efficiently, often using compiled (C-based) code. A Monte Carlo sampler is included for simulating biological variability and measurement limitations. Functions are also provided for exporting "PBTK" models to "SBML" and "JARNAC" for use with other simulation software. These functions and data provide a set of tools for in vitro-in vivo extrapolation ("IVIVE") of high throughput screening data (e.g., ToxCast) to real-world exposures via reverse dosimetry (also known as "RTK").         Version:       1.2         Imports:       deSolve, msm         Suggests:       ggplot2         Published:       2015-05-11         Author:       John Wambaugh and Robert Pearce, Schmitt method implementation by Jimena Davis, dynamic model adapted from code by R. Woodrow Setzer, Rabbit parameters from Nisha Sipes         Maintainer:       John Wambaugh <wambaugh.john at="" epa.gov="">         License:       GPL-3         NeedsCompilation:       yes</wambaugh.john>	← → C 🖍 🔒 https://	cran.r-project.org/web/packages/httk/index.html	@.☆ ≡	
from relatively high throughput, in vitro studies. Both physiologically-based ("PBTK") and empirical (e.g., one compartment) "TK" models can be parameterized for several hundred chemicals and multiple species. These models are solved efficiently, often using compiled (C-based) code. A Monte Carlo sampler is included for simulating biological variability and measurement limitations. Functions are also provided for exporting "PBTK" models to "SBML" and "JARNAC" for use with other simulation software. These functions and data provide a set of tools for in vitro-in vivo extrapolation ("IVIVE") of high throughput screening data (e.g., ToxCast) to real-world exposures via reverse dosimetry (also known as "RTK"). Version: 1.2 Imports: deSolve, msm Suggests: ggplot2 Published: 2015-05-11 Author: John Wambaugh and Robert Pearce, Schmitt method implementation by Jimena Davis, dynamic model adapted from code by R. Woodrow Setzer, Rabbit parameters from Nisha Sipes Maintainer: John Wambaugh <a href="https://wambaugh.john">wambaugh.john</a> at epa.gov> License: <u>GPL-3</u> NeedsCompilation: yes	httk: High-Thro	oughput Toxicokinetics	A	
Imports:deSolve, msmSuggests:ggplot2Published:2015-05-11Author:John Wambaugh and Robert Pearce, Schmitt method implementation by Jimena Davis, dynamic model adapted from code by R. Woodrow Setzer, Rabbit parameters from Nisha SipesMaintainer:John Wambaugh <wambaugh.john at="" epa.gov="">License:GPL-3NeedsCompilation:yes</wambaugh.john>	from relatively high throughput, in vitro studies. Both physiologically-based ("PBTK") and empirical (e.g., one compartment) "TK" models can be parameterized for several hundred chemicals and multiple species. These models are solved efficiently, often using compiled (C-based) code. A Monte Carlo sampler is included for simulating biological variability and measurement limitations. Functions are also provided for exporting "PBTK" models to "SBML" and "JARNAC" for use with other simulation software. These functions and data provide a set of tools for in vitro-in vivo extrapolation ("IVIVE") of high throughput screening data (e.g., ToxCast) to real-world exposures via reverse dosimetry			
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License: <u>GPL-3</u> NeedsCompilation: yes	Author:		vnamic	
NeedsCompilation: yes	Maintainer:	John Wambaugh <wambaugh.john at="" epa.gov=""></wambaugh.john>		
. ,	License:	<u>GPL-3</u>		
CDAN absolve: http://www.http://ww	NeedsCompilation: yes			
CRAIV checks. <u>Intra results</u>	CRAN checks:	httk results		
Downloads:	Downloads:			
Reference manual: <u>httk.pdf</u>	Reference manual:	<u>httk.pdf</u>		
Package source: <u>httk 1.2.tar.gz</u>	Package source:	httk_1.2.tar.gz		

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#### https://cran.r-project.org/web/packages/httk/

Can access this from the R GUI: "Packages" then "Install Packages"



## Steady State Concentrations with httk R Package

library(httk)

#Steady-state concentration (uM) for 1 mg/kg/day for 0.95 quantile for human for Acetochlor (published value): get\_wetmore\_css(chem.cas="34256-82-1")

```
# Should produce error:
get_wetmore_css(chem.name="34256-82-1")
```

```
#Capitalization shouldn't matter:
get_wetmore_css(chem.name="acetochlor")
get_wetmore_css(chem.name="Acetochlor")
```

```
# What's going on?
help(get_wetmore_css)
```

```
# What chemicals can I do?
get_wetmore_cheminfo()
```



## Version history for the "httk" R Package

The publicly available R package contains code and data that has been part of peer-reviewed publications

- Version 1.1 accompanied "Toxicokinetic Triage for Environmental Chemicals" Wambaugh et al. (2015) Tox. Sci.
- Version 1.2 accompanied "httk: R Package for High-Throughput Toxicokinetics" Pearce et al., submitted to Journal of Statistical Software
- Version 1.3 is in development and will be released to accompany "Incorporating High-Throughput Exposure Predictions with Dosimetry-Adjusted *In Vitro* Bioactivity to Inform Chemical Toxicity Testing" Wetmore et al., Tox. Sci. *in press*

While we develop research we maintain internal versions containing data and code that has yet to be peer reviewed.

#### **Released version**

```
sessionInfo() $otherPkgs$httk$Version
[1] "1.2"
all.data <-
get cheminfo(info=c("CAS", "Compound", "Clint", "Funbound.plasma"))
all.data[sapply(all.data$Compound,function(x) regexpr("conazole",x)!=-1),]
CAS
         Compound Human.Clint Human.Funbound.plasma
53 94361-06-5 Cyproconazole 1.537980
                                                   0.108556089
69 119446-68-3 Difenoconazole 24.210384
                                                   0.004255109
                               2.586969
75
   83657-24-3 Diniconazole
                                                   0.019467923
93 114369-43-6 Fenbuconazole 11.191873
                                                   0.005000000
119 79983-71-4 Hexaconazole 16.220043
                                                   0.040999740
214 112281-77-3 Tetraconazole
                              0.000000
                                                   0.024838039
234 131983-72-7 Triticonazole
                               6.117502
                                                   0.137256517
304 65277-42-1 Ketoconazole 55.000000
                                                   0.062000000
```

#### **Development version**

```
sessionInfo() $otherPkgs$httk$Version
[1] "1.2-4"
all.data <- get cheminfo(info=c("CAS","Compound","Clint","Funbound.plasma"))
all.data[sapply(all.data$Compound,function(x) reqexpr("conazole",x)!=-1),]
           CAS
                     Compound Human.Clint Human.Funbound.plasma
53
    94361-06-5 Cyproconazole 1.537980
                                                   0.108556089
69 119446-68-3 Difenoconazole 24.210384
                                                   0.004255109
75 83657-24-3
                 Diniconazole 2.586969
                                                   0.019467923
93 114369-43-6 Fenbuconazole 11.191873
                                                   0.005000000
119 79983-71-4 Hexaconazole 16.220043
                                                   0.040999740
215 112281-77-3 Tetraconazole
                                0.000000
                                                   0.024838039
235 131983-72-7 Triticonazole
                                6.117502
                                                   0.137256517
242 60207-90-1 Propiconazole 18.400000
                                                   0.029491801
356 65277-42-1 Ketoconazole 55.000000
                                                   0.062000000
```



## Steady State Concentrations with httk R Package

#Steady-state concentration (uM) for 1 mg/kg/day for 0.95 quantile for human for Acetochlor (calculated value): calc\_mc\_css(chem.cas="34256-82-1")

#Steady-state concentration (uM) for 1 mg/kg/day for 0.95 quantile for rat for Acetochlor (should produce errors since there is no published value, 0.5 quantile only):

get\_wetmore\_css(chem.cas="34256-82-1",species="Rat")

#Steady-state concentration (uM) for 1 mg/kg/day for 0.95 quantile for rat for Acetochlor (calculated value): calc\_mc\_css(chem.cas="34256-82-1",species="Rat")

#Steady-state concentration (uM) for 1 mg/kg/day for 0.5 quantile for rat for Acetochlor (published value): get\_wetmore\_css(chem.cas="34256-82-1",species="Rat",which.quantile=0.5)

#Steady-state concentration (uM) for 1 mg/kg/day for 0.5 quantile for rat for Acetochlor (calculated value): calc\_mc\_css(chem.cas="34256-82-1",species="Rat",which.quantile=0.5)

#Steady-state concentration (uM) for 1 mg/kg/day for 0.95 quantile for mouse for Acetochlor (should produce error since there is no published value, human and rat only):

get\_wetmore\_css(chem.cas="34256-82-1",species="Mouse")

#Steady-state concentration (uM) for 1 mg/kg/day for 0.95 quantile for mouse for Acetochlor (calculated value): calc\_mc\_css(chem.cas="34256-82-1",species ="Mouse")



## Oral Equivalent Doses with with httk R Package

#State-state oral equivalent dose (mg/kg BW/day) to produce 0.1 uM serum concentration for human, 0.95 quantile, for Acetochlor (published value):

```
get_wetmore_oral_equiv(0.1,chem.cas="34256-82-1")
```

#State-state oral equivalent dose (mg/kg BW/day) to produce 0.1 uM serum concentration for human, 0.95 quantile, for Acetochlor (calculated value):

```
calc_mc_oral_equiv(0.1,chem.cas="34256-82-1")
```

#State-state oral equivalent dose (mg/kg BW/day) to produce 0.1 uM serum concentration for human, 0.05, 0.5, and 0.95 quantile, for Acetochlor (published values):

get\_wetmore\_oral\_equiv(0.1,chem.cas="34256-82-1",which.quantile=c(0.05,0.5,0.95))

#State-state oral equivalent dose (mg/kg BW/day) to produce 0.1 uM serum concentration for human, 0.05, 0.5, and 0.95 quantiles, for Acetochlor (calculated value):

```
calc_mc_oral_equiv(0.1,chem.cas="34256-82-1",which.quantile=c(0.05,0.5,0.95))
```

#State-state oral equivalent dose (mg/kg BW/day) to produce 0.1 uM serum concentration for rat, 0.95 quantile, for Acetochlor (calculated value):

calc\_mc\_oral\_equiv(0.1,chem.cas="34256-82-1",species="Rat")





- Toxicokinetics (TK) provides a bridge between HTS and HTE by predicting tissue concentrations due to exposure
- HTTK methods developed for pharmaceuticals have been adapted to environmental testing
- A primary application of HTTK is "Reverse Dosimetry" or RTK
- New R package "httk" freely available on CRAN allows statistical analyses
  - Analysis has been submitted



### Chemical Safety for Sustainability (CSS) Rapid Exposure and Dosimetry (RED) Project

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High Throughput Toxicokinetics Researchers NRMRL Yirui Liang<sup>\*</sup> Xiaoyu Liu

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Craig Barber

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