

Advancing Dose-Response Analyses and Tools

**Board of Scientific Counselors (BOSC) Subcommittee
Chemical Safety for Sustainability (CSS) and
Health and Environmental Risk Assessment (HERA) National Research Programs
Virtual Meeting on February 2-5, 2021**

**John Vandenberg (Vandenberg.john@epa.gov)
*Director, Health and Environmental Effects Assessment Division (HEEAD)***

February 5, 2021



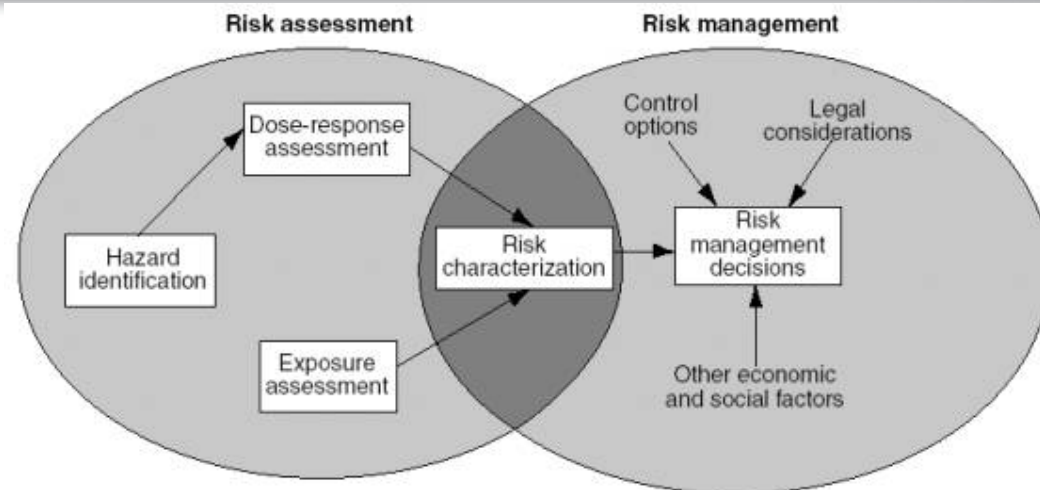
Advancing Dose-Response Analysis and Tools

Characterizing dose-response relationships is fundamental to health risk assessment

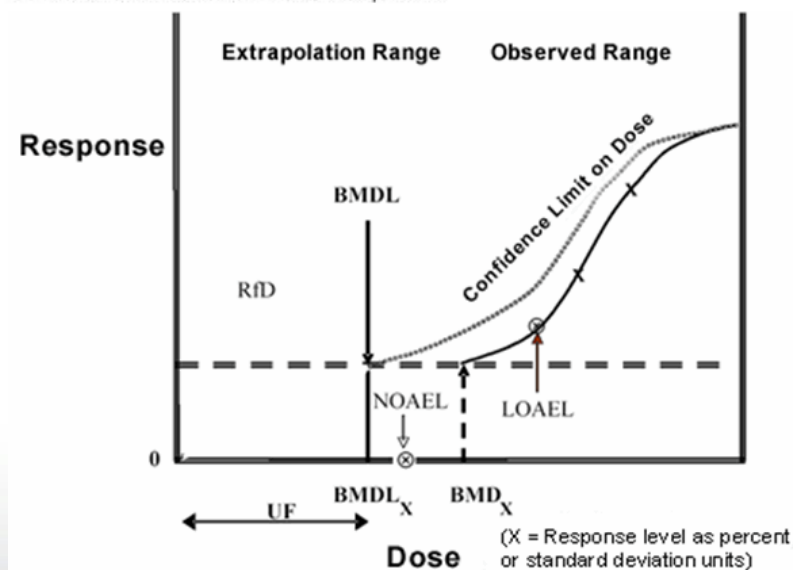
Models are used to extrapolate from the observed range, often well above environmental levels, to the range representing human exposures

Characterizing dose is of key importance, but very challenging for inhalation exposures

Characterizing uncertainty is essential for effective interpretation of model results



Source: EPA Office of Research and Development.





Coordinated Approach to Research, Tools, and Training

- **Research – Focused in Output 3.5**

- 3.5.1 - Research in support of informative parameter priors used in Bayesian model averaging of dichotomous and continuous endpoint (Allen Davis's presentation)
- 3.5.2 - Development of multivariate benchmark dose modeling for traditional toxicological and toxicogenomic data
- 3.5.3 - Development of a unified model suite for dichotomous and continuous toxicological data
- 3.5.4 - Advancement of nested dose-response modeling for developmental toxicity data
- 3.5.5 - Characterizing Determinants of Risk: Concentration, Duration and Timing of Exposure
- 3.5.6 - Case Studies and Advancements in Uncertainty Analysis (Todd Blessinger's presentation)
- And more in Appendix B, Part 2

- **Tools – Focused in Output 4.1**

- 4.1.3 Development, Operation, and Interoperability of Existing and Implementation of Planned Critical Components of BMDS and CatReg (Allen Davis's presentation)
- 4.1.5 - All Ages Lead Model
- 4.1.6 - Evaluation of the Integrated Exposure Uptake Biokinetic (IEUBK) Model version 2.0
- 4.1.7 - Multi-path Particle Dosimetry (MPPD) model (Annie Jarabek's presentation)

- **Training – Focused in Output 4.2**

- 4.2.1 - Risk Assessment Training to Improve the Harmonization and Collaboration between ORD and EPA Regional/Program Offices, State/Local/Tribal Agencies, and International Organizations (Annie Jarabek's presentation)
- 4.2.3 - Development and Maintenance of BMDS and CatReg Documentation and Training Manual



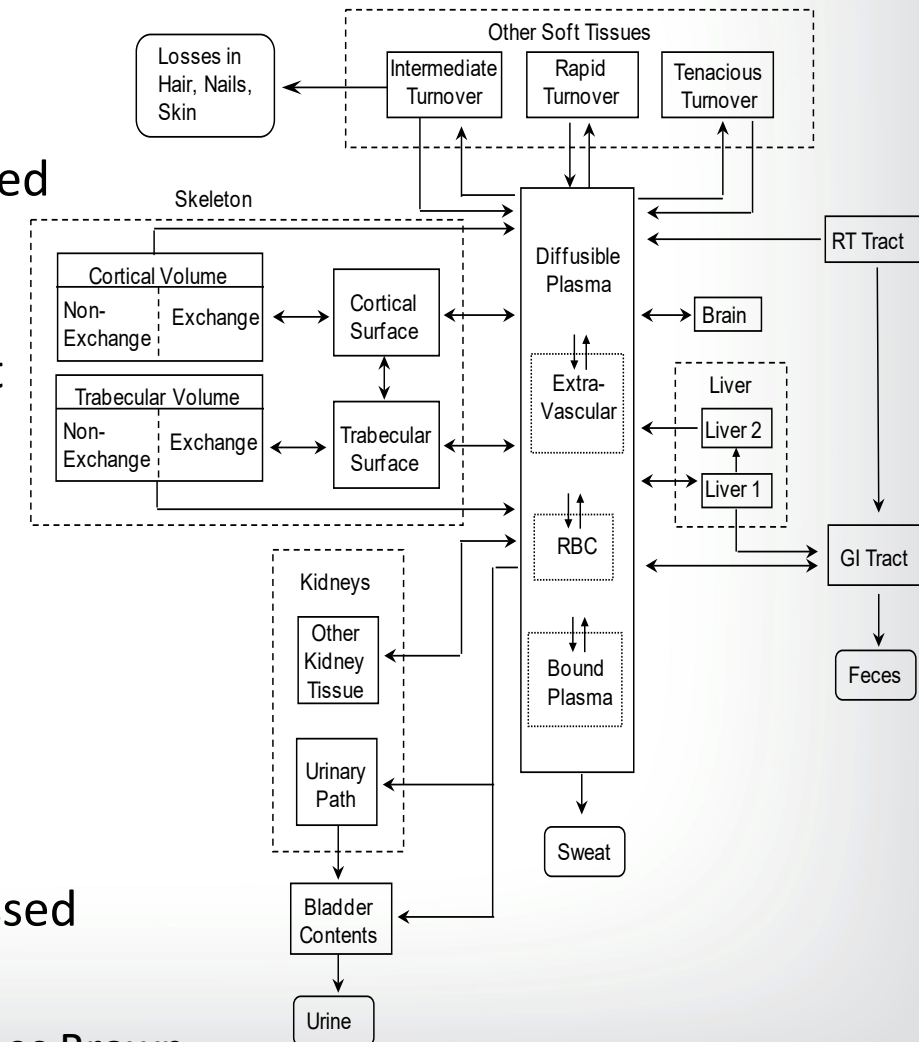
EPA Pb exposure and biokinetic models led by HERA scientists

Integrated Exposure Uptake Biokinetic (IEUBK) model

- Estimates Pb in blood of children up to 7 years of age
- Steady state exposure that can vary by year of life
- Recommended risk assessment tool to support residential lead-related site cleanups
- Used to support OPPT Lead Dust Hazard Standard, OLEM soil Pb guidance, OW Lead and Copper Rule, and Superfund risk assessment tool to support residential lead site cleanups

All Ages Lead Model (AALM)

- Estimates Pb in blood and other tissues (e.g., bone)
- Extends modeling capabilities for people up to 90 years of age
- Allows acute, transiently reoccurring, and/or chronic exposures
- Underwent EPA SAB review October 2019
- Public release planned after SAB comments (finalized 8/3/20) addressed
- Anticipated wide use after public release



modified from slides courtesy of James Brown

Advancing Dose-Response Analyses and Tools

12:20 – 12:35	Advancing Dose-Response Intro with Charge Question	John Vandenberg, CPHEA
12:35 – 12:55	Multi-path Particle Dosimetry Model	Annie Jarabek, CPHEA
12:55 – 1:15	Bayesian Model Averaging and BMDS 3.2	Allen Davis, CPHEA
1:15 – 1:35	Approximate Probabilistic Analysis (APROBA)	Todd Blessinger, CPHEA
1:35 – 2:10	BOSC Subcommittee discussion and Qs/As	Katrina Waters, Chair



Charge Question

Dose-response modeling is a critical step in human health assessment. Existing methods have improved upon older methodologies; however, unresolved issues, uncertainties, and complications remain that require targeted research. HERA has proposed research products that will result in dose-response methods that are more precise, robust, and meet varied needs. Is it clear how these planned products address important issues in dose-response modeling with an application to risk assessment? What suggestion(s) or recommendation(s) does the Subcommittee offer to continue to advance methods in dose-response modeling with an application to risk assessment? [Research Area 3, Output 3.5 and Research Area 4, Output 4.1]



EPA Multi-path Particle Dosimetry (MPPD) Model 2021 (v. 1.01) Technical Support Documentation & User's Guide

Annie M. Jarabek

Human and Environmental Effects Assessment Division

CPHEA

Advancing Dose-Response Analyses and Tools

HERA Session 3

US EPA Board of Scientific Counselors Chemical Safety Subcommittee
Meeting

February 5, 2021



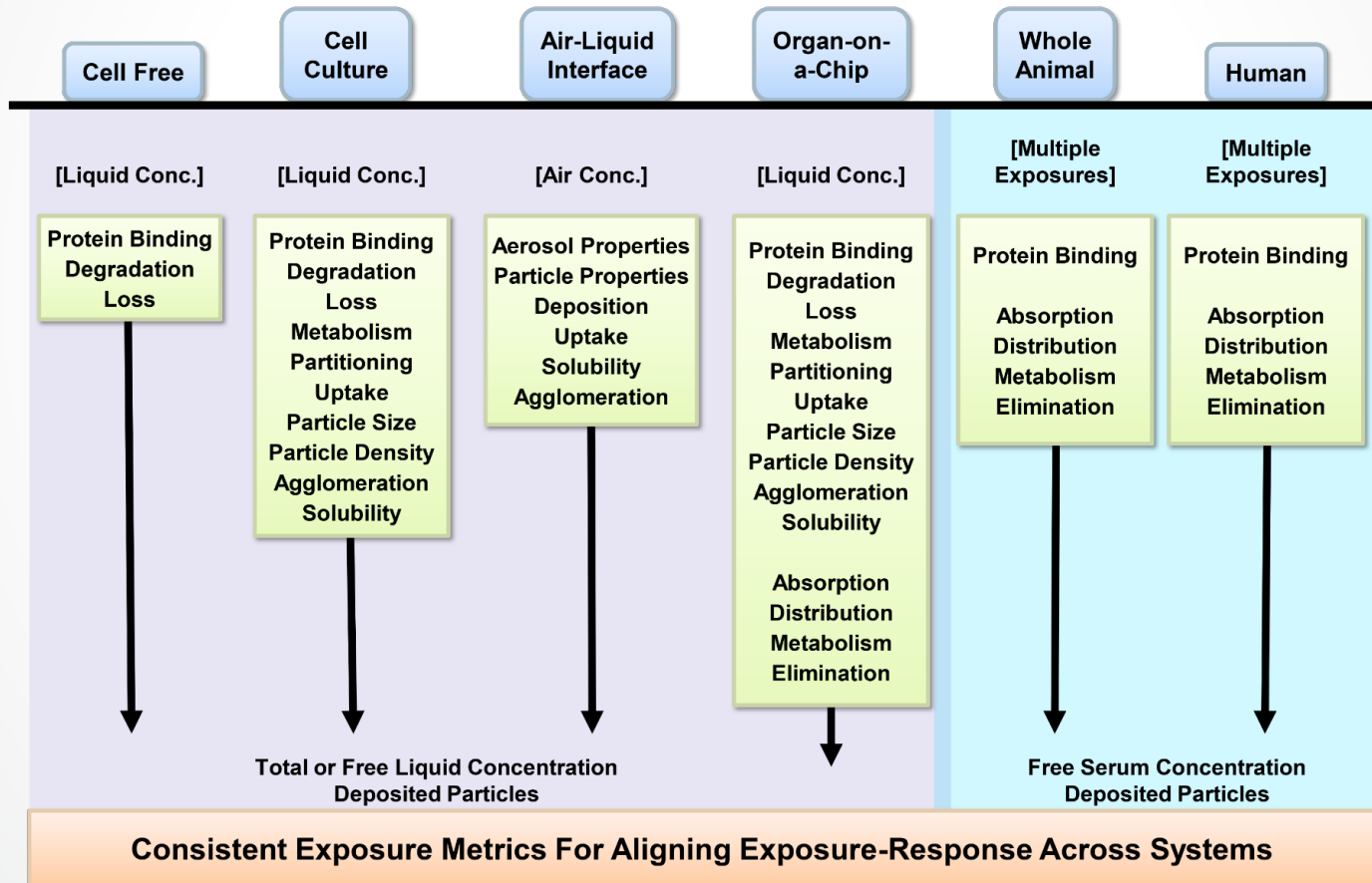
- Modernizing our assessment toolbox: Advancing evidence integration
- Why: Choice of MPPD model
- What: New EPA-specific software with technical support & user's guide
- Who: Collaborators in development and application
- How: Role in risk assessment – use case applications in OCSPP
- When: Next steps
- Roll out: Risk assessment training
- Summary

Disclaimer: These views are those of the author and do not represent US EPA policy.



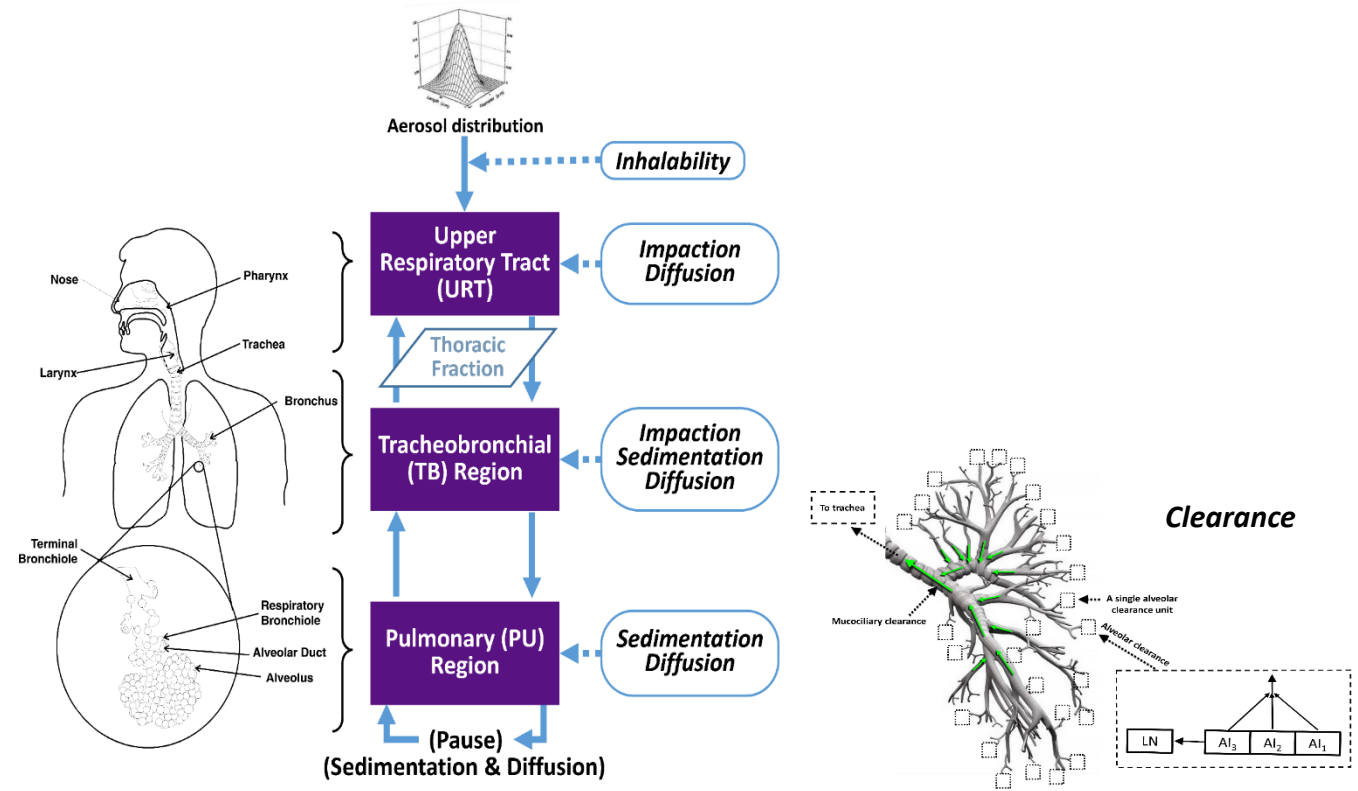
Modernizing our Assessment Toolbox

➤ *Advancing evidence integration: Dosimetry adjustments aid exposure alignment*

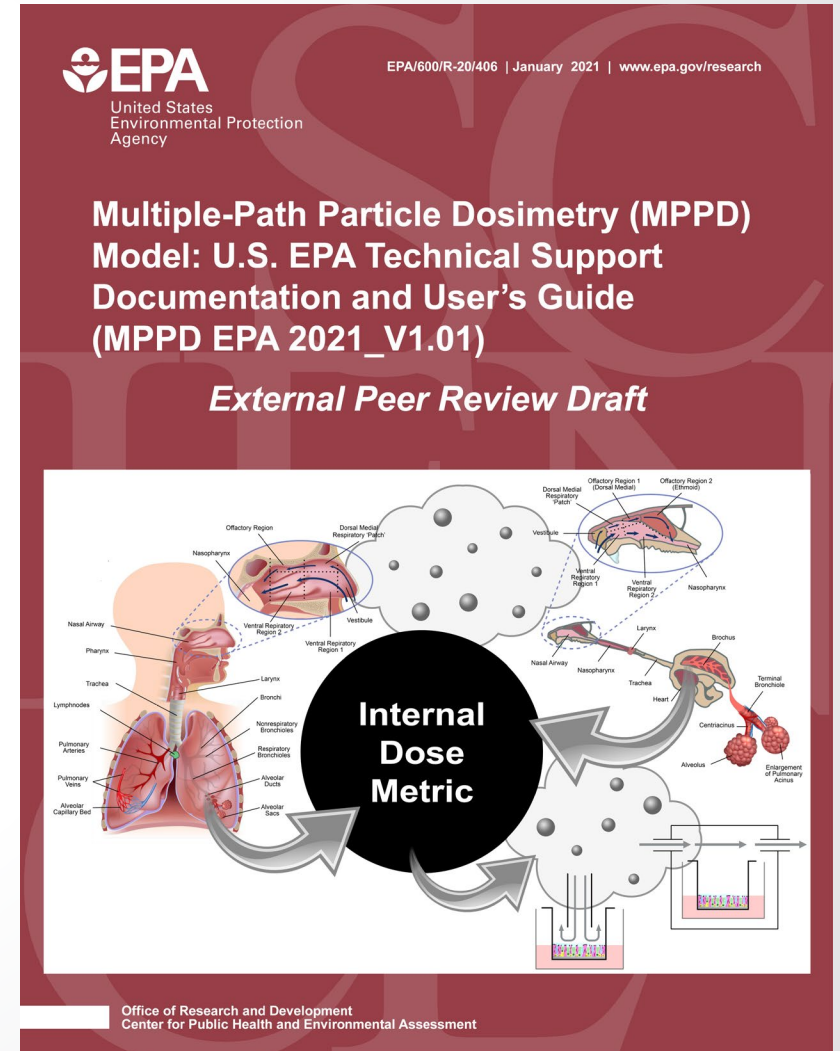


Why: Choice of MPPD Model

- Particle model in current use was developed in 1994 as part of the inhalation reference concentration (RfC) methodology
- Particle dosimetry modeling has matured
 - Additional algorithms: Inhalability
 - Move from empirical to mechanistic description of deposition
 - Clearance to predict retained dose
- Applied Research Associates, Inc. has developed many versions for various clients
 - EPA users confused regarding differences
 - External partners (states, NGO, academics) also need guidance
- Support consistent use across the Agency programs

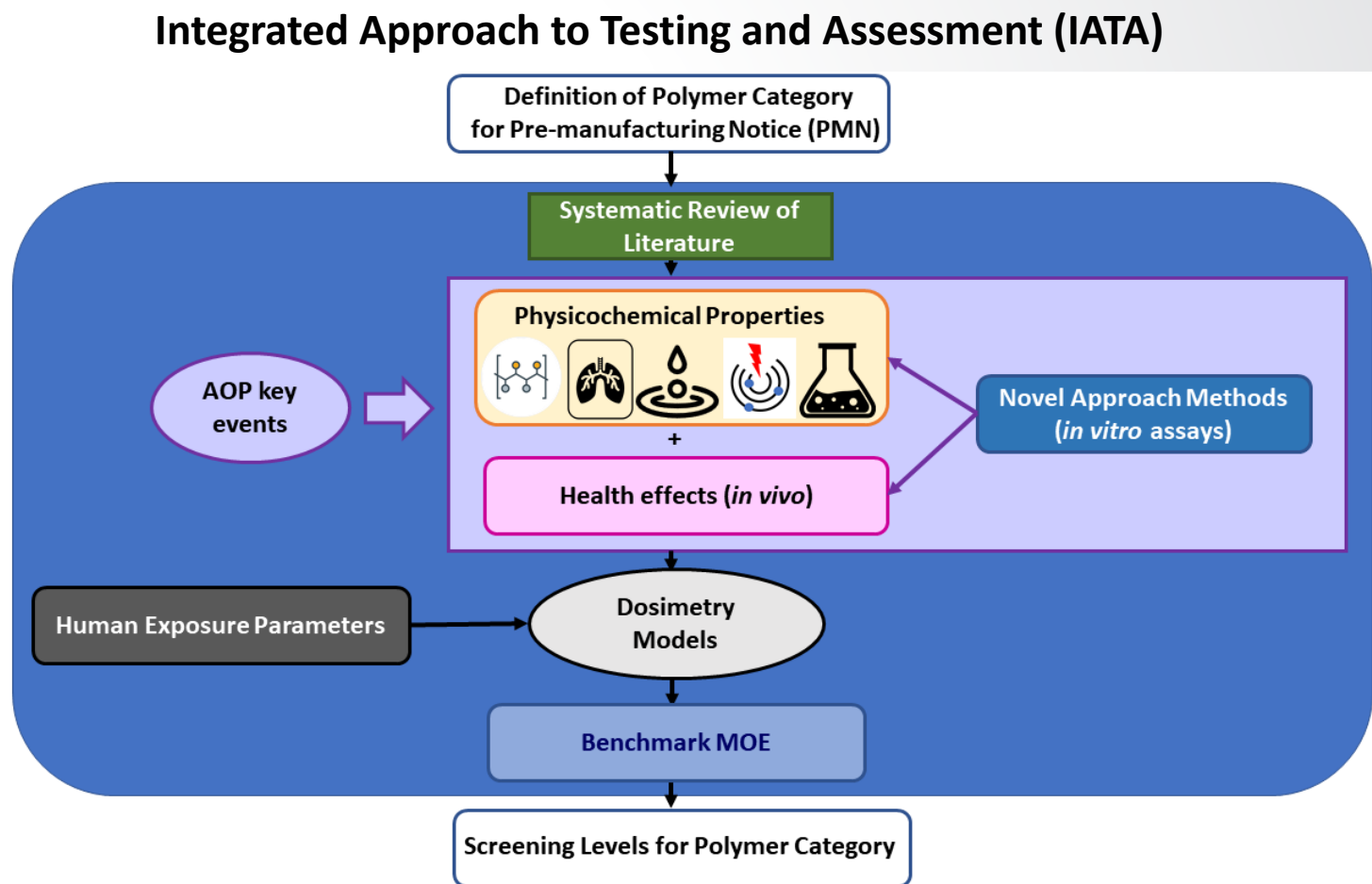


- New EPA version of the MPPD model software
 - Revised graphical user's interface (GUI)
 - Some updated algorithms
- Technical support documentation and user's guide
 - Introduction to inhalation dosimetry
 - Step-by-step explanation of input fields
 - Guidance on input parameters and procedures
 - Specific use case illustrations
- Agency deployment requires external peer review



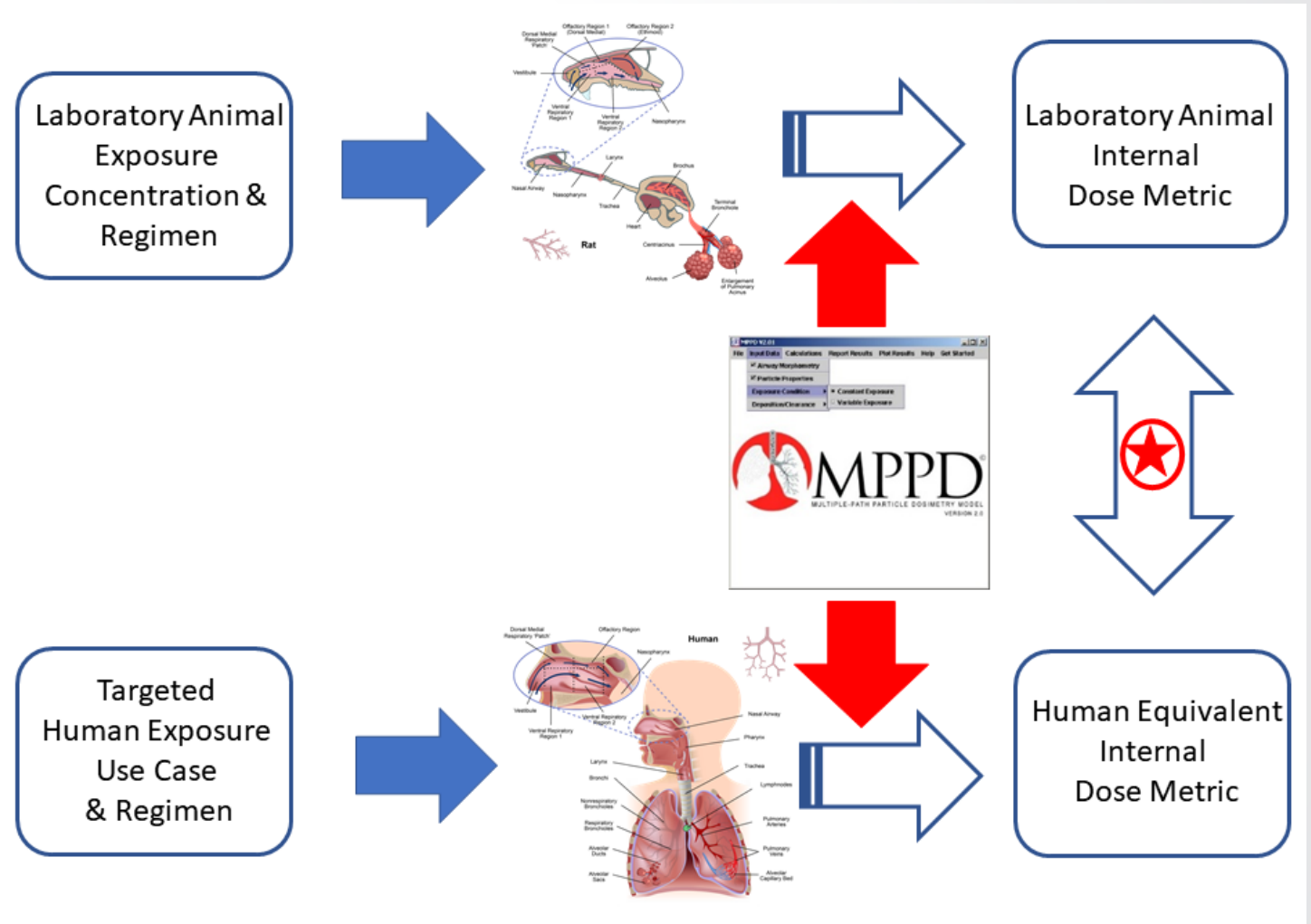
Who: Development and Application

- Collaborative development
 - Annie M. Jarabek, US EPA ORD, CPHEA HEEAD
 - Bahman Asgharian, Applied Research Associates, Inc., Raleigh, NC
 - Fred Miller, Fred Miller, LLC
 - Owen Price, Applied Research Associates, Inc., Arlington, VA
- Application: Pre-manufacturing notice (PMN) program in OPPT
- Additional use case applications: TSCA



How: Role in Risk Assessment

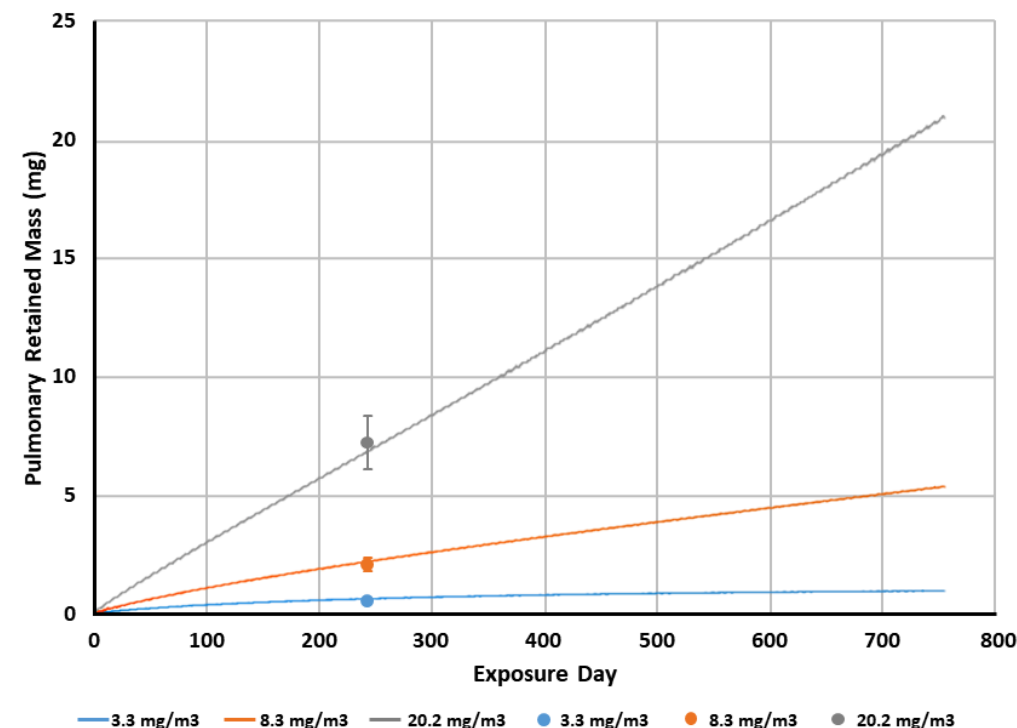
- Interspecies extrapolation: human equivalent concentration (HEC)
- Improved characterization of target human scenario
 - Specific size, distribution, density and exposure data
 - Replace default parameters and equations





Use Case: Particle Overload

- **Particle overload:** When particles **overwhelm ability of alveolar macrophages** to clear from pulmonary region
- A **kinetic phenomenon that creates context** for consideration of observed toxicity, e.g., rat lung tumors not considered relevant due to differences in clearance rates
- **Novel deployment of MPPD** to demonstrate overload occurrence to define **new pre-manufacturing notice (PMN) categories for read-across** and inform risk evaluations (manuscripts accepted in *ACS' Chem Res Toxicol* and abstracts submitted for presentations at SOT 2021)
 - **Poorly Soluble, Low Toxicity (PSLT) Polymer Category: An Integrated Approach to Testing and Assessment (IATA) Including New Approach Methods (NAMs) under the Toxic Substances Control Act (TSCA).** A. M. Jarabek¹, T. Stedeford², G. S. Ladics³, O. T. Price⁴, A. Tveit⁵, M. P. Hayes⁶, R. T. Tremblay⁷, S. A. Snyder⁸, K. D. Salazar², S. Osman-Sypher⁹, W. Irwin², M. Odin¹⁰, J. Melia¹⁰, H. Carlson-Lynch¹⁰, M. Sharma¹¹, A. J. Clippinger¹¹, A. O. Stucki¹¹, and T. R. Henry². ¹US EPA, Research Triangle Park, NC; ²US EPA, Washington, DC; ³Dupont Nutrition and Biosciences, Wilmington, DE; ⁴Applied Research Associates Inc., Arlington, VA; ⁵BASF Corporation, Florham Park, NJ; ⁶Procter & Gamble, Mason, OH; ⁷Procter & Gamble, Strombeek-Beaver, Belgium; ⁸Covestro LLC, Pittsburgh, PA; ⁹American Chemistry Council, Washington, DC; ¹⁰SRC Inc., North Syracuse, NY; and ¹¹PETA International Science Consortium Ltd., London, United Kingdom.
 - **Surfactants Category: An Integrated Approach to Testing and Assessment (IATA) Including New Approach Methods (NAMs) for Assessing Inhalation Risks under the Toxic Substances Control Act (TSCA).** T. R. Henry¹, K. D. Salazar¹, M. P. Hayes², W. Kennedy³, A. M. Keene³, A. M. Jarabek⁴, O. T. Price⁵, S. Moors⁶, L. Jovanovich⁷, J. L. Rose⁸, A. Tveit⁹, R. T. Tremblay¹⁰, R. A. Becker¹¹, S. Osman-Sypher¹¹, P. D. McMullen¹², S. D. Slattery¹², W. Irwin¹, M. Odin¹³, J. Melia¹³, M. Sharma¹⁴, A. J. Clippinger¹⁴, A. Stucki¹⁴, and T. Stedeford¹. ¹US EPA, Washington, DC; ²Procter & Gamble, St. Bernard, OH; ³Afton Chemical Corporation, Richmond, VA; ⁴US EPA, Research Triangle Park, NC; ⁵Applied Research Associates, Inc., Arlington, VA; ⁶BASF Corporation, Duesseldorf, Germany; ⁷Stepan Company, Northfield, IL; ⁸Procter & Gamble, Mason, OH; ⁹BASF Corporation, Florham Park, NJ; ¹⁰Procter & Gamble, Strombeek-Beaver, Belgium; ¹¹American Chemistry Council, Washington, DC; ¹²ScitoVation, Durham, NC; ¹³SRC Inc., North Syracuse, NY; and ¹⁴PETA International Science Consortium Ltd., London, United Kingdom.



Jarabek, Stedeford et al. (*accepted*)

Rollout: Risk Assessment Training

- Didactic and experiential modules with experts to convey scientific subject matter and methodologies
- Collaboration with OCSPP to update for TSCA applications
- MPPD training will be one module within a set covering inhalation issues:
 - Inhalation toxicology
 - Inhalation dosimetry modeling (both particles and different gases)
 - MPPD model for particle dosimetry

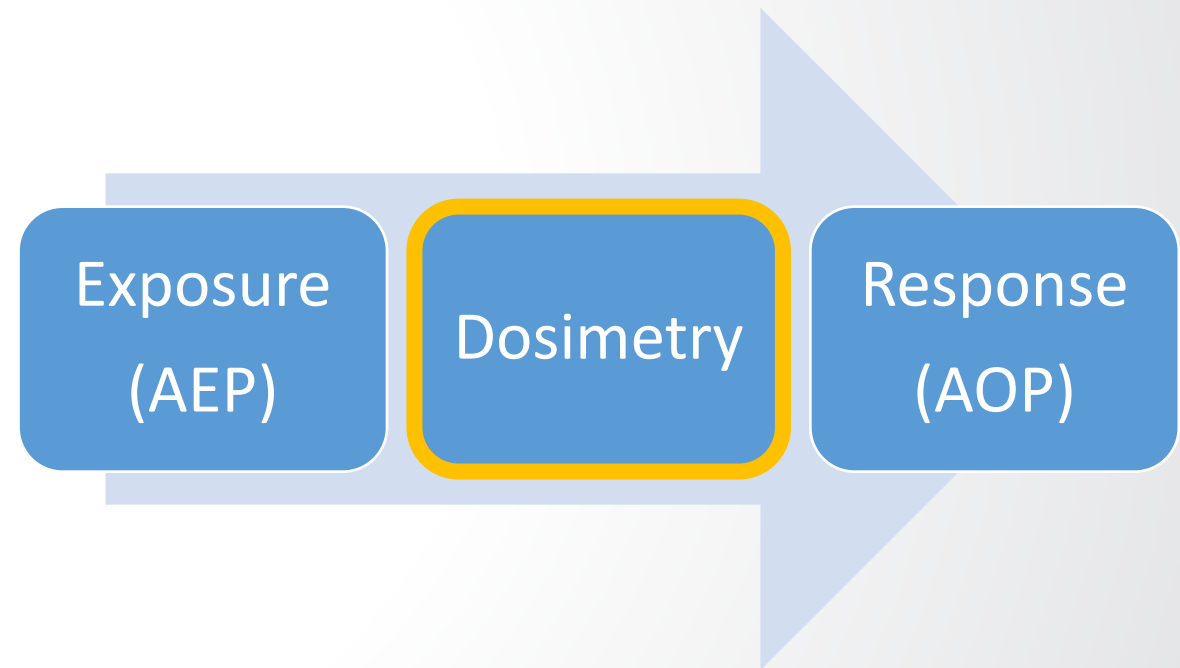


When: Next Steps

- Release of external peer review document: March 2021
- Virtual public comment and peer review meeting: April 2021
- Revision based on peer panel recommendations: Fall 2021
- Development and deployment of training: Spring 2021
- On-going maintenance and support as part of HERA program
- Build workbench to integrate dosimetry into assessment workflows



- Dosimetry modeling is **critical link** to translate exposure to internal dose for response analysis
 - Exposure alignment for evidence integration
 - Application of NAMs
- MPPD model in toolbox will build **capacity** to bring dosimetry directly into assessment workflows
- **Collaborative development** by use cases and training ensures consistent and coherent application in assessments across programs

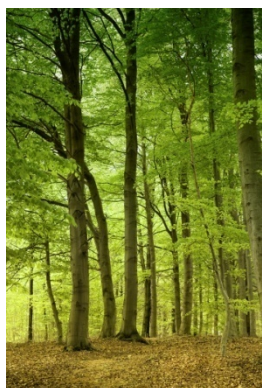




Benchmark Dose Modeling – Bayesian Model Averaging

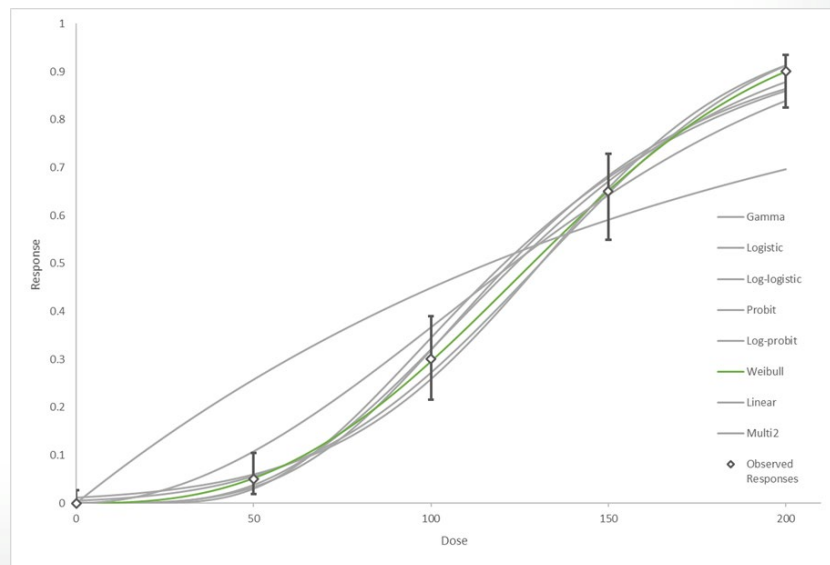
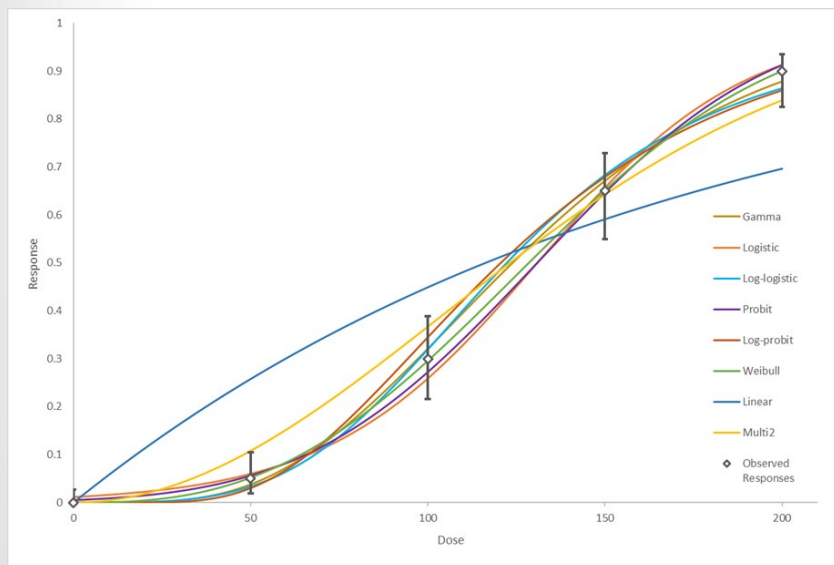
Allen Davis, MSPH

Center for Public Health and Environmental Assessment, U.S. EPA



The views expressed in this presentation are those of the author(s) and do not necessarily reflect the views or policies of the US EPA.

- **When fitting multiple models to a single dataset, many models can (*and often will*) statistically fit the data well**
 - So, is there a compelling reason (toxicology, MOA, etc.) to pick one model over any other?
 - Or (most commonly) is the model selected based on pure statistical fit?
 - This is *model uncertainty*



- **Multiple approaches have been developed for addressing and/or characterizing model uncertainty**
 - Flexible parametric models – some research has indicated that some models (Exponential 5) are flexible enough to fit the majority of dose-response shapes observed in the literature
 - Semi- or non-parametric models – completely data-driven models that are hyper-flexible
 - **Model averaging** – methods by which the results of a suite of individual models are averaged together to give one estimate of the BMD and BMDL; consistent with modeling approaches recommended by WHO and implemented by European partners (EFSA, RIVM)



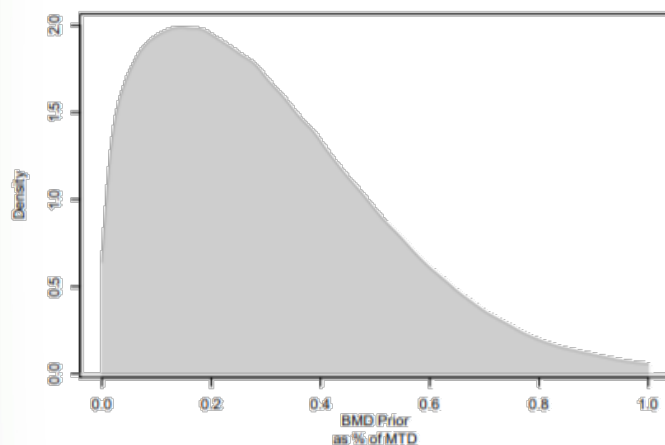
BMDS Bayesian Model Averaging

- **Unique aspects of EPA/NIOSH model averaging approach:**
 - Informed priors
 - Based on where the BMD_{10} estimates are most likely to be relative to a study's maximum dose
 - Disentangle issues related to models that “degenerate” to other models (Weibull, etc.)
 - Prevent over-fitting of individual models
 - Provides a single standard set of priors in BMDS (i.e., Excel version) that gives reasonable, health-protective, consistent, and reproducible results
 - Research on priors is part of current HERA research portfolio (HERA 3.5.1)
 - Laplace approximation of posterior density
 - Minimal loss of accuracy or reliability
 - Substantial increase in speed (~10-fold faster than MCMC approaches implemented in other platforms)
 - Increases in speed are critically important for batch analyses of many datasets

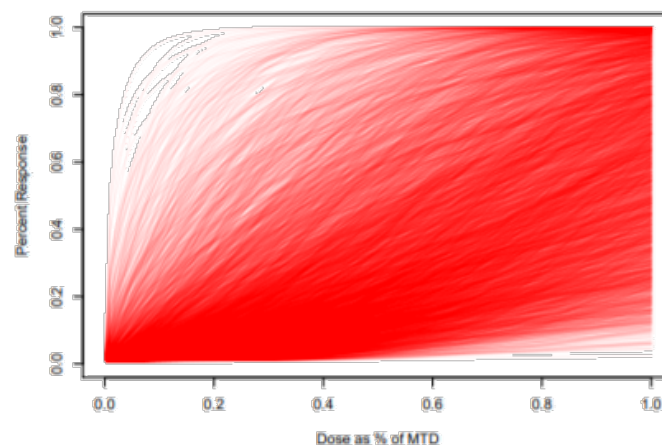
- **Instead of looking at priors over all model parameters, or specific parameters, place a reasonable prior over the value of ultimate interest, the BMD**
- **Benefits:**
 - All models are wrong, so the parameters are abstract entities. We are ultimately interested in the value of the BMD
 - In terms of maximum tolerated dose and dose-response study design, the value of the BMD can be expressed as a percentage of the maximum tolerated dose.
- **Issues:**
 - Can be perceived as subjective in terms of what is “right”
 - Significant prior impact in low data cases
 - Might change based upon target quantity (i.e., may be different for BMR = 10% vs. BMR = 1%)

- Here, assuming a prior on the **BMD** such that the majority is between 0 and 0.5 of the **MTD**

Prior on the BMD



Prior on the DR curves.



- In **BMDs 3.2** (and later versions) priors for dichotomous models are set such that the **BMD** is expected to fall within $\sim 0.2-0.5$ of the **MTD**



BMDS Bayesian Model Averaging

- **Maximum a-posteriori – find the maximum of the posterior distribution and use a normal like approximation**
 - Very fast computationally compared to Markov Chain Monte Carlo (MCMC)
 - Accurate for the right-sized posterior
- **BMDS uses the Laplace approximation to the marginal density of the data as model weights**
- **The model-averaged BMD point estimate is the weighted average of the MAP estimates from individual models**
- **BMDL and BMDU values are estimated similar to the profile likelihood approach except that the posterior density is profiled**
- **Method is fully described in Risk Analysis paper: Quantitative risk assessment: developing a Bayesian approach to dichotomous dose-response uncertainty. Risk Analysis 40(9). DOI: 10.1111/risa.13537**



BMDS Bayesian Model Averaging

- **EPA/NIOSH BMA approach was extensively tested against 1) MCMC Bayesian MA approach with uninformative priors; 2) BMDS using 2012 model selection criteria; and 3) flexible non-parametric model**
- **34 separate “true-dose” curves used to test approaches**

Percentage of Times BMDL Coverage is >90% than True BMD Value




True BMD	BMA	BMDS	NP	MCMC
All templates	70.6%	41.2%	76.5%	47.1%
True BMD < 0.2x max dose	63.2%	26.3%	57.9%	36.8%
True BMD > 0.2x max dose	80%	60%	100%	60%
True BMD < 0.1x max dose	60%	30%	20%	40%
True BMD > 0.1x max dose	75%	45.8%	100%	50%



BMDS User Interface

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   **BMDS 3.1.2**

Define Analysis

Analysis Name

Select Output Directory

Analysis Description

Select Model Type

	MLE		Alternatives		
	Frequentist Restricted	Frequentist Unrestricted	Bayesian	Bayesian Model Average	
Model Name	Enable <input type="checkbox"/>	Enable <input type="checkbox"/>	Enable <input type="checkbox"/>	Enable <input type="checkbox"/>	Prior Weights
Dichotomous Hill	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Gamma	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Logistic	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Log-Logistic	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Log-Probit	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Multistage	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Probit	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Quantal Linear	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Weibull	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Total Weight					0.000%

Option Set #	Risk Type	BMR	Confidence Level	Background
1	Extra Risk	0.1	0.95	Estimated

☒ DataSets

Main Data Report Options Logic ModelParms




Ready



Using BMDS – Enter Dose-Response Data

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   **BMDS 3.1.2**

Data

Insert New Dataset <-- Click here to insert a new dataset manually

Import Dataset <-- Click here to import an existing BMDS dataset (*.dax file)

Edit BMDS - training - Dichotomous

[Add user notes here]

Dose	N	Incidence
0	25	0
5	25	1
20	25	3
80	25	9
200	25	16

← **Name dataset**

Manually enter dose-response data

Edit BMDS - training - Continuous

[Add user notes here]

Dose	N	Mean	Std. Dev.
0	25	10.56	1.56
5	25	10.26	1.26
20	25	8.98	1.35
80	25	7.56	1.21
200	25	6.99	1.33

Ready Main **Data** Report Options Logic ModelParms



Using BMDS – Select Models to Run

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BMDS 3.1.2

Define Analysis

Analysis Name

Select Output Directory

Analysis Description

Select Model Type

Model Name	MLE		Alternatives		Prior Weights
	Frequentist Restricted	Frequentist Unrestricted	Bayesian	Bayesian Model Average	
Dichotomous Hill	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Gamma	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Logistic	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Log-Logistic	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Log-Probit	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Multistage	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Probit	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Quantal Linear	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Weibull	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	<input type="checkbox"/>	0.0000%
Total Weight					0.0000%

DataSets

Enable	DataSet
<input checked="" type="checkbox"/>	BMDS - training - Dichotomous
<input type="checkbox"/>	BMDS - training - Continuous

Option Set # **Risk Type** **BMR** **Confidence Level** **Background**

1	Extra Risk	0.1	0.95	Estimated
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Ready | Main | Data | Report Options | Logic | ModelParms | 100%






Using BMDS – Select Specific Models

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Share Comments

   **BMDS 3.1.2**

Define Analysis

Analysis Name BMDS Training Analysis

Select Output Directory C:\Users\adavis10\BMDS312

Analysis Description Examples of how to use BMDS 3.2 for dichotomous, continuous, dichotomous cancer, and nested developmental toxicity data

Select Model Type Dichotomous

Load Analysis **Save Analysis** **Run Analysis**

Model Name	MLE		Alternatives		Prior Weights
	Frequentist Restricted	Frequentist Unrestricted	Bayesian	Bayesian Model Average	
Dichotomous Hill	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Gamma	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Logistic	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Log-Logistic	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Log-Probit	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Multistage	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Probit	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Quantal Linear	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Weibull	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Total Weight					100.000%

Option Set #	Risk Type	BMR	Confidence Level	Background
1	Extra Risk	0.1	0.95	Estimated

Add Option Set

DataSets

Enable	
<input checked="" type="checkbox"/>	BMDS - training - Dichotomous
<input type="checkbox"/>	BMDS - training - Continuous

Ready



Using BMDS – Add Option Sets

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BMDS 3.1.2

Define Analysis

Analysis Name

Select Output Directory

Analysis Description

Select Model Type

Model Name	MLE		Alternatives		Prior Weights
	Frequentist Restricted	Frequentist Unrestricted	Bayesian	Bayesian Model Average	
Dichotomous Hill	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Gamma	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Logistic	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Log-Logistic	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Log-Probit	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Multistage	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Probit	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Quantal Linear	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Weibull	<input checked="" type="checkbox"/>	<input type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	11.1111%
Total Weight					100.000%

Option Set #	Risk Type	BMR	Confidence Level	Background
1	Extra Risk	0.1	0.05	Estimated

Enable	DataSets
<input checked="" type="checkbox"/>	BMDS - training - Dichotomous
<input checked="" type="checkbox"/>	BMDS - training - Continuous

Ready | Main | Data | Report Options | Logic | ModelParms | 100%



Using BMDS – Modeling Summary Results

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Summary

BMDS 3.1.2

[Add user notes here]		
Dose	N	Incidence
0	25	0
5	25	1
20	25	3
80	25	9
200	25	16

Examples of how to use BMDS 3.2 for dichotomous, continuous, dichotomous cancer, and nested developmental toxicity data

Recommended frequentist model
Model averaging

Option set #1 (Hover for details) | Scroll right to see summary plot ->

Model	Analysis Type	Restriction	RiskType	BMRF	BMD	BMDL	BMDU	P Value	AIC	Unnormalized Log Posterior Probability	Scaled Residual for Dose Group near BMD	Scaled Residual for Control Dose Group	BMDS Re
Dichotomous Hill	frequentist	Restricted	Extra Risk	0.1	15.85789	7.897094	29.144824	0.6945918	100.2357564	-	-0.067756196	-0.00062655	Viable
Gamma	frequentist	Restricted	Extra Risk	0.1	19.33104	14.31304	30.451431	0.9892812	94.36832933	-	0.274786155	-0.000617049	Viable
Log-Logistic	frequentist	Restricted	Extra Risk	0.1	15.85789	9.584413	29.14482	0.9258161	98.23575637	-	-0.06775637	-0.000623993	Viable
Multistage Degree 4	frequentist	Restricted	Extra Risk	0.1	19.33104	14.31258	36.20013	0.9892812	94.36832933	-	0.274786146	-0.000617049	Viable
Multistage Degree 3	frequentist	Restricted	Extra Risk	0.1	19.33104	14.31291	36.19987	0.9892812	94.36832933	-	0.274786164	-0.000617049	Viable
Multistage Degree 2	frequentist	Restricted	Extra Risk	0.1	19.33104	14.31285	36.20013	0.9892812	94.36832933	-	0.274786165	-0.000617049	Viable
Multistage Degree 1	frequentist	Restricted	Extra Risk	0.1	19.33104	14.3126	27.640746	0.9584861	96.36832933	-	0.274786131	-0.000619786	Viable
Weibull	frequentist	Restricted	Extra Risk	0.1	19.33104	14.31304	30.842788	0.9584861	96.36832933	-	0.274786138	-0.000617187	Viable
Logistic	frequentist	Unrestricted	Extra Risk	0.1	55.96478	43.343	71.751775	0.1446606	103.0027252	-	1.599996726	-1.362557141	Viable
Log-Probit	frequentist	Unrestricted	Extra Risk	0.1	14.77268	6.267248	25.715773	0.8052972	98.50125481	-	-0.227261455	-0.000617142	Viable - F
Probit	frequentist	Unrestricted	Extra Risk	0.1	51.68302	40.63077	65.781622	0.1906352	102.1597545	-	1.485406079	-1.271842634	Viable



Using BMDS – Option Set Results

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Option set #1 (Hover for details) Scroll right to see summary plot ->														
Model Analysis Type Restriction RiskType BMRF BMD BMDL BMDU P Value AIC Unnormalized Log Posterior Probability Scaled Residual for Dose Group near BMD Scaled Residual for Control Dose Group BMDS Re														
Dichotomous Hill frequentist Restricted Extra Risk 0.1 15.85789 7.897094 29.144824 0.6945918 100.2357564 - -0.067756196 -0.00062655 Viable														
Gamma frequentist Restricted Extra Risk 0.1 19.33104 14.31304 30.451431 0.9892812 94.36832933 - 0.274786155 -0.000617049 Viable														
Log-Logistic frequentist Restricted Extra Risk 0.1 15.85789 9.584413 29.14482 0.9258161 98.23575637 - -0.06775637 -0.000623993 Viable														
Multistage Degree 4 frequentist Restricted Extra Risk 0.1 19.33104 14.31258 36.20013 0.9892812 94.36832933 - 0.274786146 -0.000617049 Viable														
Multistage Degree 3 frequentist Restricted Extra Risk 0.1 19.33104 14.31291 36.19987 0.9892812 94.36832933 - 0.274786164 -0.000617049 Viable														
Multistage Degree 2 frequentist Restricted Extra Risk 0.1 19.33104 14.31285 36.20013 0.9892812 94.36832933 - 0.274786165 -0.000617049 Viable														
Multistage Degree 1 frequentist Restricted Extra Risk 0.1 19.33104 14.3126 27.640746 0.9584861 96.36832933 - 0.274786131 -0.000619786 Viable														
Weibull frequentist Restricted Extra Risk 0.1 19.33104 14.31304 30.842788 0.9584861 96.36832933 - 0.274786138 -0.000617187 Viable														
Logistic frequentist Unrestricted Extra Risk 0.1 55.96478 43.343 71.751775 0.1446606 103.0027252 - 1.599996726 -1.362557141 Viable														
Log-Probit frequentist Unrestricted Extra Risk 0.1 14.77268 6.267248 25.715773 0.8052972 98.50125481 - -0.227261455 -0.000617142 Viable - F														
Probit frequentist Unrestricted Extra Risk 0.1 51.68302 40.63077 65.781622 0.1906352 102.1597545 - 1.485406079 -1.271842634 Viable														
Quantal Linear frequentist Unrestricted Extra Risk 0.1 19.33104 14.31304 27.651236 0.9892812 94.36832933 - 0.274786154 -0.000617049 Viable														
Dichotomous Hill bayesian - Extra Risk 0.1 23.88815 10.16551 49.501684 - - -53.13822466 0.221120126 -0.855817711														
Gamma bayesian - Extra Risk 0.1 26.35795 12.17972 48.908685 - - -53.19703399 0.224861091 -0.943529669														
Logistic bayesian - Extra Risk 0.1 55.08391 42.76971 70.818903 - - -54.63520879 1.438333356 -1.511623327														
Log-Logistic bayesian - Extra Risk 0.1 25.56395 10.75297 52.479612 - - -52.62188353 0.226231253 -0.938424257														
Log-Probit bayesian - Extra Risk 0.1 31.95571 14.35433 64.871016 - - -54.19902941 0.617271666 -1.062742709														
Multistage Degree 4 bayesian - Extra Risk 0.1 29.70898 19.85807 46.212977 - - -56.57595301 0.336005674 -0.952118967														
Multistage Degree 3 bayesian - Extra Risk 0.1 28.01126 18.92686 43.324266 - - -53.23693243 0.282571943 -0.936621658														
Multistage Degree 2 bayesian - Extra Risk 0.1 25.65265 17.68658 39.31189 - - -52.2005658 0.194760923 -0.914261291														
Multistage Degree 1 bayesian - Extra Risk 0.1 21.897 16.01266 31.302091 - - -50.234466 0.016111501 -0.878679404														
Probit bayesian - Extra Risk 0.1 51.63621 40.60085 65.922653 - - -53.92133341 1.435643794 -1.355140023														
Quantal Linear bayesian - Extra Risk 0.1 21.51394 15.51436 31.550179 - - -50.98970952 0.000679148 -0.870151483														
Weibull bayesian - Extra Risk 0.1 29.45342 12.79312 58.812027 - - -53.25948327 0.343704366 -0.984798192														
Model Average bayesian MA - Extra Risk 0.1 24.88928 14.74397 49.453226 - - - - -														
Summary Abbreviations freq-dhl-rest-opt1 freq-gam-rest-opt1 freq-lnl-rest-opt1 freq-mst4-rest-opt1 ...														






Using BMDS – Individual Model Results

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Dichotomous Results

BMDS 3.1.2

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User Input

Info	
Model	frequentist Log-Probit v1.1
Dataset Name	BMDS - training - Dichotomous
User notes	[Add user notes here]
Dose-Response Model	$P[\text{dose}] = g + (1-g) * \text{CumNorm}(a+b*\text{Log}(\text{Dose}))$

Model Options	
Risk Type	Extra Risk
BMR	0.1
Confidence Level	0.95
Background	Estimated

Model Data	
Dependent Variable	Dose
Independent Variable	Incidence
Total # of Observations	5

Model Results

Scroll right to see BMD Cumulative Distribution Function (CDF) table →
Scroll down to see Dose Response Plot ↓

Benchmark Dose	
BMD	14.77267729
BMDL	6.267247615
BMDU	25.71577276
AIC	98.50125481
P-value	0.805297172
D.O.F.	2
Chi ²	0.433087824

Model Parameters	
# of Parameters	3
Variable	Estimate
g	1.52346E-08
a	-2.890901327
b	0.597653782

Goodness of Fit					
Dose	Estimated Probability	Expected	Observed	Size	Scaled Residual
0	1.52346E-08	3.80864E-07	0	25	-0.000617
5	0.026864535	0.671613383	1	25	0.4061993
20	0.135559224	3.388980604	3	25	-0.227261
80	0.392823885	9.820597129	9	25	-0.33605

Ready | freq-wei-rest-opt1 | freq-log-unrest-opt1 | **freq-lnp-unrest-opt1** | freq-pro-unrest-opt1 | freq-qln-unrest-opt1 | 100%






Using BMDS – Bayesian Model Averaging Results

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   **Model Average Results**

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[Return to Summary](#)

BMDS 3.1.2

User Input

Info	
Model	Bayesian Model Averaging v1.0
Dataset Name	BMDS - training - Dichotomous
User notes	[Add user notes here]

Model Options	
Risk Type	Extra Risk
BMR	0.1
Confidence Level	0.95
Background	Estimated

Model Data	
Dependent Variable	Dose
Independent Variable	Incidence
Total # of Observations	5

Model Results

Scroll right to see BMD Cumulative Distribution Function (CDF) table →
Scroll down to see Dose Response Plot ↓

Benchmark Dose	
BMD	24.88927534
BMDL	14.74397033
BMDU	49.45322573

MA - Individual Models				
Model	Posterior Probability	BMD	BMDL	BMDU
Dichotomous Hill	0.060032456	23.88815284	10.1655111	49.50168
Gamma	0.056603794	26.35795176	12.1797211	48.90868
Logistic	0.01343551	55.08390665	42.7697092	70.8189
Log-Logistic	0.100607467	25.56394637	10.7529737	52.47961
Log-Probit	0.020781862	31.95570707	14.3543258	64.87102
Multistage	0.153322601	25.65264702	17.6865846	39.31189
Probit	0.027433821	51.6362071	40.6008542	65.92265
Quantal Linear	0.514605448	21.51394337	15.5143648	31.55018
Weibull	0.053177041	29.45342064	12.7931193	58.81203

bayes-mst2-opt1 | bayes-mst1-opt1 | bayes-pro-opt1 | bayes-qln-opt1 | bayes-wei-opt1 | **DichoMA-option1** ...

100%

- **Research into model averaging methods currently included in HERA research portfolio:**
 - HERA 3.5.1 – Research in support of informative parameter priors and used in Bayesian model averaging of dichotomous and continuous endpoints
 - HERA 3.5.3 – Development of an updated model suite for dichotomous and continuous toxicological data
 - Outputs of these HERA products directly informs improvement of dichotomous model averaging and development of continuous model averaging
- **Development of Bayesian continuous model averaging is undergoing**
 - Allows for averaging across distributions if individual animal data is available
 - Allows for averaging across variance models when assuming Normal distribution
 - Planned for release with BMDS 3.4

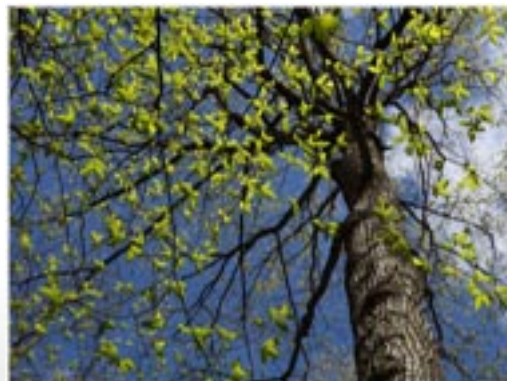


Quantitative Uncertainty Analysis & APROBA

Todd Blessinger, PhD

Center for Public Health and Environmental Assessment
U.S. Environmental Protection Agency

*Presentation at BOSC 2021
February 5, 2021*



- Quantitative uncertainty analysis, such as moving towards “risk-specific dose” estimates, in human health assessments recommended by National Research Council (2009, 2014), especially for reference value derivation
 - Increases transparency
 - Allows greater flexibility
- WHO/IPCS developed a method to estimate the uncertainty in a “target human dose” (2017 guidance document)

Current Reference Value Derivation

➤ Reference values:

➤
$$RfV = \frac{POD}{UF_1 \times \dots \times UF_k}$$

➤ POD: point of departure; UF: uncertainty factor

➤ Currently derived as point value (“deterministically”)

➤ WHO/IPCS approach is probabilistic.

Alternative Reference Value Derivation based on HD_M^I

- HD_M^I = the **human dose** at which a **fraction (or incidence) I** of the **population** experiences an effect of **magnitude (or severity) M or greater** for the critical effect considered.
- Ex: Endpoint of interest: relative liver weight decrease
 - HD_{05}^{01} = dose at which 1% of the population experiences a decrease in relative liver weight of 5% or greater.

Alternative Reference Value Derivation based on HD_M^I

- $HD_M^I = \frac{POD}{AF_1 \times \dots \times AF_k}$
 - POD: point of departure
 - AF_i 's: "assessment factors"
- Point of departure and assessment factors are treated as random variables with probability distributions.
- HD_M^I is a random variable.

Alternative Reference Value Derivation based on HD_M^I

- Allows estimation of “risk-specific dose”
 - Ex: HD_{05}^{05} = dose at which 5% of the population experiences a decrease in relative liver weight of 5% or greater.

Approximate Probabilistic Analysis

- Approximate Probabilistic Analysis (APROBA)
 - Excel-based tool for applying HD_M^I method
 - The point of departure and assessment factors are assumed to be independent and lognormally distributed.
 - HD_M^I is lognormally distributed.

Approximate Probabilistic Analysis

- Provisional lognormal parameter values provided for commonly used assessment factors
 - Interspecies, duration extrapolation, interhuman variability
 - Based on empirical data
- Other parameter values can be entered

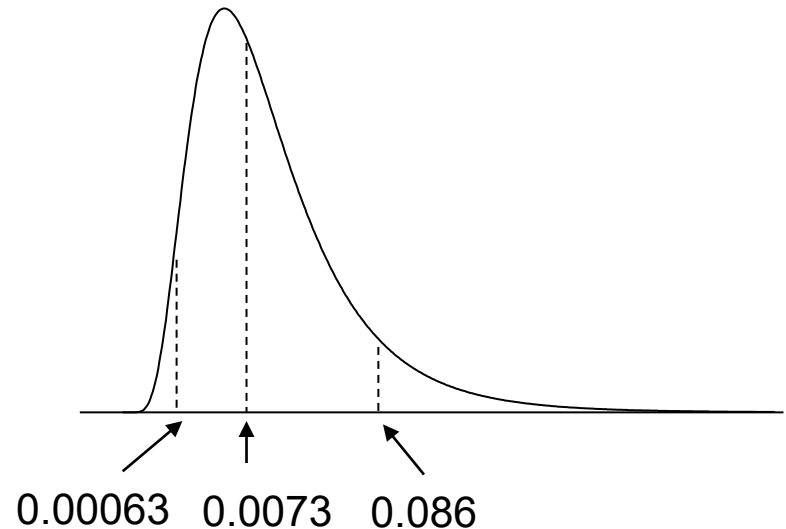
Application of APROBA to acrolein

- Application of APROBA to acrolein (Blessinger et al., 2020)
 - Endpoint: Incidence of lesions in the nasal respiratory epithelium, from Dorman et al. (2008)
 - Subchronic inhalation study in rats
 - Used $I = 1\%$ for incidence of nasal lesions of minimal severity
- HD_{minimal}^{01} = concentration that results in lesions of at least minimal severity in the nasal respiratory epithelium in 1% of a general human population.

Application of APROBA to acrolein

- HD_{minimal}^{01} has lognormal distribution with estimated percentiles:

Percentile	Value (mg/m ³)
5 th	0.00063
50 th (median)	0.0073
95 th	0.086



- 5th percentile = 0.00063 mg/m³ can be considered a “probabilistic reference concentration”.

Risk-Specific Concentration

- Risk-specific concentration: human incidence I can be adjusted
 - Ex: For $I = 5\%$, HD_{minimal}^{05} percentiles

Percentile	Value (mg/m ³)
5 th	0.0015
50 th (median)	0.014
95 th	0.133

- Expansion of APROBA and HD_M^I method:
 - Update provisional parameter values by collecting and analyzing additional toxicology data (current distributions too narrow?).
 - Relax the restriction of independent, lognormally distributed components (use numerical methods to estimate HD_M^I distribution).
 - Incorporate database uncertainty.

Summary & Conclusions

- HDMI method allows explicit, quantitative estimation of risk-specific dose.
- Quantitative uncertainty analysis methods require continual advancement.
- Input from users is crucial.

- WHO/IPCS (2017): Guidance document on evaluating and expressing uncertainty in hazard characterization, 2nd ed.
- Chiu WA & Slob W (2015): A unified probabilistic framework for dose-response assessment of human health effects. Environ Health Perspect 123(12):1241–1254.
- Blessinger et al. (2020): Application of a unified probabilistic framework to the dose-response assessment of acrolein Environ Int 143. Available online