

Quantifying Uncertainty in Dose

From the perspective of a person who evaluates uncertainty in environmental pathway models

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Let's start with an example problem

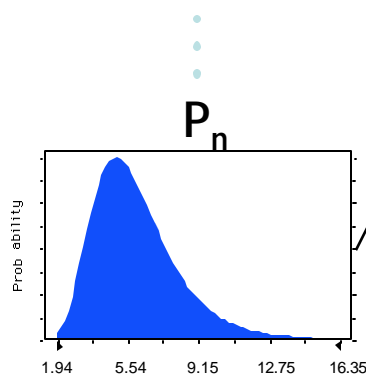
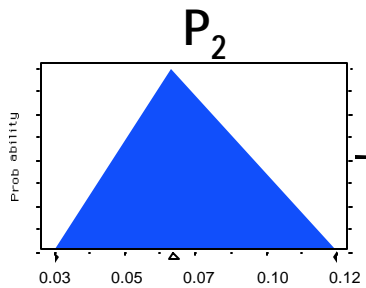
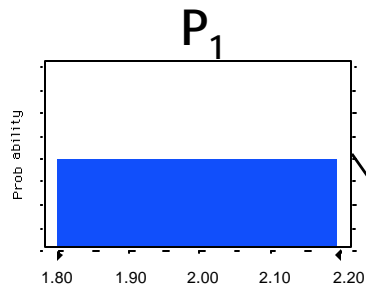
- Dose reconstruction for an accident decades ago in which a short-term release occurred of a short-lived radionuclide.
- There are no environmental measurements
- Meteorological dispersion must be inferred from present-day information
 - 90% chance the wind blew to the north
 - Excess disease seen in the south
- What do we conclude?

History of Uncertainty Analysis in Dose Reconstruction

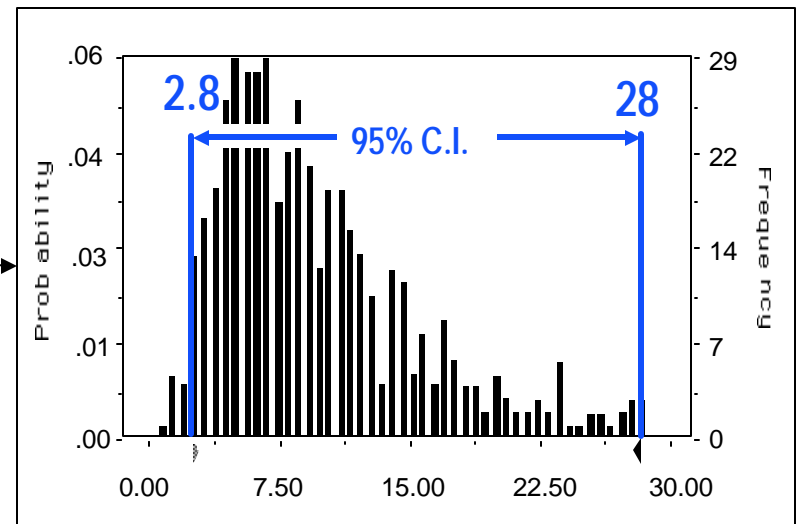
- **Quantification of uncertainty in environmental pathway models**
 - First efforts began more than 30 years ago
 - Algebraic solutions used for multiplicative and additive terms in equations
 - Monte Carlo simulation
 - with uncertain model parameters described as discrete or continuous probability distributions

Estimating Uncertainty for an Individual

Parameters 1, 2, ..., n \longrightarrow Model \longrightarrow Model Result



$$R = f(P_1, P_2, \dots, P_n)$$



Dose

The First Epidemiological Study to Address Uncertainty in Dose Reconstruction

University of Utah Investigation of Childhood Cohorts Exposed to NTS Fallout
(Stevens et al, 1992, Kerber et al, 1993)

- Dose uncertainties estimated for each individual in the cohort
 - Algebraic solutions used for equations described by products and summations,
 - Monte Carlo used for some model components
 - Dose per person given as a GM and GSD from which an arithmetic mean was obtained.

**The First Epidemiological Study
to Address Uncertainty in
Dose Reconstruction
for Every Individual in the Cohort**

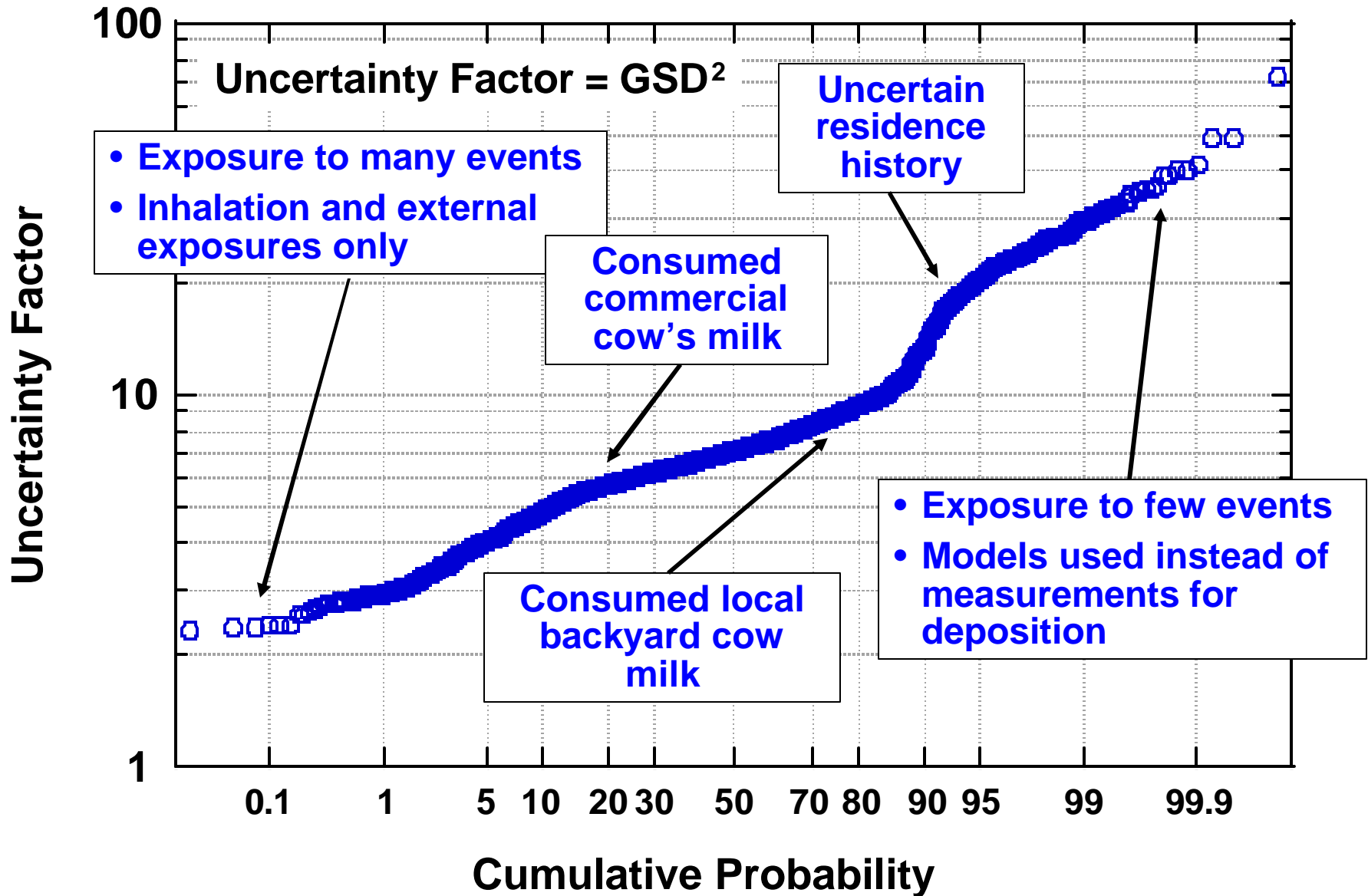
**University of Utah Investigation of
Childhood Cohorts Exposed to NTS Fallout**
(Stevens et al, 1992, Kerber et al, 1993)

University of Utah Investigation of Childhood Cohorts Exposed to NTS Fallout

(Stevens et al, 1992, Kerber et al, 1993)

- Primary dose response based on arithmetic mean dose for each individual
 - Initial analysis of dose uncertainty treated all uncertainty as classical measurement error
- Mixtures of classical, Berkson and shared uncertainties considered in more recent work
 - (Malick et al 2004, Lyon et al 2006, and Li et al., 2007)

Uncertainty Factors for NTS Cohort



Hanford Thyroid Disease Study was the First to Use a Full Monte Carlo-Based Uncertainty Analysis of Dose

(Davis et al., 2002, 2004)

- The median individual dose used for primary dose-response analysis
- Amount of dose uncertainty per person somewhat less than for NTS cohort
- Analysis of dose response assumed dose uncertainty as 100% Berkson error
 - Evaluation of varying degrees of potential bias in model parameters not evident

Estimating Dose Uncertainty for an Epidemiological Cohort

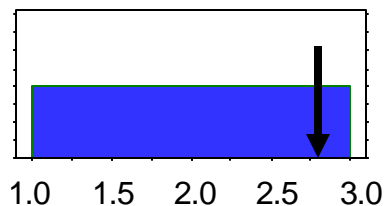
- Stochastic variability of true dose among individuals should be separated from lack of knowledge about true but unknown fixed quantities
- Individuals within and across subgroups in the cohort may be affected by potential bias in model parameters
 - Examples: Amount released, Wind direction, GM and GSD of milk transfer factor

Separation of random variability of true exposure and dose from quantities that are fixed (true) but unknown

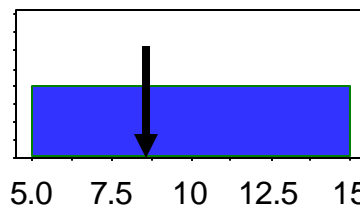
- **Requires a 2-Dimensional Monte Carlo Approach**
 - First described in the 1980's
 - The terms “Type A” and “Type B” uncertainty introduced
 - (IAEA Safety Series No. 100, 1989)

Result = Y × Z

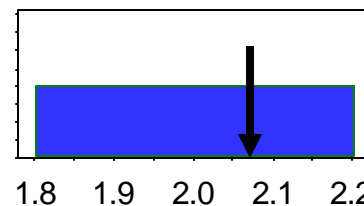
$Y_1 = 2.78$



$Z_{GM1} = 8.56$

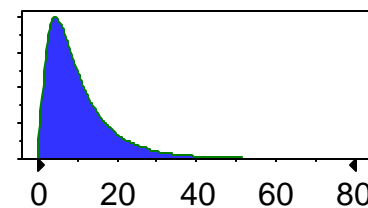


$Z_{GSD1} = 2.07$

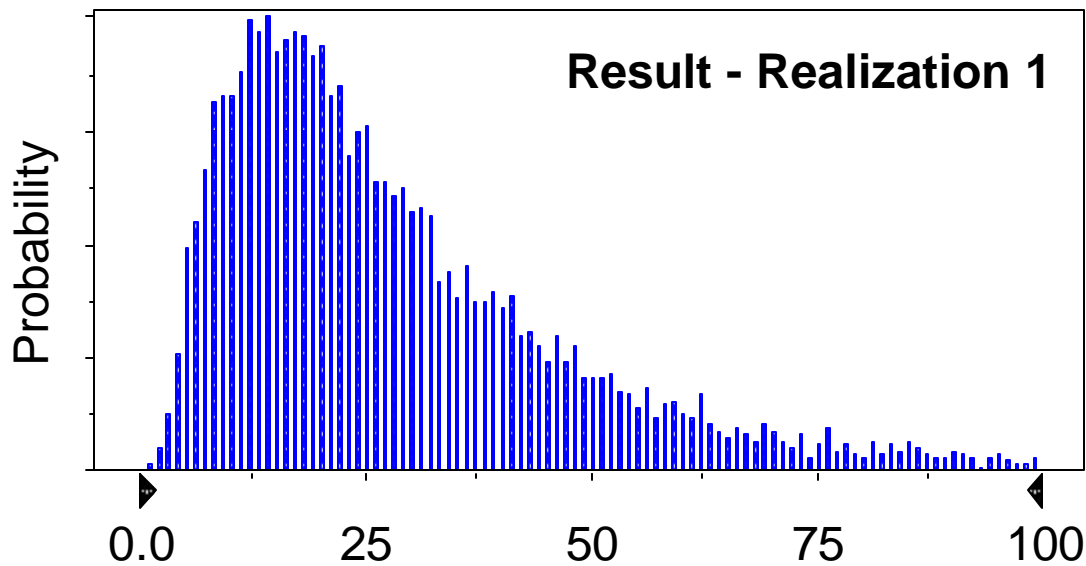


2.78

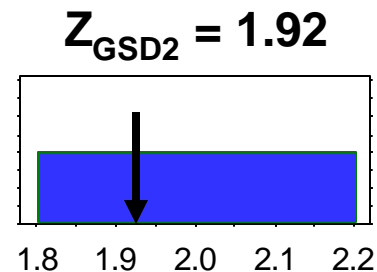
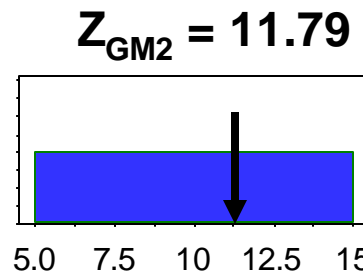
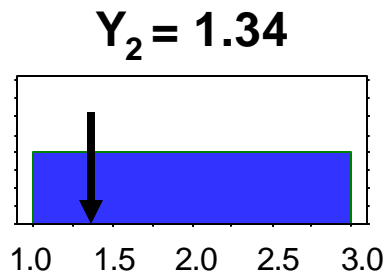
Z = Lognormal (8.56, 2.07)



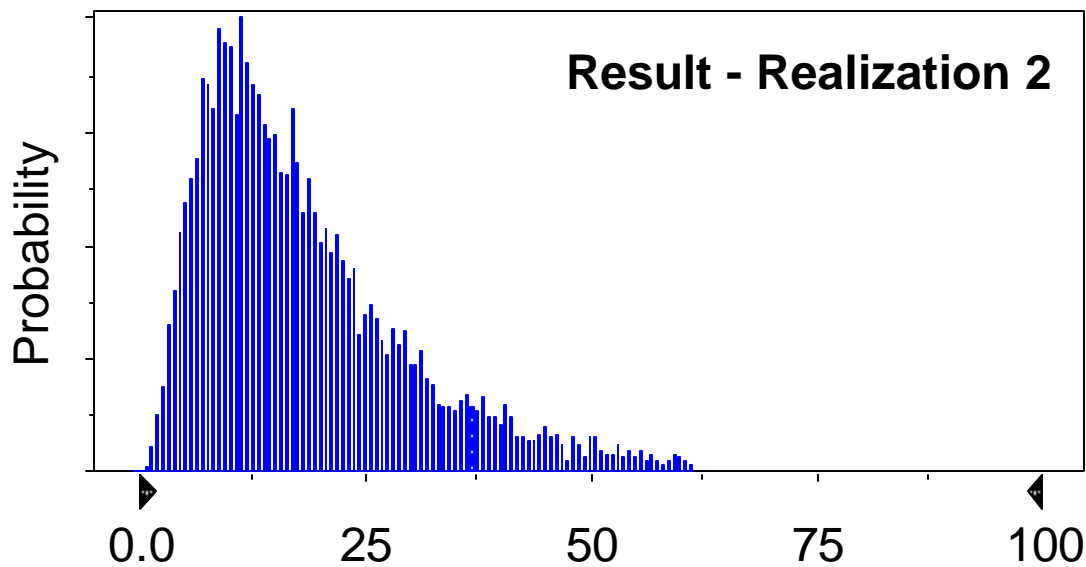
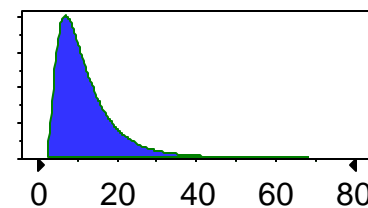
Result - Realization 1



Result = Y × Z

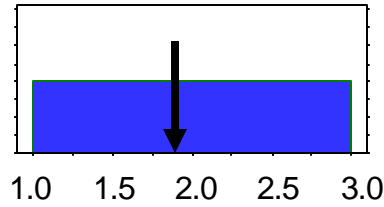


$Z = \text{Lognormal}(11.79, 1.92)$

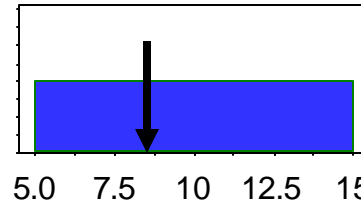


Result = Y × Z

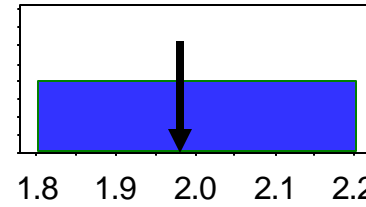
$Y_3 = 1.95$



$Z_{GM3} = 8.58$

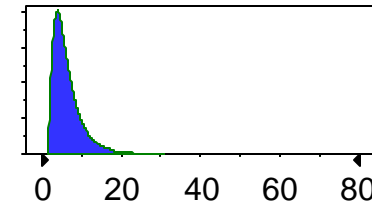


$Z_{GSD3} = 1.96$

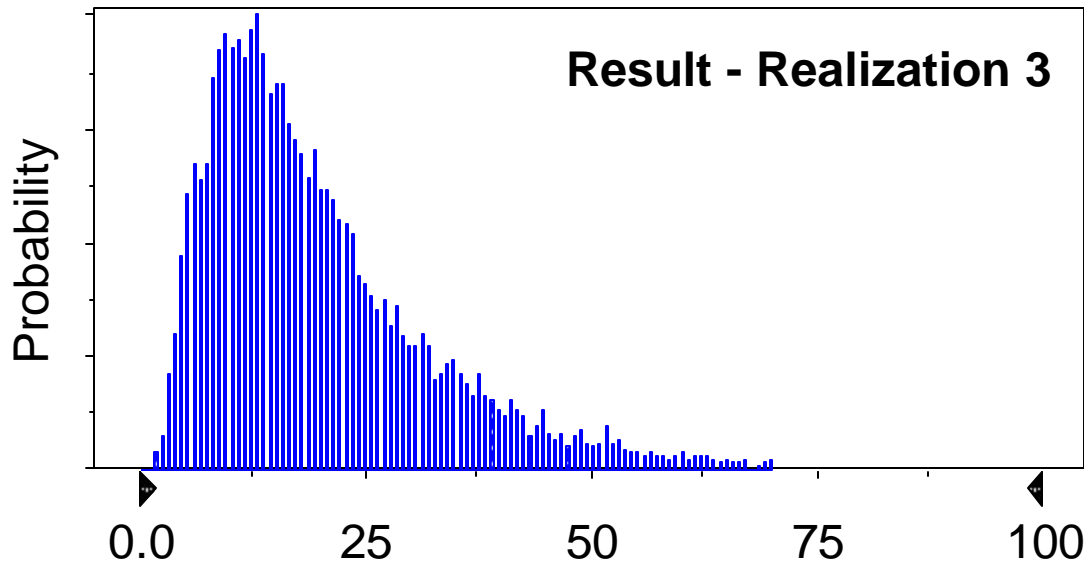


1.95

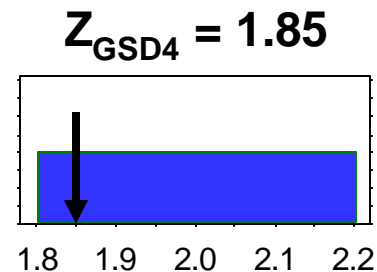
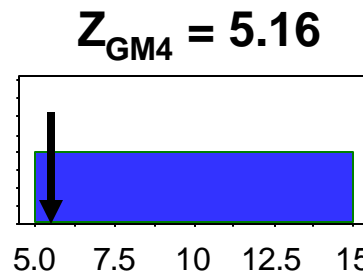
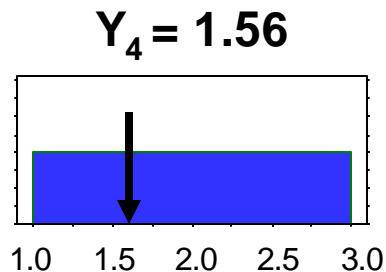
$Z = \text{Lognormal}(8.58, 1.96)$



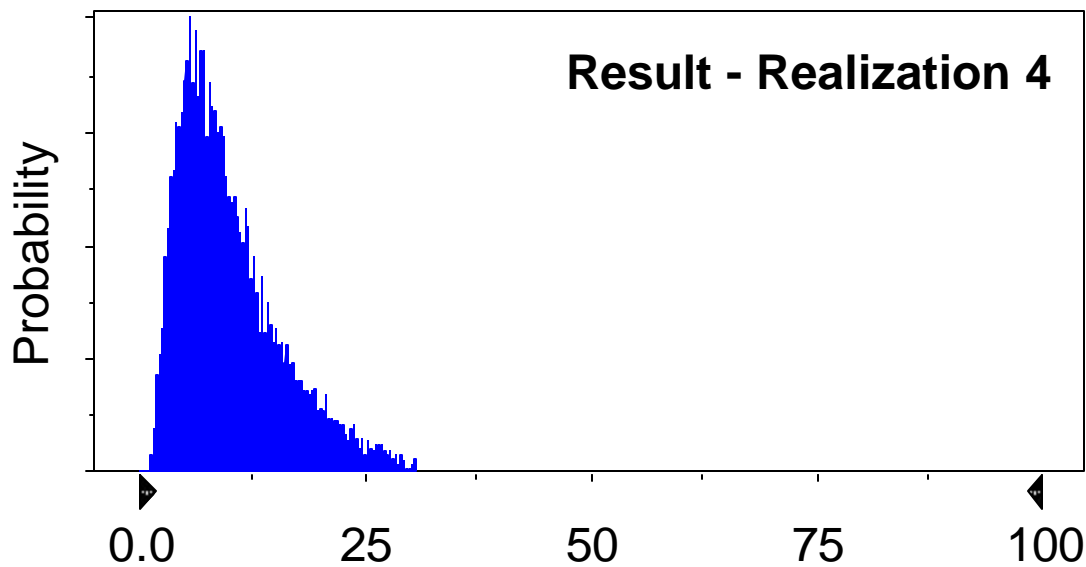
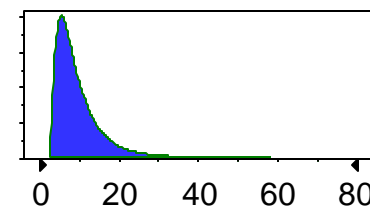
Result - Realization 3



Result = Y × Z

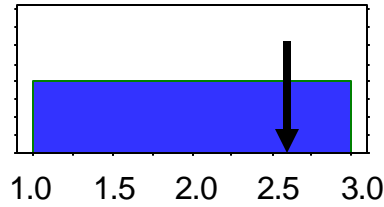


$Z = \text{Lognormal}(5.16, 1.85)$

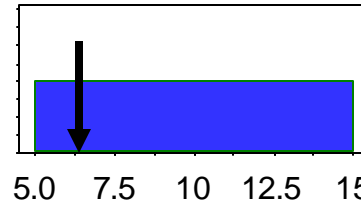


Result = Y × Z

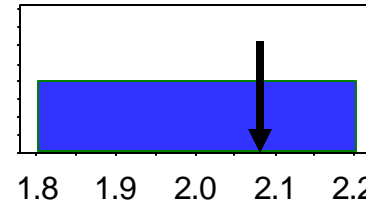
$Y_5 = 2.56$



$Z_{GM5} = 6.01$

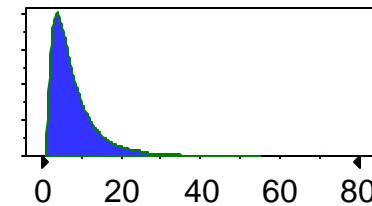


$Z_{GSD5} = 2.09$

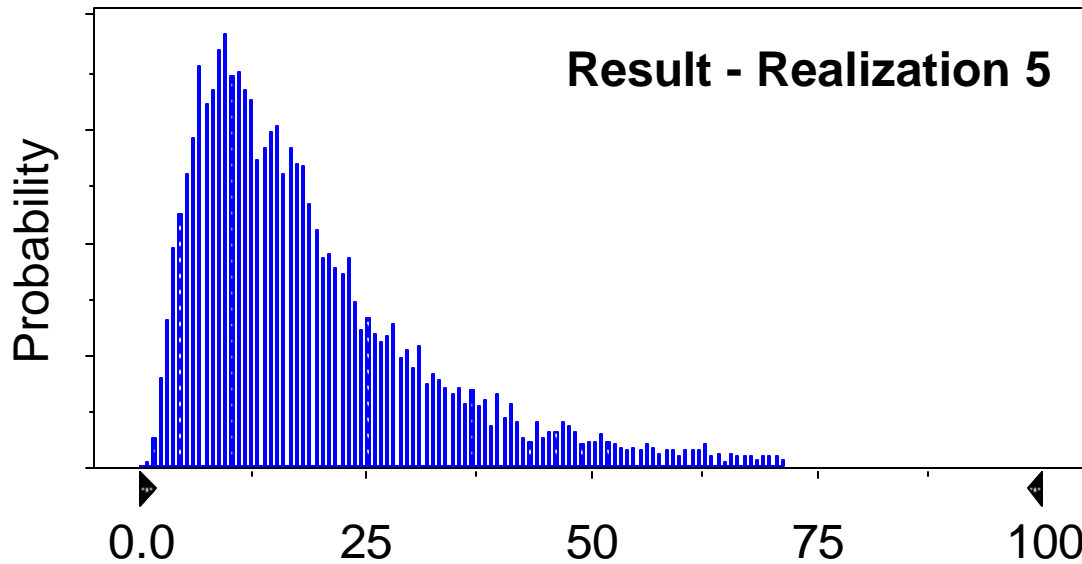


1.56

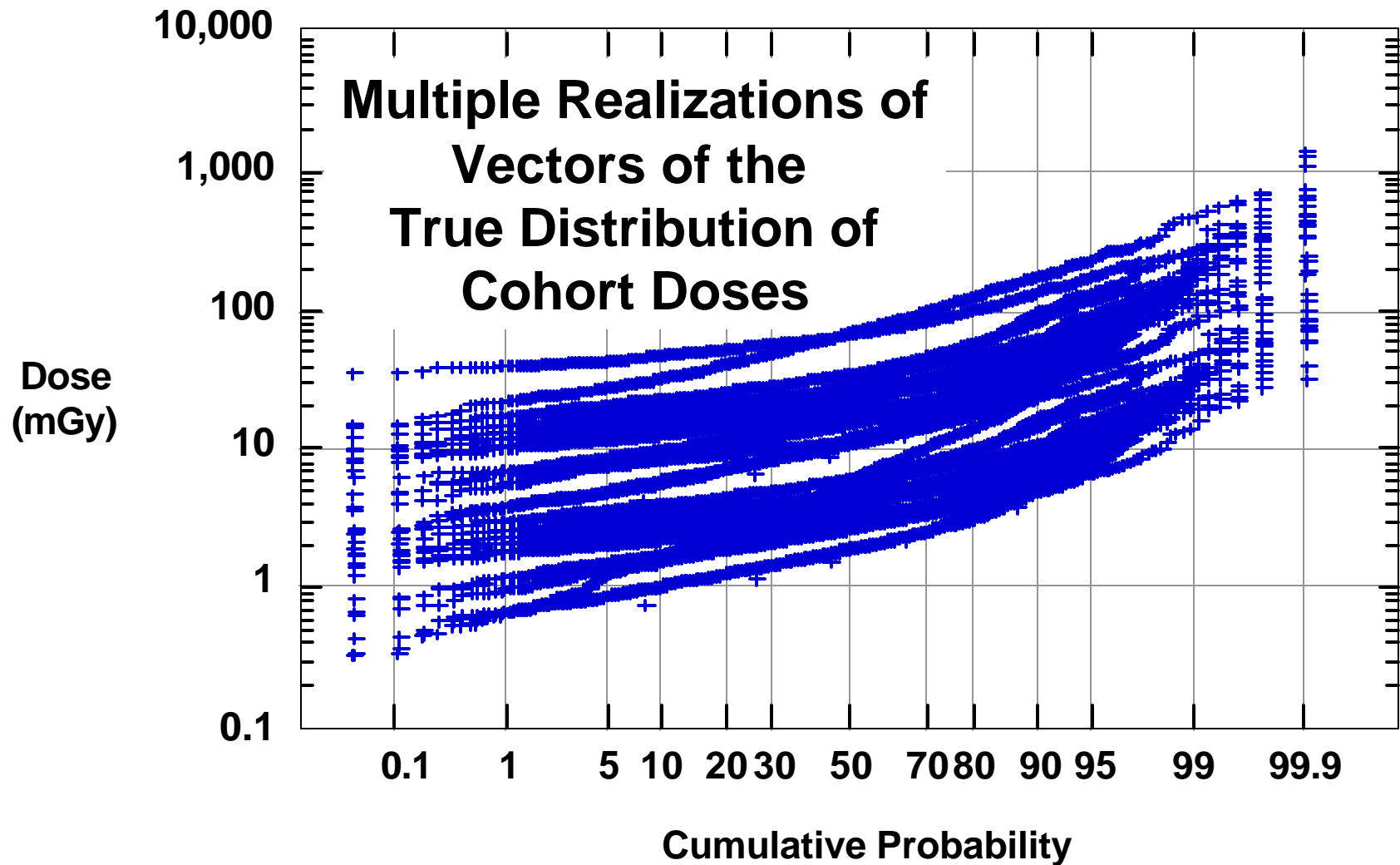
$Z = \text{Lognormal}(6.01, 2.09)$



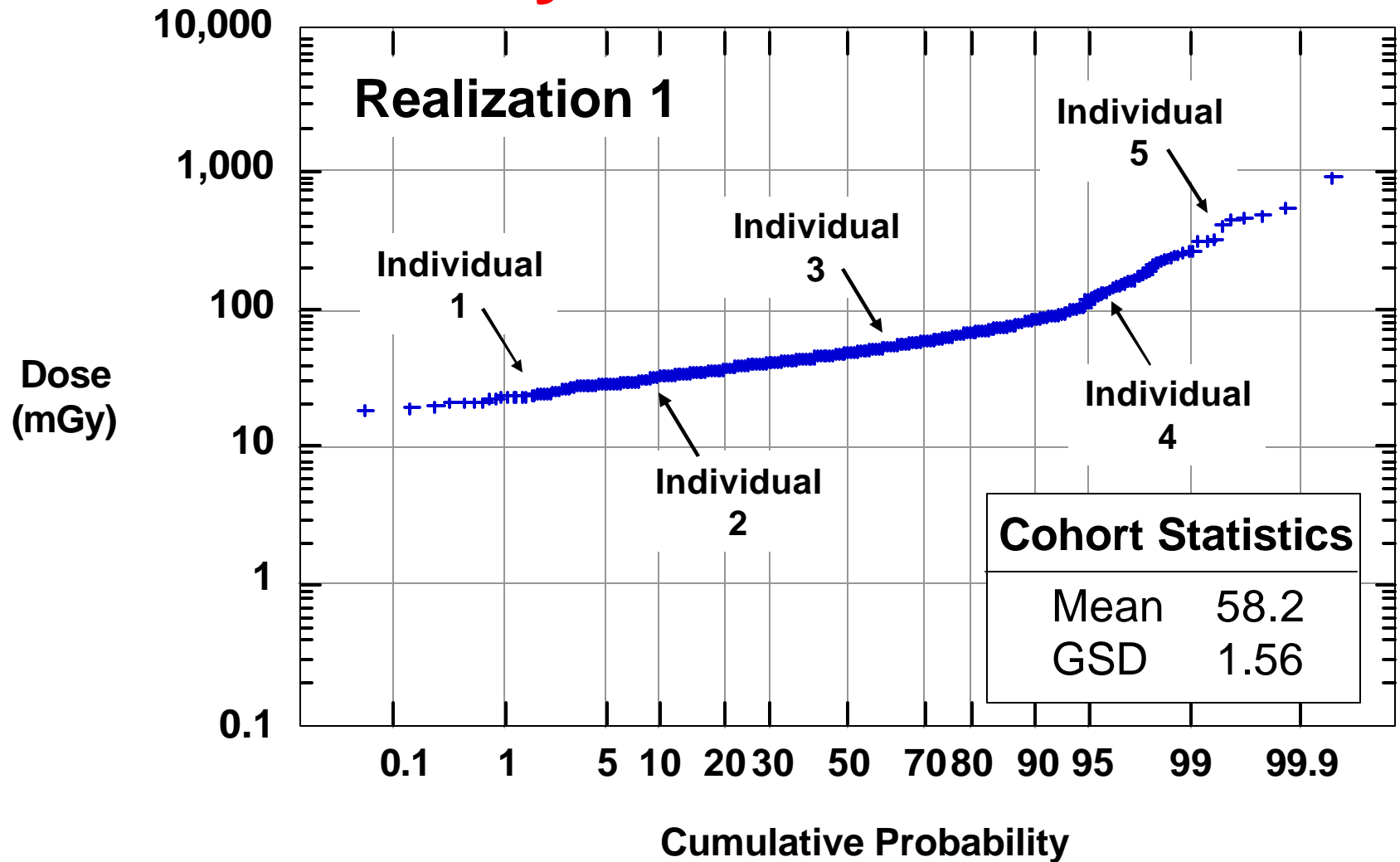
Result - Realization 5



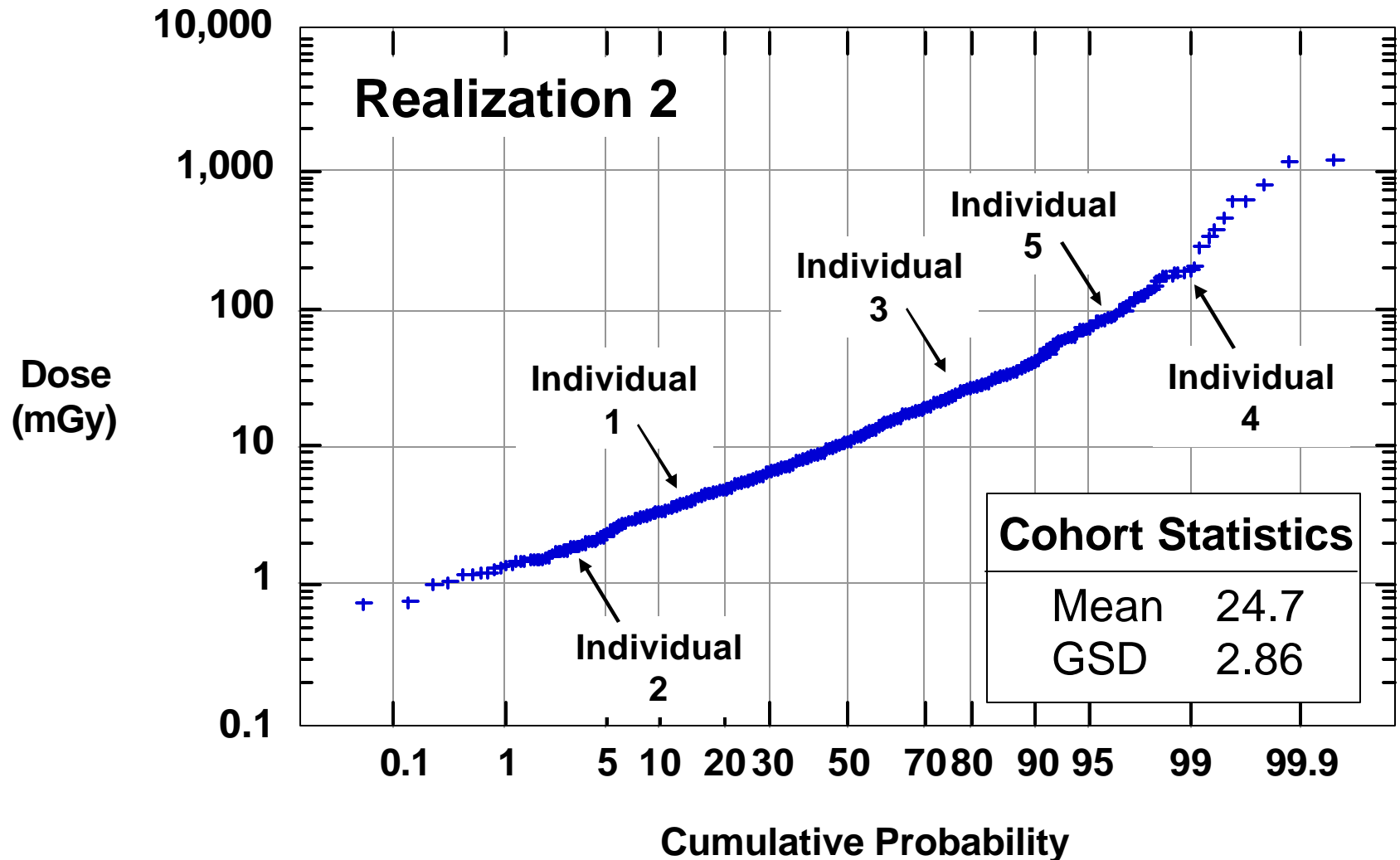
A 2-D Monte Carlo dose reconstruction for a cohort



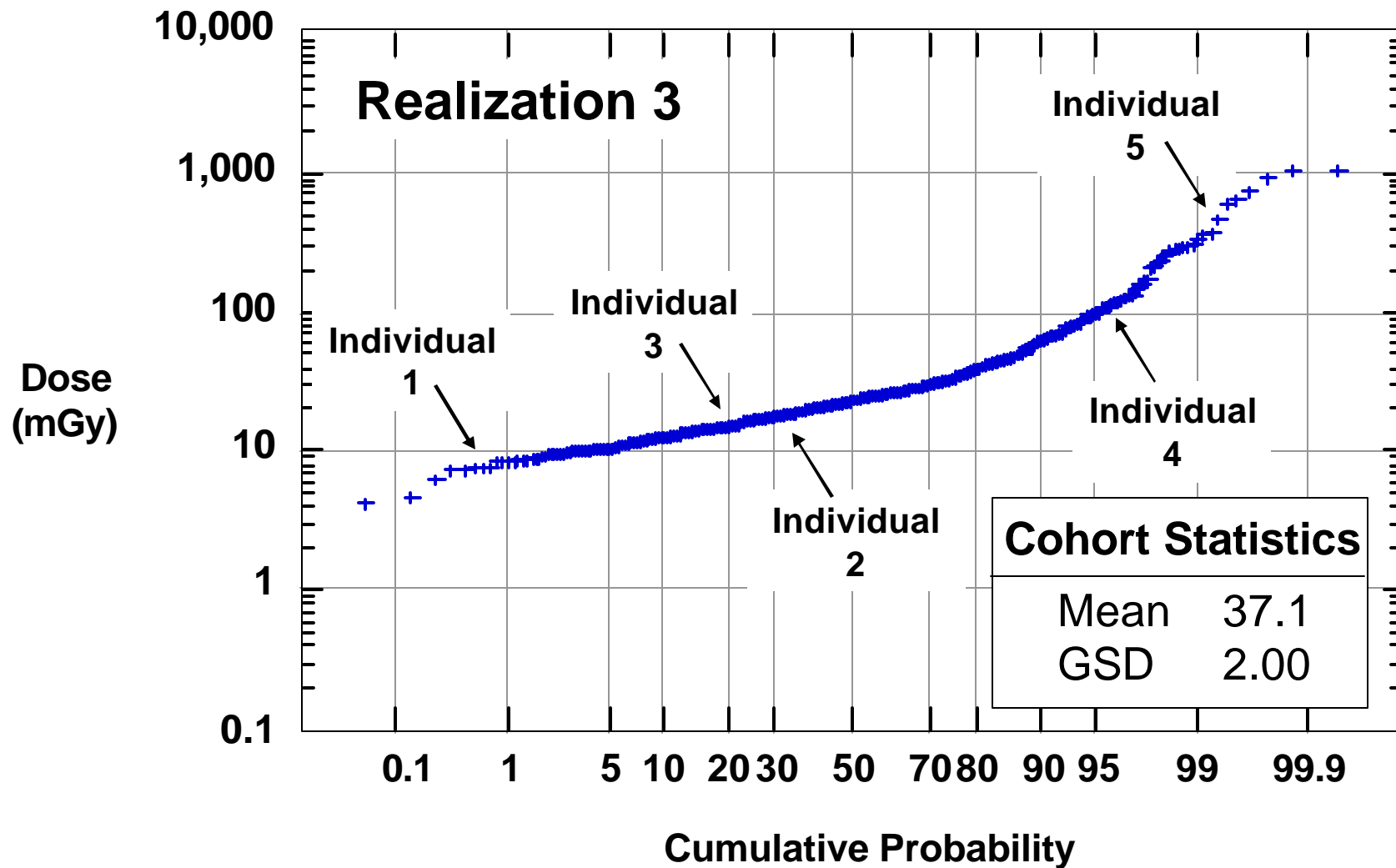
An example of a single realization from a 2-D Monte Carlo uncertainty analysis for a cohort



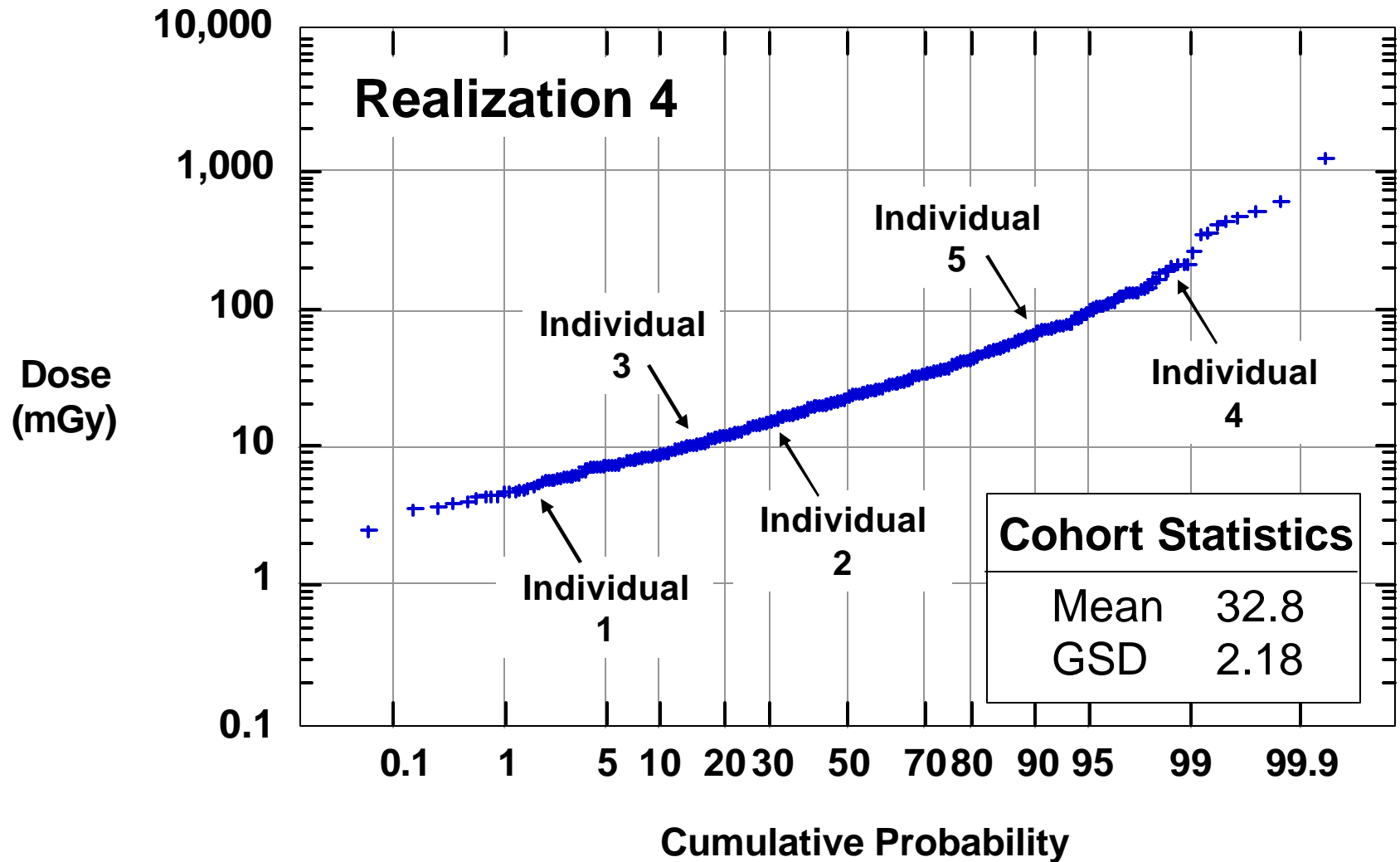
An example of a 2-D Monte Carlo uncertainty analysis for a cohort



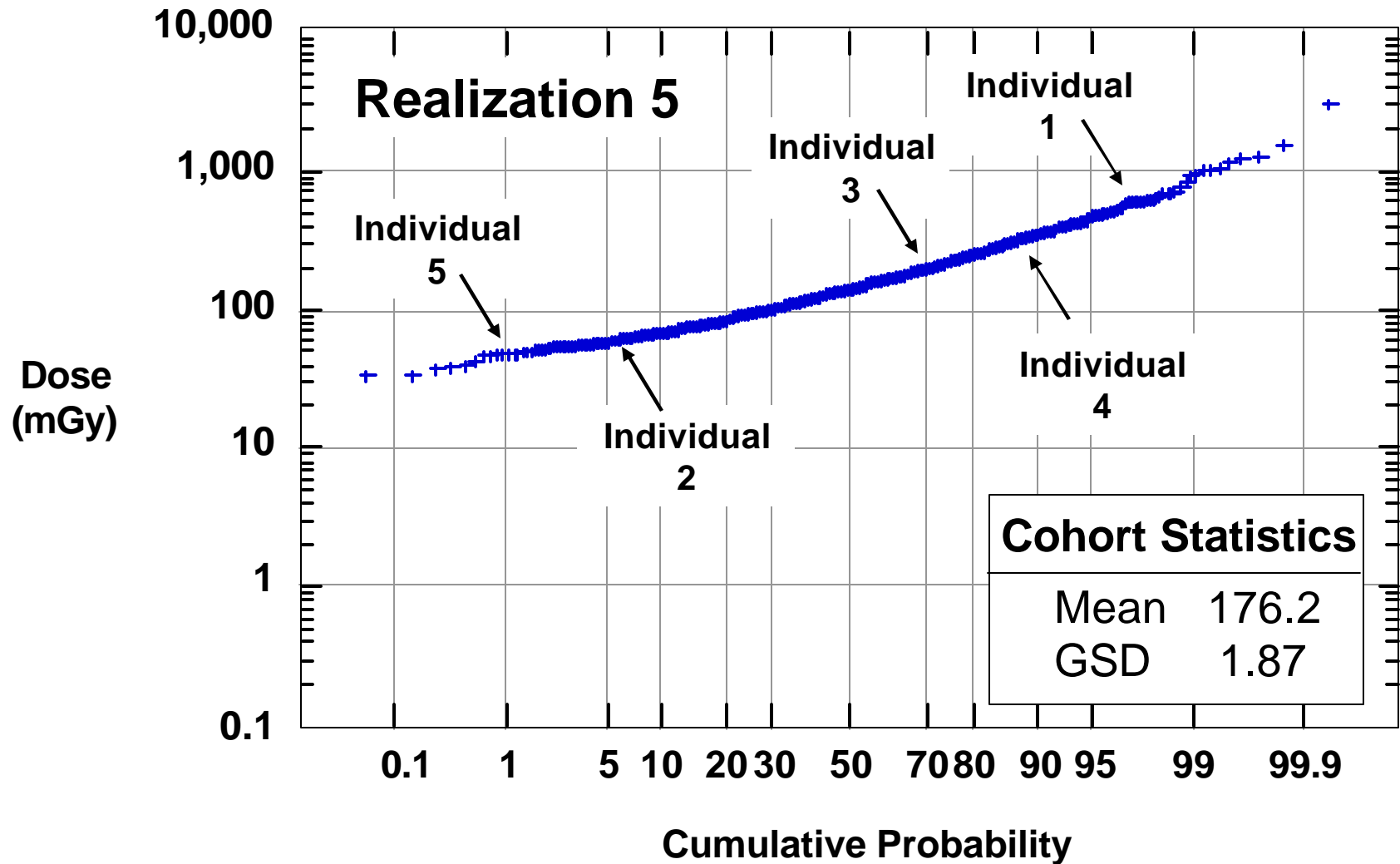
An example of a 2-D Monte Carlo uncertainty analysis for a cohort



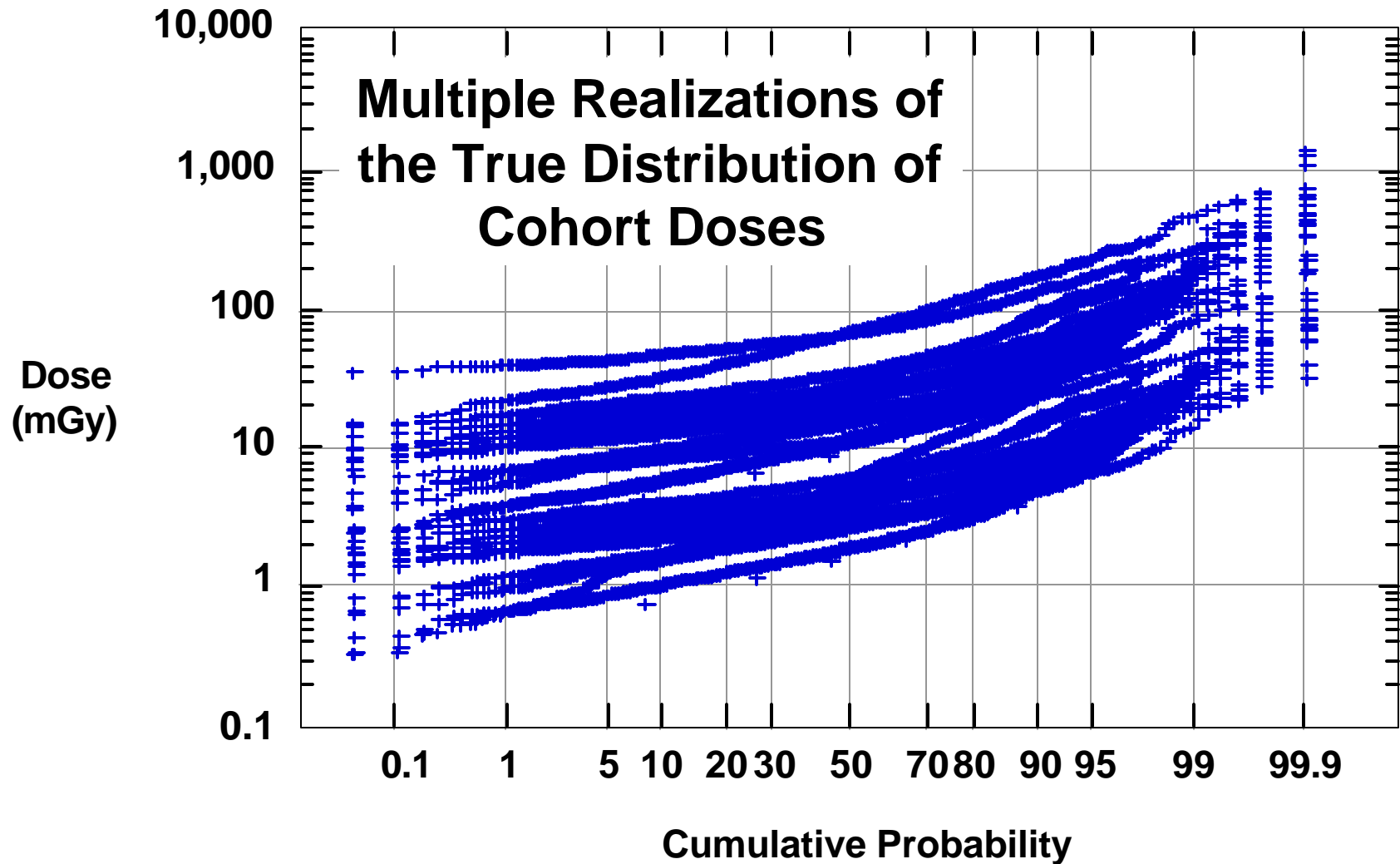
An example of a 2-D Monte Carlo uncertainty analysis for a cohort



An example of a 2-D Monte Carlo uncertainty analysis for a cohort



What do Epidemiologists do with results from a 2-D Monte Carlo Simulation?



What do Epidemiologists do with output from a 2-D Monte Carlo Simulation?

- Use mean dose as a surrogate for the true dose and ignore uncertainty (regression calibration)?
- Assign equal weight to each alternative dose response and average?
- Obtain the likelihood of each realization of dose response and average?
- Perform more complex analyses as in Li et al (2007)?
 - To account for mixtures of classical and Berkson errors
 - To account for shared uncertainties due to possible bias in parameter values and model structure

Important considerations when evaluating a 2-D Monte Carlo Dose Reconstruction

- All realizations of dose vectors may be “wrong”, hopefully some will be “less wrong” than others
- When there is the potential for a high degree of systematic error (bias) shared by subgroups of the cohort
 - Only a few alternative realizations of dose vectors might approximate the true distribution of individual doses
- Under these conditions, it is not recommended
 - To use the mean dose as a surrogate for the true dose
 - To give equal weight to each alternative dose response

Recommendation

- Perform a dose response analysis
 - On each realization of the cohort distribution of doses
- Evaluate and weight each alternative dose response for degrees of plausibility
 - Use goodness of fit to disease outcomes
 - Perform investigations to further justify why some realizations of dose and dose response are more plausible than others
- If an unbiased mean dose per person is desired,
 - Produce alternative vectors of possibly unbiased mean doses

Don't overlook QA/QC of the dose reconstruction model and input data

- Complex computer codes and data bases frequently contain errors
- Consider redundant computational platforms programmed independently
 - Compare intermediate results and dose estimates
 - Compare estimates of uncertainty
 - Resolve discrepancies

“The biggest problems are caused by things you think you know for sure, that just ain't so”

Conclusions

- When dose uncertainties are high and complex, be skeptical of dose -response analyses that conclude:
 - “Dose uncertainties did not affect the dose-response”
- Use 2-D Monte Carlo to separate random variability from lack of knowledge about true fixed quantities
 - Weight each realization of a cohort distribution of true doses by degrees of plausibility
 - Dosimetrists should participate in this evaluation
- Resist averaging doses across realizations
- Explicit evaluation of each realization of a possibly true vector of cohort doses will identify
 - situations associated with low statistical power
 - situations that give an improved fit to health outcomes