

# Predicting Low Dose Effects for Chemicals in High Through-Put Studies

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# Outline

1. Introduction
2. The Example and Study Data
3. Testing Hypotheses about Low Dose Response
4. Estimating Response at Low Doses
5. Better Estimates: Predictors of Realized Random Effects
6. Conclusions

# Introduction

- High through-put studies common
- Can data be ‘harvested’?
  - Cons: Ad-hoc, Post-hoc, Low Power, Bias, not significant
  - Pros: Low cost, Common in ‘observational’ studies, Can learn something
- Illustrate use to assess response at low doses-
- Focus: Statistical Models and Interpretation

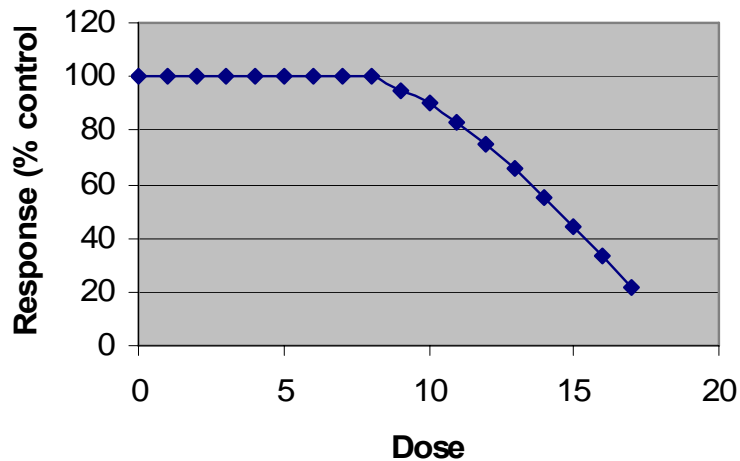
# Example and Study Data

NCI Yeast data- 2189 chemicals, 13 yeast strains

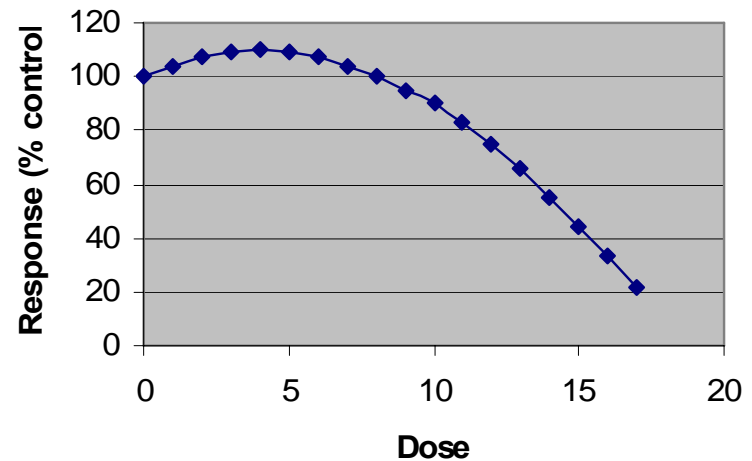
5 doses x 2 replications-

Focus on doses in Low Dose Region

**Threshold Dose-Response**

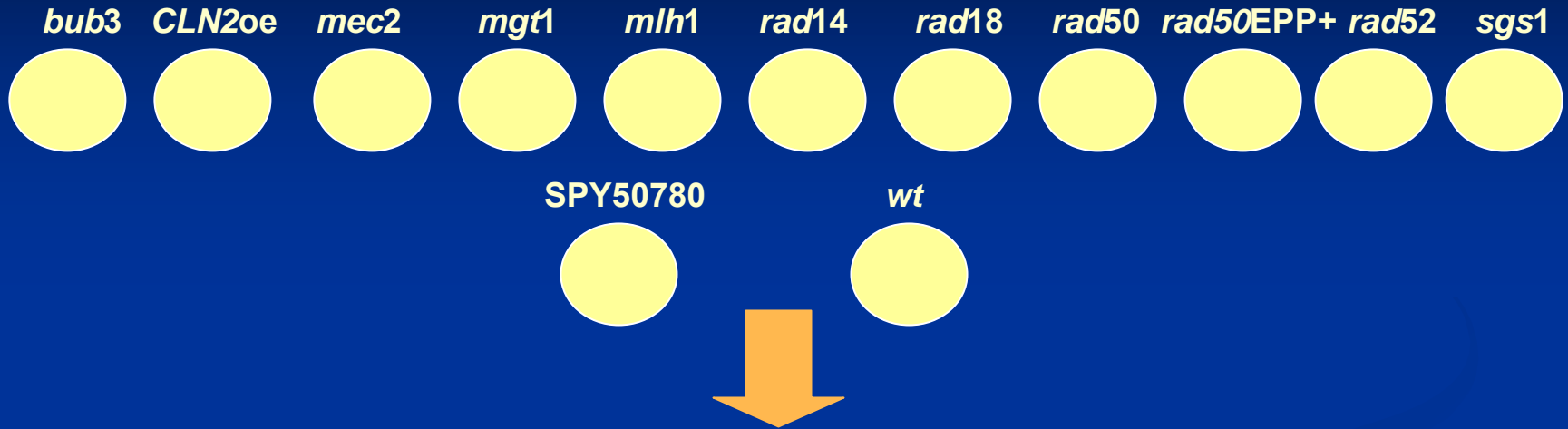


**Hormetic Dose-Response**



# NCI Methods

## 13 Yeast Strains



**2,189 Anti-tumor Agents**  
(1.2, 3.7, 11, 33 & 100  $\mu$ M + Control)

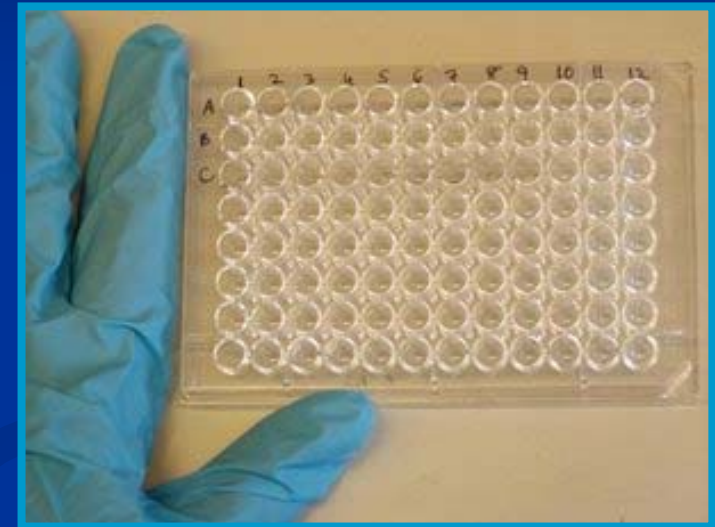


**28,457 Replicated Dose-Response Experiments**

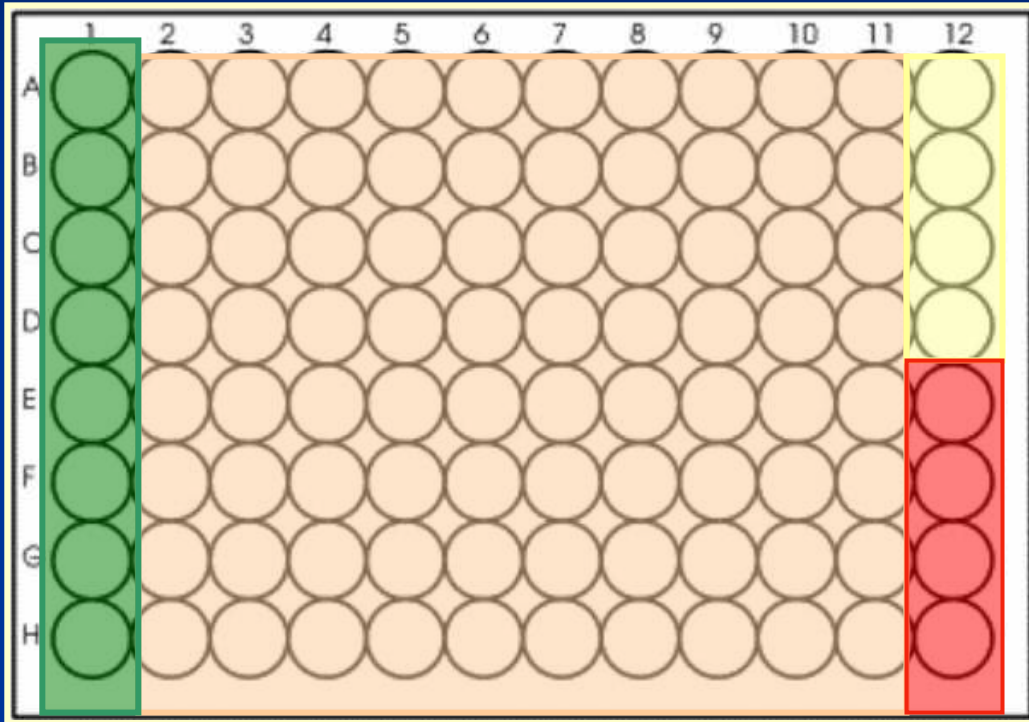






# Methods: Cell Line

- Yeast cells in exponential growth inoculated onto microtiter plates
- $10^4$  cells/well with 200 $\mu$ l of medium
- 80 different anti-tumor agents added per plate
- Each plate run at one concentration (1.2, 3.7, 10, 33, or 100  $\mu$ M)
- Each (of 5) conc. incubated over the same 12 hrs.
- Cell number measured with spectrophotometer ( $OD_{600}$ )



# Methods: 96-well Plate

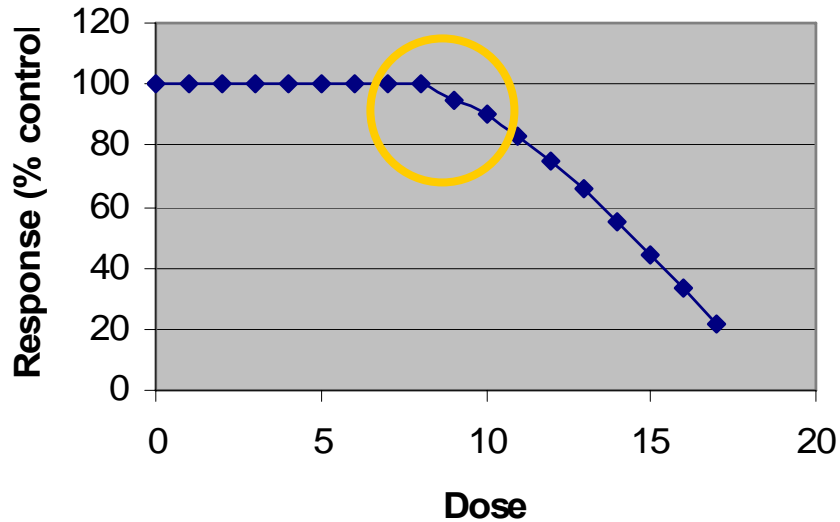


-  80 different chemicals at same concentration
-  8 wells solvent (DMSO) controls
-  4 wells cycloheximide controls
-  4 wells unexposed controls

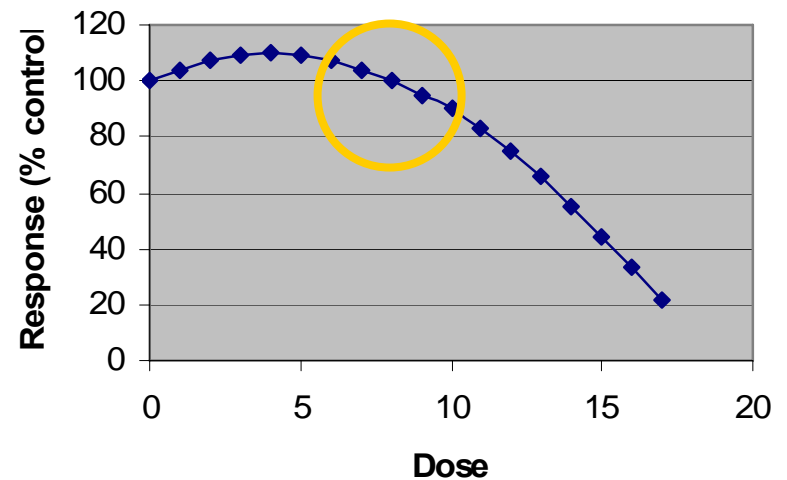
Reported response is the growth of yeast in respective concentration relative to growth in solvent control

# Where is the Low Dose Region?

## Threshold Dose-Response

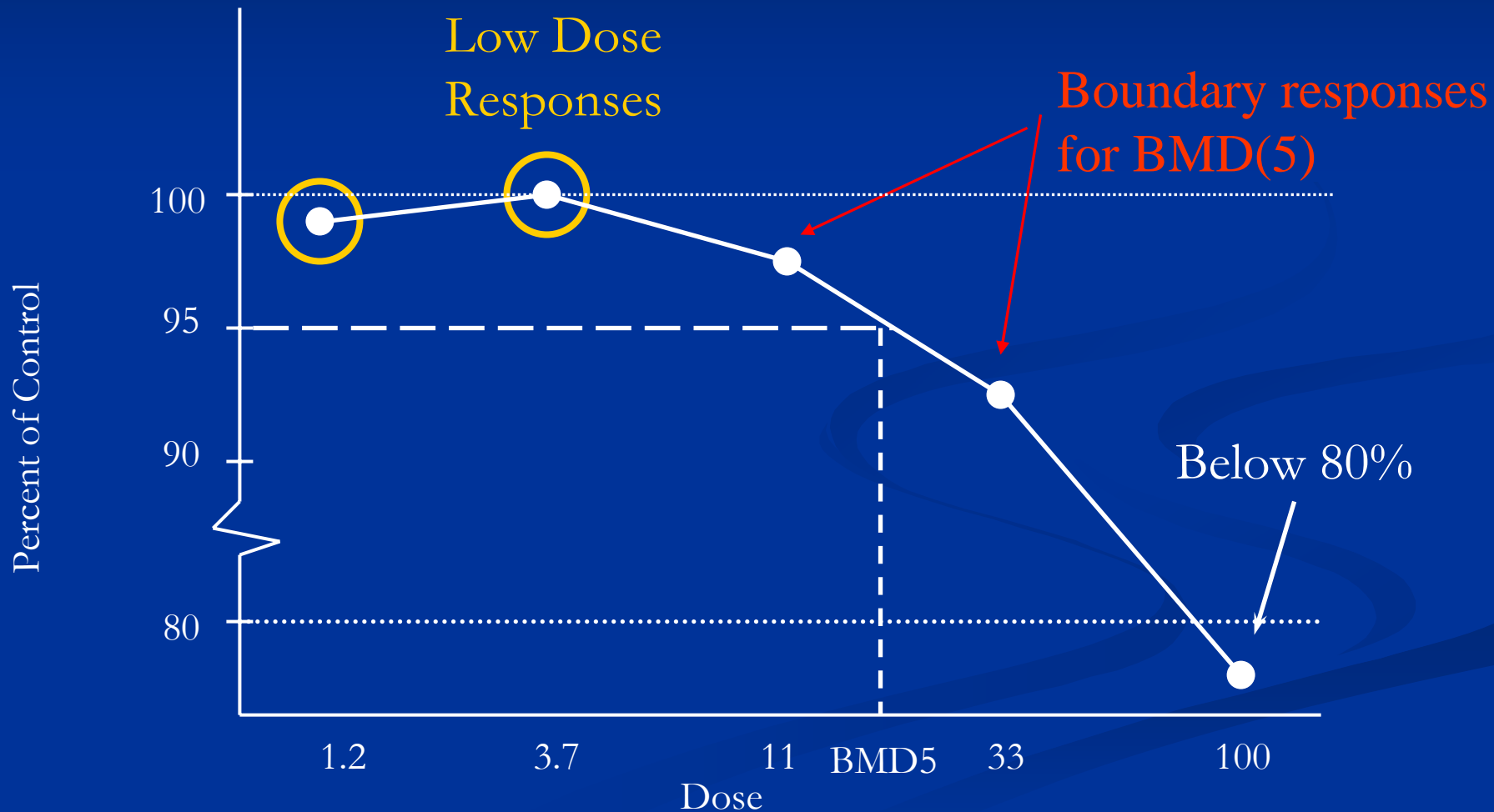


## Hormetic Dose-Response





# Figure 1. General Scheme Used for the Derivation of the BMD(5)



**START**

# Selection Process

**28,457** dose-responses included in the database:  
2,189 anticancer agents x 13 yeast strains

├ 6,480 removed (23%)  
↓

**21,977** (77%) have an inhibitory response  $\leq 80$  of the control

├ 9,375 removed (43%)  
↓

**12,602** (44%) also have a dose below the BMD(5)

├ 10,151 removed  
↓

**2,451** (9%) also have 3 doses below the BMD(5)

**FINISH**

**253** Wild Type (wt)

# Example Data

## Wild Type- NSC#1928

Dose	Ave Response	Difference
1.2	97	0
3.7	98	12
11	91	15

# Example Data

## Wild Type- NSC#1928

Dose	Rep	Ave Response	Difference	Response (% Diff from 100)
1.2	1	97	0	-3
1.2	2	97	0	-3
3.7	1	98	12	4
3.7	2	98	12	-8
11	1	91	15	-1.5
11	2	91	15	-16.5

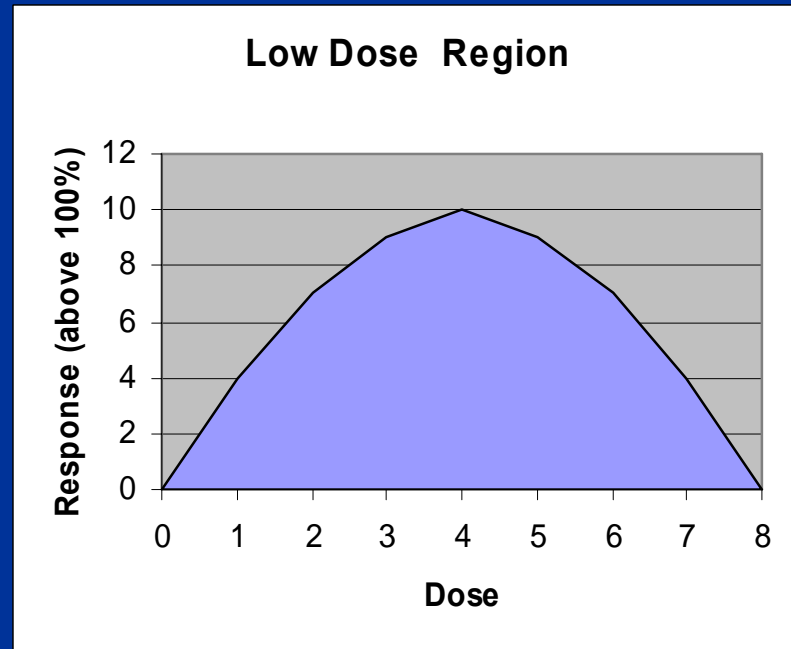
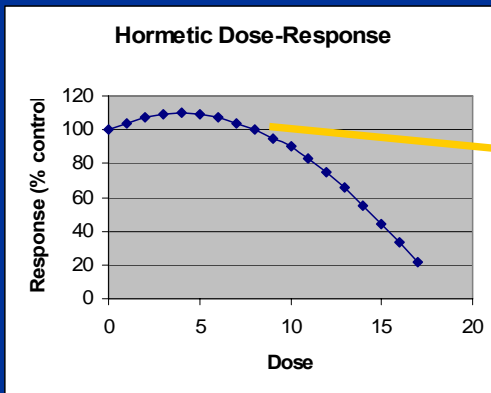
# How should we evaluate Response in Low Dose Region?

## Data:

Selected Set of chemicals (n=253)

b) For each chemical, 3 doses

c) For each dose, replicate responses



Approach 1: Test a Hypothesis

Approach 2: Estimate ‘True Response’

Need a statistical Model: Response

Chemical *s* is  
“NSC#1928”



Dose (t)	Rep (k)	$Y_{stk}$
1	1	-3
1	2	-3
2	1	4
2	2	-8
3	1	-1.5
3	2	-16.5

# What is 'True Response'?

Expected Response  
Response

$$Y_{stk} = \mu_{st} + E_{stk}$$

$$E_R(Y_{stk}) = \mu_{st}$$

True Response:

$$\mu_s = \frac{1}{m} \sum_{t=1}^m \mu_{st}$$

Dose (t)	Rep (k)	$Y_{stk}$	$\mu_{st}$
1	1	-3	$\mu_{s1}$
1	2	-3	$\mu_{s1}$
2	1	4	$\mu_{s2}$
2	2	-8	$\mu_{s2}$
3	1	-1.5	$\mu_{s3}$
3	2	-16.5	$\mu_{s3}$

# Model of Response for Chemical “s”

Chemical  $s = 1, \dots, n$

Dose  $t = 1, \dots, m$

Replication  $k = 1, \dots, r (= 2)$

where

$$\begin{aligned} Y_{stk} &= \mu_{st} + E_{stk} \\ &= \mu_s + \delta_{st} + E_{stk} \end{aligned}$$

$$\mu_s = \frac{1}{m} \sum_{t=1}^m \mu_{st}$$



# Hypothesis Testing

Model: 
$$Y_{stk} = \mu_s + \delta_{st} + E_{stk}$$

Test the Null Hypothesis:  $H_0 : \mu_s = 0$

$$H_A : \mu_s \neq 0$$

“NSC#1928”  $\hat{\mu}_s = -4.67$

$$se(\hat{\mu}_s) = 3.20$$

$$t = -1.46$$

$$p - value = 0.241$$

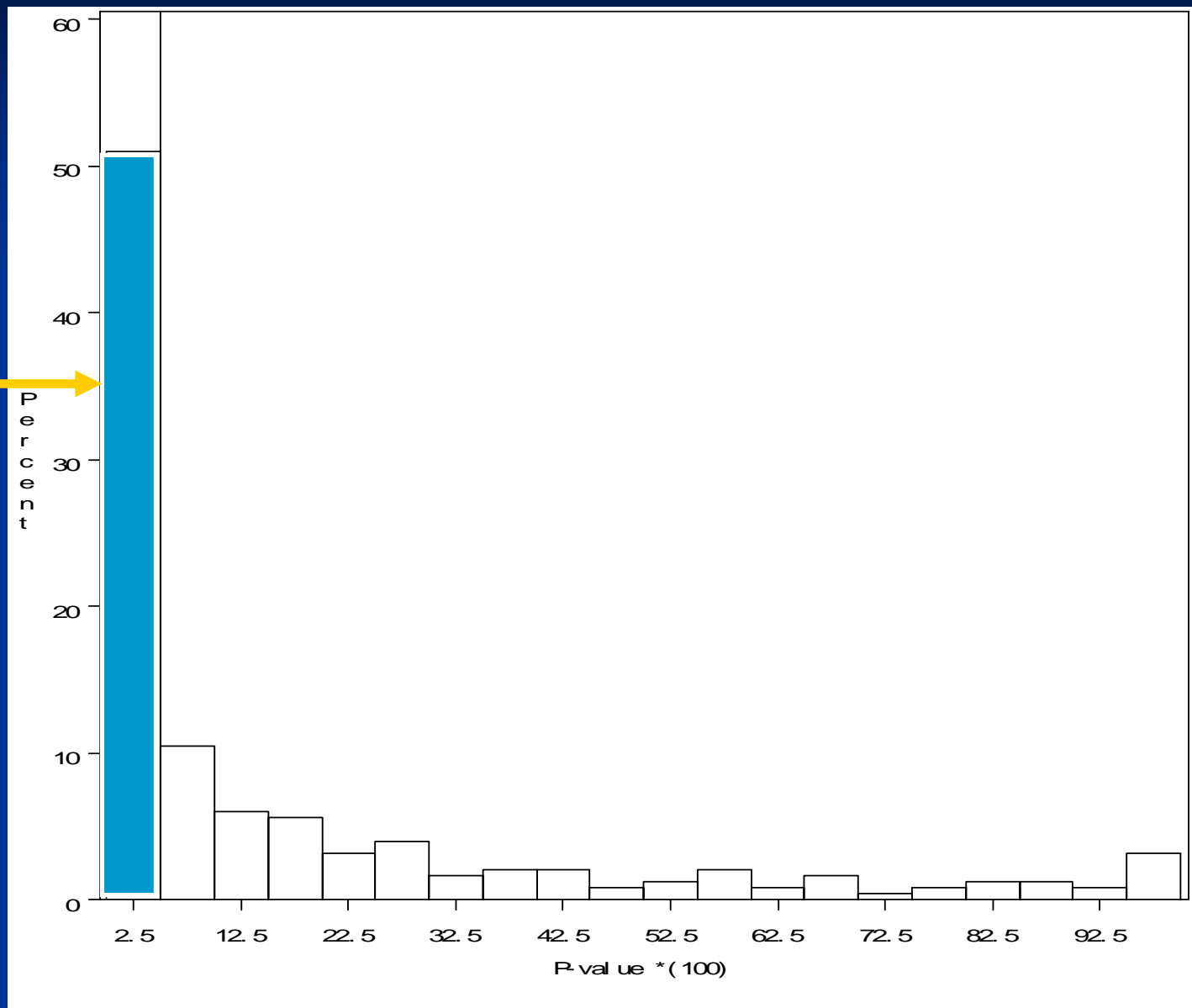
Decision: Fail to  
reject  $H_0$

# Histogram of P-values for 249 Chemicals:

P-value < 0.05  
127 Chemicals  
(50.8%)



Excluding 4  
Chemicals with  
“0” stderr



# Interpretation of Tests

## All 249 Chemicals

Expected # Significant by Chance: 12 to 13

Observed 127 Significant: Mean Response

3 Negative

124 Positive

If 1 Sided test:  $H_0 : \mu_s \leq 0$

$H_A : \mu_s > 0$

P-value < 0.05

146 Chemicals

(58.6%)

# Multiple Comparisons?

Preserve Type I Error:

Prob(False Rejection of  $H_0$ )

Change the critical value to  $\alpha = \frac{0.05}{249} = 0.0002$

P-value  $< 0.0002$

11 Chemicals (4.4%)

Mean response:

0 Negative

11 Positive

# Conclusions:

- Studies are not designed for low-dose range
- Low power to evaluate response
- If controlling for multiple comparisons, few chemicals reject null hypothesis (4.4%)
- Data are not suitable to learn about response at low doses

## Approach 2: Estimate Response at Low Doses

Fit the model to chemical “s”:

$$Y_{stk} = \mu_s + \delta_{st} + E_{stk}$$

Estimate  $\mu_s$

Estimator

$$\bar{Y}_s = \frac{1}{m} \sum_{t=1}^m \left( \frac{1}{r} \sum_{k=1}^r Y_{stk} \right)$$

Estimate a 95% Confidence Interval for  $\mu_s$

# Example: Wild Type

Dose (t)	Rep (k)	$Y_{stk}$
1	1	-3
1	2	-3
2	1	4
2	2	-8
3	1	-1.5
3	2	-16.5

Response

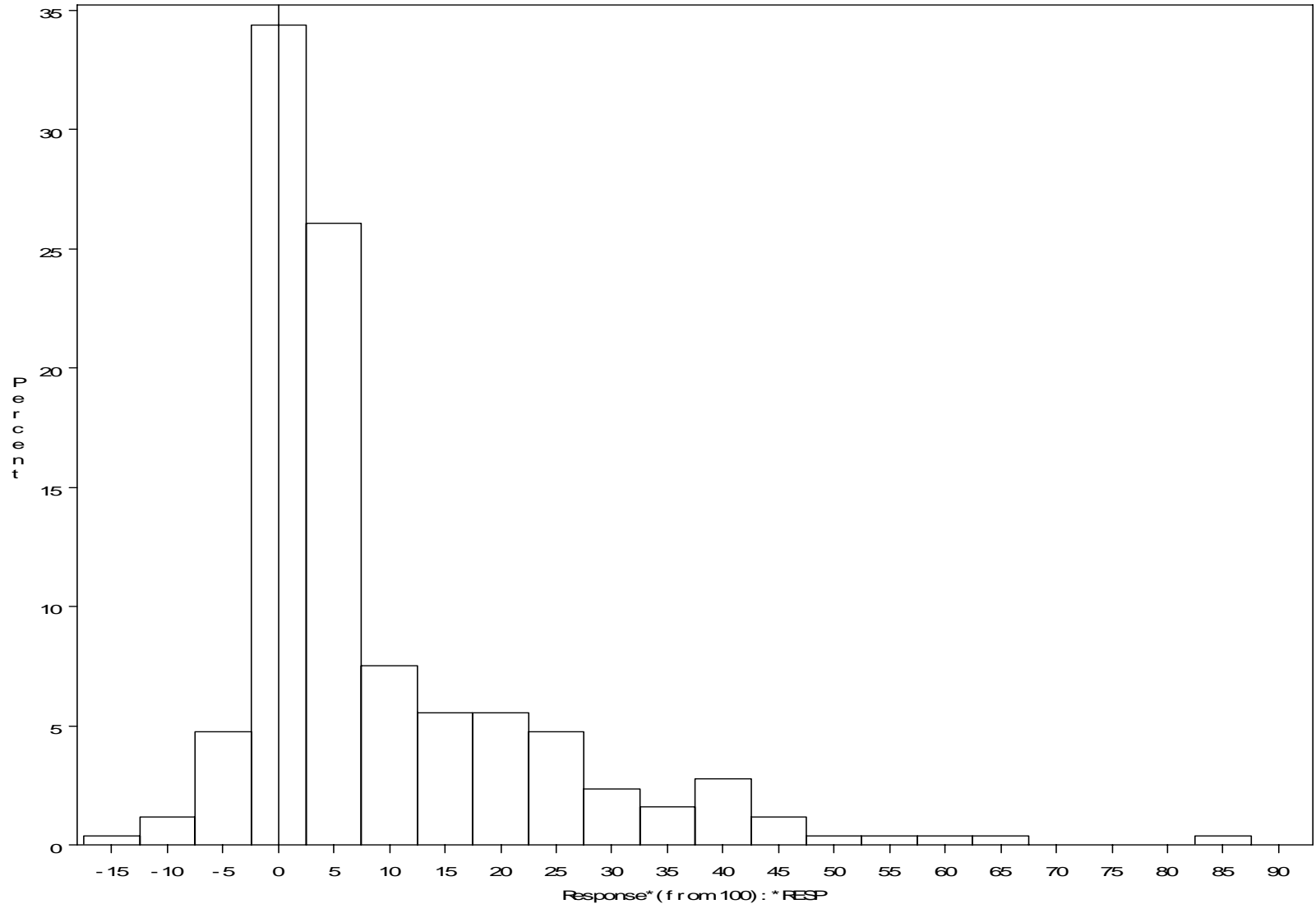
“NSC#1928”

$$\bar{Y}_s = -4.67$$

$$se(\bar{Y}_s) = 3.20$$

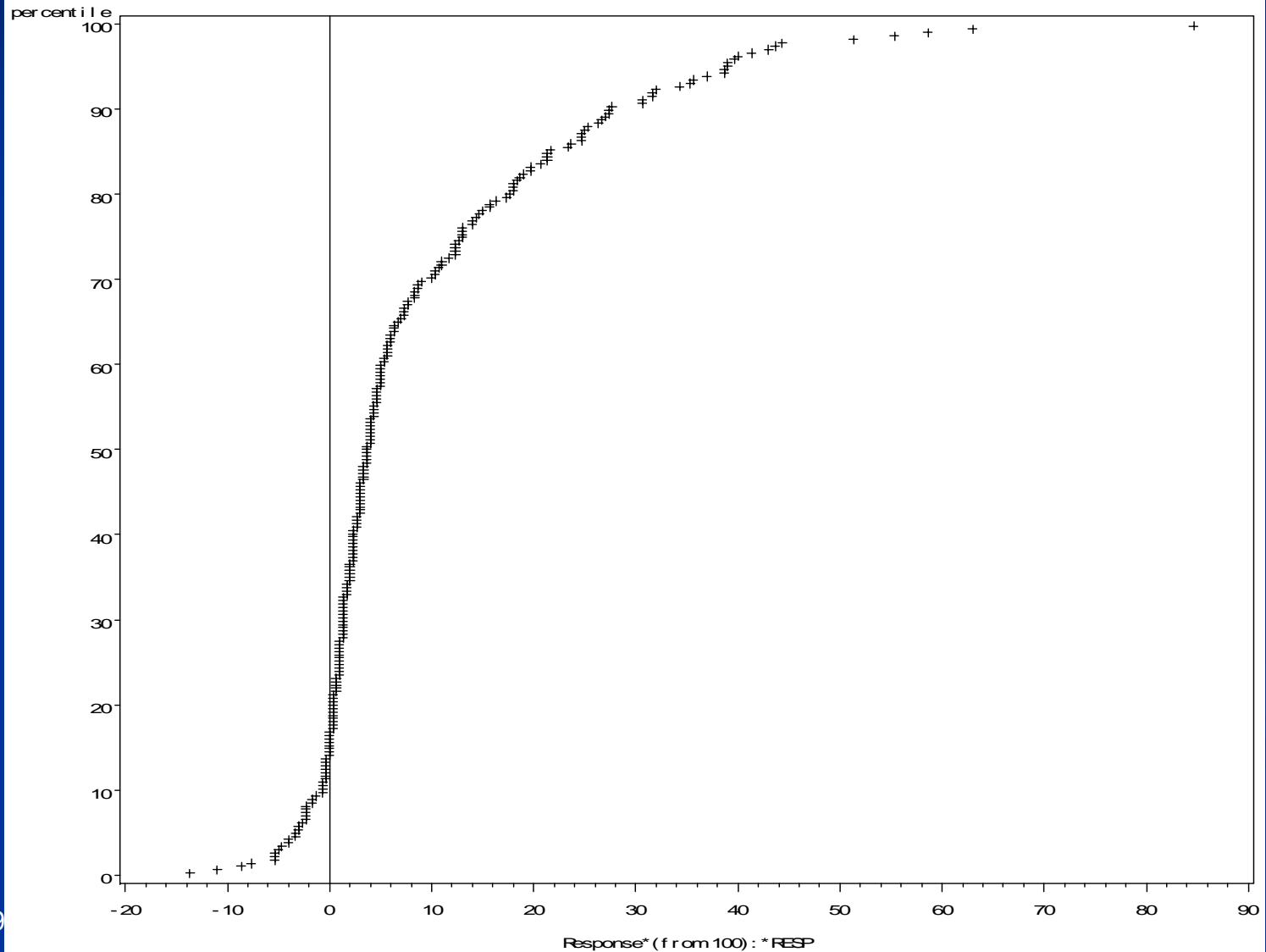
$$\left( \bar{Y}_s - t_{df, \left(1 - \frac{\alpha}{2}\right)} se(\bar{Y}_s), \bar{Y}_s + t_{df, \left(1 - \frac{\alpha}{2}\right)} se(\bar{Y}_s) \right) \quad (-14.9, 5.5)$$

# Histogram of Estimates (n=253 chemicals)





# Cumulative Distribution of Estimates (n=253 chemicals)



# What about Control for Multiple Chemicals?

$$\left( \bar{Y}_s - t_{df, \left(1 - \frac{\alpha}{2}\right)} se(\bar{Y}_s), \bar{Y}_s + t_{df, \left(1 - \frac{\alpha}{2}\right)} se(\bar{Y}_s) \right) \quad (-14.9, 5.5)$$

Should we use  $\alpha = \frac{0.05}{253} = 0.0002$

Using  $t_{3, (0.0001)} = -22.2 \quad (-75.8, 66.4)$

Conclusion: CI is so large, the estimate seems useless.

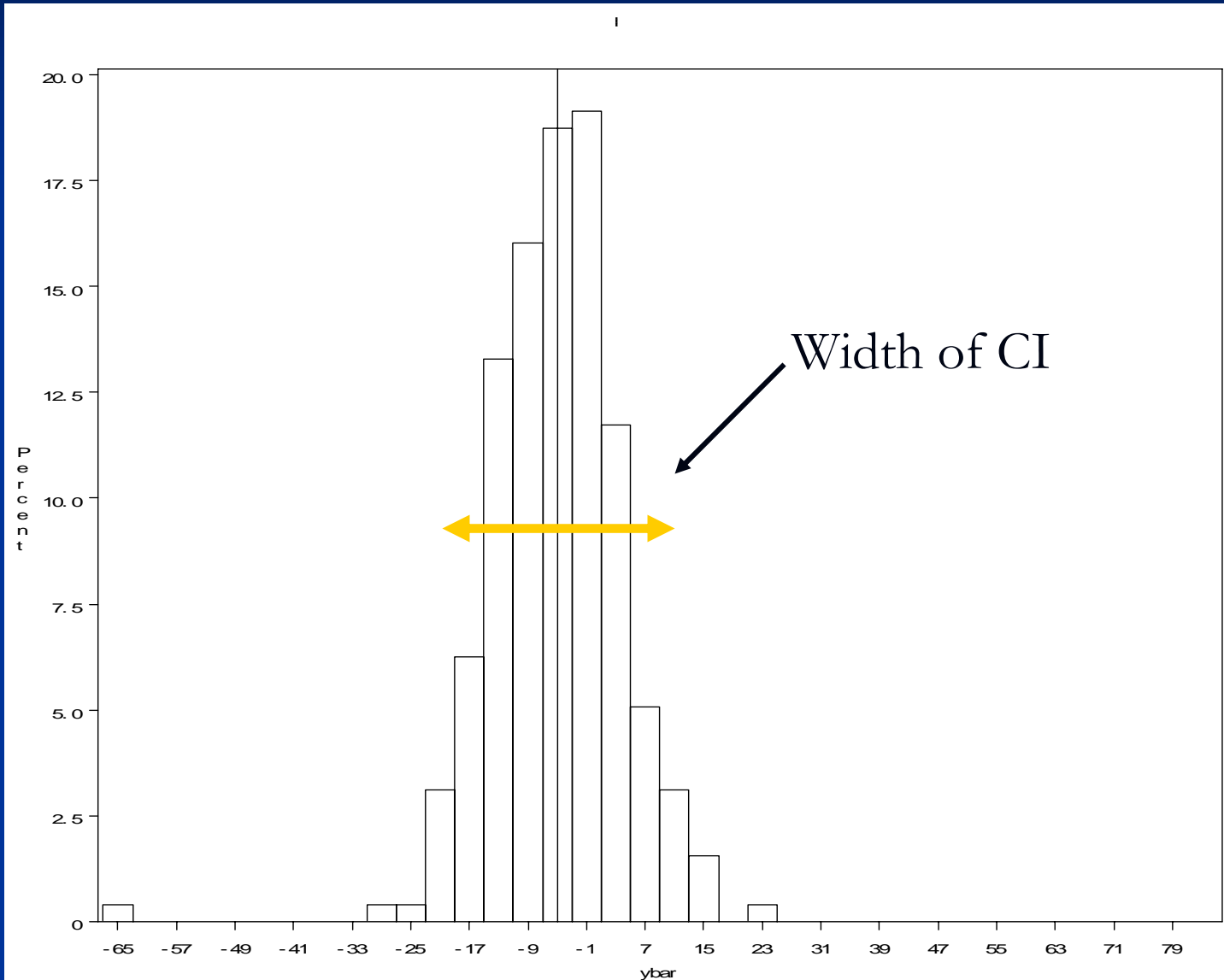
# Review: What is a Confidence Interval?

Assume  $\mu_s = -5.0$   
 $se(\bar{Y}_s) = 8.0$

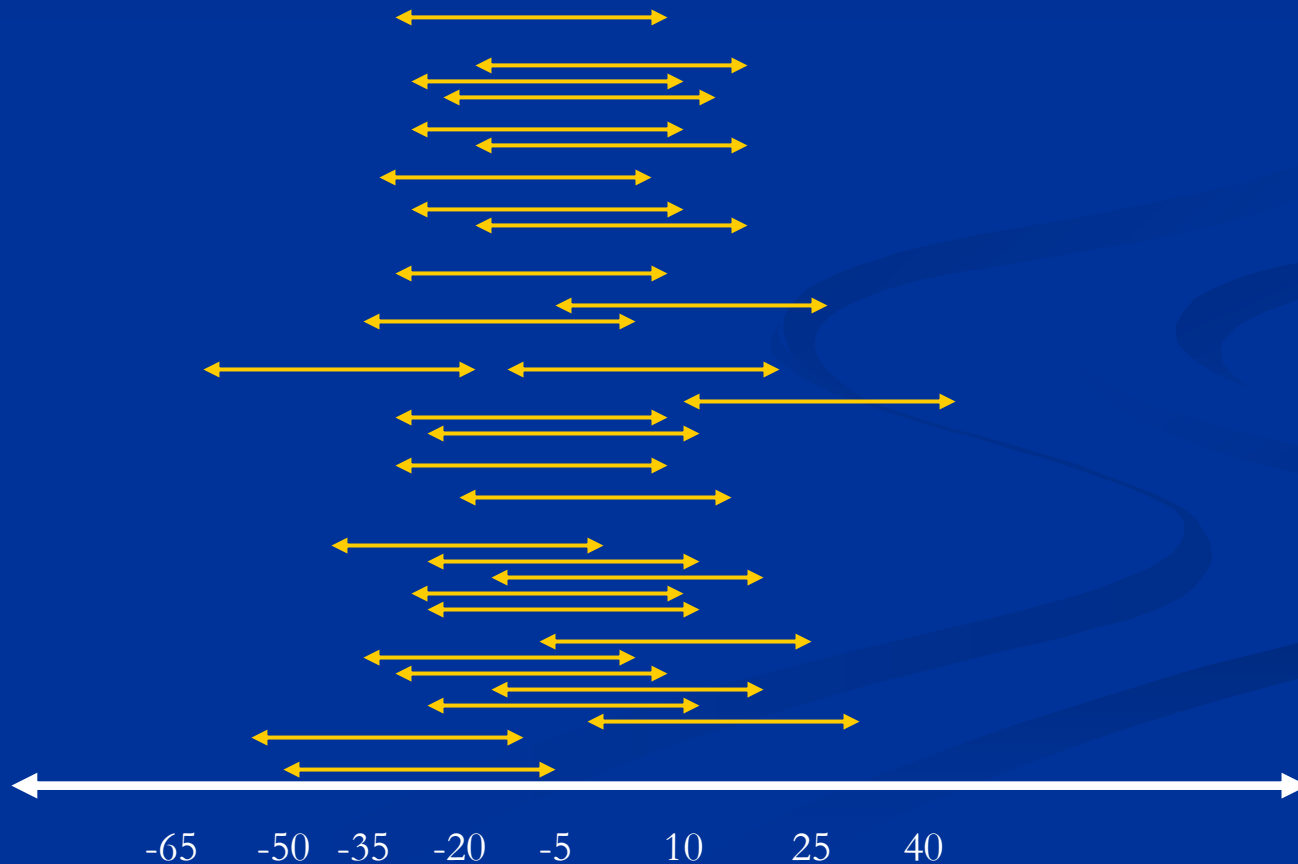
Suppose we repeat the study  
many times and summarize  
the Estimator

$$\bar{Y}_s$$

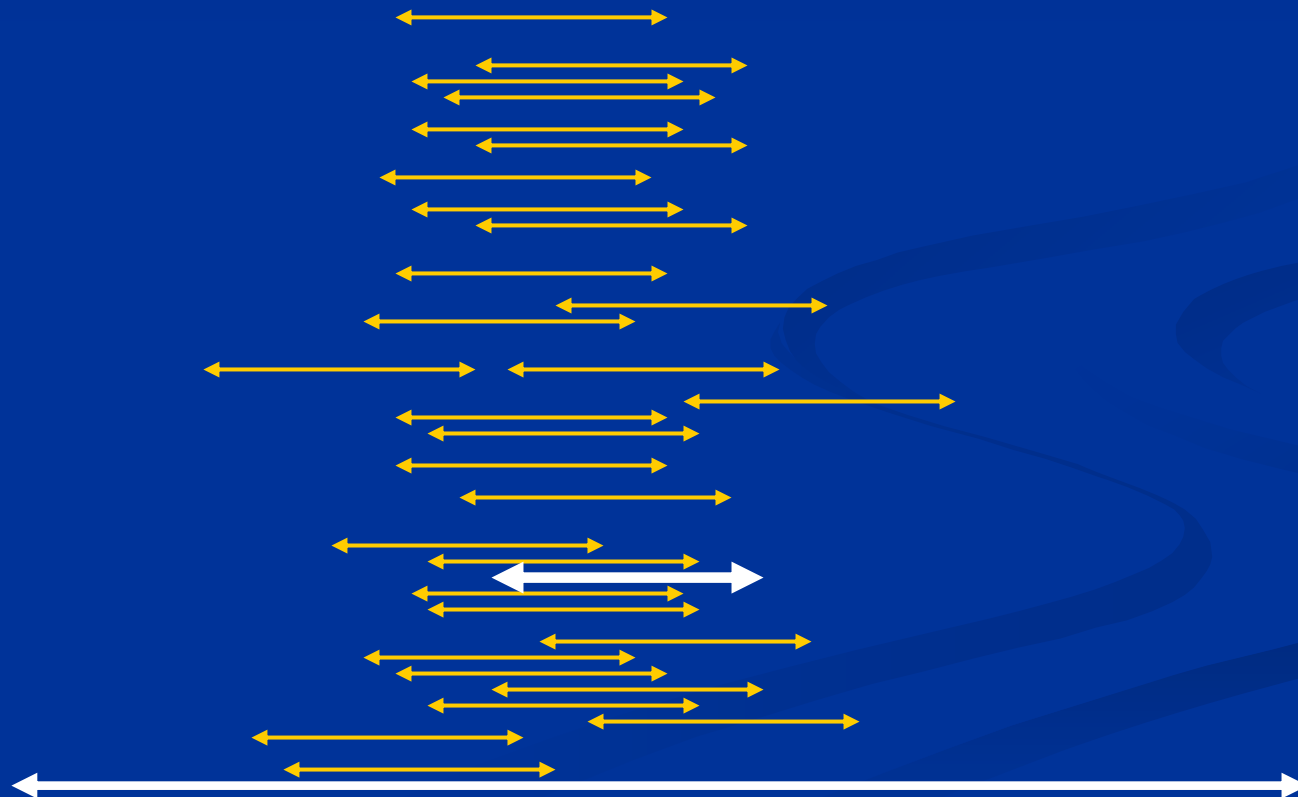
# Distribution of Estimator



# Distribution of CI Estimator



# Observed Confidence Interval



# Observed Confidence Interval

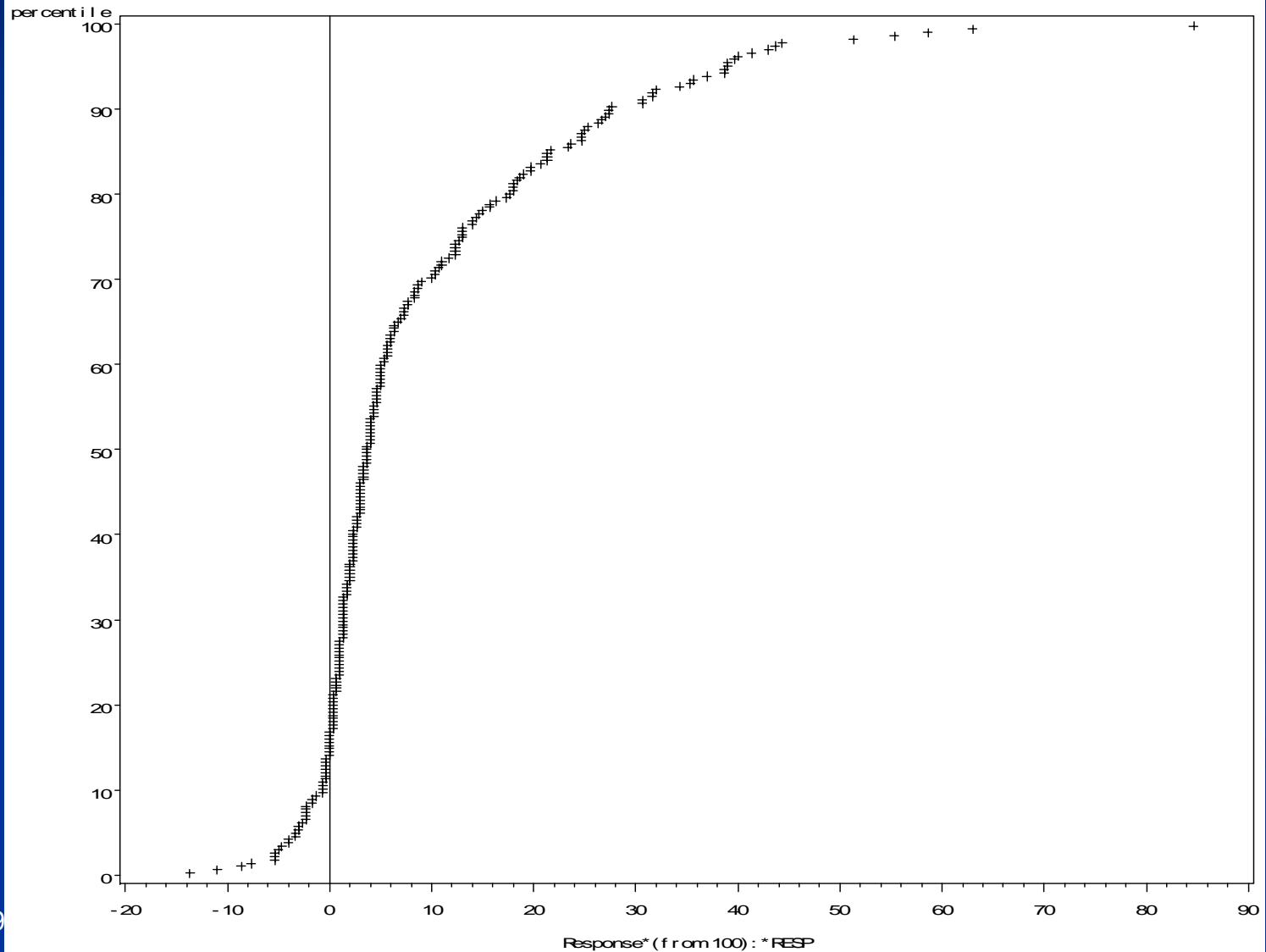


# Conclusion:

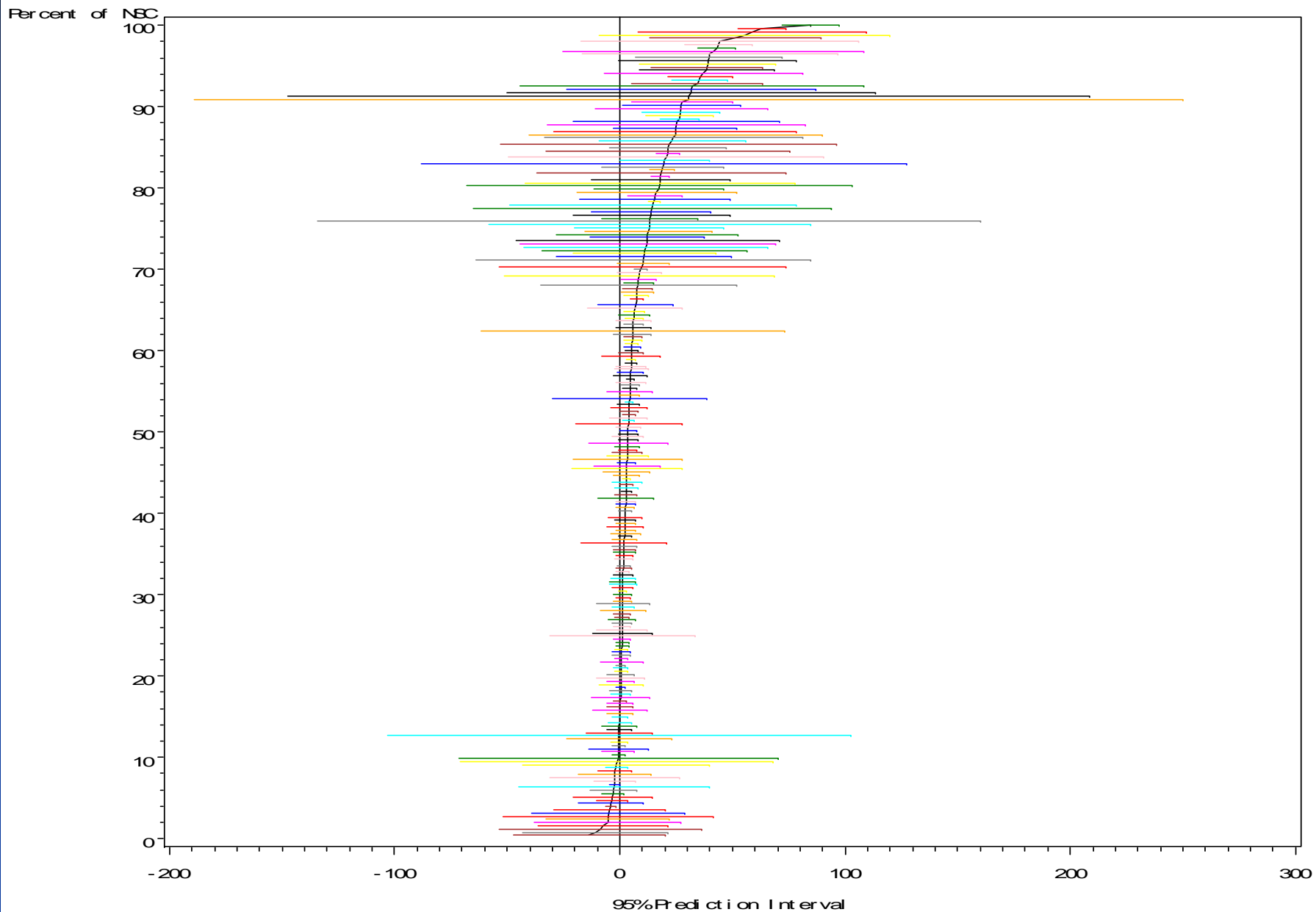
- Confidence Intervals indicate the ‘central’ width of the distribution of the Estimator
- The ‘width’ doesn’t change when Estimating different chemicals
- No adjustment for the length of confidence intervals is needed.
- Different from Hypothesis testing.



# Cumulative Distribution of Estimates (n=253 chemicals)



# Wild Type: Confidence Intervals by Chemical with 3 Doses below BMD(5)(n= 253)



# What would we expect if Threshold Model was the True Model?

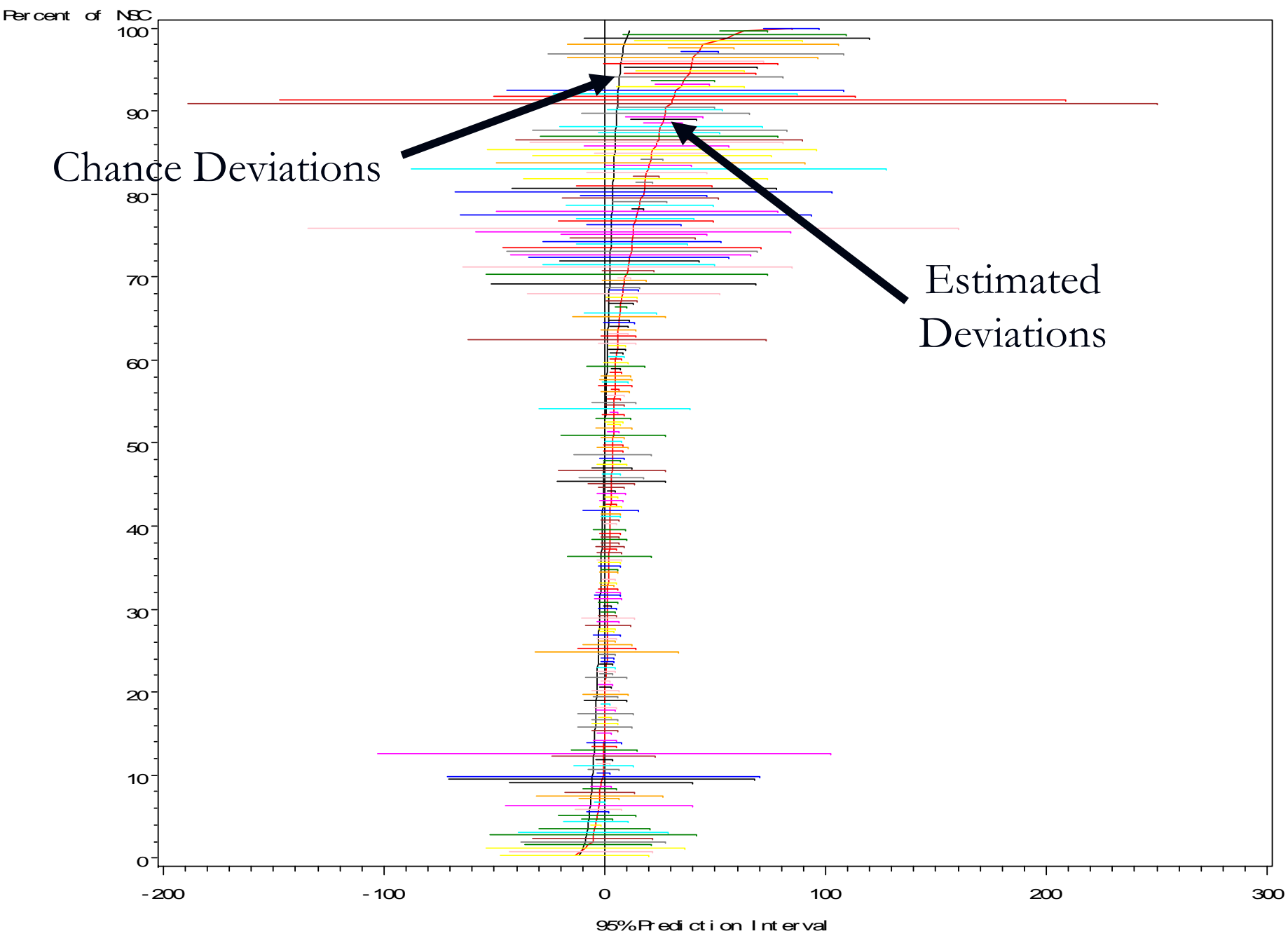
- Pool Estimates of the Response Error Variance across Chemicals

$$\hat{\sigma}_e = 10.49$$

Use the estimate to estimate the standard error of the estimator

$$se(\bar{Y}) = \frac{\hat{\sigma}_e}{\sqrt{n}} = 4.28$$

# Wild Type: Confidence Intervals by Chemical with 3 Doses below BMD(5) (n= 253)



# Can we improve the estimates?

- Account for higher Response Error at extremes.
- Improve Accuracy of estimates-

YES:

- Motivate Estimator by Considering Study Chemicals to have arisen from a sample

Mixed model

# Model of Response for Chemical “s”

Chemical  $s = 1, \dots, n$

Dose  $t = 1, \dots, m$

Replication  $k = 1, \dots, r (= 2)$

$$Y_{stk} = \mu_{st} + E_{stk}$$
$$= \mu_s + \delta_{st} + E_{stk}$$

Define a Population of  
Chemicals  $s = 1, \dots, N$

Define Mean  
for Population

$$\mu = \sum_{s=1}^N \mu_s$$

↑  
Mean for  
Chemical “s”

↓

$$Y_{stk} = (\mu + \beta_s) + \delta_{st} + E_{stk}$$

# Consider Study Chemicals as a sample

Population of Chemicals  $s = 1, \dots, N$

Sample of Chemicals  $i = 1, \dots, n$

For Chemical “s”

$$Y_{stk} = (\mu + \beta_s) + \delta_{st} + E_{stk}$$

For the  $i^{\text{th}}$  Selected  
Chemical

$$Y_{itk} = (\mu + B_i) + D_{it} + E_{itk}$$

Fixed Effects

Random  
Effects

# Summary of Mixed Model

$$Y_{ijk} = (\mu + B_i) + E_{ijk}$$

↑  
Fixed

↙ ↘  
Random

$i$  = chemical

$J$  = dose

$k$  = replication



# Best Linear Unbiased Predictor (BLUP) from Mixed Model

$$\mu + B_i$$

Latent Response of ith Selected  
Chemical

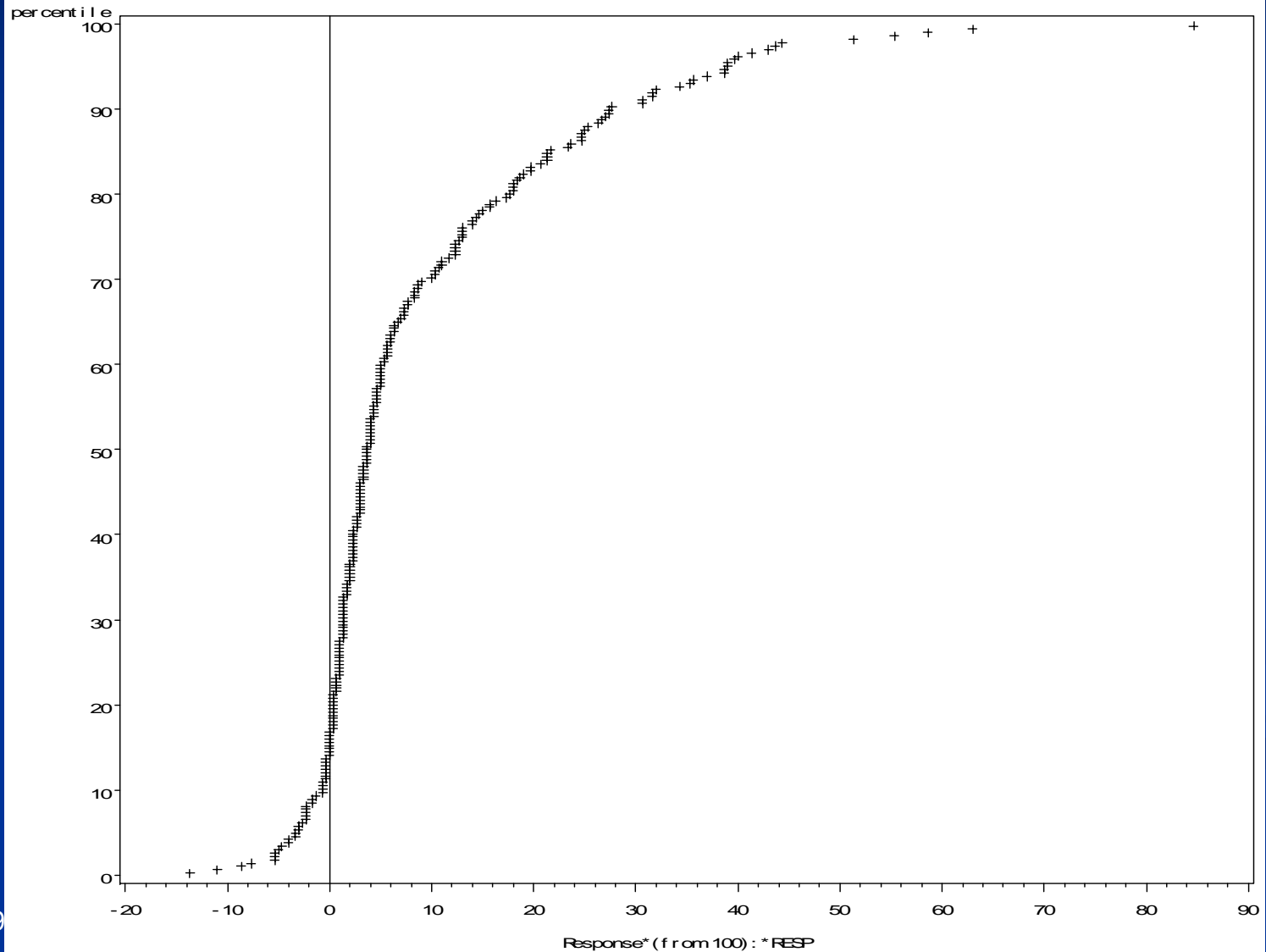
**Predictor**

$$\hat{P}_i = \hat{\mu} + k \left( \bar{Y}_i - \hat{\mu} \right)$$

**Shrinkage Constant**

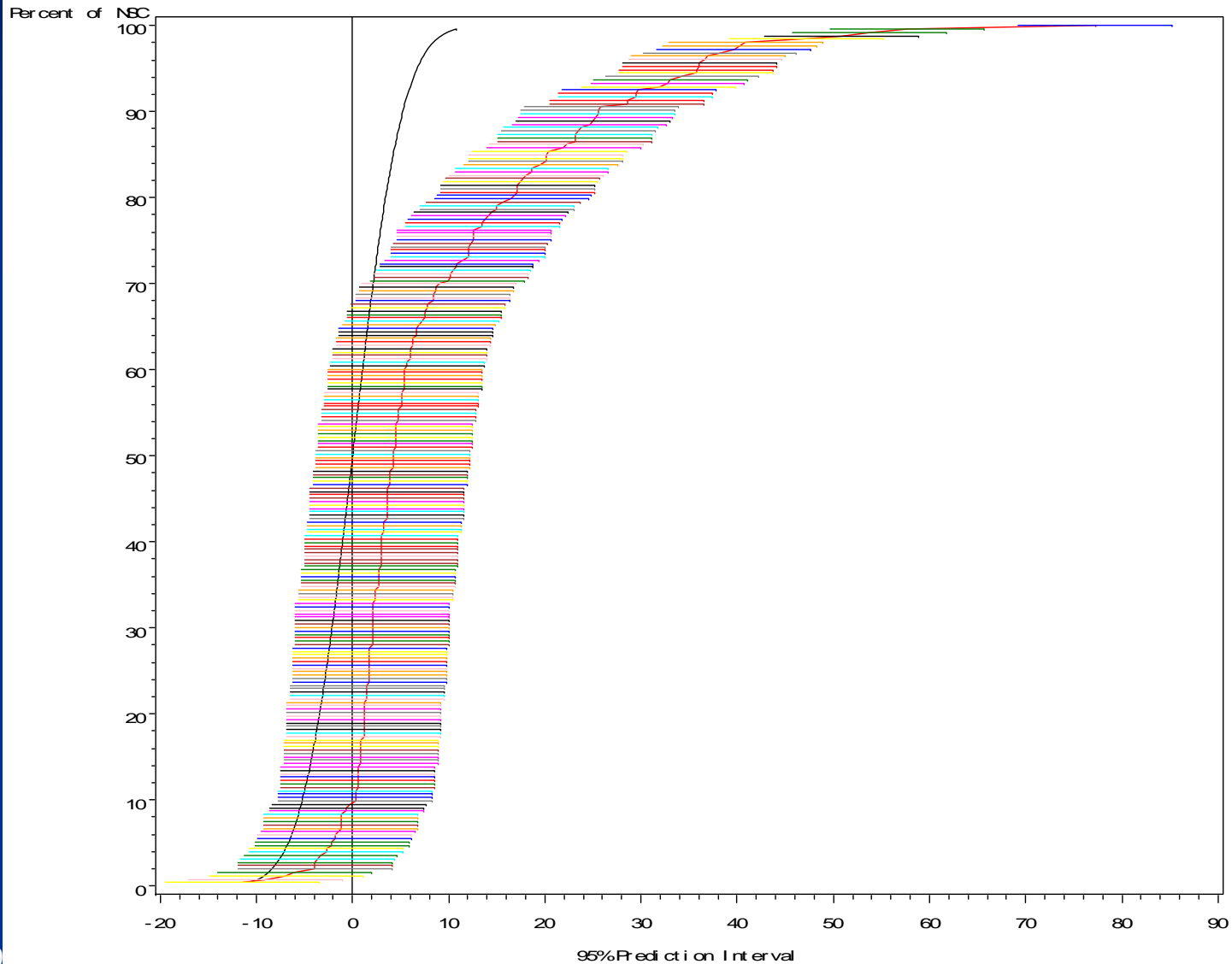
$$k = \frac{\sigma^2}{\sigma^2 + \sigma_e^2 / m}$$

# Cumulative Distribution of Simple Mean (n=253 chemicals)



# Cumulative Distribution of BLUP Estimates (n=253 chemicals)

Figure 1a. Strain=Wt in BMD(5) Range with 3 Responses (n= 253)



# Conclusions

- We can learn from data- even if it isn't an experiment.
- Estimation is the key- Hypothesis testing may lead us astray.
- More accurate estimates are possible with Mixed models
  - Require minimal additional assumptions
  - Dampen some of the response error
- Predictors are related to Bayesian methods

# Thanks

■ Questions?