

# **Directly testing the linearity assumption for assay validation**

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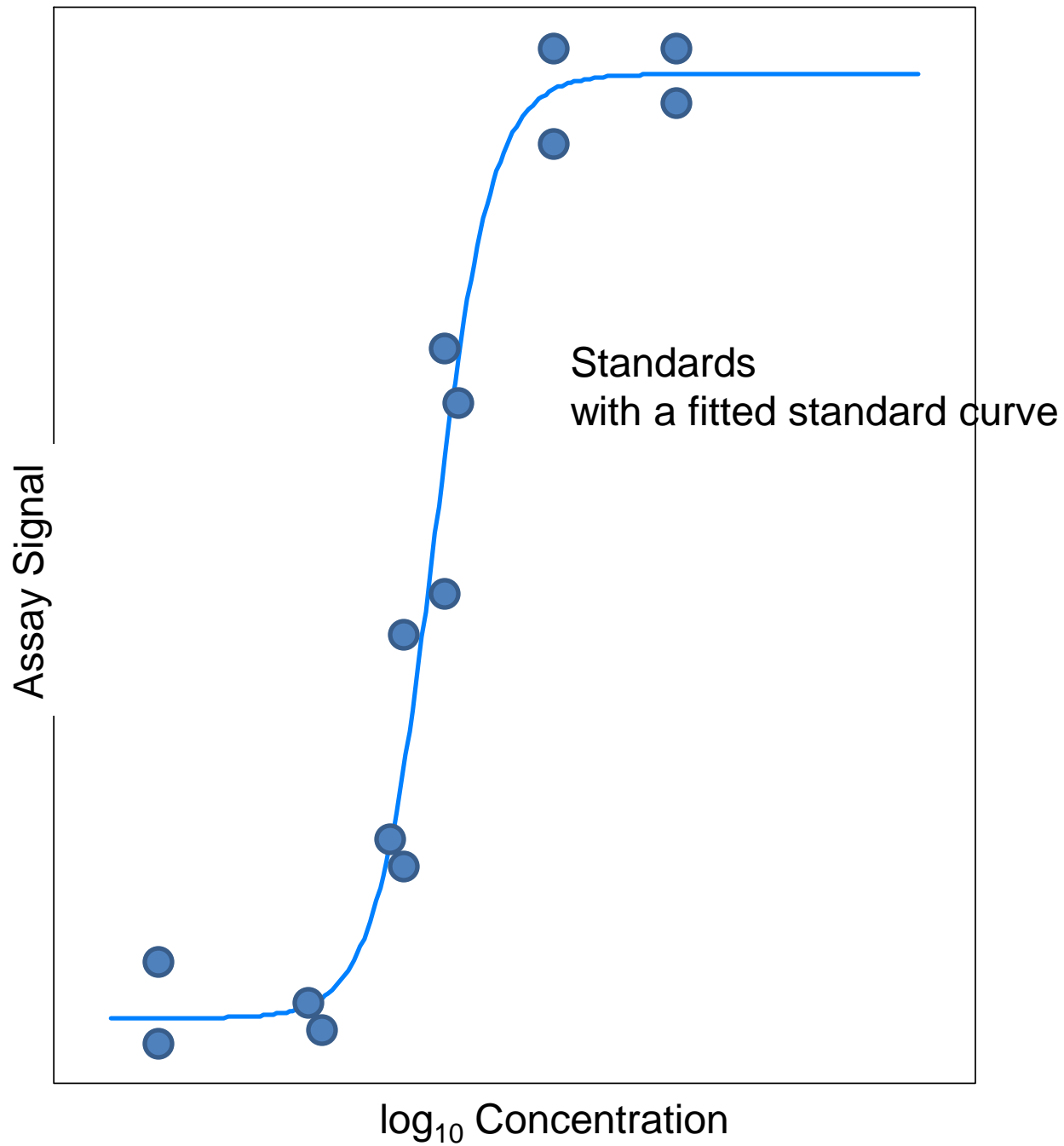
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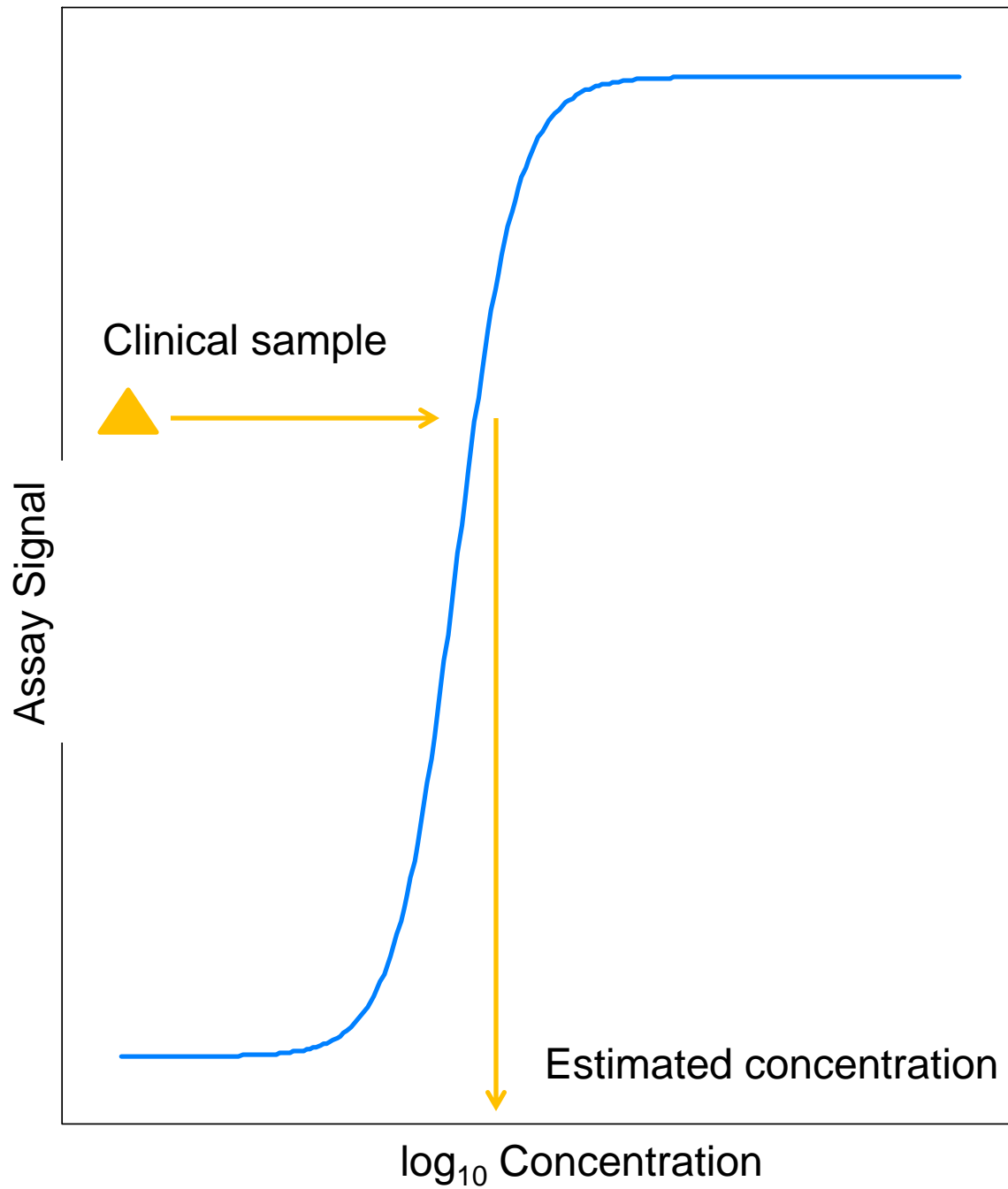
# Purpose

- Illustrate a novel method to test for linearity in an analytical assay.

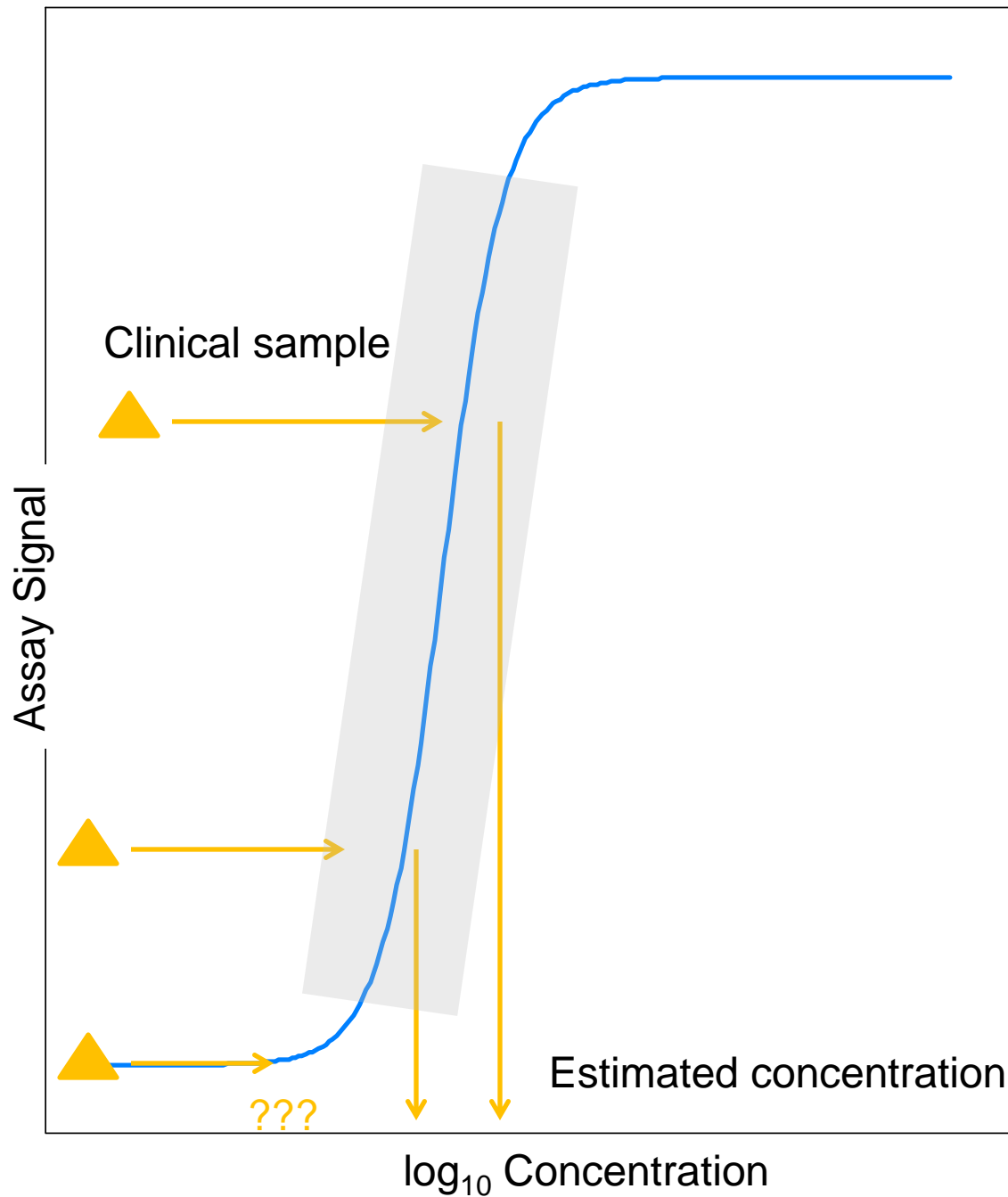
# Analytical assay – standard curve

- We wish to measure the concentration of an analyte (e.g., a protein) in clinical sample.
- Standards = known concentrations of an analyte.
- To estimate the concentration, we create a standard curve.





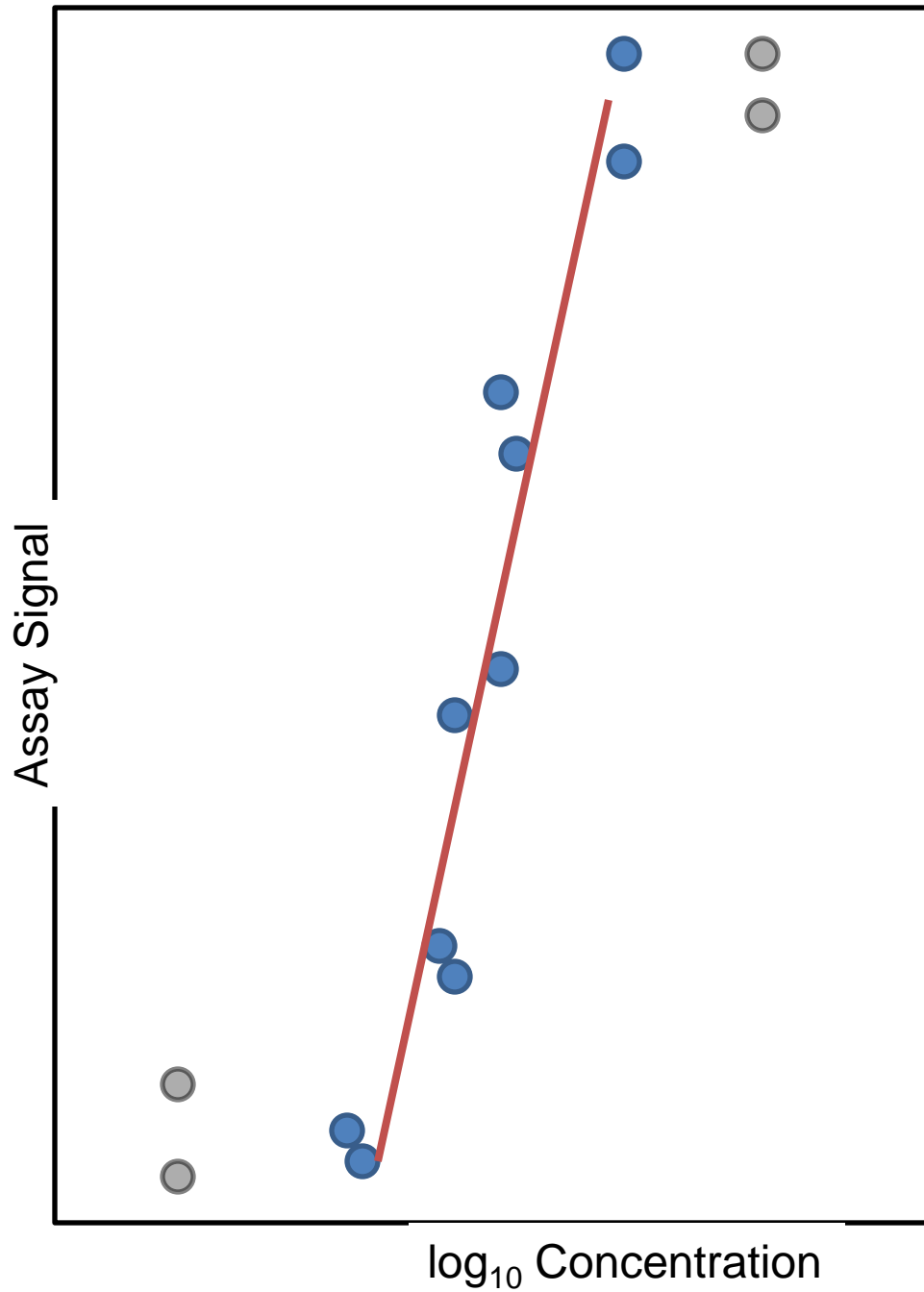
To many, the only interest lies in the linear portion of the curve.



# ICH Q2(R1) guideline

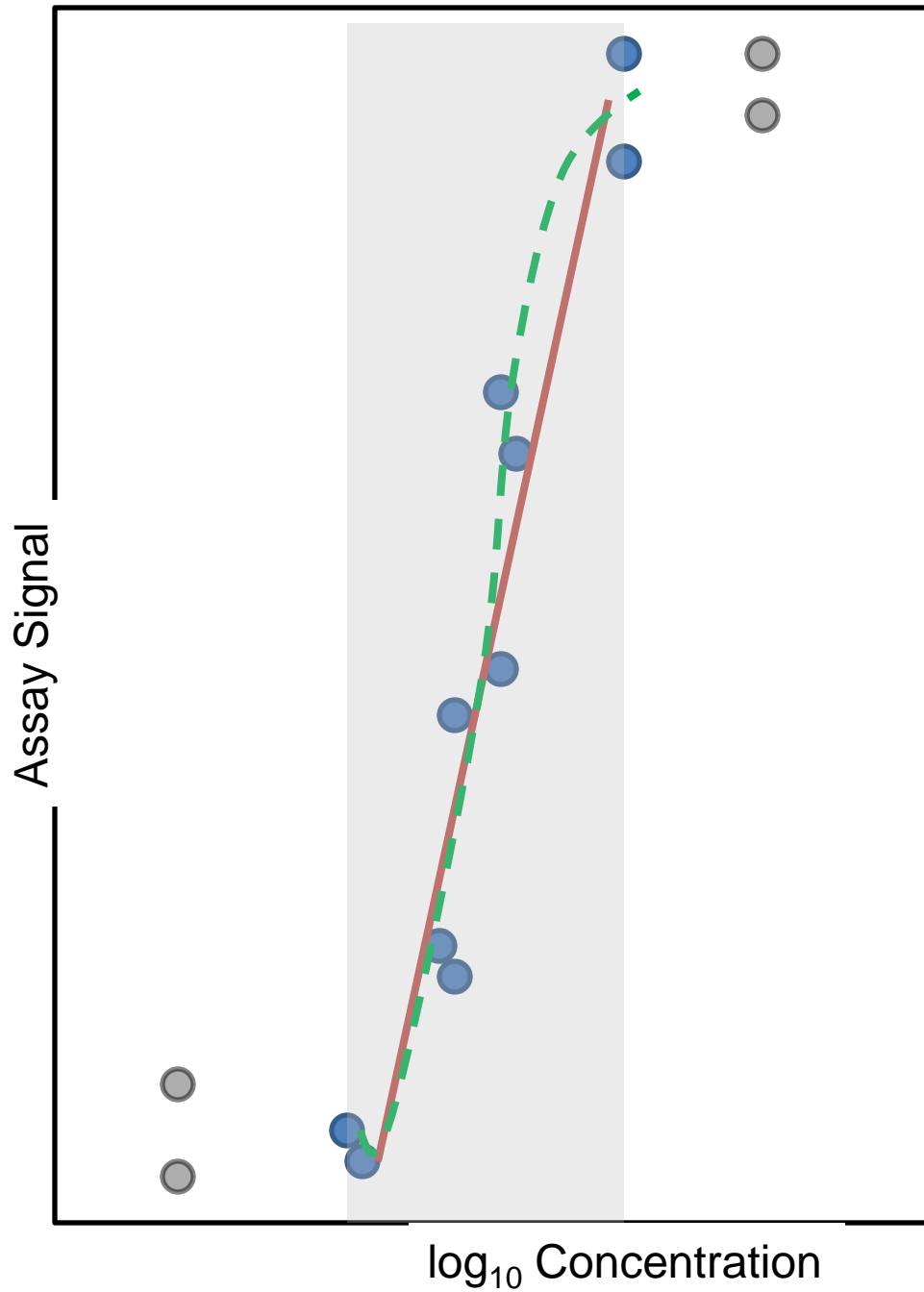
- [http://www.ich.org/fileadmin/Public\\_Web\\_Site/ICH\\_Products/Guidelines/Quality/Q2\\_R1/Step4/Q2\\_R1\\_Guideline.pdf](http://www.ich.org/fileadmin/Public_Web_Site/ICH_Products/Guidelines/Quality/Q2_R1/Step4/Q2_R1_Guideline.pdf)
- Evaluate linearity by visual inspection





# The EP6-A guidelines

- Clinical and Laboratory Standards Institute
  - <http://www.clsi.org/source/orders/free/ep6-a.pdf>
- Compare straight-line to higher-order polynomial curve fits
  - Recommendation: Test higher-order coefficients.



# Notation

$$E[ Y_{ij} | X_i ] = a_1 + b_1 X_i = g_1(X_i)$$

$$E[ Y_{ij} | X_i ] = a_2 + b_2 X_i + c_2 X_i^2 = g_2(X_i)$$

$$E[ Y_{ij} | X_i ] = a_3 + b_3 X_i + c_3 X_i^2 + d_3 X_i^3 = g_3(X_i)$$

- Assume IID normally distributed errors with equal variance.

$i = 1, 2, \dots, L$  concentrations

$j = 1, 2, \dots, n_i$  replicates

# Orthogonal polynomials

- The orthogonal polynomial of degree  $k$  is of the form

$$\sum_{r=0}^k \theta_r f_r(x),$$

$\theta_r = (k+1)$  constants (to be estimated)

$f_r(x) = (k+1)$  orthogonal polynomials

$x =$  concentration

# Orthogonal polynomials properties

$$\sum_{i=1}^L n_i f_r(X_i) = 0$$

$$\sum_{i=1}^L n_i f_r(X_i) f_s(X_i) = 0$$

$$\sum_{i=1}^L n_i f_r^2(X_i) = 1 \quad \text{ortho-normal property}$$

# OLS: orthogonal polynomials

$$E[\mathbf{Y}] = \mathbf{F}\boldsymbol{\theta} \quad \text{Var}[\mathbf{Y}] = \sigma^2 \mathbf{I}_N, \quad N = \text{total sample size}$$

$$\mathbf{Y} = \begin{pmatrix} Y_{11} \\ Y_{12} \\ \vdots \\ Y_{1n_1} \\ \vdots \\ Y_{L1} \\ Y_{L2} \\ \vdots \\ Y_{Ln_L} \end{pmatrix}, \quad \mathbf{F} = \begin{pmatrix} f_0(X_1) & f_1(X_1) & \dots & f_k(X_1) \\ \vdots & \vdots & \dots & \vdots \\ f_0(X_{Ln_L}) & f_1(X_{Ln_L}) & \dots & f_k(X_{Ln_L}) \end{pmatrix}, \quad \boldsymbol{\theta} = \begin{pmatrix} \theta_0 \\ \theta_1 \\ \vdots \\ \theta_k \end{pmatrix}$$

$$\hat{\boldsymbol{\theta}} = \mathbf{F}^T \mathbf{Y} \quad \text{and} \quad \text{Var}[\hat{\boldsymbol{\theta}}] = \sigma^2 \mathbf{I}_{k+1}.$$

$$\widehat{\boldsymbol{\theta}}[\text{cubic}] = \begin{pmatrix} f_0(X_1) & f_0(X_2) & \cdots & f_0(X_{Ln_L}) \\ f_1(X_1) & f_1(X_2) & \cdots & f_1(X_{Ln_L}) \\ f_2(X_1) & f_2(X_2) & \cdots & f_2(X_{Ln_L}) \\ f_3(X_1) & f_3(X_2) & \cdots & f_3(X_{Ln_L}) \end{pmatrix} \mathbf{Y}$$

**F<sup>T</sup>**

$$\widehat{\boldsymbol{\theta}}[\text{linear}] = \begin{pmatrix} f_0(X_1) & f_0(X_2) & \cdots & f_0(X_{Ln_L}) \\ f_1(X_1) & f_1(X_2) & \cdots & f_1(X_{Ln_L}) \end{pmatrix} \mathbf{Y}$$



Intercept and slope estimates are same for both!



# Literature

- Krouwer and Schlain (1993)
  - Assume linearity, except at last concentration
  - Ha:  $\mu_{\max} - (a_1 + b_1 X_{\max}) \neq 0$  Assumes linearity?
  
- EP6-A
  - Ha: One or both of  $c_3, d_3 \neq 0$  Wrong power profile
  - Ha:  $|g_k(X_i) - g_1(X_i)| < \delta$  for all  $i = 1, 2, \dots, L$  Tested without use of inference
  
- Kroll et al. (2000)  $\sum_{i=1}^L \{g_k(X_i) - g_1(X_i)\}^2 / X_i$ 
  - Composite statistic, ADL Wrong power profile
  - Ha:  $ADL > \lambda$  Difficult to choose  $\lambda$

# More literature

- Hsieh and Liu (2008) Choice of concentrations?
  - Ha: I-U tests  $|g_k(X_i) - g_1(X_i)| < \delta$  for all  $i = 1, 2, \dots, L$
- Hsieh, Hsiao, and Liu (2009) Difficult to choose  $\psi$   
Generally  $\psi \ll \delta^2$  !
  - Composite statistic,  $SSDL = L^{-1} \sum_{i=1}^L \{g_k(X_i) - g_1(X_i)\}^2$ .
  - Ha:  $SSDL < \psi$
  - Generalized pivotal quantity (GPQ) method

# Our proposed hypothesis

- $H_a: |g_k(x) - g_1(x)| < \delta$  for all  $x \in [x_L, x_U]$
- Similar to I-U testing, but instead of individual concentrations, performed across a range of interesting concentrations.
- Bayesian(or GPQ) methods.
  - Linear models
  - Test is function of linear contrast

# Our proposed test statistic

$$|g_k(x) - g_1(x)| < \delta \quad \text{for all } x \in [x_L, x_U]$$

$$\Rightarrow \max_{x \in [x_L, x_U]} \left| \sum_{r=2}^k \theta_r f_r(x) \right| < \delta$$

$$p(\delta, x_L, x_U) = Pr \left\{ \max_{x \in [x_L, x_U]} \left| \sum_{r=2}^k \theta_r f_r(x) \right| < \delta \mid data \right\}$$

- Accept linearity if  $p > p_0$  (e.g.,  $p > 0.9$ ).



# How to: with Jeffrey's prior (or GPQ)

- Given  $\mathbf{Y}$  (responses) and  $\mathbf{F}$  (orth poly design matrix). Assume a  $k$ th-degree polynomial.
- Let  $\hat{\theta}$  and  $\hat{\sigma}$  be OLS estimates
- Error degrees of freedom =  $N-(k+1)$

- Generate two random variables

$$Z \sim N_{k+1}(\mathbf{0}, \mathbf{I})$$

$$U \sim \chi^2(N-k-1)$$

$$\theta_{Bayes} = \hat{\theta} - \hat{\sigma} \frac{Z}{\sqrt{U / (N - k - 1)}}$$

} Generate "B" of these

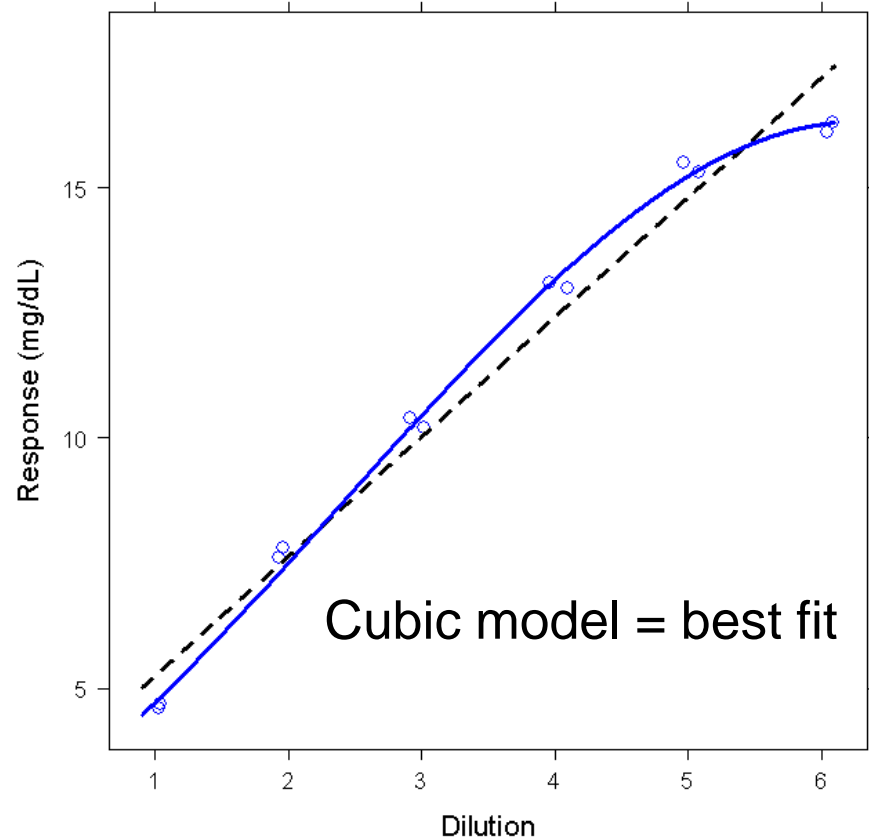
To estimate  $p(\delta, x_L, x_U)$ ,  
count the proportion of times:

$$\max_{x \in [x_L, x_U]} \left| \sum_{r=1}^k \theta_{Bayes,r} f_r(x) \right| < \delta$$

# Calcium Assay example

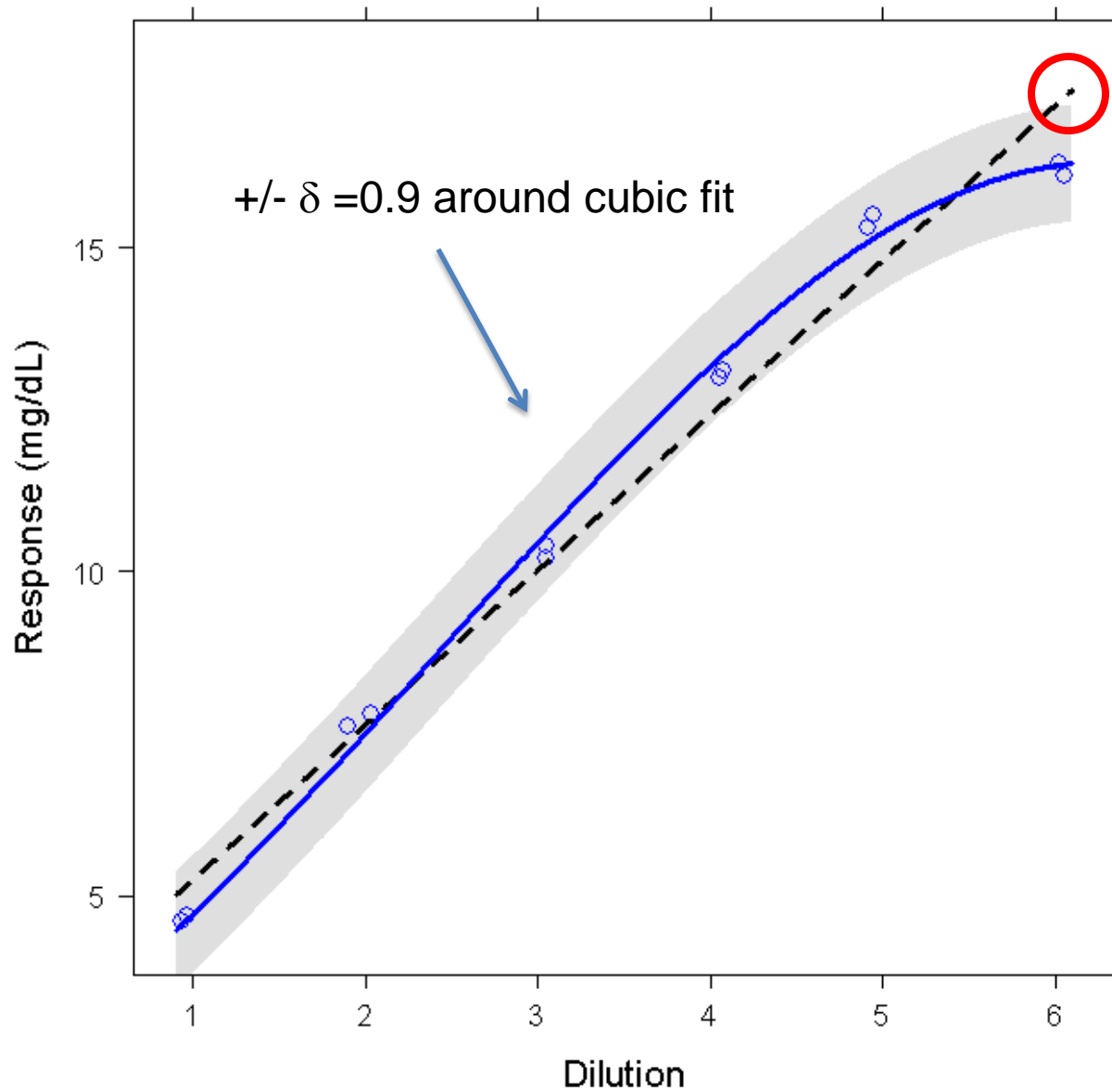
- From NCCLS EP6-A, Appendix C, ex. 2

Dilution	Replicate 1	Replicate 2
1	4.7	4.6
2	7.8	7.6
3	10.4	10.2
4	13	13.1
5	15.5	15.3
6	16.3	16.1



Compare cubic to linear from Dilution =1 to Dilution = 6:  $\delta = 0.9$  for testing.





Dilution	Mean difference: Cubic – Linear
1	-0.53
2	-0.13
3	0.42
4	0.74
5	0.42
6	<b>-0.93</b>

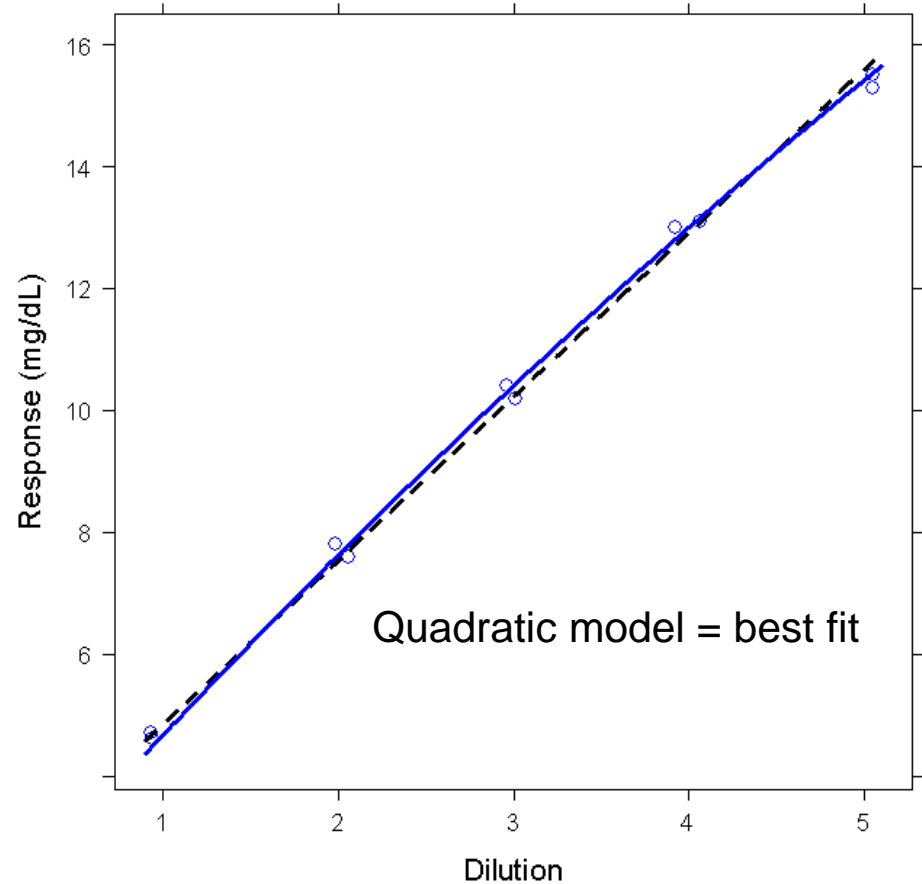
# Clear failure at Dilution = 6

- Hsieh and Liu I-U test p-value = 0.61
  - $|g_3(X_i) - g_1(X_i)| < \delta$  for all  $i = 1, 2, \dots, 6$
  - Linear fit not adequate
- $p(\delta, x_L, x_U) = 0.39$ 
  - $|g_3(x) - g_1(x)| < \delta$  for all  $1 \leq x \leq 6$
  - Linear fit not adequate
- Probability  $SSDL = \left(\frac{1}{6}\right) \sum_{i=1}^6 \{g_3(X_i) - g_1(X_i)\}^2 < \delta^2 > 0.99$ 
  - *Linear fit is equivalent to cubic fit*

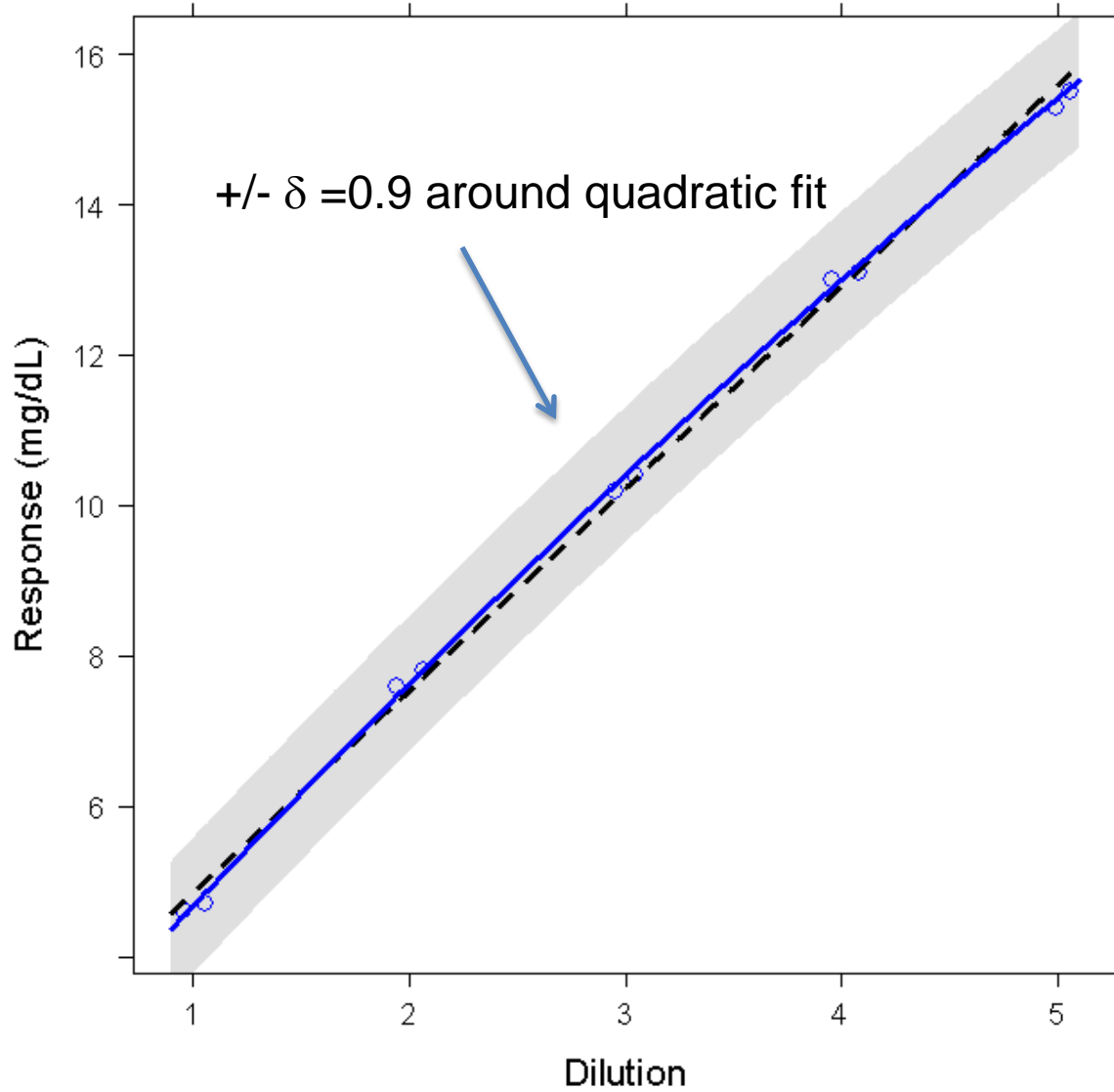
# Try again without last dilution

- From NCCLS EP6-A, Appendix C, ex. 2

Dilution	Replicate 1	Replicate 2
1	4.7	4.6
2	7.8	7.6
3	10.4	10.2
4	13	13.1
5	15.5	15.3
6		



Compare quadratic to linear from Dilution =1 to Dilution = 5 with  $\delta = 0.9$  for testing.



Dilution	Mean difference: Quad – Linear
1	-0.18
2	0.09
3	0.18
4	0.09
5	-0.18
6	

# Linear and quadratic fits equivalent

- Hsieh and Liu I-U test p-value  $< 0.01$
- $p(\delta, x_L, x_U) > 0.99$
- SSDL probability  $> 0.99$
- *Linear fit is equivalent to quadratic fit for dilutions between 1 and 5.*

# Simulation

# Quadratic vs. Linear Simulation

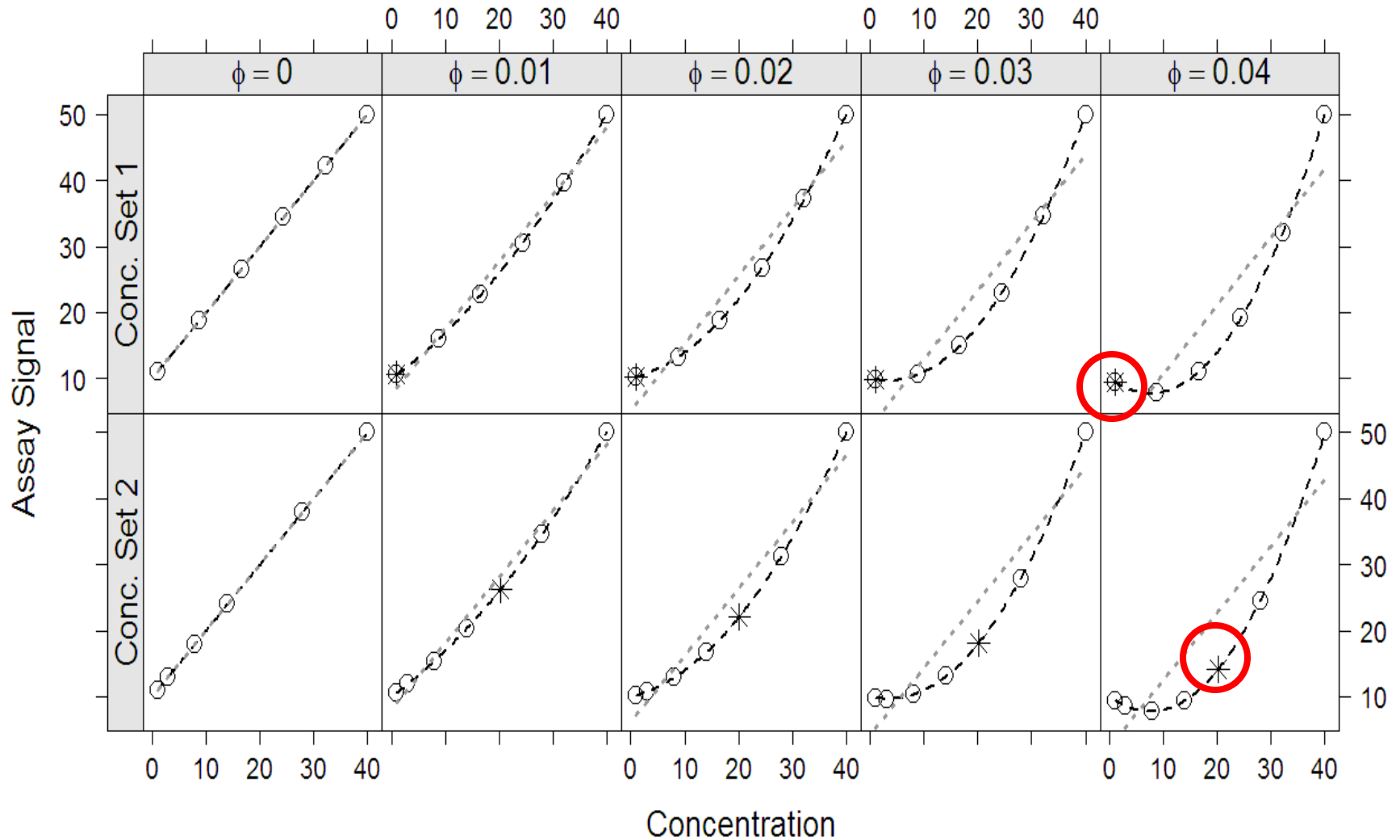
- $g_2(x) = 10 + (1-40\phi)x + \phi x^2, \quad 1 \leq x \leq 40$
- $\phi = 0 =$  linear:  $g_2(x) = 10 + x$
- $\phi = 0.04 =$  large quadratic component.
- $Y \sim N( g_2(x), \sigma=3 ). \quad 10 \leq g_2(x) \leq 50$
- 6 concentrations with two replicates each
- Testing limit:  $\delta = 6.1$

Two sets of concentrations

Set 1: Maximum deviation occurs at  $x = 1$

Set 2: Maximum deviation occurs between points

This case is H0/H1 border



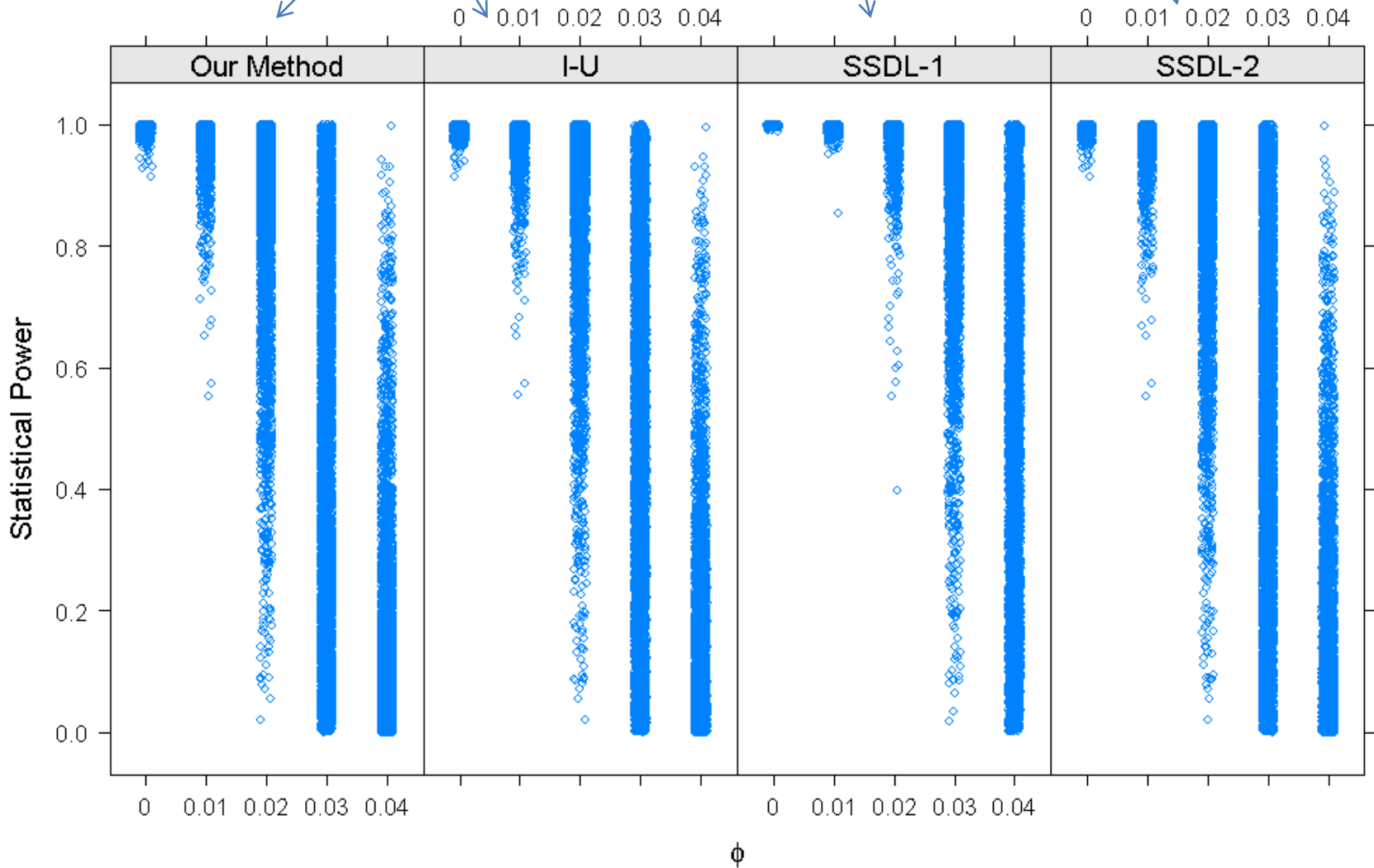


# Maximum difference occurs at $x = 1$

Because max diff occurs at design point, these tests are very similar

Test is generally Too powerful

SSDL can be tuned with knowledge of unknown curve.

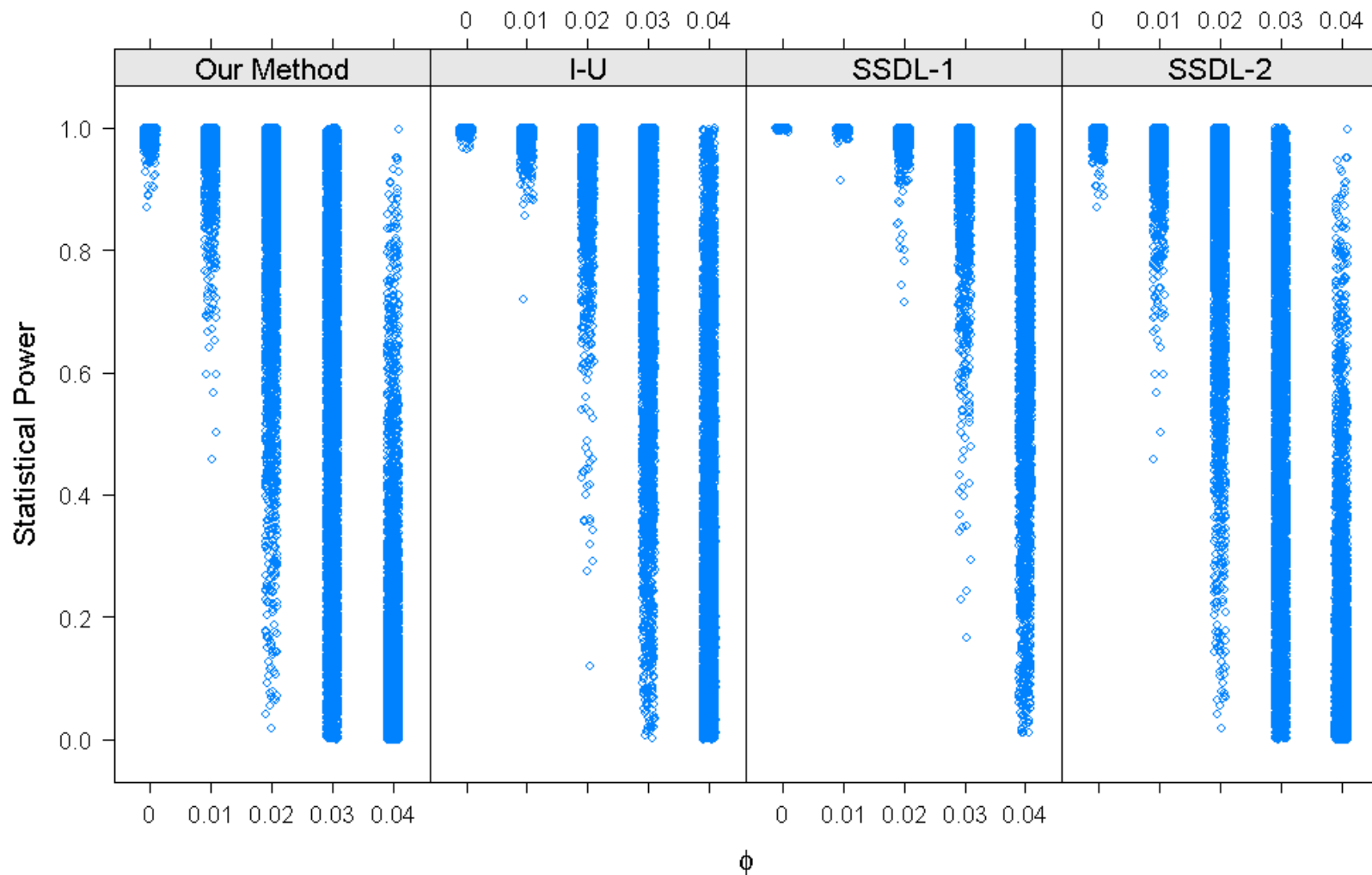


# Maximum difference occurs between points

Our method retains similar power profile

Tests are generally Too powerful

SSDL can be tuned with knowledge of unknown curve.



# Inverting the test

- Consider the true mean response  $g_k(x)$  and the reduced model  $g_1(x)=a+bx$ .
- Let  $z_k(x) = \{ g_k(x) - a \} / b$ 
  - This is polynomial Y-value back-calculated with best-fitting straight line.
- How close is  $z_k(x)$  to  $x$  ?

# A few hypotheses to consider

- $| z_k(x) - x | < \delta$  for all  $x_L \leq x \leq x_U$

- $| 100\% \{ z_k(x) - x \} / x | < \delta$

Many others!

- $| \log\{z_k(x)\} - \log(x) | < \delta$

- We can compute conditional probability

$$\Pr\{ | \log\{z_k(x)\} - \log(x) | < \delta \mid \text{data} \}$$

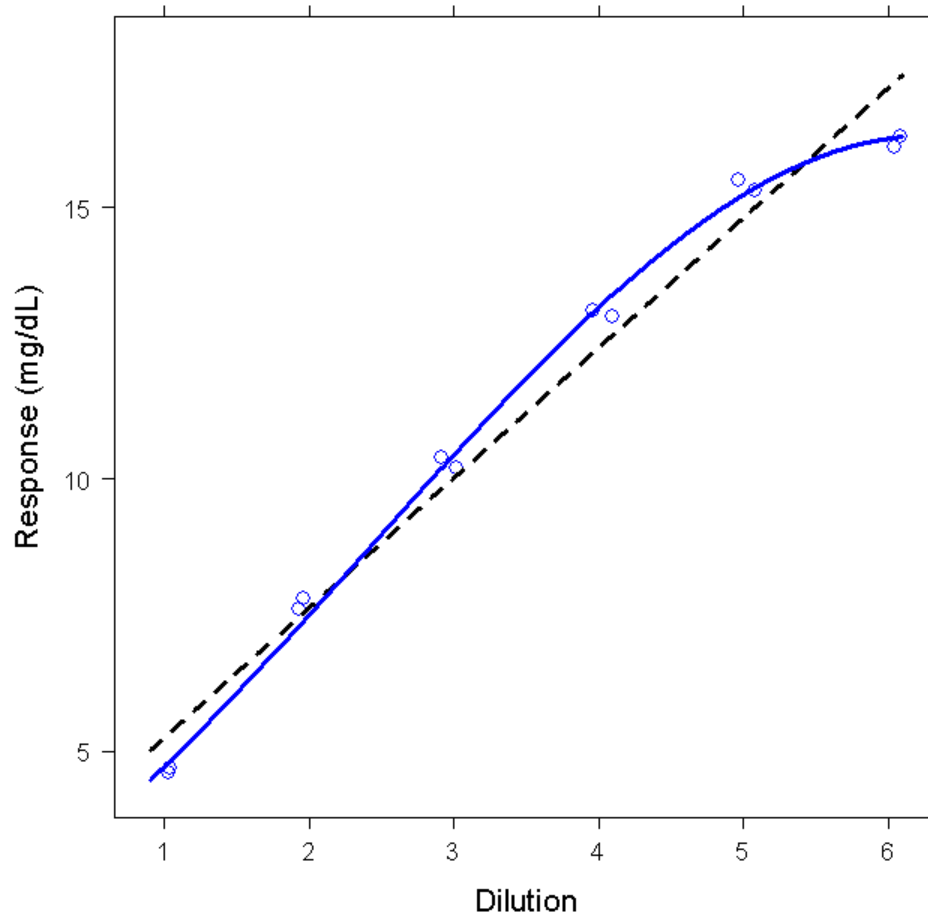
Or

- Find  $\delta$  such that

$$\Pr\{ | \log\{z_k(x)\} - \log(x) | < \delta \mid \text{data} \} = 0.95$$

$$\Pr\{|\log\{z_3(x)\} - \log(x)| < 0.15 \mid \text{data}\} = 0.95$$

for all  $1 \leq x \leq 6$

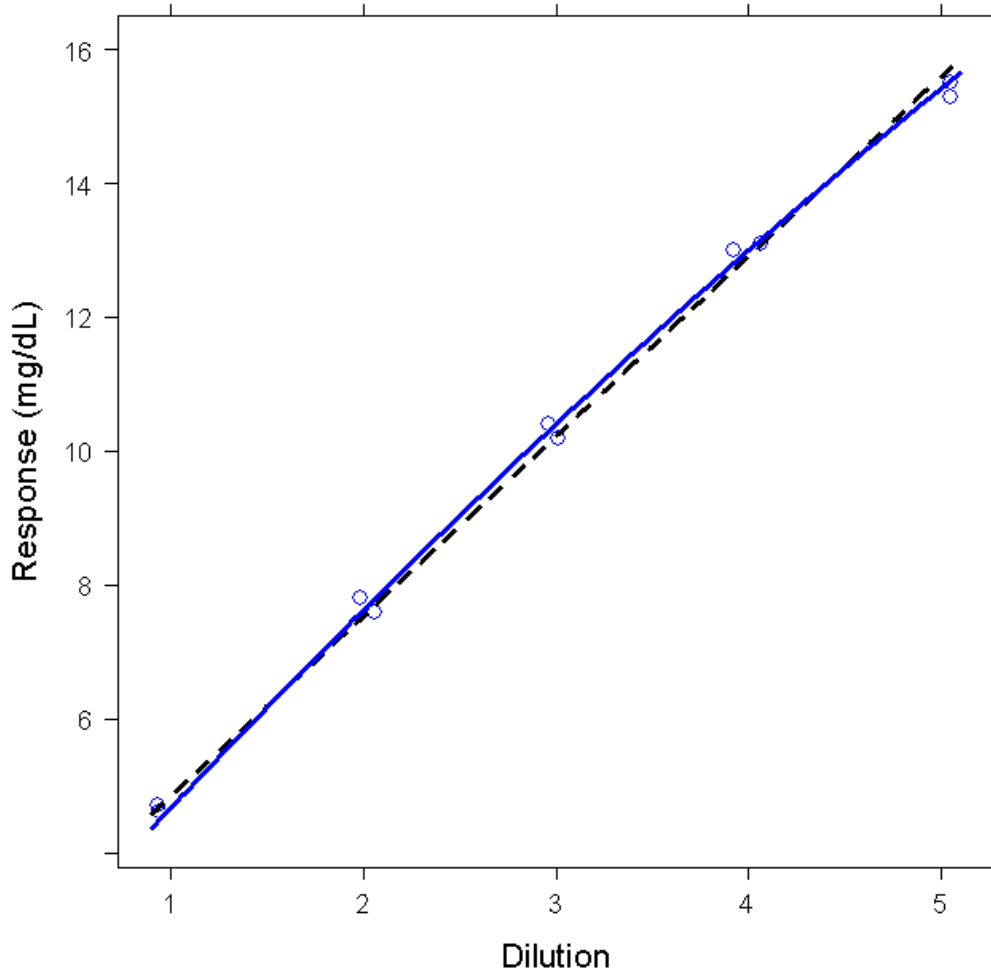


Back-calculated values are within 0.15 log<sub>10</sub> units of true value.

Back-calculated values are within  $100\%(10^{0.15}-1) = 40\%$  of true value.

$$\Pr\{|\log\{z_2(x)\} - \log(x)| < 0.05 \mid \text{data}\} = 0.95$$

for all  $1 \leq x \leq 5$



Back-calculated values are within 0.05 log<sub>10</sub> units of true value.

Back-calculated values are within  $100\%(10^{0.05}-1) = 12\%$  of true value.

# Extra bits

- When  $k=2$  (quadratic vs. linear), the proposed test statistic is central T distributed.
- When  $k = 2$ , by altering the testing limits, the I-U, SSDL, and proposed test methods can be made equal.
- From simulations, test size for the proposed test statistic appears to be  $\leq \alpha$ , depending on the experimental design.



# Summary

- Test method extends idea of NCCLS EP6-A by computing probability that best-fit curve is equivalent to a linear fit.
- Testing performed across a range of concentrations and *not* at experimental design points.

# References: Guidelines

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Thank you!

Questions?