

Modeling Error in Equilibrium-Restricted Factor Analysis

Nathanael Kazmierczak and Douglas A. Vander Griend, Ph.D., Department of Chemistry & Biochemistry, Calvin College

Introduction

- Factor analysis: a data mining technique that expresses a matrix as the product of two factor matrices.
- Application: UV-vis spectrophotometric titrations of equilibrium metal-ligand and host-guest complexation systems.
- Multiple error sources with non-normal distributions.
- Error propagation poorly understood.
- Technique often treated as a “black box.”
- Beer’s law: $A = \epsilon \times L \times c$
- Factor matrices obtained through constrained nonlinear least-squares optimization:

$$\text{Absorbance } (n \times p) = (\epsilon \text{ } (n \times m)) \times (\text{Equilibrium Concentrations } (m \times p)) + \text{Residual } (n \times p)$$

Diagnosing Error

1) Composition error

- Source: balance imprecision, user error
- Remedy: “Wiggle” stock solution concentration and addition volume.

2) Transmittance error

- Source: spectrometer imprecision
- Remedy: no method for elimination

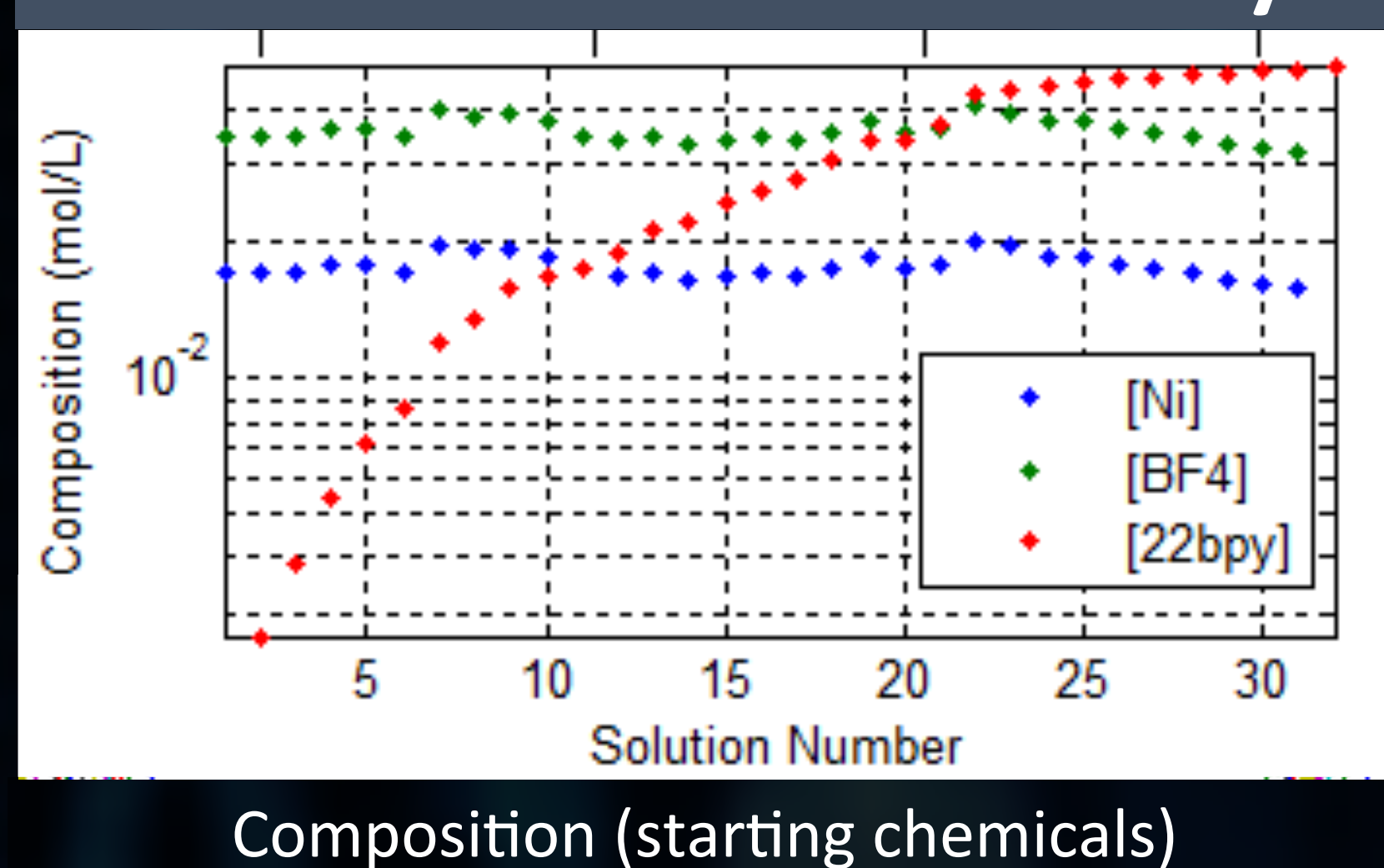
3) Stoichiometry error

- Source: user-defined equilibrium model
- Remedy: search all possible stoichiometries with a computer.

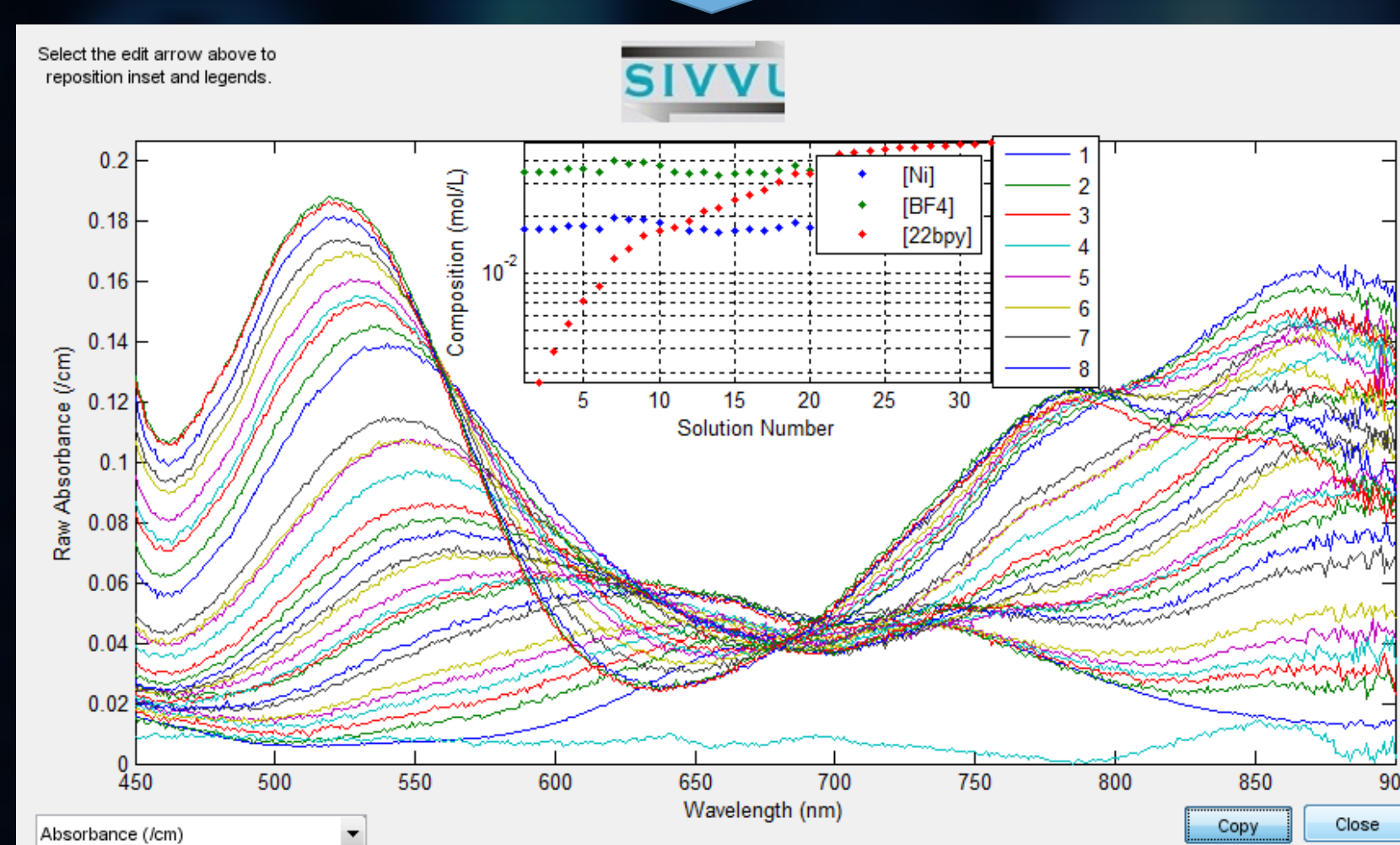
Simulating Error

- Artificial data:** absorbance data sets obeying the laws of chemical equilibrium are generated from concentration and absorptivity factors *in silico*.
- Monte Carlo simulations:** random error patterns of a given standard deviation are added to identical artificial data sets between 30 and 100 times and the model optimized for each. The calculated results form a distribution illustrating error bars.

Titration Anatomy

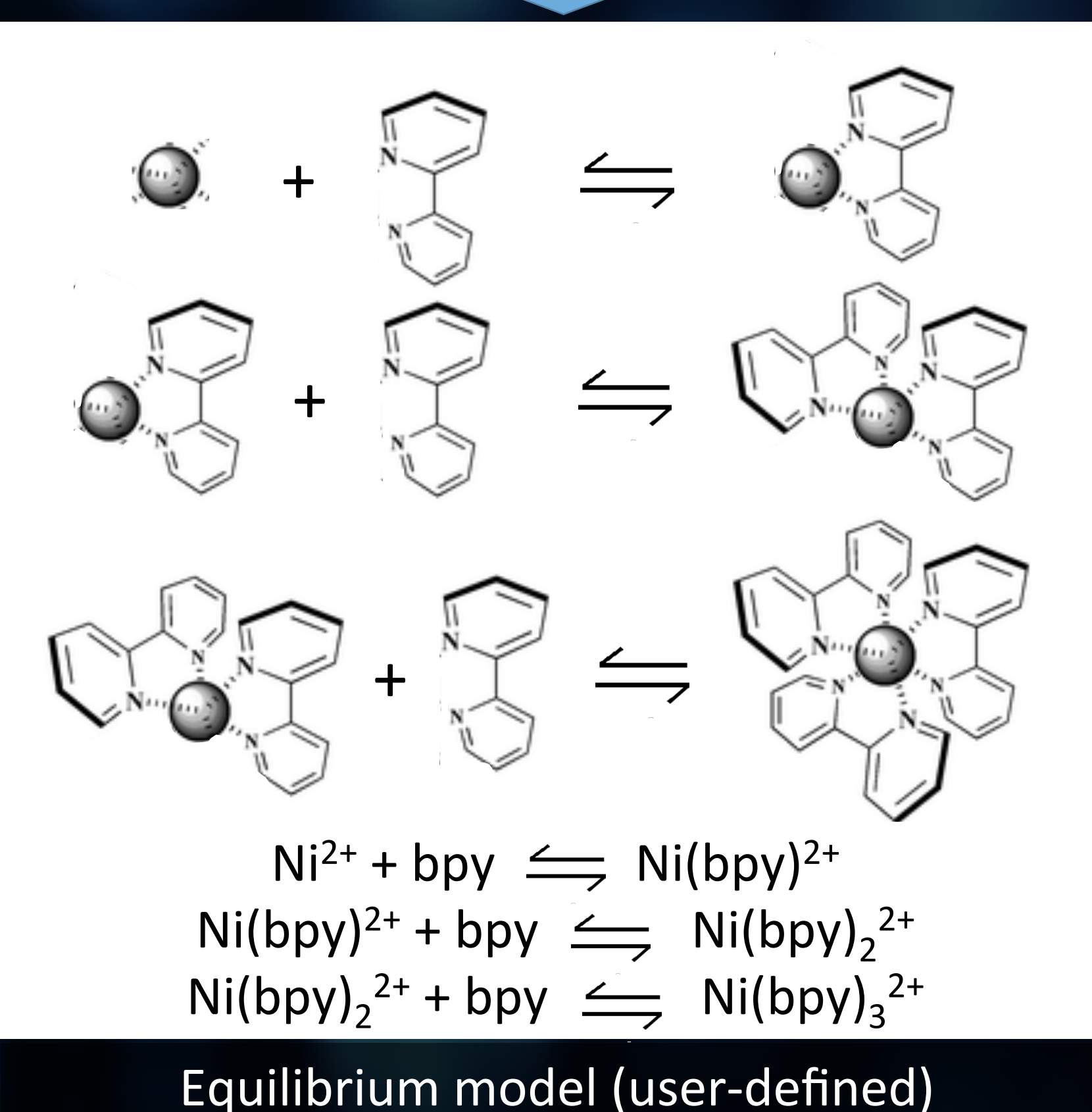


Equilibrates

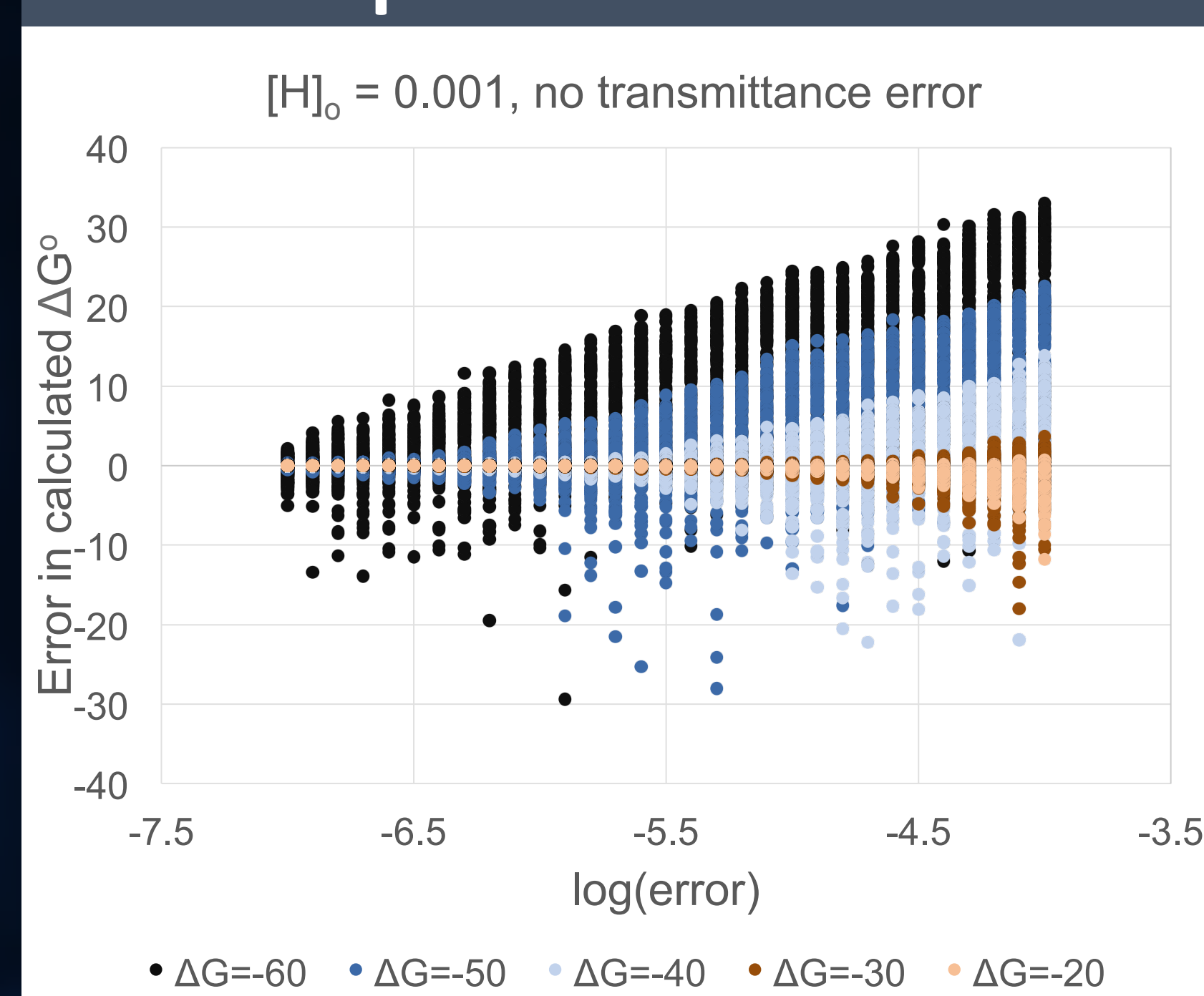


Spectrometer measures absorbance data

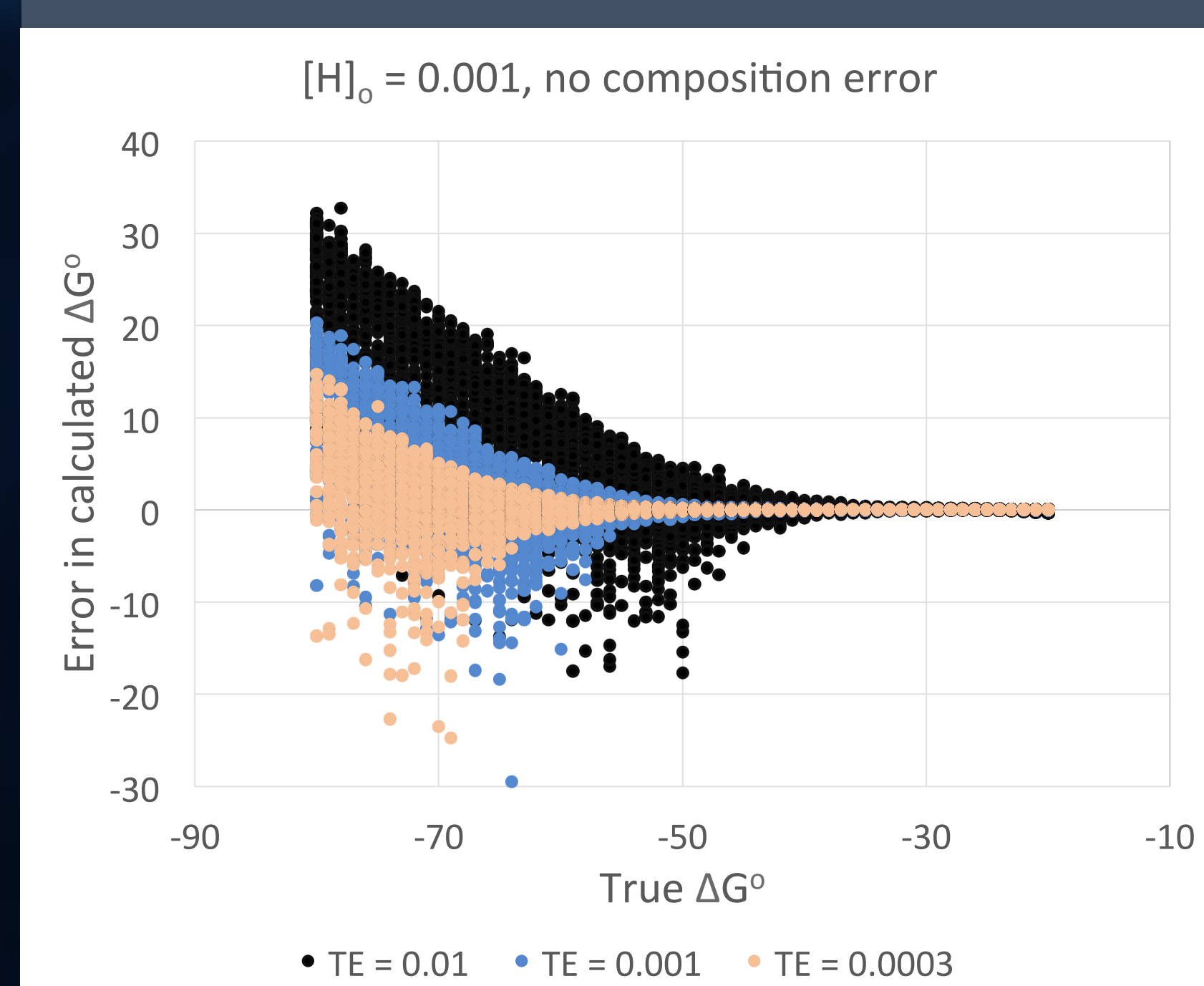
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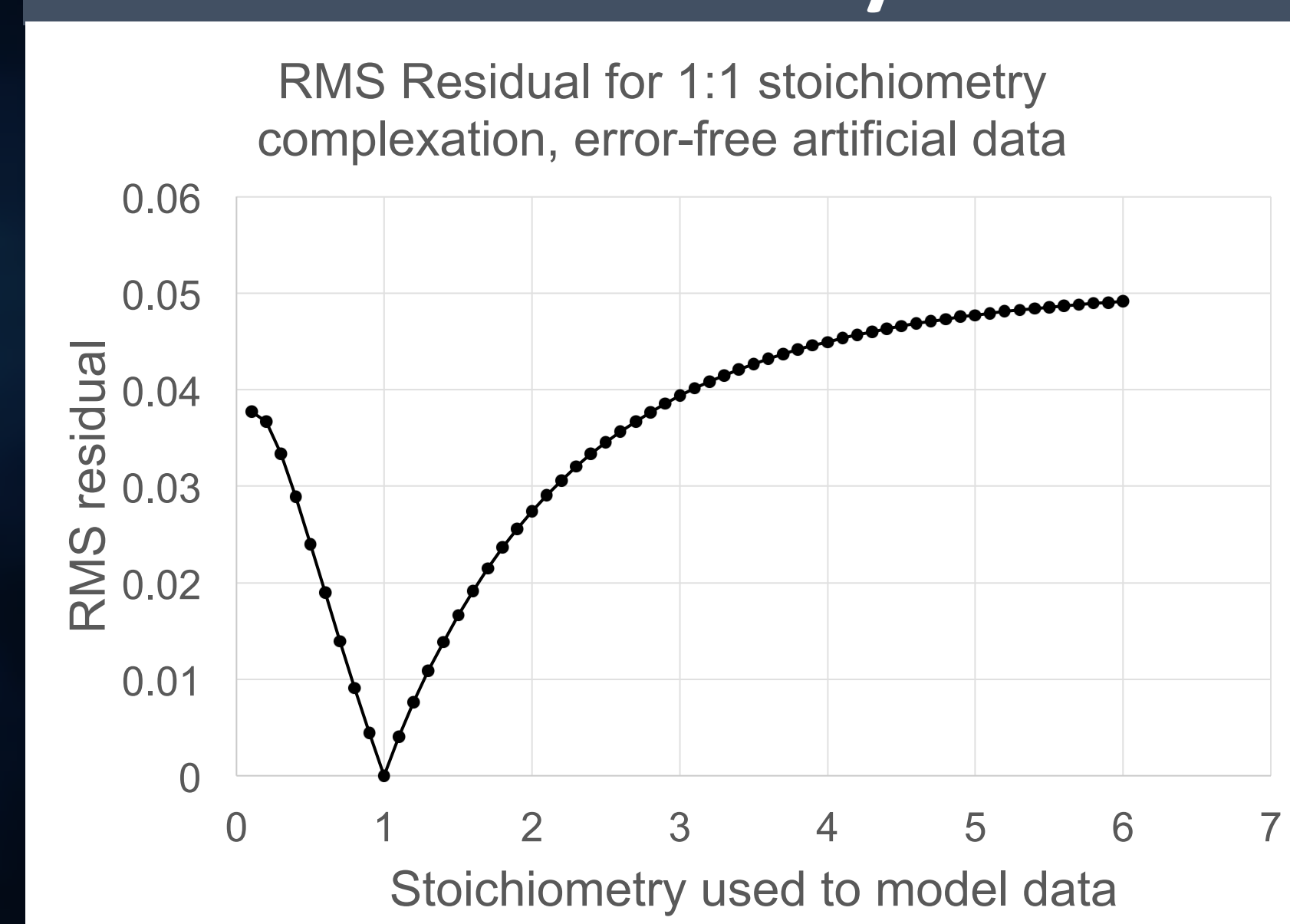
Composition Error



Transmittance Error



Stoichiometry Error



1:1 Exergonic Limits

| $[\text{H}]_0$ | ΔG° lower bound (kJ/mol) at 298 K | K | $K[\text{H}]_0$ |
|--------------------|--|--------------------|-----------------|
| 1×10^{-6} | -84 | 5×10^{14} | 5×10^8 |
| 1×10^{-5} | -79 | 7×10^{13} | 7×10^8 |
| 1×10^{-4} | -72 | 4×10^{12} | 4×10^8 |
| 1×10^{-3} | -68 | 8×10^{11} | 8×10^8 |
| 1×10^{-2} | -62 | 7×10^{10} | 7×10^8 |
| 1×10^{-1} | -56 | 6×10^9 | 7×10^8 |
| 1 | -51 | 8×10^8 | 8×10^8 |

- 0.0003 transmittance error added (best-case spectrometer error)

Conclusions

- Monte Carlo adaptations of SIVVU™ software allowed high-volume error studies.
- Each type of error has a distinct signature.
- Each type of error can be diagnosed separately.
- Composition and transmittance error effects are amplified for highly exergonic reactions owing to a loss of model sensitivity to the ΔG° value.
- There exists a smooth RMS minimum at the correct model stoichiometry, potentially allowing optimization to find the correct model.
- For 1:1 reactions, there exists a fundamental lower bound to the ΔG° value that can be reliably calculated in the presence of spectrometer transmittance error.

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