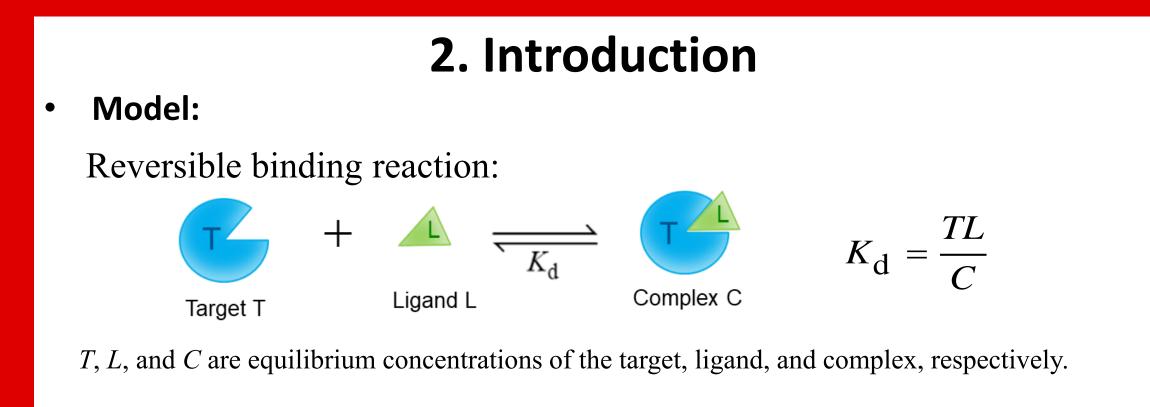


Accuracy Assessment for Equilibrium Dissociation Constant Using a Single Binding Isotherm

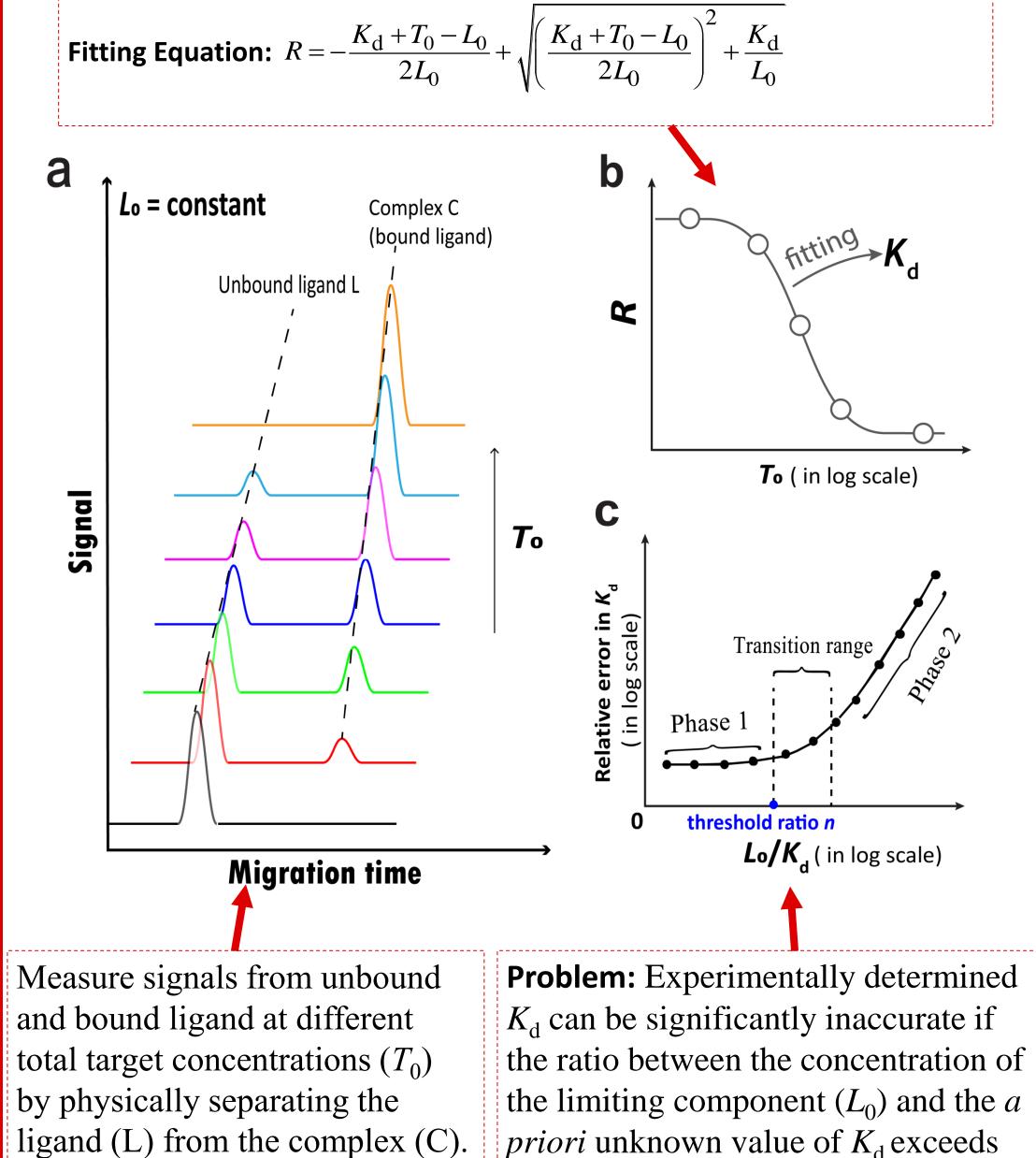
1. Abstract

The equilibrium dissociation constant (K_d) characterizes stability of non-covalent molecular complexes. Determining K_d of complexes may be extremely inaccurate if the ratio between the concentration of the limiting component (L_0) and the *a priori* unknown value of K_d exceeds an unknown threshold ratio. The only known approach to reveal this kind of inaccuracy in K_d requires building multiple experimental binding isotherms; it is resource intensive and, therefore, used very rarely. We introduced a single-isotherm approach for assessing $K_{\rm d}$ accuracy via determining the value of L_0/K_d , estimating the threshold ratio, and comparing L_0/K_d to the threshold ratio. In this proof-ofconcept work, we present the theoretical basis and develop a step-bystep algorithm for our single-isotherm approach. We also demonstrate the experimental use of the developed algorithm.



The classical approach of determining K_d in equilibrium methods and the problem faced:¹

- Calculate the fraction of unbound ligand $R = L/L_0$ based on the measured signals.
- Build the binding isotherm of "R vs T_0 ".
- Fit the binding isotherm with theoretical dependence of R on T_0 to determine $K_{\rm d}$.



an unknown threshold value.

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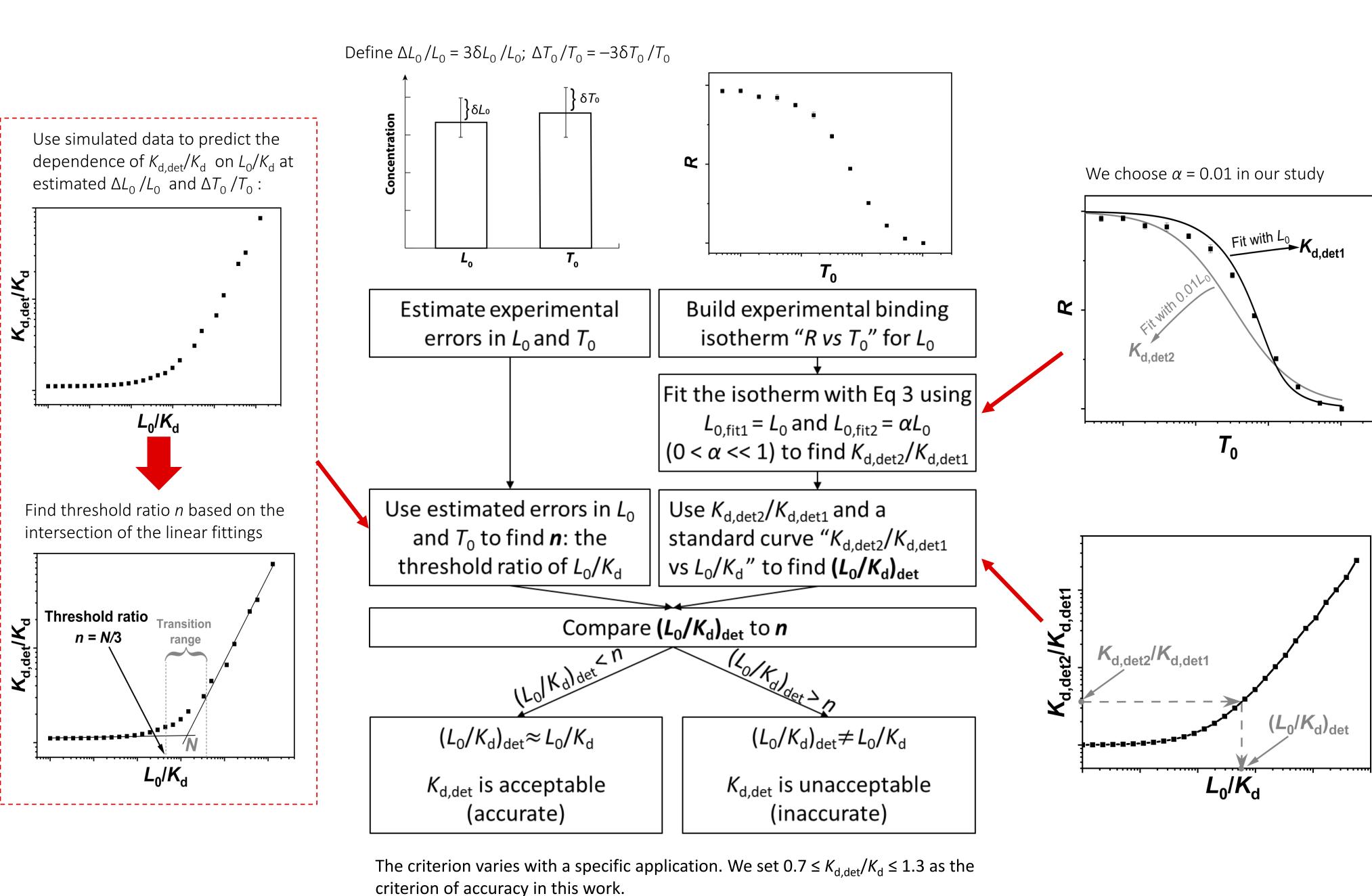
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3. Previous solution

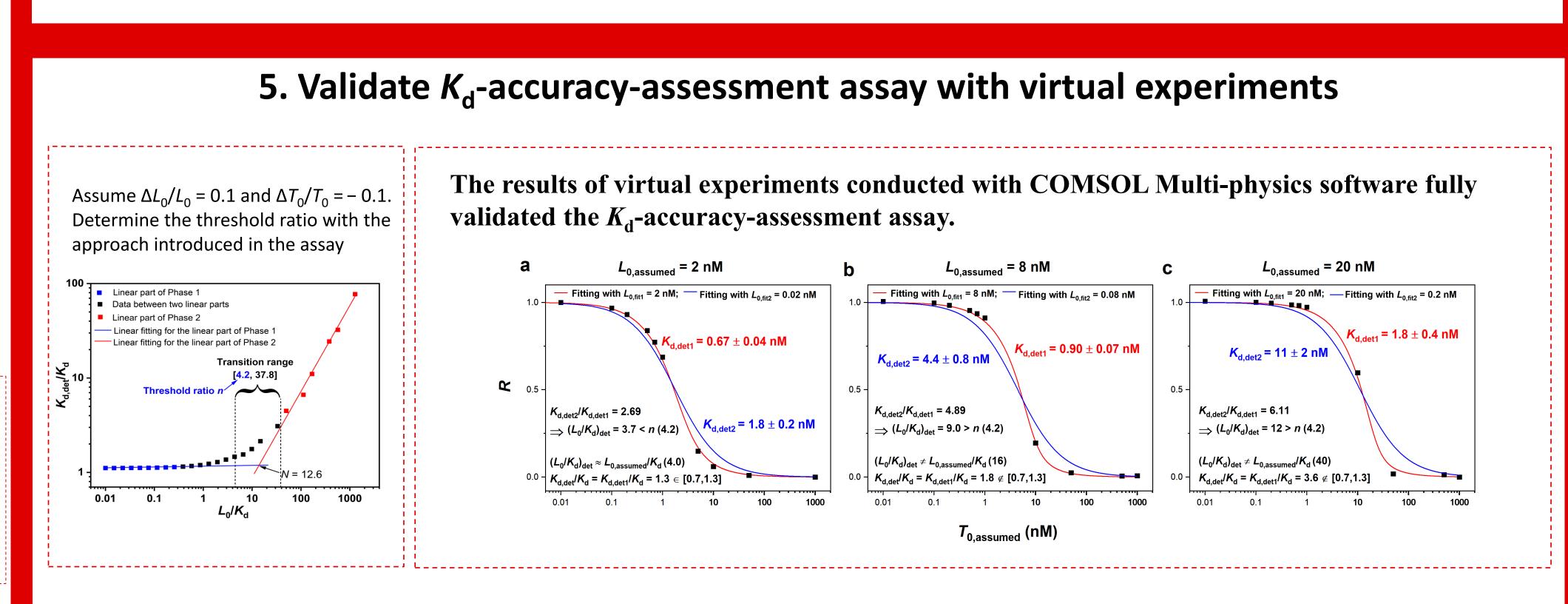
Experimentally obtain multiple binding isotherms within a range of L_0 values. If determined K_d ($K_{d,det}$) is found to be insensitive to changing L_0 , then it is assumed that determined $K_{d,det} \approx K_d$ (real K_d value).

Disadvantage: This approach drastically increases the amount of required experimental work and has been reported only in approximately 5% of publications concerned with the determination of $K_{\rm d}$.²

4. Our Solution



Advantages: This assay does not require additional binding isotherms. It will equip researchers with a "no-cost" analytical tool for assessing the accuracy of K_d values determined in equilibrium experiments.



K_d-accuracy-assessment assay using a single binding isotherm

Separation (ACTIS)³ ----- Fitting with $L_{0.fit1}$ = 10 μ M; ----- Fitting with $L_{0.fit2}$ = 0.1 μ M $K_{ m d,det1}$ = 30 ± 1 μM $\Rightarrow (L_0/K_d)_{det} = 0.3 < n (3.0)$ $(L_0/K_d)_{det} \approx L_0/K_d \ (0.36 \pm 0.08)$ **0.0** $- K_{d,det}/K_d = K_{d,det1}/K_d = 1.1 \in [0.7, 1.3]$ assay. 😢 Kd Accuracy Assessment Program * 0.023 or* 0.082 Upper Phase* <mark>6</mark> ower Phase* 4 threshold.

Acknowledgement

Reference

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- Int. Ed. 2019, 131(20), 6707-6711.

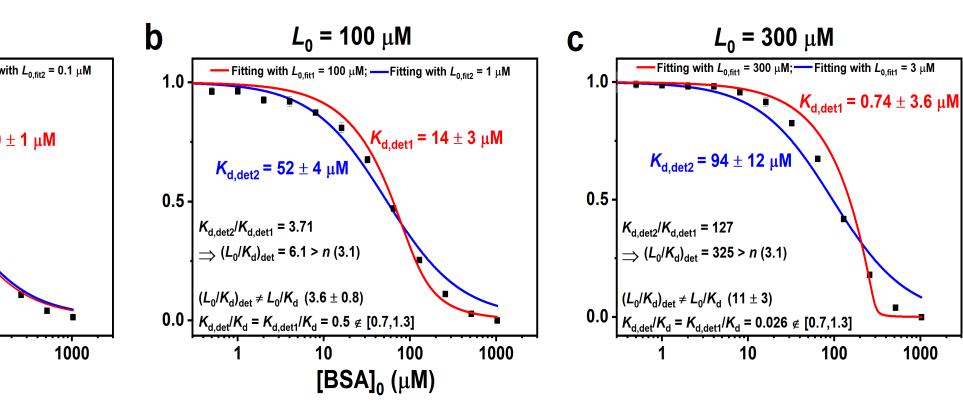




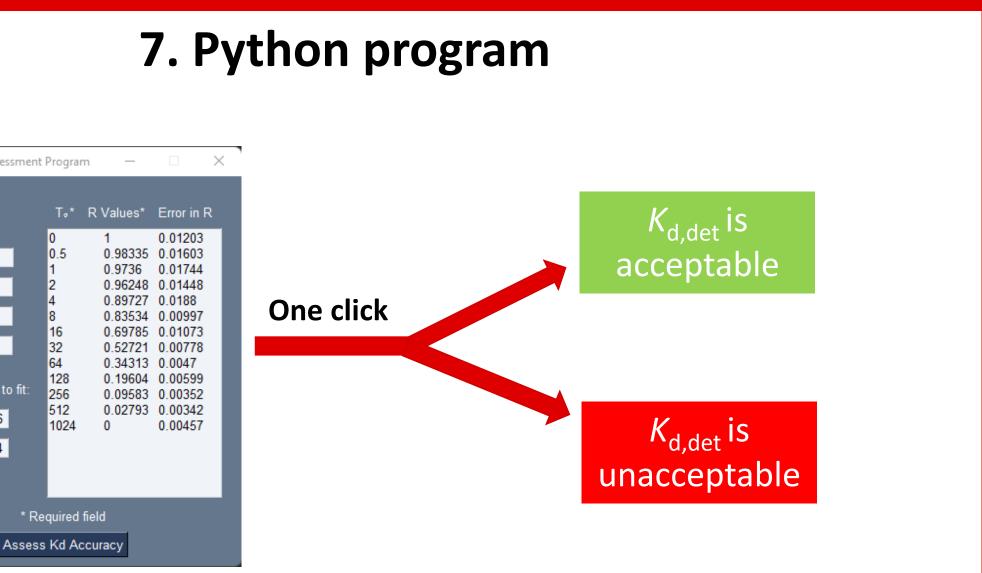
6. Validate K_d-accuracy-assessment assay with real experiments

• K_d -determination method: Accurate Constant via Transition Incomplete • Molecular pair: BSA-fluorescein

• Reference K_d value: $28 \pm 6 \,\mu\text{M}$



The results of real also confirmed the validation of the K_d -accuracy-assessment



8. Conclusion

In this proof-of-principle study, we introduced an approach for assessing the accuracy of a determined K_{d} value from a single experimental binding isotherm, which was named as K_d -accuracy-assessment assay.

 $K_{\rm d}$ -accuracy-assessment assay can assess the accuracy of determined $K_{\rm d}$ by solving two problems directly: (i) determining the threshold value for L_0/K_d under estimated errors in L_0 and T_0 and (*ii*) finding out if L_0/K_d is smaller than the

The assay was validated with both virtual and real experiments.

Using the algorithm of K_d -accuracy-assessment assay, we developed an Phyton program that can assess the accuracy of determined K_d in a few seconds.

Thanks for Jessica Latimer (orcid.org/0000- 0001-7961-6286) for writing the Python script for the K_{d} -accuracy-assessment assay.

1. Wang, T.Y.; Rukundo, J.-L.; Krylova, S. M.; Krylov, S. N. Accuracy Assessment for Equilibrium Dissociation Constant Using a Single Binding Isotherm. ChemRxiv (Analytical Chemistry). May 15, 2023, Ver. 2. DOI: 10.26434/chemrxiv-2023-zb599-v2 (accessed May 29,

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