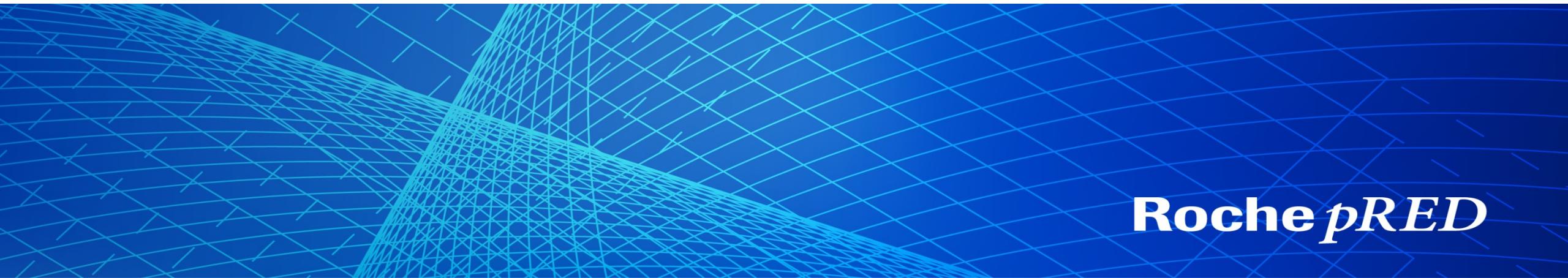

Are immunogenicity assay results really “incomparable”?

The critical role of bioanalysis to bring immunogenicity testing to the next level

Roland F. Staack

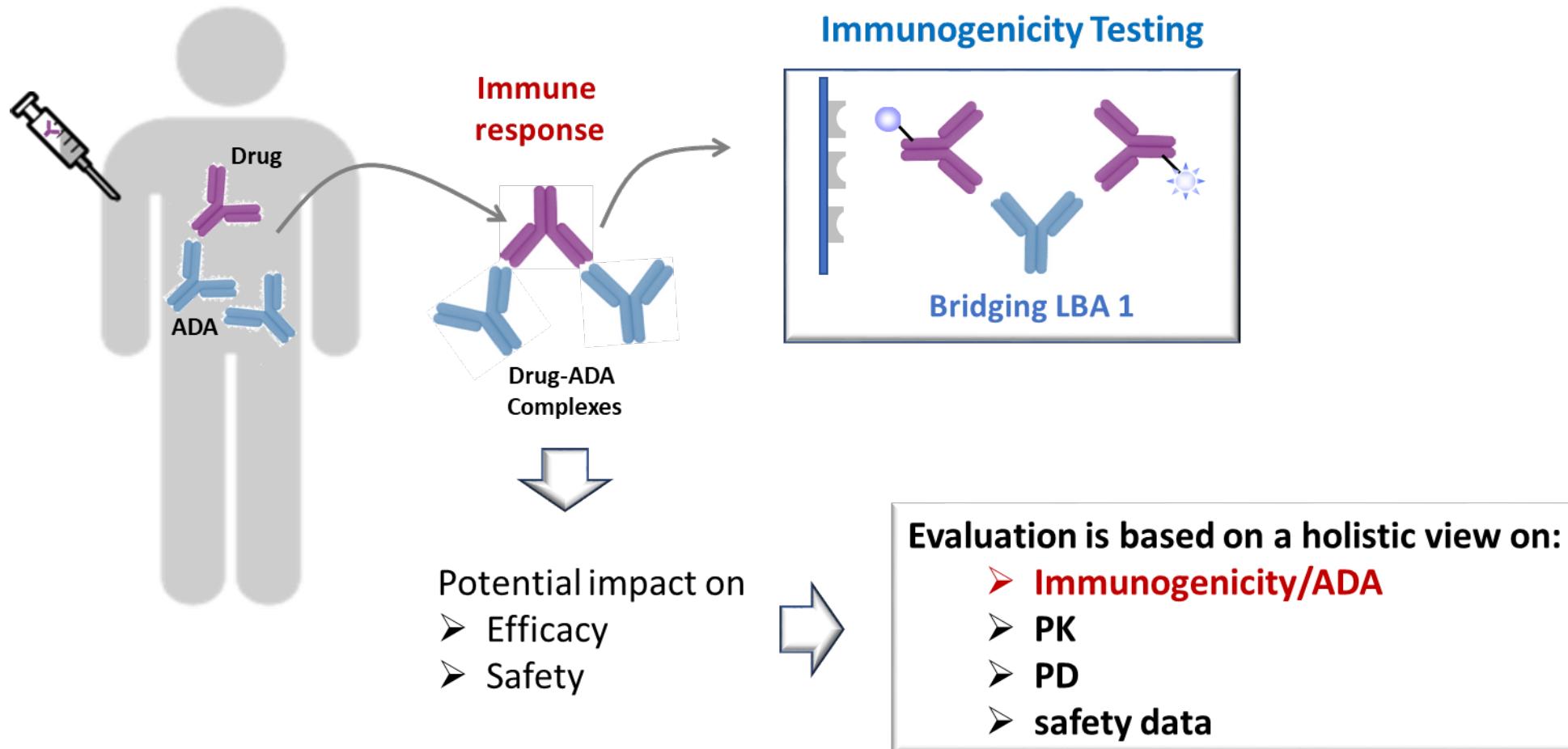
Pharmaceutical Sciences, Roche Pharma Research and Early Development, Roche Innovation Center Munich

13th EBF Open Symposium, November 17-20, 2020

A large, abstract graphic at the bottom of the slide features a blue grid that curves and slopes downwards from left to right, creating a sense of depth and perspective.

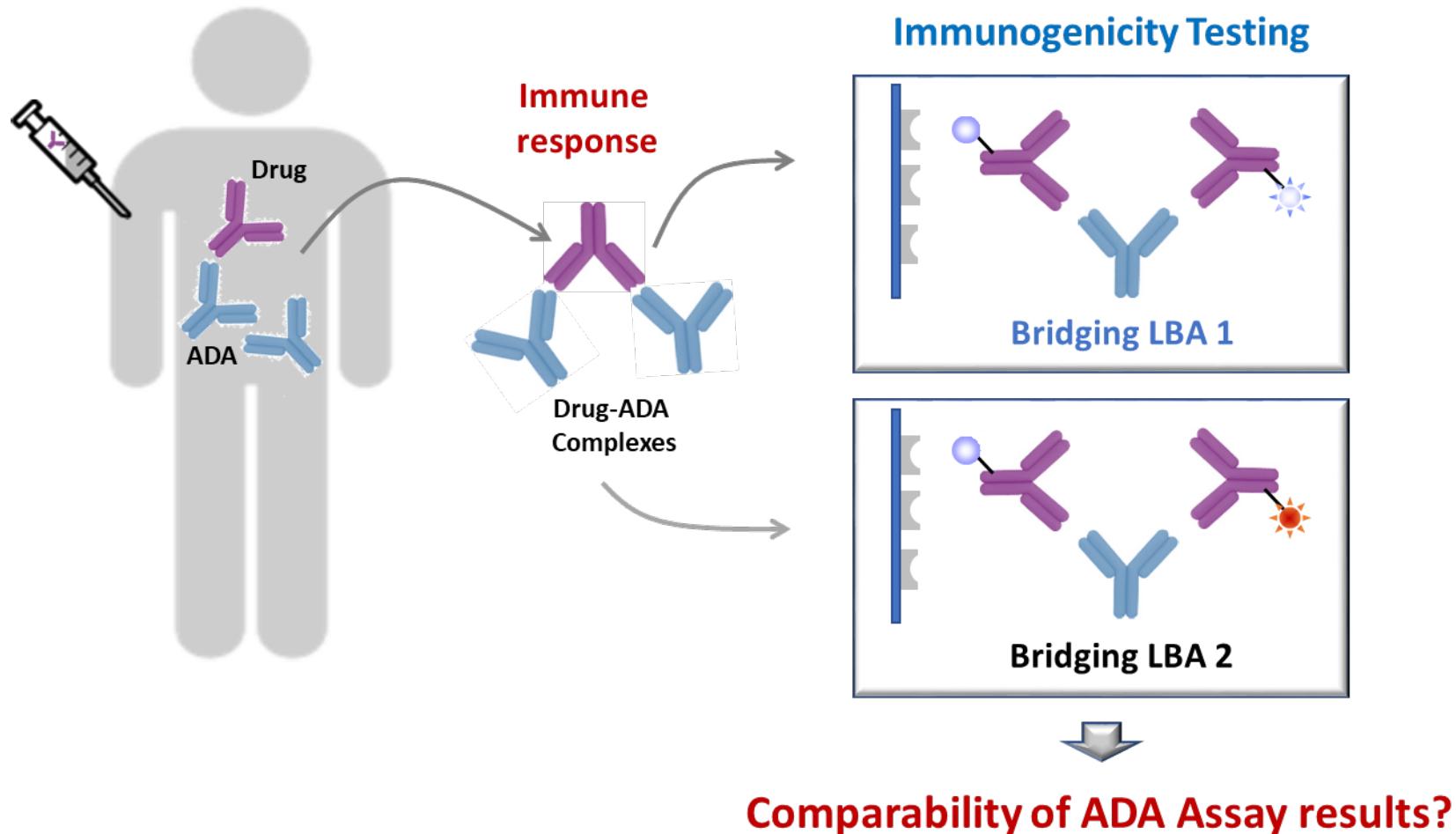
Development of therapeutic proteins

The immunogenicity challenge

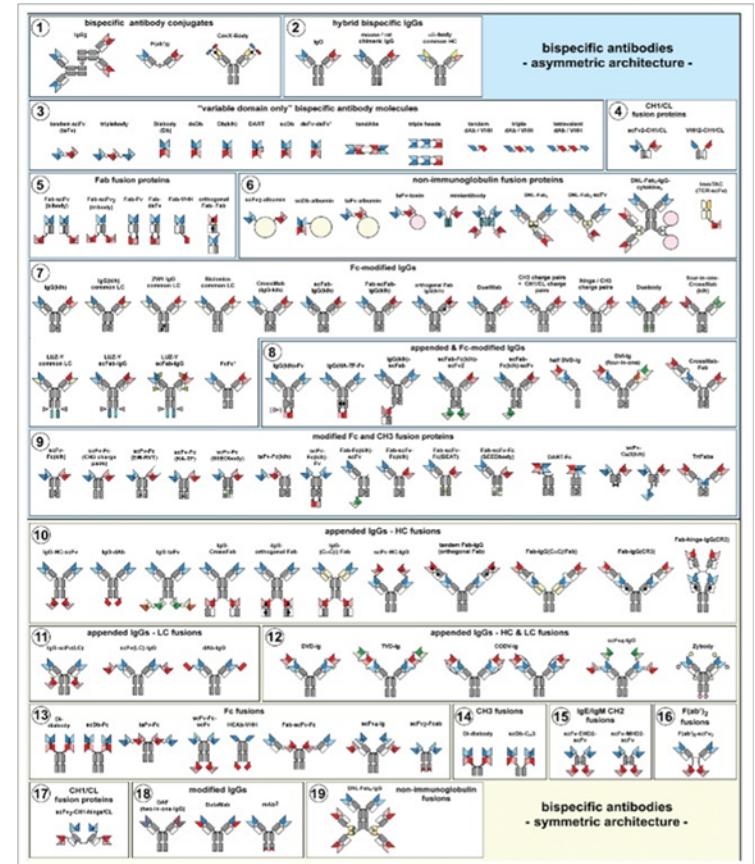
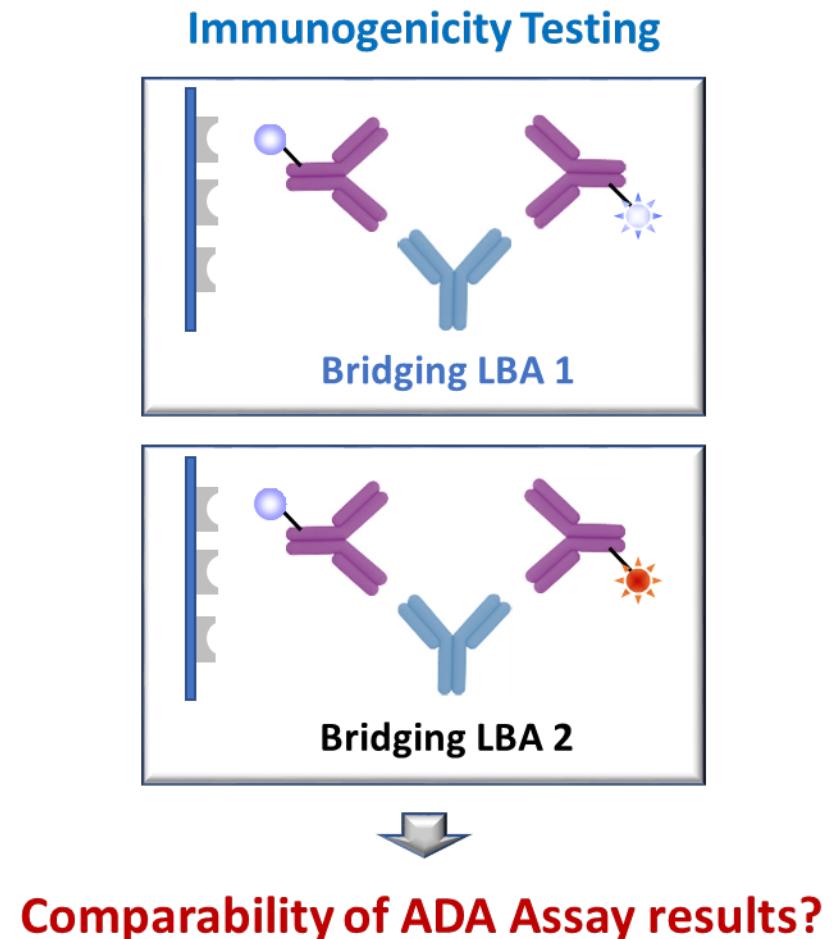
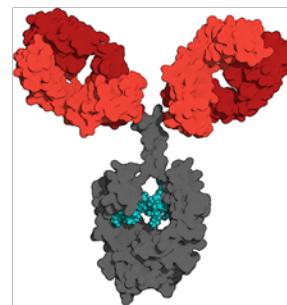
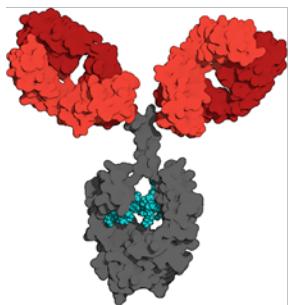


Development of therapeutic proteins

The immunogenicity challenge



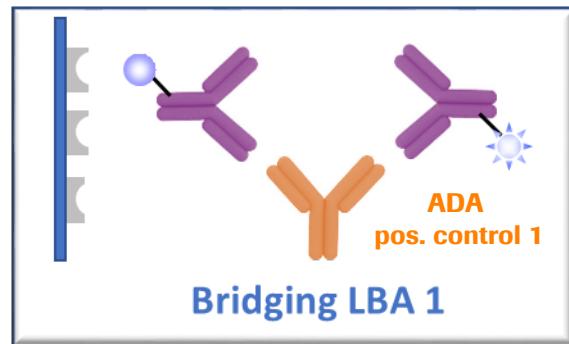
Development of therapeutic proteins



Brinkmann U & Kontermann RE, MAbs. 2017 Feb/Mar;9(2):182-212

Is the performance of the ADA assay the key to comparability?

The role of the „positive control“



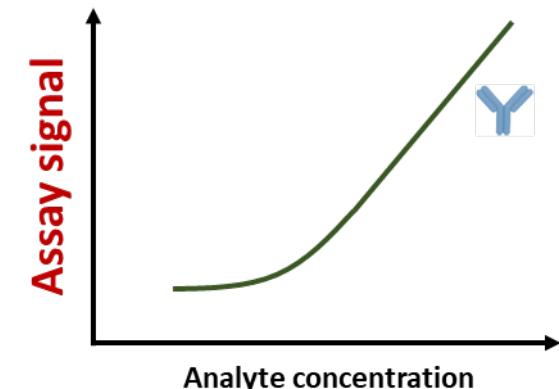
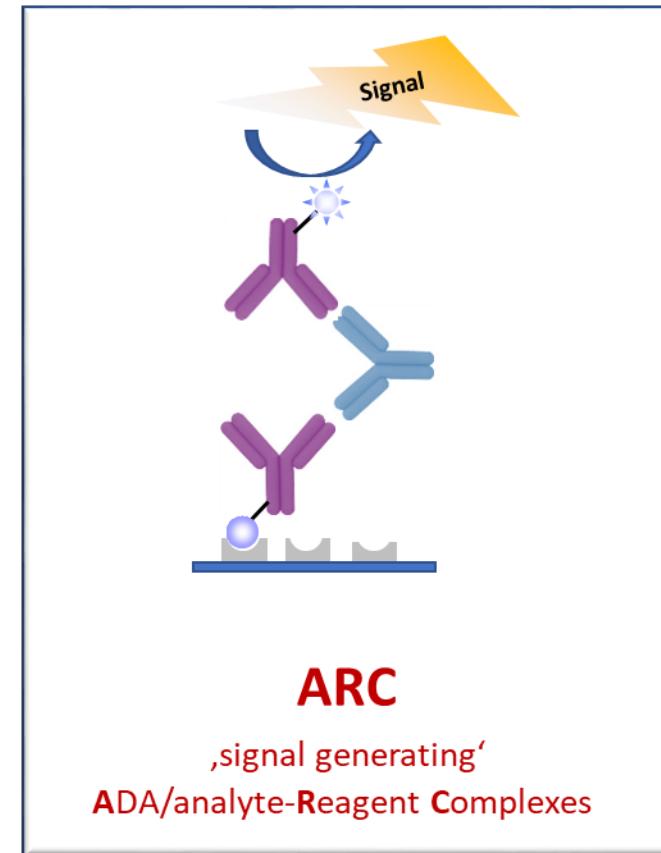
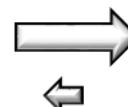
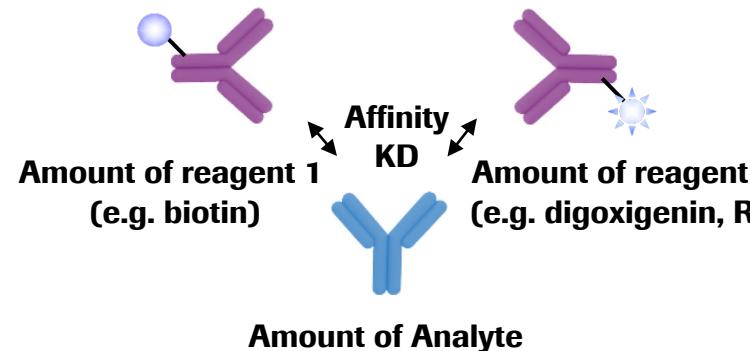
- Bioanalytical Immunogenicity/ADA data
= crucial piece of information for evaluation of clinical immunogenicity
 - Performance of BioA assay the critial parameter, particularly:
 - **Sensitivity**
 - **Drug Tolerance**
 - **Comparability** of different assays is **deemed highly difficult/not possible**
due to performance differences determined by different positive controls
- "Positive control = crucial issue**
- FDA proposal: „universal ADA-positive control“



Enabling comparable ADA assay performance = BioA contribution to achieve scientific sound comparability of the clinical immunogenicity of different drugs/biosimilars/scaffolds

Bioanalytical Assay Performance

What do we actually see?

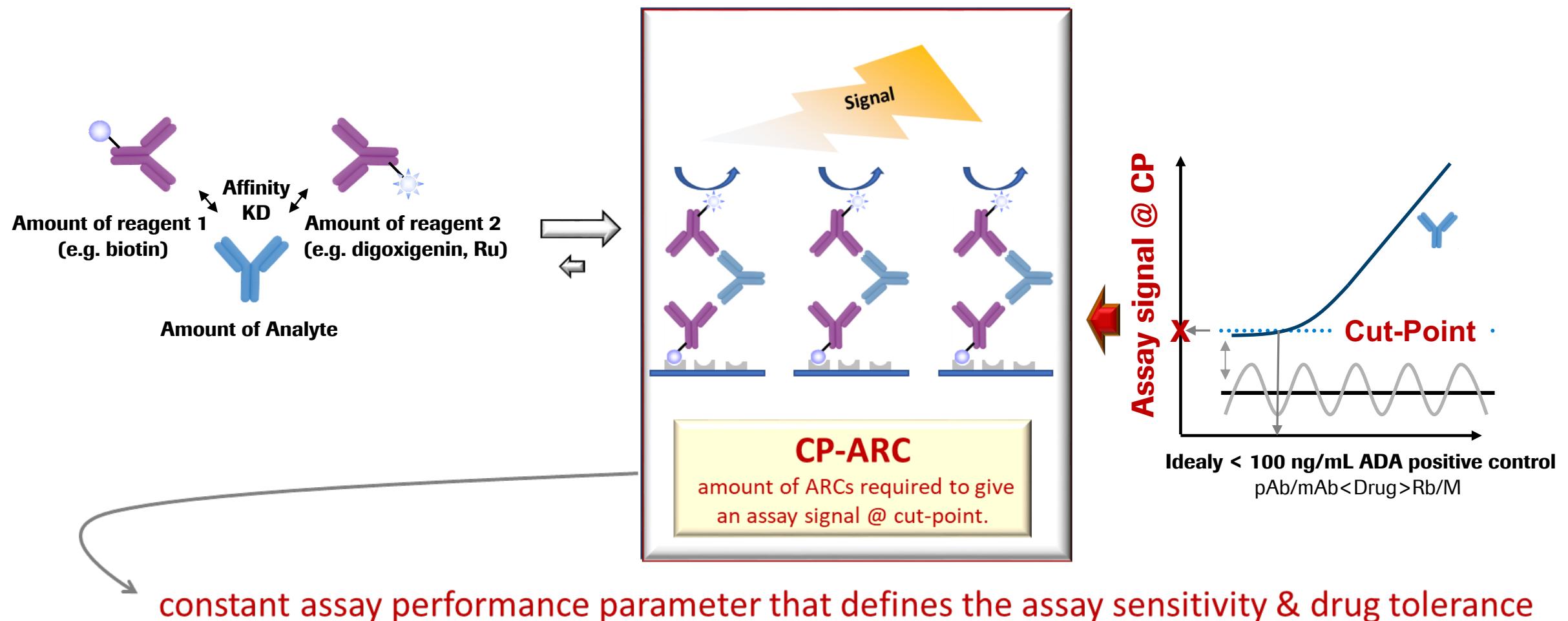


Intensity of the assay signal depends on:

- Amount of the analyte/ADA
- Amount of the reagents
- Affinity of analyte-reagent interaction

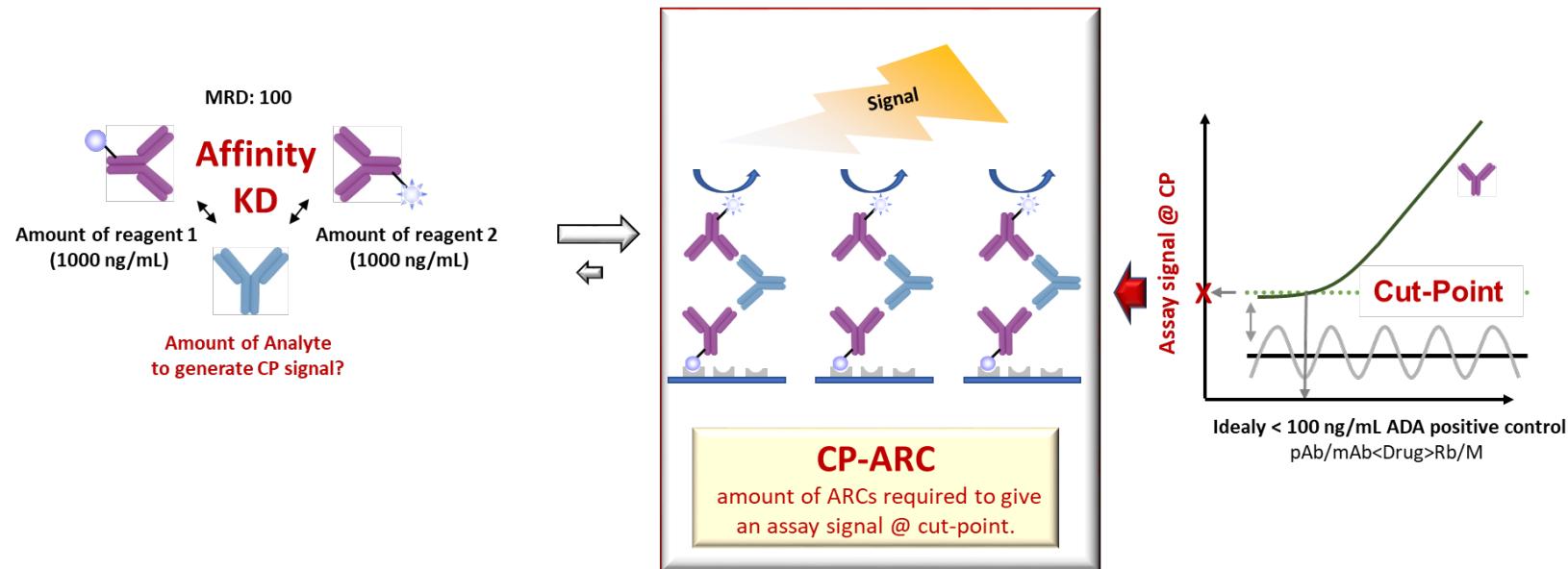
The role of formed ADA/Analyte-Reagent-Complexes at CP (CP-ARC)

... a constant assay performance parameter?



Impact of affinity on CP-ARC and on assay sensitivity

A theoretical consideration

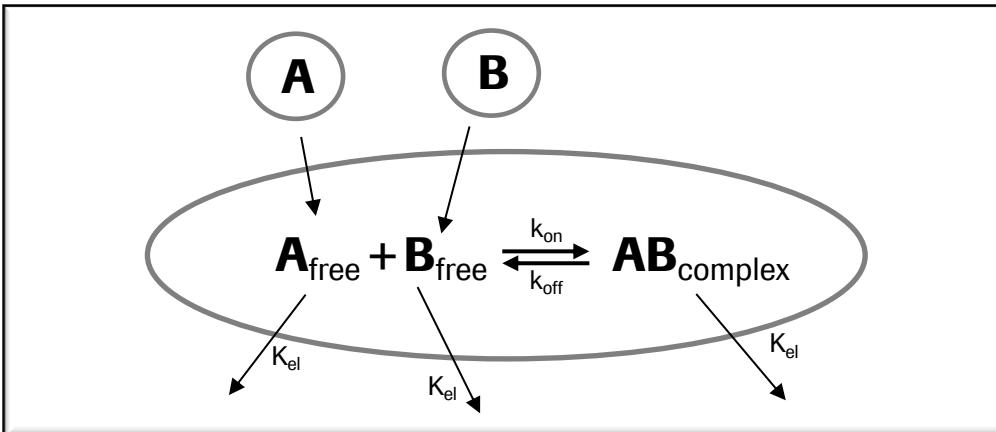


| | | | |
|--------------------------|--|---|----------------------------|
| | „Lower“ affinity ($KD = 10 \text{ nM}$) | $\rightarrow 230 \text{ ng/mL}$ | 2.3 fold lower sensitivity |
| 100-fold lower affinity | Positive control Ab ($KD = 0.1 \text{ nM}$) | $\rightarrow 100 \text{ ng/mL}$ (CP-ARC = 3 pM) | |
| 100-fold higher affinity | „Higher“ affinity ($KD = 0.001 \text{ nM}$) | $\rightarrow 97 \text{ ng/mL}$ | ~ identical |

Calculation of binding interactions

A nice gimmick or valuable for BioA?

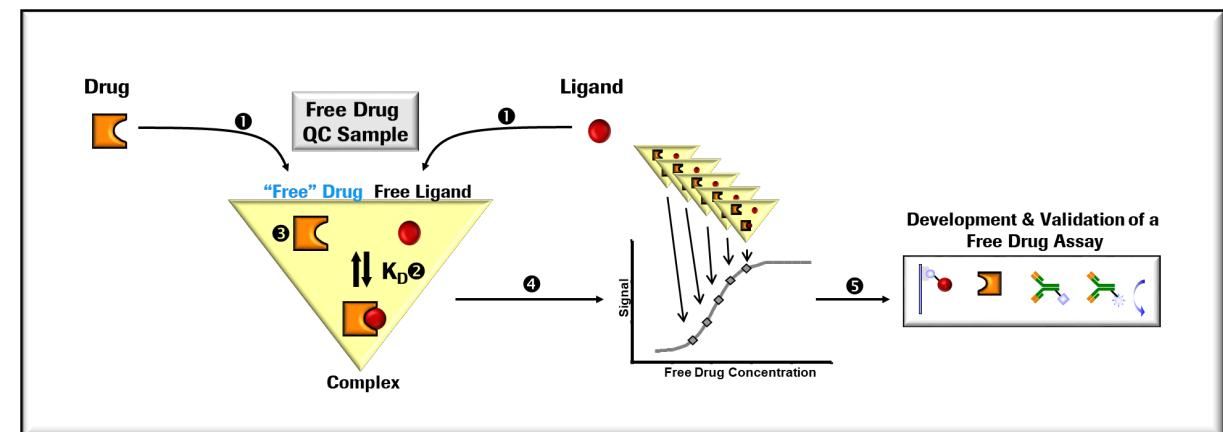
Modelling & Simulation



Chirmule, Jawa, Meibohm. *AAPS J.* 2012 Jun;14(2):296-302.

Gómez-Mantilla, Trocóniz, Parra-Guillén, Garrido. *J Pharmacokinet Pharmacodyn.* 2014 Oct;41(5):523-36
Roskos, Schneider, Vainshtein, Schwikart, Lee, Lu, Faggioni, Liang. *Bioanalysis.* 2011 Mar;3(6):659-75

BioA: „Free Analyte QC Concept“



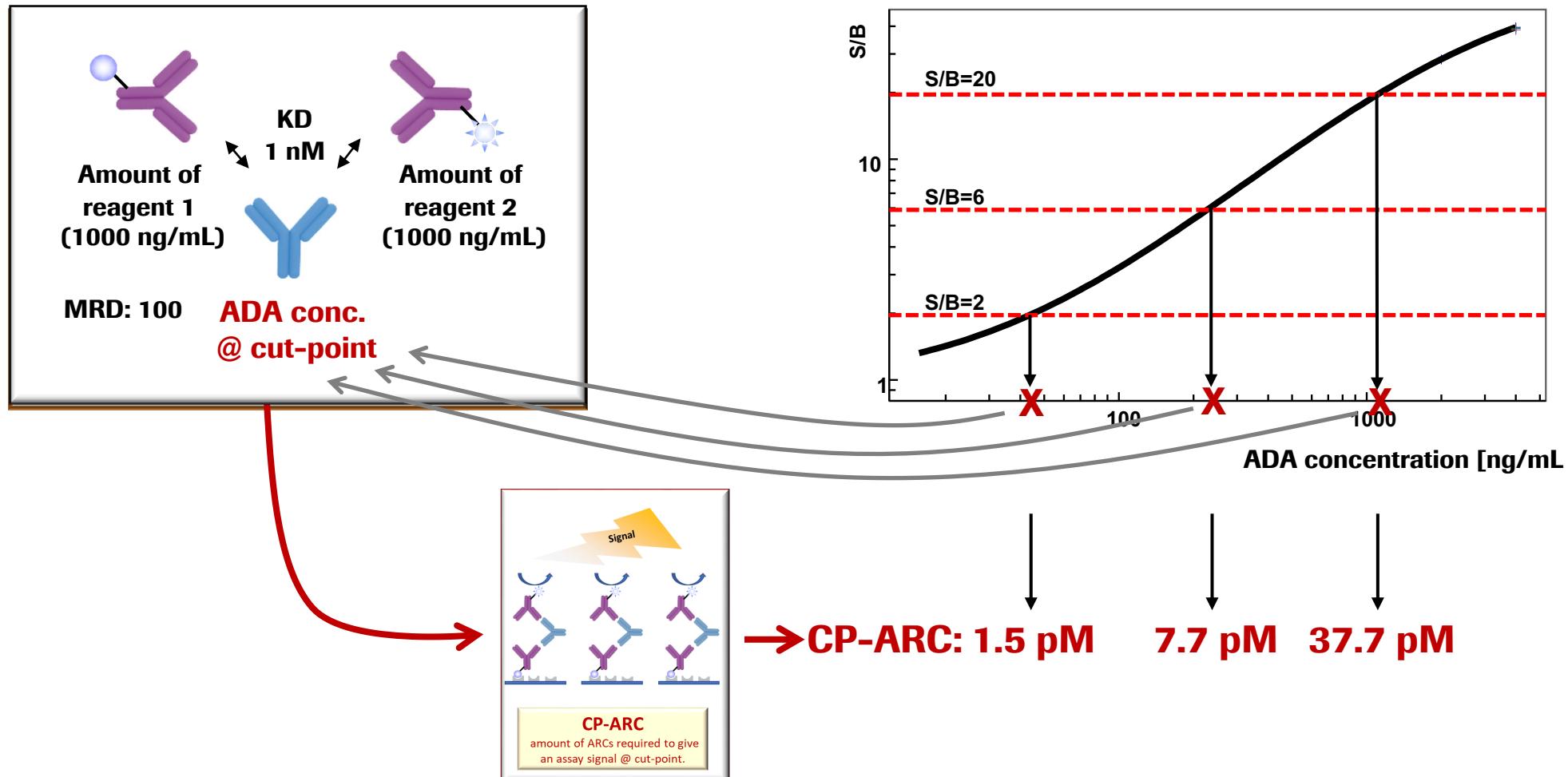
Staack, Jordan, Dahl, Heinrich. *Bioanalysis.* 2014 Feb;6(4):485-96

Schick, Staack, Haak, Jordan, Dahl, Heinrich, Birnboek, Papadimitriou; *Bioanalysis.* 2016 Dec;8(24):2537-2549
Jordan, Onami, Heinrich, Staack; *Bioanalysis.* 2017 Nov;9(21):1705-1717.

➤ **Calculation of protein interactions is a routine task**
...but needs broader implementation into BioA

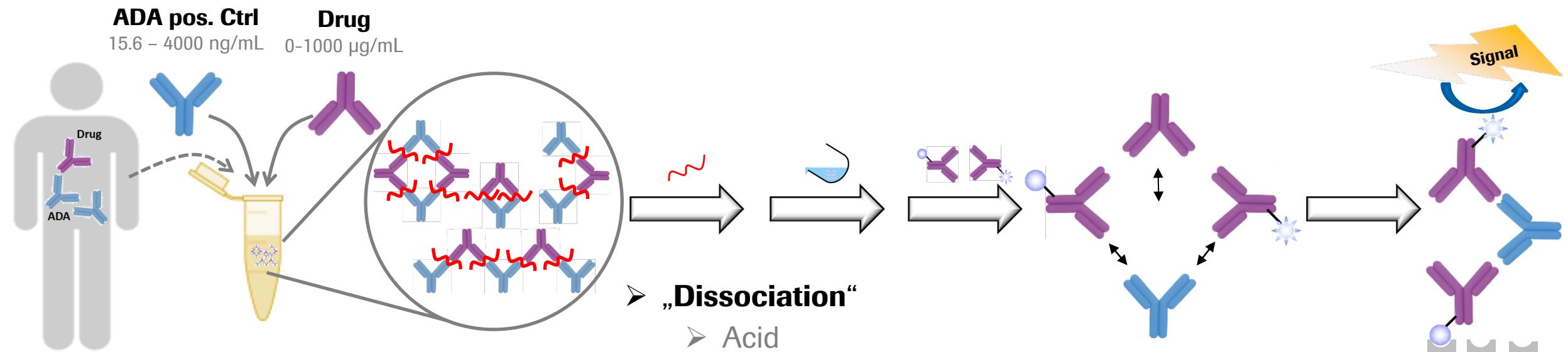
Determination of CP-ARC

→ “real” example



Challenge of „real life“ samples

→ Presence of residual drug



- „Dissociation“

- Acid

Bourdage JS, Cook CA, Farrington DL, Chain JS, Konrad RJ et al. J Immunol Methods. 2007 Oct 31;327(1-2):10-7.
Smith HW, Butterfield A, Sun D. Regul Toxicol Pharmacol. 2007 Dec;49(3):230-7

- HISDA

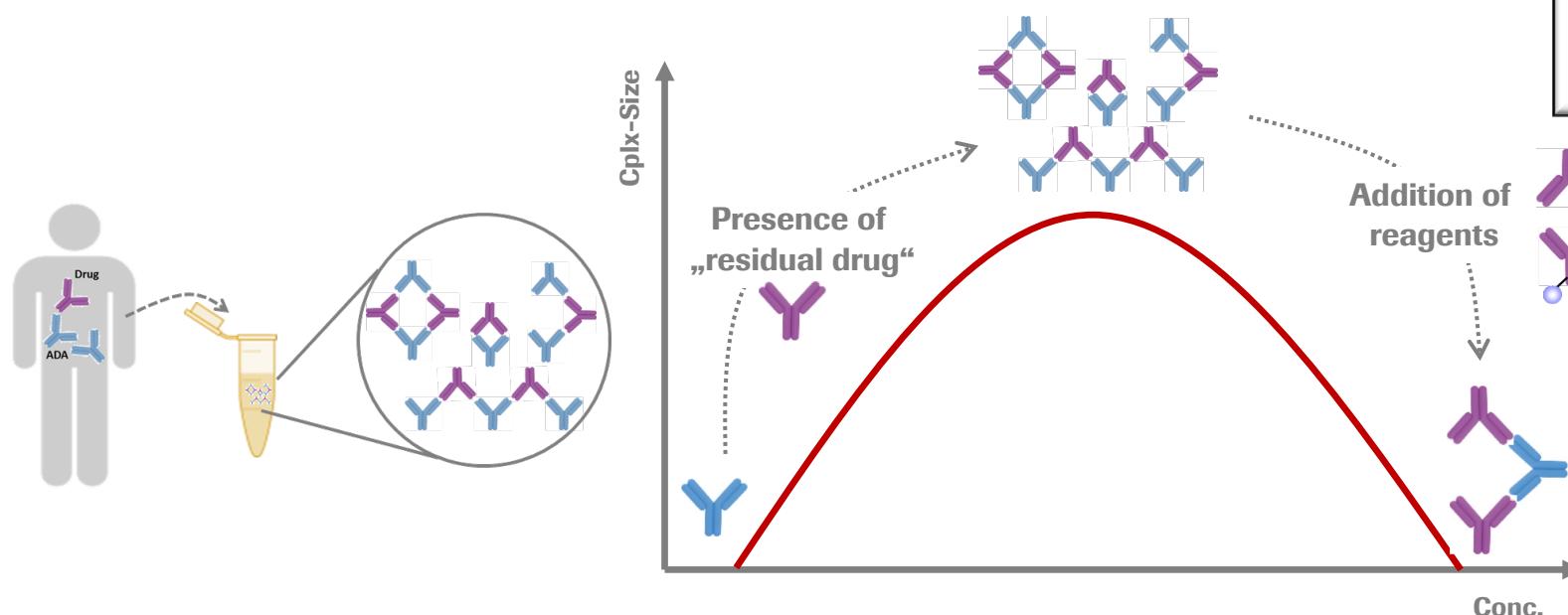
Jordan, Pöhler, Guilhot, Zaspel, Staack. Bioanalysis. 2020 Jun;12(12):857-866 → Presentation Gregor Jordan @ EBF20

- Dilution (MRD 100)

- Reagents = new binding partners
(1000ng/mL, each)

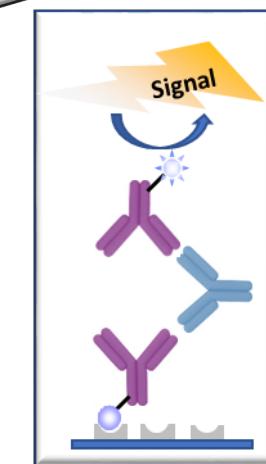
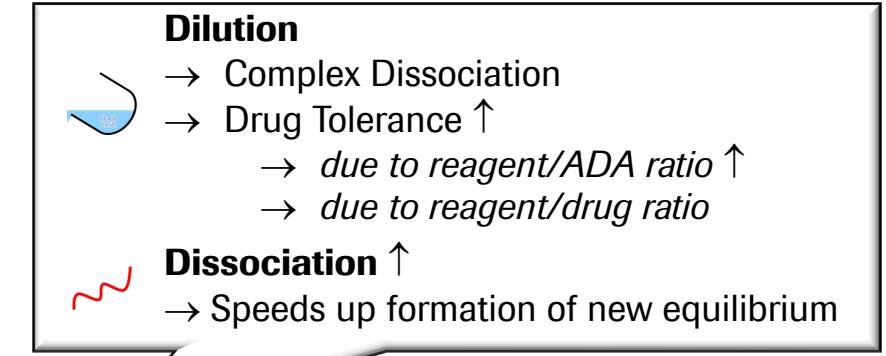
ADA-Analysis – from the ADA in the sample to an assay signal

...nothing than equilibrium shifts?



Binding Partner A = **Constant** concentration → ADA

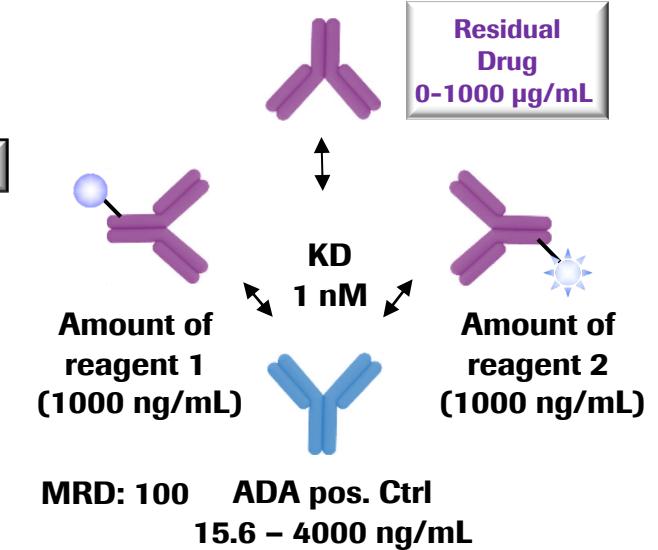
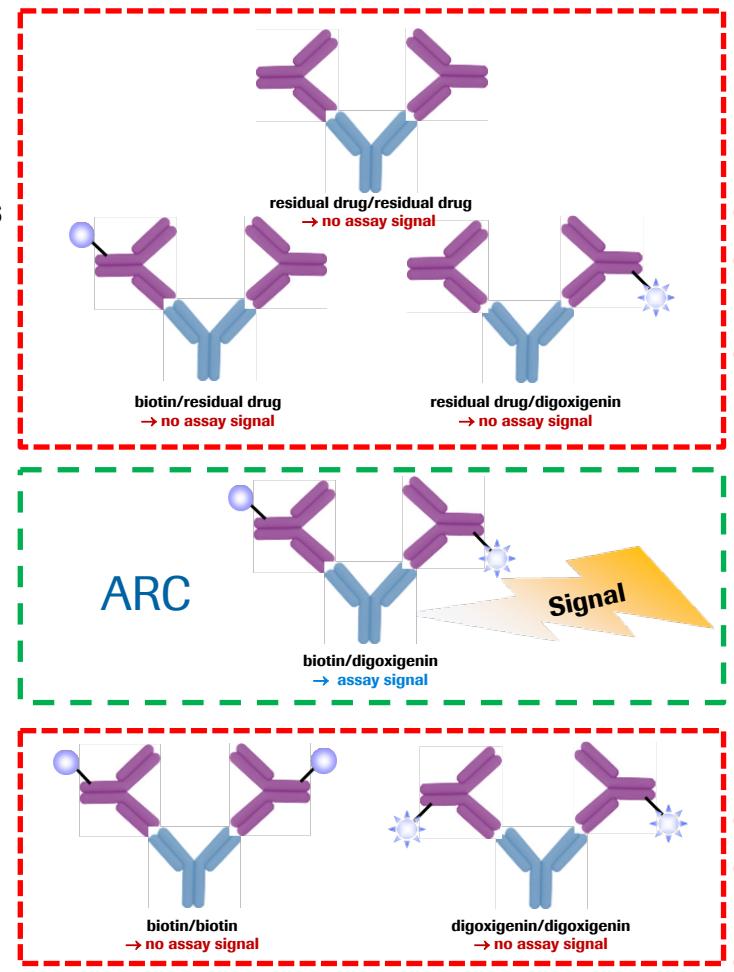
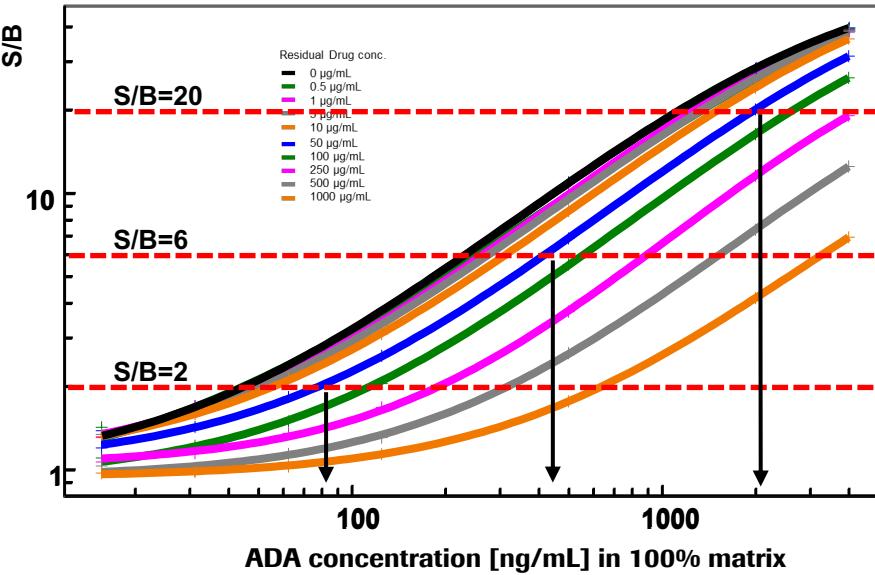
Binding Partner B = **Increasing** concentration → Drug & Reagents



ADA Analysis @ presence of residual drug

„Reduced Drug Tolerance-Effect“

- Formation of additional non-signal generating complexes

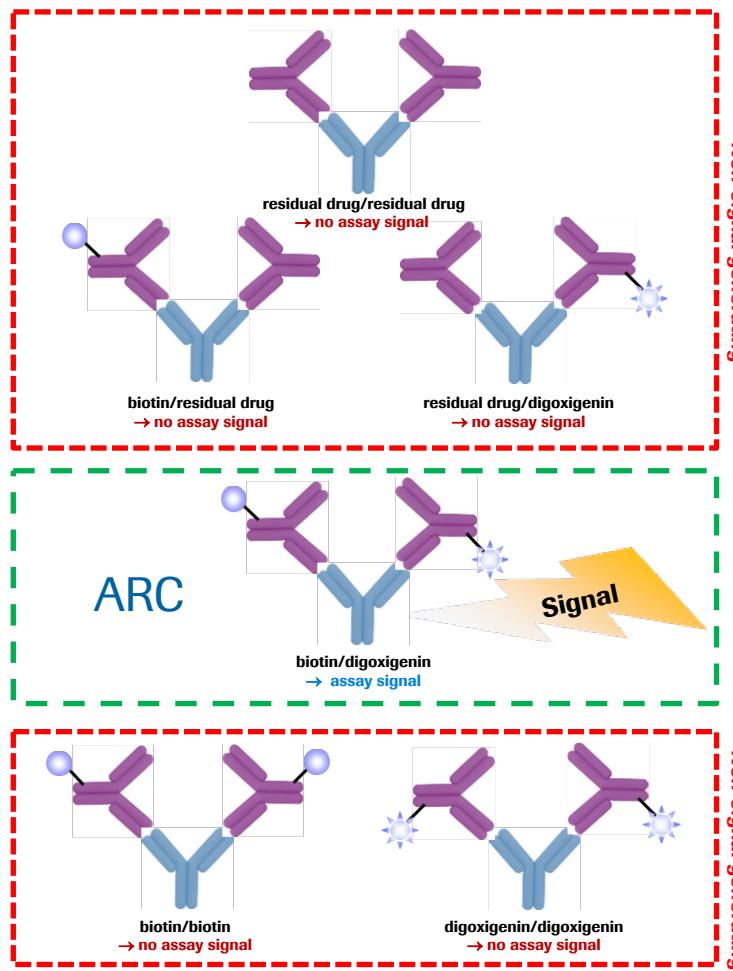
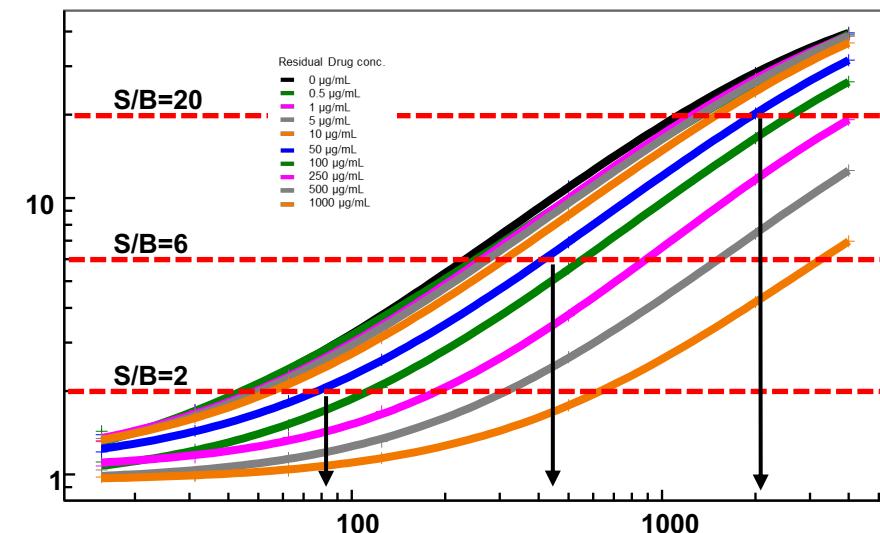


ADA Analysis @ presence of residual drug

Better understanding of drug tolerance performance thanks to CP-ARC?

Analytical experiment:

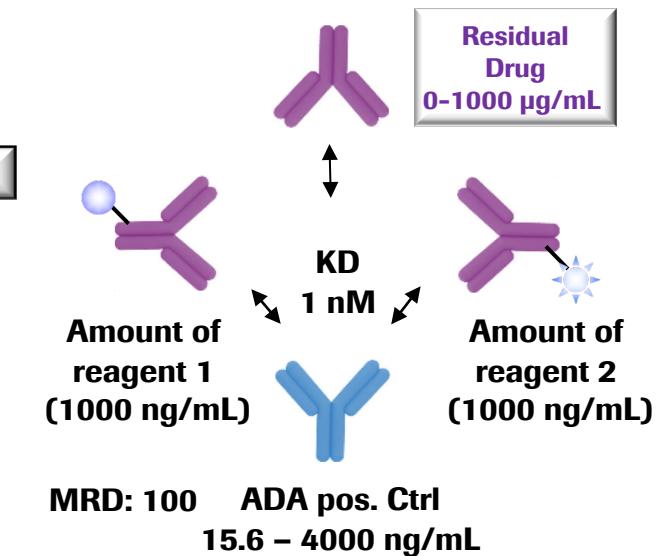
Determination of **ADA conc.** that generated a **pos. assay signal @ S/B 2, 6 & 20** in the presence of increasing residual drug concentrations



Calculation:

ADA conc. required to form sufficient amounts of **CP-ARCs @ S/B 2, 6 & 20** (1.5, 7.7, 37 pM)

in the presence of increasing residual drug concentrations



Prediction of „real“ assay performance by CP-ARC?

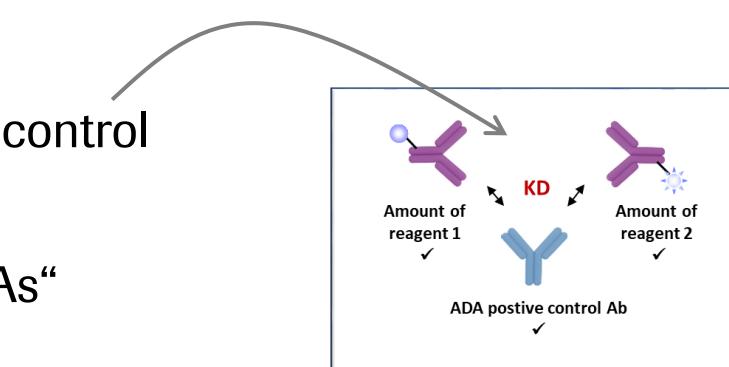


Summary & Conclusion

- Example **confirms the CP-ARC hypothesis**
 - knowledge of assay CP-ARC can predict assay performance
- **CP-ARC = constant assay parameter that defines the sensitivity and drug tolerance of a given assay**
 - Sensitivity: clear understanding of assay sensitivity independent of the positive control
 - FDA 2019 : „high affinity positive controls may overestimate the sensitivity of the assay...“

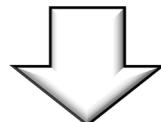
What do we need to reach goal?

- **A joint effort of the entire BioA community from assay development to regulators!**
- **Characterisation of binding properties** of ADA positive control
- Expansion of the concept on other assay formats
- Better understanding of the binding properties of „real ADAs“



Benefit for the BA community

- Application of CP-ARC concept enables **comparison of ADA-Assay performance/results**



➤ Improved ADA reporting

- No „inconclusive“ results, since the assay performance is fully understood
- **Clear statement what ADA concentrations are detectable**
- *Example: <100 ng/ml ADA with an affinity of xyz nM*

- **BioA contribution to enable valid comparison of „clinical immunogenicity“ of different compounds**



Acknowledgement

Gregor Jordan & Team

Eugenia Hoffmann

Kay Stubenrauch

Julia Heinrich

Thomas Singer

All colleagues at Bioanalytical R&D, Roche pRED

@ Roche Innovation Center Munich & Roche Innovation Center Basel

Doing now what patients need next