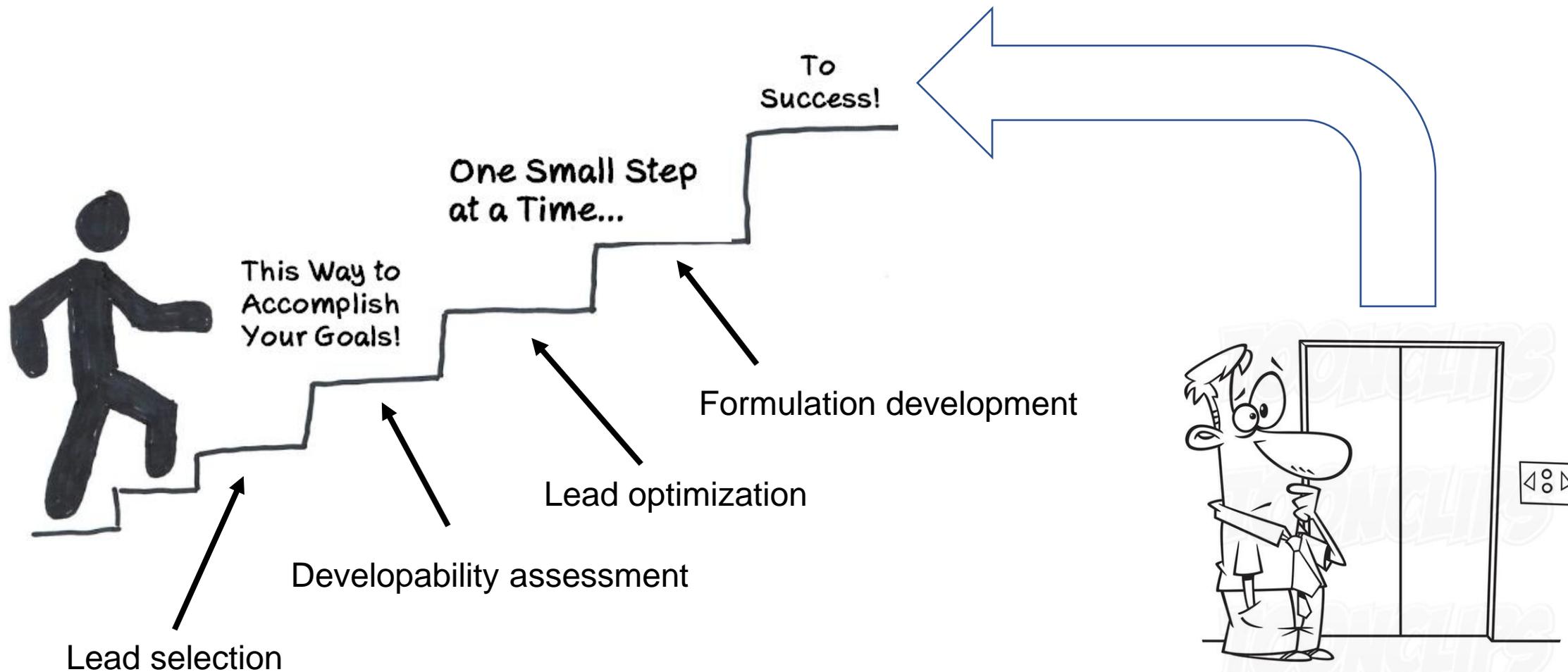


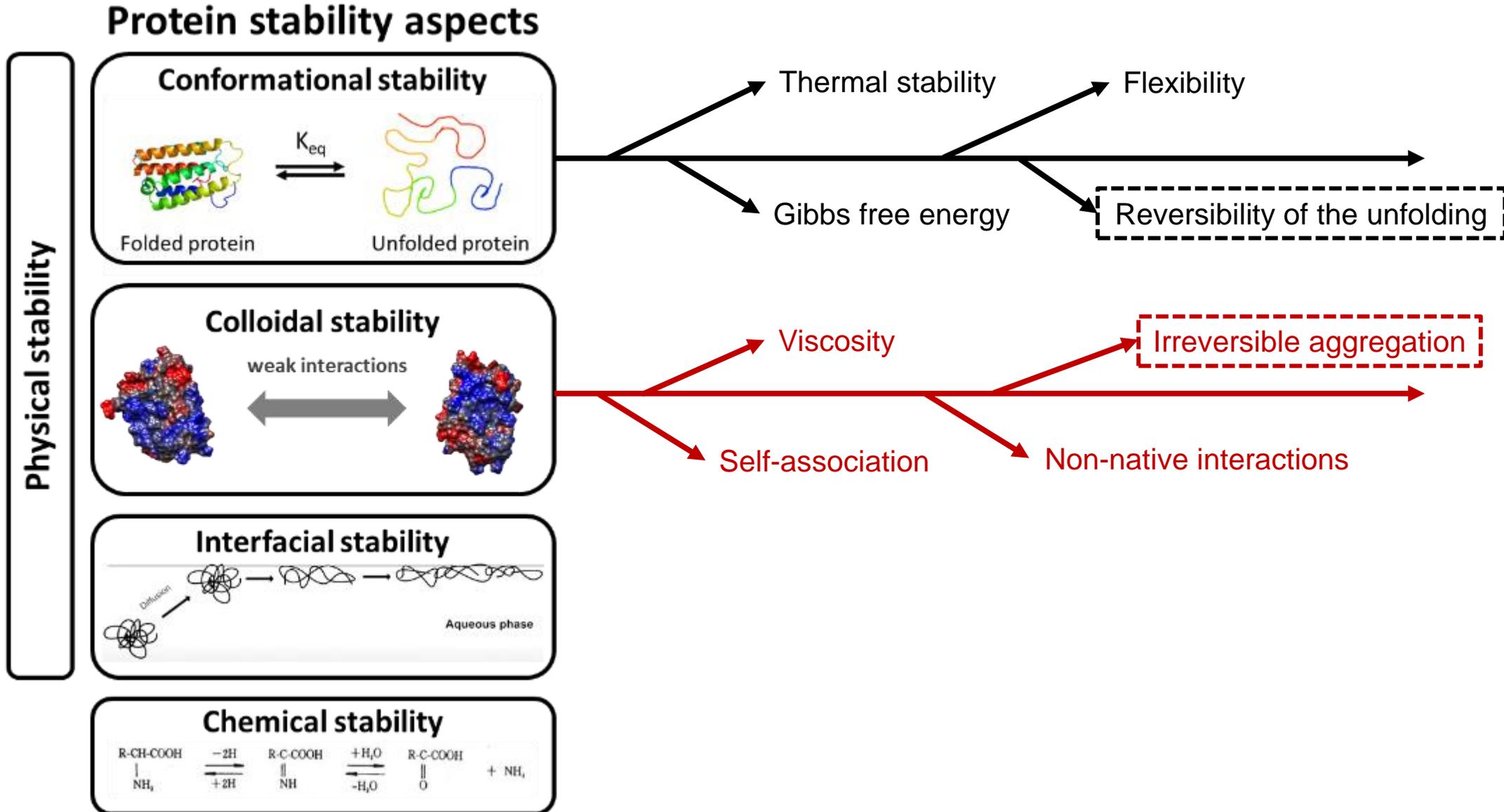
Assessing refoldability to select therapeutic proteins and formulations with lower aggregation propensity during storage

Dr. Hristo L. Svilenov
Technische Universität München

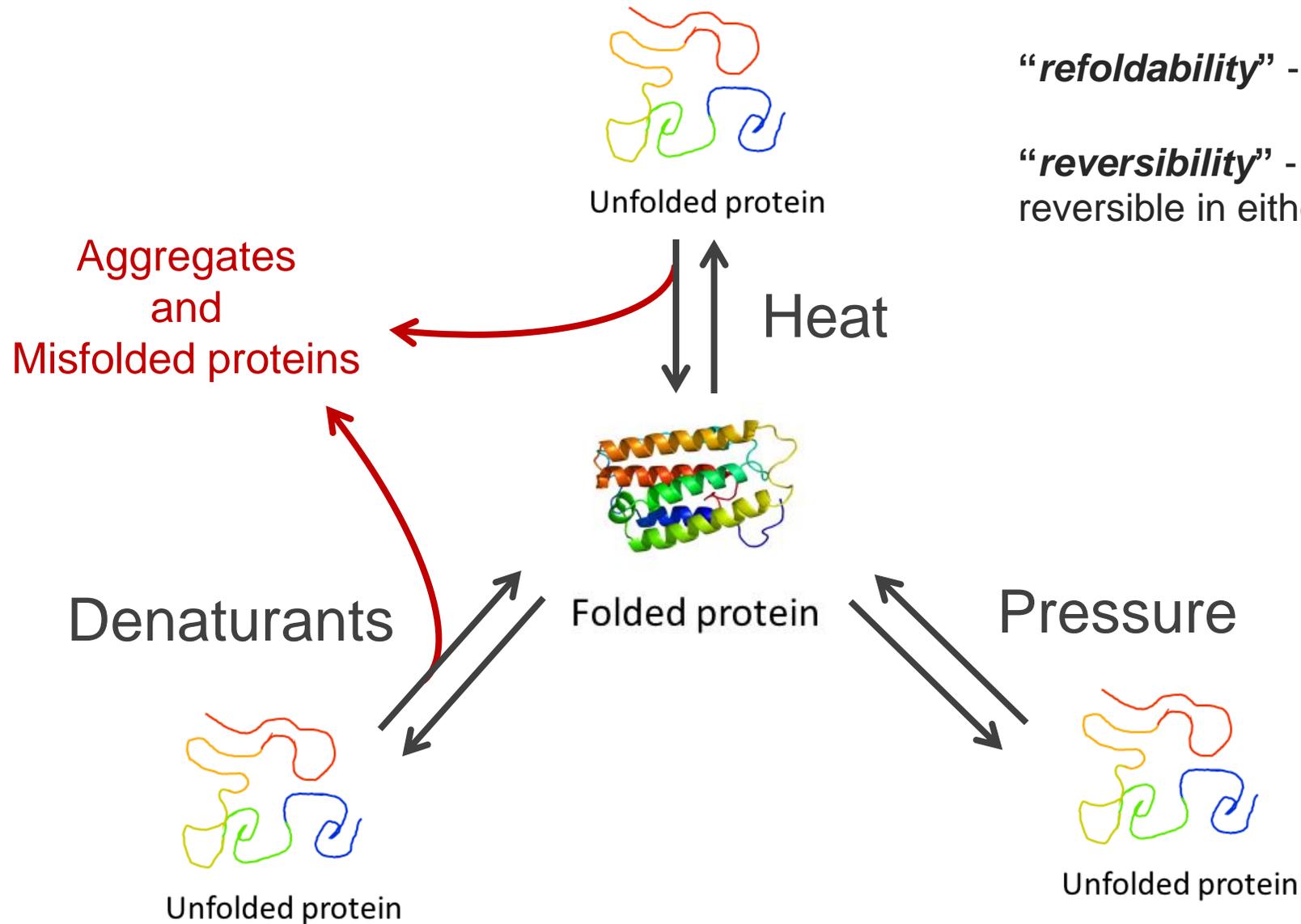
The staircase of antibody development



Getting a broad perspective on protein stability



Approaches to unfold proteins and study unfolding reversibility



“**refoldability**” - the ability to be refolded

“**reversibility**” - the quality of being reversible in either direction

Why bother about refoldability?

> Nat Biotechnol. 2004

Aggregation-resistant domain antibodies selected on phage by heat denaturation

Laurent Jespers, Oliver Schon, Kristoffer Famm, Greg Winter

> J Mol Biol. 2008

Thermodynamically stable aggregation-resistant antibody domains through directed evolution

Kristoffer Famm¹, Lars Hansen, Daniel Christ, Greg Winter

“This process appeared to select for domains with denatured states that resisted aggregation, but the domains only had low free energies of folding...”

- Unfolding reversibility is related to low aggregation propensity of antibody domains
- However, many therapeutically relevant proteins do not unfold reversibly

Consecutive heating scans on an aggregation-resistant VH

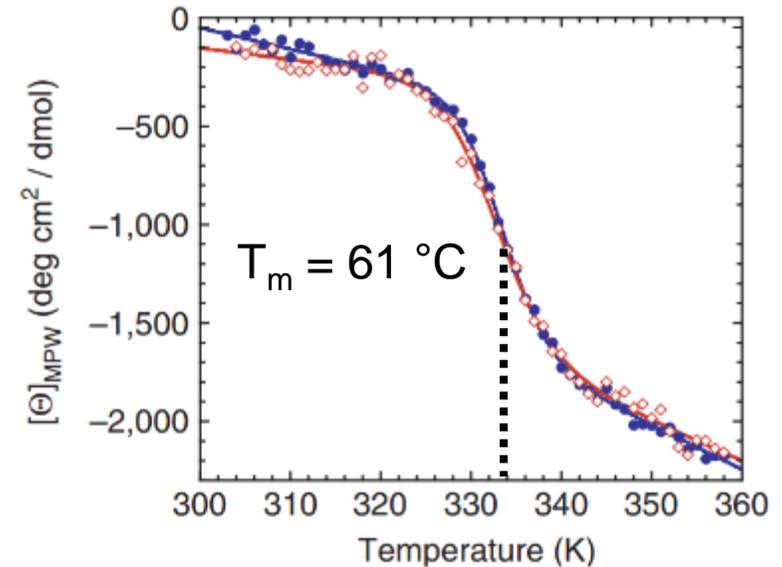
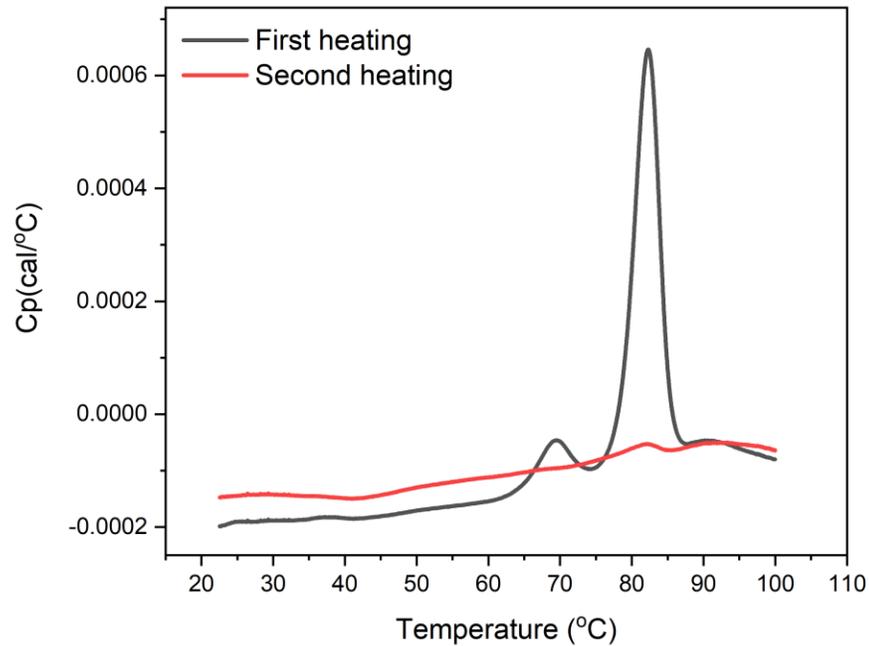


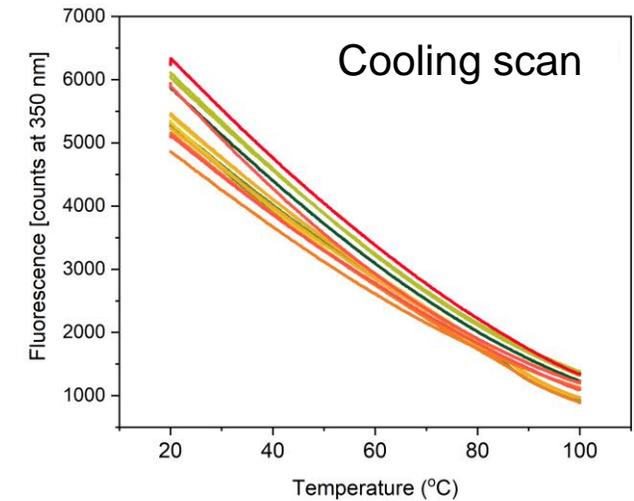
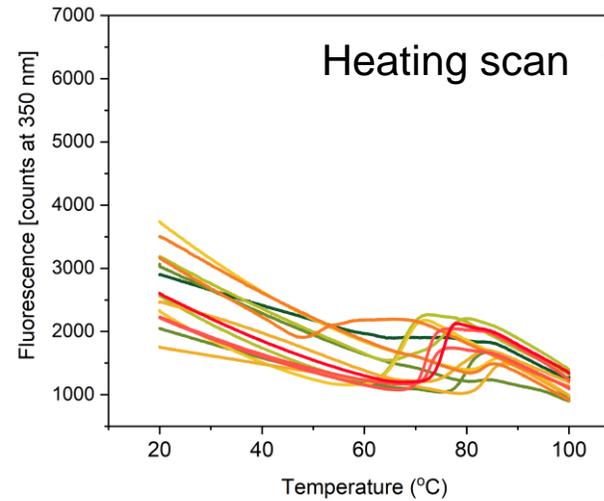
Image adapted from Jespers et al.

The traditional approach to test for reversibility after heating

Consecutive heating scans on a mAb



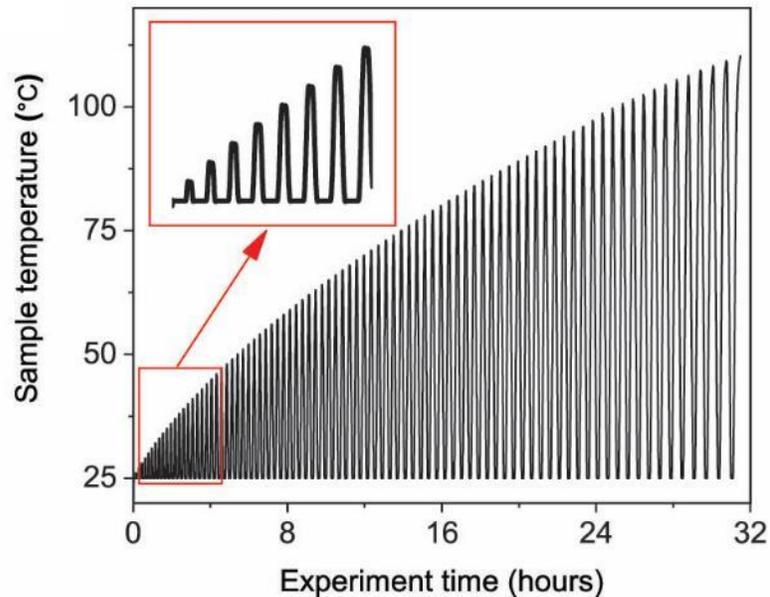
Heating and cooling scans on 14 different proteins



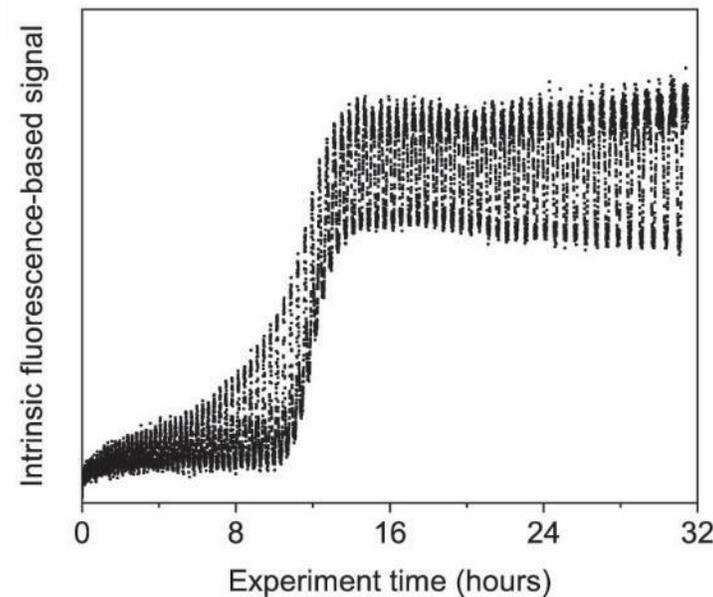
- Rapid aggregation at high temperatures → zero unfolding reversibility
- Overheating masks unfolding reversibility differences

Another approach – use incremental heating and cooling cycles

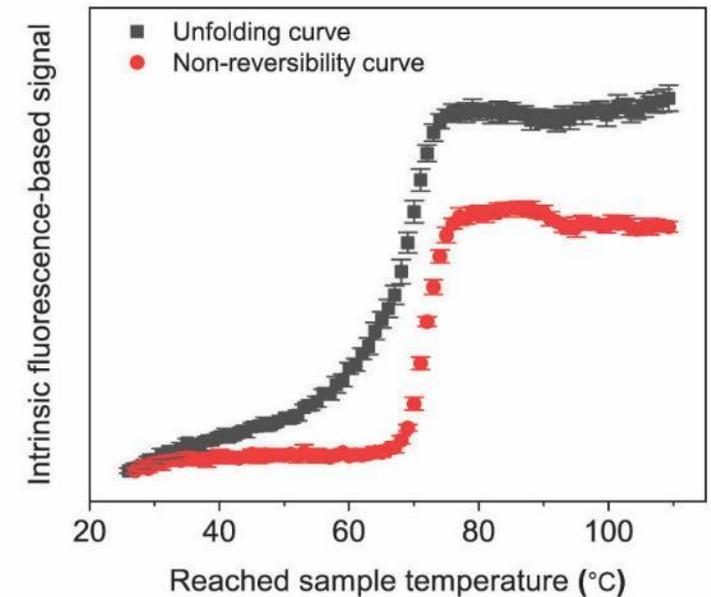
Heating program



Raw data



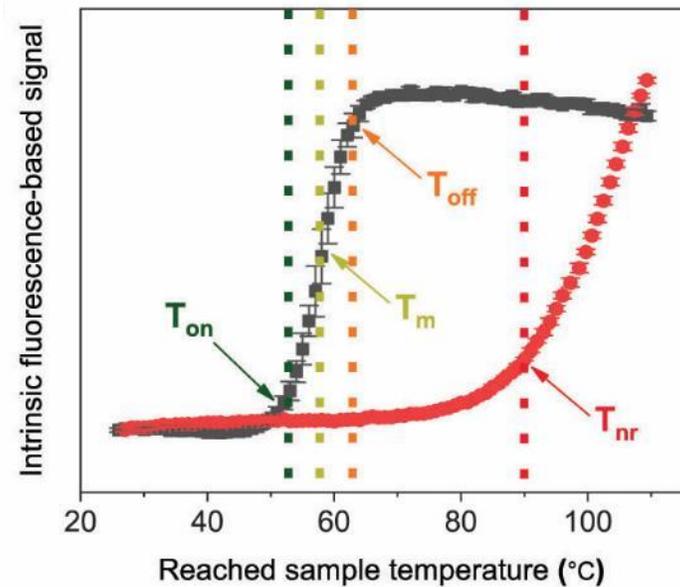
Deconvoluted data



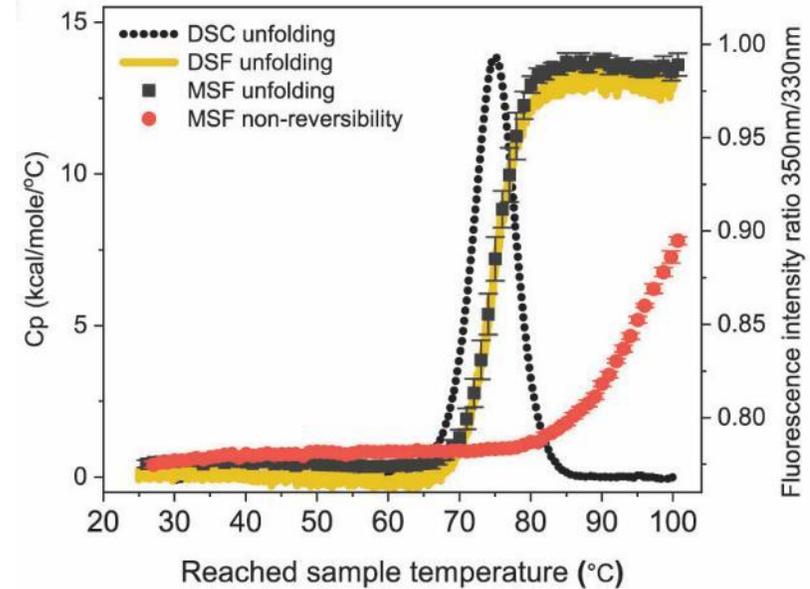
- We call it modulated scanning fluorimetry (MSF)
- The MSF analyser software from Klaus Richter deconvolutes the raw data
- Distinct unfolding and non-reversibility curves are obtained

Information obtained with modulated scanning fluorimetry

Parameters derived from the MSF experiment

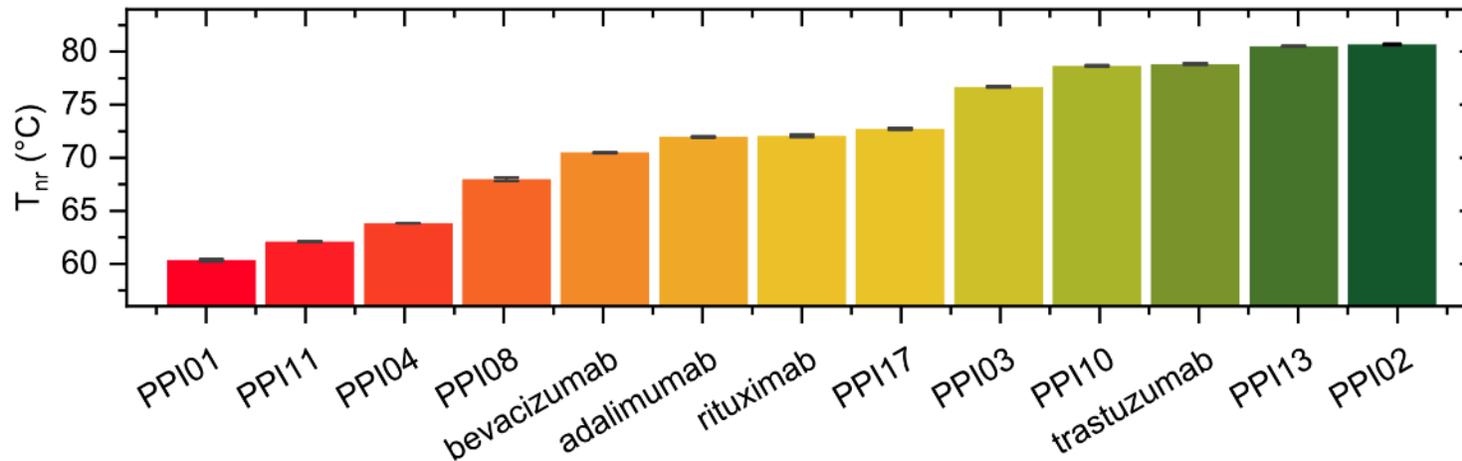
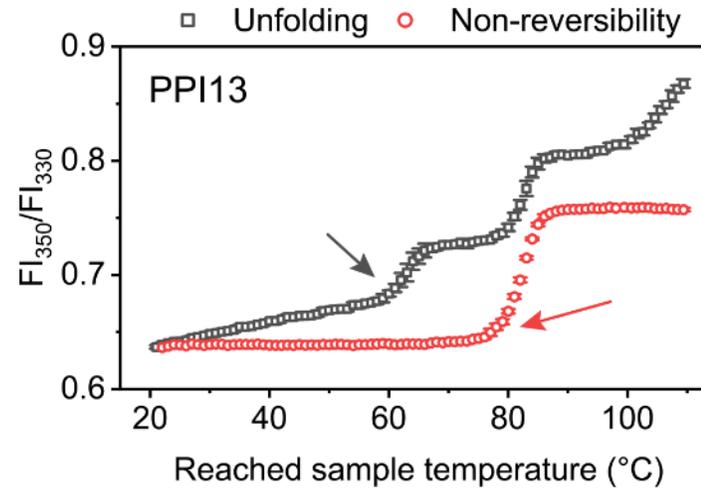
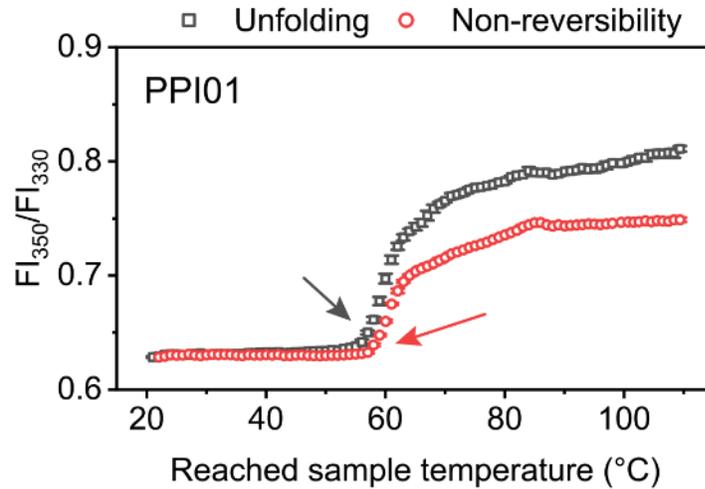


Overlay of thermal denaturation data from different methods



- Non-reversibility onset temperature (T_{nr}) can be obtained with MSF
- The non-reversibility curve provides orthogonal data to DSC and DSF
- Applicable to various proteins and concentrations

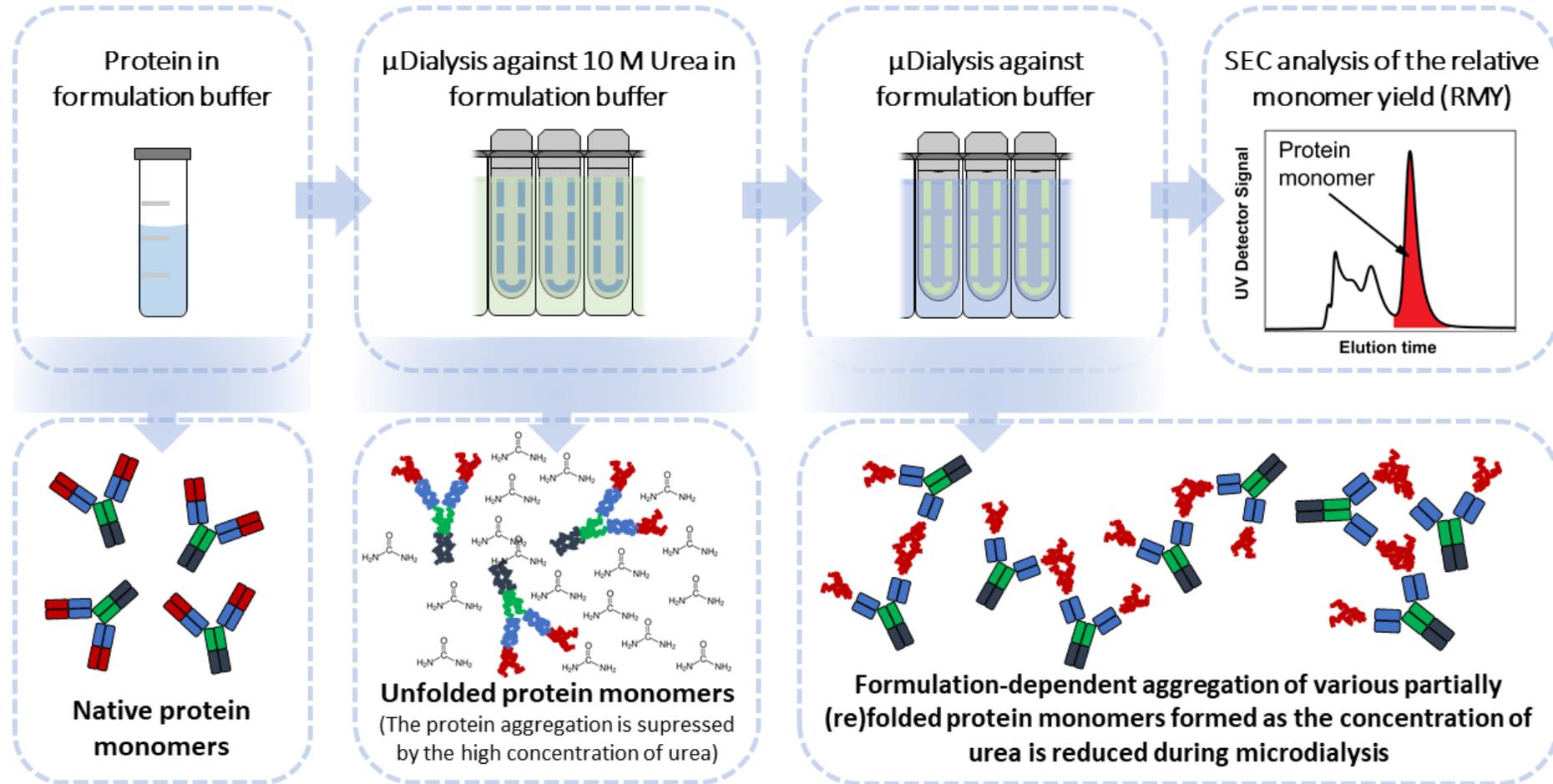
Using MSF to study and rank antibody candidates



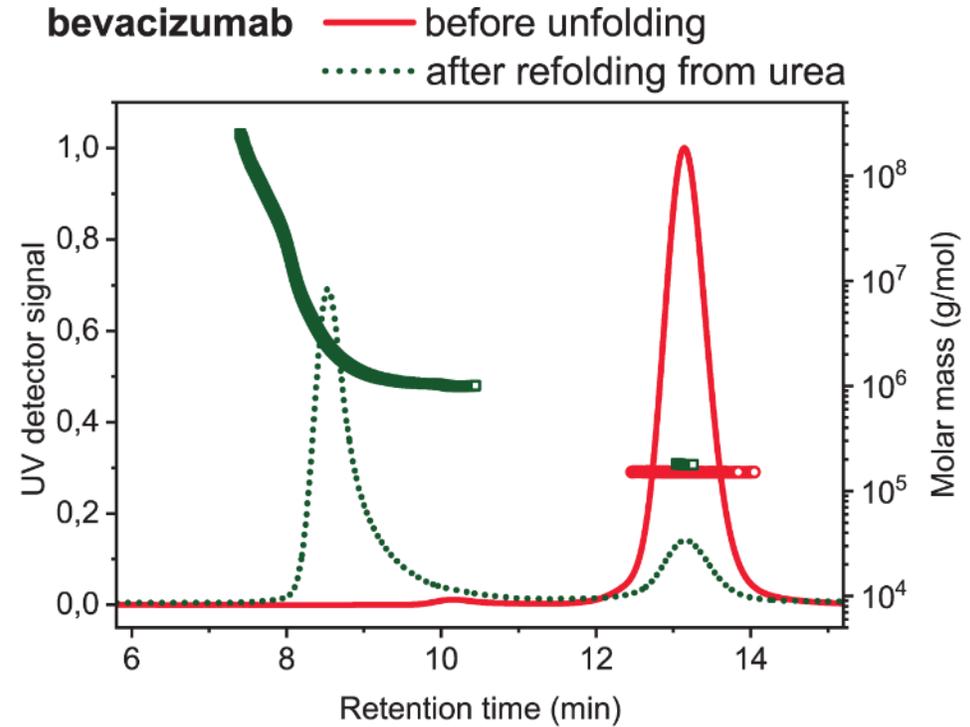
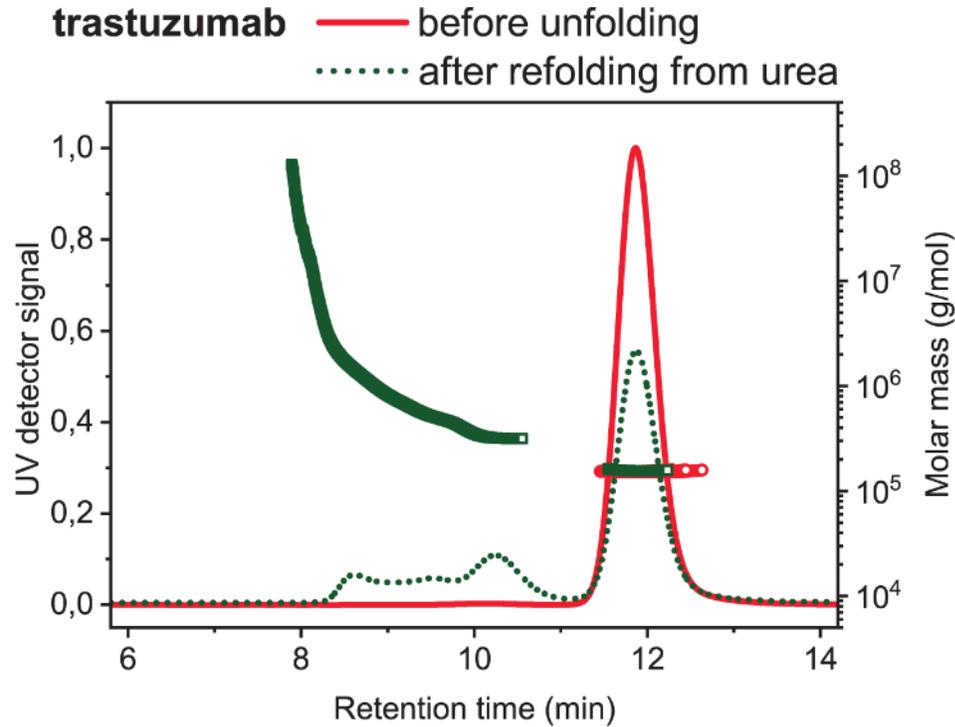
- Unique unfolding and non-reversibility traces are obtained for different antibodies
- Very different non-reversibility onset temperature (T_{nr}) of the candidates

Studying refoldability after unfolding with chemical denaturants

Schematic diagram of the ReFOLD assay

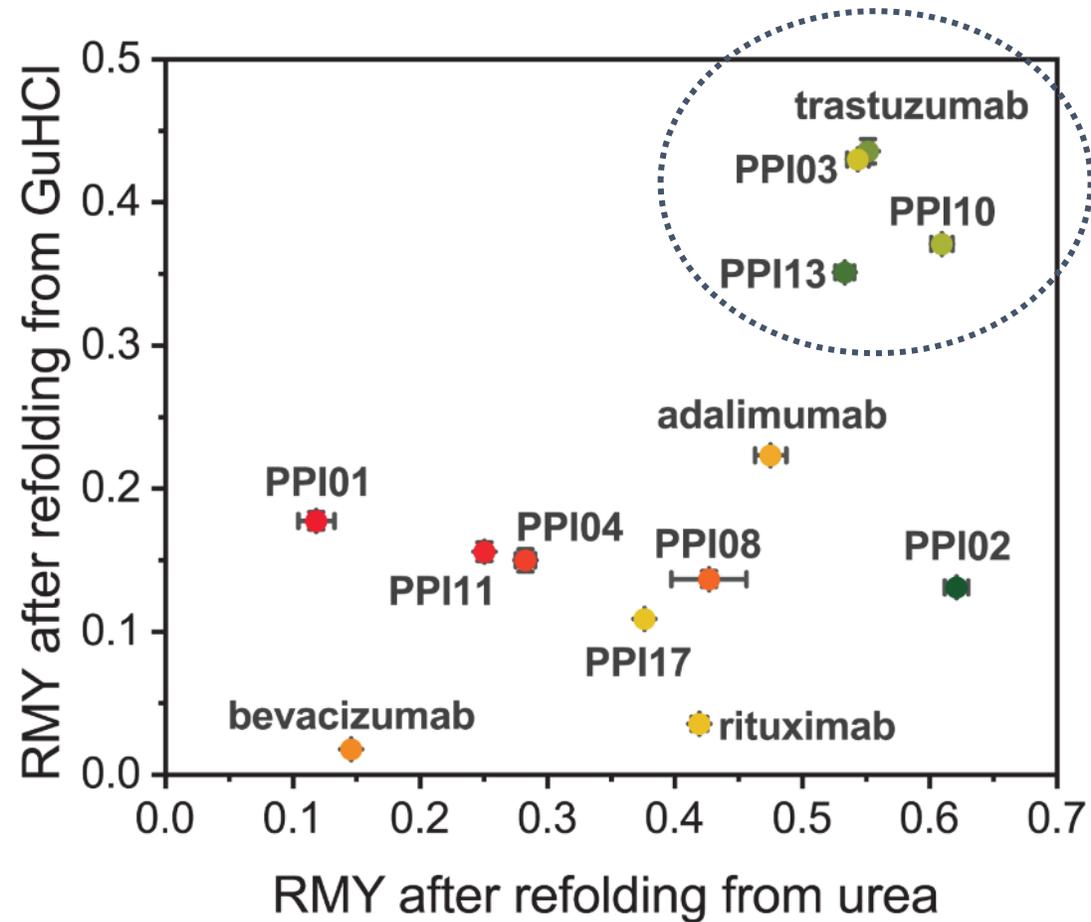


SEC-MALS analysis on native and refolded mAbs



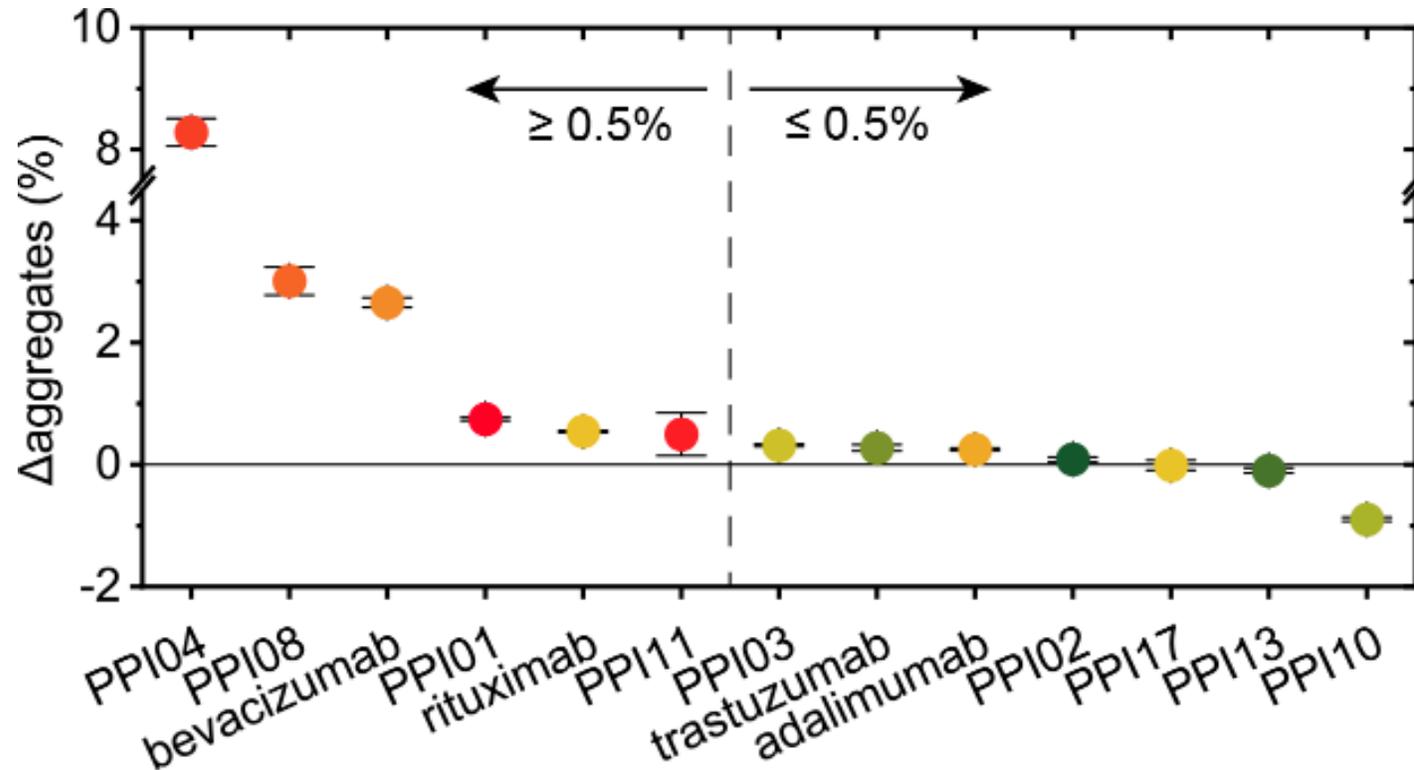
- Antibodies have different relative monomer yield after refolding from denaturants
- Aggregate distribution after refolding is protein-specific

RMY after unfolding with chemical denaturants to rank antibodies



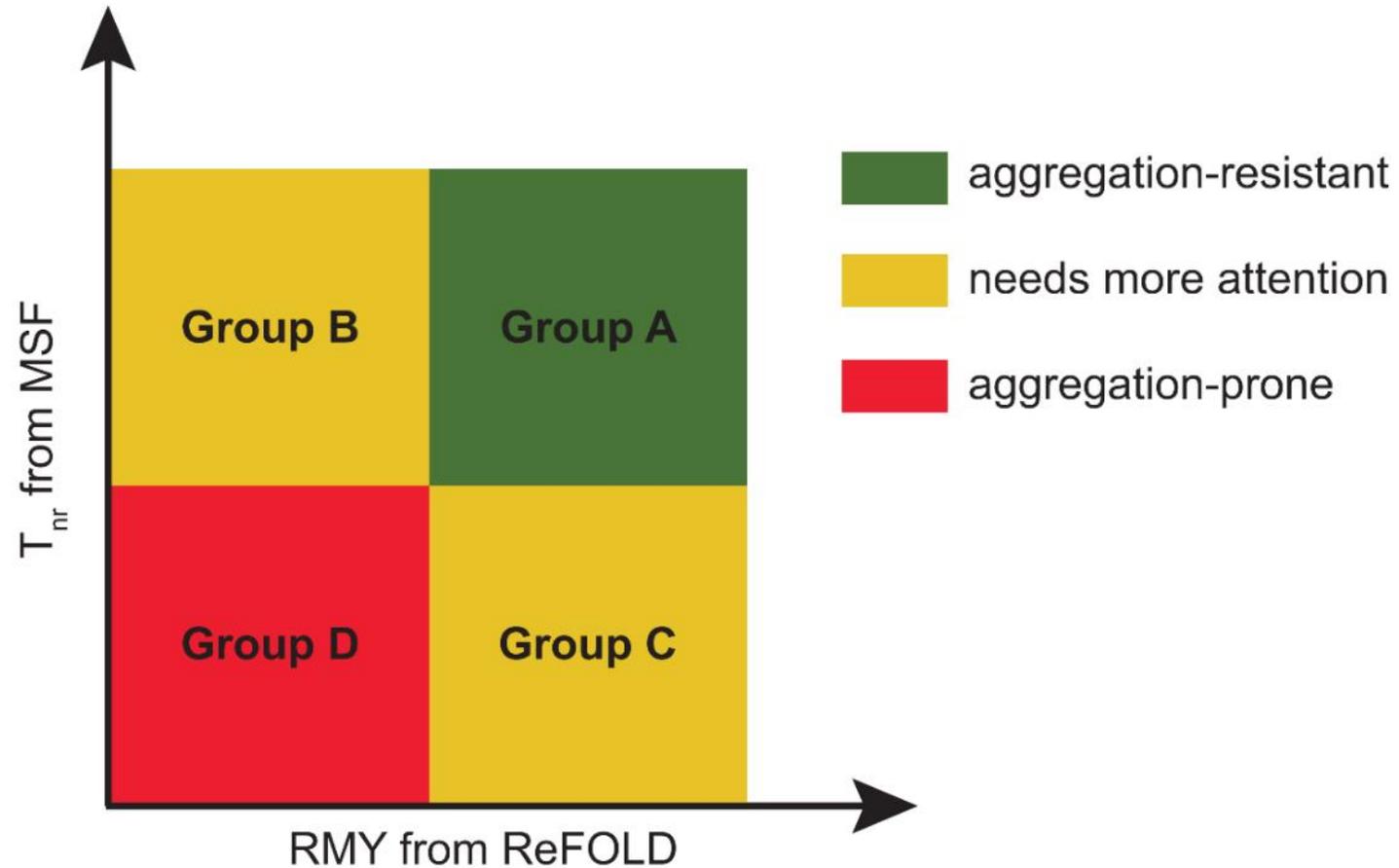
- Some antibodies exhibit high RMY after refolding from either urea or GuHCl

Aggregates formed by the antibodies during storage at 40 °C



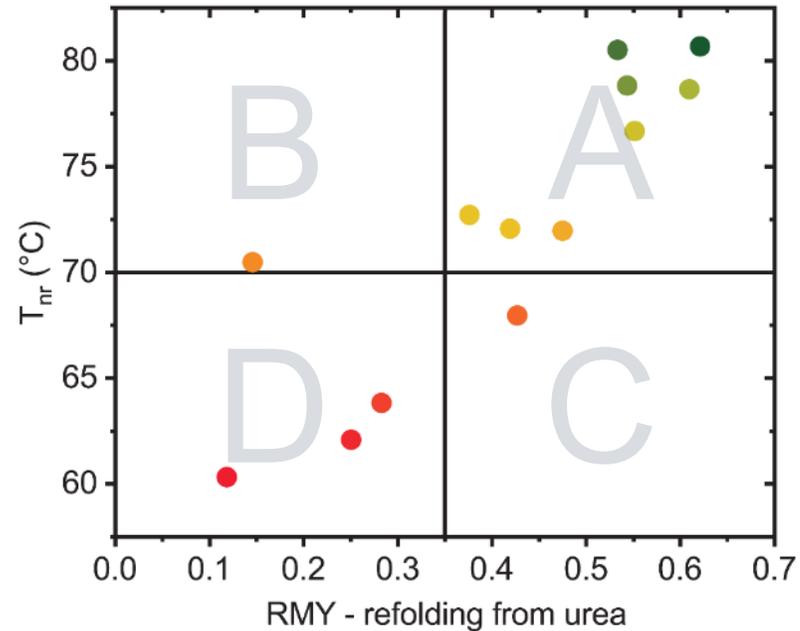
- The increase in aggregates was analyzed with size-exclusion chromatography
- The antibodies exhibit very different aggregation during storage for 3 months at 40 °C
- Only PPI11 aggregated at 4 °C (ca. 1.5% aggregates after 12 months)

Classifying proteins and formulations based on MSF and ReFOLD

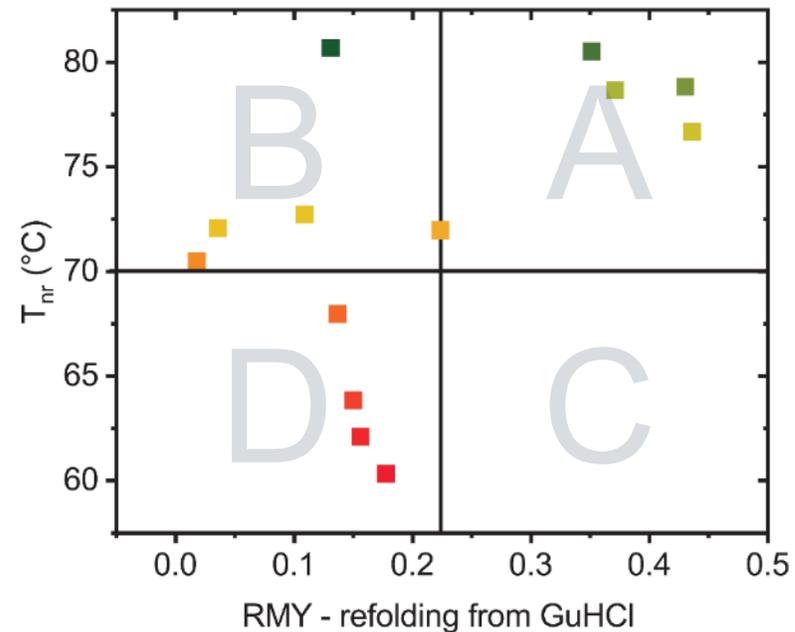


Classifying proteins and formulations based on MSF and ReFOLD

Experiment with urea



Experiment with GuHCl



- Aggregation-resistant antibodies cluster in Group A
- Antibodies that aggregated during storage cluster in Group D

Wrap up and take-home messages

- **Two approaches to study protein refoldability were presented**
- **MSF indicates what temperatures cause non-reversible structural changes**
- **ReFOLD gives you the fraction of protein that remains monomeric after refolding from denaturants**
- **The two approaches are complementary**
- **Ideally an aggregation-resistant antibody will have two properties:**
 - 1. High temperature of non-reversibility onset**
 - 2. High relative monomer yield after refolding from chemical denaturants**

Acknowledgements

Gerhard Winter

Tim Menzen

Klaus Richter

Carolin Berner

Pernille Harris

Robin Curtis

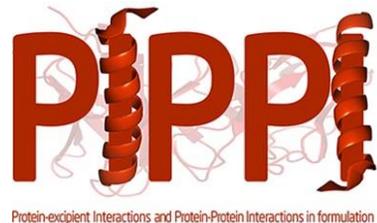
Sasha Golovanov

Matja Zalar

Dierk Roessner

Lorenzo Gentiluomo

Funding for some of the presented work:



All other colleagues in PIPPI