

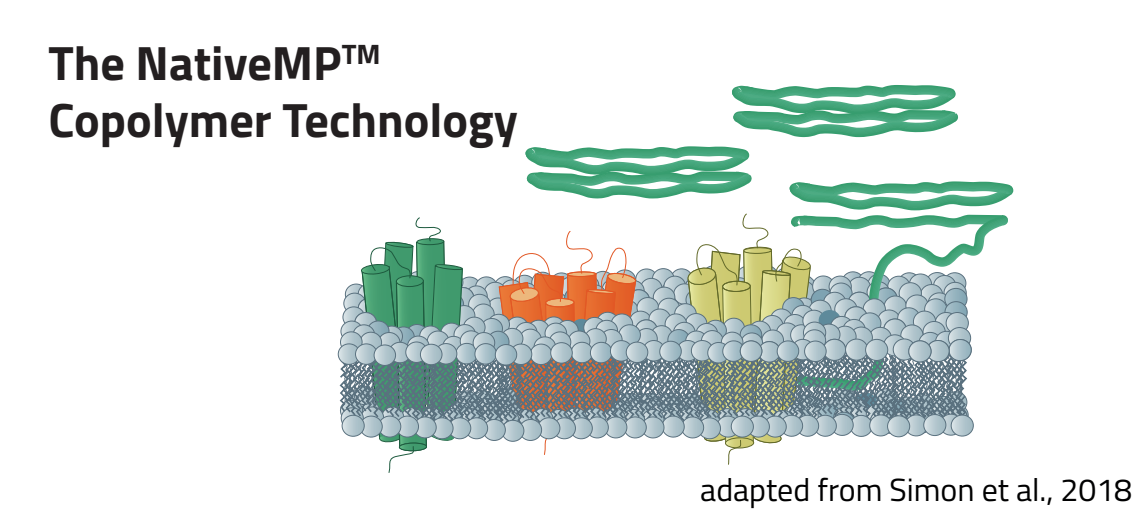
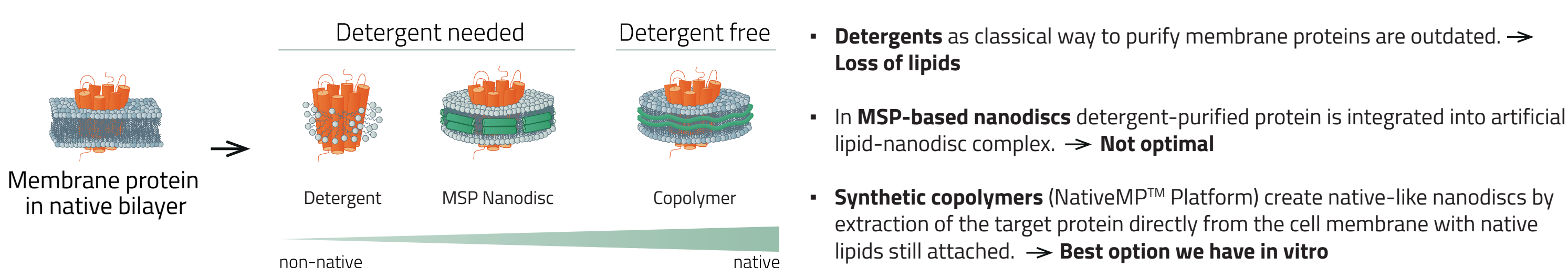
NativeMP™ Platform

Next Generation Copolymers for the Characterization of Membrane Proteins in Near-Native Conditions

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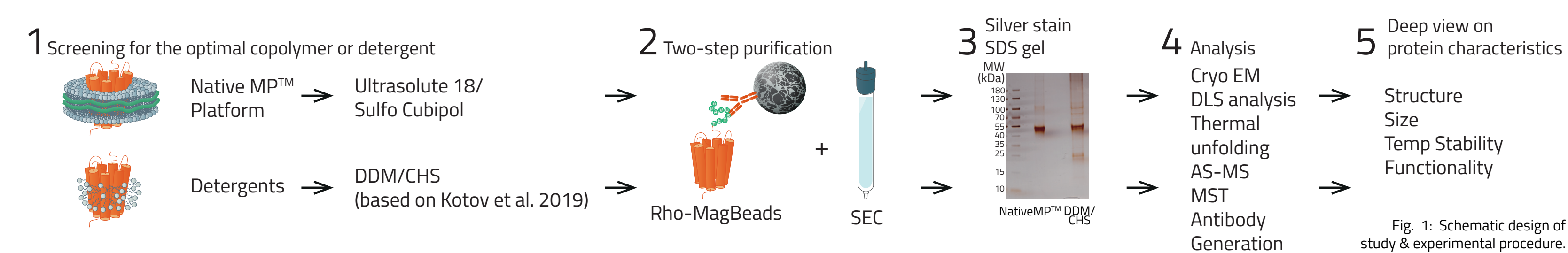
Abstract: The study of membrane proteins has been revolutionized by the development of next-generation copolymers, offering profound advancements in purification and structural characterization. Membrane proteins, particularly G protein-coupled receptors (GPCRs) and ion channels, are critical for cellular signaling and homeostasis, and represent approximately 60% of current drug targets. Research on these proteins is essential to combat a broad range of diseases, including cardiovascular disorders, neurological conditions, and cancers. Traditional detergents, while useful, often strip away essential lipids, leading to protein instability and loss of function. In contrast, copolymers utilized in Cube Biotech's unique NativeMP™ platform maintain a lipid-rich environment that preserves the native state of these proteins. We present a comprehensive workflow, from gene to structure, utilizing next-generation polymers for full-length membrane protein purification. We demonstrate the efficacy of this approach with a case study on an ion channel, showcasing the retention of functional lipids and the maintenance of protein activity throughout the purification process. The introduction highlights the pivotal role of lipids in membrane protein folding and function, as corroborated by studies from Dorr et al. (2016), Bersch et al. (2017), and Smirnova et al. (2020). Our methodology starts with gene cloning and expression in a suitable host, followed by solubilization with our NativeMP™ platform. These copolymers enable the extraction of membrane proteins in a lipid environment that closely mimics their native conditions. Subsequent steps include purification, functional assays, and structural analysis using cryo-electron microscopy (cryo-EM). Results reveal that proteins retain their functional integrity and are amenable to high-resolution structural determination. This approach represents a significant leap forward in membrane protein research, offering a viable alternative to detergent-based methods. By preserving the lipid milieu, next-generation copolymers facilitate the study of membrane proteins in a state that closely resembles their physiological context. This enables more accurate insights into their mechanisms and interactions, paving the way for advancements in drug discovery. Our findings underscore the potential of copolymer-based techniques to transform the landscape of membrane protein characterization and functional analysis, thereby enhancing our ability to develop targeted therapies for a wide range of diseases.

1. Introduction Part One Stabilizing Membrane Proteins

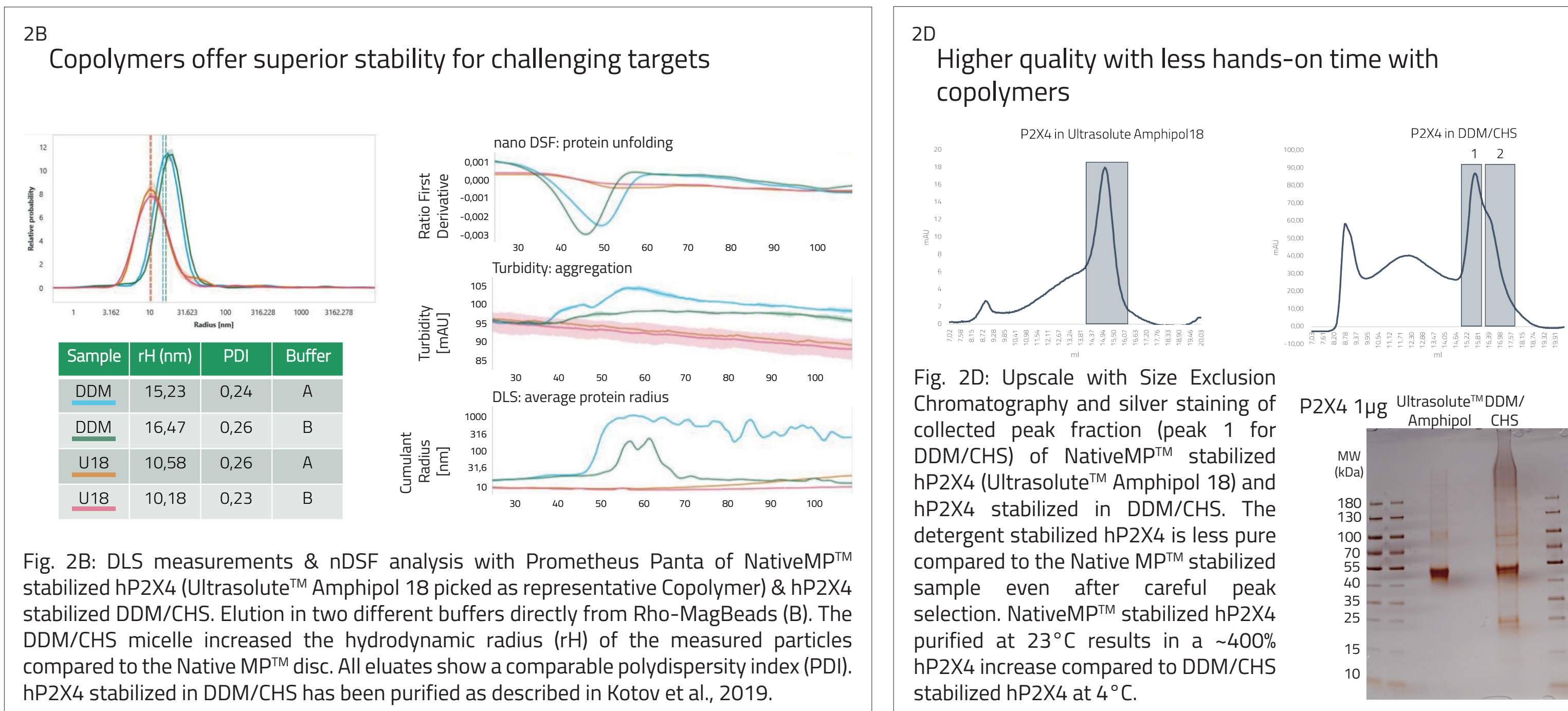
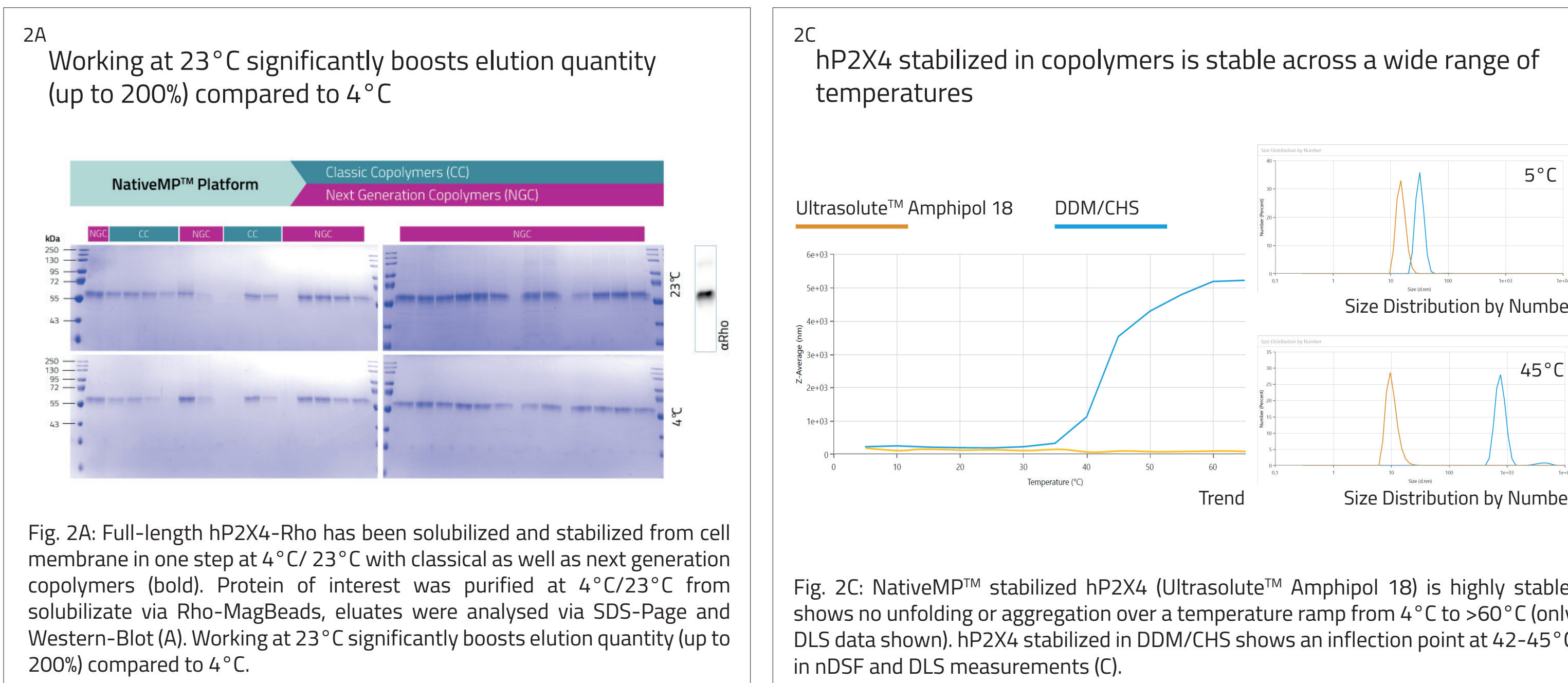


1. Introduction Part Two Our Workflow

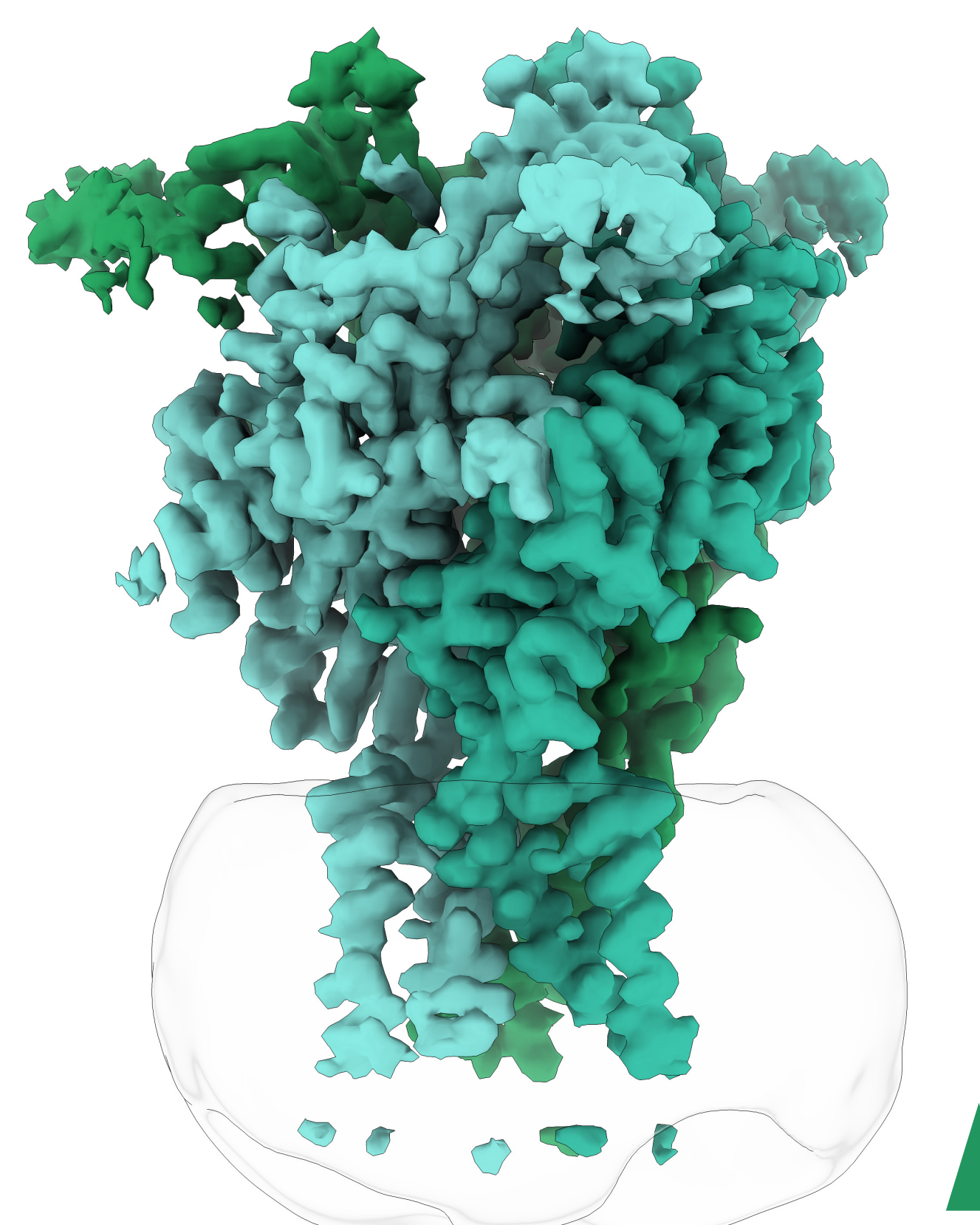
- Significant cost/time reduction for cryo-EM**
- Single symmetrical SEC peak of P2X4 in U-18
- Sample use at higher temperature possible**
- Temperature stability of P2X4 in U-18



2. Screening is Key The NativeMP™ Platform vs Detergent

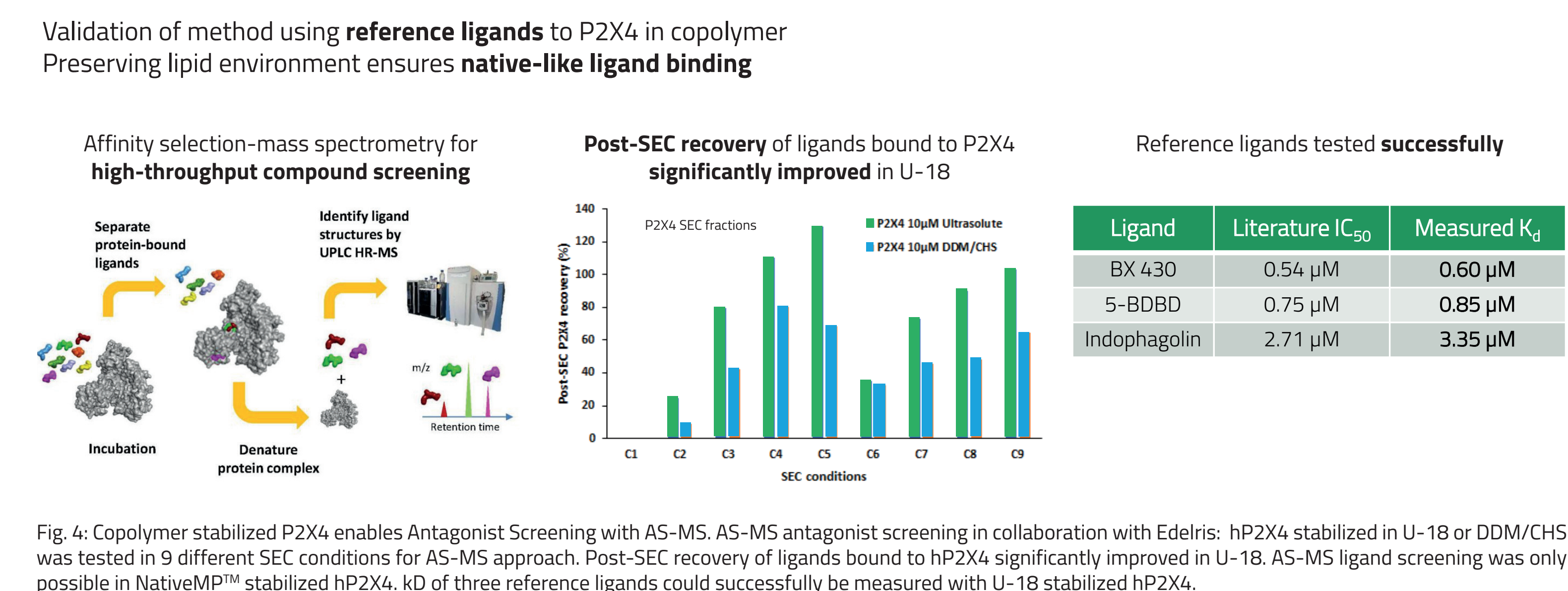


Finding The Perfect Match hP2x4 Full Length Structure



Cube Biotech

4. Yet Again Impossible in Detergent – Possible in NativeMP™ First Ever Ion-Channel Ligand Screened in AS-MS



5. Reproduction in Method Variety Copolymer enables Antagonist Screening with MST/ Spectral Shift

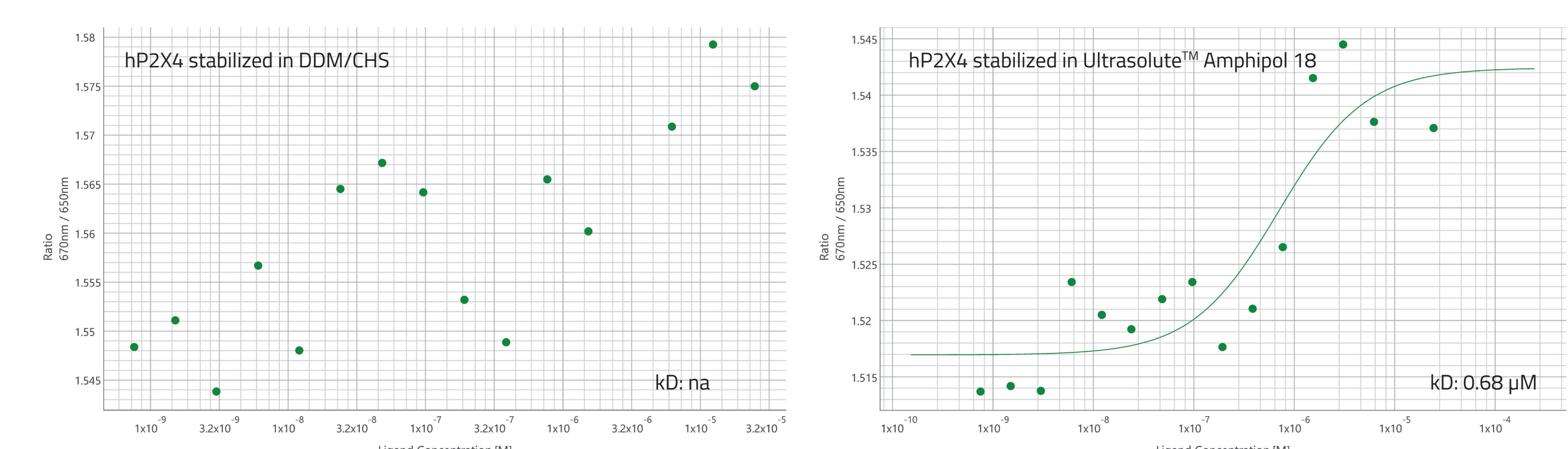
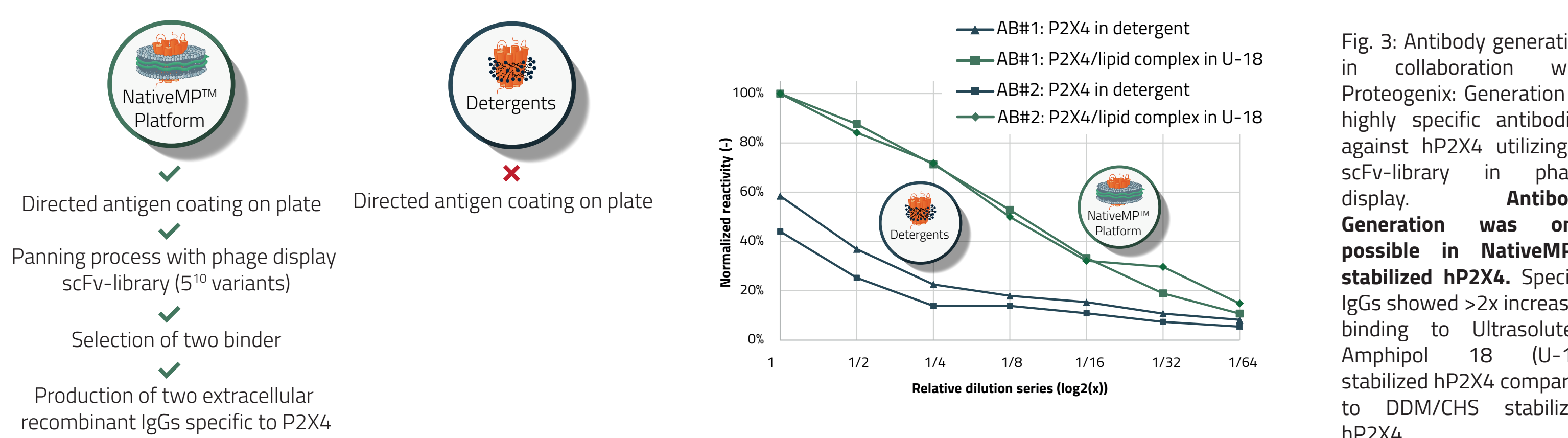


Fig. 5: hP2X4 stabilized in U-18 or DDM/CHS was labeled with a 647-NHS dye and mixed with increasing amounts of 5-BDBD. The binding affinity was measured in solution using a Nanotemper Monolith and standard capillaries after binding check and buffer optimization. A K_d measurement was only possible when hP2X4 was stabilized in synthetic copolymer (U-18, K_d: 0.68 μM). DDM/CHS stabilized hP2X4 showed no measurable K_d when mixed with 5-BDBD.

3. Making the Impossible Possible Copolymer Stabilized P2X4 Enables Antibody Generation



6. Finding The Perfect Match Structure Determination

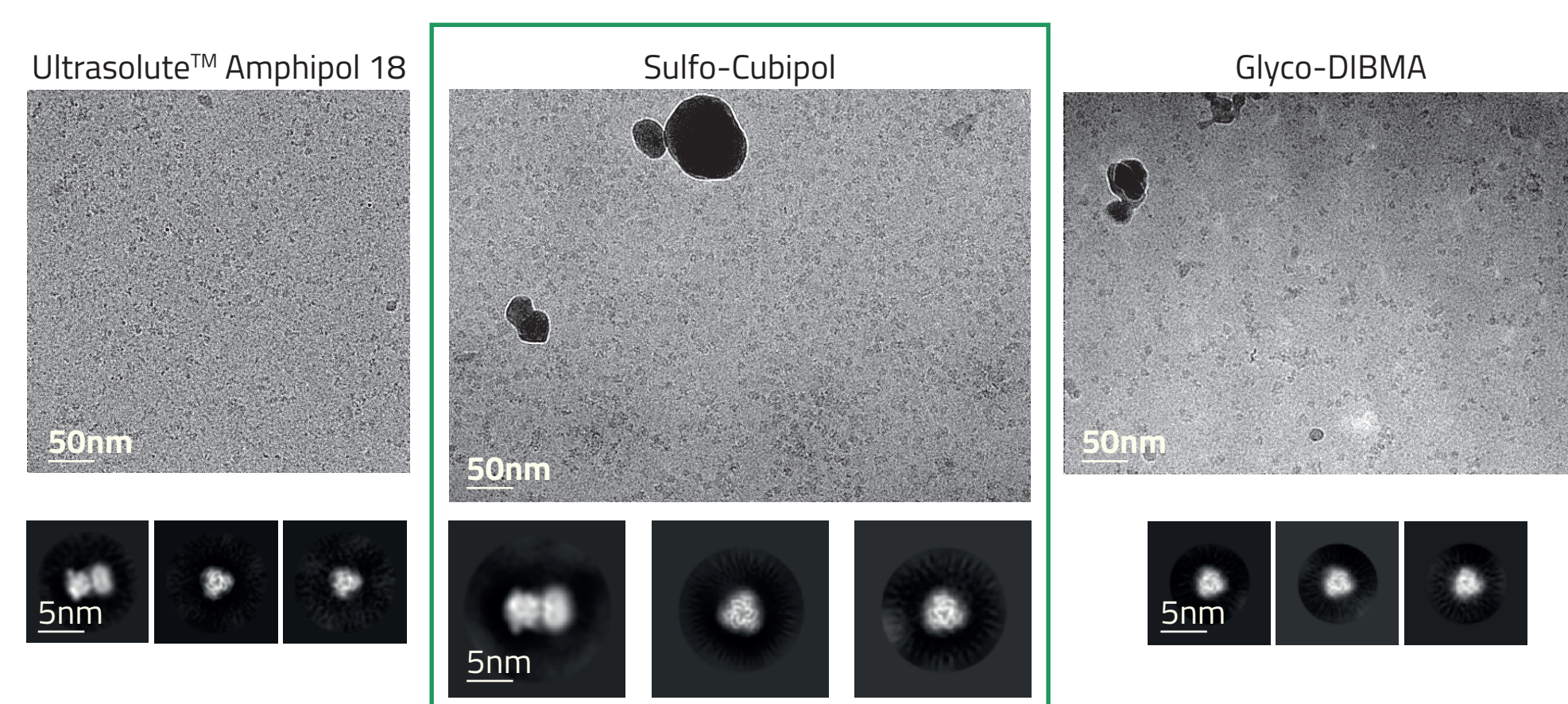


Fig. 6: Careful copolymer selection and an extensive screening process is the key to success when trying to obtain optimal results in cryo-EM. Sulfo-Cubipol produces an optimal particle orientation distribution for hP2X4. **Early refinement cycles already produce a reconstruction of < 3.5 Å global resolution.**

7. Conclusion Universal Applicability Beyond hP2X4

