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**POSTER ABSTRACT PRESENTATIONS**

**SESSION TITLE: STRUCTURE OF MEMBRANE PROTEINS**

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**Abstract P-5: Modern Membrane Mimetic Systems for Cryo-EM Based Structural Analysis**

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**Background:** Single particle approach that is used to image the sample by an electron beam requires receptors to be extracted from the cell membrane and placed into the sample grid in the solubilized form. Conventional use of detergents for GPCR solubilization often leads to poor stability of the receptor removed from its natural lipid environment. Therefore, scientists use membrane mimetic systems (MMS) such as amphipols and various types of nanodiscs, including styrene maleic acid (SMA) derivatives - SMALP and DISMALP. The last one are among the most prospective and interesting MMS and could be used as a tool in Cryo-EM studies. SMALP and DIBMALP have better thermal stability than other types of nanodiscs. SMA could negatively affect the order of phospholipids but DIBMA has a weaker hydrophilic part that is thought to be a reason of weaker impact on phospholipids.

**Methods:** Thermal stability of the nanodiscs was tested using microscale thermophoresis and thermal shift assay.

**Results:** In this work, adenosine A<sub>2a</sub> GPCR was solubilized into SMALP and DIBMALP. Then its stability was tested.

**Conclusion:** SMALP and DIBMALP are useful in Cryo-EM studies. It is better to stabilise membrane proteins in SMALP and DIBMALP for structural analysis because of their high thermal stability.

**Key Words:** nanodiscs • SMALP • DIBMALP • Cryo-EM

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