

12 July 2019

TITLE: Stabilization of human multidrug resistance protein 4 (MRP4/ABCC4) using novel solubilization agents.

AUTHORS:

David HARDY^{1,2}, Roslyn BILL², Alice ROTHNIE^{2¶} & Anass JAWHARI^{1¶}

¹ CALIXAR, 60 avenue Rockefeller 69008, Lyon, France

² Life & Health Sciences, Aston University, Aston Triangle, Birmingham, B4 7ET, UK

¶Corresponding authors:

a.rothnie@aston.ac.uk, ajawhari@calixar.com

12 July 2019

ABSTRACT

Membrane proteins (MP) are important drug discovery targets for a wide range of diseases. However, elucidating the structure and function of native MP is notoriously challenging as their original structure has to be maintained once removed from the lipid bilayer. Conventionally detergents have been used to solubilize MP with varying degrees of success concerning MP stability. To try to address this, new more stabilizing agents have been developed such as calixarene-based detergents and styrene maleic acid co-polymer (SMA). Calixarene based detergents exhibit enhanced solubilizing and stabilizing properties compared to conventional detergents, whereas SMA is able to extract membrane proteins with their surrounding lipids forming a nanodisc structure. Here we report a comparative study using classical detergents, calixarene based detergents and SMA to assess the solubilization and stabilization of the human ABC transporter MRP4 (multidrug resistance protein 4/ABCC4). We show that both SMA and calixarene based detergents have a higher solubility efficiency (at least 80%) than conventional detergents and show striking overstabilization features of MRP4 (up to 70°C) with at least 30°C stability improvement in comparison to the best conventional detergents. These solubilizing agents were successfully used to purify aggregate free homogenous and stable MRP4, with 7-fold higher yield for C4C7 Calixarene detergent in comparison to SMA. This work paves the way to MRP4 structural and functional investigations and illustrates once more the high value of using Calixarene based detergent or SMA as versatile and efficient tools to study MP and eventually enable drug discovery of challenging and highly druggable targets.

INTRODUCTION

Membrane proteins (MP) are capable of controlling what enters and exits cells through transporters and channels, modulating cell signalling with receptors, catalysing reactions by utilizing enzymes and maintaining structure with anchoring proteins. Membrane proteins are highly valuable pharmaceutical targets and understanding the structure and function of these important proteins is vital in the production of beneficial pharmaceutical drugs. Observing membrane proteins in their native state is the most useful for producing effective pharmaceutical drugs, as there have been no modifications to the native structure. Unfortunately, membrane proteins are not stable outside of the native lipid environment. In order to gain a true understanding of how membrane proteins are structured and function they must be viewed in isolation (purified). This means producing a highly stable membrane protein that retains its native structure through purification processes and is able to be used in functional and structural studies. To overcome this stability problem membrane proteins have often been altered either by changing the native amino acid sequence through protein mutagenesis or engineering¹. Antibodies have also been used in efforts to stabilize membrane protein² but all these methods come with a price, they might alter the native conformation of the membrane protein. Investigating membrane proteins in their native state is challenging as they often become highly unstable when solubilized with. Solubilization reagents are available in a range of strengths from harsh anionic detergents such as SDS and Sarkosyl, mild zwitterionic detergents, Fos-Cholines (FC) and CHAPS, to weaker non-ionic detergent such as dodecylmaltoside (DDM) and octyl glucoside (OG). All these contain a similar structure with a hydrophobic acyl tail and a hydrophilic head group allowing them to act as amphiphiles removing the lipids surrounding the membrane protein. In doing so, they are removing the lateral pressure exerted by the lipids keeping the membrane protein in its correct conformation and replacing it with much less stable detergents. Although these conventional detergents have been used for membrane protein studies, success rate is variable and is largely protein dependent. Novel detergents have therefore been produced in an effort to stabilize a much greater variety of membrane proteins.

Many of these novel detergents are built on previous conventional detergents utilizing their solubilization capabilities but also enhancing their stabilizing properties. MNG

12 July 2019

(Maltose-Neopentyl Glycol) and GNG (Glucose-Neopentyl Glycol) are two novel detergents that have very similar structures to DDM and OG respectively, through the addition of a central quaternary carbon atom derived from neopentyl glycol, two hydrophobic and hydrophilic groups can be connected³. Facial amphiphiles represent a slightly different approach to maintaining membrane protein stability. These molecules are comprised of a hydrophobic sterol backbone capable of solubilization attached to different head groups many of which are maltose based³. Facial amphiphiles have been shown to solubilize and aid in the crystallisation membrane proteins⁴ and have further been modified to create tandem facial amphiphiles and are able to span the width of a lipid bilayer³.

Calixarene based detergents have been shown to have a greater ability to stabilize membrane proteins⁵⁻⁶. They contain a calixarene platform comprising of four aromatic rings, three in the para position and the fourth attached to the hydrocarbon chain⁶. By altering the length of the hydrocarbon tail the solubilization properties are affected, increasing their efficiency. Different head groups such as carboxylate or sulfonate groups can be attached to the calixarene platform. These head groups can interact, through the formation of salt bridges, with the aromatic or charged residues at the lipid-protein interface creating a more stable membrane protein-detergent complex⁵. A bacterial ATP Binding Cassette (ABC) transporter, BmrA has previously been extracted with C4C7 (calixarene containing 7 carbon length hydrocarbon chain) and it maintained 90% function whereas solubilization with DDM or FC12 resulted in a 90-99% loss of function. Calixarenes enhance the stability of solubilized membrane proteins⁵⁻⁶.

A different approach to solubilizing and stabilizing membrane proteins is through the use of styrene maleic acid (SMA) co-polymer. This polymer is comprised of alternating units of styrene and maleic acid in varying ratios. They are able to insert into membranes and solubilize bi-layers forming SMA lipid particles (SMALPs)⁷⁻⁸. Membrane proteins can become trapped inside these SMALPs, solubilizing the membrane protein and encapsulating it in a disc of its native lipids⁸⁻⁹ (**Figure 1A**). This is very different from detergent solubilization (**Figure 1B**). Once solubilized in SMA, the membrane protein can be purified without the need for detergents in any of the buffers, making it much more cost effective¹⁰⁻¹¹. These SMA lipid particles (SMALP) can also be used in functional studies as both sides of membrane protein

12 July 2019

are accessible and have been shown to produce highly stable membrane proteins^{10, 12}. SMA has successfully been used to extract a variety of ABC transporters from different expression systems^{10, 13} along with ligand binding studies of ABC transporters and GPCRs¹², functional reconstitution of KscA¹¹ as well as lipid analysis. These examples show the versatility of the SMA polymer in aiding functional understanding of membrane proteins. SMALP purified proteins have been used for structural studies by both crystallography¹⁴ and electron microscopy^{10, 15-17}.

Most reported MP structure and function studies describe the use of one main reagent. Therefore, a comparative study is very often missing. In the current work, we have applied classical detergents, calixarene based detergents and SMA polymers to investigate in parallel the solubilization and stabilization of the same MP-membrane protein target. We have chosen a member of the ABC transporter family (ABCC4/MRP4) as a model membrane protein. ABC transporters are integral membrane proteins that are found in all types of organisms from prokaryotes to humans. They utilize energy from ATP binding and hydrolysis to transport a variety of substrates across the biological lipid bilayer¹⁸. MRP4 is made of four core domains: two transmembrane domains (TMDs) and two nucleotide binding domains (NBD). Each of the two TMDs comprises 6 transmembrane alpha helices and the two NBDs have binding sites for ATP which are homologous throughout the super family unlike the TMDs¹⁹. As its name suggests MRP4 can confer resistance to drugs including cancer chemotherapy, antivirals and antibiotics²⁰. It can also transport signalling molecules such as cyclic nucleotides and eicosanoids making it a drug target for inflammation, pain²¹ and cardiovascular disease²², and it has also been implicated in the development of cancer²³⁻²⁴. To date there is no known structure of human MRP4. To investigate MRP4 structure and function, it is critical to be express it, solubilize it and purify it while maintaining its functionality and structural integrity. Here we show that using calixarene based detergent or SMA, allowed the purification of very stable, homogenous MRP4.

12 July 2019

MATERIAL & METHODS

Expression and Membrane preparation

Expression of the recombinant human MRP4-his₆ within *Sf9* cells was conducted using a baculovirus encoding for recombinant MRP4 generated from a pFastBac-MRP4-his₆ construct with a C-terminal hexa-his tag. Typically, *Sf9* insect cells expression was using the Bac-to-Bac Baculovirus system for 48 hours at MOI of 2. Infected *Sf9* cells were harvested by centrifugation at 6000 g for 10 minutes. The cell pellet was resuspended in homogenisation buffer (50mM Tris-HCl pH 7.4, 250mM Sucrose, 0.25mM CaCl₂ and protease inhibitors (1.3μM benzamidine, 1.8μM leupeptin and 1μM pepstatin). Homogenisation of *Sf9* cells was carried out using nitrogen cavitation at 500 psi for 15 minutes on ice. The cell lysate was centrifuged at 750 g for 10 minutes, the supernatant was then subsequently centrifuged at 100,000 g for 20 minutes. The membrane pellet was resuspended in TSB buffer (50mM Tris-HCl pH 7.4, 250mM Sucrose) and stored at -80°C.

Solubilization

SMA polymers were obtained from Cray Valley (SMA 2000) or Polyscope (SZ25010) as styrene-maleic anhydride polymers and were hydrolysed in 1M NaOH, refluxed and freeze dried to form the styrene-maleic acid form as described in ¹³.

Dot blot analysis was carried out by solubilizing MRP4 *Sf9* membranes at 5 mg/mL total protein using 10x critical micellar concentration (CMC) of each detergent (calixarene based detergents, CALIXAR or conventional detergents, VWR) or 2.5%(w/v) for SMA polymers, for 2 hours at 4°C. 200μL was loaded into the Dot blot apparatus (Biorad), filtered through nitrocellulose and washed with 3 x 200μl PBS to remove insoluble material. An anti-his HRP antibody (3:2000, R&D Systems) was used to detect MRP4 and quantified on a ChemiDoc imaging system. Solubility efficiency was calculated by comparing the density of the dot to an SDS control. Western Blot analysis was performed to assess the solubilization efficiency. *Sf9* MRP4 membranes at 5 mg/mL total protein were solubilized using 10x CMC detergent, or 2.5%(w/v) SMA polymer at 4°C for 2 hours then centrifuged at 100,000 g. The insoluble pellet was resuspended in 1% (w/v) SDS. Samples were run on SDS-

12 July 2019

PAGE and transferred to PVDF membrane and a primary anti-his antibody was used to detect MRP4. Solubility efficiency was calculated via densitometry analysis using ImageJ or Image Studio analysis software by calculating the intensity of the solubilised band as a percentage of the total (soluble + insoluble). Optical density (OD) readings were measured on an Ultrospec 2000 spectrophotometer (Pharmacia Biotech) at 600nm.

Western blot based thermal shift assay

Thermostability of solubilized or purified MRP4 was measured by heating samples for 10 minutes at different temperatures (4 – 90°C) then centrifuged at 14, 000g for 10 minutes. The supernatant was removed and analysed by Western Blot. Data fitting was performed by fitting a dose (temperature) vs normalised response curve using GraphPad Prism. The method was previously described and applied to GPCR solubilization ²⁵.

MRP4 Purification

Sf 9 MRP4 membranes solubilized with 10x CMC Calixarenes were mixed with Ni-NTA affinity resin at a volumetric ratio of 10 : 1 (soluble MRP4 : resin) for 2 hours at 4°C. Resin was washed with 3 x 10 column volumes (CV) 5mM Imidazole and eluted in 5 x 1 CV using 200 mM imidazole. All purification buffers were supplemented with 2 CMC Calixar detergents. MRP4 was concentrated using a 50kDa MWCO spin concentrator.

Affinity purification using SMA polymer was performed using an adapted protocol from ¹³. *Sf* 9 MRP4 membranes solubilized with 2.5% SMA 2000 were mixed with Ni-NTA resin overnight at 4°C a volumetric ratio of 20:1 (soluble MRP4 : resin). Resin was washed with 5 x 10 CV 20mM imidazole and eluted in 5 x 1 CV using 200 mM imidazole. MRP4 was concentrated using a 30 MWCO spin column.

RESULTS

MRP4 solubilization

We wanted to find out which detergents or polymers were best for both solubilizing and stabilizing MRP4. We started by measuring their solubilization properties by screening a large range of detergents and SMA polymers monitored by dot blot (**Figure 2A**). All conventional detergents screened were able to solubilize around 50% of MRP4 from *Sf* 9 cells membranes with the exception of harsh anionic Sarkosyl which was equal to SDS. Calixarene detergents tested here had the same calix[4]arene platform onto which three acidic methylene-carboxylate groups have been grafted at the *para* position while the other face bears a single aliphatic chain of different lengths. Typically, C4C5, C4C7 and C4C8 correspond to detergents with 5, 7 and 8 carbons length, respectively. C4C5, C4C7 and C4C8 were able to solubilize over 90% of the MRP4. Anything below (C4C3) or above (C4C10 C4C11 and C4C12) decreased the solubilization efficiency by around half and were comparable to conventional detergents. The length of the acyl tail in calixarene detergents therefore plays a key role in solubilization efficiency. The SMA polymers also had high solubilization efficiency with SMA 2000 and SZ25010 both at 100%. These results show that novel solubilizing agents are capable of greatly increasing the solubilization efficiency of MRP4 when compared to conventional detergents. The top conditions with the highest solubility efficiency revealed by dot blot were chosen to measure and confirm the solubility efficiency using western blot analysis (**Figure 2B**). C4C5 was shown to have a very high solubility efficiency of 90%; C4C7 was at 76% and C4C8 was at 65% solubility again showing the length of the acyl tail can affect solubility efficiency. SMA 2000 and SZ25010 solubility efficiency was at 83 and 73%, respectively. Only detergent mixtures with harsh conventional detergents such as Sarkosyl and FC12 allowed higher solubilization efficiency than Calixarene detergents alone. Interestingly the kinetics of MRP4 solubilisation by SMA 2000 were much faster than is typically reported for SMA polymers ¹⁰ (Supplementary Figure 1) and it can be seen that the vast majority of solubilisation events occur within the first 15 minutes.

Thermostability of solubilized MRP4

Before moving onto purification, the ability of these solubilizing agents to stabilize MRP4 was assessed using a previously established western blot based thermal shift assay ²⁵. This assay relies on the assumption that unstable heated proteins will aggregate and after ultracentrifugation and western blot the band intensity corresponding to the protein will decay proportionally to its instability. By measuring the amount of MRP4 that remains soluble at increasing temperatures the denaturing point (T_m), 50% soluble, can be estimated. MRP4 solubilized using either C4C5 or C4C7 showed a very high T_m of around 70°C (**Figure 3A**). SMA 2000 solubilised MRP4 also showed a very high T_m of around 75°C (**Figure 3C**). The conventional detergents chosen all have a similar solubilization percentage and had previously been used in studies involving the solubilization of ABC transporters. The T_m for conventional detergents ranged from 28 to 40°C with FC 12 being the highest and C12E8 the lowest (**Figure 3B**). Thus, the T_m for MRP4 in the calixarene detergents (C4C5 and C4C7) or SMALPs was 30°C or 35°C higher than the best conventional detergent for MRP4 stability, respectively.

MRP4 Purification

To evaluate the impact of solubilization reagents on protein purification, his-tag affinity purification was performed for MRP4 solubilized with either C4C7, SMA 2000 or DDM. **Figure 4A** shows that MRP4 was specifically loaded and eluted from the Ni-NTA affinity column. SDS-PAGE gels demonstrate a relatively pure MRP4 after one affinity purification step for both C4C7 and SMA 2000 (**Figure 4B**). However, with DDM, the MRP4 is not at all pure, with multiple contaminating bands which are equally as intense as the MRP4 band (Figure 4B). Notably C4C7 consistently gave higher yields of pure protein (184±8 µg/l cell culture) compared to SMA 2000 (26±7 µg/l cell culture).

Native PAGE Western blot analysis (**Figure 5**) confirmed that one main non-aggregated population of MRP4 was obtained for both C4C7 and SMA 2000, demonstrating the homogeneity and good behavior in solution when SMA 2000 or C4C7 were used for solubilization and purification. MRP4 solubilized and purified in

12 July 2019

C4C7 could easily be concentrated using centrprep centrifugal filter unit without generating any aggregates since the MRP4 band became more intense upon concentration with no aggregate species observed on the well of the gel (**Figure 5A**, compare lane 2 to 1). For both C4C7 and SMA 2000 storage for 7 days at 4°C had no effect on MRP4 aggregation. Similarly, MRP4 showed no changes after freezing and thawing steps in C4C7 or SMA 2000. The same aggregate free behavior in solution was observed even in absence of cryoprotectant (10% glycerol) (**Figure 5**).

The thermostability of purified MRP4 was also examined using the same western blot thermal shift assay as previously described. Interestingly, the same tendencies were noticed after solubilization and purification of MRP4 (**Figure 6**). In fact, SMA 2000 and C4C7 maintained high stability of MRP4 during purification and the T_m remained high, 71°C and 65°C respectively (**Figure 6A** and **6B**), whereas MRP4 solubilized and purified in DDM has a T_m of 49°C (**Figure 6C**).

DISCUSSION

Here we show that MRP4 was successfully solubilized and purified using both C4C7 and SMA 2000. Concerning calixarene based detergent, the length of the hydrophobic chain controls solubilization efficiency. MRP4 was kept in a very stable state compared to the best of the conventional detergents DDM or FC12. Indeed, the stabilizing properties of C4C7 and SMA 2000 were demonstrated by a dramatic thermostability improvement of 30 and 35°C, respectively at the solubilization step in comparison to the best conventional detergents. This stabilization shift is very high considering that the MRP4 investigated was full length, wild type without any single point mutation. It is very common to heavily mutate membrane proteins to improve their thermostability²⁶. This was the case for the Adenosine receptor which was mutated at 8 residues and had 96 amino-acids at the carboxy-terminal deleted leading to ~15°C stability improvement²⁷. Our results illustrate the fact that there is no need to systematically mutate or truncate MPs in order to stabilize them. Using favorable chemical environments can impact positively the stability and functionality of MPs. The fact that C4C7 detergent exhibits a comparable over-stabilizing features in comparison to SMA is outstanding considering that in contrast to detergents, SMALP particles contain lipids and it is well accepted that lipids exert stabilizing effect on MP²⁸. The fact that stability improvement was reduced (to +16 and +22°C for C4C7 and SMA, respectively in comparison to DDM) when MRP4 was solubilized and purified in comparison to solubilized only, is maybe due to the loss of some key stabilizing lipids during the purification process. Further studies including mass spectrometry analysis are required to confirm that. Differences in the degree of cooperativity of the thermal shift curves were noticed. This was also observed for other MPs using other detergents²⁹. Further studies using other thermostability assays are required to explain the contribution of the chemical environment (lipid/ detergent/ polymer) and the MP dynamics (in detergent or in SMA) on the shape of thermal shift curves.

Now that good solubilization and stabilization conditions have been found, structural investigation of MRP4 can begin. The next steps would be to use cryo-EM to investigate MRP4 structure in solution. Preliminary negative stain electron microscopy images showed isolated particles of MRP4 (data not shown). SMA and Calixarene based detergent have both previously been shown to be compatible with electron microscopy^{15-17, 30}. It has previously been reported that SMA somehow

12 July 2019

interferes with binding to Ni²⁺ resin^{13, 31-32}, and this might explain the difference in protein yield. It has also been previously reported that SMA 2000 results in much more pure protein samples than conventional detergents^{10, 33}. Here it is shown that SMA 2000 gave a good degree of purity following a single step affinity purification, but, in contrast to more conventional detergents, C4C7 also gave a comparable degree of purity. There were however some limitations found for each approach. Due to its calixarene platform, Calixarene detergent absorbs at 280 nm which makes protein quantification difficult and limits the use of some biophysical characterization such as Circular Dichroism or tryptophan fluorescence. To address this limitation, new classes of compounds with similar architecture but without the calixarene platform have been designed and applied successfully to MP stabilization³⁴, and this could be a promising future direction for MRP4 studies. The addition of mild groups such as saccharide heads or cholesterol like groups provide more diversity for the such class of stabilizing detergent^{6, 35}. Current limitations to the SMA approach include the disc size. The typical size is 10-12 nm diameter, which may mean some large proteins or protein complexes will not fit. However, recent reports have suggested there is some flexibility with this¹⁶. SMA is also sensitive to divalent cations such as magnesium and calcium³³. This is particularly problematic for proteins like the ABC transporters which require Mg²⁺ for ATP hydrolysis. Alternative polymers such as styrene maleimide SMI and diisobutylene -maleic acid (DIBMA) have also been developed which are reported to overcome the divalent cation sensitivity by replacing the maleic acid with either maleimide (SMI) or replacing the styrene with diisobutylene (DIBMA)³⁶⁻³⁷.

Taken together, we report here that if used for solubilization and purification, C4C7 and SMA maintain the structural integrity of MRP4. In addition to structural studies, these findings open up the possibility of functional and drug discovery approaches with MRP4. Calixarenes based detergents and SMA represent important versatile tools as part of the fast-growing toolbox to help unlock the drug discovery potential of challenging membrane protein targets. This is and will undoubtedly be the case for antibody discovery, structure-based drug design and small molecule screening of highly druggable membrane proteins

FIGURES LEGEND

Figure 1- Schematic representation of membrane proteins in detergent (A) or in Styrene maleic acid (B). Membrane protein helices are represented in blue. SMA, phospholipids and detergent are also indicated.

Figure 2- Solubilization of MRP4.

A- Screening of conventional detergents, Calixarenes and SMA polymers for MRP4 solubilization analysed by Dot blot using a His-tag antibody. All dots intensities were compared to an SDS control. Data are the mean and variation between duplicates. B- Solubilization efficiency assessed by western blot for the selected conditions from dot blot analysis.

Figure 3- Thermostability of MRP4.

Thermostability was measured by the percent of soluble MRP4 present after heating. A- thermostability of MRP4 solubilized with calixarenes C4C5 and C4C7 from 45 to 95°C (n=2). B- Conventional detergent thermostability from 4 to 90°C (n=2). C- thermostability results for MRP4 solubilized with SMA2000 from 40 to 89°C (n=2). The T_m was calculated as the temperature at which 50% remained soluble. Dose (temperature) vs normalised response curve fitted for all graphs. Data represent the mean and the variation between duplicates.

Figure 4- MRP4 purification.

A- Western blot showing His-tag affinity purification of MRP4 after solubilization using C4C7, SMA 2000 or DDM. B- Instant Blue stained SDS-PAGE of purified MRP4 using C4C7, SMA 2000 or DDM.

Figure 5- Native gel analysis.

Native PAGE and western blot analysis to assess the behavior in solution of MRP4 purified with C4C7 (A) or SMA 2000 (B). Examined after protein concentration using centrifugal filter concentrators, storage at 4°C for 7 days or after freezing/ thawing steps. +Gly corresponds to the addition to 10% Glycerol before freezing and thawing steps.

Figure 6- Thermostability of purified MRP4.

Thermostability curves of MRP4 purified in SMA (A), C4C7 (B) and DDM (C) respectively, based on percent soluble after heating and centrifugation (n=3 ±SEM). The blue line represents the transition/melting point (T_m), the temperature at which 50% of MRP4 is still soluble as previously described²⁵. Western blots below graph A, B and C show examples of amounts of soluble MRP4 at each temperature in SMA, C4C7 and DDM respectively. The density of each of the bands was taken and compared to the 4°C control to calculate the percent soluble.

12 July 2019

ACKNOWLEDGMENTS

This work was funded by a Biotechnology and Biological Sciences Research Council Industrial Case Studentship [BB/L015846/1]. A.J.R. was also the recipient of a Royal Society Research Grant [RG110156]. We thank the CALIXAR team for helpful discussion and Emmanuel DEJEAN for continuous support.

DISCLOSURES

AJ is an employee of CALIXAR that have patents applications that covers some of CALX detergent described in this manuscript.

REFERENCES

1. Scott, D. J.; Kummer, L.; Tremmel, D.; et al. Stabilizing membrane proteins through protein engineering. *Current Opinion in Chemical Biology* **2013**, *17*, 427-435.
2. Kobilka, B. K., G protein coupled receptor structure and activation. *Biochimica Et Biophysica Acta-Biomembranes* **2007**, *1768*, 794-807.
3. Chae, P. S.; Rasmussen, S. G. F.; Rana, R. R.; et al. Maltose-neopentyl glycol (MNG) amphiphiles for solubilization, stabilization and crystallization of membrane proteins. *Nature Methods* **2010**, *7*, 1003-1008.
4. Lee, S. C.; Bennett, B. C.; Hong, W. X.; et al. Steroid-based facial amphiphiles for stabilization and crystallization of membrane proteins. *Proceedings of the National Academy of Sciences of the United States of America* **2013**, *110*, 1203-1211.
5. Matar-Merheb, R.; Rhimi, M.; Leydier, A.; et al. Structuring Detergents for Extracting and Stabilizing Functional Membrane Proteins. *Plos One* **2011**, *6*, 1-10.
6. Hardy, D.; Desuzinges Mandon, E.; Rothnie, A. J.; et al. The yin and yang of solubilization and stabilization for wild-type and full-length membrane protein. *Methods* **2018**, *147*, 118-125.
7. Scheidelaar, S.; Koorengel, M. C.; Pardo, J. D.; et al. Molecular Model for the Solubilization of Membranes into Nanodisks by Styrene Maleic Acid Copolymers. *Biophysical Journal* **2015**, *108*, 279-290.
8. Dorr, J. M.; Scheidelaar, S.; Koorengel, M. C.; et al. The styrene-maleic acid copolymer: a versatile tool in membrane research. *Eur Biophys J* **2016**, *45*, 3-21.
9. Jamshad, M.; Lin, Y.-P.; Knowles, T. J.; et al. Surfactant-free purification of membrane proteins with intact native membrane environment. *Biochemical Society Transactions* **2011**, *39*, 813-818.
10. Gulati, S.; Jamshad, M.; Knowles, T. J.; et al. Detergent-free purification of ABC (ATP-binding-cassette) transporters. *Biochemical Journal* **2014**, *461*, 269-278.
11. Dorr, J. M.; Koorengel, M. C.; Schafer, M.; et al. Detergent-free isolation, characterization, and functional reconstitution of a tetrameric K⁺ channel: the power of native nanodisks. *Proc Natl Acad Sci U S A* **2014**, *111*, 18607-18612.
12. Jamshad, M.; Charlton, J.; Lin, Y. P.; et al. G-protein coupled receptor solubilization and purification for biophysical analysis and functional studies, in the total absence of detergent. *Biosci Rep* **2015**, *35*, 1-10.
13. Rothnie, A. J., Detergent-Free Membrane Protein Purification. *Methods Mol Biol* **2016**, *1432*, 261-267.
14. Broecker, J.; Eger, B. T.; Ernst, O. P., Crystallogenesis of Membrane Proteins Mediated by Polymer-Bounded Lipid Nanodisks. *Structure* **2017**, *25*, 384-392.
15. Postis, V.; Rawson, S.; Mitchell, J. K.; et al. The use of SMALPs as a novel membrane protein scaffold for structure study by negative stain electron microscopy. *Biochimica Et Biophysica Acta-Biomembranes* **2015**, *1848*, 496-501.
16. Sun, C.; Benlekbir, S.; Venkatakrishnan, P.; et al. Structure of the alternative complex III in a supercomplex with cytochrome oxidase. *Nature* **2018**, *557*, 123-126.

17. Qiu, W.; Fu, Z.; Xu, G. G.; et al. Structure and activity of lipid bilayer within a membrane-protein transporter. *Proc Natl Acad Sci U S A* **2018**, *115*, 12985-12990.
18. Wilkens, S., Structure and mechanism of ABC transporters. *F1000prime reports* **2015**, *7*, 14-14.
19. Linton, K. J., Structure and Function of ABC Transporters. *Physiology*. **2007**; *22*, 122-130.
20. Russel, F. G.; Koenderink, J. B.; Masereeuw, R., Multidrug resistance protein 4 (MRP4/ABCC4): a versatile efflux transporter for drugs and signalling molecules. *Trends in pharmacological sciences* **2008**, *29*, 200-207.
21. Lin, Z. P.; Zhu, Y. L.; Johnson, D. R.; et al. Disruption of cAMP and prostaglandin E2 transport by multidrug resistance protein 4 deficiency alters cAMP-mediated signaling and nociceptive response. *Molecular pharmacology* **2008**, *73*, 243-251.
22. Belleville-Rolland, T.; Sassi, Y.; Decouture, B.; et al. MRP4 (ABCC4) as a potential pharmacologic target for cardiovascular disease. *Pharmacological research* **2016**, *107*, 381-389.
23. Henderson, M. J.; Haber, M.; Porro, A.; et al. ABCC multidrug transporters in childhood neuroblastoma: clinical and biological effects independent of cytotoxic drug efflux. *Journal of the National Cancer Institute* **2011**, *103*, 1236-1251.
24. Copsel, S.; Garcia, C.; Diez, F.; et al. Multidrug resistance protein 4 (MRP4/ABCC4) regulates cAMP cellular levels and controls human leukemia cell proliferation and differentiation. *The Journal of biological chemistry* **2011**, *286*, 6979-6988.
25. Igonet, S.; Raingeval, C.; Cecon, E.; et al. Enabling STD-NMR fragment screening using stabilized native GPCR: A case study of adenosine receptor. *Sci Rep* **2018**, *8*, 8142.
26. Magnani, F.; Serrano-Vega, M. J.; Shibata, Y.; et al. A mutagenesis and screening strategy to generate optimally thermostabilized membrane proteins for structural studies. *Nat Protoc* **2016**, *11*, 1554-1571.
27. Magnani, F.; Shibata, Y.; Serrano-Vega, M. J.; et al. Co-evolving stability and conformational homogeneity of the human adenosine A2a receptor. *Proc Natl Acad Sci U S A* **2008**, *105*, 10744-10749.
28. Gupta, K.; Donlan, J. A. C.; Hopper, J. T. S.; et al. The role of interfacial lipids in stabilizing membrane protein oligomers. *Nature* **2017**, *541*, 421-424.
29. Guillet, P.; Mahler, F.; Garnier, K.; et al. Hydrogenated Diglucose Detergents for Membrane-Protein Extraction and Stabilization. *Langmuir* **2019**, *35*, 4287-4295.
30. Agez, M.; Schultz, P.; Medina, I.; et al. Molecular architecture of potassium chloride co-transporter KCC2. *Scientific Reports* **2017**, *7*, 1-14.
31. Smirnova, I. A.; Sjostrand, D.; Li, F.; et al. Isolation of yeast complex IV in native lipid nanodiscs. *Biochimica et biophysica acta* **2016**, *1858*, 2984-2992.
32. Pollock, N. L.; Lee, S. C.; Patel, J. H.; et al. Structure and function of membrane proteins encapsulated in a polymer-bound lipid bilayer. *Biochimica et biophysica acta* **2018**, *1860*, 809-817.
33. Morrison, K. A.; Akram, A.; Mathews, A.; et al. Membrane protein extraction and purification using styrene-maleic acid (SMA) copolymer: effect of variations in polymer structure. *Biochemical Journal* **2016**, *473*, 4349-4360.

12 July 2019

34. Nguyen, K. A.; Peuchmaur, M.; Magnard, S.; et al., Glycosyl-Substituted Dicarboxylates as Detergents for the Extraction, Overstabilization, and Crystallization of Membrane Proteins. *Angew Chem Int Ed Engl* **2018**, *57*, 2948-2952.
35. Dauvergne, J., Mandon Desuzinges, E., Faugier, C., et al. Glycosylated Amphiphilic Calixarene-Based Detergent for Functional Stabilization of Native Membrane Proteins. *ChemistrySelect* **2019**, *4*, 5535-5539.
36. Hall, S. C. L.; Tognoloni, C.; Charlton, J.; et al. An acid-compatible copolymer for the solubilization of membranes and proteins into lipid bilayer-containing nanoparticles. *Nanoscale* **2018**, *10*, 10609-10619.
37. Oluwole, A. O.; Klingler, J.; Danielczak, B.; et al. Formation of Lipid-Bilayer Nanodiscs by Diisobutylene/Maleic Acid (DIBMA) Copolymer. *Langmuir* **2017**, *33*, 14378-14388.

12 July 2019

Figure 1

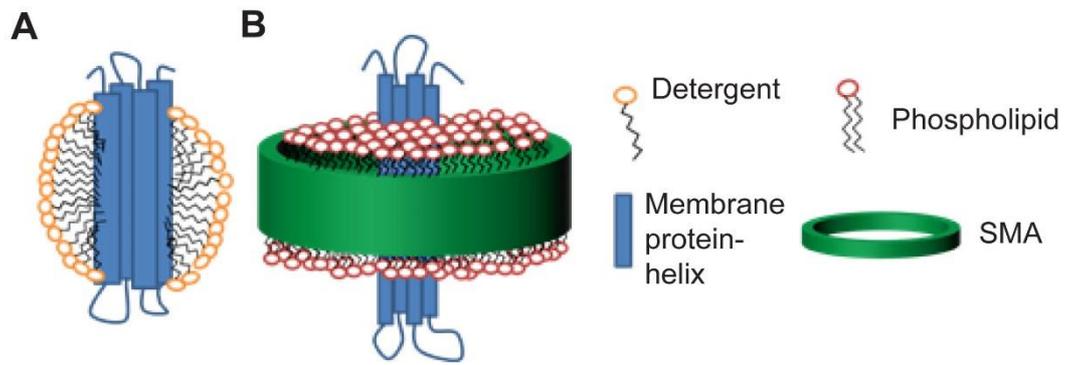


Figure 2

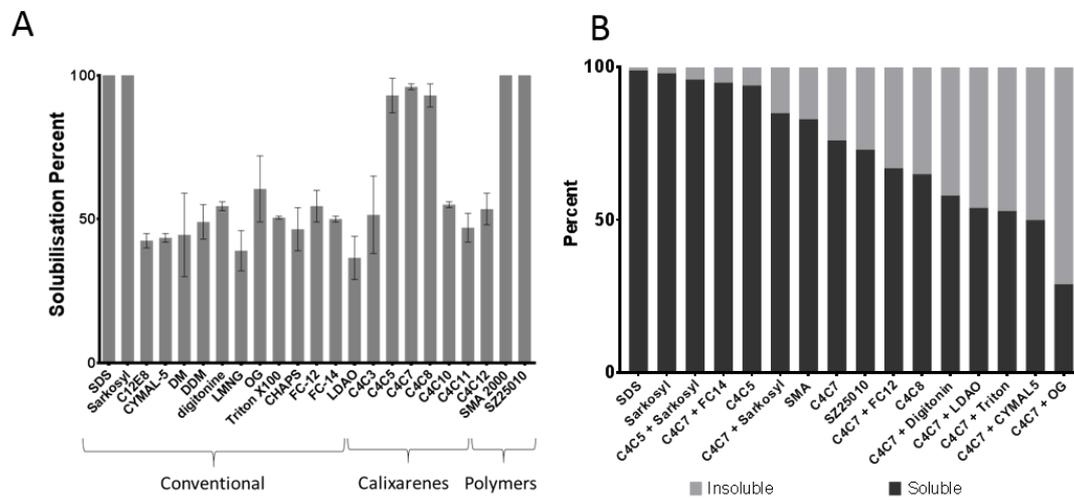


Figure 3

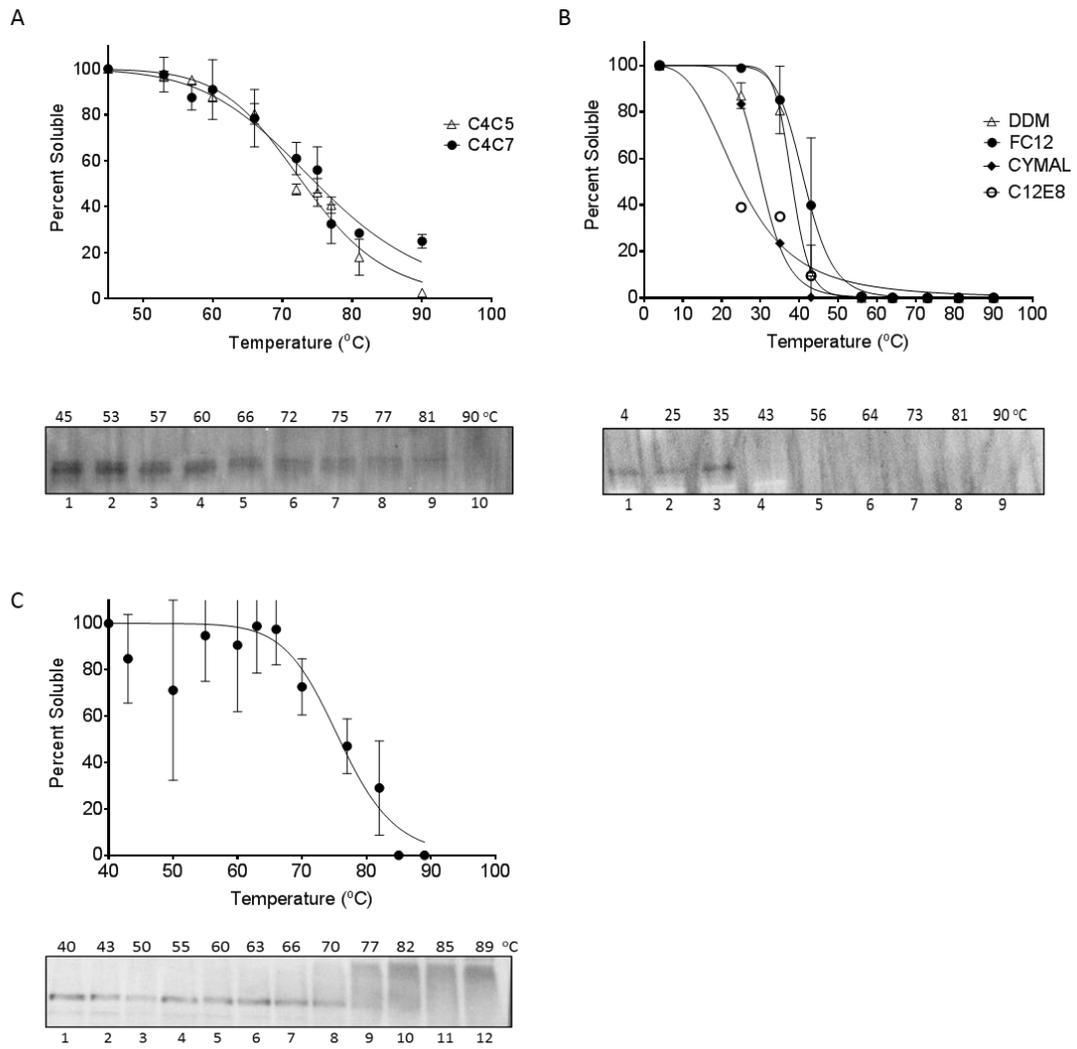


Figure 4

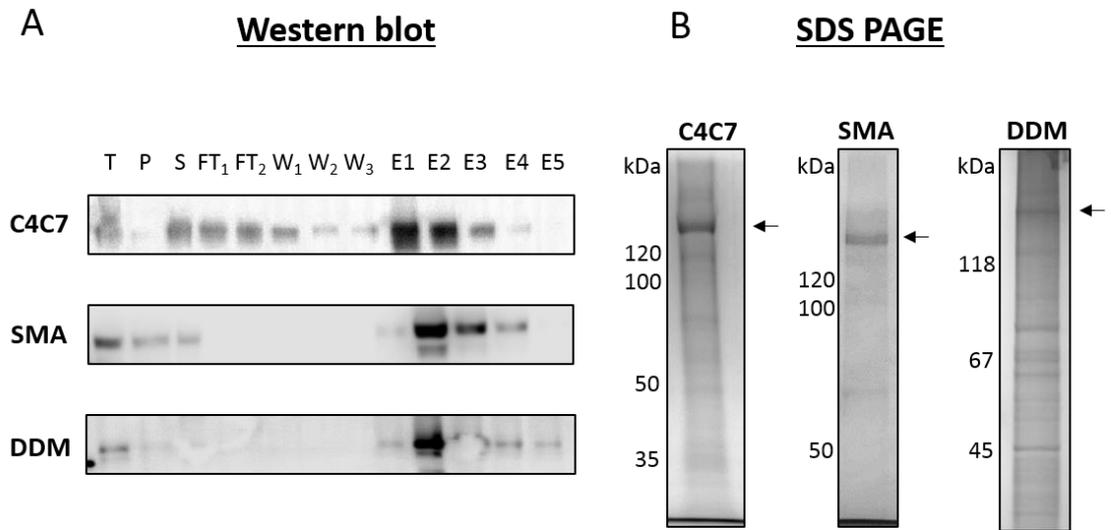
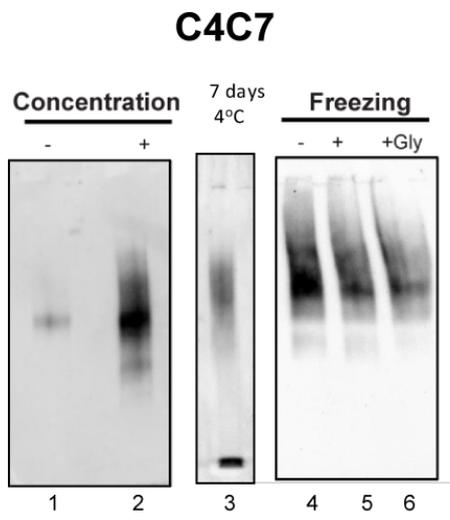


Figure 5

A



B

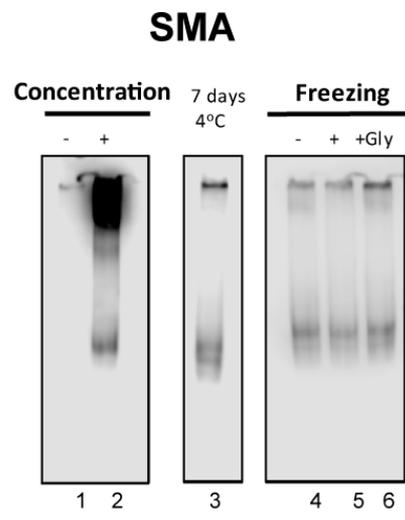
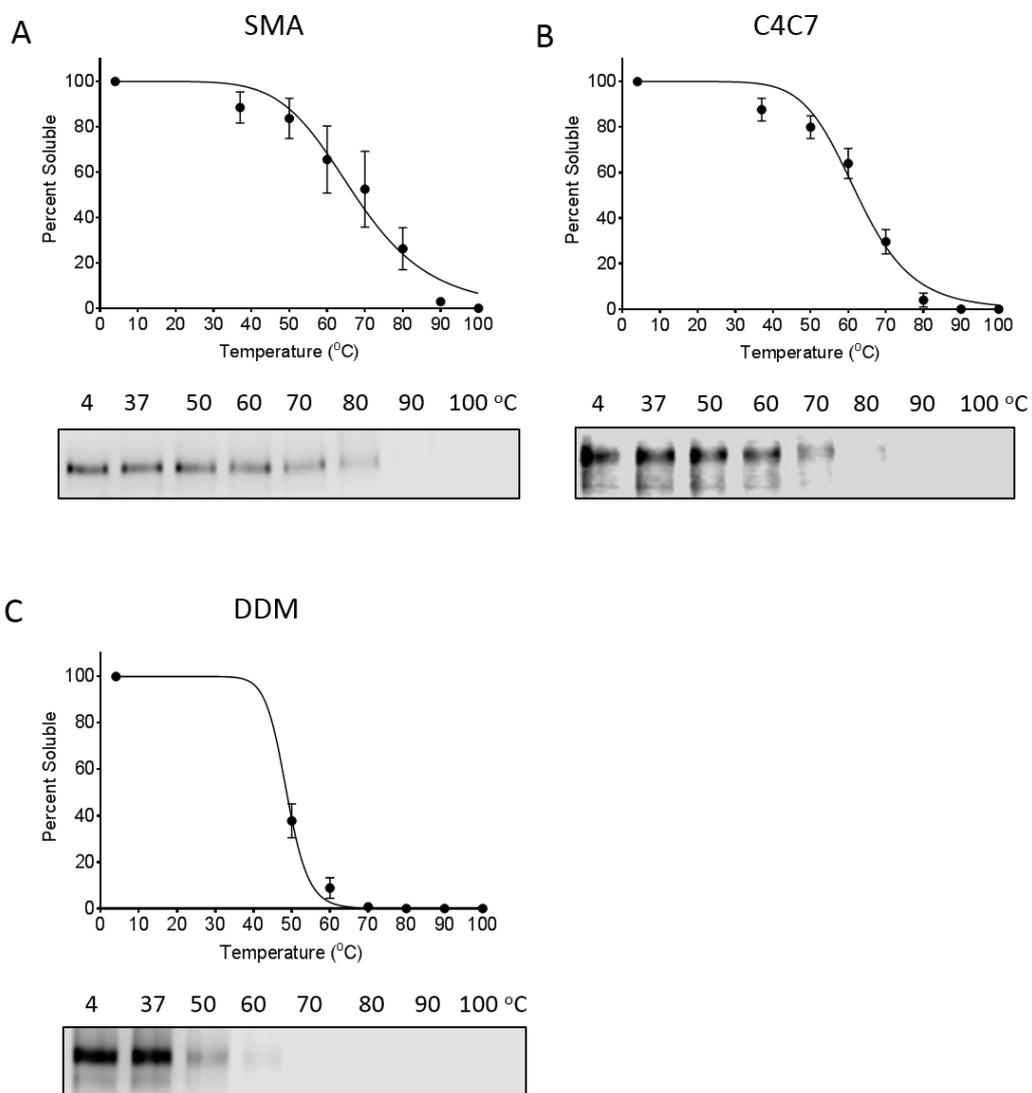


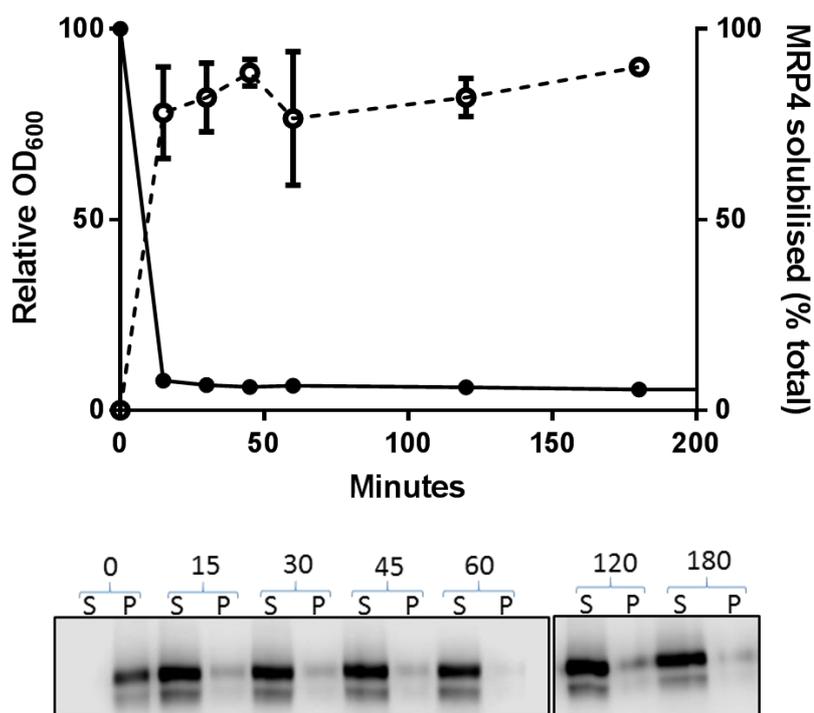
Figure 6



Stabilization of human multidrug resistance protein 4 (MRP4/ABCC4) using novel solubilization agents.

David Hardy, Roslyn Bill, Alice Rothnie & Anass Jawhari

Supplementary Information



Supplementary Figure 1: Kinetics of MRP4 solubilisation with SMA 2000.

The solubilisation of MRP4 expressing Sf9 cell membranes at room temperature was monitored over time by measuring both the turbidity/OD₆₀₀ (closed circles) to monitor general membrane disruption, and Western blotting to measure MRP4-specific solubilisation (open circles). It can be seen that the solubilisation was quick, and essentially complete after 15 minutes.