FISHVIER

Contents lists available at ScienceDirect

# Biochimica et Biophysica Acta

journal homepage: www.elsevier.com/locate/bbamem



# Characterizing the structure of lipodisq nanoparticles for membrane protein spectroscopic studies



Rongfu Zhang, Indra D. Sahu, Lishan Liu, Anna Osatuke, Raven G. Comer, Carole Dabney-Smith, Gary A. Lorigan \*

Department of Chemistry and Biochemistry, Miami University, Oxford, OH 45056, USA

#### ARTICLE INFO

Article history: Received 7 February 2014 Received in revised form 28 April 2014 Accepted 4 May 2014 Available online 20 May 2014

Keywords: lipodisq nanoparticle POPC/POPG vesicle 3:1 SMA polymer dynamic light scattering SSNMR spectroscopy

#### ABSTRACT

Membrane protein spectroscopic studies are challenging due to the difficulty introduced in preparing homogenous and functional hydrophobic proteins incorporated into a lipid bilayer system. Traditional membrane mimics such as micelles or liposomes have proved to be powerful in solubilizing membrane proteins for biophysical studies, however, several drawbacks have limited their applications. Recently, a nanosized complex termed lipodisq nanoparticles was utilized as an alternative membrane mimic to overcome these caveats by providing a homogeneous lipid bilayer environment. Despite all the benefits that lipodisq nanoparticles could provide to enhance the biophysical studies of membrane proteins, structural characterization in different lipid compositions that closely mimic the native membrane environment is still lacking. In this study, the formation of lipodisq nanoparticles using different weight ratios of POPC/POPG lipids to SMA polymers was characterized via solidstate nuclear magnetic resonance (SSNMR) spectroscopy and dynamic light scattering (DLS). A critical weight ratio of (1/1.25) for the complete solubilization of POPC/POPG vesicles has been observed and POPC/POPG vesicles turned clear instantaneously upon the addition of the SMA polymer. The size of lipodisq nanoparticles formed from POPC/POPG lipids at this weight ratio of (1/1.25) was found to be about 30 nm in radius. We also showed that upon the complete solubilization of POPC/POPG vesicles by SMA polymers, the average size of the lipodisq nanoparticles is weight ratio dependent, when more SMA polymers were introduced, smaller lipodisq nanoparticles were obtained. The results of this study will be helpful for a variety of biophysical experiments when specific size of lipid disc is required. Further, this study will provide a proper path for researchers working on membrane proteins to obtain pertinent structure and dynamic information in a physiologically relevant membrane mimetic environment. This article is part of a Special Issue entitled: NMR Spectroscopy for Atomistic Views of Biomembranes and Cell Surfaces. Guest Editors: Lynette Cegelski and David P. Weliky.

© 2014 Elsevier B.V. All rights reserved.

# 1. Introduction

Magnetic resonance spectroscopic studies of membrane proteins remain highly challenging due to the requirement of a membrane-mimicking environment that maintains the integrity and stability of membrane proteins outside their native cellular environment [1]. The most commonly applied method to solubilize membrane proteins in aqueous solution is detergent-formed micelles [2], which are widely used to solve high resolution three-dimensional structures of membrane proteins [2,3]. However, the lack of a lipid bilayer and the limitation of the size of micelles may not preserve the membrane proteins' structural and dynamic integrities under physiological conditions [1, 3–5].

To better maintain the structural integrity of membrane proteins, several classes of membrane mimics (e.g. liposomes, bicelles, or

nanodiscs) comprising lipid bilayer have been developed previously [6–9]. Despite the advantages, several drawbacks have limited their applications for spectroscopic studies. In the case of liposomes, homogenous liposome samples are not readily obtainable, especially when membrane proteins are to be incorporated. Also, the inaccessibility of liposomal interior raises challenges for the cytoplasmic domain studies of membrane proteins [10]. Furthermore, it is not easy to incorporate large amounts of proteins into liposomes which is problematic given that sometimes a higher protein to lipid molar ratio is required for biophysical studies [11,12]. A second alternative bicelles, are artificial lipid bilayer discs formed by the mixture of long-chain phospholipids (e.g. DMPC) and short-chain phospholipids (e.g. DHPC). Bicelles are favorable for the study of interactions within membrane proteins that are not retained in micelles [13,14]. Bicelles are able to provide accessibility for the interaction study of both extracellular and cytoplasmic domains of membrane protein [10]. However, the specific types of lipids amenable to bicelle formation limit its applications as the lipid compositions in the membrane have been shown to influence the function of antimicrobial peptides and amyloid peptides [10,15,16]. The introduction of nanodiscs has provided a great tool to study membrane proteins in a native-like

<sup>↑</sup> This article is part of a Special Issue entitled: NMR Spectroscopy for Atomistic Views of Biomembranes and Cell Surfaces. Guest Editors: Lynette Cegelski and David P. Weliky.

<sup>\*</sup> Corresponding author. Tel.: +1 513 529 3338. E-mail address: gary.lorigan@miamioh.edu (G.A. Lorigan).

membrane environment [17,18]. Nanodiscs consist of lipids surrounded by a membrane scaffold protein such as apolipoprotein to form a discoidal bilayer. Nanodiscs can be formed with different types of lipids, which gives nanodiscs great advantages over bicelles since many membrane proteins require specific types of lipids for structural folding and functional reconstitution [19–21]. Also, it has been reported that membrane scaffold proteins helped to improve the stability of nanodiscs compared to other membrane-mimicking systems [22]. The drawbacks of using nanodiscs are that it requires detergent for protein incorporation and the absorbance properties of the membrane scaffold protein may interfere with the membrane protein of interest [9,17].

An alternative membrane mimic is highly desirable for the proper functional and structural characterization of membrane proteins. In this study, we characterize the recently developed lipodisq nanoparticle system (Fig. S1 A) as a potential membrane mimic system [23-25]. Unlike nanodiscs, lipodisq nanoparticles are formed from lipids solubilized by polymers instead of membrane scaffold proteins, thus they should not interfere with the absorbance properties of the membrane proteins of interest [23,26,27]. Also, no detergent is needed for protein incorporation into the lipodisq nanoparticles. The polymer for solubilizing lipids consists of styrene and maleic acid (SMA) at a molar ratio of 3:1 (Fig. S1 B). Though lipodisg nanoparticles show a high potential to become a good membrane mimetic to enhance biophysical studies of membrane proteins, its structural characterization in a native-membrane mimetic environment is still lacking. Here, we use POPC/POPG lipids at a molar ratio of 9/1 and SMA polymers to form lipodisg nanoparticles. POPC is the most common phospholipid found in mammalian cell membranes, and POPG was chosen to mimic the 10-20 molar percent anionic phospholipids typically found in mammalian membranes [28,29]. The weight ratio of lipid to polymer was varied and different lipodisq nanoparticle samples were characterized by solid-state NMR (SSNMR) and dynamic light scattering (DLS). Transmission electron microscopy (TEM) was also used to confirm the size and homogeneity of vesicles and lipodisq nanoparticle samples.

# 2. Materials and methods

# 2.1. Materials

1-Palmitoyl-2-oleoyl-sn-glycero-3-phosphocholine (POPC), 1-palmitoyl-2-oleoyl-sn-glycero-3-phospho-(1'-rac-glycerol) (sodium salt) (POPG), 1-myristoyl-2-hydroxy-sn-glycero-3-phospho-(1'-rac-glycerol) (sodium salt) (LMPG), 1,2-dimyristoyl-sn-glycero-3-phosphocholine (DMPC), and 1,2-diheptanoyl-sn-glycero-3-phosphocholine (DHPC) were purchased from Avanti Polar Lipids (Alabaster, Alabama, USA). 3:1 SMA polymer was purchased from Malvern Cosmeceutics (Tewkesbury, Gloucestershire, UK). *N*-[2-Hydroxyethyl]piperazine-*N*'-2-ethanesulfonic acid (HEPES) was purchased from Sigma-Aldrich (St. Louis, MO, USA). Sodium chloride (NaCl) was purchased from Fisher Scientific (Pittsburgh, PA, USA).

# 2.2. Preparation of POPC/POPG vesicles and lipodisq nanoparticles

POPC/POPG vesicles were composed of POPC and POPG with molar ratio of [POPC]/[POPG] = 9/1. POPC and POPG powdered lipids were suspended in the buffer (100 mM NaCl, 20 mM HEPES, pH 7.0) to a final concentration of 25 mM. Lipid slurry was vortexed vigorously to mix completely and vesicles were formed spontaneously after ten freeze/sonication cycles. SMA polymers were dissolved in the buffer (100 mM NaCl, 20 mM HEPES, pH 7.0) to a final concentration of 5% (m/v) followed by 1 min of water-bath sonication to a complete dissolution. Lipodisq nanoparticles with different weight ratio of lipid to polymer (1/0.25, 1/0.5, 1/0.75, 1/1.25, 1/1.75, and 1/2.25) were formed by adding SMA polymers dropwise to the POPC/POPG vesicles. Samples were then mixed well by vortexing and equilibrated through two freeze/sonication cycles. Samples were

allowed to equilibrate at room temperature overnight. For solid state NMR experiment, sample was concentrated to around 100  $\mu$ l at 14,000 g using Amicon concentrator at cutoff of 3 kDa according to manufacturer's instructions.

# 2.3. Preparation of LMPG micelles and DMPC/DHPC bicelles

LMPG micelles were formed by dissolving LMPG in buffer (100 mM NaCl, 5 mM HEPES, pH 7.0) to a final concentration of 42 mM (20 mg/ml). DMPC/DHPC bicelles were composed of DMPC and DHPC with molar ratio of q = [DMPC]/[DHPC] = 3.2. DMPC and DHPC powdered lipids were dissolved in 5 mM HEPES buffer (pH 7.0) to a final concentration of 12% (m/v). Three cycles of the following: heating at 40 °C for 15 min, vortexing for 1 min, cooling on ice for 15 min and again vortexing for 1 min, were performed until the lipid suspension was clear.

## 2.4. Dynamic light scattering (DLS)

DLS measurements were performed on a Zetasizer nano series (Malvern Instruments) at 25  $^{\circ}$ C in disposable 40  $\mu$ l micro cuvettes. Data were collected for 20 s and averaged for 10 scans. The size distribution in radius is shown on a log scale using Igor Pro (WaveMetrics).

### 2.5. Solid-state NMR spectroscopy

A 500 MHz WB Bruker Avance solid-state NMR spectrometer and a Bruker 4 mm triple resonance NMR probe (Billerica, MA, USA) were used to collect the  $^{31}P$  solid-state NMR spectra.  $^{31}P$  NMR spectra were recorded with  $^{1}H$  decoupling using a 7  $\mu s$   $\pi/2$  pulse for  $^{31}P$  and a 3 s recycle delay, 2 K scans were averaged, and the free induction decay was processed using 100 Hz of line broadening. The spectral width was set to 500 ppm. All of the data were collected at 25 °C except for DMPC/DHPC bicelle, which was collected at 42 °C.  $^{31}P$  NMR spectra were plotted using Igor Pro (WaveMetrics).

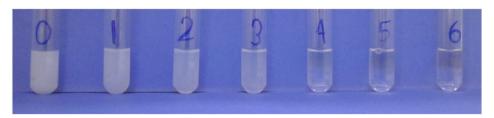
# 2.6. Transmission electron microscopy (TEM)

One drop of control POPC/POPG vesicle and lipodisq nanoparticle (1/1.25) samples were adsorbed to 200 mesh copper carbon-coated grids for 10 s for fully absorbance. Grids were stained with two drops of 1.5% ammonium molybdate. Images were recorded using a Joel-1200 electron microscope operated at 120 kV.

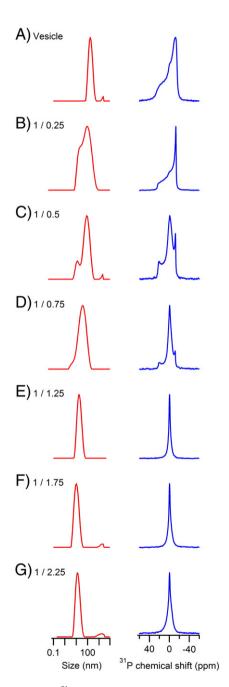
### 3. Results and discussions

Fig. 1 shows the visible observation of the transparency of POPC/POPG vesicles titrated with different amounts of SMA polymers. From left to right, the weight ratio of POPC/POPG lipid to SMA polymer gradually increased from (1/0) (tube 0) to (1/2.25) (tube 6). At a weight ratio of (1/0), the sample was opaque and cloudy. As the weight ratio increased, the sample appeared less cloudy. When the critical weight ratio of (1/1.25) (tube 4) was reached, the sample turned to clear, which indicated the formation of a homogenous sample. The addition of the SMA polymer instantaneously turned the sample clear, which indicates the potential of using lipodisq nanoparticles for efficient sample preparation. These results revealed a qualitative observation of the solubilization of POPC/POPG vesicles and demonstrated the optimal amount of SMA polymers needed to fully dissolve POPC/POPG vesicles at a weight ratio of (1/1.25). After the formation of liposomes, SMA lipodisq nanoparticles are quick and easy to prepare.

<sup>31</sup>P SSNMR and DLS experiments were performed to investigate the formation of lipodisq nanoparticles from POPC/POPG vesicles and SMA polymers. <sup>31</sup>P SSNMR spectra provided unique structural and dynamic information on the phospholipid head groups, and DLS measured the



**Fig. 1.** Visible observation of lipodisq nanoparticles formation through the combination of POPC/POPG vesicles and SMA polymers at different lipid to polymer weight ratios and normal POPC/POPG vesicles as a control: (0) control POPC/POPG vesicles, (1) lipodisq nanoparticles (1/0.25), (2) lipodisq nanoparticles (1/0.5), (3) lipodisq nanoparticles (1/0.75), (4) lipodisq nanoparticles (1/1.25), (5) lipodisq nanoparticles (1/1.75), (6) lipodisq nanoparticles (1/2.25).



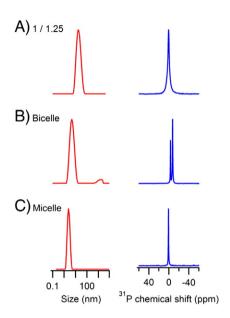
**Fig. 2.** DLS data (left, red) and <sup>31</sup>P SSNMR data (right, blue) for control POPC/POPG vesicles and lipodisq nanoparticles generated from different lipid to polymer weight ratios: (A) control POPC/POPG vesicles, (B) lipodisq nanoparticles (1/0.25), (C) lipodisq nanoparticles (1/0.5), (D) lipodisq nanoparticles (1/0.75), (E) lipodisq nanoparticles (1/1.25), (F) lipodisq nanoparticles (1.75), (G) lipodisq nanoparticles (1/2.25).

size distribution of each sample. Both experiments were performed and the results were compared side by side. Fig. 2 shows DLS data (left, red) and the corresponding <sup>31</sup>P SSNMR data (right, blue) for control POPC/ POPG vesicles and lipodisq nanoparticles formed with different weight ratios of lipid to polymer. Both DLS and <sup>31</sup>P SSNMR results clearly indicate a change in the size of POPC/POPG vesicles from larger to smaller upon adding SMA polymers. As observed in Fig. 2 A through E, there was a gradual shift of average size in radius from about 250 nm to about 30 nm (see Table 1). Furthermore, a broader size distribution was observed when the weight ratio of lipid to polymer was greater than (1/1.25), which indicated an incomplete dissolution of POPC/ POPG vesicles at these weight ratios and the existence of heterogeneity in these samples (Fig. 2 B-D, The right most small peak may be due to small amounts of sample inhomogeneity which is not significant when compared to the main peak). This transition was supported by the corresponding <sup>31</sup>P SSNMR results. An obvious <sup>31</sup>P powder pattern shape was observed in Fig. 2 A, and this powder pattern gradually decreased from Fig. 2 B through 2 D, accompanied by a gradual increasing  $^{31}$ P isotropic peak. When the weight ratio of (1/1.25) was reached, as shown in Fig. 2 E, the <sup>31</sup>P powder pattern disappeared and only the isotropic peak was left, suggesting the formation of small lipodisg nanoparticles corresponding to fast isotropic motion. This formation of an isotropic peak when the weight ratio of lipid to polymer was (1/1.25) corresponded to the complete dissolution of the lipids as observed in Fig. 1. Furthermore, inspection of the DLS and SSNMR results from samples with weight ratios of (1/0.75) to (1/1.25) revealed a change in particle size in radius from about 90 nm to about 30 nm on average and a change in chemical shift width from 45 ppm to 6 ppm (see Table 1), clearly indicating the formation of small and homogenous lipodisg nanoparticles at the specific weight ratio of (1/1.25). These results indicated that the weight ratio was crucial in the formation of the lipodisg nanoparticles, and a threshold of weight ratio has to be reached in order to make a homogenous lipodisg nanoparticle suspension. We observed that the samples with weight ratios smaller than (1/1.25) (Fig. 2 F and G), the uniformity of the lipodisq nanoparticle sample was unchanged, though the size of the particle was smaller (see Table 1). This result suggested that by modifying the weight ratio at the critical dissolution point, the lipodisq nanoparticles with a specific size can be formed. The DLS and <sup>31</sup>P NMR studies are useful to help characterize lipodisq nanoparticles and for studying lipid systems when a specific size is required.

In order to further characterize the lipodisq nanoparticles as a model membrane mimic, <sup>31</sup>P powder spectrum and size distribution of the lipodisq nanoparticles at the weight ratio of (1/1.25) were compared with other model membrane mimics (LMPG micelles and DMPC/DHPC bicelles). Fig. 3 shows DLS data (left, red) and <sup>31</sup>P SSNMR data (right, blue) of these three membrane mimics. DLS analysis of the particles demonstrated that the lipodisq nanoparticles had a narrow size distribution, which is comparable to micelles and bicelles. Under these sample conditions, the lipodisq nanoparticles are larger in size than both the micelle and bicelle samples. This is advantageous for incorporating larger integral membrane proteins. The isotropic <sup>31</sup>P NMR linewidths are broader for the lipodisq nanoparticle sample when compared to the micelle and bicelle samples. This is because of the larger size and the corresponding slower motion of the lipodisq nanoparticles

**Table 1** Particle size in nm and  $^{31}$ P NMR spectrum line shape width in ppm of each sample.

Sample names	Vesicles	Lipodisq nanoparticles						Bicelles	Micelles
		1/0.25	1/0.5	1/0.75	1/1.25	1/1.75	1/2.25		
Particle size (r. nm)	250	190	40-180	90	31	19	15	9	3
<sup>31</sup> P Line shape width (ppm)	42	38	35	34	6	8	10	1	1.5



**Fig. 3.** Comparison of DLS data (left, red) and SSNMR data (right, blue) of lipodisq nanoparticles (1/1.25) at 25  $^{\circ}$ C, DMPC/DHPC bicelles at 42  $^{\circ}$ C, and LMPG micelles at 25  $^{\circ}$ C.

in solution. Also, the <sup>31</sup>P NMR spectrum of the lipodisq nanoparticles is comparable to another commonly utilized powerful membrane mimic system called nanodiscs that have been reported previously in the literature [30]. This result indicated that the homogeneity of the lipodisq nanoparticles has strong potential to serve as an alternative membrane mimic. Most importantly, the larger size and lipid bilayer properties of the lipodisq nanoparticles shows great promise over micelles and bicelles as a membrane mimic.

To confirm the size of the lipodisq nanoparticles obtained from the DLS measurements, transmission electron microscopy (TEM) experiments were conducted. Fig. 4 shows TEM micrographs of uranyl acetate stained POPC/POPG vesicles and lipodisq nanoparticles (1/1.25). The average size of lipodisq nanoparticles (1/1.25) from TEM agreed with the

analysis of the particles by DLS (Figs. 2 and 3). Also, TEM results indicated that the lipodisq nanoparticle (1/1.25) sample was more homogenous in size when compared to POPC/POPG vesicles, which is a significant improvement in membrane mimetic systems, making the lipodisq nanoparticle system very attractive as an alternative method for membrane mimicking systems.

In conclusion, the formation of lipodisq nanoparticles from POPC/POPG lipids and SMA polymers was successfully demonstrated. A critical weight ratio (lipid to polymer) of (1/1.25) or smaller was required for the complete dissolution of POPC/POPG vesicles to form homogenous lipodisq nanoparticles. Also, it was found that the lipid composition played an important role in making lipodisq nanoparticle samples. Furthermore, lipodisq nanoparticle samples of specific size can be formed by modifying weight ratio, which is useful for biophysical studies of membrane proteins when different sizes of membrane mimic are needed. Future studies will focus on the characterization of membrane proteins in the lipodisq nanoparticle system, especially on the study of protein–protein complexes. Overall, lipodisq nanoparticles are a new membrane mimic system that shows great potential to advance biophysical studies of membrane proteins.

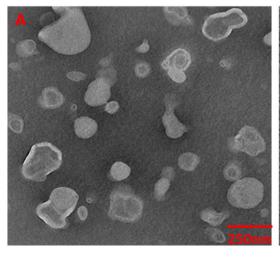
Supplementary data to this article can be found online at http://dx. doi.org/10.1016/j.bbamem.2014.05.008.

# Acknowledgement

This work was generously supported by National Institutes of Health Grant R01 GM108026. Funding was also provided by National Science Foundation (NSF) Grant CHE-1305664.

### References

- J.N. Sachs, D.M. Engelman, Introduction to the membrane protein reviews: the interplay of structure, dynamics, and environment in membrane protein function, Annu. Rev. Biochem. 75 (2006) 707–712.
- [2] G.G. Prive, Detergents for the stabilization and crystallization of membrane proteins, Methods 41 (4) (2007) 388–397.
- [3] C.G. Tate, Practical considerations of membrane protein instability during purification and crystallisation, Methods Mol. Biol. 601 (2010) 187–203.



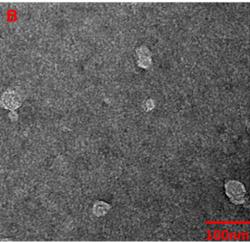


Fig. 4. TEM micrographs of uranyl acetate stained (A) control POPC/POPG vesicles and (B) lipodisq nanoparticles (1/1.25).

- [4] E. van den Brink-van der Laan, V. Chupin, J.A. Killian, B. de Kruijff, Stability of KcsA tetramer depends on membrane lateral pressure, Biochemistry 43 (14) (2004) 4240–4250.
- [5] R.S. Cantor, Lipid composition and the lateral pressure profile in bilayers, Biophys. J. 76 (5) (1999) 2625–2639.
- [6] A.M. Seddon, P. Curnow, P.J. Booth, Membrane proteins, lipids and detergents: not just a soap opera, Biochim. Biophys. Acta 1666 (1–2) (2004) 105–117.
- [7] A.A. De Angelis, A.A. Nevzorov, S.H. Park, S.C. Howell, A.A. Mrse, S.J. Opella, Highresolution NMR spectroscopy of membrane proteins in aligned bicelles, J. Am. Chem. Soc. 126 (47) (2004) 15340–15341.
- [8] A.A. De Angelis, S.J. Opella, Bicelle samples for solid-state NMR of membrane proteins. Nat. Protoc, 2 (10) (2007) 2332–2338.
- [9] T.H. Bayburt, S.G. Sligar, Self-assembly of single integral membrane proteins into soluble nanoscale phospholipid bilayers, Protein Sci. 12 (11) (2003) 2476–2481.
- [10] T. Raschle, S. Hiller, M. Etzkorn, G. Wagner, Nonmicellar systems for solution NMR spectroscopy of membrane proteins, Curr. Opin. Struct. Biol. 20 (4) (2010) 471–479.
- [11] E.R. Geertsma, N.A. Nik Mahmood, G.K. Schuurman-Wolters, B. Poolman, Membrane reconstitution of ABC transporters and assays of translocator function, Nat. Protoc. 3 (2) (2008) 256–266.
- [12] G. Fang, R. Friesen, F. Lanfermeijer, A. Hagting, B. Poolman, W.N. Konings, Manipulation of activity and orientation of membrane-reconstituted di-tripeptide transport protein DtpT of Lactococcus lactis, Mol. Membr. Biol. 16 (4) (1999) 297–304.
- [13] T.L. Lau, C. Kim, M.H. Ginsberg, T.S. Ulmer, The structure of the integrin alphallbbeta3 transmembrane complex explains integrin transmembrane signalling, EMBO J. 28 (9) (2009) 1351–1361.
- [14] D. Lee, K.F. Walter, A.K. Bruckner, C. Hilty, S. Becker, C. Griesinger, Bilayer in small bicelles revealed by lipid-protein interactions using NMR spectroscopy, J. Am. Chem. Soc. 130 (42) (2008) 13822–13823.
- [15] R.R. Vold, R.S. Prosser, Magnetically oriented phospholipid bilayered micelles for structural studies of polypeptides. Does the ideal bicelle exist? J. Magn. Reson. B 113 (3) (1996) 267–271.
- [16] U.H.N. Dürr, M. Gildenberg, A. Ramamoorthy, The magic of bicelles lights up membrane protein structure, 2012.
- [17] T.H. Bayburt, S.G. Sligar, Membrane protein assembly into nanodiscs, FEBS Lett. 584 (9) (2010) 1721–1727.
- [18] J. Borch, T. Hamann, The nanodisc: a novel tool for membrane protein studies, Biol. Chem. 390 (8) (2009) 805–814.
- [19] M.A. Hanson, V. Cherezov, M.T. Griffith, C.B. Roth, V.P. Jaakola, E.Y. Chien, J. Velasquez, P. Kuhn, R.C. Stevens, A specific cholesterol binding site is established

- by the 2.8 Å structure of the human beta2-adrenergic receptor, Structure 16 (6) (2008) 897–905.
- [20] S. Hiller, R.G. Garces, T.J. Malia, V.Y. Orekhov, M. Colombini, G. Wagner, Solution structure of the integral human membrane protein VDAC-1 in detergent micelles, Science 321 (5893) (2008) 1206–1210.
- [21] F. Hagn, M. Etzkorn, T. Raschle, G. Wagner, Optimized phospholipid bilayer nanodiscs facilitate high-resolution structure determination of membrane proteins, 2013.
- [22] I.G. Denisov, Y.V. Grinkova, A.A. Lazarides, S.G. Sligar, Directed self-assembly of monodisperse phospholipid bilayer nanodiscs with controlled size, J. Am. Chem. Soc. 126 (11) (2004) 3477–3487.
- [23] M.C. Orwick, P.J. Judge, J. Procek, L. Lindholm, A. Graziadei, A. Engel, G. Grobner, A. Watts, Detergent-free formation and physicochemical characterization of nanosized lipid-polymer complexes: lipodisq, Angew. Chem. Int. Ed. Engl. 51 (19) (2012) 4653–4657.
- [24] M. Orwick-Rydmark, J.E. Lovett, A. Graziadei, L. Lindholm, M.R. Hicks, A. Watts, Detergent-free incorporation of a seven-transmembrane receptor protein into nanosized bilayer lipodisq particles for functional and biophysical studies, Nano Lett. 12 (9) (2012) 4687–4692.
- [25] I.D. Sahu, R.M. McCarrick, K.R. Troxel, R. Zhang, H.J. Smith, M.M. Dunagan, M.S. Swartz, P.V. Rajan, B.M. Kroncke, C.R. Sanders, G.A. Lorigan, DEER EPR measurements for membrane protein structures via bifunctional spin labels and lipodisq nanoparticles, Biochemistry 52 (38) (2013) 6627–6632.
- [26] T.J. Knowles, R. Finka, C. Smith, Y.P. Lin, T. Dafforn, M. Overduin, Membrane proteins solubilized intact in lipid containing nanoparticles bounded by styrene maleic acid copolymer, J. Am. Chem. Soc. 131 (22) (2009) 7484–7485.
- [27] M. Jamshad, Y.P. Lin, T.J. Knowles, R.A. Parslow, C. Harris, M. Wheatley, D.R. Poyner, R.M. Bill, O.R. Thomas, M. Overduin, T.R. Dafforn, Surfactant-free purification of membrane proteins with intact native membrane environment, Biochem. Soc. Trans. 39 (3) (2011) 813–818.
- [28] R.B. Gennis, Biomembranes: Molecular Structure and Function, Springe-Verlag, New York, 1989.
- [29] A.T. Coey, I.D. Sahu, T.S. Gunasekera, K.R. Troxel, J.M. Hawn, M.S. Swartz, M.R. Wickenheiser, R.J. Reid, R.C. Welch, C.G. Vanoye, C. Kang, C.R. Sanders, G.A. Lorigan, Reconstitution of KCNE1 into lipid bilayers: comparing the structural, dynamic, and activity differences in micelle and vesicle environments, Biochemistry 50 (50) (2011) 10851–10859.
- [30] S.H. Park, S. Berkamp, G.A. Cook, M.K. Chan, H. Viadiu, S.J. Opella, Nanodiscs versus macrodiscs for NMR of membrane proteins, Biochemistry 50 (42) (2011) 8983–8985.