Synthesis, Characterization and Applications of a Perdeuterated Amphipol

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Abstract Amphipols are short amphipathic polymers that can substitute for detergents at the hydrophobic surface of membrane proteins (MPs), keeping them soluble in the absence of detergents while stabilizing them. The most widely used amphipol, known as A8-35, is comprised of a polyacrylic acid (PAA) main chain grafted with octylamine and isopropylamine. Among its many applications, A8-35 has proven particularly useful for solution-state NMR studies of MPs, for which it can be desirable to eliminate signals originating from the protons of the surfactant. In the present work, we describe the synthesis and properties of

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perdeuterated A8-35 (perDAPol). Perdeuterated PAA was obtained by radical polymerization of deuterated acrylic acid. It was subsequently grafted with deuterated amines, yielding perDAPol. The number-average molar mass of hydrogenated and perDAPol, \sim 4 and \sim 5 kDa, respectively, was deduced from that of their PAA precursors, determined by size exclusion chromatography in tetrahydrofuran following permethylation. Electrospray ionization–ion mobility spectrometry–mass spectrometry measurements show the molar mass and distribution of the two APols to be very similar. Upon neutron scattering, the contrast match point of perDAPol is found to be \sim 120 % D₂O. In ¹H-¹H nuclear overhauser effect NMR spectra, its contribution is reduced to $\sim 6 \%$ of that of hydrogenated A8-35, making it suitable for extended uses in NMR spectroscopy. PerDAPol ought to also be of use for inelastic neutron scattering studies of the dynamics of APol-trapped MPs, as well as small-angle neutron scattering and analytical ultracentrifugation.

Keywords Amphipol · A8-35 · Deuteration · Mass spectrometry - NMR

Abbreviations

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Introduction

NMR studies of membrane proteins (MPs) give access to such important biological information as their secondary and tertiary structure, their dynamics, the way they recognize their ligands, etc. Traditionally, these studies are performed in aqueous detergent solutions (Kang and Li [2011\)](#page-14-0). However, the destabilizing character of detergents, the high concentrations of them that is required for NMR experiments, the relatively high temperatures at which

those are carried out and their long duration combine to make it difficult to prepare stable enough MP samples. This has driven the search for more stabilizing environments, such as lipid bicelles (Czerski and Sanders [2000](#page-13-0); Sanders and Landis [1995;](#page-15-0) Sanders and Prosser [1998](#page-15-0)), lipopeptides (McGregor et al. 2003 ; Privé 2009), nanodiscs (Glück et al. [2009](#page-14-0); Hagn et al. [2013](#page-14-0); Mazhab-Jafari et al. [2013](#page-14-0); Raschle et al. [2009,](#page-15-0) [2010](#page-15-0); Shenkarev et al. [2010](#page-15-0)) or amphipols (Zoonens et al. [2005](#page-15-0)) (for reviews, see e.g., Catoire et al. [2014](#page-13-0); Elter et al. [2014](#page-14-0); Planchard et al. [2014](#page-14-0); Popot [2010](#page-14-0); Warschawski et al. [2011;](#page-15-0) Zoonens and Popot [2014](#page-15-0)). Whatever the environment, the size of the resulting complexes entails correlation times in excess of 10 ns. This renders structural and dynamics studies by NMR quite challenging. In addition, the presence of high concentrations of detergent and/or of lipids (in the case of bicelles or nanodiscs) generates interference signals from their ${}^{1}H$ or natural-abundant ce 13 C nuclei, which can hinder the observation of signals from the protein, in particular intramolecular ¹H-¹H dipolar nuclear overhauser effect (NOE) interactions. Even though some NMR pulse sequences can reduce the impact of these interference signals, the best strategy is to work with perdeuterated surfactants (see e.g. Tzitzilonis et al. [2013\)](#page-15-0). This remains true when studying small organic compounds interacting with MPs, because many ${}^{13}CH_n$ chemical shifts of ligands and surfactants overlap. When the cost of labeling isotopically the ligand is prohibitive, deuteration or perdeuteration of the surfactant(s) can be the most efficient approach.

Amphipols (APols) are comprised of a polymeric backbone carrying hydrophilic and hydrophobic moieties (Popot, [2010;](#page-14-0) Popot et al. [2011](#page-15-0); Tribet, Audebert and Popot, [1996](#page-15-0); Zoonens and Popot, [2014](#page-15-0)). Three types of APols have been validated for NMR studies of MPs: carboxylated APols, best represented by A8-35 (Fig. [1\)](#page-2-0) (Catoire et al. [2009](#page-13-0), [2010a,](#page-13-0) [b](#page-13-0); Elter et al. [2014;](#page-14-0) Etzkorn et al. [2013](#page-14-0); Raschle et al. [2010](#page-15-0); Zoonens et al. [2005](#page-15-0)), sulfonated APols (Dahmane et al. [2011](#page-14-0)) and non-ionic APols (Bazzacco et al. [2012](#page-13-0); for reviews, see Elter et al. [2014](#page-14-0); Planchard et al. [2014](#page-14-0); Popot [2010](#page-14-0); Popot et al. [2011](#page-15-0)). A8-35, to date the best characterized and most extensively used APol, is comprised of a polyacrylic acid (PAA) backbone grafted with octylamine and isopropylamine side chains (Tribet et al. [1996](#page-15-0)) (Fig. [1\)](#page-2-0). It has been used to-date for solution NMR studies of five MPs, namely the transmembrane β -barrel (tOmpA) of outer membrane proteins A from Escherichia coli (Zoonens et al. [2005](#page-15-0)) and from Klebsiella pneumoniae (Planchard et al. [2014;](#page-14-0) Renault [2008](#page-15-0)), OmpX from E. coli (Catoire et al. [2009](#page-13-0), [2010b](#page-13-0); Etzkorn et al. [2014](#page-14-0)), bacteriorhodopsin (BR) from Halobacterium salinarum (Etzkorn et al. [2013](#page-14-0), [2014;](#page-14-0) Raschle et al. [2010](#page-15-0)) and the human BLT2 G protein-coupled

Fig. 1 Structure of hydrogenated A8-35 (HAPol) (Tribet et al. [1996](#page-15-0)) and partially deuterated A8-35 (DAPol) (Gohon et al. [2006](#page-14-0), [2004](#page-14-0)). In perDAPol, all of the non-exchangeable hydrogens are deuterated

receptor of leukotriene B_4 (LTB₄) (Catoire et al. [2010a](#page-13-0)). For some of these experiments, a partially deuterated version of A8-35 (hereafter, DAPol) has been used, in which the PAA main chain is hydrogenated, but the octylamine and isopropylamine side chains are deuterated (Gohon et al. [2004](#page-14-0), [2006](#page-14-0)). The use of DAPol has permitted, in particular, mapping of the points of contact between the polymer and tOmpA, by comparing the width of the resonance peaks of amide protons in the presence of either DAPol or unlabeled, hydrogenated A8-35 (hereafter, HAPol) (Zoonens et al. [2005](#page-15-0)). Following DAPol-assisted folding of deuterated BLT2, measurements of transferred NOE signals led to the determination of the conformation adopted by hydrogenated leukotriene B_4 (LTB₄) when bound to its specific recognition site at the surface of the receptor (Catoire et al. [2010a\)](#page-13-0), an approach that may be widely applicable (Banères, Popot and Mouillac, [2011](#page-13-0); Catoire et al. [2011](#page-13-0)). Part of the magnetization transfer signals between $LTB₄$ protons could not be observed, however, because they were masked by signals originating from the hydrogenated backbone of DAPol (Catoire et al. [2010a](#page-13-0)) (reviewed in Planchard et al. [2014\)](#page-14-0).

In the present article, we describe the synthesis and properties of a perdeuterated form of A8-35 (perDAPol). Synthesis of an exact mimic of HAPol required re-examination of the average degree of polymerization of the commercial hydrogenated PAA used for HAPol and DAPol synthesis. This was followed by the synthesis of an equivalent deuterated PAA, which was obtained by radical telomerization of its perdeuterated acrylic acid precursor. The deuterated PAA was then successively grafted with deuterated octylamine and deuterated isopropylamine. Upon dissolution into aqueous buffers, the resulting perDAPol self-organizes into particles with the same behavior in aqueous SEC as those formed by HAPol and DAPol. The mass distribution of the three forms of A8-35 was studied by electrospray ionization–ion mobility spectrometry–mass spectrometry (ES-IMS-MS) (Leney et al. [2012\)](#page-14-0) and found to be comparable. The contrast match point (CMP) of perDAPol was established by small-angle neutron scattering (SANS). Homonuclear and heteronuclear NMR spectroscopy was performed on the three forms of A8-35. Partial deuteration reduces proton signals of A8- 35 by \sim 68 %, perdeuteration by \sim 94 %. The applications of perDAPol to NMR, inelastic neutron scattering (INS), SANS and analytical ultracentrifugation (AUC) studies of MPs are discussed.

Materials and Methods

Materials

Tetrahydrofuran (THF), purchased from VWR (FontenaysousBois, France), was distilled over calcium hydride before use. Poly(acrylic acid) (PAA) and other chemicals were from Sigma-Aldrich (Saint Quentin Fallavier, France). A single and identical weight-average molar mass $(\bar{M}_{w} \approx 5,000 \text{ g mol}^{-1})$ was given by the purchaser for all PAA batches used and/or cited in this paper. Acrylic acid-d4 was from Cil (Courbevoie, France) and used as received. A8- 35 (batch FGH29), A8-75 and DAPol-4 were synthesized and characterized as described in refs. (Gohon et al. [2004](#page-14-0), [2006\)](#page-14-0).

Buffer

Tris buffer: Tris/HCl 20 mM, NaCl 100 mM, pH 8.0.

Methods

Synthesis of a Perdeuterated Poly(acrylic acid) Precursor

The overall scheme of the synthesis of perdeuterated PAA and perDAPol is shown in Fig. [2](#page-3-0). In a 25-mL two-neck round-bottom flask, acrylic acid-d₄ $(1.06 \text{ g}, 14 \text{ mmol})$ was dissolved into 9 mL of a 8:1 water/THF mixture and the solution freed from O_2 with argon by multiple cycles of freeze–pump–thaw degassing under vacuum. Prior to polymerization, the solution was flushed with a stream of argon. Thioglycolic acid (TGA) (0.074 g, 0.8 mmol) and 1 mL (26.2 mg, 0.16 mmol) of a previously degassed 0.16 M 2,2'-azoisobutyronitrile (AIBN) (recrystallized twice in ethanol) solution in THF were injected into the aqueous mixture. The flask was sealed and kept under

Permethylated perDAPol

Fig. 2 Synthesis of perDAPol, starting from acrylic acid-d₄, and permethylation of polymers for SEC analysis. a TGA, AIBN in 80:20 water/THF at 65 °C, 4 h; **b** *n*-octylamine-d₁₇, DCI, NMP, 60 °C, 1 h,

vigorous stirring in an oil bath at 65° C for 4 h. It was then quickly brought to 0° C (ice bath) and the polymerization reaction quenched by flushing the mixture with air and cooling at -30 °C (dry ice in ethanol). The flask was brought to room temperature, and the solution added dropwise to 100 mL of 0.2 N NaOH in ethanol under vigorous stirring. The precipitate was filtered, rinsed with 100 mL of absolute ethanol and dissolved in 10 mL of water purified on an A10 Advantage Millipore system (mQ water). After acidification with 1.5 mL of concentrated HCl, the water was evaporated under reduced pressure and the dry PAA triturated in 20 mL of THF. The precipitated sodium chloride was removed by filtration, and the filtrate concentrated to a 10-mL final volume. The THF solution was added dropwise to 100 mL of petroleum ether (or n-heptane) under vigorous stirring, and the resulting precipitate recovered after filtration. The polymer was dried under vacuum and dissolved in 20 mL of mQ water. Freeze-drying of the solution yielded 1.0 g (94 % yield) of the expected perdeuterated PAA (PAA-d-3, batch 3).

Synthesis of perDAPol

PAA-d-3 (0.86 g) (11.46 mmol carboxylic acid) was dissolved into 17 mL of N-methylpyrrolidone (NMP) and stirred at 60 °C in an oil bath for 4 h. Octylamine-d₁₇ (0.418 g, 2.86 mmol) and dicyclohexylcarbodiimide (DCI) (0.59 g, 2.86 mmol) in solution in NMP (2 mL and 4 mL, respectively) were successively added dropwise to the reaction medium (Fig. 2). After complete addition of the

then RT 12 h, then isopropylamine-d₇, DCI, NMP, 50 °C, 1 h, then RT 12 h, then NaOH; c H_3O^+ ; d TMSCHN₂, THF:MeOH 9:1

reagents, the mixture was kept at 60° C for 1 h, then brought to room temperature and stirred overnight. The precipitate of dicyclohexylurea (DCU) was removed by filtration and 1-N-hydroxybenzotriazole (HOBt) (0.62 g, 4.73 mmol) added to the filtrate, which was heated at 50 °C. Isopropylamine-d₇ (0.3 g, 4.54 mmol) and DCI (0.994 g, 4.82 mmol) were added to the reaction medium, which was kept at 50 \degree C for a further hour after complete addition of the reactants (Gohon et al. [2004,](#page-14-0) [2006\)](#page-14-0). After overnight stirring at room temperature, followed by filtration, the filtrate was treated with sodium methoxide (1.5 g, 27.8 mmol) and stirred for 15 min before being poured into 300 mL of mQ water. The mixture was stirred for 30 min, filtered, and the filtrate added dropwise to 20 mL of 2 N HCl under vigorous stirring. The precipitate was recovered by filtration and suspended into 100 mL of water under vigorous stirring; 1 mL of a 30 % NaOH solution in mQ water was added. The resulting solution was centrifuged $(5,000 \times g)$ for 30 min), and the supernatant filtered on Millipore filters. After three further cycles of precipitation in acidic solution/dissolution in basic medium, the solution was dialyzed against mQ water for 4 h and freeze-dried to yield 1.43 g (90% yield) of the expected perdeuterated A8- 35 (perDAPol, batch 1) as a white solid.

Analysis of the Chemical Composition of the Polymers

The chemical composition of the polymers was analyzed by 1 H and 13 C NMR spectroscopy as previously described (Gohon et al. [2004](#page-14-0), [2006\)](#page-14-0). Results are reported in Table [1.](#page-4-0)

PAA	Method	Measured \overline{M}_n (kDa)	Measured \overline{M}_w (kDa)	$D = \bar{M}_{w}/\bar{M}_{n}$	\overline{DP}_n
PAA_{CT} -1*	$SECa$ (H ₂ O)	(~ 3.1) **	6.2	(~ 2) **	(~ 35) **
$PAACT-2***$	$SECa$ (H ₂ O)	7.5	12.0	1.6	80
$PAA-1$	$SECa$ (H ₂ O)	6	15.5	2.6	64
	SEC ^b (THF)	2.5	5.1	2.1	27
	SEC ^c (THE)	3.5	6.8	1.9	38
PAA-d-3 (batch 3)	SEC ^b (THF)	3.5	7.8	2.2	37
	SEC ^c (THF)	3.3	6.6	2.0	35

Table 1 Estimation of the average molar masses and dispersities of various batches of PAA by different SEC systems

SEC was performed either (1) in aqueous media by double detection ("universal" calibration with PEO) according to ^a Tribet et al. [1996](#page-15-0)* and Gohon et al. [2004](#page-14-0)^{**} or (2) in THF after permethylation and monitored by either ^bsimple (RI and calibration with PS) or ^ctriple detection (RI, dV, RALS/LALS). PAA-1 (commercial) was used as precursor for the synthesis of the batches of HAPol and DAPol used in the present work, PAA-d-3 (home made) for that of perDAPol. \bar{M}_n and \bar{M}_w are given for the polymers under their sodium salt form and neglect the contribution of deuteration, the ratio of the molar mass of the deuterated versus the hydrogenated form (\sim 1.06 for DAPol, \sim 1.08 for perDAPol) being inferior to the experimental error (\sim 10 %). *D* is the molar mass dispersity. *From Tribet et al. [1996.](#page-15-0) \bar{M}_n and \overline{DP}_n were not reported in this article; their values were deduced assuming $D \approx 2$. ** Estimated D value and resulting estimate of \bar{M}_n and \overline{DP}_n . ***From Gohon et al. [2004](#page-14-0)

Characterization of the Molar Mass of the Polymers

Estimation of the Average Molar Mass of the Polymers by SEC Analysis

• SEC analysis of PAA in aqueous buffer

SEC analyses of PAAs in aqueous buffer were carried out as described in refs. (Gohon et al. [2006;](#page-14-0) Tribet et al. [1996](#page-15-0)). In both cases, PAAs were injected under their sodium salt form on a Shodex OH pack column (Waters) calibrated with poly(ethylene oxide) (PEO) standards (''universal'' calibration) and eluted with a 0.5 M LiNO₃ aqueous solution at 40 C. Injections were monitored by double detection [refractive index (RI) and differential viscometry (dV) detection]. SEC experiments were employed to determine the weight-average molar mass \bar{M}_w and the number-average molar mass \overline{M}_n of the PAAs but, as described below (see "Results"), masses were overestimated.

• SEC analyses of PAA in organic solvent.

Prior to SEC analysis, the AA units of all PAAs were modified by methylation of the carboxylic acid groups using trimethylsilyldiazomethane according to the method described by Couvreur et al. [2003.](#page-13-0) Briefly, 50 mg of the PAA was dissolved in 10 mL of a 9:1 THF/methanol mixture and treated with 0.5 mL of trimethylsilyldiazo-methane 2 M in diethyl ether (Presser and Hüfner [2004](#page-15-0)). The solvent was evaporated under reduced pressure, and the permethylated polymer, poly(methyl acrylate), was dried under vacuum for 12 h.

Measurements were performed with a Viscotek TDAmax system from Malvern Instruments that consists of an integrated solvent and sample delivery module (GPCmax) and a Tetra Detector Array (TDA) including a differential

detector RI, a right (90 \degree) and a low (7 \degree) angle light scattering (LS) detector (RALS/LALS), a four-capillary differential viscometer and a diode array UV detector. THF was used as the mobile phase at a flow rate of 1 mL min^{-1} and toluene as a flow rate marker. All polymers were injected $(100 \mu L)$ of solution) at a concentration of 10 g L^{-1} after filtration through a 0.45-um pore-size membrane. The separation was carried out on three Polymer Laboratories columns $(3 \times$ PLgel 5-µm Mixed C; 300×7.5 mm) and a guard column (PL gel 5 µm). Columns and detectors were maintained at 40 °C. The OmniSEC 4.6.2 software was used for data acquisition and data analysis. \overline{M}_n , \overline{M}_w , and the molar mass dispersity, $D = \overline{M}_w / \overline{M}_n$, were calculated with a calibration curve based on narrow PS standards (from Polymer Standard Services), using only the RI detector (simple detection). Indeed, regarding the Mark–Houwink–Sakurada parameters in THF, the polystyrene calibration is appropriate for poly(methyl acrylate) (Couvreur et al. [2003;](#page-13-0) Velasquez et al. [2013\)](#page-15-0). In addition, \overline{M}_n was also calculated using the combined signals from RALS/LALS, RI and the viscometer (triple detection). All molar mass were recalculated and expressed for the sodium salt of each polymer (see Tables).

Estimation of the Average Molar Mass and Dispersity of the Polymers by Mass Spectrometry

ES-IMS-MS experiments were performed on the sodium salt of the polymers on a Synapt HDMS mass spectrometer (Micromass UK Ltd./Waters, Manchester, UK) as described in ref. (Leney et al. [2012\)](#page-14-0). Samples were introduced into the instrument by direct infusion using gold-coated borosilicate capillaries. A capillary voltage of 1.8 kV was

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applied, and a source temperature of 80 $^{\circ}$ C was used. A cone voltage of 70 V was used, and trap and transfer collision voltages were both set to 5 V. Ion mobility separation was performed by ramping the wave height from 4.5–28.5 V at a speed of 400 m s^{-1} . An aqueous solution of CsI was used for m/z calibration. All data were acquired over the m/z range 500–5,000 and the raw data processed by use of MassLynx $v.4.1$ and Driftscope $v.3.0$ software (Micromass UK Ltd/Waters Corp., Manchester, UK).

Average masses are expressed as the number-average molar mass, \overline{M}_n , and the weight-average molar mass, \overline{M}_w , with

$$
\bar{M}_n = \frac{\sum_i N_i \cdot M_i}{\sum_i N_i}
$$

and

$$
\bar{M}_{w} = \frac{\sum_{i} m_{i} \cdot M_{i}}{\sum_{i} m_{i}} = \frac{\sum_{i} N_{i} \cdot M_{i}^{2}}{\sum_{i} N_{i} \cdot M_{i}}
$$

where N_i is the number of moles, M_i is the molar mass and m_i the mass of species *i*.

Characterization of the association behavior of APols in aqueous buffer

APols were dissolved at 100 g L^{-1} in mQ water. The stock solution was diluted at 10 g L^{-1} in Tris buffer and kept under stirring for 48 h at room temperature; 80 μ L of the resulting sample were injected at room temperature on an Akta purifier 10 FPLC system (GE Healthcare) equipped with a calibrated Superose 12 10/300 GL column (GE Healthcare) (see Gohon et al. [2004,](#page-14-0) [2006](#page-14-0), for detailed procedure). The flow rate was set at 0.5 mL min⁻¹. Injection buffer was 0.1 M NaCl, 20 mM Tris/HCl, pH 8. The elution profile was monitored at 220 nm.

Small-Angle Neutron Scattering (SANS)

Sample Preparation

PerDAPol was dissolved at 7.2 g L^{-1} either in Tris/HCl 20 mM, pH 8.0, 100 mM NaCl, 2 mM EDTA, H2O or in Tris/HCl 20 mM, pD 7.6 100 mM NaCl, 2 mM EDTA, D_2O . UV spectra of the two solutions confirmed that their concentrations were identical. The H_2O and D_2O samples were mixed in the appropriate proportions to yield the desirable ratios of H_2O/D_2O .

SANS Experiments

SANS experiments were performed on the Bio-SANS instrument of the High Flux Isotope Reactor of the Oak Ridge National Laboratory (Lynn et al. [2006\)](#page-14-0). SANS data were taken at sample-to-detector distances of 1.1 and 6.8 m. The neutron wavelength, λ , was set to 6 A^{\AA}, and the

wavelength spread, $\Delta \lambda/\lambda$, was set to 0.15. The two instrument settings provided a q-range from 0.006–0.70 \AA^{-1} . Scattered neutrons were collected by a 1 m \times 1 m twodimensional (2D) linear position-sensitive detector having 192×256 pixels (Berry et al. [2012](#page-13-0)). The solutions and associated solvent backgrounds were measured at 20 \degree C in 1-mm path-length quartz cells (Hellma U.S.A., Plainview, NY, USA). The 2D data were corrected for detector pixel sensitivity, dark current and scattering by the buffer. The 2D reduced data were azimuthally averaged to yield the 1D scattering intensity $I(q)$ versus q, where $q = 4\pi \sin(\theta)/\lambda$ and 2θ is the scattering angle from the incident beam. The 1D profiles from the two detector distances used on Bio-SANS were merged to produce the full scattering intensity profile using software developed at the National Institute of Standards and Technology (Kline [2006](#page-14-0)). A constant background representing the difference in incoherent scattering resulting from the differences in hydrogen content of the samples and backgrounds was subtracted from the 1D profile (Fig. [5](#page-9-0)a).

The radius of gyration, $R_{\rm g}$, was determined with the Guinier approximation shown in Eq. 1 (Guinier and Fournet [1955\)](#page-14-0):

$$
I(q) = I(0)e^{-q^2 R_g^2/3}
$$
\n(1)

where $I(0)$ is the forward scattering intensity. A linear fit of ln($I(q)$) versus q^2 ("Guinier plot") (Fig. [5](#page-9-0)b) provides $I(0)$ and R_g from the intercept and slope, respectively. The molar mass of the particle is provided by the Guinier fitting through the relation in Eq. 2:

$$
M = \frac{I(0) \times N_A}{C((\rho - \rho_{\text{buffer}}) \times V_s)^2}
$$
 (2)

where N_A is Avogadro's number, C is the concentration of the samples in solution, the neutron scattering length densities of the sample and buffer are ρ and ρ_{buffer} , respectively, and V_s is the specific volume of the scattering particle.

The contrast match point of perDAPol is determined by plotting the square root of the forward scattering intensities as a function of the D_2O ratio (v/v) (Fig. [5c](#page-9-0)). The intercept of the linear fitting of the plot with the y axis (zero intensity) determines the contrast match point.

Results

Re-examination of the Degree of Polymerization of the Hydrogenated PAA used for Synthesizing A8-35

The average degree of polymerization of the hydrogenated PAA (PAA-1) used for synthesizing A8-35 was determined by SEC analysis in organic solvent (Table [1](#page-4-0)). This study

Table 2 Chemical composition and expected number-average molar mass (\bar{M}_n) of various APols

APol	Precursor	$x(\%)$	$v(\%)$	$z(\%)$	Expected M_n (unrevised) ^a (kDa)	Expected M_n $(revised)^b$ (kDa)	Expected M_n $(revised)^c(kDa)$
Hydrogenated A8-75	PAA-1	75	25		7.5	3.1	4.4
HAPol (FGH29)	PAA-1	35	25	40	8.0	3.3	4.7
DAPol (batch 4)	PAA-1	35	$25^{\rm d}$	40 ^d	8.4	3.6	5.2
perDAPol (batch 1)	$PAA-d-3$	35	$25^{\rm d}$	40 ^d	-	4.7	-

x, y and z are the ratios (expressed in percentage of PAA units) of each moiety (free carboxylate, octylamide and isopropylamide, respectively) in the polymer (see Fig. [1\)](#page-2-0). \bar{M}_n is given by the relation $\bar{M}_n = \overline{DP}_n \times M_0$, where M_0 is the average molar mass of the monomer, defined by the relations $M_0 = 94 x + 183 y + 113 z$, $M_0 = 94 x + 200 y + 120 z$ and $M_0 = 97 x + 203 y + 123 z$ for HAPol, DAPol and perDAPol, respectively. \overline{DP}_n is the number-average degree of polymerization of the PAA precursor (see Table [1](#page-4-0)). The \overline{DP}_n of PAA was determined by SEC either ^ain aqueous buffer (Gohon et al. [2004](#page-14-0)) or in organic medium with either simple ^b or triple detection^c (this work; *cf.* Table I). ^dThe values of y and z for DAPol and perDAPol could not be separately determined (see "Materials and Methods") and are nominal values. The percentage of carboxylates in these polymers (x) was determined by ¹³C NMR. Expected \bar{M}_n values, based on the \overline{DP}_n of PAA and the degree of derivatization, are calculated for the sodium salt and take into account the contribution of deuteration. Because earlier SEC experiments performed in aqueous medium^(a) were not consistent with those performed later in organic medium ^(b,c), \bar{M}_n deduced from the \overline{DP}_n of the PAA estimated according to procedures (b,c) are revised \bar{M}_n

made it possible to clarify the difference between the degree of polymerization estimated in ref. (Tribet et al. [1996\)](#page-15-0) ($\overline{DP}_w \approx 70$, *i.e.* $\overline{DP}_n \approx 35$ assuming a molar mass dispersity of 2; PAA_{CT} -1 in Table [1\)](#page-4-0), using SEC in aqueous solution with a RI calibration based on PEO standards and a differential viscometry (dV) detection and that found later, using a similar procedure, $(\overline{DP}_w \approx 130, \overline{DP}_n \approx 80)$ (Gohon et al. 2004) (PAA_{CT}2 in Table [1](#page-4-0)). The present data were obtained by SEC of permethylated PAA in an apolar organic solvent (THF) using either a simple detection mode (differential RI detector) or a triple detection (possessing as a key element a static light scattering detector). The results tally with the estimate of ref. (Tribet et al. [1996](#page-15-0)), suggesting that the \overline{DP}_n of the PAA was overestimated in ref. (Gohon et al. [2004\)](#page-14-0) (see Table [1\)](#page-4-0). Several factors that may explain the inconsistency between the estimates of refs. (Tribet et al. [1996](#page-15-0)) and (Gohon et al. [2004](#page-14-0)) are indicated in the '['Discussion'](#page-10-0)' section. On the basis of the new measurements, the \overline{M}_n of the sodium salt of hydrogenated A8-35 is expected to be \sim 4 kDa, rather than 8–10 kDa as assumed in refs. (Gohon et al. [2004](#page-14-0), [2006](#page-14-0); Table 2). The lower estimate was confirmed qualitatively by MS analysis of the final products (see below). As discussed below, this revision has no practical consequence under most circumstances, where it is the mass of the APol particle or MPassociated APol that matters, not the number of molecules it is comprised of (see "[Discussion](#page-10-0)").

Synthesizing a Deuterated PAA with a Matching Length

In order to produce a perdeuterated version of A8-35 (perDAPol) that be chemically as similar as possible to the unlabeled form (HAPol), it has to be synthesized starting from a perdeuterated PAA of similar chain length (\overline{DP}_n)

 \approx 35) and molar mass dispersity ($D \approx 2$) as the unlabeled PAA and according the same protocol (hydrophobic modification). Perdeuterated PAA was synthesized by radical polymerization of the perdeuterated acrylic acid (AA_{d4}) in the presence of a telogen. The protocol used was adapted from that described by Loubat, Javidan and Boutevin, [2000.](#page-14-0) Polymerization was initiated by heat decomposition of AIBN, TGA being used as the telogen or chain transfer reagent. The use of TGA allows to limit the growth of the macroradicals formed during the propagation step of the polymerization and yields polymers of restricted length, namely telomers (see refs. Loubat et al. [2000](#page-14-0), and Pavia et al. [1992,](#page-14-0) for further details). With this approach, the only parameters involved in the control of the number-average degree of polymerization (\overline{DP}_n) are the initial concentration of TGA, $[TA]_0$, and the transfer constant C_t of the telogen. These three terms are related by Eq. 3 (Loubat et al. [2000](#page-14-0)), which is accurate only at the beginning of the reaction (degree of advancement of the reaction $\langle 10 \, % \rangle$):

$$
\frac{1}{\overline{DP_n}} = C_t \frac{[TA]_0}{[AA]_0} \tag{3}
$$

In the ideal case, $C_t = 1$ and the final length of the telomer $\overline{DP}_n \approx [AA]_0/[TA]_0$, this relation assuming that TA and AA are consumed in the same proportion during the reaction. It has been reported that this ideal condition is obtained with TGA when the polymerization of AA is performed in a 80:20 water/THF mixture (Loubat et al. 2000). Our first attempts at telomerizing AA_{d4} using these conditions gave unsatisfactory results (unexpected \overline{DP}_n and/or poor yields), but sample PAA-d-3 (Table [3\)](#page-7-0) yielded finally the desired \overline{DP}_n (~35). Subsequent experiments gave similar results as those reported by (Loubat et al. [2000](#page-14-0)) (Table [3\)](#page-7-0), illustrating that the method could be mastered only after several attempts.

Table 3 Batches of perdeuterated PAA synthesized by telomerization of perdeuterated acrylic acid

Polymer	$R_0 = (AA)_0/(TA)_0$ $(= expected \overline{DP}_n)$ (achieved)	DP _n	Measured M_n (kDa)	$D = M_{w}$ $M_{\rm n}$
$PAA-d-3$	17.5	37	3.5	2.2
$PAA-d-4$	14.0	15	1.4	1.6
$PAA-d-5$	17.5	21	2.0	1.8
$PAA-d-6$	25.0	25	2.3	2.1
$PAA-d-7$	14.0	14	1.3	17

Telomerization was carried out as described in ref. (Loubat et al. [2000\)](#page-14-0). The number-average molar mass of the resulting PAAs was estimated by SEC performed in THF after permethylation. Injections were monitored by simple (RI) detection, using PS standards. The values are calculated for the sodium salt form and neglect the contribution of deuteration. R_0 is the ratio of the initial monomer concentration, $[AA]_0$, to the initial telogen agent concentration, $[TA]_0$, corresponding to the expected \overline{DP}_n . All telomerizations were performed at the same initial concentration of initiator $[(AIBN)₀ = 1.6 \times 10^{-2} M]$ and of perdeuterated acrylic acid $[(AA)₀ = 1.4 M]$. Batch PAA-d-3 was used for the synthesis of the batch of perDAPol described in the present work (batch 1)

Synthesis, Purification and Chemical Analysis of perDAPol

The synthesis and purification of perDAPol were carried out according to the same procedure as used to synthesize HAPol (Gohon et al. [2004](#page-14-0), [2006\)](#page-14-0). Perdeuterated PAA was hydrophobically modified in two steps in NMP, first with perdeuterated octylamine and then with perdeuterated isopropylamine, in the presence of DCI as a coupling reagent. Purification of the polymer was achieved by four cycles of precipitation/dissolution in aqueous solution. Dialysis of the final basic solution followed by freezedrying yielded perDAPol quantitatively as its sodium salt. The overall degree of modification was estimated by 13 C NMR spectroscopy to be close to 65 % (leaving \sim 35 % of free carboxylates), as expected for 100 % grafting efficiency. The relative amount of grafted octylamine and isopropylamine could not be accurately determined by the same technique as for HAPol because of 13 C-D heteronuclear coupling. Indeed, the relaxation time (τ_R) of each carbon depends on its degree of deuteration (the higher this degree, the lower τ_R). This makes it difficult to derive the degree of grafting of each type of chain from the intensity of the signals, as is done for hydrogenated A8-35 (Gohon et al. 2004 , 2006). Decoupling deuterium and ¹³C could permit to circumvent this problem. This has not been implemented here. In our experience, however, the amount of grafted alkylamine has always been found to be close to that aimed for. The extent of grafting in perDAPol can therefore be safely assumed to be close to 25 % octylamine and 40 % isopropylamine, consistent with the solution behavior of the polymer (see below).

ESI-IMS-MS Comparison of the Mass Distribution of A8-35, DAPol and perDAPol

The heterogeneity of APols and other polymers makes their study challenging. However, the advent of electrospray ionization–ion mobility spectrometry–mass spectrometry (ESI-IMS-MS) has significantly improved the capability of MS to study complex polymeric mixtures (Weidner and Trimpin [2010](#page-15-0)). IMS separates ions based on their drift time though a neutral buffer gas under the influence of a weak electric field, the drift time depending on the ions' shape (collision cross-sectional area) and charge state (Kanu et al. [2008](#page-14-0)). Coupled with MS, this provides an extra dimension of separation which is particularly useful in the study of polymers.

The three-dimensional (3D) ESI-IMS-MS spectra of A8- [3](#page-8-0)5 and perDAPol are shown in Fig. 3. Ions with $+1$, $+2$, $+3$ and $+4$ charge states that overlap in MS can be separated by IMS, observed and identified in the spectra, as indicated. There are also low-intensity ions with higher charge states that cannot be assigned reliably in the mass spectra. The ESI-IMS-MS data indicate that, as expected, the two APols have broad mass distributions. Note that, as species with higher mass-to-charge ratios (m/z) are less readily desolvated and detected, the distribution observed may be skewed toward low masses (Hernandez and Robinson [2007](#page-14-0)). Nevertheless, the approach is extremely useful on a comparative basis: The similar mass distributions observed in the two spectra indicate that A8-35 and perDAPol are highly comparable in terms of their heterogeneity and composition.

The weight-average (\bar{M}_w) and number-average (\bar{M}_n) molar masses of each APol were calculated from the MS data by first converting from m/z to mass and then proceeding as described in ref. (Gohon et al. [2006\)](#page-14-0). For A8-35, \overline{M}_w = 4.8 kDa and \overline{M}_n = 3.8 kDa, corresponding to a polydispersity index $D = \overline{M}_n / \overline{M}_w \approx 1.26$. For perDApol, \overline{M}_w = 5.2 kDa, \overline{M}_n = 4.2 kDa, $D \approx 1.24$. The most abundant ions in the spectra correspond to species with molecular masses of 3.4 kDa (A8-35) and 3.6 kDa (perD-APol). These values compare well with the revised mass estimates of A8-35 (Table [2](#page-6-0)), even though they may be skewed, as outlined above, due to high mass polymer species potentially being ionized and detected less efficiently, and to the inability to assign conclusively all the low intensity high m/z species. The difference between the two values of Ð, 1.24 and 1.26, is much smaller than the experimental error, indicating a comparable molar mass dispersity. They are, however, lower than the value expected, $D \approx 2$, possibly due to the skewing effect mentioned above. Furthermore, the similarity of A8-35 and perDAPol in terms of chain length is confirmed by their

Fig. 3 ESI-IMS-MS spectra of a A8-35 and b perDAPol $(1 g L^{-1}$ in 100 mM ammonium hydrogen carbonate, pH 8.0). The contour plots indicate the drift time of ions in the IMS cell (ms) as a function of m/z

mass ratios: 1.088 for \bar{M}_w , 1.101 for \bar{M}_n , 1.059 for the most abundant ions, to be compared with the value of 1.081 expected from the extent of deuteration if the degrees of polymerization are identical. MS, therefore, confirms that HAPol and perDAPol are essentially identical but for isotopic labeling.

Self-assembly of perDAPol in Aqueous Buffer

The behavior of perDAPol in aqueous buffer was characterized by SEC on a Superose 12 column (Gohon et al. [2004,](#page-14-0) [2006\)](#page-14-0). The elution profile of perDAPol is identical to that obtained for A8-35 (Fig. 4). In both cases, one main sharp peak is followed by a smaller, broader one. The main peak at $V_e \approx 12$ mL comprises the particles formed by self-association of the polymer in aqueous buffer, whose Stokes radius R_S is \sim 3 nm (Gohon et al. [2006\)](#page-14-0). Its sharpness width at half height (WHH) \approx 1 mL—reflects the narrow size distribution of the particles. The smaller but broader peak (WHH \approx 2.1 mL) observed at $V_e \approx 14.2$ mL is characteristic of A8-35 samples and indicates the presence of smaller ($R_S \approx 2$ nm), heterogeneous species, which have not been formally identified. They may be short, probably poorly hydrophobized polymers that subsist in solution under their monomeric state, or loosely associated with the particles.

Fig. 4 Overlay of the chromatograms obtained after the injection of HAPol (unlabeled A8-35) (circles) or perDAPol (batch 1) (squares) in Tris buffer on a Superose 12 10–300 GL column. The elution profile was monitored at 220 nm. The two profiles have been normalized to the same maximum. *a.u.* arbitrary units. V_0 , V_T excluded and total volumes, respectively

Neutron Scattering contrast match point

SANS is widely used to determine the mass, shape and arrangement of components in soluble particles (Zaccai and Jacrot, [1983\)](#page-15-0). For MPs, being able to contrast match the surfactant by adjusting the H_2O/D_2O ratio in the buffer is an important asset. The contrast match point (CMP) of perDAPol was determined by measuring the intensity scattered at zero angle $[I(0)]$ by perDAPol particles as a function of the percentage of D_2O in the solution (Fig. [5](#page-9-0)). The CMP obtained by extrapolation of $I(0)$ to zero intensity, \sim 120.5 % D₂O, is in perfect agreement with that expected theoretically $(\sim 120 \%)$ on the basis of the composition of perDAPol (this work) and the specific volume of A8-35 (Gohon et al. [2004\)](#page-14-0) (G. Zaccai, personal communication).

Less expected are the smaller R_g (1.9 \pm 0.2 nm) and mass (\sim 18.8 kDa) obtained for perDAPol particles from the Guinier plots in Fig. [5](#page-9-0)b, which do not tally with those determined previously for HAPol and DAPol $(2.4 \pm$ 0.2 nm and $39-40 \pm 5$ kDa, respectively) (Gohon et al. [2006](#page-14-0)). The two measurements are consistent one with another in indicating \sim 2 \times smaller particles, but it is highly unlikely that such an effect originates from isotopic labeling. Further investigations will be needed to establish its origin (possibly, but somewhat surprisingly, the effect of one component of the buffer, EDTA, which was not present in earlier SANS studies; about the effect of EDTA on the dispersity of A8-35 particles and MP/A8-35 complexes, see Picard et al. [2006](#page-14-0); Catoire et al. [2010b\)](#page-13-0).

Fig. 5 SANS analysis of perDAPol particles. a SANS by perDAPol at different D_2O/H_2O ratios. **b** Guinier plots of the data in A. Dashed vertical lines define the region chosen ($\sim 0.6 \le (q \times R_g) \le \sim 1.7$) to determine the molar mass and R_g of the particles. All samples were at 7.2 $g L^{-1}$. c Determination of the contrast match point

Membrane Protein Trapping by perDAPol

In order to check on the ability of perDAPol to trap MPs, BLT2, a G protein-coupled receptor of leukotriene B4

Fig. 6 Comparative size exclusion chromatography analysis of human G protein-coupled receptor BLT2 folded and trapped in either fully protonated A8-35 (HAPol, dashed line) or perDAPol (solid line) at 1:5 w/w BLT2/A8-35 ratio. Injections of 100 - μ L samples onto a Superose 12 10–300GL column. Elution profiles were recorded at 220 nm. Protein concentration was ~ 1 g L⁻¹. The equilibration and running buffer for both experiments was Tris/HCl (pH 8.0) supplemented with 100 mM NaCl. V_0 , excluded volume; V_T , total volume. Asterisk indicates the elution peak of residual salts. Elution volumes correspond to apparent Stokes radii of 5.4 and 5.5 nm for BLT2/ HAPol and BLT2/perDAPol complexes, respectively

(Yokomizo et al. [2000\)](#page-15-0), was folded and trapped using either HAPol or perDAPol, according to the procedure described in refs. (Catoire et al. [2010a](#page-13-0); Dahmane et al. [2009](#page-14-0)). Comparative SEC analysis of the BLT2/HAPol and BLT2/perDAPol complexes revealed almost identical particle sizes (Stokes radii of \sim 5.4 and \sim 5.5 nm, respectively) and size distributions (peak width at mid-height \sim 1.7 mL) (Fig. 6). These experiments suggest that folding and trapping procedures that were formerly defined for various PMs using HAPol, e.g. those used for GPCRs (Dahmane et al. [2009\)](#page-14-0), can be reproduced with perDAPol without the need of a new optimization.

Comparison of HAPol, DAPol and perDAPol 2D¹H,¹H NOESY and ¹H,¹³C HSQC NMR Spectra

 $2D⁻¹H₁¹H$ NOE spectroscopy (NOESY) and $¹H₁¹³C$ heter-</sup> onuclear single quantum correlation (HSQC) spectra of HAPol, DAPol and perDAPol solutions, recorded at identical APol concentrations (20 g L⁻¹ in pure D₂O), are shown in Figs. [7](#page-10-0) and [8.](#page-11-0) In 1D and $2D⁻¹H$ homonuclear experiments (Fig. [7\)](#page-10-0), 1 H signals are reduced, as compared to HAPol, by 68 % for DAPol and by 94 % for perDAPol. In particular, between 0 and 2.5 ppm, the intensity of 1 H signals is considerably reduced in perDAPol, as shown by

Fig. 7¹H 1D a and 2D NOESY **b** homonuclear NMR spectra of HAPol, DAPol and perDAPol. In (a), the inset describes the chemical structure of APol A8-35. Circled numbers refer to chemically equivalent ¹H whose resonances are shown in the 1D spectra displayed

in a. One-dimensional spectra corresponding to the rows labeled C–F in (b) are shown to the *right (panels c-f, respectively)*. All data were acquired at 400 MHz 1 H Larmor frequency, 25 °C, at the same concentration of polymers (30 g L^{-1}) solubilized in pure D₂O

the 1D rows extracted from the 2D NOESY spectra (Fig. 7d–f).

The 2D $\rm ^1H, ^{13}C$ HSQC spectra show that substituting HAPol or DAPol with perDAPol entails a substantial reduction in the ${}^{1}H, {}^{13}C$ correlation peaks that could mask amino acid resonances, particularly above 24 ppm in the $13C$ dimension (Fig. [8\)](#page-11-0). Note, however, that in the case of methyl-TROSY-based experiments (Plevin and Boisbouvier [2012;](#page-14-0) Sprangers, Velyvis and Kay [2007](#page-15-0)), the interference by HAPol would be severe, but that by DAPol limited (areas indicated by colored ellipses in Fig. [8](#page-11-0)).

Discussion

Revisitation of the Mass Distribution of A8-35 Molecules

The aim of the present work was the synthesis of a perdeuterated version of A8-35 that would be chemically as similar as possible to hydrogenated A8-35. This implied the prior synthesis of perdeuterated PAA with the same average size as the commercial PAA used as starting material to synthesize the hydrogenated version. Previous

Fig. 8 Two-dimensional ¹H,¹³C HSQC spectra of HAPol, DAPol and perDAPol in D_2O . *Circled numbers* have the same meaning as in Fig. [7](#page-10-0)a. The areas where most ${}^{13}CH_3$ correlation peaks lie in methyl-TROSY-based experiments are indicated by colored ellipses [adapted from ref. (Plevin and Boisbouvier [2012](#page-14-0))]

estimates of the size of commercial PAA, performed by SEC in aqueous solution using double detection, RI and differential viscometry (dV) detection and a ''universal'' PEO calibration, yielded scattered results (Gohon et al. [2004,](#page-14-0) [2006](#page-14-0); Tribet et al. [1996](#page-15-0)). Here, we have reinvestigated the average molar mass and mass distribution of PAA, deuterated or not, by a more appropriate method, namely SEC in organic solvent following permethylation, and examined the molar mass distribution of the resulting APols by ESI-IMS-MS. SEC profiles monitored by two detection modes were used to calculate the average molar masses and mass distribution of the samples: A simple detection mode (differential RI detector) or a triple-detection one (possessing as a key element a static light scattering detector). It should be noted here that the accurate and reliable mass determination of low molar mass chains species is very difficult by light scattering (weak signals with strong noise), which may lead to overestimated average values. As regards the simple RI detection, it has been shown that in THF, a polystyrene calibration is valid for poly(methyl acrylate) samples (Couvreur et al. [2003](#page-13-0)), which simplifies analyses of methylated PAA samples. SEC analyses consistently indicate that the \overline{DP}_n of commercial PAA is close to \sim 35, which, assuming $D \approx 2$ (*cf.*) Table [1](#page-4-0), PAA-1), is consistent with the \overline{DP}_w of \sim 70 estimated in ref. (Tribet et al. [1996](#page-15-0)), but lower than the \overline{DP}_n of ~ 80 (deduced from $\overline{M}_n \approx 7.5$ kDa) found, also for commercial PAA, in subsequent measurements (Gohon et al. [2004](#page-14-0), [2006\)](#page-14-0). The difference may stem from (1) the difficulty of determining accurately the molar mass of PAA by SEC using a ''universal'' calibration based on a PEO standards, due to differences in interaction parameters (Gohon et al. [2004](#page-14-0); Tribet et al. [1996](#page-15-0)) (the procedure used in these works was initially developed to characterize neutral polymers and not polyelectrolytes), (2) a slight drift of the semi-logarithmic ''universal'' calibration curve, (3) the reproducibility of the preparation of the different batches of PAA at the industrial scale and in the laboratory and/or possibly other unidentified causes. The revised value of $\overline{M_n}$ is $\sim 2 \times$ smaller than that (8–10 kDa, depending on the source of PAA) assumed previously. Note that a similar revision also applies to sulfonated amphipols (SAPols), which are synthesized starting from the same PAA (Dahmane et al. [2011\)](#page-14-0).

Revising the average molar mass of A8-35 does not affect any of the conclusions previously published regarding the mass of A8-35 particles or the mass ratio of APol to MP in complexes. Simply, an A8-35 particle formed in buffer solution, whose mass was determined to be \sim 40 kDa (Gohon et al. [2006\)](#page-14-0), is now concluded to comprise \sim 9 rather than \sim 4 molecules, as had been previously estimated. The mass of A8- 35 bound by BR, which is \sim 54 kDa (Gohon et al. [2008\)](#page-14-0), corresponds to 12–13 molecules, not \sim 6. For functionalized derivatives of A8-35 such as fluorescent (Fernandez et al. [2014;](#page-14-0) Giusti et al. [2012;](#page-14-0) Opačić et al. [2014](#page-14-0); Zoonens et al. [2007\)](#page-15-0) or tagged (Charvolin et al. [2009](#page-13-0); Giusti et al. [2014;](#page-14-0) Le Bon et al. [2014a\)](#page-14-0) APols [reviewed in ref. (Le Bon, Popot and Giusti, [2014b\)](#page-14-0)], the number of fluorophores or tags per particle or complex is not affected either. There are very few circumstances where knowledge of the actual number of APol molecules in a particle or complex is needed. This would be the case, e.g., should one want to calculate the entropy of dissociation of a particle or a MP-bound A8-35 layer into individual molecules (Giusti et al. [2012](#page-14-0); Tribet et al. [2009\)](#page-15-0), as well as for some other thermodynamic calculations (Pocanschi et al. [2013](#page-14-0)). Knowledge of the actual molar mass is also necessary should one want to make sure that most of the individual molecules are labeled or tagged, which is seldom relevant but can be of use under certain circumstances.

Synthesis and Chemical Characterization of perDAPol

The first and limiting step of the synthesis of perDAPol consisted in producing a perdeuterated PAA precursor whose molar mass be close to that of the perhydrogenated PAA. Perdeuterated PAA was synthesized by free radical telomerization of commercial perdeuterated acrylic acid according to a procedure adapted from that described in ref. (Loubat et al. [2000\)](#page-14-0). Although this method is well established and described in details, a considerable period of development was needed before obtaining the desired perdeuterated PAA, which was achieved on a multi-gram scale. Perdeuterated PAA was then easily converted to

perDAPol according to the classic procedure of PAA hydrophobization (Gohon et al. [2004,](#page-14-0) [2006\)](#page-14-0). The molar mass of the polymer could be deduced from that of the perdeuterated PAA precursor, as determined by SEC. Because of the perdeuteration, however, NMR spectra did not permit to assess whether the extent of octylamine and isopropylamine grafting was exactly that aimed for. The grafting procedure, however, is very reliable and large deviations from the nominal composition highly unlikely.

ESI-IMS-MS spectra of A8-35 and perDAPol cannot ascertain absolute masses, because ionization and detection yields depend on the mass of the polymers, but, used as a comparative method, they do show that the two polymers have the same average mass, which is close to that expected from the mass of the PAA, and very similar mass distributions.

Solution Properties of perDAPol

Above a concentration of ~ 0.002 g L⁻¹ (Giusti et al. [2012\)](#page-14-0), A8-35 self-assembles in aqueous solutions into small, well-defined globular particles, with an average mass of \sim 40 kDa, which have been extensively characterized by light scattering, SEC, AUC and small-angle neutron scattering (Gohon et al. [2004,](#page-14-0) [2006](#page-14-0)), as well as modeled by molecular dynamics (Perlmutter et al. [2011](#page-14-0)). As expected from its chemical composition, perDAPol was found to behave, according to SEC, exactly as hydrogenated A8-35. Upon SANS, however, perDAPol particles appeared smaller than previously observed for A8-35 and DAPol (Gohon et al. [2006](#page-14-0)), for reasons that have not been identified yet but are unlikely to be related to isotopic labeling and may stem from differences in buffer composition. Whatever the case may be, MP trapping with perDAPol proceeded with the same efficacy as with HAPol or DAPol and, according to SEC, yielded MP/APol complexes of the same apparent size and dispersity.

Foreseeable Applications

The development of perDAPol was primarily undertaken in view of NMR applications. In previous work, determination of the structure of BLT2-bound $LTB₄$ was possible using DAPol, because a sufficient number of protons in the ligand have chemical shifts that differ from those of the backbone protons of A8-35 (Catoire et al. [2010a\)](#page-13-0). Nevertheless, the conformation of some regions of the molecule remained less precisely defined, because NOE signals between aliphatic protons of the ligand were masked by those from the APol. This problem would have been crippling for other ligands, such as the heptadecanoid 12-HHT (12S-hydroxyheptadeca-5Z, 8E, 10E-trienoic acid) (Catoire et al. [2011;](#page-13-0) Hamberg, Svensson and Samuelsson, [1974](#page-14-0)). For such studies, perDAPol appears as a promising new tool, given the very low intensity of the signals originating from the few remaining ¹H nuclei. Indeed, the protonated backbone of DAPol gives rise to intense signals between 0.8 and 2.6 ppm (Fig. [7b](#page-10-0)), creating important distortions of the baseline of the NOESY spectrum. Baseline corrections carried out after the Fourier transform can affect the volumes of weak ligand cross-peaks that are located close to DAPol ones, which affects the accuracy of peak volume measurements. With perDAPol, no such baseline treatment will be needed.

As regards MP studies, APols signals can interfere with those from α -carbon-linked protons and CH_n moieties of amino acid side chains, as illustrated in Fig. [8.](#page-11-0) The use of perDAPol can be very advantageous, especially when protein ¹³C signals need to be observed above \sim 25 ppm (Fig. [7\)](#page-10-0). Among various approaches that emerged during the past 10 years to study large proteins or protein complexes, one of the best strategies is the use of $^{13}CH_3$ methyl groups immersed in a perdeuterated environment (Plevin and Boisbouvier [2012](#page-14-0); Sprangers et al. [2007\)](#page-15-0). Unfortunately, most of the ${}^{13}CH_3$ methyl groups in amino acids resonate between 0 and 2.5 ppm and 10 and 30 ppm in the ¹H and ¹³C dimensions, respectively (Fig. [8\)](#page-11-0), *i.e.*, in a region where many signals from the CH_n groups of the alkyl chains of surfactants are localized. In the case of methyl-TROSY-based experiments, the use of perDAPol rather than DAPol would not make much of a difference, because in this region, *i.e.*, between 0.6 and 1.4 ppm in the H dimension, the remaining ${}^{1}H$ signals are more or less the same.

Previous work has shown that the resolution of $2D⁻¹H$, ¹⁵N TROSY spectra collected on APol-trapped MPs is sufficient for to obtain insights into membrane protein structure and dynamics at least up to the size of BR (27 kDa) (Bazzacco et al. [2012;](#page-13-0) Catoire et al. [2009](#page-13-0), [2010b](#page-13-0); Dahmane et al. [2011](#page-14-0); Elter et al. [2014](#page-14-0); Etzkorn et al. [2013](#page-14-0); Planchard et al. [2014;](#page-14-0) Zoonens et al. [2005](#page-15-0)). A8-35 permits the identification of most amide protons, but not, however, of those whose exchange with the solvent is too rapid at the slightly alkaline pH (typically, pH 8) that is optimal for use of this APol. Hence, sequence-specific assignment of backbone ${}^{1}H^{N}$, ${}^{15}N$ and ${}^{13}C^{\alpha}$ chemical shifts using TRO-SY-based HNCA (Salzmann et al. [1999](#page-15-0)) and HNCOCA (Kay et al. [1990](#page-14-0)) experiments should be realized with MPs trapped in a pH-insensitive APol, such as SAPol (Dahmane et al. [2011](#page-14-0)) or non-ionic APol (NAPol) (Bazzacco et al. [2012](#page-13-0)), at least for amide protons located in solvent-exposed loops. Concerning the identification of intramolecular ¹H dipolar interactions through 3D 15 N-edited and 13 C-edited NOESY experiments, the choice of the APol to be used would again depend on how exchangeable the protons investigated are. DAPol and perDAPol would be highly recommended to observe methyl–methyl interactions, for

instance, whereas NAPol or SAPol would be needed to observe NOEs involving loop ${}^{1}H^{N}$ backbone protons, provided that the NOE cross-peaks are not hidden by NAPol or SAPol signals. This begs the question of SAPol and NAPol perdeuteration. Because their solubility is ensured by sugar residues, NAPol would be extremely costly to perdeuterate. SAPol, on the other hand, could be perdeuterated following the route described here for obtaining perDAPol. A remaining concern is that SAPol, probably because of its high charge density (\sim 75 charges per 100 U), appears less stabilizing than A8-35 or NAPol (Bazzacco et al. 2012; Dahmane et al. [2011](#page-14-0); Picard et al. [2006;](#page-14-0) see however Huynh et al. [2014\)](#page-14-0). It will probably be adequate for many MPs, but a higher stabilizing power could probably be achieved by lowering its charge density to the level of that of A8-35 (*i.e.*, \sim 35 charges per 100 U). This, however, will require the development of a different synthesis protocol. Phosphocholine-based APols (Diab et al. [2007a,](#page-14-0) [b\)](#page-14-0) could in principle be perdeuterated, but their use for NMR studies remains to be established. For a more detailed discussion of the applications of APols to NMR studies of MPs, see ref. (Planchard et al. [2014](#page-14-0)).

Deuterated APols have other uses than in NMR studies. DAPol has been used, for instance, to modulate the contrast of APol with the solvent in both AUC and SANS experiments (Gohon et al. [2004,](#page-14-0) [2006,](#page-14-0) [2008](#page-14-0); Sverzhinsky et al. [2014\)](#page-15-0). The benefit of perdeuteration for AUC experiments would be limited, because the extra contrast afforded over DAPol is small, the expected hydrostatic density of perD-APol being \sim 1.336 g L⁻¹ versus 1.307 g L⁻¹ for DAPol and 1.236 g L^{-1} for HAPol (Gohon et al. [2004\)](#page-14-0). On the contrary, the CMP of perDAPol in SANS experiments is \sim 120.5 % D₂O (present work), versus 85.1 and 23.1 % for DAPol and HAPol, respectively (Gohon et al. [2004](#page-14-0), [2006](#page-14-0)). This is potentially highly interesting, because it makes it possible, by mixing perDAPol and HAPol, to achieve a CMP of 100 $\%$ D₂O, providing ideal conditions of high contrast and low inelastic scattering noise for MP SANS studies. Equally interesting would be the use of perDAPol for INS studies of the dynamics of APol-trapped MPs. This is an important issue, because damping of MP dynamics by APols seems to be involved both in MP stabilization and in some functional effects [for discussions, see refs. (Perlmutter et al. [2014;](#page-14-0) Popot et al. [2011](#page-15-0); Zoonens and Popot [2014](#page-15-0))]. Current INS data have been obtained on DAPol particles and MP/ DAPol complexes, in which case the hydrogenated backbone of DAPol contributes to the signal (Tehei et al. [2014](#page-15-0)). The use of perDAPol would make it possible to totally eliminate the contribution of APol protons.

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