



Contrast variation of micelles composed of Ca^{2+} and block copolymers of two negatively charged polyelectrolytes

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Abstract

Block copolymers were prepared with two anionic polyelectrolyte blocks: sodium polyacrylate (PA) and sodium polystyrene sulfonate (PSS), in order to investigate their phase behavior in aqueous solution in the presence of Ca^{2+} cations. Depending on the concentration of polymer and Ca^{2+} and on the ratio of the block lengths in the copolymer, spherical micelles were observed. Micelle formation arises from the specific interaction of Ca^{2+} with the PA block only. An extensive small-angle scattering study was performed in order to unravel the structure and dimensions of the block copolymer micelles. Deuteration of the PA block enabled us to perform contrast variation experiments using small-angle neutron scattering at variable ratios of light and heavy water which were combined with information from small-angle X-ray scattering and dynamic light scattering.

Keywords Block copolyelectrolytes · Self-assembly · Specific interaction · Contrast variation · Micelles · Small-angle scattering

Introduction

Polyelectrolytes and their interactions with oppositely charged species such as ions, [1] surfactants [2–4], or charged surfaces [5] are essential to understand and control a variety of industrial processes. These include waste water treatment, [6] modification of the rheological behavior of concrete through superplasticizers [7, 8], or the properties of consumer goods such as shampoo [9]. The phase behavior of polyelectrolytes with several multivalent cations such as Ca^{2+} , [10, 11] Sr^{2+} , [11–14] Pb^{2+} , [15] Al^{3+} [16, 17], and La^{3+} [18, 19] has been extensively studied during the last decades. These aqueous systems show a rich phase behavior strongly depending on the polyelectrolyte, [20] cation, [16,

17, 21], temperature [16, 22], or concentration of inert salts such as NaCl [23].

Extensive work on the polyelectrolyte sodium polyacrylate (PA) addressed the specific interaction with Ca^{2+} [10, 11, 13, 23, 24]. Upon the addition of Ca^{2+} , the cation will bind to the negatively charged moieties of the PA chain. When the Ca^{2+} concentration is increased, more cations will bind to the polyelectrolyte until a critical concentration $[\text{Ca}^{2+}]_c$ is reached. Beyond this concentration, the polymer starts to aggregate and precipitate from the solution. A phase boundary according to

$$[\text{Ca}^{2+}]_c = r_0 + m[\text{PA}] \quad (1)$$

is found for this system where [PA] is the concentration of polyacrylate monomer units, r_0 the minimum concentration of Ca^{2+} required to precipitate the polymer at infinite dilution, and m defines the stoichiometry of binding. Precipitation at infinite dilution here means an intramolecular coil collapse. Equation 1 relates to a complexation equilibrium completely on the side of the Ca^{2+} /PA complex. Specificity of the interaction between Ca^{2+} and PA was further confirmed by isothermal titration calorimetry (ITC), which revealed a large endothermic heat. This large heat of reaction was interpreted by a liberation of around ten water molecules from hydration shells which coalesce upon complex bond formation between Ca^{2+} and two COO^- [25].

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Since one Ca^{2+} binds quantitatively to two polyacrylate monomer units, this system has for example been used for the design of an efficient Ca^{2+} sensor [26].

In contrast, another common polyelectrolyte polystyrene sulfonate (PSS) does not show specific interactions with Ca^{2+} . No matter how large the concentration of Ca^{2+} is set, no precipitation of the polymer does occur. Furthermore, light scattering demonstrated that the polymer chains do not even aggregate in the presence of Ca^{2+} . Isothermal titration calorimetry revealed just a weak endothermic signal, suggesting that the addition of Ca^{2+} induces only a replacement of two Na^+ cations condensed into the polyelectrolyte domains by one Ca^{2+} and thus confirming the lack of any binding of Ca^{2+} to the sulfonic residues of PSS [20].

In a previous work, we combined these two different interaction patterns in one polymer by synthesizing a block copolymer (PA-*b*-PSS) of sodium polyacrylate and sodium polystyrene sulfonate [27]. Whereas the anionic residues of the PA block interact specifically with Ca^{2+} , no such interactions are expected for the PSS block. Indeed, it turned out that the block copolyelectrolytes form micelles in aqueous solution. Application of small-angle neutron scattering (SANS) at variable scattering contrasts, revealed the morphology of these micelles: The PA blocks change their solution behavior in water as specific interactions trigger decoration of PA with Ca^{2+} cations thereby establishing the core of the micelles. The PSS blocks keep their charges along the polymer backbone due to the lack of any specific interactions and form the stabilizing corona of the micelles. The entropic nature of the specific interactions between Ca^{2+} and PA enabled us to reversibly form and disintegrate the micelles by oscillating the temperature with an increase of temperature inducing formation and a decrease disintegration.

The present work extends the achievements published in Ref. [27] with the following new results: It interprets SANS and SAXS curves with a model [28] giving access to the morphology of the micellar corona, it outlines the benefit of combining SANS at variable contrast with SAXS, and it makes full use of Stuhrmann's approach [29, 30] in order to unravel the species in the core as well as the core dimensions independently of any model. All these features are established with three different block copolyelectrolytes: one sample with similar block lengths and the PA block fully deuterated and two samples with short PSS blocks and long PA blocks whereby in one case the PA block is fully deuterated and in the other case it is hydrogenated. This for the first time makes accessible a detailed comparative discussion.

The work is grouped in four parts: in the first part, we shortly revise the results on PA-*b*-PSS published in our preceding work, [27] in order to help the reader understanding the basis of the present work and to clearly separate it from

the previous one. The second part deals with the comparative analysis of a contrast variation SANS experiment of three different block copolymers using a form factor model for micelles based on block copolymers [28]. The result is a comprehensive structural picture of the micelles. In particular, the advantage of joint SANS and SAXS experiments on the same set of samples is demonstrated and used to unravel an isotope effect occurring for two samples. The isotope effect results in a change of aggregation numbers, which was taken into account in the data analysis. The third part investigates the origin of this isotope effect using isothermal titration calorimetry and relates it to the binding of Ca^{2+} to PA. The fourth part demonstrates a model-independent analysis, which yields quantitative structural information. The obtained information is verified against the results from the form factor fits and can be used for systems where mathematical expressions for a form factor model are not available. It is demonstrated that this approach even works on a fully hydrogenated block copolymer, which has important implications for samples where deuteration is impractical or not feasible.

Experimental details

Materials

Light water (H_2O) was purified using a Milli-Q-system (Millipore) to a final resistivity of 18.2 $\text{M}\Omega$ cm. Heavy water (D_2O , Euriso-top, France, 99.90 atom% deuterium) was filtered with 100-nm PVDF filters (Merck Millex SLVV033RS) prior to use. $\text{CaCl}_2 \cdot 2\text{H}_2\text{O}$ (Sigma Aldrich, France, >99.9%), NaCl (Sigma Aldrich, France, >99.9%), and NaOH (Sigma Aldrich, France) were used as received.

Polymer synthesis and characterization

The synthesis and characterization of the polymers were described in detail in a previous work [27]. Figure 1 shows the chemical structure of the used polymers. All

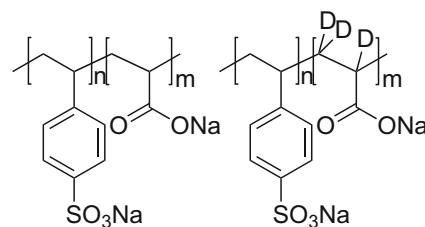


Fig. 1 Chemical structure of the used polymers. The left structure shows the fully hydrogenated polymer (h_3 -PA-*b*-PSS). The right structure shows a polymer with a PA block which is fully deuterated (d_3 -PA-*b*-PSS)

polymers were used as sodium salt. Prior to the small-angle scattering experiments the polymers and their phase behavior with Ca^{2+} were characterized using NMR and static and dynamic light scattering respectively. Detailed information can be found in the previous work [27]. Table 1 summarizes the results of the polymer characterization.

Sample preparation

The sample preparation was performed similarly as in previous works [22–24, 27]. The total number of positive charges for all samples was adjusted to 100 mmol L^{-1} . The total concentration of positive charges $[+]$ is

$$[+] = [\text{NaCl}] + 2 \cdot [\text{CaCl}_2] = 100 \text{ mmol L}^{-1} \quad (2)$$

with $[\text{NaCl}]$ being the concentration of sodium chloride and $[\text{CaCl}_2]$ the concentration of calcium chloride. A sample containing $25 \text{ mmol L}^{-1} \text{ CaCl}_2$ also contained $50 \text{ mmol L}^{-1} \text{ NaCl}$, whereas a sample with $50 \text{ mmol L}^{-1} \text{ CaCl}_2$ contains no additional NaCl.

SANS

SANS measurements were performed at the D11 small-angle scattering instrument of the Institut Laue-Langevin (Grenoble, France). Three sample to detector distances (39.0 m collimation 40.5 m, 8.0 m collimation 8.0 m, 1.4 m collimation 5.5 m) and a neutron wavelength of 0.5 nm (FWHM 9 %) were used to cover a q -range of $2 \cdot 10^{-2} \text{ nm}^{-1} - 5 \text{ nm}^{-1}$. We used a circular neutron beam with a diameter of 15 mm. Scattered neutrons were detected with a ^3He MWPC detector (CERCA) with 256×256 pixels of 3.75×3.75 -mm pixel size. Samples were filled in 2-mm Hellma 404 Quartz Suprasil cells. The sample temperature was adjusted to $25 \text{ }^\circ\text{C}$ using a circulating water bath. The detector images were azimuthally averaged, corrected to transmission of the direct beam, and scaled to absolute intensity using the LAMP software. Scaling to absolute intensity was based on a 1-mm H_2O cell as

Table 1 Overview of polymers, compositions, weight-averaged molecular weights M_w , and hydrodynamic radii R_h determined from NMR and static and dynamic light scattering in $100 \text{ mmol L}^{-1} \text{ NaCl}$ as a good solvent and in the absence of CaCl_2

Sample	PA:PSS / mol%	M_w / kg mol^{-1}	R_h / nm
h ₃ -PA ₁₁₉₀ PSS ₇₀	94:6	160 ± 2	13.8 ± 0.6
d ₃ -PA ₁₁₉₀ PSS ₇₀	94:6	181 ± 3	13.6 ± 0.5
d ₃ -PA ₃₆₀ PSS ₄₀₀	47:53	127 ± 1	10.4 ± 0.2

The subscript denotes the degree of polymerization, while d₃ and h₃ correspond to the deuteration or hydrogenation of the PA block, respectively

secondary calibration standard ($\frac{d\Sigma}{d\Omega} = 0.929 \text{ cm}^{-1}$ at a neutron wavelength of 5 Å). The scattering from the solvent and the incoherent background were subtracted from the scattering curves. Details for the data reduction can be found in Chapter 2 of Ref. [31].

SAXS

Small-angle X-ray scattering was performed at the ID02 beamline of the European Synchrotron Radiation Facility (ESRF). Two sample to detector distances (10 m and 1 m) were measured at an X-ray energy of 12.46 keV (0.0995 nm) using a Rayonix MX-170HS CCD detector to cover a q -range of $8 \cdot 10^{-3} \text{ nm}^{-1} - 6 \text{ nm}^{-1}$. Samples were filled in 2-mm quartz glass capillaries (WJM Glas Müller, Berlin, Germany). The detector images were corrected for dark and flat-field, azimuthally averaged, corrected to transmission of the direct beam, and scaled to absolute intensity using water as a secondary standard [32, 33]. The scattering from the solvent was subtracted from the scattering curves. Details can be found in Ref. [33]. Error bars were estimated as standard deviations from measurements of at least five different positions within the capillary.

Small-angle scattering data analysis

The form factor of spherical self-assembled block copolymers with excluded volume interaction of the polymer chains in the corona [28] was used to fit the SAXS and SANS data. The specific macroscopic scattering cross-section per unit solid angle $\frac{d\Sigma}{d\Omega}(q)$ of a solution of micelles can be written as [28]

$$\begin{aligned} \frac{d\Sigma}{d\Omega}(q) = N \left[N_{\text{agg}}^2 \beta_{\text{core}}^2 A_{\text{core}}^2(q) + N_{\text{agg}} \beta_{\text{corona}}^2 P'_{\text{corona}}(q) \right. \\ \left. + 2N_{\text{agg}}^2 \beta_{\text{core}} \beta_{\text{corona}} A_{\text{core}}(q) A_{\text{corona}}(q) \right. \\ \left. + N_{\text{agg}} (N_{\text{agg}} - P'_{\text{corona}}(0)) \beta_{\text{corona}}^2 A_{\text{corona}}^2(q) \right] \quad (3) \end{aligned}$$

where N is the number density of micelles, N_{agg} the aggregation number of micelles, and β_{core} and β_{corona} are the total excess scattering length of the block forming the spherical core and the corona, respectively. They are defined as

$$\beta_{\text{corona}} = V_{\text{m, corona}} \text{DP}_{\text{corona}} \Delta\rho_{\text{corona}} \quad (4)$$

and

$$\beta_{\text{core}} = V_{\text{m, core}} \text{DP}_{\text{core}} \Delta\rho_{\text{core}} \quad (5)$$

with V_{m} being the molecular volume of the respective monomer unit, $\text{DP}_{\text{corona}}$ and DP_{core} the degree of polymerization of the corona and core block, and $\Delta\rho$ the corresponding excess scattering length density.

Equation 3 consists of four different contributions: scattering from the spherical homogeneous core $A_{\text{core}}^2(q)$, scattering from the polymer chains in the corona $P'_{\text{corona}}(q)$, the term which takes into account the cross-term between core and corona $A_{\text{core}}(q) \cdot A_{\text{corona}}(q)$, and a term which takes into account interferences between chains in the corona $A_{\text{corona}}^2(q)$. $A_{\text{core}}(q)$ is the scattering amplitude of a homogeneous sphere [34] with radius R_{core}

$$A_{\text{core}}(q) = 3 \frac{\sin(qR_{\text{core}}) - qR_{\text{core}} \cos(qR_{\text{core}})}{(qR_{\text{core}})^3} \quad (6)$$

$P'_{\text{corona}}(q)$ is the form factor of a chain in the corona. It contains the self-correlation of the chain $P_{\text{exv}}(q)$ as well as the interaction between the chains, which is expressed by the interaction parameter ν [28, 35]

$$P'_{\text{corona}}(q) = \frac{P_{\text{exv}}(q)}{1 + \nu P_{\text{exv}}(q)} \quad (7)$$

where $P_{\text{exv}}(q)$ is the form factor of a semi flexible self-avoiding chain. This form factor was first derived by Pedersen and Schurtenberger [36] and later corrected [37]. In experiments, ν typically adopts values between 0 and 8 and is related to the osmotic compressibility κ by [28, 35, 38]

$$\kappa = 1 + \nu \quad (8)$$

$A_{\text{corona}}(q)$ is given by

$$A_{\text{corona}}(q) = \frac{\int \rho_{\text{corona}}(r) r^2 \frac{\sin(qr)}{qr} dr}{\int \rho_{\text{corona}}(r) r^2 dr} \quad (9)$$

with $\rho_{\text{corona}}(r)$ as the scattering length density profile in the corona. $A_{\text{corona}}(q)$ is normalized so that $A_{\text{corona}}(0) = 1$. In this work, we use a Gaussian profile, which is defined as

$$\rho_{\text{corona}}(r) = \begin{cases} 0 & \text{for } r < R_{\text{core}} \\ 1 & \text{for } r = R_{\text{core}} \\ \exp\left(\frac{-(r-R_{\text{core}})^2}{2s^2}\right) & \text{for } r > R_{\text{core}} \end{cases} \quad (10)$$

with s controlling the thickness of the corona.

In order to take into account the size distribution of micelles, we assumed a log-normal distribution of the aggregation number N_{agg}

$$p(N_{\text{agg}}) = \frac{1}{H\sqrt{2\pi}N_{\text{agg}}} \exp\left(\frac{-\log(N_{\text{agg}} - M)^2}{2H^2}\right) \quad (11)$$

where H and M define the distribution and are connected to the mean aggregation number $\overline{N_{\text{agg}}}$ and standard deviation $\sigma_{\overline{N_{\text{agg}}}}$ by

$$\overline{N_{\text{agg}}} = \exp\left(M + \frac{H^2}{2}\right) \quad (12)$$

$$\sigma_{\overline{N_{\text{agg}}}} = \sqrt{\exp(H^2 + 2M)(\exp(H^2) - 1)} \quad (13)$$

The macroscopic scattering cross-section is therefore

$$\frac{d\Sigma}{d\Omega}_{\text{polydisperse}}(q) = \int \frac{d\Sigma}{d\Omega}(q) p(N_{\text{agg}}) dN_{\text{agg}} \quad (14)$$

Instrumental resolution for SANS has been taken into account according to Ref. [39]. The macroscopic scattering function is convoluted with a resolution function $R(q, \sigma_q)$, which depends on wavelength spread, finite collimation of the beam, and detector resolution

$$\frac{d\Sigma}{d\Omega}_{\text{smearred}}(q) = \int R(q, \sigma_q) \frac{d\Sigma}{d\Omega}_{\text{polydisperse}}(q) dq \quad (15)$$

For the form factor fits, we used the SASET program [40], which allows global fitting of several contrasts at the same time. During the data analysis of the SANS curves, we took into account the instrumental resolution for each detector configuration and merged the data only for final representation. This approach allowed us to increase the number of available data points since we do not truncate the data in the region of overlapping q . The factor for merging the SANS measurements at 39 m to the other configurations was set as a global fitting parameter for all the curves as using water as an absolute calibration standard is not feasible at this sample to detector distance.

Dynamic light scattering

Dynamic light scattering (DLS) available in the ILL PSCM lab was performed using an ALV CGS-3 (ALV, Langen, Germany) equipped with a HeNe laser operating at $\lambda_0 = 632.8$ nm. The samples were filtered in cylindrical quartz glass cells using Millex PVDF filters with a pore size of $0.45 \mu\text{m}$. The temperature of the samples was kept at 25°C . DLS was measured in an angular range from 30° to 150° in steps of 10° .

The intensity-time correlation function $g_2(\tau) - 1$ was analyzed using the method of cumulants [41]

$$g_2(\tau) - 1 = B + \beta \exp(-2\Gamma\tau) \left(1 + \frac{\mu_2}{2!} \tau^2\right)^2 \quad (16)$$

B is a factor correcting the baseline, β a factor which depends on the experimental setup, and Γ the relaxation rate and μ_2 the second cumulant. The apparent diffusion coefficient $D_{\text{app}}(c, q)$ for a given q is calculated according to

$$D_{\text{app}} = \frac{\Gamma}{q^2} \quad (17)$$

The diffusion coefficient is extrapolated towards $q = 0$ according to [42, 43]

$$D_{\text{app}}(q) = D_0 \left(1 + CR_g^2 q^2\right) \quad (18)$$

where C is a constant describing the q dependence of $D_{\text{app}}(q)$. The diffusion coefficient D_0 is used to calculate

the hydrodynamic radius R_h using the Stokes-Einstein equation

$$R_h = \frac{k_B T}{6\pi\eta D_0} \quad (19)$$

where T is the temperature, k_B the Boltzmann constant, and η the viscosity of the solvent.

The viscosity of D₂O and H₂O was obtained from Ref. [44], while for mixtures of D₂O and H₂O the viscosity was calculated based on the volume fraction.

Isothermal titration calorimetry (ITC)

Isothermal titration calorimetry (ITC) was performed using a VP-ITC calorimeter from Malvern Instruments at 25.0 °C. The cell contained 1.4 mL of a solution of PA (0.2 gL⁻¹ in 100 mmol L⁻¹ NaCl). To this solution, between 4 and 24 μL of an aqueous solution containing 12 mmol L⁻¹ CaCl₂ and 76 mmol L⁻¹ NaCl was added. The time span between the injections was 300 s. The experiment was repeated twice, first using H₂O as a solvent and then D₂O. The baseline correction and integration of the peaks was performed using the software NITPIC [45]. The signals were not corrected for the heat of dilution of the polyelectrolyte and the CaCl₂ solution as these contributions were considerably smaller than the heat of binding of Ca²⁺ to PA. The data were fitted using the model of a single set of identical binding sites [46, 47].

Results and discussion

Review of micellization of PA-*b*-PSS in the presence of Ca²⁺ [27]

Four polymers at three different ratios of block lengths were synthesized and investigated in the context of the previous work: two polymers with a long PA and a short PSS block (d₃-PA₁₁₉₀PSS₇₀ and h₃-PA₁₁₉₀PSS₇₀), one polymer with a symmetric block ratio between PA and PSS (d₃-PA₃₆₀PSS₄₀₀), and one polymer with a long PSS and a short PA block (h₃-PA₁₀₀PSS₇₅₀). In the presence of Ca²⁺, d₃-PA₁₁₉₀PSS₇₀ and d₃-PA₃₆₀PSS₄₀₀ form spherical block copolymer micelles arising from the specific interaction of Ca²⁺ with the PA block. In contrast, micelle formation was not observed for h₃-PA₁₀₀PSS₇₅₀ due to the considerably shorter PA block, not rendering the polymer chain hydrophobic enough to form micellar aggregates. Phase diagrams of the two polymers forming micelles were established and are presented in Fig. 2.

Contrast variation small-angle neutron scattering of the micelles was performed at variable H₂O/D₂O ratios using the two block copolymers with the deuterated PA

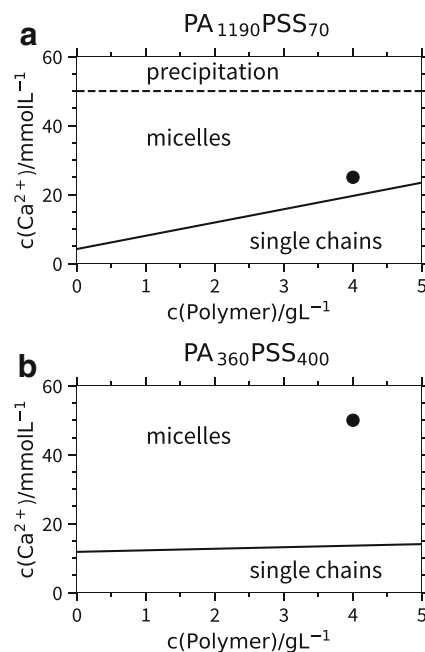


Fig. 2 Phase diagrams of **a** d₃-PA₁₁₉₀PSS₇₀/h₃-PA₁₁₉₀PSS₇₀ and **b** d₃-PA₃₆₀PSS₄₀₀ in the presence of Ca²⁺ at 25 °C. The black points indicate the states where the contrast variation experiments were performed

block, labelled as d₃-PA. The black dots in Fig. 2 indicate the sample composition where the contrast variation experiments were conducted. A preliminary analysis of the scattering data revealed that d₃-PA blocks complexed by Ca²⁺ form the hydrophobic core of the micelles, while PSS is located in the corona and stabilizes the aggregates.

Furthermore, it was shown that an increase of temperature promotes the binding of Ca²⁺ to PA and was therefore used to induce micelle formation of a sample close to the phase transition. It was demonstrated that micelle formation is reversible and can be induced and reverted by consecutive additions of Ca²⁺ and a complexing agent such as Na₂EDTA and by consecutive cycles of temperature increase and drop respectively.

Scattering analysis with a form factor model

The main motivation for performing a contrast variation experiment on the presented system is to figure out whether PSS or PA forms the hydrophobic core of the anticipated micellar aggregates. This was already elucidated in the previous work [27]; however, fitting the data with a form factor model also makes accessible information about the exact size of the micelle core, the water content inside the micelle core, and the thickness of the corona as well as the interaction of the polymer chains in the corona. In order to

assess such information, the model of a block copolymer micelle with self-avoiding chains in the corona (Eq. 3) was used to fit the experimental data [28]. In addition to the polymers d_3 -PA₃₆₀PSS₄₀₀ and d_3 -PA₁₁₉₀PSS₇₀, a third fully hydrogenated polymer h_3 -PA₁₁₉₀PSS₇₀ is investigated in the current work. Deuteration is not always feasible as deuterated monomers do not exist for a wide range of polymers and deuteration of biological molecules such as proteins or sugars is often difficult and very cost intensive. Nevertheless, a contrast variation experiment is still possible if the scattering length densities of two components are sufficiently different. h_3 -PA₁₁₉₀PSS₇₀ will be used as a showcase to illustrate the feasibility of performing a contrast variation study on a fully hydrogenated block copolymer. The present section will present a comparative and exhaustive interpretation of the SANS and SAXS curves of all three block copolyelectrolyte samples with suitable model form factors [28].

Figure 3 shows the SANS profiles of d_3 -PA₃₆₀PSS₄₀₀ micelles at variable H₂O/D₂O ratios. As has been outlined in Ref. [27], the overall shape and the forward scattering of the SANS profiles presented in Fig. 3 drastically change upon variation of the solvent ratio H₂O/D₂O. SAXS on the same samples was also performed to show that changing from light to heavy water does not influence the aggregation number, size, and morphology of the aggregates. Figure 4 a shows these SAXS profiles. All SAXS curves overlap perfectly (Fig. 4b), when plotted without the multiplication factor used in Fig. 4a. The perfect overlay of the SAXS curves clearly demonstrates that the variation in the D₂O/H₂O ratio does not affect the nature of the aggregates.

This finding simplifies the interpretation of the scattering data from d_3 -PA₃₆₀PSS₄₀₀ considerably.

Figures 5 and 7 show the SANS profiles of d_3 -PA₁₁₉₀PSS₇₀ and h_3 -PA₁₁₉₀PSS₇₀ micelles in six different H₂O/D₂O mixtures, and Figs. 6a and 8a show the SAXS profiles of the corresponding solutions. As observed for d_3 -PA₃₆₀PSS₄₀₀, the forward scattering of the SANS curves changes with varying D₂O content. For d_3 -PA₁₁₉₀PSS₇₀, the forward scattering continuously decreases with increasing content of D₂O, while for h_3 -PA₁₁₉₀PSS₇₀ the forward scattering decreases until 54.0% D₂O and subsequently increases again.

Moreover, the slope at high q changes. This becomes particularly evident for the partly deuterated polymer when comparing Fig. 5c and f. In Fig. 5c, the PSS block is matched and a slope of -4 is found. The contrast in Fig. 5f is the closest possible one to the match of d_3 -PA and shows a less steep slope, which is closer to -2 . The data at high q for the fully hydrogenated polymer show a higher statistical error than the data from the two other polymers. This can be attributed to the higher ¹H content of the fully hydrogenated polymer. Hence, the subtraction of the higher incoherent background introduces more noise to the data. This makes the determination of slopes at high q for this polymer difficult.

The SAXS profiles of d_3 -PA₁₁₉₀PSS₇₀ and h_3 -PA₁₁₉₀PSS₇₀ in Figs. 6a and 8a show well-defined form factor oscillations; however, a closer look at the position of the first minimum of $P(q)$ shows that q_{\min} shifts to slightly higher values when the D₂O content is increased. This corresponds to a slightly changing micelle size when

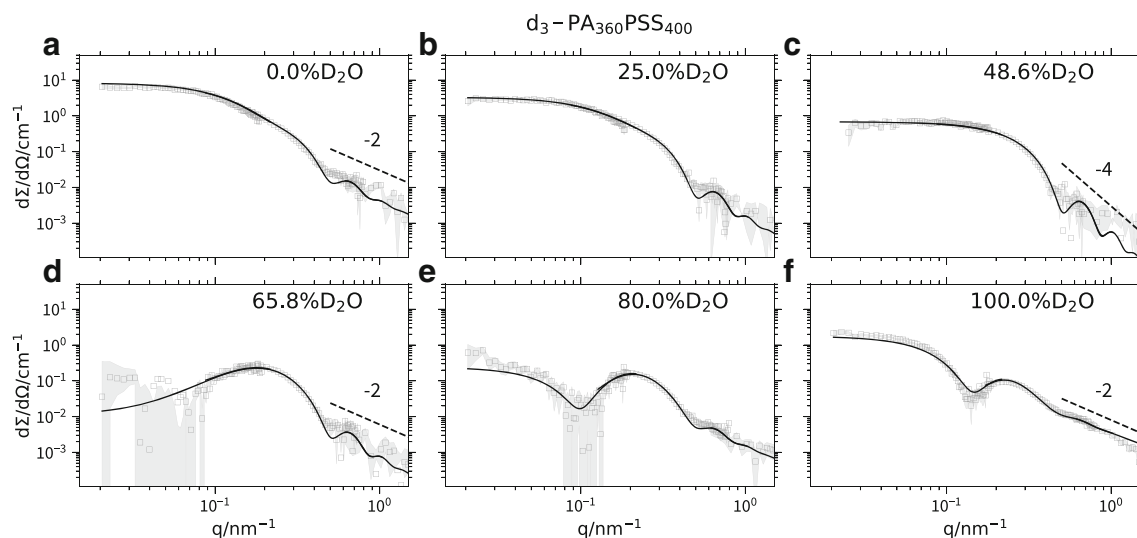


Fig. 3 SANS profiles of d_3 -PA₃₆₀PSS₄₀₀ ($c_{\text{poly}} = 4\text{gL}^{-1}$, $c_{\text{Ca}^{2+}} = 50\text{mmolL}^{-1}$) in 0.0% D₂O (a), 25.0% D₂O (b), 48.6% D₂O (c), 65.8% D₂O (d), 80.0% D₂O (e), and 100.0% D₂O (f). The solid lines

represent fits to the form factor of a polydisperse block copolymer micelle [28]. An overview of the fit results can be found in Table 2 and Table S3 of the Supporting Information

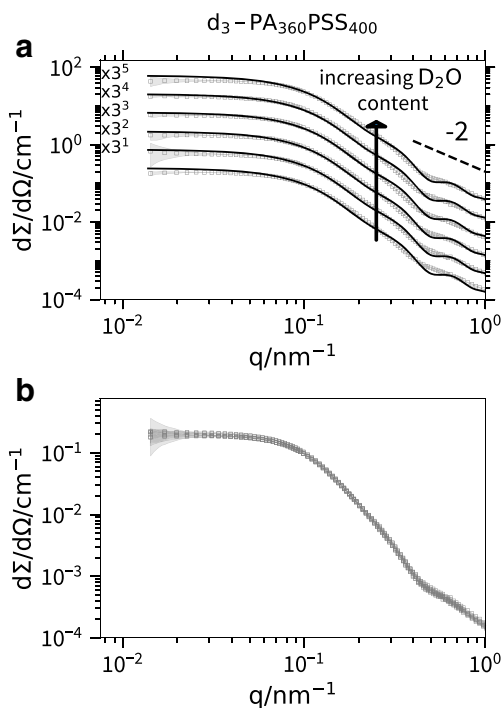


Fig. 4 SAXS profiles of d_3 -PA₃₆₀PSS₄₀₀ ($c_{\text{poly}} = 4\text{gL}^{-1}$, $c_{\text{Ca}^{2+}} = 50\text{ mmol L}^{-1}$) in 0.0% D₂O, 25.0% D₂O, 48.6% D₂O, 65.8% D₂O, 80.0% D₂O, and 100.0% D₂O. **a** The curves multiplied with an offset. **b** The curves without the multiplication factor. The solid lines represent fits to the form factor of a polydisperse block copolymer micelle [28]. An overview of the fit results can be found in Table 2 and Table S3 of the Supporting Information

changing the D₂O/H₂O ratio. As this feature is very hard to be identified with SANS curves only, it highlights the relevance of a combined SANS and SAXS experiment for all samples of a contrast variation study.

The scattering curves of all three samples were analyzed with two possible configurations: Ca_{0.5}-d₃-PA forming the core and PSS the corona as well as Ca_{0.5}-PSS forming the core and PA the corona. SANS profiles at different contrasts as well as the SAXS profiles of the same samples were analyzed simultaneously in a global fit. Furthermore, the fit was heavily constrained due to the knowledge of the polymer composition, polymer concentration, molecular volumes [48], and scattering length densities of the blocks and the solvent.

A structure factor was not considered during the data analysis since the data did not show any evidence for the presence of particle interaction such as an up- or downturn at low q . This is rather unusual for highly charged systems since these systems usually show long range interactions or aggregation [49].

However, for the present case, the absence of a structure factor contribution can be explained by the high ionic strength of the respective polymer solution, which was introduced by adding a certain amount of NaCl as “inert salt” (see Sample preparation for details). This addition of NaCl leads to a strong screening of the charges of the polyelectrolyte. We estimated the ionic strength and Debye length of the solution by assuming that every PA monomer unit binds 0.5 Ca²⁺ cations. For d_3 -PA₃₆₀PSS₄₀₀, we obtain an ionic strength of 141 mmol L⁻¹ and a Debye length of 0.81 nm and for d_3 -PA₁₁₉₀PSS₇₀/h₃-PA₁₁₉₀PSS₇₀ an ionic strength of 98 mmol L⁻¹ and a Debye length of 0.97 nm. Previous works [12–16] showed that this approach can be used to investigate the single chain behavior of homopolyelectrolytes as the added “inert salt” effectively screens interactions between the chains. In the case of the present work, this approach was used analogously and explains the lack of any significant impact of inter-micelle repulsions on the present scattering data.

The strong screening also results in a conformation of the PSS chains in the corona, which can be well described by a self-avoiding polymer. In previous works, the use of “inert salt” such as NaCl allowed to study the single chain behavior of homopolyelectrolytes such as PA and PSS [16, 23].

The best fit obtained from analyzing the data of d_3 -PA₃₆₀PSS₄₀₀ with Ca_{0.5}-PSS forming the core and PA the corona is shown in Figure S4 of the Supporting Information. It poorly describes the features of the scattering curves. In contrast, the solid lines in Figs. 3 and 4a show the fits for d_3 -PA₃₆₀PSS₄₀₀ with Ca_{0.5}-d₃PA forming the core. The main features of the experimental data are considerably better described by the model with a core consisting of d₃-PA [27, 28]. This nicely agrees with the conclusions drawn from our previous work [27], confirming once again the fact that d₃-PA/Ca²⁺ forms the core of the micellar aggregates, while PSS is located in the corona. The form factor model with PA/Ca²⁺ in the core was also used for the analysis of d_3 -PA₁₁₉₀PSS₇₀ and h₃-PA₁₁₉₀PSS₇₀. The solid lines in Figs. 5, 6a, 7, and 8a show the form factor fits to the SANS and SAXS profiles. The model describes the data well for all contrasts. Unlike to the form factor analysis of the data from d_3 -PA₃₆₀PSS₄₀₀, where the aggregation number turned out to be independent of the ratio H₂O/D₂O, the slightly changing micelle size with increasing D₂O content observed in the case of the two samples with the long PA block is taken into account by assuming individual aggregation numbers for each contrast. Therefore, each SANS and SAXS profile for a given contrast is attributed to an individual N_{agg} . All the other

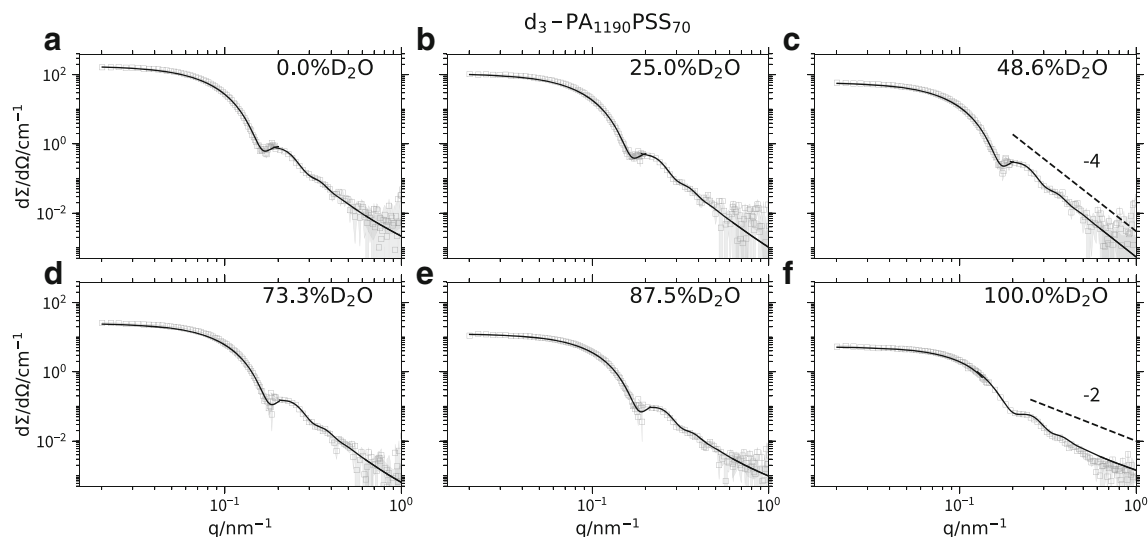


Fig. 5 SANS profiles of d_3 -PA₁₁₉₀PSS₇₀ ($c_{\text{poly}} = 4\text{gL}^{-1}$, $c_{\text{Ca}^{2+}} = 25\text{ mmol L}^{-1}$) in 0.0% D₂O (a), 25.0% D₂O (b), 48.6% D₂O (c), 73.3% D₂O (d), 87.5% D₂O (e), and 100.0% D₂O (f). The solid lines

represent fits to the form factor of a polydisperse block copolymer micelle [28]. An overview of the fit results can be found in Table 2 and Table S3 of the Supporting Information

fit parameters ($\frac{\sigma N_{\text{agg}}}{N_{\text{agg}}}$, $R_{g,\text{corona}}$, s , ν , h , $V_{m,\text{Ca}^{2+}}$) are fitted simultaneously for the whole data set. Table 2 summarizes the resulting fit parameters. The given aggregation numbers and core sizes for d_3 -PA₁₁₉₀PSS₇₀ and h_3 -PA₁₁₉₀PSS₇₀

are shown representatively for the sample in 100 % D₂O. Table S3 in the Supporting Information summarizes the aggregation numbers N_{agg} for the other contrasts.

d_3 -PA₃₆₀PSS₄₀₀ micelles have an aggregation number of 19 polymers per micelle. Compared to this the aggregation numbers of $168 \geq N_{\text{agg}} \geq 126$ for d_3 -PA₁₁₉₀PSS₇₀ and $145 \geq N_{\text{agg}} \geq 117$ for h_3 -PA₁₁₉₀PSS₇₀ are considerably bigger. This increase in aggregation number can be nicely reconciled with the increase of the PA block length as the block responsible for the formation of the poorly soluble core.

During the data analysis, it turned out that it is necessary to take into account a considerable amount of water in the core. Assuming a dry core consisting of polymer does not fit the data on absolute scale since aggregation number N_{agg} and number density of micelles N are strictly linked to the known polymer concentration. This heavily constrains the fit and shows the importance of calibration to absolute intensity. Therefore, we allowed the core to be swollen by solvent and defined the scattering length density ρ_{core} and the molecular volume of the core as

$$\rho_{\text{core}} = \frac{b_{\text{core}}}{V_{m,\text{core}}} = \frac{b_{\text{PA}^-} + hb_{\text{solvent}} + 0.5b_{\text{Ca}^{2+}}}{V_{m,\text{PA}^-} + hV_{m,\text{solvent}} + 0.5V_{m,\text{Ca}^{2+}}} \quad (20)$$

with b_{core} being the scattering length and $V_{m,\text{core}}$ the molecular volume of a monomer in the core. h is the number of solvent molecules per monomer unit. We assumed that every Ca^{2+} binds to two PA^- monomer units. Previous works using isothermal titration calorimetry and calcium selective electrodes [25] suggest that Ca^{2+} binds quantitatively to PA.

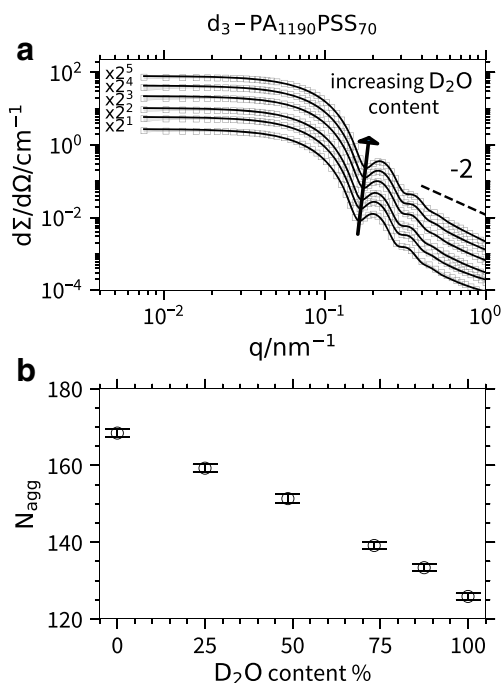


Fig. 6 a SAXS profiles of d_3 -PA₁₁₉₀PSS₇₀ micelles ($c_{\text{poly}} = 4\text{gL}^{-1}$, $c_{\text{Ca}^{2+}} = 25\text{ mmol L}^{-1}$) in 0.0%, 25.0%, 48.6%, 73.3%, 87.5%, and 100.0% D₂O. The solid lines represent fits to the form factor of a polydisperse block copolymer micelle [28]. An overview of the fit results can be found in Table 2 and Table S3 of the Supporting Information. b Aggregation number of the micelles as a function of D₂O content

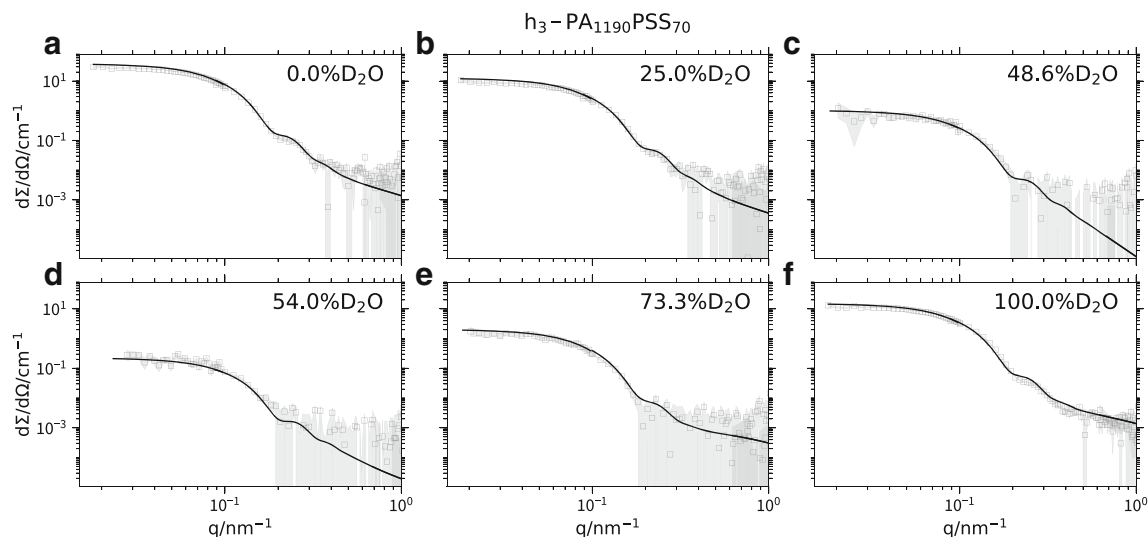


Fig. 7 SANS profiles of h_3 -PA₁₁₉₀PSS₇₀ micelles ($c_{\text{poly}} = 4\text{gL}^{-1}$, $c_{\text{Ca}^{2+}} = 25\text{mmol L}^{-1}$) in 0.0% D₂O (a), 25.0% D₂O (b), 48.6% D₂O (c), 54.0% D₂O (d), 73.3% D₂O (e), and 100.0% D₂O (f). The solid

lines represent fits to the form factor of a polydisperse block copolymer micelle [28]. An overview of the fit results can be found in Table 2 and Table S3 of the Supporting Information

The volume fraction of water in the core can be calculated by

$$f_{\text{solvent}} = 1 - \frac{(V_{m, \text{PA}^-} + 0.5V_{m, \text{Ca}^{2+}}) \text{DP}_{\text{core}} N_{\text{agg}}}{\frac{4}{3}\pi R_{\text{core}}^3} \quad (21)$$

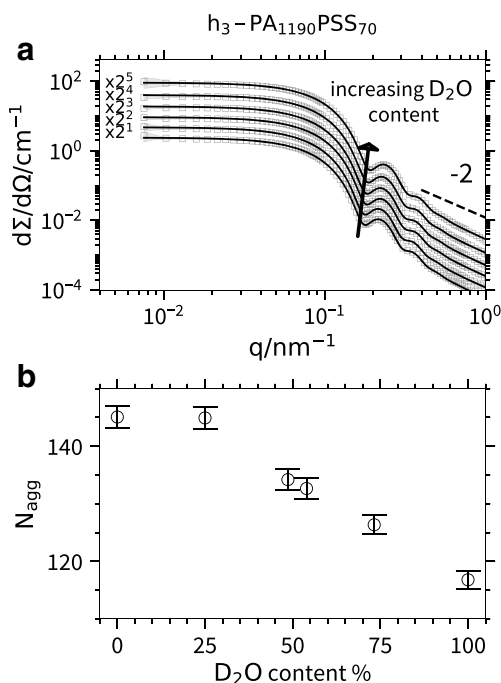


Fig. 8 a SAXS profiles of h_3 -PA₁₁₉₀PSS₇₀ micelles ($c_{\text{poly}} = 4\text{gL}^{-1}$, $c_{\text{Ca}^{2+}} = 25\text{mmol L}^{-1}$) in 0.0%, 25.0%, 48.6%, 54.0%, 73.3%, and 100.0% D₂O. The solid lines represent fits to the form factor of a polydisperse block copolymer micelle [28]. An overview of the fit results can be found in Table 2 and Table S3 of the Supporting Information. b Aggregation number of the micelles as a function of D₂O content

where DP_{core} is the degree of polymerization of the core block.

The molar volume of condensed Ca^{2+} $V_{m, \text{Ca}^{2+}}$ was left free as a fit parameter since the value based on the ionic radius ($2.5\text{cm}^3\text{mol}^{-1}$) [50] and the solvated ion ($-28.9\text{cm}^3\text{mol}^{-1}$) [50] did not give satisfactory results. From the fits of the three block copolymer micelles, an averaged value of $V_{m, \text{Ca}^{2+}} = 17.0 \pm 2.8\text{cm}^3\text{mol}^{-1}$ was obtained. Unlike the negative molar volume of the solvated ion, the value we found is positive, which is attributed to the release of hydrated water molecules upon complexation with PA.

From the fits, a quite high volume fraction of 83–85% water in the core of the micelles is obtained. This corresponds to roughly 12 water molecules per monomer unit. In order to achieve the same core volume without solvent in the core, an aggregation number of 874 would be necessary for d_3 -PA₃₆₀PSS₄₀₀, which can be safely excluded from fitting the data on absolute scale and given polymer concentration. Similar water contents have been observed for the core of Pluronic P123 micelles [51].

The core size results from the aggregation number, degree of polymerization of the PA block, molar volumes of PA, Ca^{2+} , and the water molecules in the core. For d_3 -PA₃₆₀PSS₄₀₀, a rather small core radius of $R_{\text{core}} = 8.8\text{nm}$ together with a well-defined standard deviation of 6.9% for the log-normal distribution was found. In contrast, for d_3 -PA₁₁₉₀PSS₇₀ and h_3 -PA₁₁₉₀PSS₇₀, a considerably larger micelle core of 23.7 nm and 22.3 nm was found. This arises from the longer PA block and the higher aggregation numbers of those micelles compared to the polymer with equal ratios of d_3 -PA and PSS. The distribution in aggregation numbers and consequently the distribution of the core radius for d_3 -PA₁₁₉₀PSS₇₀ and h_3 -PA₁₁₉₀PSS₇₀ is

Table 2 Structural parameters resulting from the analysis of the SANS and SAXS profiles with the form factor of a polydisperse block copolymer micelle [28]. An extended version of the table including error bars can be found in the Supporting Information

	d ₃ -PA ₃₆₀ PSS ₄₀₀	d ₃ -PA ₁₁₉₀ PSS ₇₀	h ₃ -PA ₁₁₉₀ PSS ₇₀
N_{agg}	19.1	125.9 ^a	116.8 ^a
$R_{\text{core}}/\text{nm}$	8.8	23.7 ^a	22.3 ^a
$\frac{\sigma_{R_{\text{core}}}}{R_{\text{core}}}$	0.069	0.102	0.105
$R_{g,\text{corona}}/\text{nm}$	8.55	2.12	2.07
s/nm	8.9	4.05	3.53
ν	1.19	0.0	0.0
f_{solvent}	0.85	0.85	0.83
$V_{m,\text{Ca}^{2+}}/\text{cm}^3\text{mol}^{-1}$	19.6	17.3	14.0
R_h/nm^b	26.2 ± 0.9	30.9 ± 1.0	34.0 ± 1.0

^aQuantity shown for the sample in 100 % D₂O

^bHydrodynamic radius obtained from dynamic light scattering

also narrow but slightly broader than that of sample d₃-PA₃₆₀PSS₄₀₀, resulting in a value of $\frac{\sigma_{R_{\text{core}}}}{R_{\text{core}}} = 10.2\%$ and 10.5 %, respectively. This is in particular visible in the SAXS profiles, which show well-defined oscillations.

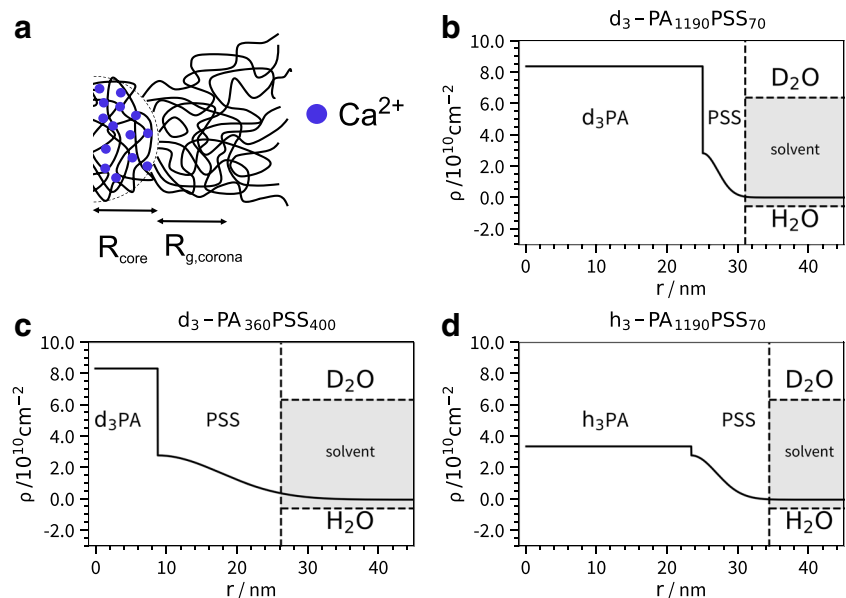
The radial scattering length density profiles of the micelles are shown in Fig. 9b–d. The density profile of the PSS corona was assumed to be Gaussian and is determined by the standard deviation s . It was shown previously that the overall dimensions of the corona of block copolymer micelles containing polyelectrolytes cannot safely be determined by small-angle scattering [49], as it is not trivial to define a reasonable cutoff criteria for the scattering length density of the corona. Hence, Förster [49] suggested using the hydrodynamic radius to estimate the dimension of the corona including the part with very low volume fractions on the surface of the micelle. Therefore, the scattering length density profiles shown in Fig. 9b–d also indicate the hydrodynamic dimensions of the micelles measured by dynamic light scattering.

Using this approach, the corona covers nearly $2 \cdot R_{g,\text{corona}}$, with $R_{g,\text{corona}}$ the radius of gyration of the polymer chain in the corona resulting from the Pedersen-Schurtenberger form factor [36]. We conclude that this method is a reasonable approach to estimate the overall dimensions of the corona.

The scattering length density profiles in Fig. 9 show the homogeneous micelle core composed of d₃-PA/h₃-PA, which is complexed by Ca²⁺ and considerably bigger for the h₃-PA₁₁₉₀PSS₇₀/d₃-PA₁₁₉₀PSS₇₀ micelles. Vice versa, the profiles highlight the substantially bigger corona for d₃-PA₃₆₀PSS₄₀₀ compared to h₃-PA₁₁₉₀PSS₇₀/d₃-PA₁₁₉₀PSS₇₀. The parameter s controlling the thickness of the corona is similar for d₃-PA₁₁₉₀PSS₇₀ (4.05 nm) and h₃-PA₁₁₉₀PSS₇₀ (3.53 nm), being in agreement with the radius of gyration $R_{g,\text{corona}}$ of 2.12 nm (d₃-PA₁₁₉₀PSS₇₀) and 2.07 nm (h₃-PA₁₁₉₀PSS₇₀).

The relatively small dimensions of the corona for h₃-PA₁₁₉₀PSS₇₀/d₃-PA₁₁₉₀PSS₇₀ pose the question if it is

Fig. 9 **a** Sketch of the d₃-PA₃₆₀PSS₄₀₀ micelles. **b** Scattering length density profile of d₃-PA₁₁₉₀PSS₇₀ micelles. **c** Scattering length density profile of d₃-PA₃₆₀PSS₄₀₀ micelles. **d** Scattering length density profile of h₃-PA₁₁₉₀PSS₇₀ micelles. The vertical dashed lines indicate the hydrodynamic dimensions of the micelles. The horizontal dashed lines indicate the scattering length density of H₂O ($-0.558 \cdot 10^{-10} \text{cm}^{-2}$) and D₂O ($6.355 \cdot 10^{-10} \text{cm}^{-2}$)



necessary to consider the PSS corona in the data analysis. Therefore, SANS curves of two selected contrasts (48.6 % and 100 % D₂O) and a SAXS curve of d₃-PA₁₁₉₀PSS₇₀ were analyzed with the model of a polydisperse sphere and are shown in Figure S5 of the Supporting Information. The model describes the contrast, where PSS is matched well (48.6 % D₂O). However, for the SANS curve with the contrast in 100 % D₂O and the SAXS curve, the model fails to describe the high q part as here the corona scattering contributes to the signal.

Besides the corona dimension, the interaction of the polymer chains in the corona can be derived from ν [28]. It has been demonstrated by Monte Carlo simulations that the corona of block copolymer micelles can be treated as quasi-two-dimensional dilute or semi-dilute polymer solution with an osmotic compressibility κ related to ν by $\kappa = 1 + \nu$ [35]. Moreover, ν depends on the reduced surface coverage $\frac{\Sigma}{\Sigma^*}$

$$\frac{\Sigma}{\Sigma^*} = \frac{N_{\text{agg}} R_{g, \text{corona}}^2}{4(R_{\text{core}} + R_{g, \text{corona}})^2} \quad (22)$$

according to a power law [28]. The reduced surface coverage is the two-dimensional analogue of $\frac{c}{c^*}$, where c^* is the overlap concentration [28]. The reduced surface coverage for d₃-PA₃₆₀PSS₄₀₀ is 1.3, which is above the critical value of unity for the overlap of the chains. This is compatible with the value of $\nu = 1.19$ inferred from the analysis of the scattering profiles. We thus conclude that the PSS chains in the corona of d₃-PA₃₆₀PSS₄₀₀ micelles are overlapping.

When calculating the reduced surface coverage for h₃-PA₁₁₉₀PSS₇₀ and d₃-PA₁₁₉₀PSS₇₀, one obtains 0.23 and 0.22, which is significantly smaller than 1.0, indicating that the chains in the corona do not overlap. This is in agreement with the values of ν , which converged to the lower boundary of 0.0 for both samples. The coronas of d₃-PA₁₁₉₀PSS₇₀ and h₃-PA₁₁₉₀PSS₇₀ micelles contain PSS chains which are not interacting and do not overlap. To confirm this hypothesis, we also analyzed the scattering profiles with a block copolymer form factor, assuming Gaussian chains grafted to the surface of a hard sphere [52]. This form factor is less general than the one used before since it is only valid for low surface coverage. The details of the model can be found in the Supporting Information (Figures S1–S3). We were able to describe the scattering data of d₃-PA₁₁₉₀PSS₇₀ and h₃-PA₁₁₉₀PSS₇₀ micelles with similar fit quality and micelle dimensions. However, when the simpler model [52] was applied to the profiles of d₃-PA₃₆₀PSS₄₀₀, we were not able to fit the contrast shown in Fig. 3d–f satisfactorily. The best fit with this model is also shown in Figure S3 of the Supporting Information. The simpler model [52] does not allow to take into account interactions between the chains in the corona by a ν parameter and thus

fails to reproduce the scattering signal which is dominated by the corona. Accordingly, the polymer chains in the corona for d₃-PA₁₁₉₀PSS₇₀/h₃-PA₁₁₉₀PSS₇₀ micelles are not overlapping, whereas PSS chains in d₃-PA₃₆₀PSS₄₀₀ micelles are overlapping.

We conclude the section by stressing again that the H₂O/D₂O ratio influences the aggregation number of d₃-PA₁₁₉₀PSS₇₀/h₃-PA₁₁₉₀PSS₇₀. This effect deserves further consideration. Figures 6b and 7b show that the obtained aggregation numbers of the micelles decrease with increasing content of heavy water. This slight decrease in aggregate size is attributed to an isotope effect occurring upon change of the solvent from light to heavy water, irrespective of the use of deuterated or hydrogenated PA and will be addressed in detail in the next section.

Isotope effect

In the previous section, it was found that the size of d₃-PA₁₁₉₀PSS₇₀ and h₃-PA₁₁₉₀PSS₇₀ micelles slightly decreases with increasing D₂O content. For the form factor fits, this changing micelle size was taken into account by attributing an individual aggregation number to each contrast and using this N_{agg} for the corresponding SANS and SAXS curves at the respective contrast.

This change in aggregation number is caused by an isotope effect occurring upon change of the solvent from light to heavy water and is observed for a polymer with a deuterated PA block as well as a hydrogenated PA block. Preceding works have shown that the solvent quality and the phase behavior of polymer-solvent systems can change if hydrogenated solvents are replaced by deuterated solvents [53]. Moreover, it has been shown that the phase boundaries of proteins in the presence of multivalent cations may change when going from light to heavy water [54]. These effects arise from the slightly different interaction of the deuterated solvent with the macromolecule. This seems to be also the case for the present system.

The isotope effect only appears for d₃-PA₁₁₉₀PSS₇₀ and h₃-PA₁₁₉₀PSS₇₀ but not for d₃-PA₃₆₀PSS₄₀₀. As the effect only appears for the polymers with the long PA block, it is possible that the binding of Ca²⁺ to PA plays an important role in the process. In order to clarify if the isotope effect is accompanied by a difference in binding of Ca²⁺ to PA in H₂O or D₂O isothermal titration calorimetry was performed in light and heavy water. This technique measures the heat of binding upon addition of Ca²⁺ to a solution of PA. Figure 10 shows the titration curves in H₂O and D₂O as well as the fit to the model of a single set of identical binding sites [46, 47]. Table 3 summarizes the obtained enthalpy of binding ΔH° , entropy of binding $T\Delta S^\circ$, binding constant K , and stoichiometry of binding n . From the titration curves, it is clear that the binding in H₂O and D₂O have identical

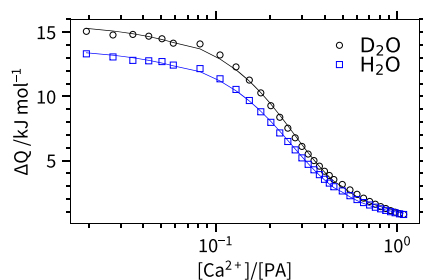


Fig. 10 Isothermal titration curves of PA homopolymer in the presence of Ca^{2+} in H_2O and D_2O . The curve shows the heat of binding per injection as a function of Ca^{2+}/PA ratio. The solid lines are fits with the model of a single set of identical binding sites. Table 3 summarizes the results of the data analysis

binding constants K ; however, the enthalpy of binding ΔH° and the entropy of binding $T\Delta S^\circ$ is slightly larger for the titration in D_2O . We attribute this slight difference also to an isotope effect and conclude that this isotope effect controls the binding of Ca^{2+} and the aggregation number of the micelles.

Stuhrmann analysis

In Section 2, it was shown that the SANS profiles from the contrast variation experiment can be well described by a suitable form factor model [28]. However, such an analytical expression for a form factor model is not always available, in particular for systems which are not as well-defined as block copolymer micelles. Moreover, deuteration is not always feasible as deuterated monomers do not exist for a wide range of polymers and deuteration of biological molecules such as proteins or sugars is often difficult and very cost intensive. Nevertheless, a contrast variation experiment is still possible if the scattering length densities of two components are sufficiently different.

In order to be able to analyze contrast variation experiments, Stuhrmann [29, 30] developed a model-independent approach, the so called Stuhrmann analysis. In the present section, this approach is employed for $\text{d}_3\text{-PA}_{1190}\text{PSS}_{70}$ as well as $\text{d}_3\text{-PA}_{360}\text{PSS}_{400}$ and compared with the results from the form factor fits. Eventually, the potential of the method is demonstrated by applying it

Table 3 Results of the ITC experiments probing the binding of Ca^{2+} to PA in H_2O and D_2O at 25°C

	H_2O	D_2O
$K/\text{L mol}^{-1}$	$6.3 \cdot 10^3 \pm 0.3 \cdot 10^3$	$6.4 \cdot 10^3 \pm 0.5 \cdot 10^3$
$\Delta H^\circ/\text{kJ mol}^{-1}$	17.45 ± 0.34	19.76 ± 0.56
$T\Delta S^\circ/\text{kJ mol}^{-1}$	39.14 ± 0.46	41.49 ± 0.75
$\Delta G^\circ/\text{kJ mol}^{-1}$	-21.69 ± 0.12	-21.73 ± 0.19
n	0.231 ± 0.002	0.239 ± 0.003

to the data of $\text{h}_3\text{-PA}_{1190}\text{PSS}_{70}$, nicely demonstrating that reliable structural information can be obtained even from a fully hydrogenated polymer and more importantly without necessarily using form factor fits.

Figure 11a shows the square root of the coherent SANS scattering intensity at $q \rightarrow 0$ for the measured contrasts as a function of D_2O content for all three samples. We used N_{agg} (determined from the form factor fits) for the samples $\text{d}_3\text{-PA}_{1190}\text{PSS}_{70}$ and $\text{h}_3\text{-PA}_{1190}\text{PSS}_{70}$ to normalize the coherent forward intensity from SANS in order to account for the effect of a change in aggregation number. The data follow a linear behavior as first shown by Stuhrmann [29, 30] which is expected when the contrast term $\Delta\rho$ is changed. A linear fit of the data shows that the match points of the block copolymer micelles is at $64.3 \pm 0.4\%$ D_2O ($\text{d}_3\text{-PA}_{360}\text{PSS}_{400}$), $125.6 \pm 0.7\%$ D_2O ($\text{d}_3\text{-PA}_{1190}\text{PSS}_{70}$), and $57.4 \pm 0.6\%$ D_2O ($\text{h}_3\text{-PA}_{1190}\text{PSS}_{70}$). Based on the chemical composition of the block copolymer and the scattering length densities of the two blocks, the theoretical match point of the block copolymers and their micelles can be calculated. The theoretical matchpoint for all three polymers is close to the experimentally found, which confirms the successful synthesis and preliminary characterization of the block copolymer.

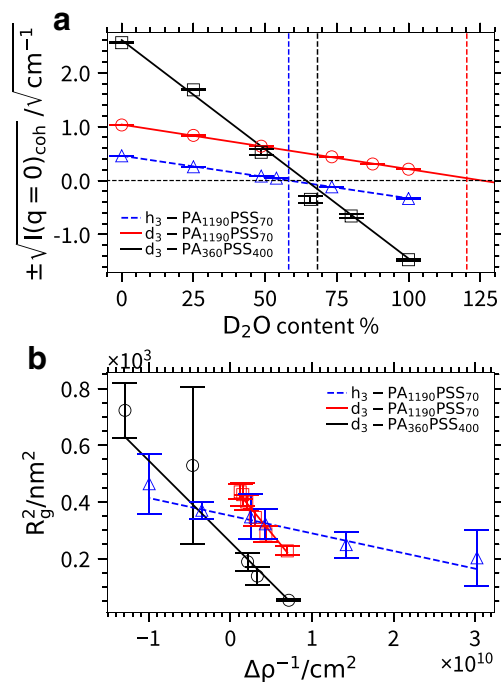


Fig. 11 a Coherent forward scattering for $\text{d}_3\text{-PA}_{360}\text{PSS}_{400}$, $\text{d}_3\text{-PA}_{1190}\text{PSS}_{70}$, and $\text{h}_3\text{-PA}_{1190}\text{PSS}_{70}$ from SANS as a function of D_2O content. The solid line represents a linear fit to the data. The data for $\text{d}_3\text{-PA}_{1190}\text{PSS}_{70}$ and $\text{h}_3\text{-PA}_{1190}\text{PSS}_{70}$ are normalized by the aggregation number N_{agg} . The vertical dashed lines indicate the theoretical match points calculated from the chemical composition. **b** Squared radius of gyration R_g^2 as a function of inverse scattering contrast for $\text{d}_3\text{-PA}_{360}\text{PSS}_{400}$, $\text{d}_3\text{-PA}_{1190}\text{PSS}_{70}$ and $\text{h}_3\text{-PA}_{1190}\text{PSS}_{70}$. The straight lines are fit to Eq. 23 [29, 30]

Further, more quantitative information can be extracted from the scattering curves by plotting the squared apparent radius of gyration R_g^2 from a Guinier analysis at low q as a function of the inverse scattering contrast $\Delta\rho^{-1}$. Figure 11 b shows the so called Stuhrmann plot [29, 30]. Ibel and Stuhrmann [29, 30] showed that for a two-component system R_g^2 depends on $\Delta\rho^{-1}$ as

$$R_g^2 = R_m^2 + \alpha\Delta\rho^{-1} + \gamma\Delta\rho^{-2} \quad (23)$$

with R_m being the radius of gyration at infinite contrast. The coefficient α is related to the second moment of the scattering length density fluctuations about its mean value. The coefficient γ is the square of the first moment of the density fluctuations about the mean. If α is positive, the component of the lower scattering length density is located more towards the inside of the objects. For a negative value of α , the component with the higher scattering length density is concentrated in the inside. γ is proportional to the squared separation of the two components, with $\gamma = 0$ indicating that the centers-of-mass are coinciding.

Figure 11 b shows R_g^2 as a function of $\Delta\rho^{-1}$ for the three samples. For all three polymers, the points follow a straight line with γ being close to zero and with a negative slope α .

For all three samples, a negative value for α indicates that the PA blocks are always located in the core of the micelles. It may not have been clear at the outset of the experiment that this also holds for the fully hydrogenated sample h₃-PA₁₁₉₀PSS₇₀. However, calculation of the scattering length densities ρ for PSS ($2.82 \cdot 10^{10} \text{cm}^{-2}$) and h₃-PA ($4.21 \cdot 10^{10} \text{cm}^{-2}$) reveals that they are sufficiently different also for the fully hydrogenated sample, with h₃-PA having a larger ρ . This nicely demonstrates the benefit of additionally analyzing a fully hydrogenated sample.

The fact that γ is close to zero shows that the center-of-mass of PSS and d₃-PA/h₃-PA coincide. Such a situation is only compatible with a highly symmetrical core-shell structure of the micelles. Furthermore, the scattering curves do not show any indication for the presence of cylindrical or vesicular micelles, which would also be compatible with a value of γ close to zero. Accordingly, we can already conclude from the model free analysis and without using information from the form factor fits that the micelles are spherical with a core composed of d₃-PA/h₃-PA and a corona of PSS, being in full agreement with the form factor fits.

Using the parallel axis theorem, one can relate α and R_m^2 to the radius of gyration of the core $R_{g,\text{core}}$ and the radius of gyration of the shell $R_{g,\text{shell}}$ [55]

$$R_m^2 = f_{\text{core}}R_{g,\text{core}}^2 + f_{\text{shell}}R_{g,\text{shell}}^2 \quad (24)$$

$$\alpha = (\rho_{\text{core}} - \rho_{\text{shell}})f_{\text{core}}f_{\text{shell}}(R_{g,\text{core}}^2 - R_{g,\text{shell}}^2) \quad (25)$$

with f_{core} and f_{corona} being the volume fraction of core and corona block and ρ_{core} and ρ_{corona} the scattering length densities of the core and corona block, respectively. All the information is available from the chemical composition and molar volumes of the monomer units. Equations 24 and 25 can be solved for $R_{g,\text{core}}$ and $R_{g,\text{shell}}$, making the dimensions of the micelles directly accessible from a model-independent analysis of the scattering data. Table 4 summarizes the values of R_m , α , $R_{g,\text{core}}$, and $R_{g,\text{shell}}$ for all three samples.

For d₃-PA₃₆₀PSS₄₀₀, a value of $R_{g,\text{core}} = 7.1$ nm and $R_{g,\text{shell}} = 18.0$ nm is found. This indicates that the micelles contain a d₃-PA core which is surrounded by a rather thick shell of PSS, in perfect agreement with the scattering length density profiles obtained from the form factor fits. The core radius R_{core} from the form factor fits can be used to calculate a value for $R_{g,\text{core}}$ by [31]

$$R_{g,\text{core}} = \sqrt{\frac{3}{5}}R_{\text{core}} \quad (26)$$

$R_{g,\text{core}}$ calculated accordingly is 6.8 nm, in perfect agreement with the value obtained from the model independent analysis (7.1 ± 2.5 nm).

Using Eqs. 24–25 for d₃-PA₁₁₉₀PSS₇₀ yields $R_{g,\text{core}} = 19.9$ nm and $R_{g,\text{shell}} = 29.7$ nm, nicely illustrating the considerably bigger d₃-PA core for this sample and the rather thin PSS corona. Comparing the core radius from the form factor analysis with $R_{g,\text{core}}$ using Eq. 26 yields a value of 18.4 nm from the form factor fit. This shows a good agreement between both methods, in particular when taking into account that the isotope effect and therefore the slightly changing micelle dimensions is not considered in Fig. 11b of the Stuhrmann analysis.

Finally, the micelle dimensions for the fully hydrogenated polymer h₃-PA₁₁₉₀PSS₇₀ are calculated. The values of $R_{g,\text{core}} = 16.8$ nm and $R_{g,\text{shell}} = 30.1$ nm are very similar to the results found for d₃-PA₁₁₉₀PSS₇₀ micelles, showing that the h₃-PA₁₁₉₀PSS₇₀ micelles also consist of a

Table 4 Micelle dimensions obtained from the Stuhrmann plot (R_g^2 versus $\Delta\rho^{-1}$)

Polymer	R_m / nm	α	$R_{g,\text{core}}$ / nm	$R_{g,\text{shell}}$ / nm
d ₃ -PA ₃₆₀ PSS ₄₀₀	16.1 ± 1.4	$(-2.9 \pm 0.3)10^6$	7.1 ± 2.5	18.0 ± 1.7
d ₃ -PA ₁₁₉₀ PSS ₇₀	21.6 ± 1.0	$(-3.5 \pm 0.4)10^6$	19.9 ± 1.1	29.7 ± 4.9
h ₃ -PA ₁₁₉₀ PSS ₇₀	18.7 ± 1.2	$(-6.2 \pm 2.5)10^5$	16.8 ± 2.3	30.1 ± 5.1

large PA core and a thin PSS corona. The radius of gyration of the core according to Eq. 26 and the form factor fit is 17.3 nm, which is in perfect agreement with the result from the Stuhmann plot (16.8 ± 2.3 nm).

The quantitative discussion presented above is a nice verification of the power of the Stuhmann analysis, for which we could provide a highly suitable model system. The model-independent analysis does not only offer values for important size parameters of the micelles, but provides evidence for a core-shell structure of these micelles, with the PA blocks located at the core of the micelles. These features and conclusions could be consistently and independently derived from investigating three different block copolyelectrolyte samples of the type PA-*b*-PSS. This nicely shows the potential of the model free analysis given in the present section and at the same time confirms the form factor fits.

Conclusion

We investigated self-assemblies of block copolymers of the two polyelectrolytes polyacrylate (PA) and polystyrene sulfonate (PSS) by a combination of contrast variation small-angle neutron scattering, small-angle X-ray scattering, and dynamic light scattering. In a previous work [27], we showed that such block copolymers form well-defined micelles in the presence of Ca^{2+} . In order to investigate the morphology of these micelles, the PA block was selectively deuterated.

First and foremost, SANS and SAXS reveal a spherical micellar shape for the aggregates with a well-defined size and a low polydispersity for three different block copolyelectrolytes of the type PA-*b*-PSS, using the model of a block copolymer micelle with a homogeneous scattering length density in the core and self-avoiding chains in the corona [28]. Model interpretation confirmed the micellar shape of the aggregates and unambiguously proved that Ca^{2+} cations are binding to PA and that this complex forms the hydrophobic core of the micelles. This core contains up to 85 vol% of water. PSS forms the corona of the micelles and adopts the conformation of self-avoiding polymer chains. Furthermore, we found that the PSS chains in the corona do not overlap for a block copolymer with a high fraction of PA, while they start to show excluded volume effects once they overlap due to an increase of the PSS fraction. As the interactions of the Ca^{2+} cations with the PSS chains are only non-specific being dominated by electrostatically based screening of electrostatic forces, the resulting PSS corona stabilizes the micelles and keeps the phase separation otherwise observed with pure PA and Ca^{2+} confined to the core of the micelles.

For a set of block copolymers with a long PA and a short PSS block, a small change in dimensions was

observed in the SAXS profiles when going from H_2O to D_2O . This was attributed to an isotope effect arising from a slightly different binding enthalpy of Ca^{2+} to PA in H_2O and D_2O . Having SAXS and contrast variation SANS data available, it was possible to take this into account in the data analysis as a change in aggregation number. Such effects are at risk to be neglected in SANS based contrast variation experiments and the present example stresses the importance of combining SAXS and SANS in order to elucidate such effects and take them into account during the data analysis.

Finally, we demonstrated that contrast variation can also be performed on a fully hydrogenated block copolymer as long as the scattering length densities of the blocks are sufficiently different. In combination with the SAXS data, it was possible to reliably determine the size, structure, and morphology of the micelles. This is particularly interesting for samples where deuteration is not possible and nicely illustrates the future potential of novel, more powerful neutron sources.

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Compliance with Ethical Standards

Conflict of interests The authors declare that they have no conflict of interest.

Dedication The authors are happy to dedicate this work on small-angle scattering of block copolyelectrolytes to Matthias Ballauff. The block copolyelectrolytes reversibly form micelles in response to interactions with Ca^{2+} cations. It is a happy coincidence that this work nicely fits into the broader scope of Matthias' scientific work. This does not come as a surprise though as Matthias' accomplishments inspired the present as well as some of our other work. Above all, the authors are grateful to him for having introduced them to the power of ASAXS. They wish Matthias Ballauff all the best for his future!

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