

Chain Dynamics and Intermolecular Association in Dilute Aqueous Solutions of Isotactic and Atactic Poly(Methacrylic Acid): Effect of NaCl Concentration

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Abstract

Dynamic and static light scattering measurements were performed on isotactic and atactic forms of poly(methacrylic acid), iPMA and aPMA, respectively, in order to perform a wide survey of their molecular properties and chain dynamics in water as functions of tacticity, degree of neutralization, α_N , and salt concentration, c_s . The molecular parameters of PMA chains were analyzed at low α_N and chain dynamics (diffusion coefficients and in this connection the polyelectrolyte slow mode behavior) at higher α_N . The data revealed that both PMAs form microgel-like aggregates with a core-shell structure at low α_N ($= 0$ and 0.25 for aPMA and iPMA, respectively). The distribution of the hydrophilic and hydrophobic functional groups within the aggregates and their compactness depended considerably on chain tacticity and for aPMA also on c_s . Further, the effect of c_s on the polyelectrolyte slow diffusion coefficient, D_s , of partly ($0.25 \leq \alpha_N < 1$) or completely charged PMA polyions ($\alpha_N = 1$) was studied. In iPMA solutions, D_s was detected up to $c_s = 0.1$ M, regardless of α_N , whereas in aPMA solutions this c_s value gradually decreased with decreasing α_N . These differences were also related to tacticity and through that to rigidity of the studied forms of PMA. It is argued that the segregation of uncharged and charged carboxyl groups is operative over the entire α_N region in iPMA solutions.

1. Introduction

Response of polymers in solution to changes in environment is the basis for their use as advanced materials in numerous technological and biochemical applications [1,2,3]. In all these cases, polymers respond to certain external stimuli and often self-associate into nano-structured soft materials with a huge variety of properties and potential functionalities. The stimuli to achieve or manipulate polymer self-association (self-assembly) in aqueous solutions are most often changes in pH, ionic strength, temperature, and/or even mechanical stress [4]. Some polymers respond to a combination of two or more stimuli [5], which enables very effective creation of soft materials. Situation is even more complex in case that a polymer chain can acquire charge through ionization of its functional groups, which may lead to additional options for customizing properties, for example through viscosity increase.

Poly(methacrylic acid), PMA, is a synthetic homopolymer that responds to all of the above stimuli and is a potential polyelectrolyte as well. Historically, the reason for the interest in PMA is that its chain exhibits a reversible and cooperative change in conformation as a result of the change in solution's pH, which affects ionization of carboxyl groups [6-10]. In acidic conditions, when unionized, PMA chain is in a compact form, whereas in basic conditions, when fully ionized, it adopts a more extended coil-like conformation, thus enabling the release of ligands bound within the compact PMA core. This principle is widely exploited in various pharmaceutical and biomedical applications [3,11-13].

Much less studied is the fact that behavior of PMA depends significantly on so-called tacticity or stereoregular structure of the polymer chain. Stereoregularity refers to the chain microstructure, *i.e.* to the spatial distribution of functional groups along the chain, which can be highly regular (isotactic and syndiotactic polymers) or irregular (heterotactic or so-called atactic polymers). The effect of tacticity on properties of polymers is an important issue in polymer chemistry [14]. However, it is surprising that almost no attention is devoted to the

effect of tacticity on solution behaviour of polyelectrolytes, which is also PMA when its carboxyl groups are ionized. This contribution aims at filling this gap by studying the behavior of isotactic and atactic PMA, iPMA and aPMA, respectively, in the whole range of degrees of neutralization, α_N , of carboxyl groups, from $\alpha_N = 0$, when PMA chains are uncharged and can potentially form nanoparticles through self-association, to $\alpha_N = 1$, when they are fully charged and electrostatic interactions become important. As far as the authors are aware, such a wide survey has not yet been carried out for PMA isomers in relation to chain tacticity and ionic strength of solutions. The ionic strength is normally varied by adding a low molar mass salt. The added salt is important not only when the polymer is highly charged (in electrostatic screening), but also at low charge through the competition between the added ions and the polymer chain for hydration with water molecules. This issue is well known in protein research.

Properties of the highly regular iPMA are relatively poorly investigated, both at low and in particular at gradually increasing charge. Until recently, it was known only that iPMA does not dissolve in water unless around 20% of its carboxyl groups are ionized (charged), whereas aPMA dissolves already in the pure acid, and that the well-known conformational transition of iPMA takes place along different paths when the polyacid is either protonated or deprotonated [8-13], whereas both paths coincide in the aPMA case [8-10,14-21]. Possible molecular interpretation of these observations was recently offered by a complete thermodynamic analysis of the conformational transition [10]. It was shown therein that iPMA and aPMA respond oppositely to the change in temperature, as clearly proven by the opposite sign of the heat capacity change of the conformational transition. In agreement with these thermodynamic result, van den Bosch *et al.* [19] have shown by rheology that iPMA at $\alpha_N \approx 0.2$ is associated and forms gel upon cooling, whereas Eliassaf and Silberberg [22] demonstrated that the unionized aPMA ($\alpha_N = 0$) does that upon heating. Another striking

difference is the response of PMA isomers to shear: aPMA chains associate when subjected to shear [23-25], whereas the aggregates between iPMA chains decompose when the solution is sheared and reform in solution at rest [20,21].

These observations apply to the low α_N limit and devote no attention to the role of the added salt, *i.e.* to the ionic strength of solutions. Moreover, the effect of tacticity on the state of the highly charged PMA polyions (high α_N) is mostly not disclosed, although it is expected that the ionic strength should strongly affect the interactions between polyions themselves and also their interactions with oppositely charged small ions in solution. For example, Jerman *et al.* [26] have determined the osmotic coefficients of iPMA and aPMA in dependence on degree of ionization, but only in salt free solutions. They concluded that iPMA chain has a higher local charge density than the aPMA one and thus binds counterions more strongly [26,27], which leads to lower osmotic coefficients. The effect of tacticity should also be reflected in diffusion coefficients (chain dynamics) of PMA polyions in solution, which can be determined by dynamic light scattering, DLS. In DLS, strong electrostatic correlations between polyions in low salt conditions lead to a characteristic feature known as the polyelectrolyte, PE, mode. Two relaxation processes are observed by DLS in PE solutions [28-33,3537]. The fast relaxation process stems from a coupled diffusion of polyions and their counterions and the slow relaxation process (revealed as an “apparent” slow diffusion coefficient, D_s) is attributed to hindered diffusion of polyions due to their high charge. It is labelled by various terms: as an “extraordinary” [28] or even “anomalous” [34] mode, “slow” mode, as “multimacroion-domain” [41] (or “cluster”)-formation, polyion-lattice-formation [41] etc.

It was realized as early as 1976 in the DLS study of poly(L-lysine) in aqueous NaBr solutions by Lin *et al.* [28] that salt concentration, c_s , significantly affects Brownian motion of polyions. The authors identified a rather sharp c_s value below which the PE-slow mode

(D_s) in PE solutions suddenly appeared and simultaneously light scattering intensity dropped substantially. Sedláč has extensively investigated this feature in PE solutions, including solutions of the usual (or conventional) aPMA form [39-41], and attributed the PE-slow mode to the formation of so-called multimacroion domains in PE solutions with low ionic strength. It was shown that when the concentration of added salt (*i.e.* the ionic strength) is sufficiently high, the PE-slow mode disappears. We should stress that studies of chain dynamics (slow mode diffusion coefficients) in relation to PMA tacticity are scarce [20,21]. Specifically, there are no reports on the effect of c_s and α_N on D_s of iPMA and aPMA polyions in aqueous solutions.

Consequently, in order to fully understand the behavior of PMA isomers in water, investigations of the effect of ionic strength on the molecular state of PMAs in relation to their tacticity are needed, both at low α_N as well as at higher α_N . In this contribution we therefore discuss two subjects: (i) the tendency of iPMA and aPMA towards intermolecular association (low α_N conditions) and of (ii) polyion chain dynamics in dependence on polyion charge (high α_N conditions, *i.e.* up to $\alpha_N = 1$) in a broad range of salt (NaCl) concentrations, which served to vary the ionic strength of the solutions. Both, static light scattering, SLS, and DLS were used to determine the size and diffusion properties of polyions. We first investigate the effect of c_s on the hydrodynamic radius, R_h , and the radius of gyration, R_g , of PMA particles in solution at low α_N ($\alpha_N = 0$ for aPMA or somewhat higher, *i.e.* $\alpha_N = 0.25$, for iPMA, due to solubility issues) and then treat the effect of c_s on the diffusion coefficients of partly ($\alpha_N < 1$) or completely charged polyions ($\alpha_N = 1$). Such a wide survey of tacticity effects on chain dynamics (in particular on so-called polyelectrolyte slow mode) in solutions of iPMA and aPMA is presented for the first time.

2. Experimental section

2.1 Materials

PMA samples with well-defined stereoregular composition (tacticity) were from Polymer Source Inc. and were the same as used previously [20,21]. The molecular properties of iPMA were the following: the weight and number average molar mass, M_w and M_n , respectively, were $M_w = 32000 \text{ g mol}^{-1}$ and $M_n = 11000 \text{ g mol}^{-1}$ and the tacticity was 93% of isotactic, 3% of syndiotactic, 4% of atactic triads. The aPMA sample (Polymer Source Inc., sample no. P14096-MAA: $M_w = 189000 \text{ g mol}^{-1}$ and $M_n = 165000 \text{ g mol}^{-1}$) was rich in syndiotactic triads (almost 70% syndiotactic). Additional details on properties and preparation of polymers are reported in our previous publications [20,21]. NaCl, NaOH (Fixanal), and HCl (Fixanal) were from Merck KGaA. Triple distilled water was used to prepare all solutions. All solvents (water and aqueous NaCl solutions) were filtered through 0.1 μm filters prior to solution preparation.

2.2 Preparation of solutions

Stock solutions of iPMA and aPMA in water at various α_N values were prepared as reported previously [20] by bearing in mind that iPMA is not soluble in water or in aqueous NaCl solutions unless α_N exceeds ~ 0.2 . Thus, solid iPMA was first suspended in water and neutralized to around $\alpha_N \approx 0.5$ by adding 1 M NaOH so that the polymer dissolved completely. The α_N values below 0.5 were then prepared by adding 1 M HCl and those above 0.5 by adding 1 M NaOH. Salt (NaCl) concentration, c_s , in these stock solutions was adjusted to the desired value by adding a calculated amount of either 0.1 M ($c_s < 0.07 \text{ M}$) or 1 M NaCl ($c_s \geq 0.07 \text{ M}$).

Solid aPMA, which is soluble in water at any α_N including $\alpha_N = 0$, was dissolved directly in water. The stock solutions with $\alpha_N > 0$ were prepared by adding a calculated amount of 1

M NaOH to this solution. As above, required amounts of 0.1 M or 1 M NaCl were added to stock solutions to adjust NaCl concentration to the desired value. Experiments for both, iPMA and aPMA, were performed in the range $0.01 \text{ M} < c_s < 0.5 \text{ M}$. Polymer concentration, c_p , was set to $c_p = 2 \text{ g L}^{-1}$ (at $\alpha_N = 0$), which corresponds to 0.023 moles of COOH groups per volume (abbreviated as M for mol COOH/L). To make sure that the studied polymer concentration corresponds to dilute region, the overlap concentration, c_p^* ($= \frac{M_w}{N_A(4\pi R_h^3/3)}$, where N_A is the Avogadro constant), was estimated from the measured R_h values of particles (single polymer chains, designated as $R_{h,1}$, *c.f.* Table S1 in Supporting Information, SI, were used for this estimation). The calculated c_p^* was in the range (11-16) and (33-51) g L^{-1} for iPMA and aPMA, respectively, showing that measurements were performed well below c_p^* , *i.e.* under dilute conditions.

2.3 Dynamic and Static Light Scattering

DLS and SLS measurements were conducted with a 3D-DLS spectrometer from LS Instruments GmbH (Fribourg, Switzerland). The instrument operates in the 3D cross-correlation mode at a wavelength of incident radiation, λ_o , equals to 632.8 nm, which is provided by the 20 mV He-Ne laser (Uniphase JDL 1145P). The 3D cross-correlation technique was especially developed for studying strongly scattering samples in order to suppress multiple scattering [42]. Other details regarding the instrumentation were reported previously [20,21] and can be found in SI.

All LS studies were performed at 25 °C. Solutions were carefully filtered through hydrophilic and low protein binding Millex-HV filters (diameter 13 mm, pore size 0.22 μm) directly into the sealable dust-free cylindrical quartz sample cell. The cell was thoroughly purified before filling it with the sample. The samples were allowed to equilibrate for 30 min

in the decalin bath before the measurement was initiated. Scattered light was collected in the angular range from 40° to 150° with a step of 10° .

Details about the methodological aspects of DLS and SLS can be found in SI and in literature [43-45], so as all other aspects of the data analysis used in this paper [20,21]. Shortly, the correlation functions of the intensity of scattered light, $G_2(q,t)$, were recorded simultaneously with the integral time-averaged LS intensities, $I_\theta \equiv I_q$, where q ($= (4\pi n_0/\lambda_0)\sin\frac{\theta}{2}$) is the scattering vector, θ is the angle, and n_0 is the refractive index of the medium). The measured intensities were normalized with respect to the Rayleigh ratio, R , of toluene. From the measured $R(q)$ for solutions, the radius of gyration, R_g , of scattering particles was evaluated (Eqs. S1a and S1b, SI). In order to determine R_g of particles, two approaches were used: (i) for small particles (with sizes that fulfill the criterion $qR_g < 1$) the Zimm function (Eq. S2) was employed and (ii) for larger particles the Debye-Bueche scattering function (Eq. S3) was found to be the most suitable for R_g determination. For additional comments on these functions and for particle topologies that they describe, see SI and references 46-52. In order to determine the diffusion coefficients, D , and hydrodynamic radius, R_h , of particles the $G_2(q,t)$ function were converted into the time correlation function of the scattered electric field, $g_1(q,t)$, by means of the Siegert's relationship (Eq. S4, SI). Most of the $g_1(q,t)$ functions for the herein studied systems were bi-modal (see Eqs. S5), having two well-separated peaks in the calculated relaxation times, τ , distributions. The analysis of the bi-modal $g_1(q,t)$ functions was performed as reported previously [21] (see SI) and resulted in the relative distribution functions, $A_1(q,\tau_1)$ and $A_2(q,\tau_2)$, of the corresponding mean relaxation times, τ_1 and τ_2 , for species 1 and 2 (example of such procedure is demonstrated in Figure S1 together with the procedure for splitting the total LS intensity into separate contributions of both populations). When appropriate, the distributions over τ were converted into size (R_h) distributions by means of the relationship $\tau^{-1} = Dq^2$ and the Stoke-Einstein

equation (Eq. S6). The evaluated fast translational diffusion coefficient, D_f , is associated with individual aPMA chains (small particles with $R_{h,1}$), whereas the slow translational diffusion coefficient, D_s , represents intermolecular associates (large particles with $R_{h,2}$). In other cases, when the polyelectrolyte effect is observed, the fast and slow relaxation rates, Γ_f and Γ_s (where $\Gamma = \tau^{-1} = Dq^2$), respectively, do not represent translational diffusion. Consequently, we do not use terminology of sizes but speak about slow and fast diffusive processes. Finally, the determined $R_{g,2}$ and $R_{h,2}$ values for the aggregates are combined to give so-called shape parameter [44], $\rho (= \frac{R_{g,2}}{R_{h,0}}$, where $R_{h,0}$ is the hydrodynamic radius extrapolated to $\theta = 0^\circ$ or $q = 0$).

2.4 Dn/dc measurements

A Wyatt Optilab rEX differential refractometer ($\lambda_0 = 658$ nm) was used to determine the refractive index increment, dn/dc , of aPMA and iPMA solutions. The dn/dc values were measured for aPMA at $\alpha_N = 0$ and $c_s = 0.01, 0.02, 0.04, 0.05, 0.07, 0.1, 0.15, 0.2,$ and 0.5 M NaCl, and for iPMA at $\alpha_N = 0.25$ and $c_s = 0.02, 0.05, 0.07, 0.1, 0.2,$ and 0.5 M NaCl. In these experiments, the polymer concentration was $0.5 - 4$ g L⁻¹.

2.5 pH measurements

pH values were measured with a combined glass electrode (Mettler Toledo Inlab) connected to the pH meter. The electrode was calibrated with borate and phthalate buffers having pH = 9.180 and 4.006 at 25 °C, respectively.

3. Results and discussions

A natural way to present the results for PMA isomers is to start with the low α_N limit and continue with studies at higher α_N values. The first part is therefore devoted to intermolecular association, which is observed at low polyion charge (low α_N), and its dependence on c_s and

the second one to chain dynamics and the polyelectrolyte slow mode behavior at higher polyanion charge in dependence on both, α_N and c_s .

3.1 Intermolecular association at $\alpha_N = 0.25$ (iPMA) and $\alpha_N = 0$ (aPMA) and structural characteristics of the aggregates

LS was first used to determine the c_s dependence of R_h , R_g , and ρ values in aqueous iPMA and aPMA solutions at the lowest possible α_N , which allowed solubility of both PMAs: aPMA could be studied at $\alpha_N = 0$, while for iPMA α_N had to be higher (see Experimental section) and so $\alpha_N = 0.25$ was used in this case. The reason why $\alpha_N = 0.25$ was not employed for aPMA is that aggregation is no longer observed in aPMA solution with $\alpha_N \geq 0.25$, irrespective of c_s (see plots of LS intensity in Figure S6b: the sharp drop in LS intensity is interpreted as the disappearance of the aggregates; for more detail see discussion below).

Multi exponential CONTIN analysis of $G_2(t)$ functions (shown in Figure S2) results in bimodal distributions of the relaxation times τ (*c.f.* insets in Figure S2) at all scattering angles and all c_s values. At the same time, the LS intensity from these solutions is very high (*c.f.* Figure S6), suggesting that large particles are present in solution. The corresponding fast and slow diffusive processes (or modes) are therefore associated with small (individual) PMA chains and with large intermolecular aggregates between chains, respectively. Figure S3 shows the corresponding dependencies of Γ_f and Γ_s on q^2 for small and large particles, respectively. Both curves go through the center of coordinates and thus represent a true translational diffusion (*i.e.* real particles in solution). Consequently, two dynamic modes can be analyzed individually and the corresponding relative amplitudes of both modes and D and R_h ($R_{h,1}$ and $R_{h,2}$) values can be calculated.

The Γ_f versus q^2 dependences (small particles; Figure S3, the upper panels) are all perfectly linear, showing that small particles have narrow size distributions, whereas the Γ_s versus q^2 curves for the aggregates (Figure S3, the lower panels) show positive deviations

from linearity at higher q^2 , suggesting that aggregates are much more polydisperse in size. The diffusion coefficients for the small and large particles are determined from the initial slopes of these curves and therefrom the $R_{h,1}$ and $R_{h,2}$ values are calculated. In case of the aggregates, the radius of gyration, designated as $R_{g,2}$, can be determined from the angular dependency of the scattered light intensity (for details see SI). On the other hand, individual chains are too small to display any angular dependency and consequently $R_{g,1}$ could not be evaluated. The obtained results for both PMAs are summarized in Table S1 and plotted in Fig. 1 in dependence on c_s . Note that the $R_{h,2}$ data apply to zero angle.

The hydrodynamic radius of the smaller particles ($R_{h,1}$ ~10 nm for iPMA and ~11-13 nm for aPMA), is independent of c_s , whereas the hydrodynamic radius of the aggregates, $R_{h,2}$, depends on c_s and the data display a distinctively different dependence for iPMA and aPMA (Fig. 1b). In the iPMA ($\alpha_N = 0.25$) case, $R_{h,2}$ ($R_{g,2}$) gradually increases with increasing c_s from ~100 (~65) to ~155 (~97) nm. Obviously, increase in the ionic strength promotes formation of increasingly larger iPMA aggregates, which is observed also as a strong increase in the absolute LS intensity (Fig. 1a). Taking into account that iPMA chains are charged, the growth of the aggregates with increasing c_s is expected and results from the screening effect of added salt. In addition, Fig. 1c shows ρ as a function of c_s . Similarly to $R_{h,2}$ and $R_{g,2}$, ρ is practically independent of c_s in the iPMA case ($\rho = 0.60 - 0.63$).

For aPMA ($\alpha_N = 0$), on the other hand, a pronounced maximum in the aggregate size is seen at $c_s \approx 0.07$ M and more or less constant (or even decreasing) values are observed for $c_s \geq 0.15$ M, whereas $R_{h,2}$ and $R_{g,2}$ increase in this c_s range for iPMA. The lowest $R_{h,2}$ (≈ 120 nm) and $R_{g,2}$ (≈ 85 nm) are found for aPMA at $c_s = 0.01$ M and the highest ones, $R_{h,2}$ (≈ 310 nm) and $R_{g,2}$ (≈ 250 nm) at $c_s \approx 0.07$ M. The same pattern in the c_s dependency applies to ρ values, which are higher for aPMA ($\rho = 0.64-0.8$) in comparison with iPMA solutions and display a clear maximum ($\rho = 0.80$) at $c_s \approx 0.07$ M.

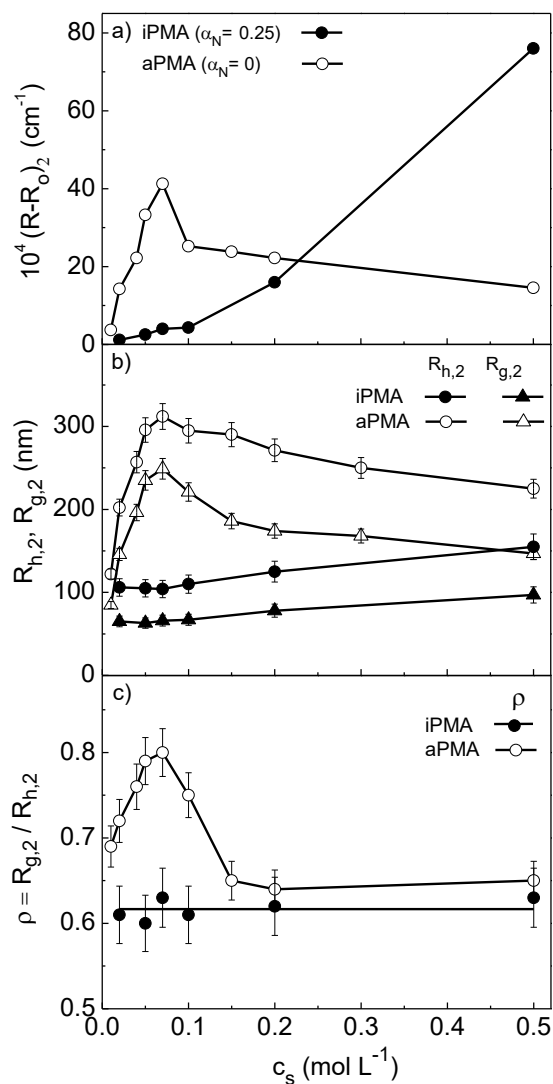


Fig. 1. a) The contribution of the aggregates (index 2) to the total LS intensity (represented as the difference $(R - R_o)_2$, where R_o is the contribution of the solvent to R at $\theta = 0$); b) the $R_{h,2}$ and $R_{g,2}$, and c) ρ ($= R_{g,2}/R_{h,2}$) for the aggregates as a function of c_s in aqueous iPMA ($\alpha_N = 0.25$; full symbols) and aPMA ($\alpha_N = 0$; open symbols) solutions at 25 °C. The contribution of individual chains to the total LS intensity is very small and is not plotted in Fig. 1a. Also, $R_{h,1}$ values are not included in the plot because they are small and independent of c_s (for $R_{h,1}$ see Table S1). The error bars in Fig. 1a do not exceed the size of the data points.

Parameter ρ is a structure sensitive factor that can be used for qualitative interpretation of the aggregate structure in terms of mass distribution within the aggregates. The ρ values have

been reported for various particle topologies [43,44,53]. For example, ρ would be ≥ 2 for fairly stiff and expanded objects (rigid rods), around 1.5 for random coils in a θ -solvent, 1 for hollow spheres and 0.778 for homogeneous hard spheres. The value of ρ for the herein studied iPMA and aPMA aggregates is considerably lower than 1 (in the range 0.6-0.8, signifying that $R_{h,2}$ is much larger than $R_{g,2}$), which points to structures with a rather high packing density in their interior in comparison to the outer shell/corona. Although ρ for aPMA is close to the value for a homogeneous (impenetrable) hard sphere (e.g. $\rho = 0.80$ in 0.07 M NaCl), this shape is not anticipated for aPMA aggregates due to the large size of these objects and to the hydrophobic character of the (methyl) side groups on the chain.

Burchard *et al.* report very low ρ values ($\rho \approx 0.60$ [54,55] or even as low as $\rho \approx 0.35 - 0.55$ [56,57]) for the chemically cross-linked microgel-like particles formed in ethanol solutions of poly-disperse hyper-branched poly(N-vinylimidazole), PVI. In the PVI case, such low ρ is attributed to chains dangling out of the highly contracted core, meaning that mass is concentrated in the center of the particle. An analogous structure is proposed for PMA aggregates in our study. They are similar to microgel-like particles with a core-shell structure, in which the core is considered to be compact and undrained by the solvent, whereas the shell or corona is penetrable for the solvent and thus swollen. In contrast to PVI, PMA chains are not chemically cross-linked but are held within the aggregate by physical forces. These forces are an interplay of rather strong (and in case of iPMA cooperative [10,19]; see detailed discussion below) hydrogen bonds, so-called hydrophobic interactions and van der Waals forces.

A question arises why do iPMA ($\alpha_N = 0.25$) chains form aggregates with a more compact core than do aPMA ($\alpha_N = 0$) ones, although they are substantially more charged. Intuitively, one would expect a more loose association and therefore higher ρ for iPMA with $\alpha_N = 0.25$ due to larger repulsion between charges on the partly ionized chains. However, important is

first of all chain tacticity and based on this different solute-solvent interactions or affinity of the PMA isomer for the aqueous environment. It has been demonstrated through the analysis of temperature dependence of the thermodynamic functions accompanying ionization of carboxyl groups [10] that iPMA behaves as a typical polar (hydrophilic) and aPMA as a nonpolar (hydrophobic) solute in water. These differences in water affinity can be interpreted by proposing a different distribution of hydrophilic (COOH and also COO⁻) and hydrophobic (CH₃) groups within the aggregate, which is schematically depicted in Scheme S1. The aggregates formed between iPMA chains ($\alpha_N = 0.25$) have most of their ionized COO⁻ groups situated on the outside of the aggregate and the unionized ones (COOH), together with the hydrophobic CH₃ groups, are buried in the aggregate interior. That is, a segregation of COO⁻ and COOH groups within the aggregate is proposed: its corona consists of charged chain portions extending into the solution, whereas the core is deficient in charges. The iPMA ($\alpha_N = 0.25$) aggregate is thus actually hydrophilic outwards. This picture agrees with pyrene fluorescence measurements [27]. It has to be noted that normally (for a single isolated chain) an even distribution of COOH and COO⁻ groups along the polyion chain is imagined, which results in a certain average charge density for the whole chain. On the other hand, by permitting segregation of charged and uncharged groups the local degree of ionization within the core, α_N^{core} , can easily be close to 0, which then allows strong and in case of iPMA also cooperative intermolecular H-bonding between the “core” COOH groups, as also highlighted in Scheme S1. The basis for this cooperativity is the regular isotactic arrangement of carboxyl groups [10,19-21] that enables a “zipper-like” mechanism of inter-chain connection. That is, when one H-bond is formed the others follow simultaneously, as in cooperative conformational changes. It can be imagined that such an interaction results in a much contracted iPMA aggregate core.

Along with a lower α_N within the iPMA core, its value in the corona, α_N^{corona} , must be higher than 0.25 in order to assure an average α_N of 0.25. For example, if one quarter (or one third) of the COOH groups on the iPMA chain is incorporated in the core of the aggregate (with $\alpha_N^{core} = 0$), then α_N^{corona} for the remaining three quarters (or two thirds) of the chain in the corona, with the residual COOH and all COO⁻ groups, increases to $\alpha_N^{corona} \approx 0.34$ (or 0.38). It is therefore not surprising that the water exposed parts of iPMA chains swell substantially, because ions are usually strongly hydrated. Hydration (swelling) of the corona and simultaneous cooperative H-bonds within the core lead to very low ρ values for the iPMA ($\alpha_N = 0.25$) aggregates. The described situation (segregation of COOH and COO⁻ groups within the aggregate) is consistent with the tendency that the concentration of charges on partially ionized polyions is always larger at chain ends, a trend that may even be larger with the isotactic chains.

Similar findings were reported previously for poly(ethacrylic acid), PEA [58], where the strong H-bonding between suitably positioned COOH groups was even shown to prevent the ionization of the “core” COOH groups by NaOH. Tao et al. [59] have demonstrated by density functional theory calculations that an electron-donating (or electron-withdrawing) capability of the substituent R (= CH₃, C₂H₅, ...) bound next to COOH affects the acidity (and other molecular properties, particularly those related to hydrogen bonding) of monomeric carboxylic acids relative to the simplest analog, *i.e.* formic acid (H–COOH). The electron-donating groups (such as the hydrophobic CH₃ or C₂H₅ groups) enhance the electron density at the proton, which results in a shorter O–H bond and in lower acidity of the monomeric carboxylic acid, meaning that it is more difficult to detach the proton from COOH. In agreement with the findings for monomeric acids, it was experimentally observed in case of polymeric carboxylic acids that PMA and PEA (R = CH₃ and C₂H₅, respectively) are weaker acids than poly(acrylic acid), PAA (R = H) [6,8-10,16-18,58]. With polymeric

carboxylic acids, one has to take into account also the effect of multiple and adjacent acidic sites and the favorable (or unfavorable) microstructure of the chain (tacticity) on the acidity and the strength of eventual intermolecular H-bonds. Van den Bosch et al. [19] have for example argued that intermolecular association between iPMA chains through cooperative H-bonding is very strong and can only be disturbed by solvents (such as DMF and DMSO) capable of forming stronger H-bonds with COOH groups than water or COOH themselves. We conclude that this cooperativity in case of iPMA is the key to understanding its complex behavior in aqueous solutions. The interplay of three factors is crucial: (i) highly regular micro-structure of the polymer chain, (ii) the presence of (hydrophobic and electron-donating) CH₃ groups bound next to COOH groups, and (iii) the sequence and proximity of many functional sites.

The distribution of functional groups on the atactic chain is irregular, which prevents the favorable interplay of the above factors. In addition, aPMA ($\alpha_N = 0$) is almost un-charged. The compact interior of the aPMA aggregates at $\alpha_N = 0$ is considerably more nonpolar than the interior of the iPMA ones at $\alpha_N = 0.25$, as demonstrated by fluorescence measurements [27]. This is a result of more CH₃ groups residing in the interior of the aPMA aggregates (Scheme S1), which can relatively easily be incorporated within the core due to their irregular positions on the chain and due to more chain flexibility of the aPMA relative to the iPMA chain. The irregular distribution of COOH groups on aPMA also decreases the H-bonding strength. The cooperativity is absent and thus the core of the aPMA aggregates is less contracted, whereas the corona is less swollen in comparison with iPMA, as it contains only a few COO⁻ groups. Clearly, we cannot speak about segregation in this case. All this leads to higher ρ values in the aPMA case.

To get an additional confirmation of the proposed microgel-like structure of iPMA and aPMA aggregates, we have carried out a detailed analysis of the angular dependency of light

scattered by the aggregates. The particle scattering function, $P(q)$, was calculated and is plotted as $(qR_g)^2 P(\theta)$ versus qR_g in Fig. 2a (for iPMA at $\alpha_N = 0.25$) and Fig. 2b (for aPMA at $\alpha_N = 0$) for some selected c_s values. Solid lines in Figure 2 show the calculated particle scattering functions for some typical particle topologies [60] and points are the experimental data.

The data points in the aPMA ($\alpha_N = 0$) case extend well above $qR_g \approx 1.5$ (the small particle limit) and agree with the Debye-Bueche scattering function for most c_s values. Only for $c_s = 0.07$ M, where all size parameters in aPMA solutions display a pronounced maximum (*c.f.* Fig. 1), the deviation towards the Debye function is observed. The Debye function is generally used to describe the coil conformation, which is penetrable for the solvent. Mass distribution within the aPMA aggregates in solution with $c_s = 0.07$ M thus resembles that in solvent-drained polymer coils. These results show that the concentration of added NaCl regulates the compactness of the aggregates between aPMA chains with $\alpha_N = 0$.

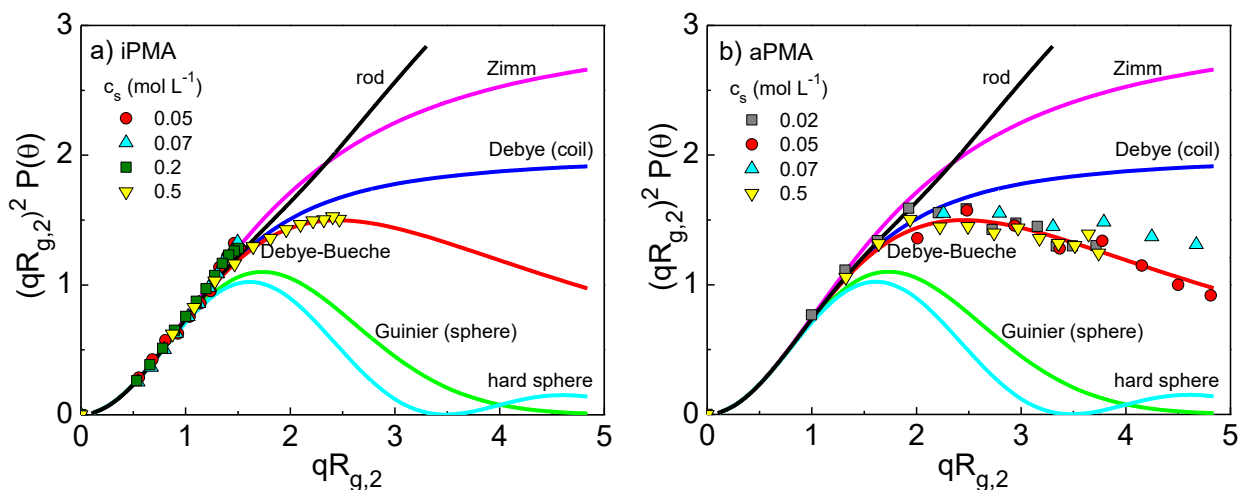


Fig. 2. The dependency of $(qR_g)^2 P(\theta)$ on qR_g for six selected particle topologies (solid lines) and the experimental data for intermolecular aggregates (index 2) in a) iPMA ($\alpha_N = 0.25$) and b) aPMA ($\alpha_N = 0$) solutions at various c_s values; polymer concentration is $c_p = 0.023$ M.

An unambiguous classification of the particle topology in iPMA solutions ($\alpha_N = 0.25$), such as with aPMA, is not so straightforward because the measured data points for most c_s values remain in the region of low qR_g values ($qR_g < 1.5$) where all topologies (except that for a hard sphere) agree. Only the data series for the highest c_s ($= 0.5$ M) extend to qR_g values close to 3 due to the largest size of the aggregates in this case (see $R_{h,2}$ and $R_{g,2}$ values in Table S1 and Fig. 1b). These data points clearly fit the Debye-Bueche function, so as those for aPMA for most c_s values (compare Fig. 2a and b). On the basis of these observations and of the low ρ values we conclude that both PMA isomers form similar aggregates in aqueous NaCl solutions at $T = 25$ °C as far as mass distribution is concerned. However, the particular distribution of functional groups is different, as discussed in detail above, and contributes to different degree of compaction on one hand and swelling on the other. As a result, the iPMA and aPMA aggregates differ considerably in the detailed dependence of size parameters on the ionic strength of the medium, regulated by c_s . The primary role of added NaCl is to screen electrostatic repulsions. This is important for iPMA ($\alpha_N = 0.25$) but not so much for aPMA, since its α_N is 0. However, the added Na^+ ions may be able to partly replace hydrogen atoms from the unionized COOH groups and thus increase the degree of ionization, α_i , of aPMA at $\alpha_N = 0$. In order to follow the effect of NaCl concentrations on α_i the pH of PMA solutions ($c_p = 0.023$ M) was measured in the studied c_s range. From pH, α_i was calculated using the relationship $\alpha_i = \alpha_N + \frac{[\text{H}^+] + [\text{OH}^-]}{c_p}$ (where $[\text{H}^+]$ and $[\text{OH}^-]$ are activities of hydronium and hydroxide ions, respectively), which follows from the electro-neutrality condition. The calculated α_i values are plotted in Fig. 3a. Evidently, the ionization of carboxyl groups in aPMA solutions with $\alpha_N = 0$ depends a great deal on c_s . It increases from around 1.6 % in 0.01 M NaCl to approximately 3 % in 0.5 M NaCl, *i.e.* by a factor of almost 2. On the contrary, α_i ($\cong 0.25$) is independent of c_s in iPMA solutions with $\alpha_N = 0.25$. The plot in Fig. 3a shows that the increase of α_i with c_s in aPMA ($\alpha_N = 0$) solutions is steeper for c_s values

below ≈ 0.08 M, a limit that agrees surprisingly well with the maximum in the LS parameters ($c_s \approx 0.07$ M; *c.f.* Fig. 1). Interestingly, the measured dn/dc values (*c.f.* Fig. 3b) also show a distinctive maximum in the same c_s range. We suggest that the increase in α_i (decrease in pH) of aPMA stems from a partial substitution of hydrogen atoms, which are covalently bound to the aPMA chain, with Na^+ ions. Through this, the aPMA chains acquire a higher charge which leads to increased repulsion and affects also the hydration state of the polymer chain due to more charged COO^- groups. Simultaneously with changed hydration, the contrast between scattering particles and the solvent medium (and thus dn/dc values) changes as well.

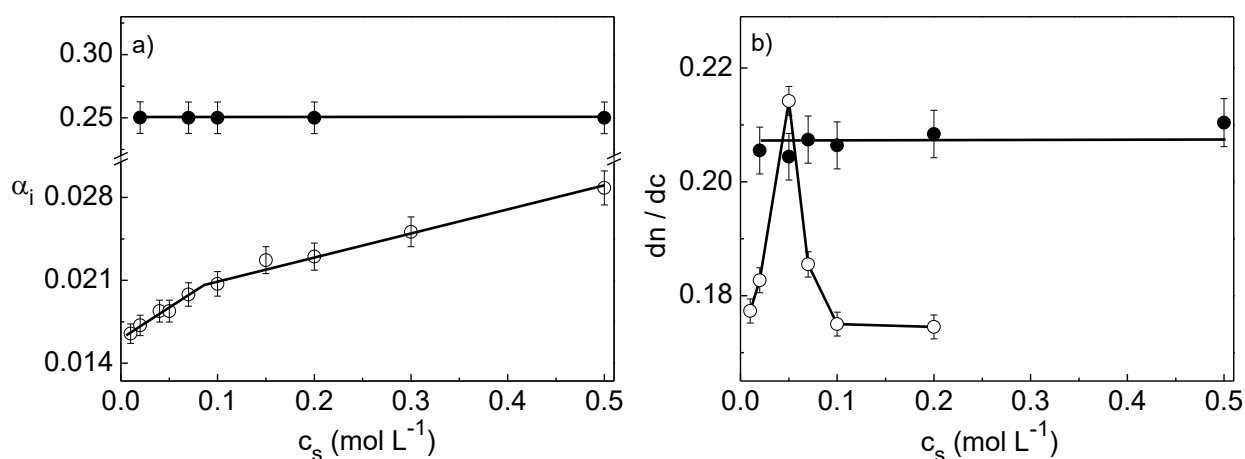


Fig. 3. a) The degree of ionization, α_i , and b) the dn/dc values in iPMA ($\alpha_N = 0.25$; full symbols) and aPMA ($\alpha_N = 0$; open symbols) solutions with $c_p = 0.023$ mol L⁻¹ in dependence on salt concentration, c_s .

Because ions are more strongly hydrated than their uncharged counterparts (vis. COO^- vs. COOH), more water enters into the physical network of the aPMA aggregates and particles swell, which causes some loosening of the whole structure and results in higher $R_{h,2}$, $R_{g,2}$, and also ρ values. For c_s up to 0.07 M, the hydration and water uptake work hand in hand. However, at $c_s \approx 0.07$ M their delicate balance is destroyed. The screening effect of salt on electrostatic repulsion between COO^- groups becomes more efficient, the repulsion decreases

and the associates can again contract in order to reduce the unfavorable contact of the hydrophobic CH₃ groups in the core with water. Simultaneously, the aggregates become less permeable for water, which is observed as a decrease in the size parameters.

On the other hand, the effect of NaCl on the ionization of COOH groups on iPMA chains at $\alpha_N = 0.25$ is negligible (*c.f.* Fig. 3a). This is in agreement with previous observations of the effect of other cations (*e.g.* cationic surfactants) on the ionization of several poly(alkylcarboxylic acids), where it was found that the binding of surfactant cations effects ionization of COOH groups only for $\alpha_N < 0.2$ [61]. We conclude that the delicate effect of NaCl on intermolecular association plays a role only at $\alpha_N = 0$ (for aPMA) and has a negligible effect in the case of partly ionized acid, as is the case with iPMA ($\alpha_N = 0.25$).

We followed also the effect of the polymer concentration on the aggregate structure of iPMA ($\alpha_N = 0.25$) and aPMA ($\alpha_N = 0.0$) and arrived at similar conclusions. The data for $c_p = 1.5, 2.0,$ and 2.4 g L^{-1} are collected in Table S2 and Figure S4 (SI). The main conclusion of those results is that the aggregates remain similar to microgel-like particles in the investigated c_p range. The data for aPMA show that the compactness of aggregates increases with increasing c_p (ρ decreases), whereas with iPMA c_p has no significant effect on ρ .

3.2 Chain dynamics in dependence on α_N and c_s

In the second part, iPMA and aPMA solutions were studied at higher α_N values ($\alpha_N \geq 0.4$ for iPMA and $\alpha_N \geq 0.25$ for aPMA), meaning that polyions are partly or completely ($\alpha_N = 1$) charged and aggregation is prevented. In this case, $G_2(t)$ is used to evaluate diffusion coefficients of polyions in order to study the fast and slow mode behavior of charged PMAs. The focus is on the PE-slow mode, which is a typical feature of highly charged polyions in conditions of low simple salt concentration when their mutual electrostatic interactions are not effectively screened [32,39,41]. Examples of the measured $G_2(q,t)$ functions for iPMA

and aPMA solutions with $\alpha_N = 1$, $c_p = 0.023 \text{ mol L}^{-1}$, and several c_s values are shown in Figure S5 (SI), together with the calculated τ -distributions in the insets.

As noted earlier [31], the LS intensity of salt-free (or low c_s) PE solutions is low due to the small osmotic compressibility. This is demonstrated for the herein studied iPMA and aPMA solutions in Figure S6 for several c_s values. The total LS intensity drops steeply with increasing polyion charge (increasing α_N), suggesting that the aggregates between chains disappear for $\alpha_N > 0.25$ (iPMA) or $\alpha_N > 0$ (aPMA) and only small particles exist in solution (see the R_h data in Tables S3 and S4). Nevertheless, just like in the low α_N case, the measured correlation functions at the chosen PMA concentration ($c_p = 0.023 \text{ mol L}^{-1}$) show two relaxation modes (see insets in Figure S5), *i.e.* two diffusion coefficients (two relaxation processes). The higher D value relates to the fast diffusion of single chain, described by D_f (index “f” for fast diffusion) and the lower one to the PE-slow mode feature, described by D_s (index “s” for slow diffusion). The PE-slow mode is not to be mistaken with the slow mode discussed in the previous section, which is related to the slow diffusion of real particles, in the present case to large inter-chain aggregates.

The PE-slow mode at low c_s and high α_N is most clearly detected in $G_2(t)$ for aPMA with the highest polyion charge (see for example the case with $c_s = 0.02 \text{ M}$ and $\alpha_N = 1$; Figure S5b). In the iPMA case, the PE-slow mode is present up to 0.1 M NaCl for all $\alpha_N \geq 0.4$, whereas for aPMA it is observed up to progressively higher c_s values as α_N increases. This will be in more detail discussed in relation to the slow diffusion coefficients, D_s .

Fig. 4 shows some Γ vs. q^2 curves for $\alpha_N = 1$ and different c_s values for both PMA isomers. The upper panels apply to the fast diffusion and the lower ones to the PE-slow diffusion mode. Clearly, Γ_f is a linear function of q^2 and goes through the center of coordinates, thus representing true translational diffusion individual polyion chains. The slope of these curves is proportional to D_f and gives their R_h (*c.f.* $R_{h,1}$ values in Tables S3 and S4). The size ($R_{h,1}$) of

single iPMA (aPMA) chains increases from around 9 (10) nm at $\alpha_N = 0.4$ (0.25) to 11 (20) nm for $\alpha_N \geq 0.5$. Higher $R_{h,1}$ values for $\alpha_N > 0$ are consistent with some chain expansion at higher polyion charge. However, the well-known conformational transition of PMA, which is usually observed in the range $0.2 < \alpha_N < 0.5$, was not studied in here.

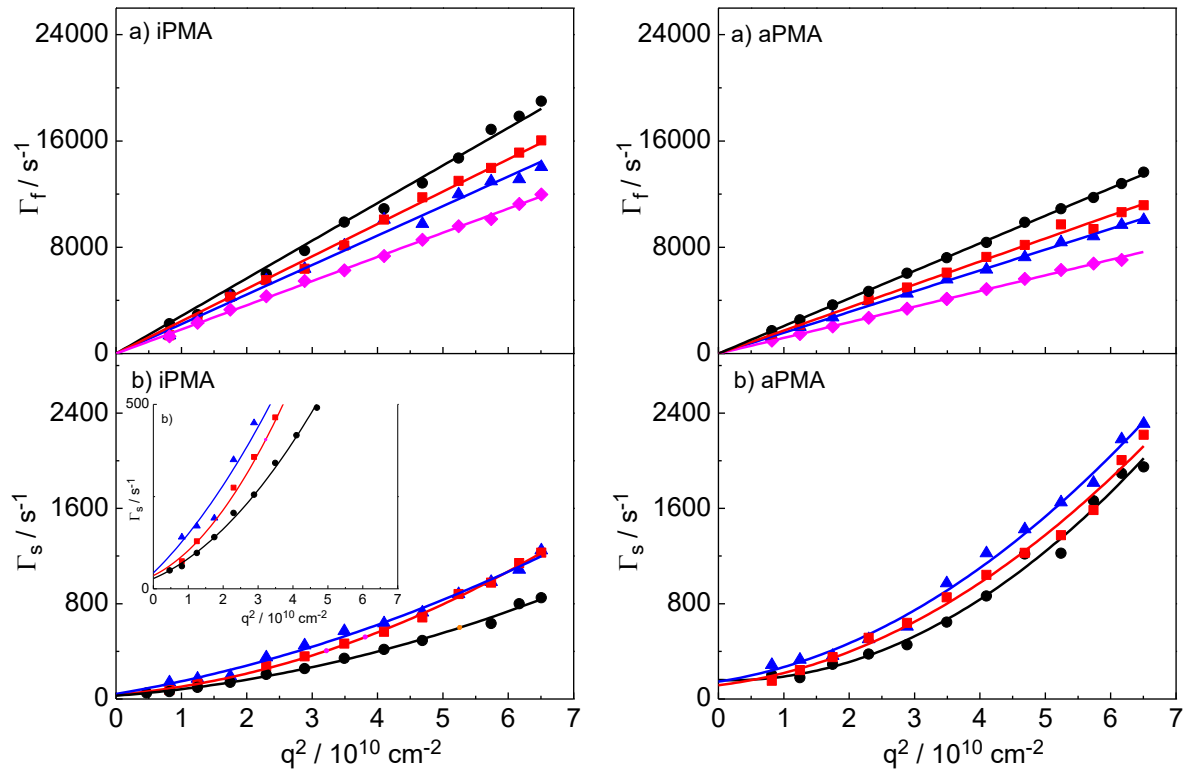


Fig. 4. Decay rates, Γ , as a function of the square of the scattering vector, q^2 , for $\alpha_N = 1$ and various salt concentration: (●) 0.05, (■) 0.07, (▲) 0.1, and (◆) 0.5 M NaCl; a) fast mode (Γ_f) and b) slow mode (Γ_s). For more details, the Γ_s vs. q^2 curves for iPMA are enlarged in the inset of Fig. 4b (*c.f.* left lower panel). Note that the slow mode is no longer observed in 0.5 M NaCl, therefore these Γ_s vs. q^2 curves are not presented in Fig. 4b.

The slow-mode relaxation rate curves (Γ_s vs. q^2) do not start from the origin of the coordinate system (*c.f.* lower panels in Fig. 4; compare those with lower panels in Figure S3 for the low α_N case), meaning that this mode is not originating from translational diffusion but owes to the polyelectrolyte effect [45]. This presumption follows also from the fact that

the total LS intensity is very low (*c.f.* Figure S6 and the discussion above) and that the angular dependency of LS intensity gives physically meaningless results (not shown). The initial slope of the I_s vs. q^2 curves is therefore used simply to evaluate the value of D_s .

The combined slow and fast mode behaviors for both PMAs are shown in Fig. 5 as D vs. c_s plots. The D_f values (single chains) span from $\alpha_N = 0$ to $\alpha_N = 1$, whereas the D_s ones are limited to higher α_N , *i.e.* to the region between $\alpha_N = 0.25$ (aPMA) or 0.4 (iPMA) and $\alpha_N = 1$, clearly because the slow mode is observed only at sufficiently high polyion charge. The D_s values are more than one order of magnitude lower than the D_f ones. Besides, their dependence on c_s is different. D_f initially steeply decreases with increasing c_s and remains more or less constant for higher c_s , with the exception of the aPMA ($\alpha_N = 0$) case, where it is practically constant (note that in this case, the D_s was not detected). Contrary to D_f , the D_s values increase with increasing c_s for both PMAs and disappear at high enough salt concentrations. Interestingly, the c_s limit where this takes place in iPMA solutions is the same for all α_N , *i.e.* $c_s > 0.1$ M, while in aPMA solutions it clearly depends on α_N . In aPMA solutions with $\alpha_N = 1$, D_s disappears for $c_s > 0.1$ M, for $\alpha_N = 0.75$ and 0.5 the limit is the same ($c_s > 0.07$ M) and for $\alpha_N = 0.25$ D_s is no longer detected for $c_s > 0.05$ M NaCl. Above the quoted c_s values, only the fast diffusion coefficient, D_f , is observed. For aPMA, this c_s dependency of D_s on α_N is expected and is due to the lower polyion charge and consequently to weaker electrostatic interactions at lower α_N . Based on these results, we suggest that the charge of the aPMA chains changes continuously with α_N , *i.e.* no segregation of ionized and unionized groups is anticipated. By decreasing α_N , the overall (effective) charge of the polyion decreases and this leads to weaker electrostatic interactions, which are responsible for the PE-slow mode.

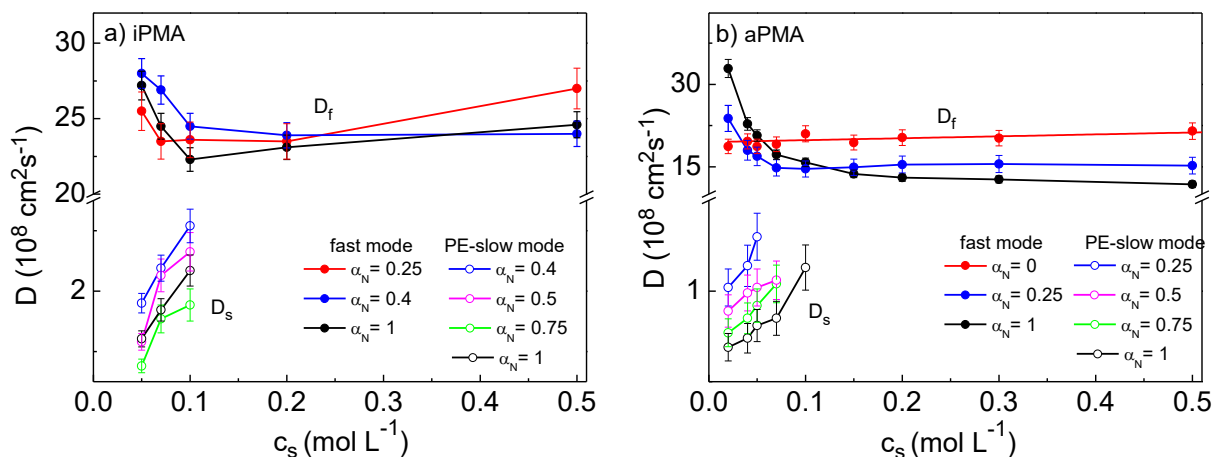


Fig. 5. The slow mode, D_s , and fast mode diffusion coefficients, D_f , in dependence on c_s for various α_N : a) iPMA and b) aPMA, both with $c_p = 0.023 \text{ mol L}^{-1}$. In order to clearly present the D_s values, the y-axis contains a brake and is enlarged in the region $D_s \approx (1-3) \times 10^{-8} \text{ cm}^2 \text{ s}^{-1}$.

A question arises why the value of c_s , where the PE-slow mode (D_s) disappears in iPMA solutions, is independent of α_N , in obvious contrast to aPMA. We propose that this is because of the isotactic chain configuration and features associate with it, as discussed already above in relation to inter-chain aggregation. Let's propose that the segregation of COOH and COO⁻ groups, as proposed above for the low α_N case, persists also at higher α_N where the aggregates have already decomposed. This can be imagined in a way that portions of iPMA chains contain only the ionized COO⁻ (the situation corresponding to a local α_N value $\alpha_N^{loc} \approx 1$) and others predominately the unionized COOH groups (the situation corresponding to $\alpha_N^{loc} = 0$). As mentioned above, this seems realistic as the concentration of charges is always larger at chain ends; besides, isotacticity may additionally contribute to this tendency. The repulsion between the charged portions of the chains would then be approximately the same, irrespective of their average charge (average α_N value). Because the main factor that leads to the PE-slow mode and the related D_s is the polyion charge, this phenomenon in iPMA

solutions is detected up to the same c_s value at all α_N . This situation resembles the case of cooperative binding of charged surfactants to oppositely polyelectrolytes [61]. The surfactant micelles, which cooperatively form at the polyion, block charges on occupied portions of the chain, but leave the rest of the polyion free and with an unchanged charge density. The charge density is the property that controls various features of PEs, such as counterion binding and also the PE-slow mode (or D_s), which thus becomes independent of α_N .

The dependency of D_s on α_N at a constant c_s value of aPMA solutions shows that D_s increases as the polyion charge (α_N) decreases. This also makes sense, because a lower charge means weaker electrostatic interactions and less pronounced slow mode. For iPMA, however, the $\alpha_N = 0.75$ and 1.0 cases are interchanged: D_s at $\alpha_N = 1$ is higher than D_s at $\alpha_N = 0.75$. To clarify this observation, more systematic investigations are necessary on the α_N dependency of the PE-slow mode.

4. Conclusions

A systematic LS analysis of iPMA and aPMA in aqueous solutions with added NaCl is performed in order to investigate intermolecular association, limited to low α_N values, and the polyelectrolyte slow mode, observed at high α_N values. These two phenomena are studied in dependence on the ionic strength of the solutions and chain tacticity at a constant polyion concentration ($c_p = 2 \text{ g L}^{-1}$ at $\alpha_N = 0$, which corresponds to 0.023 moles of COOH groups per volume). The distinction between the two phenomena, aggregation and PE-slow mode, is made on the basis of (i) the total LS intensity, which drops dramatically upon increasing α_N above 0 (for aPMA) or above 0.25 (for iPMA) as a result of the disappearance of the aggregates, and (ii) physically meaningless angular dependency of the LS intensity of the corresponding PE-slow diffusive mode.

In the case of intermolecular association/aggregation, the geometrical parameters for the small scatterers (single chains) and for large aggregates are determined for iPMA at $\alpha_N = 0.25$

and for aPMA at $\alpha_N = 0$. While the size of the smaller particles (individual chains) is more or less independent of the ionic strength, the size and shape of the aggregates depends strongly on the salt (NaCl) concentration. For iPMA, the size of the aggregates gradually increases with increasing c_s , whereas aPMA shows quite a peculiar dependence of the aggregate size on c_s : the R_h (and R_g) values initially steeply increases with increasing ionic strength, reach a maximum and thereafter gradually decreases. Thus, the aggregates swell at a particular ionic strength. This result may be relevant for protein solutions, where interaction with water (swelling) controls protein function. From the calculated ρ values, it is concluded that aPMA and iPMA aggregates show characteristics of microgel-like particles with a core-shell structure, in which the core is much denser in comparison with the shell. However, the precise distribution of the polar (hydrophilic) and nonpolar (hydrophobic) functional groups and the associated compactness of the aggregates differs considerably for both isomers. It is suggested that the charged and uncharged carboxyl groups segregate in the iPMA case, which is the basis for very strong and cooperative formation of H-bonds between uncharged chain portions. This is suggested for iPMA in our study for the first time and may be the underlying reason for so-called irreversibility of the conformational transition known for iPMA, which was observed years ago.

Different chemical composition of the aggregate interior (and at the same time aggregate surface) dictates the affinity of PMA isomers for the solvent, *i.e.* water. Compact core of iPMA is proposed to consist of un-ionized COOH groups, which are strongly and cooperatively H-bonded, leaving the hydrophobic CH₃ groups exposed to water and thus preventing iPMA to dissolve for $\alpha_N < 0.2$. In the aPMA case, cooperativity in H-bonding is absent and an opposite distribution of polar and nonpolar groups is proposed: the methyl groups are assembled in the interior of the aggregate and the hydrophilic COOH groups are

mostly oriented towards the polar aqueous environment, thus granting solubility of aPMA already at $\alpha_N = 0$.

At higher α_N values ($\alpha_N \geq 0.4$ for iPMA and $\alpha_N \geq 0.25$ for aPMA), where intermolecular association ceases, the electrostatic forces become operative and the PE-slow diffusive mode is detected. In iPMA solutions, the PE-slow mode is present at all $\alpha_N \geq 0.4$ up to $c_s = 0.1$ M, whereas in aPMA solutions this phenomena is most pronounced for the highest polyion charge ($\alpha_N = 1$), namely up to 0.1 M NaCl, whereas for lower α_N (≤ 0.75) it is seen up to gradually lower c_s values, which is to be expected, the PE-slow mode is to a large extent dependent on the polyion charge: at lower charge (lower α_N) less salt is needed to suppress electrostatic interactions. In iPMA solutions, on the other hand, the value of c_s where the PE-slow mode disappears is surprisingly independent of α_N . This remarkable result is again attributed to the isotactic chain configuration. It is presumed that the electrostatic interactions between charged chain parts, which are responsible for the PE-slow mode, may become independent of c_s , if the segregation of charged and uncharged carboxyl groups is allowed also at higher α_N .

To conclude, isotacticity, which is the reason for the segregation of charged and uncharged carboxyl groups within the iPMA chains, has a far-reaching effect on solution behavior of iPMA in the whole α_N region. The segregation is operative over the entire α_N region and affects both, aggregate formation involving exceptionally strong and cooperative H-bonding between COOH groups and the polyelectrolyte slow diffusion mode, which is demonstrated by independence of the polyion diffusion coefficients on ionic strength.

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Appendix A. Supplementary material

Detailed aspects of the data analysis, examples of data treatment, and additional experimental results are given in supporting information, SI.

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