

RESEARCH ARTICLE

View Article Online
View Journal | View IssueCite this: *Mater. Chem. Front.*,
2020, 4, 905pH-Dependent morphology and optical properties
of lysine-derived molecular biodynamers†Sangeun Lee, ^{ab} Cansu Kaya, ^{ac} Hongje Jang, ^d Marcus Koch, ^e Brigitta Loretz, ^b
Eric Buhler, ^f Claus-Michael Lehr*^{bc} and Anna Katharina Herta Hirsch ^{ac}

Polymerization of carbazole dicarboxaldehydes and lysine derivatives by imine and acylhydrazone formation afforded peptide-derived molecular biodynamers. Characterization of their physicochemical properties revealed an interesting morphology change upon polymerization from monomers forming submicrometer spherical micelles to nanometer-sized rigid-rod-shaped polymeric particles. A combination of light-scattering methods, small-angle neutron scattering, and transmission electron microscopy enabled a detailed investigation of this morphological change. Moreover, we investigated by dynamic and static light scattering how the pH affects the fluorescence and size of the biodynamers. These morphological and pH-dependent changes are expected to open the door to a myriad of applications of molecular biodynamers.

Received 22nd October 2019,
Accepted 23rd December 2019

DOI: 10.1039/c9qm00651f

rsc.li/frontiers-materials

Molecular biodynamers combine the advantages of biopolymers and dynamic covalent chemistry.¹ Monomers are connected through covalent bonds that are reversible under specific conditions such as different pH and temperature. The resulting materials are stimuli-responsive systems and self-healing.^{2,3} Therefore, the polymerization and degradation of the molecular biodynamers are reversible depending on conditions. Moreover, when the biodynamers mimic biological biopolymers such as nucleotides, peptides, and polysaccharides, these molecular biodynamers are expected to feature advantages inherent to biopolymers (biocompatibility, biodegradability, and biofunctionality).^{4,5}

In a previous study, we reported a biodynamer composed of carbazole dicarboxaldehyde **1** and lysine hydrazide **2** (Fig. 1A).⁶ Acylhydrazone and imine bond formation between hexaethylene glycol conjugated carbazole dicarboxaldehyde and lysine hydrazide, both of which are water-soluble, afforded proteoid biodynamers at slightly acidic conditions. Interestingly, the

resulting biodynamers form rigid nanorods in aqueous solution due to numerous π - π -stacking interactions. A number of features of each monomer make this biodynamer particularly attractive for various applications. The carbazole is a fluorophore, thus, its derivatives are widely used for bioimaging and sensors.^{7,8} The other monomer L-lysine hydrazide, is a derivative of an amino acid, which is used as a building block for pharmaceutical ingredients, as food additive, and cosmetics. Especially, polymerized L-lysines (poly-L-lysine) are versatile biomedical materials and are widely used for tissue culture and drug delivery.⁹ In this study, we have evaluated the physicochemical properties of the biodynamers, and examined pH-dependent changes in view of further application of the polymer.

We set out to investigate the morphological and optical changes upon polymerization. To do so, we synthesized each monomer following the reported methods (Fig. S1 and S2, ESI†).^{10,11} Each of the monomers was dissolved in 100 mM acetate buffer with a final concentration of 10 mM for biodynamer synthesis. After incubation at r.t. for 19 hours, we observed successful polymerization confirmed by following the disappearance of the aldehyde protons by ¹H-NMR spectroscopy (Fig. S2, ESI†). After filtration, we monitored the chemical, optical, and morphological changes.

Using dynamic light scattering (DLS) and transmission electron microscopy (TEM), we examined the size and morphology changes upon polymerization (Fig. 1B). The dialdehyde monomer **1** itself formed spherical micelles ($D_h = 255$ nm; polydispersity index (PDI): 0.077) (Fig. 1C) in aqueous solution (Fig. 1D, right). This micelle formation *via* self-assembly is driven by hydrophobic and π - π -stacking interactions between the monomers and stabilized in the aqueous solution by the hydrophilic shells

^a Department for Drug Design and Optimization, Helmholtz Institute of Pharmaceutical Research Saarland (HIPS) – Helmholtz Centre for Infection Research (HZI), Campus E 8.1, 66123 Saarbrücken, Germany.
E-mail: Anna.Hirsch@helmholtz-hips.de

^b Department for Drug Delivery, HIPS – HZI, Campus E8.1, 66123 Saarbrücken, Germany. E-mail: Claus-Michael.Lehr@helmholtz-hips.de

^c Department of Pharmacy, Saarland University, 66123, Saarbrücken, Germany

^d Third Institute of Physics, Georg-August-Universität, Friedrich-Hund-Platz 1, 37077 Göttingen, Germany

^e INM-Leibniz-Institute for New Materials, Campus D2.2, Saarland University, 66123 Saarbrücken, Germany

^f Laboratoire Matière et Systèmes Complexes (MSC), UMR 7057, University of Paris (Paris Diderot), Bâtiment Condorcet, 75205 Paris Cedex 13, France

† Electronic supplementary information (ESI) available. See DOI: 10.1039/c9qm00651f

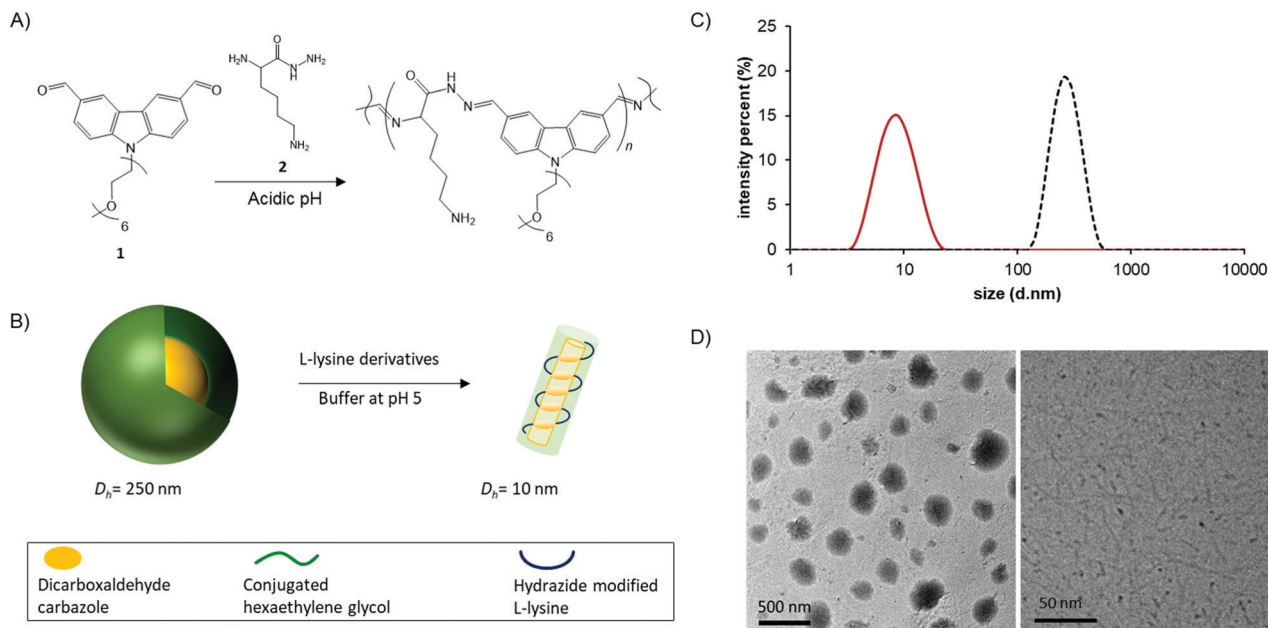


Fig. 1 Chemical structure of the monomers **1** and **2** and the corresponding molecular biodynamer (A) and illustration of carbazole-based dialdehyde **1** micelles ($D_h = 250$ nm) transferring to biodynamer ($D_h = 10$ nm) (B). Size of the carbazole micelle (dotted line, $Z_{ave} = 255$ nm, polydispersity index (PDI) = 0.077) and biodyn timers (solid line, $Z_{ave} = 8$ nm, PDI = 0.099) obtained by DLS (C). TEM image of carbazole-based dialdehyde **1** micelles (D, left) and cryo-TEM image of biodyn timers (D, right).

of the hexaethylene glycol substituents. Upon addition of hydrazide **2**, the nanoparticle size decreased to 8 nm (D_h , PDI: 0.099) (Fig. 1C) as a result of the polymerization, affording biodyn timers, in line with the nanorods we had previously observed. In the cryo-TEM image of the biodyn timers (Fig. 1D, right), the polymer has a rod-like shape, similar to reported supramolecular polymers.¹² The shown morphology also corresponds well with our previously reported biodyn timers.^{6,13} Note that, the darker parts of the nanorods in the images are the results of the increased density due to the vertically oriented nanorods.^{14,15} However, the cryo-TEM image gives limited information on its morphology. Therefore, the morphology, such as the shape of the section, the stacking, and the local structures were further investigated.

Solution-state polymer morphologies were investigated in depth using complementary small-angle neutron (SANS, beamline D11 at ILL)¹⁶ and static light scattering (SLS) experiments. A significant advantage of this combination is its ability to provide information on characteristic sizes, local structure and overall shape of objects in solution across a broad range of length scales varying between ~ 1 and 300 nm. Fig. 2 shows the variation of the product $M_w P(q)$ as a function of the scattering vector q for a 10 mg mL⁻¹ solution at pD 7.4, where M_w and $P(q)$ are respectively the weight-averaged molecular weight and the form factor of the objects. The SLS and SANS scattered intensities are here normalized by their respective contrast terms and concentration, allowing a representation of $M_w P(q)$ without an arbitrary shift to scale SLS data on SANS data (see Methods in ESI†). The formation of cylindrical structures identified using cryo-TEM is in agreement with the overall behavior observed here. The following sequence describes the behavior: (i) a low- q plateau in the SLS q -range associated with

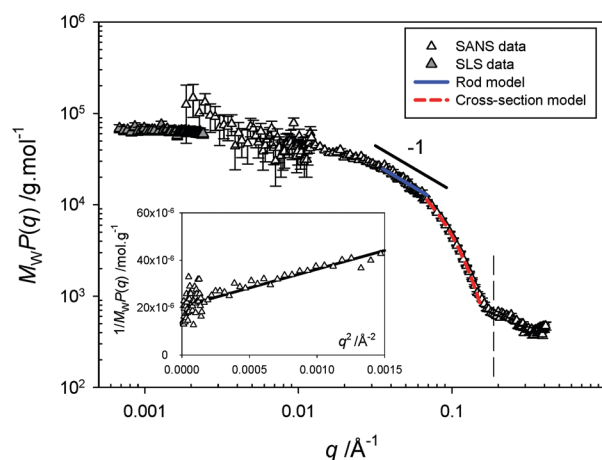


Fig. 2 Scattering profile obtained from combined static light (SLS) and small-angle neutron scattering (SANS) experiments for a 10 mg mL⁻¹ solution at pD 7.4. The product of the weight-averaged molecular weight and form factor, $M_w P(q)$, is obtained by normalization of the scattered intensity by concentration and contrast term of the technique (see ESI†), allowing a rescaling of the SLS and SANS data without an arbitrary shift. The continuous and dashed lines represent the best fit of the data in the intermediate q ⁻¹ regime by a rod model and in the high- q range by a Guinier expression for the form factor of the cross-section, respectively. The inset represents the variation of $1/M_w P(q)$ with q^2 in the low- q SANS domain.

the finite mass of the polymers and allowing a direct determination of M_w by extrapolation to zero- q , followed by (ii) a Guinier regime associated to the finite size (radius of gyration R_g) of the objects, (iii) a small intermediate rod-like regime, in which the variation of the signal can be described by a power-law with an exponent close to -1 , (iv) another Guinier regime at

the upper q -range, corresponding to the cross-section of the cylinders, and finally (v) an oscillation associated with the shape-dependent form factor of the section of the cylinders.

The data obtained at low q corresponding to large spatial scales have been fitted by a classical Guinier law (see eqn (S5) in ESI†). From the best linear fit to the curve representing $1/M_W P(q)$ versus q^2 (see inset of Fig. 2), one obtains a radius of gyration equal to $R_g = 54.4 \text{ \AA}$ giving a relatively short contour length of $L = \sqrt{12} \times R_g = 188 \text{ \AA}$ for the rods and explaining the narrowness of the q^{-1} regime. In the intermediate q regime, the scattering profile can be fitted satisfactorily using a rigid-rod model using the form factor derived for rod-shaped particles, $P(q) = \pi/qL$. The high q data can be fitted by a Guinier expression for the form factor of the section, $V_p P(q) = \pi a_c/q \times \exp(-q^2 r_c^2/2)$. By fitting the two equations above to the experimental data, one can determine the linear mass density (μ), the cross-sectional area (a_c), and the R_g of the section of the cylinders (r_c). From the fits of Fig. 2, we obtain $\mu = 279.4 \text{ g mol}^{-1} \text{ \AA}^{-1}$, $a_c = 444 \text{ \AA}^2$, and $r_c = 14 \text{ \AA}$. The contour length of the cylinders can also be obtained from the value of M_W and μ : $L = M_W/\mu = 253 \text{ \AA}$, in a relatively good agreement with that determined directly from the R_g .

The cylindrical morphology is also confirmed by the so-called ρ ratio derived from the combination of SANS and DLS. The correlation function characterized by a single diffuse relaxation (*i.e.*, single population, see ESI†, Fig. S3) allows determination of the hydrodynamic radius R_h of the objects. Therefore, the ratio showing a value, $\rho = R_g/R_h = 1.52$, larger than one is a characteristic of elongated objects such as cylinders. While R_g , R_h , and L give information on the overall structure of the cylinders, the experimental values of μ , a_c , and r_c provide important indications on the local particle topology and molecular stacking. Assuming a full disk-shaped cross-section with $a_c = \pi R^2$ leads to conflicting results as one obtains a geometrical radius $R = 11.9 \text{ \AA}$ for the disk smaller than $r_c = 14 \text{ \AA}$. A possible explanation is the formation of a core-shell structure with a cross-sectional area given by $a_c = \pi(R_{\text{ext}}^2 - R_{\text{int}}^2)$ and $r_c = \sqrt{2/2} \times \sqrt{(R_{\text{ext}}^2 + R_{\text{int}}^2)}$, where R_{ext} and R_{int} are the external and internal radii of the section of the cylinders, respectively. Using the experimental values for a_c and r_c , one finds $R_{\text{ext}} = 16.3 \text{ \AA}$ and $R_{\text{int}} = 11.2 \text{ \AA}$; the thickness of such a shell section being equal to $R_{\text{ext}} - R_{\text{int}} = 5.1 \text{ \AA}$. Such a structure is compatible with the stacking of the carbazole moieties in the core and hexaethylene glycol as a shell in agreement with the linear mass density value of $279.4 \text{ g mol}^{-1} \text{ \AA}^{-1}$.

Lastly, we observed optical changes of the biodynamers by polymerization. The carbazole monomer **1** has shown absorption and emission at $\lambda_{\text{abs}}/\lambda_{\text{em}} = 350/470 \text{ nm}$ (Fig. S5, ESI†). By adding the lysine monomers, the absorption around 425 nm increased. Due to the changes in absorption, the observed color of the polymer solution changed from yellow to orange over 24 hours (Fig. 3A). Besides, the intensity of the emission spectrum of the polymeric solution decreased in the first 4 hours of polymerization. The decrease is presumably ascribed to a substantially reduced distance between the carbazoles upon polymerization, resulting in quenching. However, after 10 hours of polymerization,

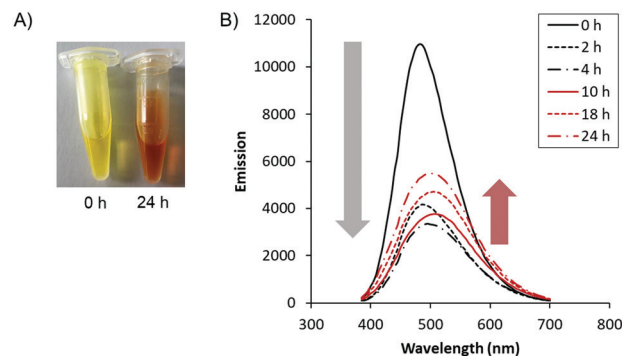


Fig. 3 Optical changes by molecular biodynamer formation. (A) The colorimetric changes of monomer mixture by reaction time (left: 0 hours, right: 24 hours at r.t.) (B) emission changes as a function of reaction time. The peak top shifted from 485 nm to 505 nm.

the emission intensity slightly increased. A possible explanation is that the initially randomly stacked and quenched carbazole moieties reorganized during nanorod formation. This rearrangement and nanorod formation possibly increase the distance between the carbazole moieties by cation- π interaction between the carbazole and lysine and by steric repulsion of hexaethylene glycols, both leading to a slight recovery of the quenched fluorescence.

Having confirmed the polymerization, we evaluated the pH effect on the optical properties of the polymer. We prepared the polymer solution (1 mg mL^{-1}) at pH 3 to 13 (pH 3, 5, 6, 7.4, 9, 11, 13) and measured the absorption and emission of the polymeric solutions at different pH values (Fig. 4A). As shown in the spectra, λ_{max} of the emissions shifts from 470 nm at pH 3 to 535 nm at pH 13 (Fig. 4B), corresponding to a gradual redshift as the pH increase. Interestingly, each of the emission peaks shifts by a ratio change between two different emission peaks of $\lambda_{\text{max}} = 470 \text{ nm}$ and $\lambda_{\text{max}} = 535 \text{ nm}$. It is more obvious at pH 9 and 11 that the peaks have two peak tops composed of two emission peaks. It suggests that a ratio change of the two components in the polymer affects the fluorescence. Considering the peak shifts are pH-dependent, the proton concentration affects the fluorophores, *i.e.*, the carbazole groups, in the polymer. We speculate that protonation of the tertiary amine of the carbazole induced the emission peak shifts. The protonation of the tertiary amine shortens the conjugated system of the fluorophore. On the other hand, the unprotonated form has a longer conjugated system, resulting in an increased redshift. Another explanation is that pH-dependent protonation on the lysine side chains affects the electron conjugation on the fluorophore by photo-induced electron transfer (PET). However, even if the fluorophore can behave either as an electron acceptor (a-PET) or a donor (d-PET) towards the side chains, the PET affects the emission intensity by pH but not the emission shifts. Moreover, considering the pK_a of the ϵ -lysine side chain, the gradual peak shift between pH 3 to 13 is not explainable by side chain protonation. (Note that, the pK_a value of the ϵ -lysine side chain is 10.53). Therefore, we believe that the blueshift under acidic conditions is a result of protonation on the backbone.

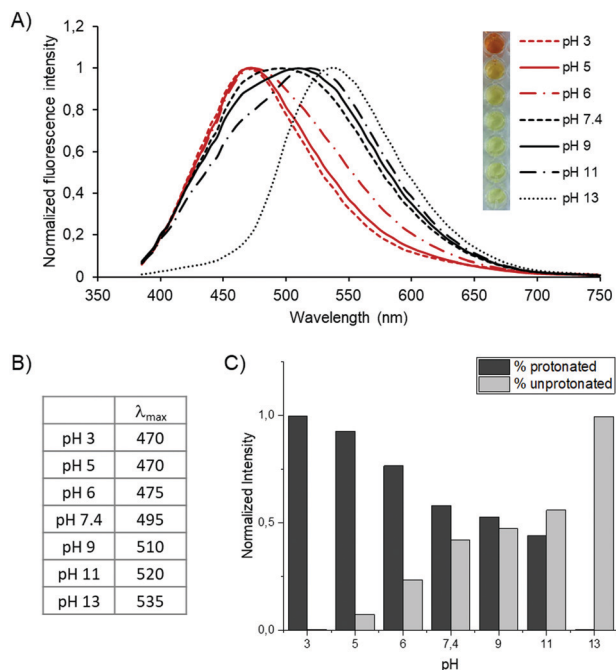


Fig. 4 pH-dependent optical changes (A) emission shift of the biodynamers as a function of pH. The concentration of the polymer is set to $100 \mu\text{g mL}^{-1}$. 100 mM Acetate solution was used for pH 3, and pH 5. 100 mM Phosphate solution was used for pH 7.4 to pH 13. The colors of each solution are shown on the left. (B) Table of the λ_{max} obtained from the emission spectrum. (C) The intensity ratio of two fluorescence components at various pH. The ratio is calculated based on the fitting process using two model curves.

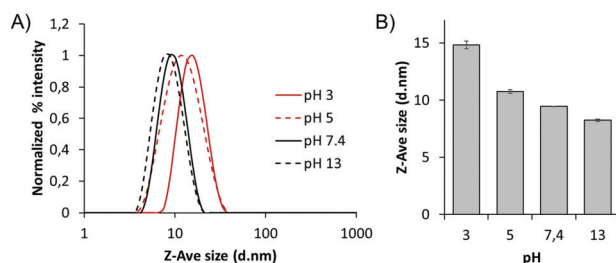


Fig. 5 pH-dependent size changes (A) normalized percent intensity of the size of the biodynamers (1 mg mL^{-1}) as a function of pH. (B) Zeta average (Z-Ave) size difference of the biodymer (1 mg mL^{-1}) at pH 3, 5, 7.4, and 13. Each size value is the mean value of DLS measurements performed in triplicate.

To evaluate the ratio of these two components (protonated/unprotonated) at different pH, we calculated it with a

fluorescence peak deconvolution method (Fig. 4C). The two spectra at pH 3 and 13 are utilized as model spectra of the protonated and deprotonated states of the molecule. Each spectrum was composed of two Gaussian curves to fit the curves precisely. The height and width ratio of the Gaussian functions of the model spectrum was fixed to keep the shape of each basic spectrum. By a linear combination of the two model spectra of pH 3 and 13, the protonation ratio at other pH conditions was calculated from each spectrum. Consequently, we revealed clearly that the emission spectrum between pH 5 and 11 is composed of the summation of the two primary signals. The ratio of unprotonated carbazole gradually increases when increasing the pH and eventually is reversed between pH 9 and 11. Note that we obtained the ratio based on the assumption that the emission peak at pH 3 is purely protonated, and it is an entirely unprotonated spectrum at pH 13. Therefore, further investigation is required to quantify absolute protonation values.

We expected that this pH-dependent protonation also affects the size of the polymeric nanorods. To verify it, we measured the size of the polymer at pH 3, 5, 7.4, and 13 by DLS using the cumulant method, based on the backscattering at 173° (Fig. 5A and B). As a result, the size of the nanorods was larger in a more acidic environment. The size of the nanorods at pH 3 was 14.5 nm, while it was 8.8 nm at pH 13. This means that the hydrodynamic radius of the biodymer increased by up to 65% upon increasing the proton concentration 10^{10} times. However, measurement using a single scattering angle may not be sufficient to support these changes and also does not allow a mass determination.

Therefore, we have also confirmed the pH-dependent size and mass differences using combined DLS and SLS with scattering angles from 30° to 140° . From the SLS data, the M_w of the polymer has shown a negligible difference between pH 5 and pH 7.4 (Table 1). At pH 5, the measured molecular weight of the polymer is $67\,939 \text{ g mol}^{-1}$, and the polymerization degree derived from the M_w is about 109. At pH 7.4, the M_w is $66\,552 \text{ g mol}^{-1}$, and the degree of polymerization is 106, which is not significantly different compared to the values obtained at pH 5. In both of the calculation methods, however, the size of the polymer at pH 5 is 35% (by the Contin method, and 31% by the cumulant method) larger, indicating a change within the structure and the compactness of the objects. In other words, the nanorod size changed as a function of pH despite the molecular weight remaining consistent. A possible explanation is that the increment in electrostatic repulsion upon protonation on the polymer backbone affects the packing of the nanostructure.

Table 1 Structural parameters of the biodymer obtained by SLS, DLS and SANS measurements

Buffer acidity	c	ϕ	M_w	DP	R_g	R_h^a	R_h^b	k_2/k_1^2	$\rho = R_g/R_h$	r_c	a_c	μ
pD 7.4	10	6.83×10^{-3}	70 570	113	54.4	39.4	35.7	0.095	1.52	14	444	279.4
pH 7.4	1	6.83×10^{-4}	66 552	106	—	39.8	32.1	0.1153	—	—	—	—
pH 5	1	6.83×10^{-4}	67 939	109	—	61.5	47	0.1434	—	—	—	—

c = polymer concentration (mg mL^{-1}), $\phi = c/d$ volume fraction (monomer density $d = 1.46 \text{ g cm}^{-3}$), M_w = weight-averaged molecular weight (g mol^{-1}), DP = degree of polymerization, R_g = radius of gyration (\AA), R_h = hydrodynamic radius obtained using the Contin^a and the cumulant^b analysis (\AA), k_2/k_1^2 = polydispersity index, r_c = cross-sectional radius of gyration (\AA), a_c = cross-sectional area (\AA^2), μ = linear mass density of cylinders ($\text{g mol}^{-1} \text{\AA}^{-1}$). (SLS, DLS and SANS measurements determined the values with $\sim 10\%$ error, see ESI.)

In this study, we have investigated molecular biodynamers composed of carbazole and lysine derivatives. We evaluated morphological, optical-property, and size changes of the biodynamers upon polymerization under acidic conditions. The spherical micelles of the carbazole monomer transformed into rigid nanorods in the course of the polymerization, as shown through a thorough analysis by SANS and SLS. pH-dependent changes in fluorescence emission and size of the biodynamers are expected to broaden the further application of the molecular biodynamers.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

Authors deeply acknowledge the Institut Laue Langevin (ILL Grenoble, France) for beamtime allocation as well as G. Mariani and S. Prevost for their precious help during the SANS experiments, and Jörg Schmauch at Saarland University, Experimentalphysik INM-Leibniz-Institute for New Materials for the TEM experiments. Funding from the Helmholtz Association's Initiative and Networking Fund (A. K. H. H.) and the SFB937 Collective behavior of soft and biological matter (A11, H. J.) is gratefully acknowledged. Dedicated to J.-M. Lehn on the occasion of his 80th birthday.

References

- 1 E. Kolomiets and J. M. Lehn, Double dynamers: molecular and supramolecular double dynamic polymers, *Chem. Commun.*, 2005, 1519–1521.
- 2 H. Ying, Y. Zhang and J. Cheng, Dynamic urea bond for the design of reversible and self-healing polymers, *Nat. Commun.*, 2014, 5, 3218.
- 3 F. Garcia and M. M. Smulders, Dynamic covalent polymers, *J. Polym. Sci., Part A: Polym. Chem.*, 2016, 54(22), 3551–3577.
- 4 Y. Liu, M. C. A. Stuart, E. Buhler and A. K. H. Hirsch, Dynamic Proteoids Generated From Dipeptide-Based Monomers, *Macromol. Rapid Commun.*, 2018, 39(13), 1800099.
- 5 Y. Liu, M. C. A. Stuart, M. D. Witte, E. Buhler and A. K. H. Hirsch, Saccharide-Containing Dynamic Proteoids, *Chem. – Eur. J.*, 2017, 23(64), 16162–16166.
- 6 Y. Liu, M. C. A. Stuart, E. Buhler, J.-M. Lehn and A. K. H. Hirsch, Proteoid Dynamers with Tunable Properties, *Adv. Funct. Mater.*, 2016, 26(34), 6297–6305.
- 7 Y. Liu, J. M. Lehn and A. K. Hirsch, Molecular Biodynamers: Dynamic Covalent Analogues of Biopolymers, *Acc. Chem. Res.*, 2017, 50(2), 376–386.
- 8 X. J. Feng, P. L. Wu, F. Bolze, H. W. C. Leung, K. F. Li, N. K. Mak, D. W. J. Kwong, J.-F. Nicoud, K. W. Cheah and M. S. Wong, Cyanines as New Fluorescent Probes for DNA Detection and Two-Photon Excited Bioimaging, *Org. Lett.*, 2010, 12(10), 2194–2197.
- 9 I. L. Shih, Y. T. Van and M. H. Shen, Biomedical applications of chemically and microbiologically synthesized poly(glutamic acid) and poly(lysine), *Mini-Rev. Med. Chem.*, 2004, 4(2), 179–188.
- 10 J. F. Folmer-Andersen, E. Buhler, S.-J. Candau, S. Joulie, M. Schmutz and J.-M. Lehn, Cooperative, bottom-up generation of rigid-rod nanostructures through dynamic polymer chemistry, *Polym. Int.*, 2010, 59(11), 1477–1491.
- 11 D. A. Patrick, D. W. Boykin, W. D. Wilson, F. A. Tanious, J. Sychala, B. C. Bender, J. E. Hall, C. C. Dykstra, K. A. Ohemeng and R. R. Tidwell, Anti-Pneumocystis carinii pneumonia activity of dicationic carbazoles, *Eur. J. Med. Chem.*, 1997, 32(10), 781–793.
- 12 P. Besenius, G. Portale, P. H. Bomans, H. M. Janssen, A. R. Palmans and E. W. Meijer, Controlling the growth and shape of chiral supramolecular polymers in water, *Proc. Natl. Acad. Sci. U. S. A.*, 2010, 107(42), 17888–17893.
- 13 A. K. Hirsch, E. Buhler and J. M. Lehn, Biodynamers: self-organization-driven formation of doubly dynamic proteoids, *J. Am. Chem. Soc.*, 2012, 134(9), 4177–4183.
- 14 E. Buhler, S. J. Candau, J. Schmidt, Y. Talmon, E. Kolomiets and J.-M. Lehn, Fibrillar structure of self-assemblies formed from heterocomplementary monomers linked through sextuple hydrogen-bonding arrays, *J. Polym. Sci., Part B: Polym. Phys.*, 2007, 45(1), 103–115.
- 15 Y. Ruff, E. Buhler, S. J. Candau, E. Kesselman, Y. Talmon and J. M. Lehn, Glycodynamers: dynamic polymers bearing oligosaccharides residues—generation, structure, physicochemical, component exchange, and lectin binding properties, *J. Am. Chem. Soc.*, 2010, 132(8), 2573–2584.
- 16 G. Marinani, E. Buhler, J.-R. Colard-Itté, D. Dattler, N. Giuseppone, E. Moulin and S. Prevost, *Effects of polymer molecular weight on the structure and on the efficiency of light-driven contractile gels*, Institut Laue-Langevin (ILL), 2019, DOI: 10.5291/ILL-DATA.9-11-1845.