

Electronic Supplementary Information

A comparison of the network structure and inner dynamics of homogeneously and heterogeneously crosslinked PNIPAM microgels with high crosslinker content

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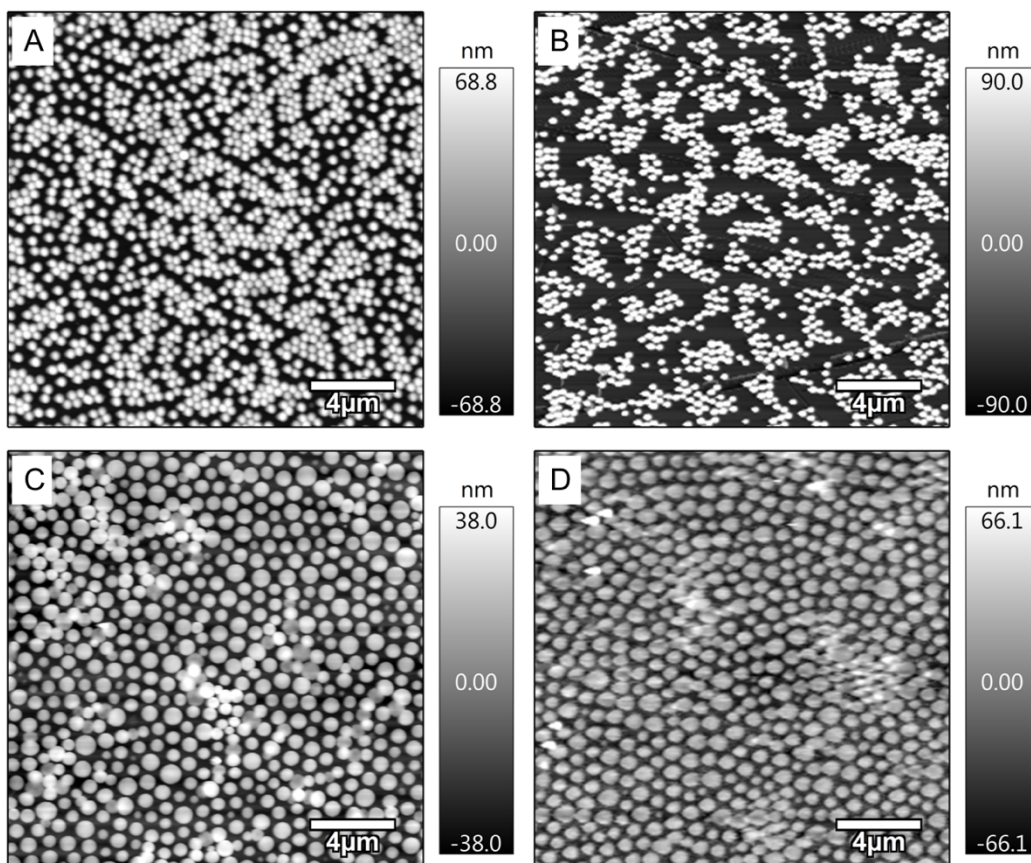


Figure S. 1: AFM images of A) HetMG10 under ambient conditions, B) HetMG10 swollen in H_2O , C) HomMG10 under ambient conditions and D) HomMG10 swollen in H_2O .

Table S. 1: Dry and swollen widths and heights of HetMG10 and HomMG10. Cross sections of 5 individual particles were analyzed and the standard deviation was calculated.

Microgel	Width _{dry} (nm)	Height _{dry} (nm)	Width _{H₂O} (nm)	Height _{H₂O} (nm)
HetMG10	626 ± 57	134 ± 8	650 ± 45	237 ± 15
HomMG10	885 ± 134	48 ± 1	1020 ± 138	106 ± 12

Images under ambient conditions were scanned in tapping mode with AC160TS cantilevers (Olympus). Images in H_2O were scanned with TR800PSA cantilevers (Olympus).

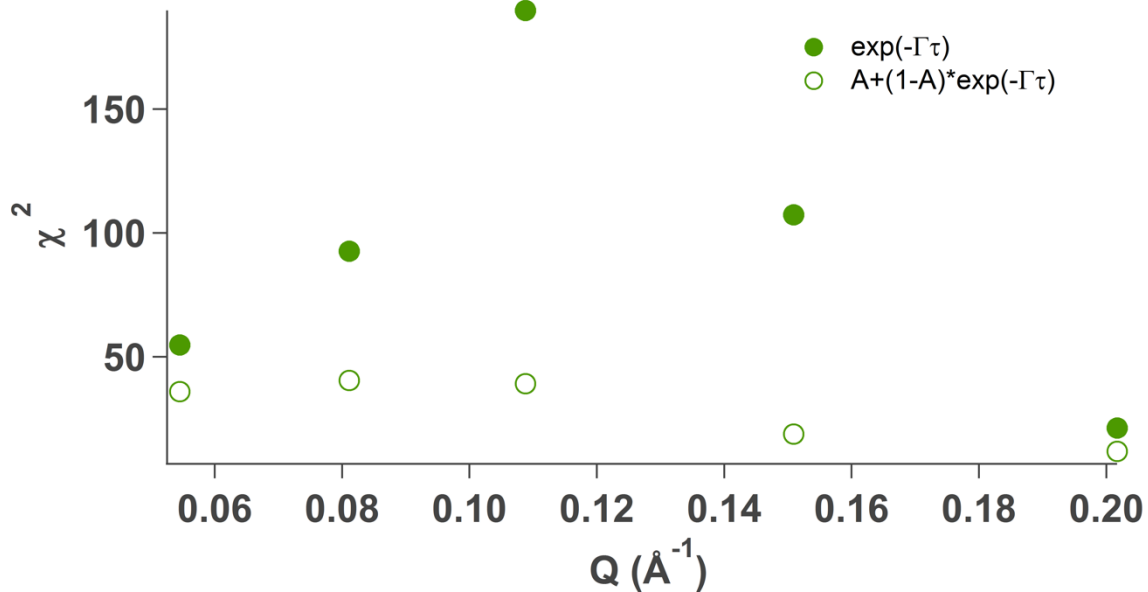


Figure S. 2: χ^2 for fits of the normalized intermediate scattering functions of HetMG10 with and without the elastic contribution.

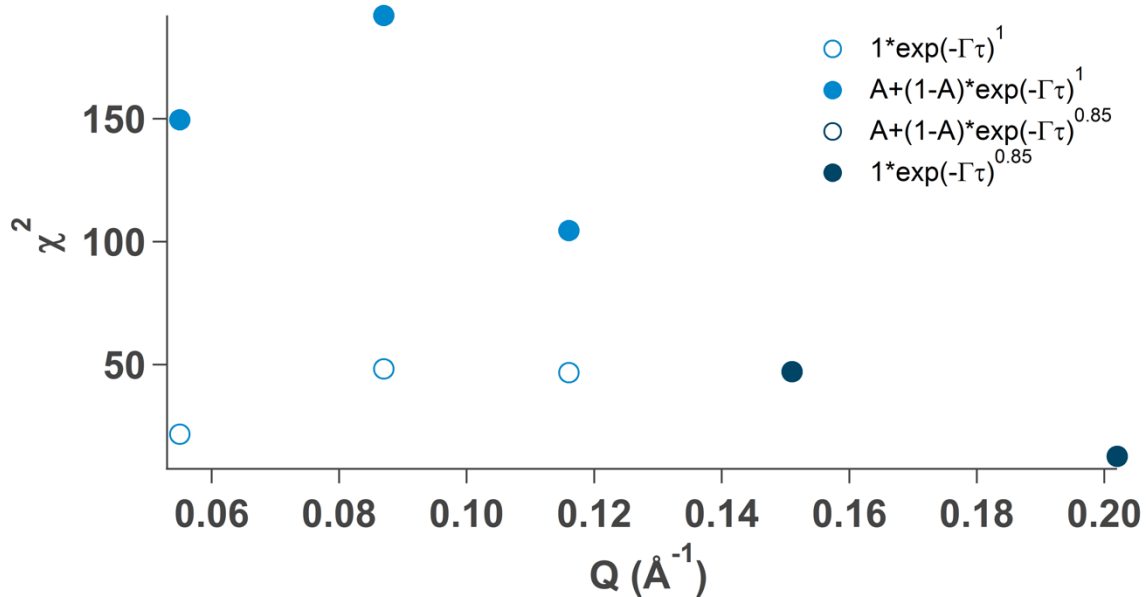


Figure S. 3: χ^2 for fits of the normalized intermediate scattering functions of HomMG10 with and without the elastic contribution. $Q = 0.15 \text{ \AA}^{-1}$ and $Q = 0.20 \text{ \AA}^{-1}$ were fitted with a stretched exponential with exponent $\beta = 0.85$.

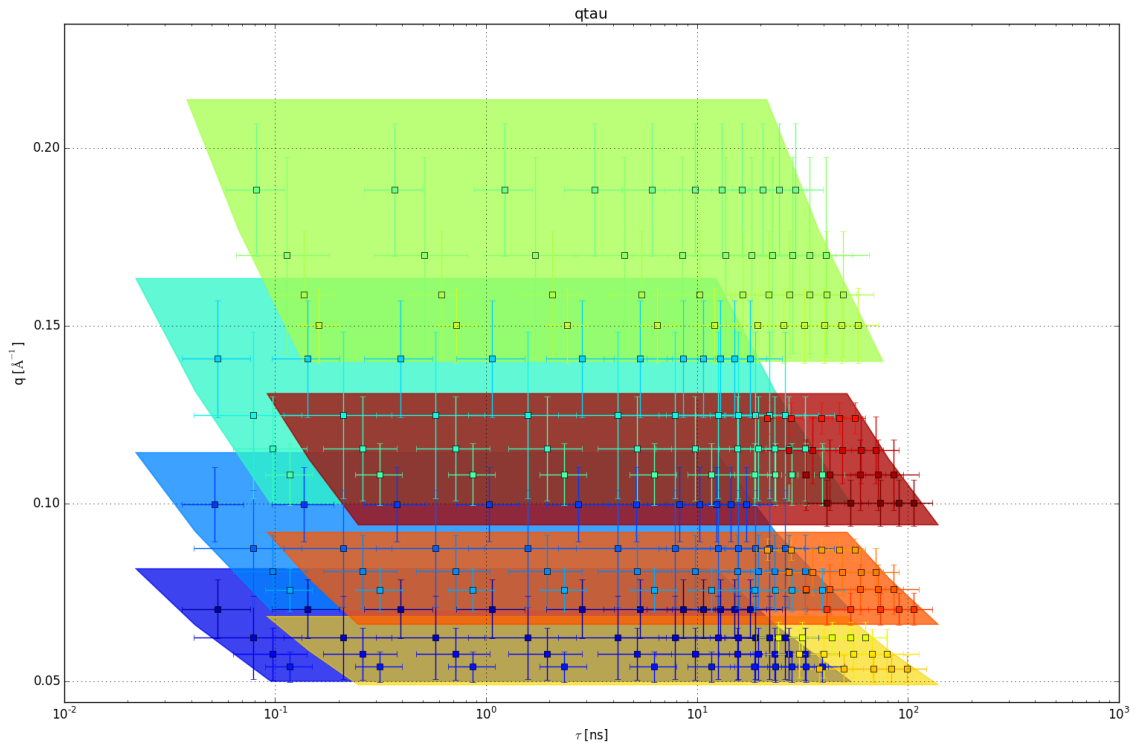


Figure S.4: Q - τ map for the measurements at the SNS-NSE at the spallation neutron source in Oak Ridge, TN.

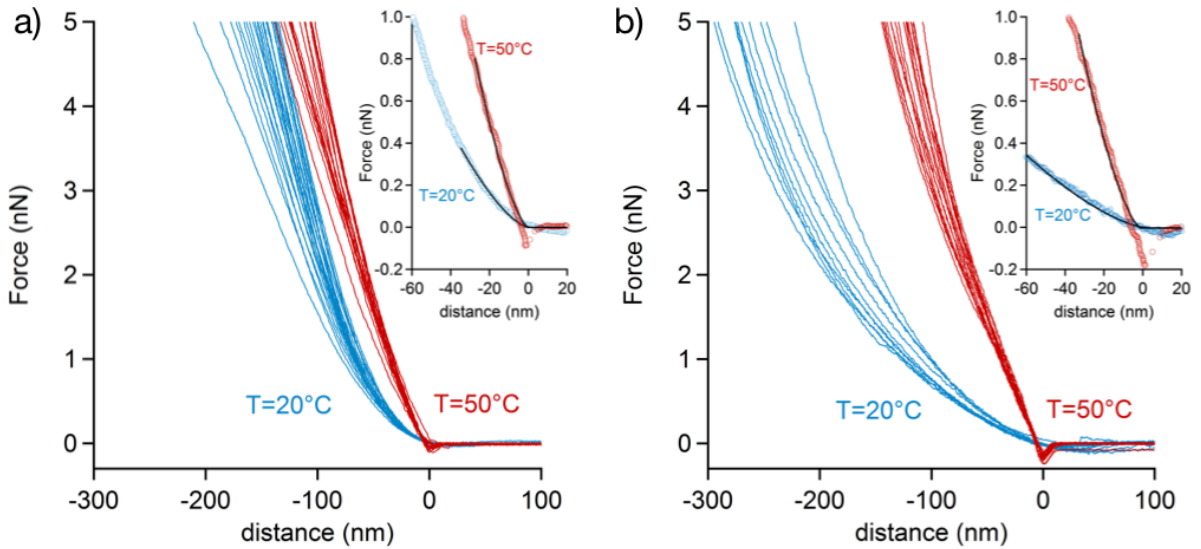


Figure S. 5: AFM force-distance curves for a) $MG_{10}P$ and b) $MG_{10}F$ at $T = 20^\circ\text{C}$ (blue) and $T = 50^\circ\text{C}$ (red). Insets show an exemplary fit according to the Hertz model (equation 1 in main text) at $T = 20^\circ\text{C}$ (blue) and $T = 50^\circ\text{C}$ (red). The positive indentation depth values were transferred into negative tip-sample-distances.

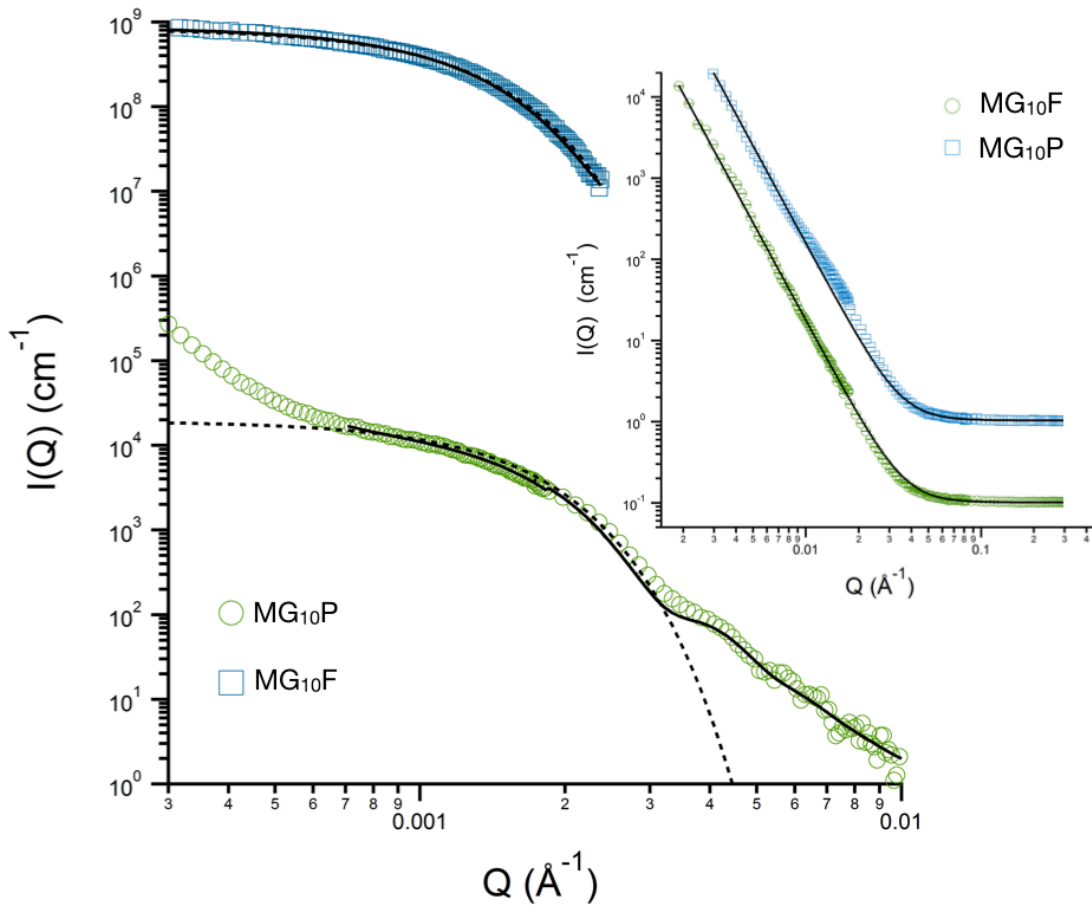


Figure S. 6: SANS curves of MG10P and MG10F measured at 50°C in a wide Q range. A Guinier analysis of the data yields the radius of gyration R_g at 50°C. The dashed line in the figure are the Guinier fits to the data. Solid lines are fits to the fuzzy sphere model. Results are given in Table 2 and 3 in section 4.4 of the paper. The inset shows the medium and high Q-range. For both samples, the intensity decreases in this range according to the Porod law, $I \propto Q^{-4}$ until the incoherent background level is reached.

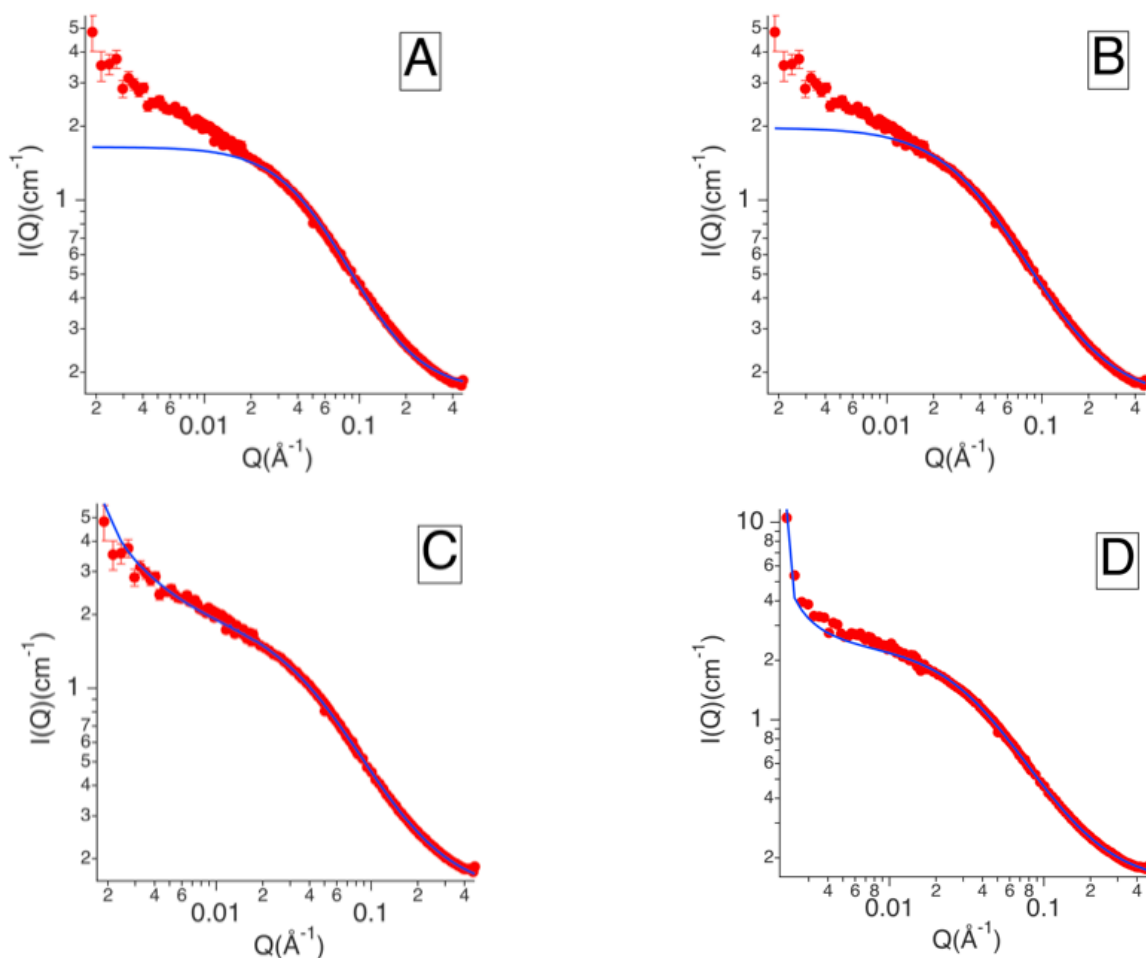


Figure S. 7: A: Ornstein Zernike fit with $m=2$, no clustering term

B: modified fit equation with $m=1.7$, according to Rubinstein and Colby for semi dilute polymers in good solvent
 C: correlation length fit with $m=1.7$ and a low Q exponent of $n=1.2\pm 0.002$ indicating clustering within the PNIPAM solution as described by Hammouda et al. (**J. Chem. Phys.** **133**, 084901 (2010))

—> low exponent, no scattering from an interface as in Porod scattering ($1/q^4$), but structural inhomogeneities in the semi dilute solution on larger length scales at low Q's

D: data at 20°C: correlation length fit with $m=1.7$ and a low Q exponent of $n=2.0 \pm 0.01$ indicating clustering within the PNIPAM solution as described by Hammouda et al. The scattering intensity seems to be increased compared to 15°C (10 vs. 5) and the clustering exponent is higher.

	A	B	C	D
m	2	1.7 ± 0.001	1.7 ± 0.001	1.7 ± 0.001
n			1.2 ± 0.002	2.1 ± 0.01
ξ (nm)	2.1 ± 0.03	2.7 ± 0.03	2.4 ± 0.02	2.9 ± 0.01

Figure 7.A-C displays data at 15°C and Figure 7.D at 20°C. As described by Hore and Hammouda we also observe that the low-Q feature becomes steeper with increasing temperature.

However, for our purposes we are concerned with the intermediate and high-Q region, which give us information about the correlation length ξ . As can be seen in the table the values for ξ are similar independent of the chosen fit (between 2.1 and 2.9) with an average value of 2.5 nm and a standard deviation of 0.35 nm.

The exponent $m=1.7$ was used as suggested by Rubinstein and Colby for polymers in good solvent. (M. Rubinstein, R. H. Colby, *Polymer Physics*, Oxford University Press, New York, 2003, pp. 189-190.)

Since water is a good solvent for PNIPAM below the LCST of 32°C and we measured at 15°C and 20°C, this assumption is valid for our data.