

Table 1. Fit parameters for bead models of unfolded proteins

Protein and conditions	$\epsilon = 0.0$, σ varies	$\epsilon = 0.1$, σ varies	$\sigma = 7.48$, λ varies	Gaussian chain
	σ, Å	σ, Å	λ	$\langle R^2 \rangle^{1/2}$, Å
Protein L, 2.5 M GdmCl	4.70	6.72	0.120	58.78
CspTm, 2.5 M GdmCl	5.10	7.00	0.114	61.21
Protein L/CspTm, 4 M GdmCl	5.50	7.14	0.108	64.22
Protein L/CspTm, 7.2 M GdmCl	6.10	7.48	0.100	70.87