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Supplementary Information for

Unusual Denaturation Trajectory of Bovine Gamma Globulin Studied by Fluorescence Correlation Spectroscopy

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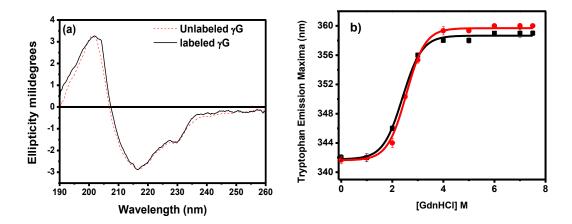


Fig. S1 a) Far-UV CD spectra of γG in the Rhodamine-labeled and unlabeled states. b) Tryptophan emission maxima (excitation at 295 nm) of labeled (and unlabelled () γG in solutions of increasing GdnHCl concentration, superposed with sigmoidal fit of each set of data.

Table S1. Percentages of secondary structure content of γG as function of GdnHCl concentration obtained by fitting the CD spectra of Fig. 2(a) with the CDSSTR suite of programs. The RMSD values obtained were <0.03.

| [GdnHCl] (M) | Helix | Beta sheet | Turns | Random coil |
|--------------|-------|------------|-------|-------------|
| 0 | 3 | 63 | 14 | 20 |
| 1 | 17 | 55 | 10 | 18 |
| 2 | 36 | 43 | 9 | 12 |
| 2.5 | 53 | 35 | 4 | 8 |
| 3 | 36 | 36 | 08 | 20 |
| 4 | 28 | 39 | 11 | 22 |
| 5 | 21 | 29 | 18 | 32 |
| 6 | 13 | 27 | 17 | 43 |

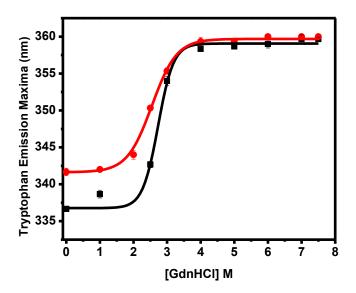


Fig. S2 Unfolding profiles of R- γG measured by fluorescence in absence (\blacksquare) and presence (\blacksquare) of 2 M NaCl. Solid lines are sigmoidal fits to the data.

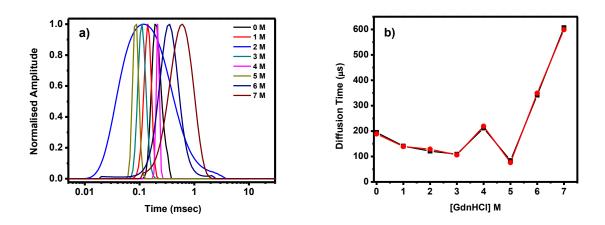


Fig. S3 a) Variation of the maximum entropy profiles of R- γG with GdnHCl concentrations. b) Comparison of diffusion times obtained by fitting the ACF with Eqn. (1) of text (\bigcirc) with the times at the peak of the distributions obtained from the MEMFCS program (\bigcirc).

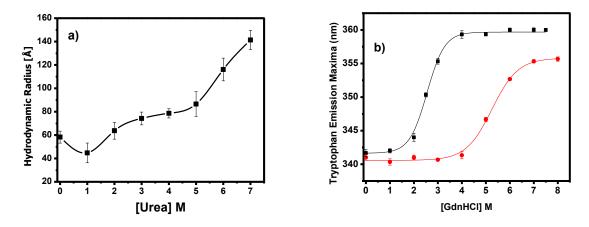


Fig. S4 a) Urea induced denaturation of R- γG studied with FCS. b) Change in wavelength of Trp emission maxima of R- γG in presence of varying concentration of GdnHCl (\blacksquare) and urea (\bullet) during unfolding.