

Supporting Information for

Exploring the inner environment of protein hydrogels with fluorescence spectroscopy toward understanding their drug delivery capabilities

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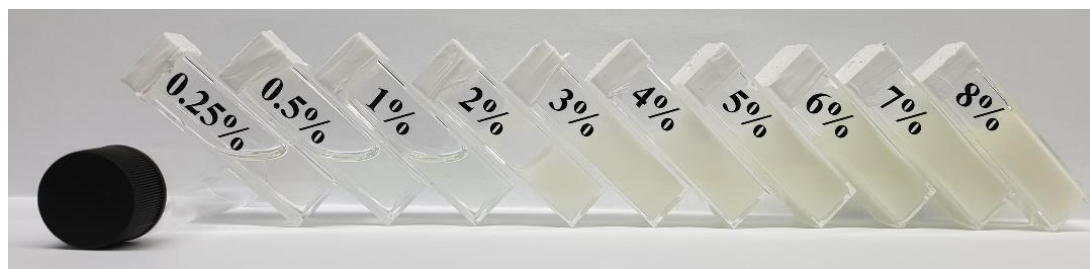


Figure S1: Images of different samples after gelation.

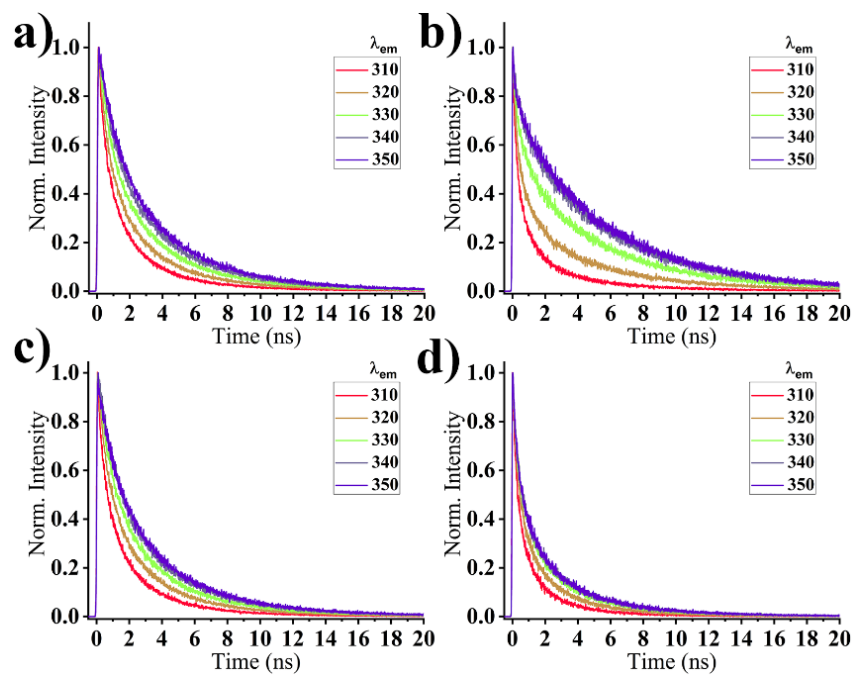


Figure S2: Wavelength dependent fluorescent transient decay of BSA solution before a) 0.25%, b) 8% and after gelation c) 0.25%, d) 8%.

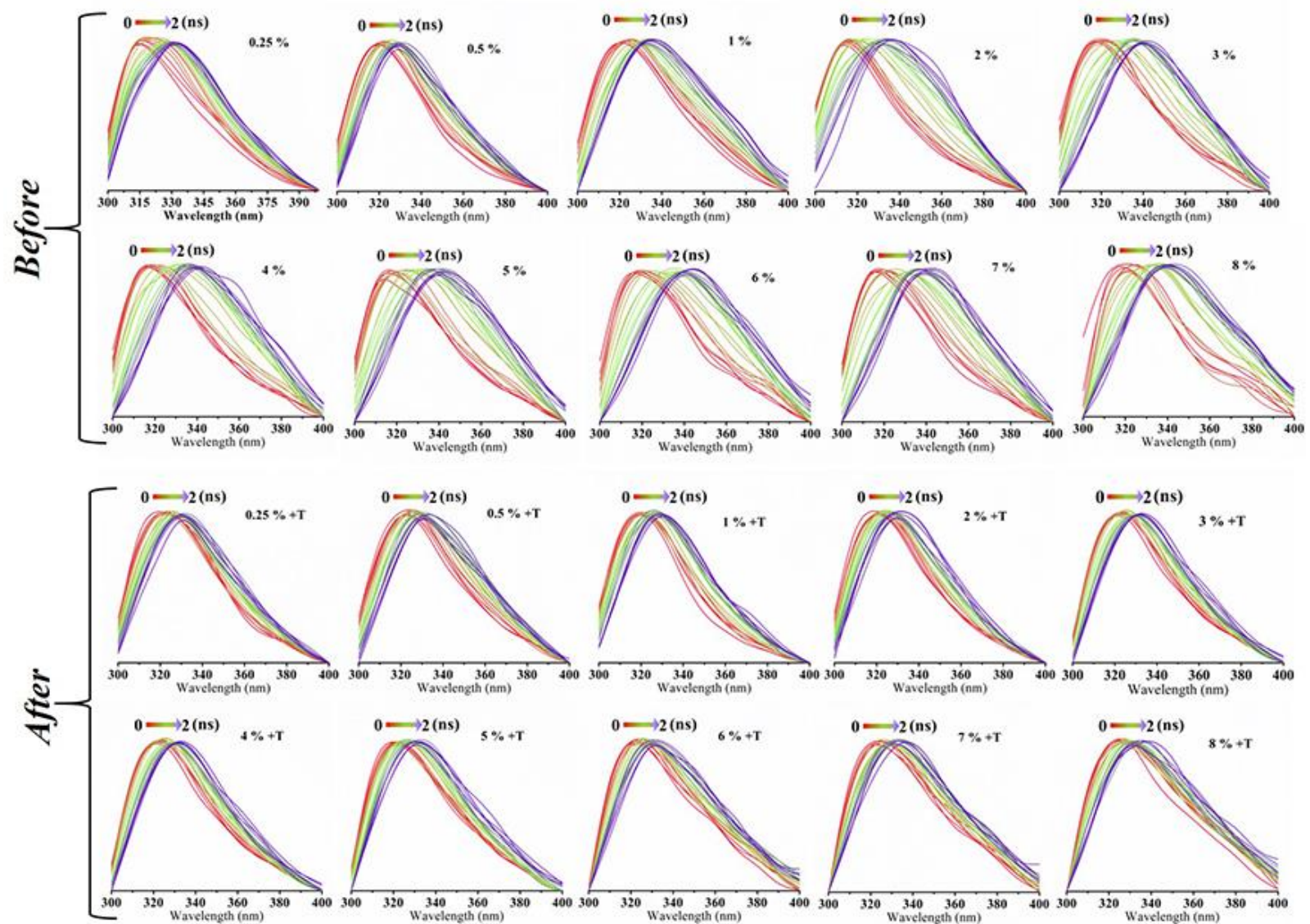


Figure S3: TRANES measurements for the first 2 ns of the emission decay of Trp for different weight percentage of BSA before and after thermal treatment.

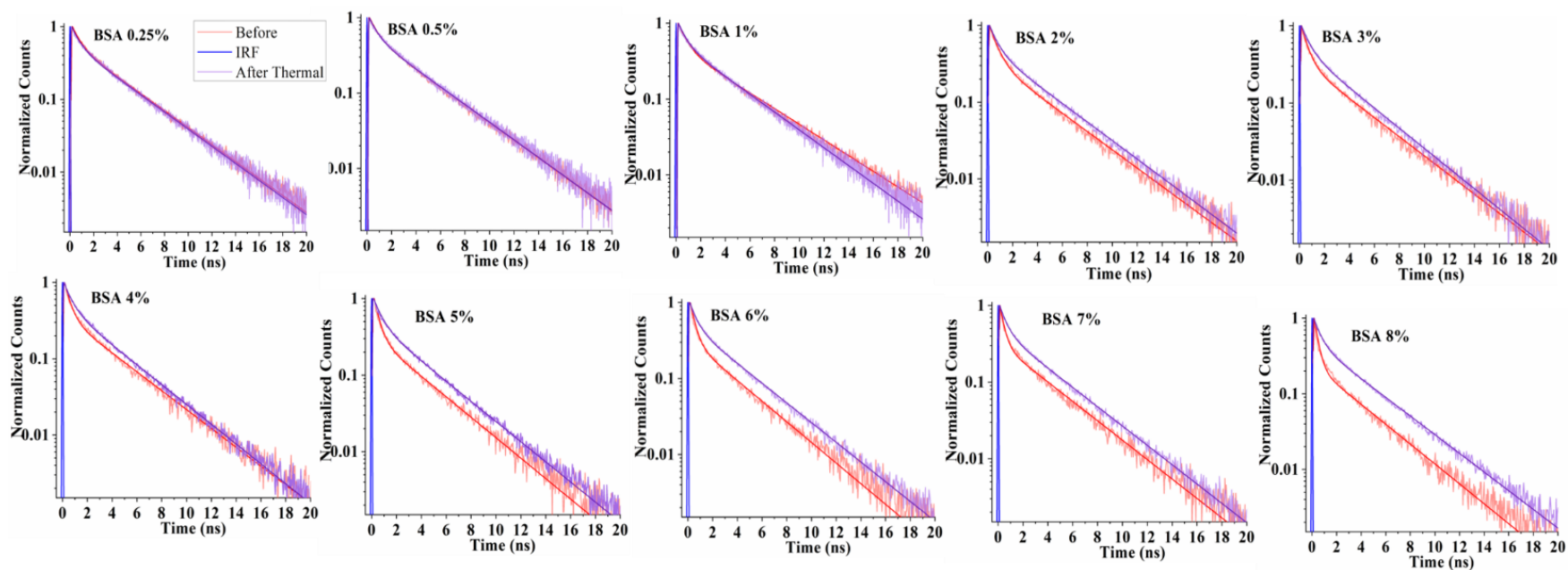


Figure S4: Two exponent fitted time resolved emission spectra of HPTS inside BSA (0.25 to 0.8% concentration) before and after thermal. Samples were excited at 390 nm and emission was monitored at 435 nm.

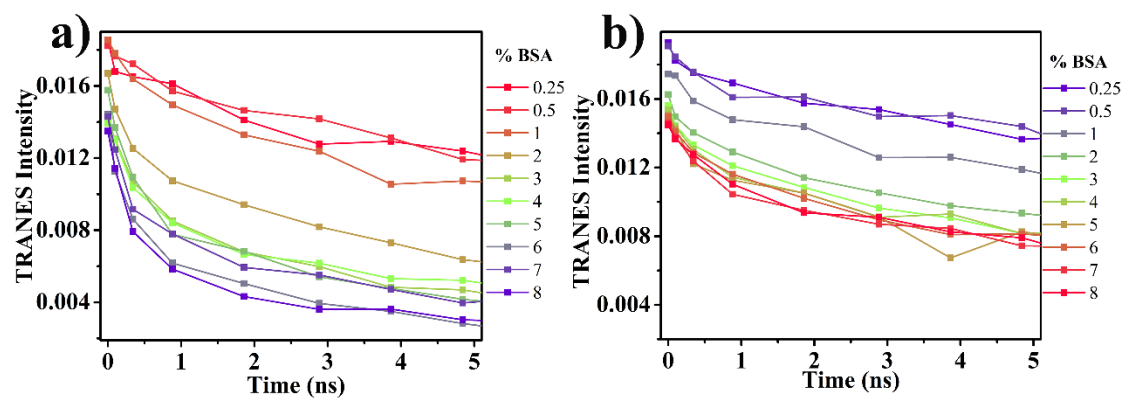


Figure S5: Change in ROH* intensity with time a) before and b) after gelation, was evaluated from TRANES of each sample.

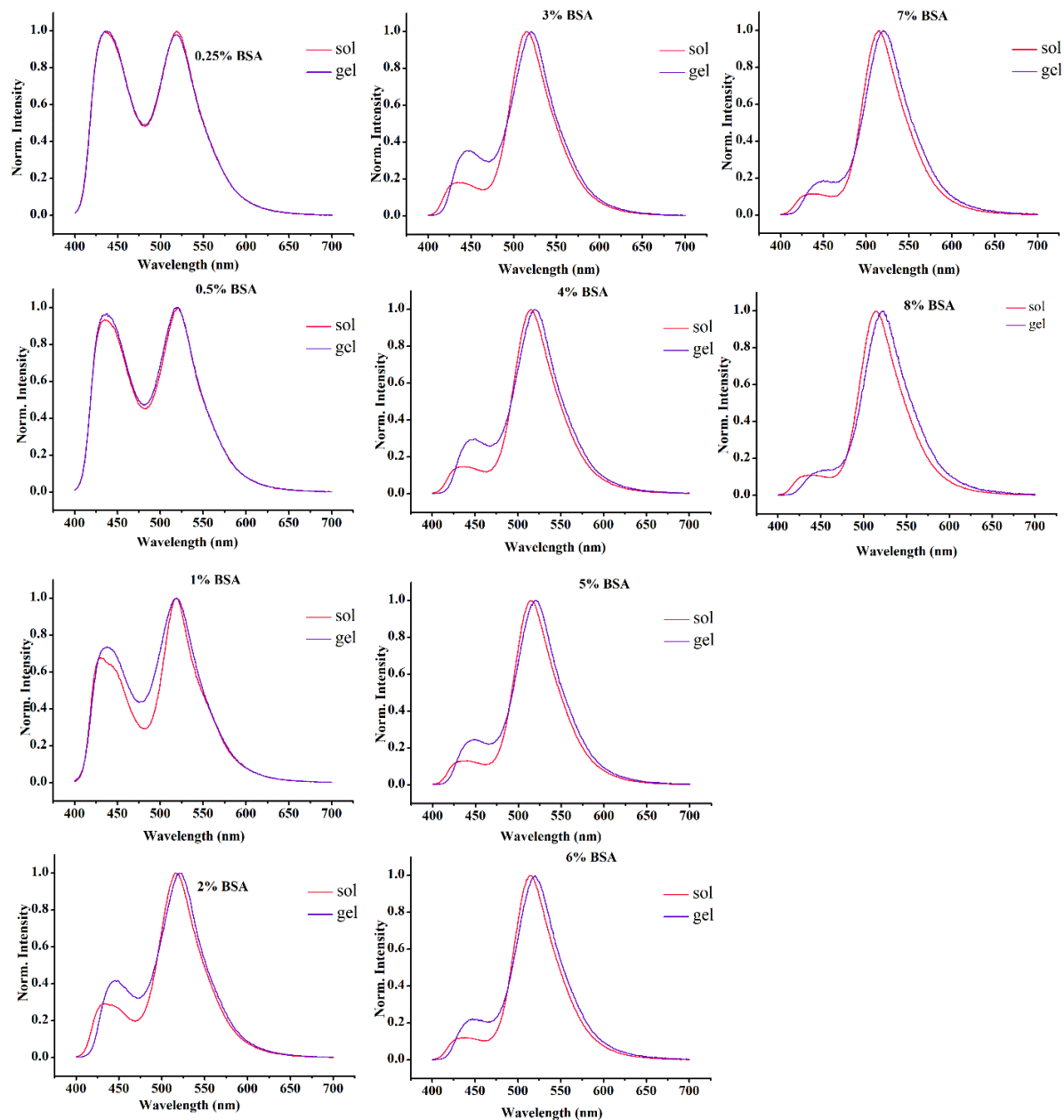


Figure S6: Steady state emission spectra of HPTS inside BSA before and after gelation normalized at RO⁻ peak position.

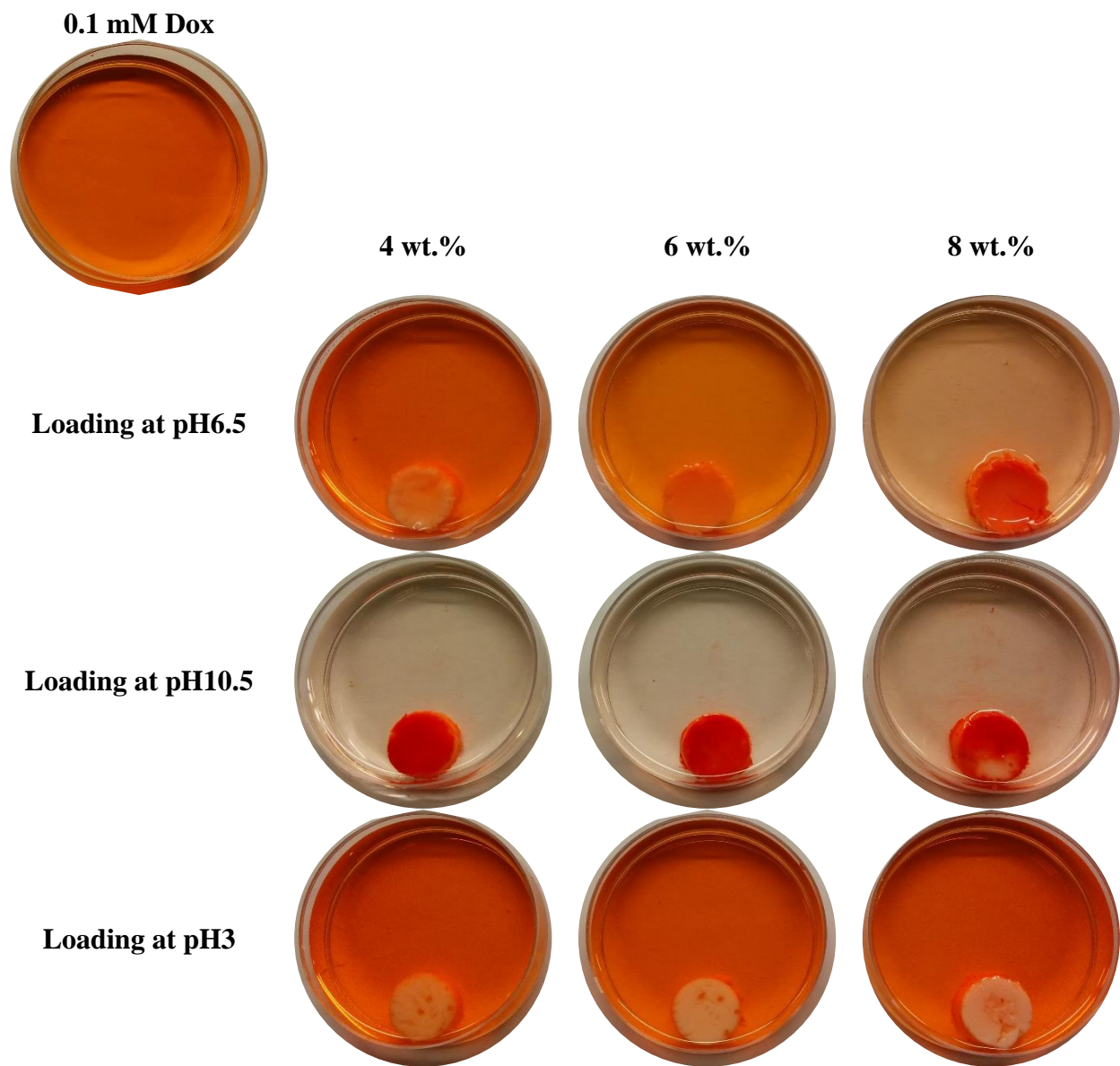


Figure S7. Images of the loading of Dox within BSA hydrogels at different weight concentrations and at different loading pH values, together with a picture of 0.1 mM Dox solution. The volume of the gels is 2 ml. The pictures were taken following 3 days of loading.

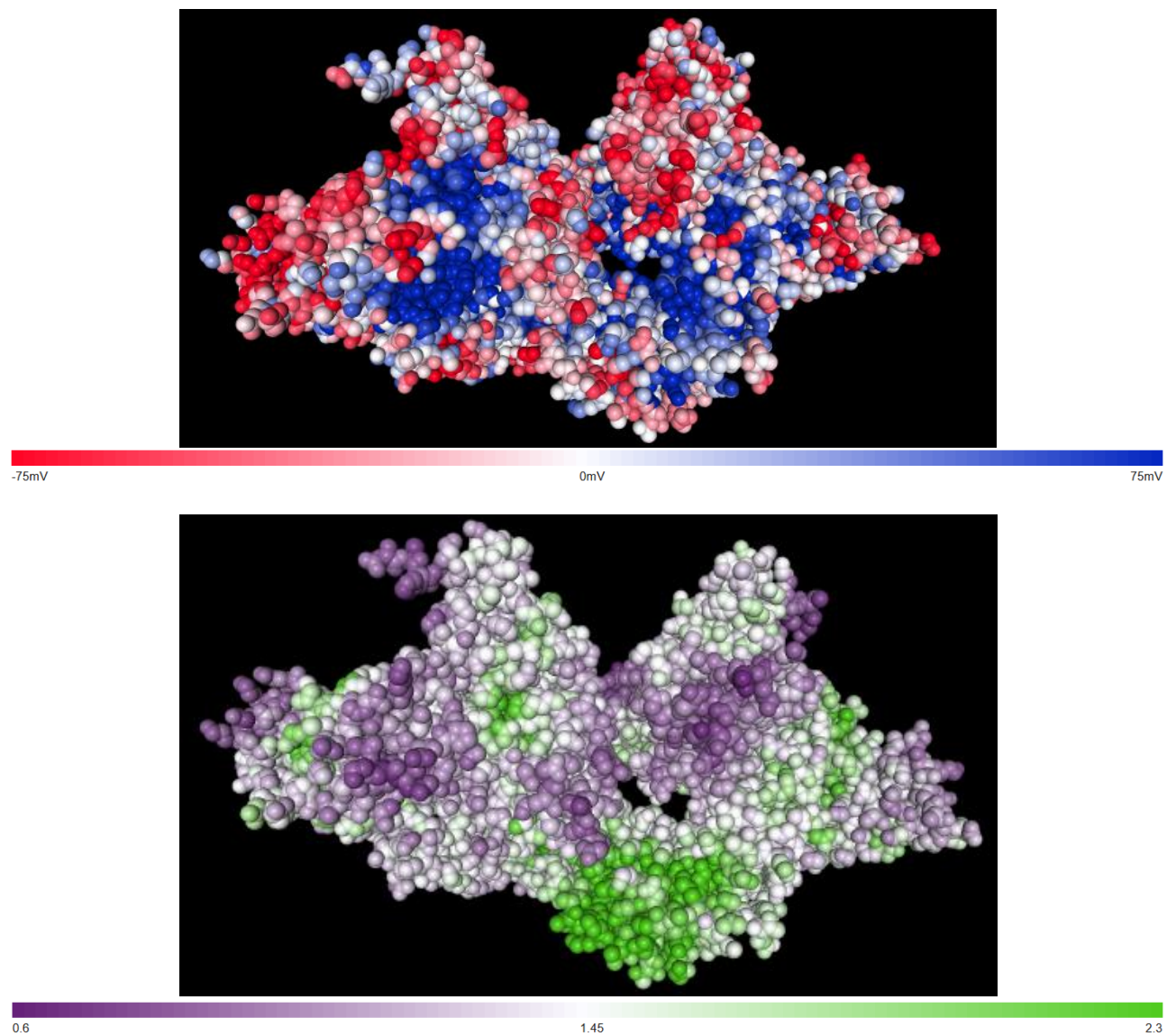


Figure S8. (a) Potential field and (b) non-polar/polar ratio (green is more hydrophobic and purple is more polar) of BSA (PDB ID: 4F5S). The maps were generated using Protein-Sol (M. Hebditch and J. Warwicker, *Sci. Rep.*, 2019, **9**, 1969).