

Supporting Information

Effects of tryptophan microenvironment, soluble domain, and vesicle size on the thermodynamics of membrane protein folding: Lessons from the transmembrane protein OmpA

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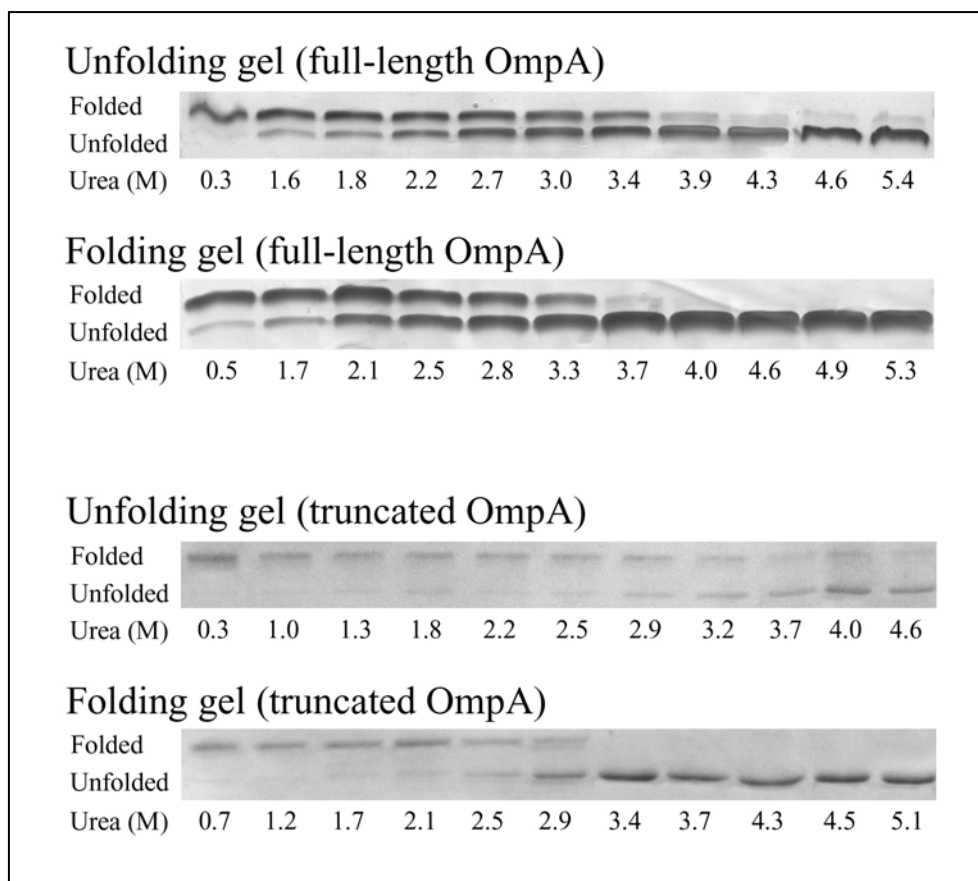


Figure S1. Unfolding and folding gel-shift assays of representative full-length (top) and truncated (bottom) single-trp mutants of OmpA. The C_m values determined from the fluorescence studies was 3.1 M urea for the full-length mutant and 2.3 M for the truncated mutant. See main text for additional details.