Supporting Information

Yuan and Bauer 10.1073/pnas.0802149105

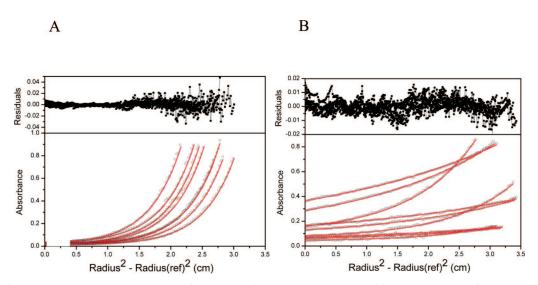


Fig. S1. Results from sedimentation equilibrium analysis of PixD alone (A) and the PixD-PixE samples (B). Absorbance values from independent experiments are indicated as open circles with the lines representing fitted data based on a single ideal macromolecular component model. Residuals of the fits are plotted at the top.