

Supporting Information, AhpF and thioredoxin reductase chimera, Reynolds and Poole

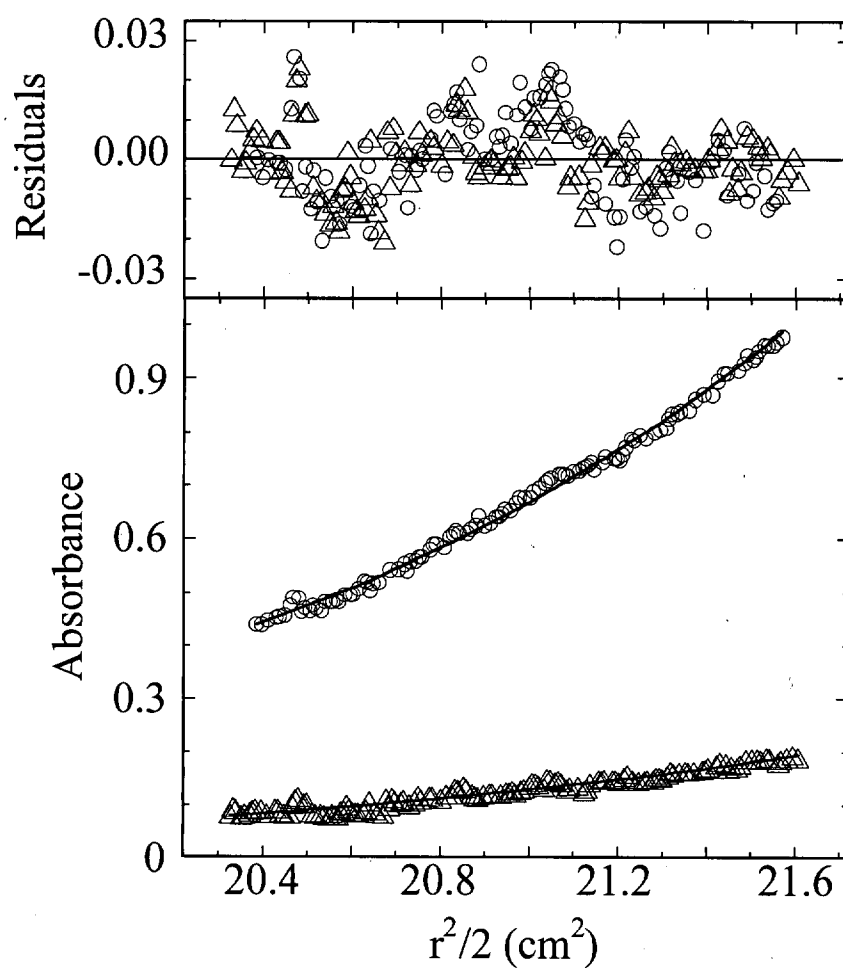


Figure S1. Sedimentation equilibrium analyses of Nt-TrR to determine molecular weight. Shown are single data sets at 280 (circles) and 450 nm (triangles) for Nt-TrR equilibrated at 7,300 rpm and 20 °C. Solid lines represent global fits to a single ideal species model of 15 data sets at 280 nm and 450 nm, and rotor speeds from 7,300 to 14,300 rpm. The upper panel depicts the deviations between experimental and calculated data (residuals) for each data set.