

Supplementary material: Light-scattering/refractive index analysis.

The protein was stored in 10 mM Tris, 5 mM DTT and the Shodex 802.5 gel-filtration column was run at 277 K in a buffer containing was: 250 ppm azide, 100 mM NaCl, 50 mM Tris 7.5 at 277 K. Data analysis revealed a single dominant monodisperse peak (A) with MW \approx 101 kDa (B,C) that corresponds to a tetramer of Pf1337 (protomer 25.7 kDa).

