



Figure S3. Structure of Crosslinked CA (Orthorhombic Crystal Form)

(A) Bias-minimized map (pink mesh, 1σ) and unbiased omit density (blue mesh, 1σ) after rigid body refinement of 23 domains from the molecular replacement solution, with the final model of the omitted domain.

(B) Crystal packing viewed perpendicular to the ab plane. The unit cell is colored magenta. The two hexamers in the asymmetric unit are colored yellow and green.

(C) Per residue real space correlation plots against the bias-minimized map.