

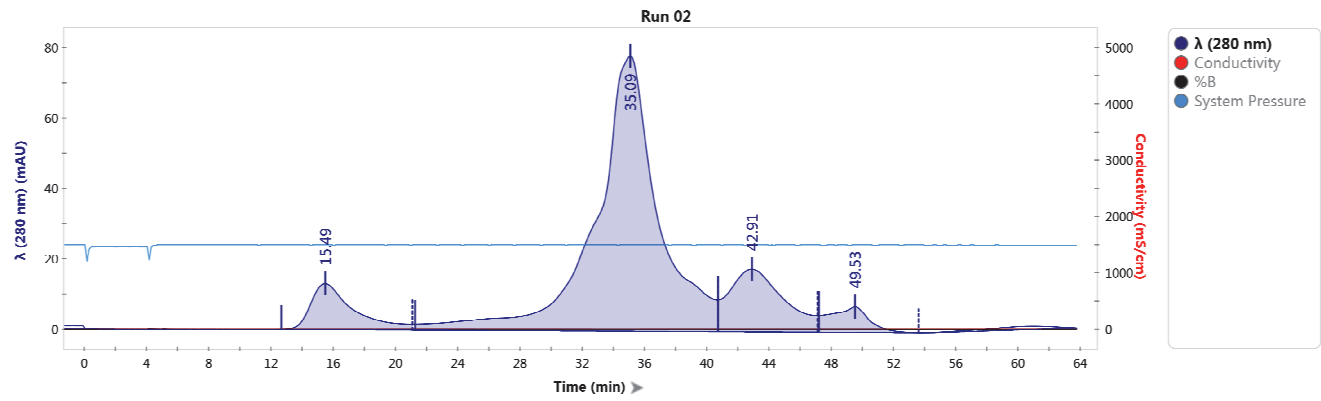


Fig. S1. Analysis of fast protein liquid chromatography of purified CelA203.

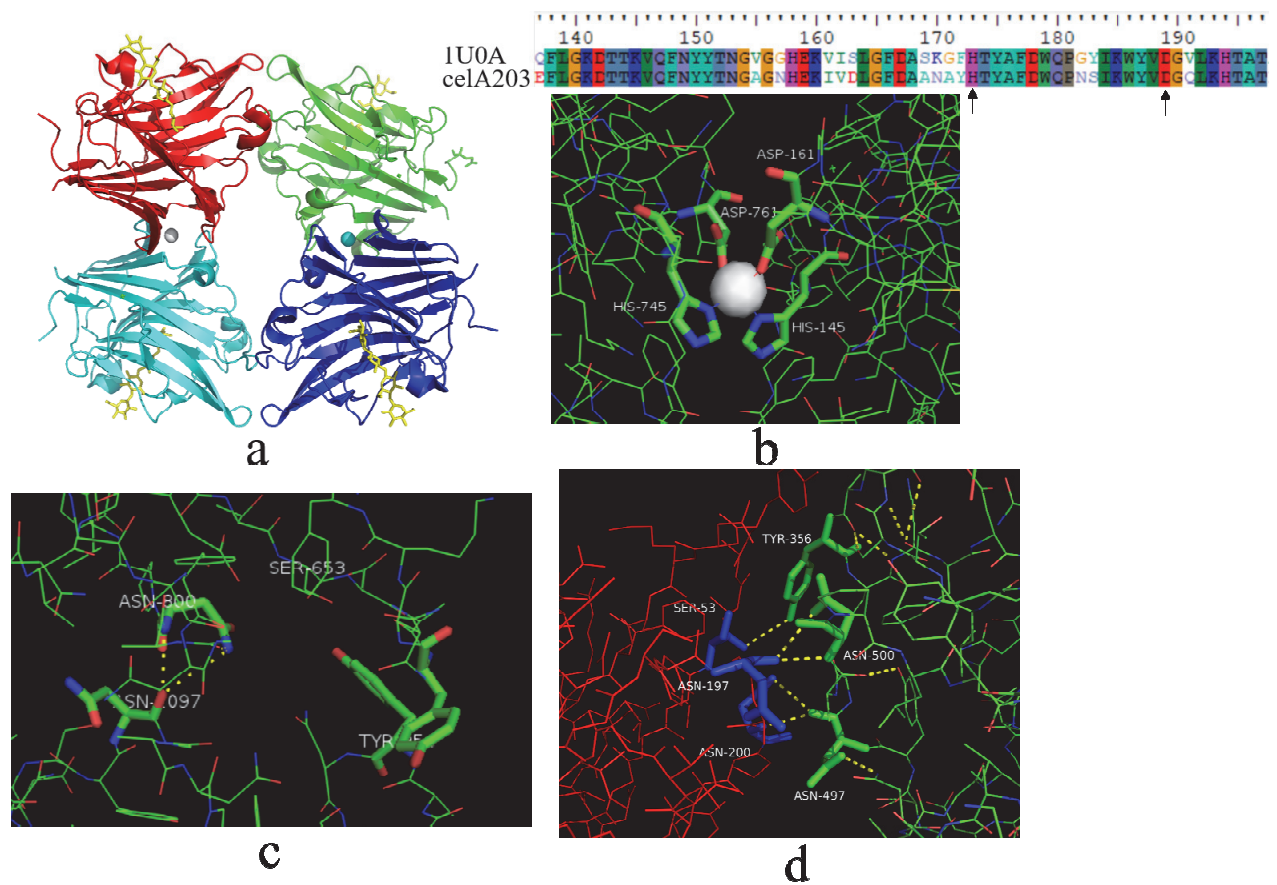


Fig. S2. 3D structure model of CelA203 tetramer protein. (A) Ribbon display of 3D structure. (B) 3D structure model of zinc ion binding amino acid. (C) (D) Amino acid sites binding between dimers.