

Molecular mechanism of the *Escherichia coli* AhpC in the function of a chaperone under heat-shock conditions

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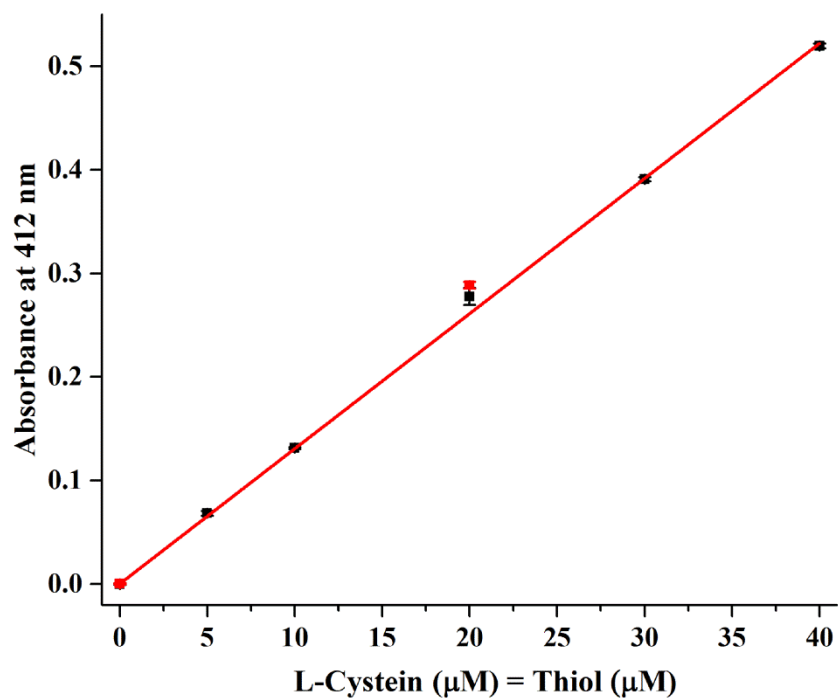
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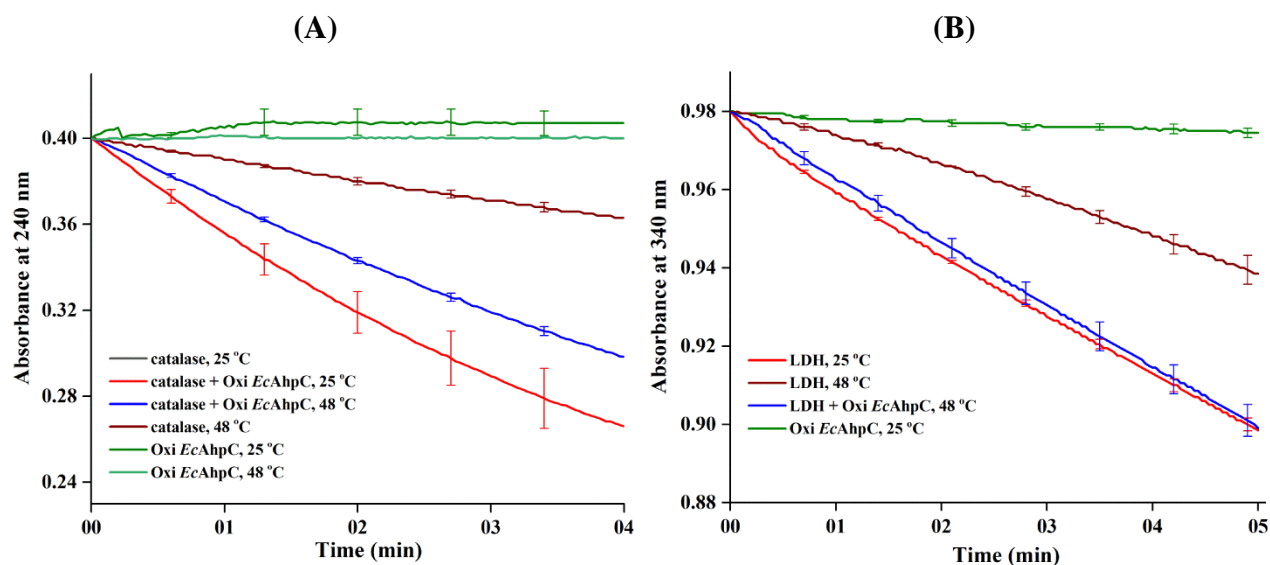
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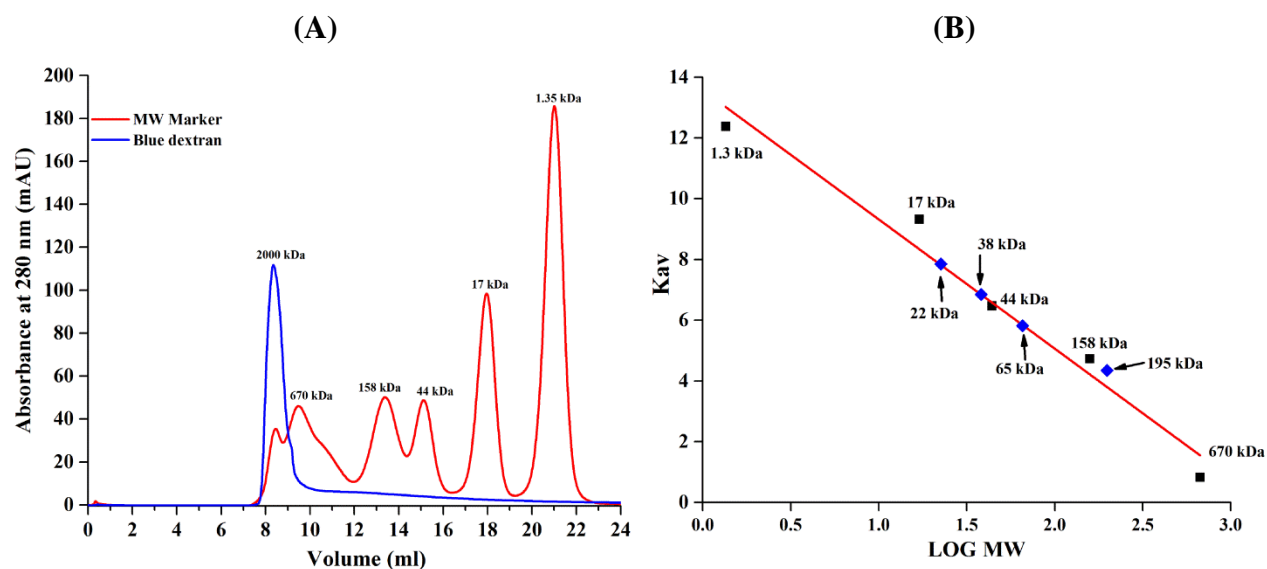
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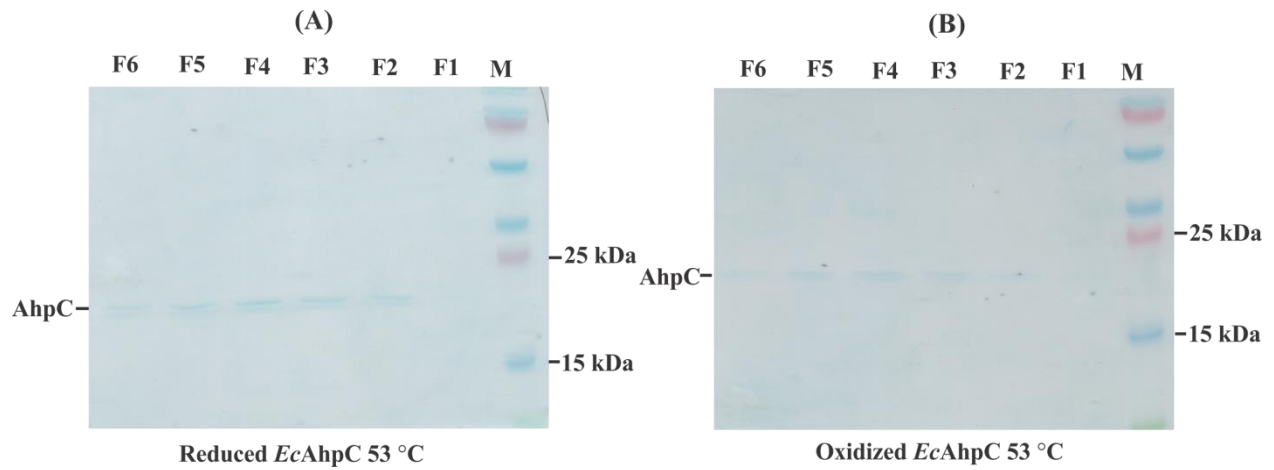
Supplementary Figure S1. *Standard curve for free thiol quantification.* The standard curve is generated by reacting DNTB with different molar concentrations of 0-40 μM L-cysteine (*black square*), followed by detection of the TNB produced at $\lambda = 412$ nm, which was used to calculate the free thiol content of 10 μM of recombinant WT *EcAhpC* (*red square*). The oxidized *EcAhpC* was estimated to contain 0 ± 0.03 thiol per monomer and the reduced *EcAhpC* contains 2.1 ± 0.3 thiols per monomer.



Supplementary Figure S2. Catalase and LDH activity assay. (A) The decrease in absorbance at $\lambda = 240$ nm resulting from decomposition of H_2O_2 was measured to assess the catalase activity in the presence of oxidized *EcAhpC* at 25 °C and 48 °C. Oxidized *EcAhpC* alone did not show H_2O_2 -decomposition activity at 25 °C and 48 °C (*green*), since no decrease in absorbance was observed. (B) The decrease in NADH absorbance at $\lambda = 340$ nm was measured to assess the enzymatic activity of LDH in the presence of oxidized *EcAhpC* at 25 °C and 48 °C. The absence of LDH did not show NADH oxidase activity at 25 °C (*green*), since no decrease in absorbance was observed at 340 nm.



Supplementary Figure S3. Calibration curve of the SEC. (A) Elution profiles from Superdex 200 10/300 GL column (GE Healthcare) for the calibration of standards (thyroglobulin 670 kDa, γ -globulin 158 kDa, ovalbumin 44 kDa, myoglobin 17 kDa and vitamin B₁₂ 1.35 kDa) and blue dextran 2000 as absorption in milliabsorption units (mAU) at $\lambda = 280$ nm versus volume in milliliters (ml). (B) The calibration curve of the Superdex 200 10/300 GL column was prepared using protein standards (*black squares*). The Kav value was calculated using the equation $K_{av} = (V_e - V_o)/(V_c - V_o)$, where V_o = column void volume = 8.3 ml, V_c = geometric column volume = 23.6 ml and V_e = elution volume for each protein: thyroglobulin (670 kDa) $V_e = 9.48$ ml, γ -globulin (158 kDa) $V_e = 13.38$, ovalbumin (44 kDa) $V_e = 15.12$, myoglobin (17 kDa) $V_e = 17.97$, and vitamin B₁₂ (1.35 kDa) $V_e = 21.03$. For each protein samples, the observed V_e was used to calculate the corresponding Kav value that was used to determine the molecular mass. The estimated molecular mass of oxidized *EcAhpC* decamer (195 kDa), LDH (65 kDa), oxidized *EcAhpC* dimer (38 kDa) and reduced *EcAhpC* monomer (22 kDa) is shown (*blue diamond*).



Supplementary Figure S4. *HMW oligomers forming ability of reduced and oxidized EcAhpC at 53 °C.* (A) SDS-PAGE analysis (full-length gel is presented) of the 8 ml elution fractions (F1 to F6) as indicated in the chromatogram (Fig. 4A) of reduced *EcAhpC* incubated at 53 °C for 60 min. (B) A similar 8 ml elution fractions (F1 to F6) of oxidized *EcAhpC* incubated at 53 °C for 60 min has been collected and tested on SDS-PAGE analysis. The amount of formed HMW oligomers of reduced *EcAhpC* at this temperature was higher compared to oxidized *EcAhpC*.