

Supplementary Materials: The following supporting information can be downloaded at: www.mdpi.com/xxx/ and supplementary materials contain: Figure S1, Figure S2 and Table S1.

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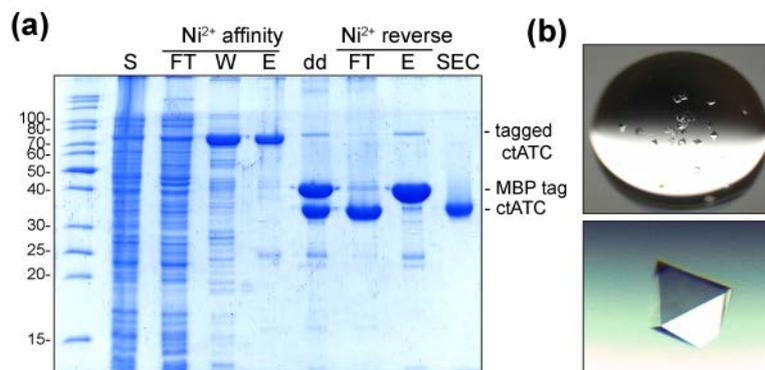


Figure S1. ctATC2Cys purification and crystallization. (a) SDS-PAGE of different steps along the purification of ctATC2Cys: S, soluble fraction of the lysate; Ni²⁺ affinity FT, W, and E, are the flowthrough, wash with 35 mM imidazole and elution of the first affinity column; dd, sample after dialysis and digestion with PreScission; Ni²⁺ reverse FT and E are the flowthrough and elution of the second affinity column; SEC, sample after size exclusion chromatography. (b) Bipyramidal-shaped crystals of ctATC2Cys.

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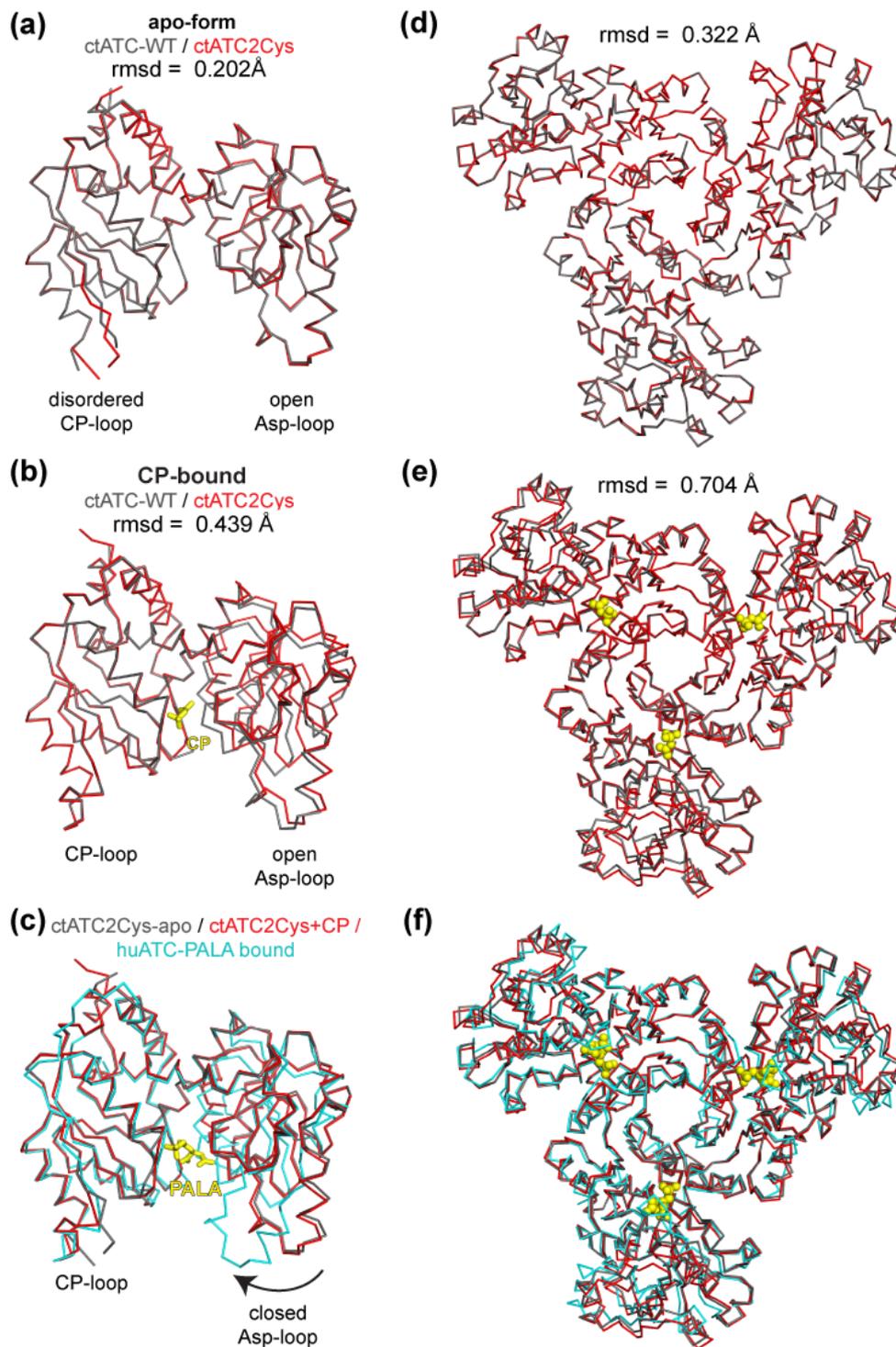


Figure S2. Structural similarity of ctATC2Cys and ctATC-WT. Ribbon superposition ctATC-WT (in grey) and ctATC2Cys (in red) subunits in apo form (a) or bound to CP (b). (c) Superposition of ctATC2Cys apo and CP-bound structures with human ATC bound to PALA (shown in yellow sticks). The arrow indicates the expected movement of the Asp-loop in ctATC2Cys upon binding to PALA. (d–f) Superpositions for the entire protein trimer.

Table S1. Oligonucleotides used for PCR amplification and mutagenesis of ctATC. Codons for Cys mutations are in red and flanking regions for In-fusion cloning are underscored. 14
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Oligonucleotide	Sequence 5'-3'
ctATC-N2045C_Fw	CTG CGA ACC CTG GCC TGC TAC TCT GAT GCT ATC
ctATC-N2045C_Rv	GAT AGC ATC AGA GTA GCA GGC CAG GGT TCG CAG
ctATC-R2238C_Fw	TAT TTT CGG CAG ATG TGC TAT GGG CTG TAT TGT
ctATC-R2238C_Rv	ACA ATA CAG CCC ATA GCA CAT CTG CCG AAA ATA
ctATC-Fw	<u>AAGTTCTGTTTCAGGGCCCTCATTCAAGAAATCTCAC</u>
ctATC-Rv	<u>ATGGTCTAGAAAGCTTTAGGACATGACCAGTGC</u>