Expanded View Figures

Figure EV1. Packing of SERCA2a E2 $-AlF_4^-$ -CPA and [Ca₂]E1-AMPPCP crystals.

- A Parallel packing of the SERCA2a structure in the E2–AIF₄⁻-CPA conformational state. SERCA2a forms contact points between the A-domains and N-domains (Met1-Asn3, His15 to Glu458-Lys460) and between the A-domain (Arg134, Arg139, Lys141) and the P-domain (Glu659, Glu667, Ser663), as well as N-domain (Thr506-Ser509, Asp567-Arg571) and L7/8 (Asp859-Arg863, Lys876-Gly884).
- B Parallel packing of the SERCA2a structure in the E2–AIF₄-CPA conformational state. Contact points are within the A-domain (Arg35-Glu40) and P-domain (Asp658-Ser663). A-, P-, and N-domains are colored in yellow, blue, and red, respectively. M1-M2 are colored in wheat, M3-M4 in brown, M5-M6 in dark gray, and M7-M10 in light gray.



Figure EV1.



Figure EV2. Coordination of ions and molecules in the SERCA2a structures.

A-C Ca²⁺ ions are shown in purple spheres. Omit maps for the CPA (A) and the AMPPCP (C) molecules contoured at 3 σ . (B) CPA coordination in [H₂₋₃]E2-AIF₄⁻-CPA state of SERCA. SERCA1a is depicted in gray and SERCA2a in orange. Coordination of Ca²⁺ ions in the [Ca₂]E1-AMPPCP state. SERCA1a is shown in gray and SERCA2a in green. Residues are marked according to the SERCA2a

D numbering.



Figure EV3. Sequence alignment of SERCA1a and SERCA2a.

Alignment of rabbit SERCA1a and pig SERCA2a protein sequences indicating the eight most conserved core sequence regions in P-type ATPases (red squares) (Axelsen & Palmgren, 1998). The post-translational modifications found by MS in this study are marked by blue (reported before) and green (identified in this study) squares.



Figure EV4. Expression levels of SERCA1a and SERCA2a constructs.

Immunoblot with a non-isoform-specific SERCA TRY2 antibody (Mountian *et al*, 2001) depicting similar protein levels of the WT and chimera constructs of SERCA2a and SERCA1a following transient expression in COS cells.