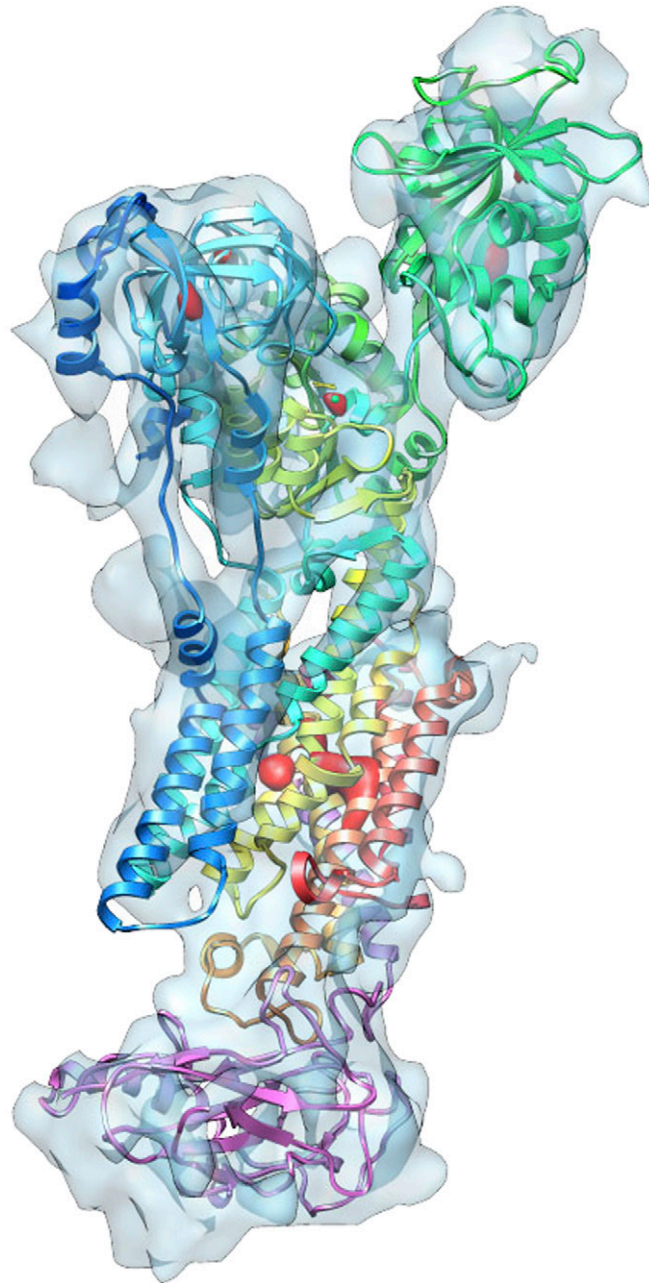


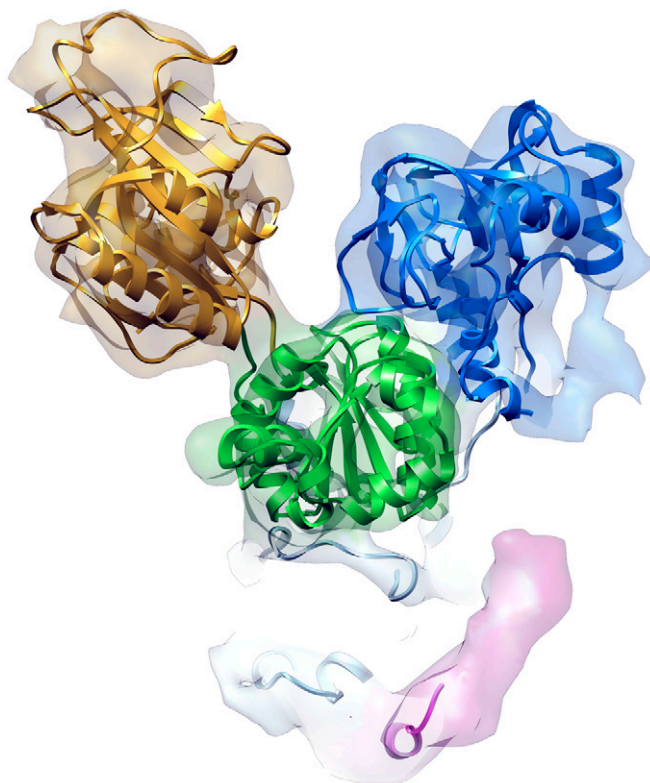
Supporting Information

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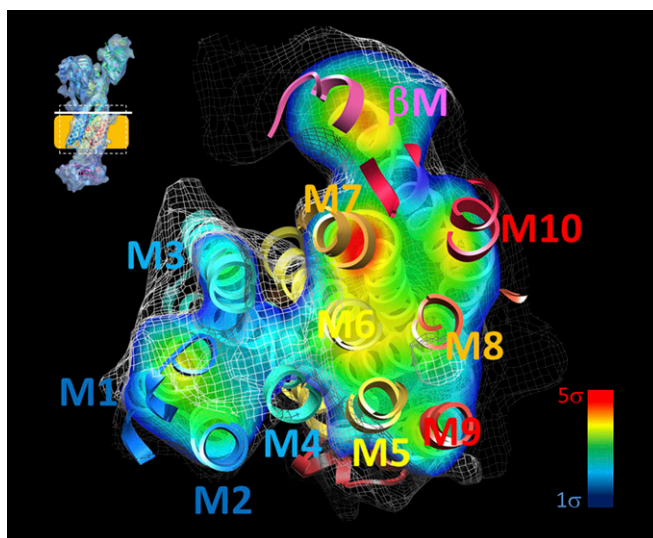
Movie S1. Cryo-EM structure of H^+,K^+ -ATPase in $(Rb^+)E2\sim A1F$ state. Molecular surface shows the EM map of $(Rb^+)E2\sim A1F$ at $1\ \sigma$ (A domain, blue; P domain, green; N domain, orange; TM, light blue; β -subunit, pink) and $5\ \sigma$ (red), with superimposed homology model [color for TM helices (M1–M10) gradually changes from blue to red].

[Movie S1](#)



Movie S2. Rb⁺-induced rearrangement of the cytoplasmic domains. Sequential appearance of E2~AIF (dark colors) and (Rb⁺)E2~AIF (light colors) shows Rb⁺-induced movement of A and N domains relative to the P domain. Structures were viewed from the same direction as in Fig. 7C.

[Movie S2](#)



Movie S3. Horizontal sections throughout the TM region of (Rb⁺)E2~AIF. Cross-sections of the TM region (as shown in Fig. S8) are sequentially displayed. Positions are indicated in the upper left. Color codes are as in Fig. S8.

[Movie S3](#)

Other Supporting Information Files

[SI Appendix \(PDF\)](#)