

Supporting Information

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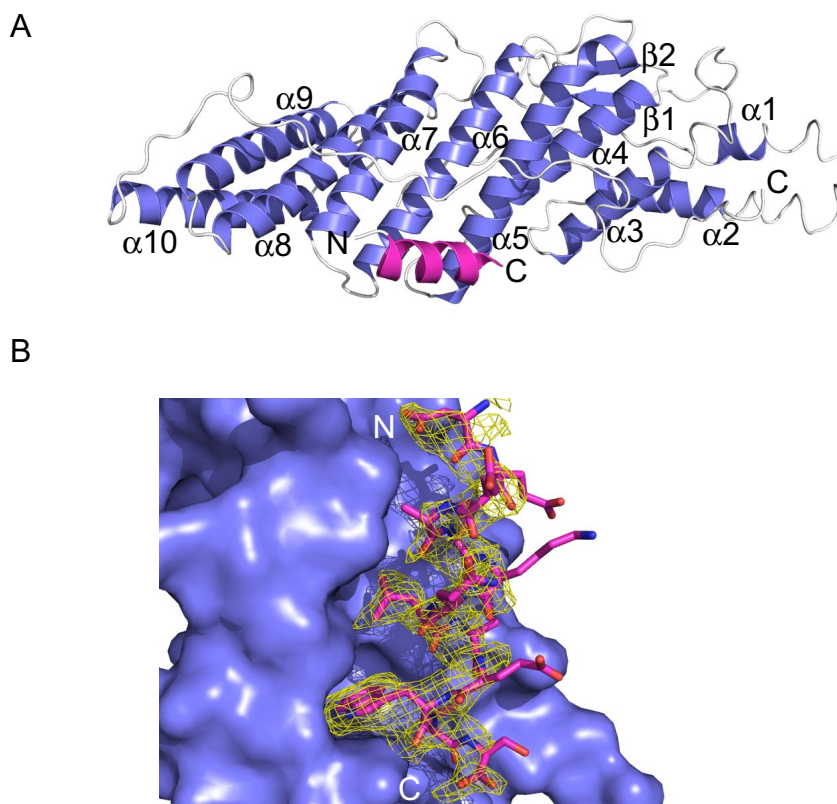


Fig. S1. ALIX_{Bro1} in complex with the C-terminal CHMP4A helix. (A) Ribbon diagram showing the complex between ALIX_{Bro1} and the C-terminal helix from CHMP4A (purple). This figure is oriented so that the CHMP4A helix in Fig. 1 *Lower* is rotated $\approx 90^\circ$ toward the viewer. (B) CHMP4A is represented in sticks against a solid ALIX surface, with the Fo-Fc peptide omit map contoured at $2 \times$ rmsd and displayed over the peptide. To generate the peptide omit map, the peptide was deleted from the model, random shifts (0.3 \AA in x, y, z) were applied throughout and the model re-refined without the peptide in REFMAC. The orientation is the same as in Fig. 3.