



**Fig. S3.** Comparative analyses of the ALIX<sub>Bro1</sub>-CHMP4B interface. (A) Overlay of ALIX<sub>Bro1</sub>-bound CHMP4A-C helices illustrating how hydrophobic residues at the N-terminal ends of the helices can make nearly equivalent contacts despite displacement of the CHMP4B helix (green). (B) Expanded view of the CHMP4B C terminus, highlighting the unique ALIX<sub>Bro1</sub> contacts made by the C-terminal Met-224 residue.