



Figure 4—figure supplement 2. Alignment of L9 and L2 from the crystal structure (L9, orange; L2, wheat; PDB code: 2AW4) and their counterparts from the cryo-EM reconstruction (L9, blue; L2, lightblue; PDB code: 5AFI). The cross-links are mapped to both structures and shown as orange and blue dashed lines, respectively. Many of the cross-linked lysine residues are far in space according to the crystal structure, but much closer according to the cryo-EM structure.