



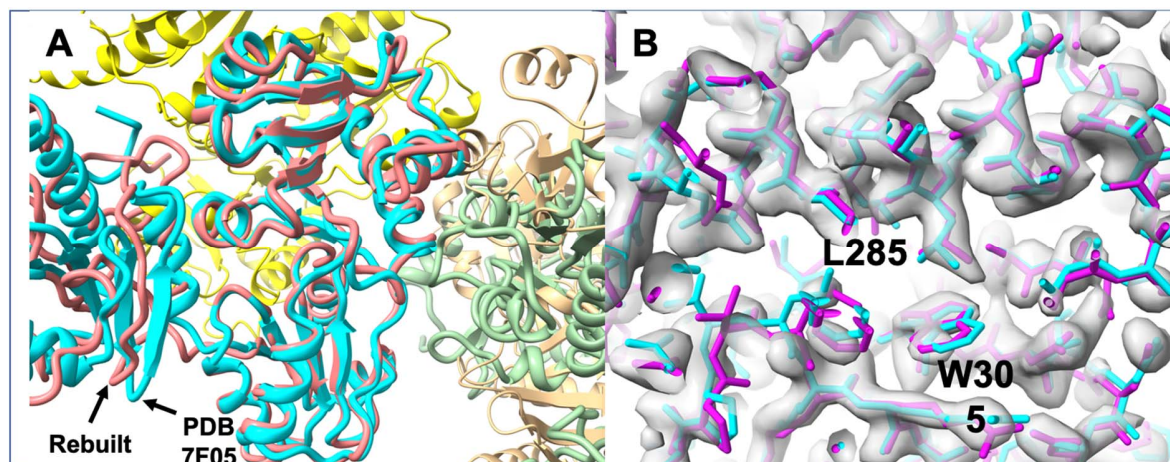
STRUCTURAL  
BIOLOGY

**Volume 79 (2023)**

**Supporting information for article:**

**Accelerating crystal structure determination with iterative AlphaFold prediction**

**Thomas C. Terwilliger, Pavel V. Afonine, Dorothee Leibschnner, Tristan I. Croll, Airlie J. McCoy, Robert D. Oeffner, Christopher J. Williams, Billy K. Poon, Jane S. Richardson, Randy J. Read and Paul D. Adams**



**Figure S1** A. Rebuilt model for PDB entry 7f05 (brown, chain B, green, chain C) and deposited model (light blue, chain A, yellow, chain B, light brown, chain D, chains superposed with space group symmetry on rebuilt model). B. Density-modified electron density density map after rebuilding, rebuilt model (magenta) and deposited model (light blue) shown in the region of chain B of rebuilt model (region of chain A of deposited model).