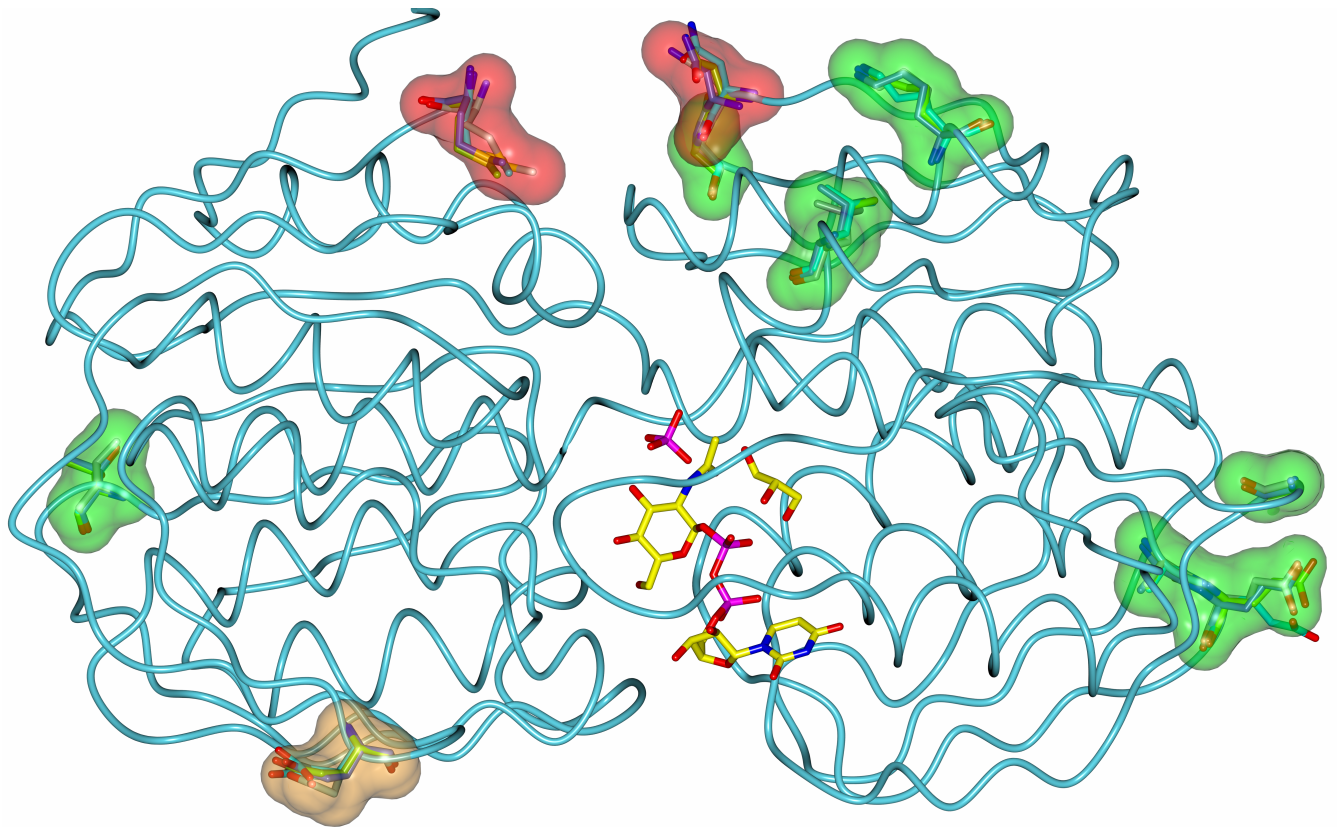


Supplementary Data 1.



SFig 1.

The four protein chains in the crystallographic asymmetric subunit have a C-alpha RMSD of 0.26 Å when superposed. Here, residues which have a deviation greater than 0.3 Å, are highlighted. Residues highlighted with a green transparent surface (K59, G69, L87, G150, A177, E178 and T286) differ by 0.4 Å. One residue, shown in orange transparent surface deviates by 0.5 Å (E349). Residues highlighted with a red transparent surface (M1 and N68) have C-alpha RMSDs between 0.7-0.8 Å. Side chains for each protein chain are shown as well. UNAG, glycerol, and phosphate are shown in yellow.