



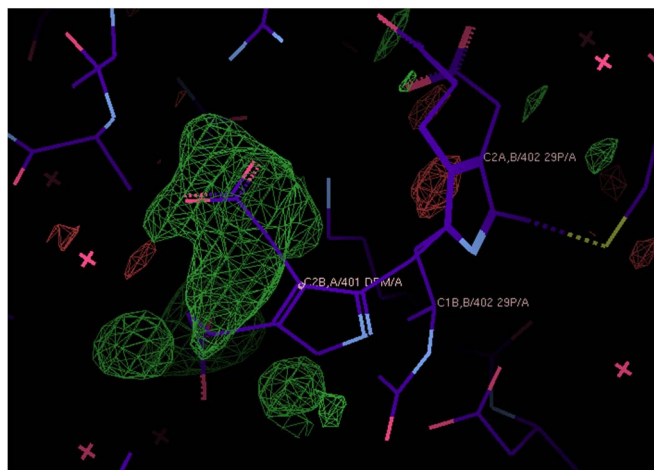
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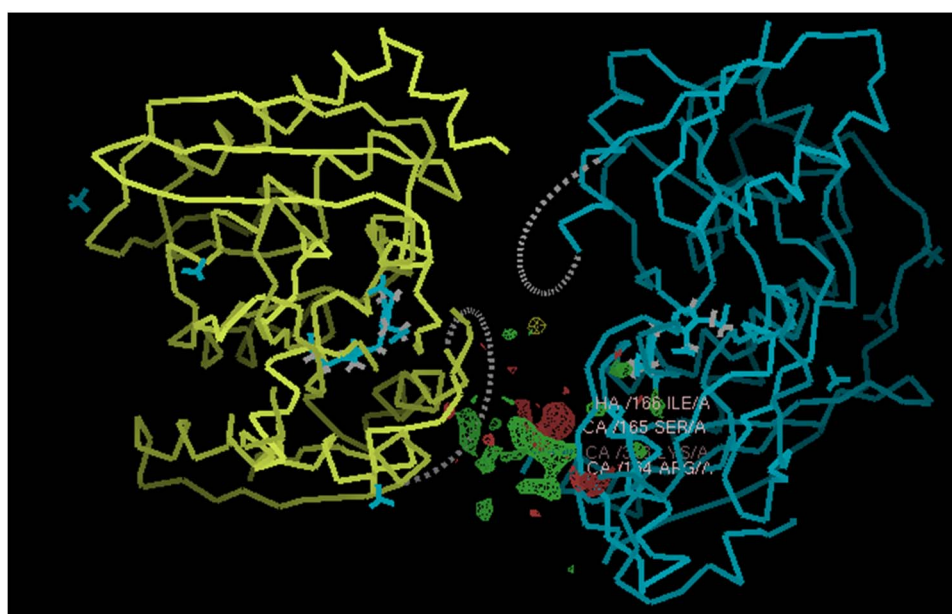
**Supporting information for article:**

**The crystal structures of the enzyme hydroxymethylbilane synthase, also known as porphobilinogen deaminase**

**John R. Helliwell**



**Figure S1** (Fo-Fc) difference electron density at DPM401A of PDB 4mlq contoured at  $3\sigma$ . Azim et al 2014 explain though that “the structure of the enzyme in which the cofactor has undergone more extensive oxidation is presented in 4mlq compared with 4mlv”.



**Figure S2** (Fo-Fc) difference electron density top peak ( $7.0\sigma$ ) for PDB 5m6r contoured at  $3\sigma$ .