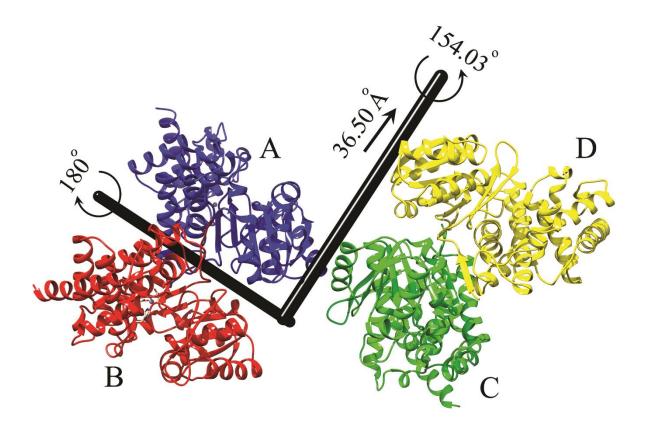


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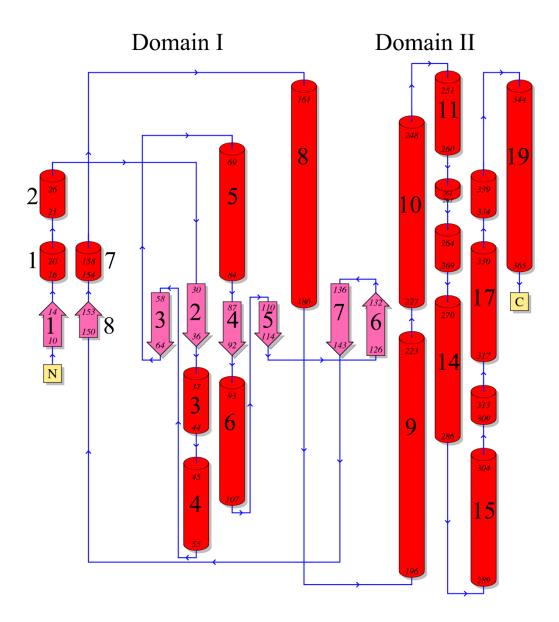
**Supporting information for article:** 

Seeing but not believing: structure of glycerol dehydrogenase initially assumed to be the structure of a survival protein from Salmonella typhimurium

Kaushik Hatti, Yamuna Kalyani Mathiharan, Narayanaswamy Srinivasan and Mathur R. N. Murthy



**Figure S1** Four protomers in the ASU are coloured red, blue, green and yellow. A perfect 2-fold symmetry is observed in dimmers formed between chains A-B and chains C-D. These two dimmers are related by a 154<sup>0</sup> rotation and 36 Å translation. Black rods are placed to indicate the axis of rotation and translation.



**Figure S2** The topology of *Entero*GlyDH. N-terminal domain is comparable to the Rossmann-fold while the C-terminal domain has an antiparallel Greek key helix bundle fold.  $\alpha$  helices and  $\beta$  sheets are numbered independently.