



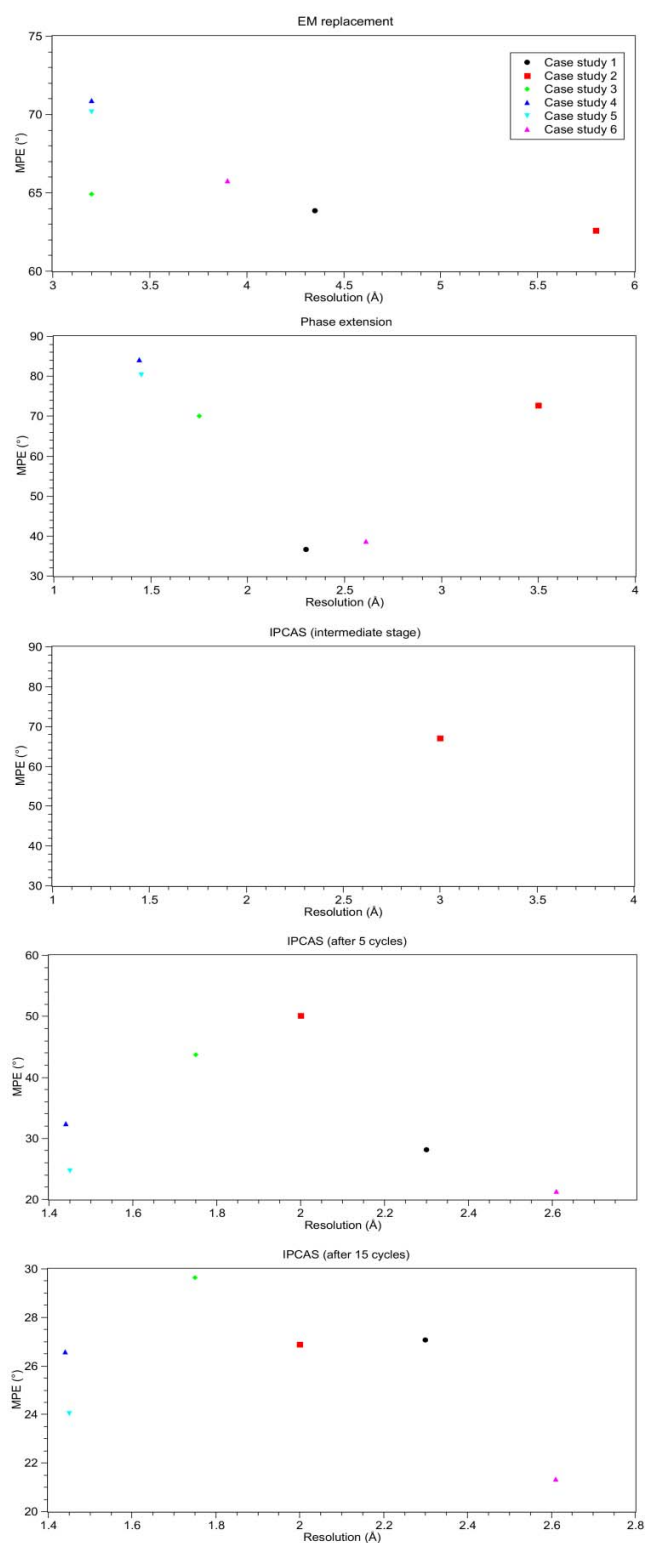
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**Supporting information for article:**

**Using cryo-electron microscopy maps for X-ray structure  
determination of homologues**

**Lingxiao Zeng, Wei Ding and Quan Hao**



**Figure S1** Mean phase errors as a function of resolution for each of the five stages of the process.