



Supplementary Fig. S3. Predicted tertiary structures of the CdCSE proteins in *Camellia drupifera*. The 3D protein models of the 17 CdCSE proteins were generated using the SWISS-MODEL server. Blue ribbons represent α -helices and β -sheets, whilst red coils denote disordered or flexible regions. Structural variations amongst the CdCSE members imply potential functional divergence within the family, which may influence their specific roles.