

Figure 1

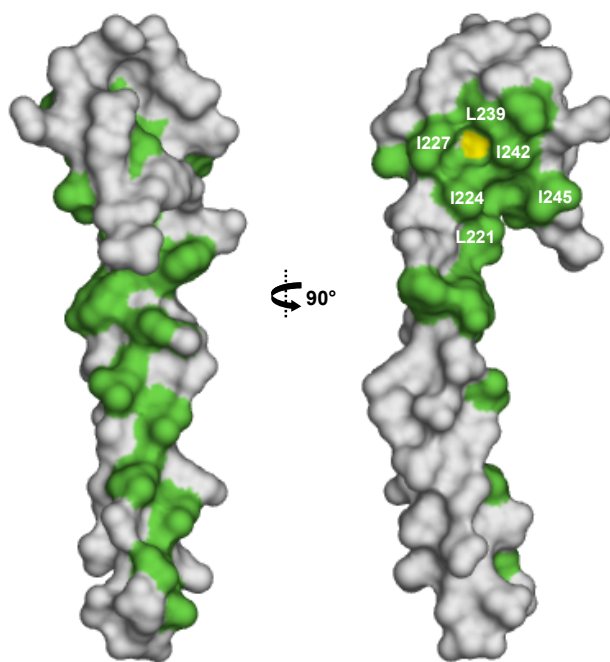


Fig. 1. Monomer A of EB1-C in surface representation. Conserved hydrophobic residue side chains engaged at the dimer interface are highlighted in green. Cys228 is shown in yellow.

SUPPLEMENTARY REFERENCE

Engh RA & Huber R (1991) Accurate bond and angle parameters for X-ray protein structure refinement. *Acta Cryst* **A47**: 392-400