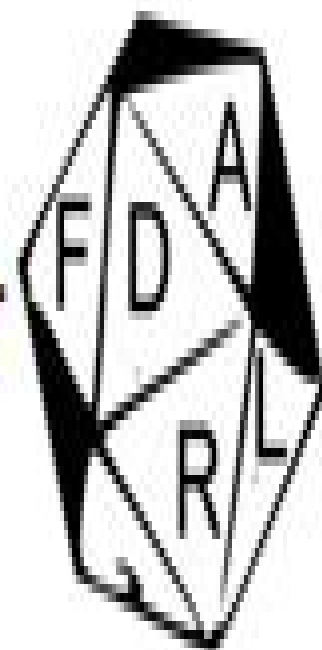
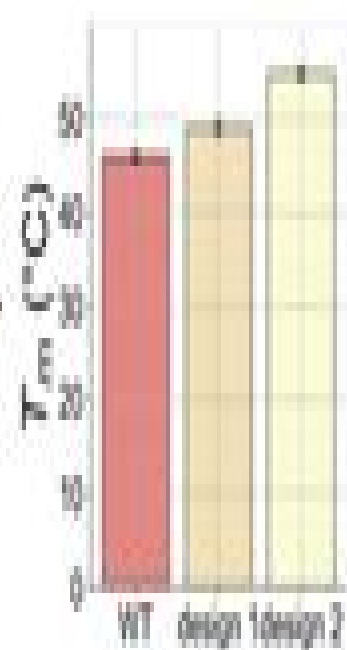
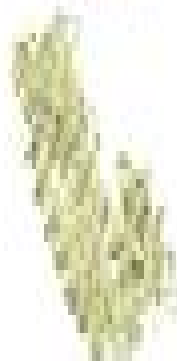


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Estimation of  
 Evolutionary  
 energy  
 $E_{EH}$



Co-evolutionary  
 identified residues

MC sequence  
 shuffling

MD  
 simulations

Experiments

# Recent Developments In Protein Folding Stability And Design

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