

Supplementary Materials: The Crystal Structure of *Bacillus cereus* HblL₁

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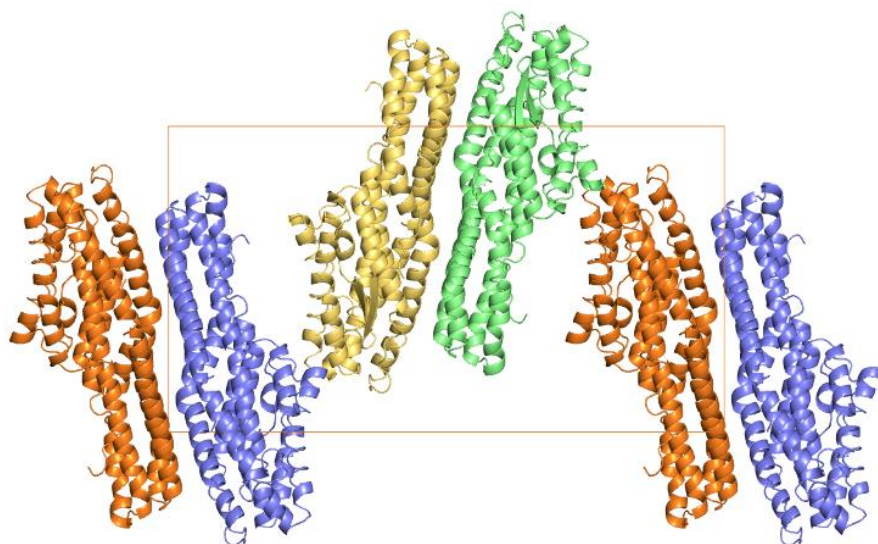


Figure S1. Crystal packing of HblL₁.

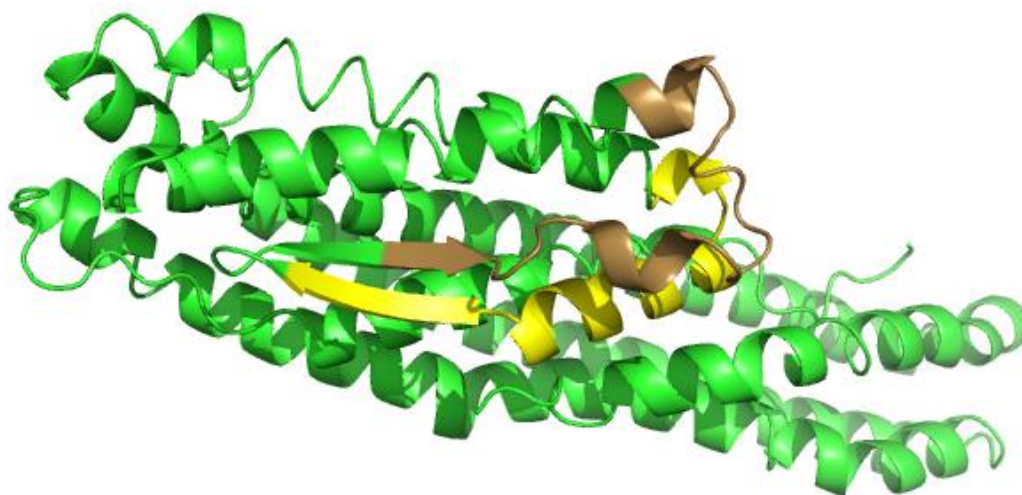


Figure S2. Predicted transmembrane regions in HblL₁. Tm region 1, residues 239-261 is shown in yellow; Tm region 2, residues 268-290 is shown in light brown.

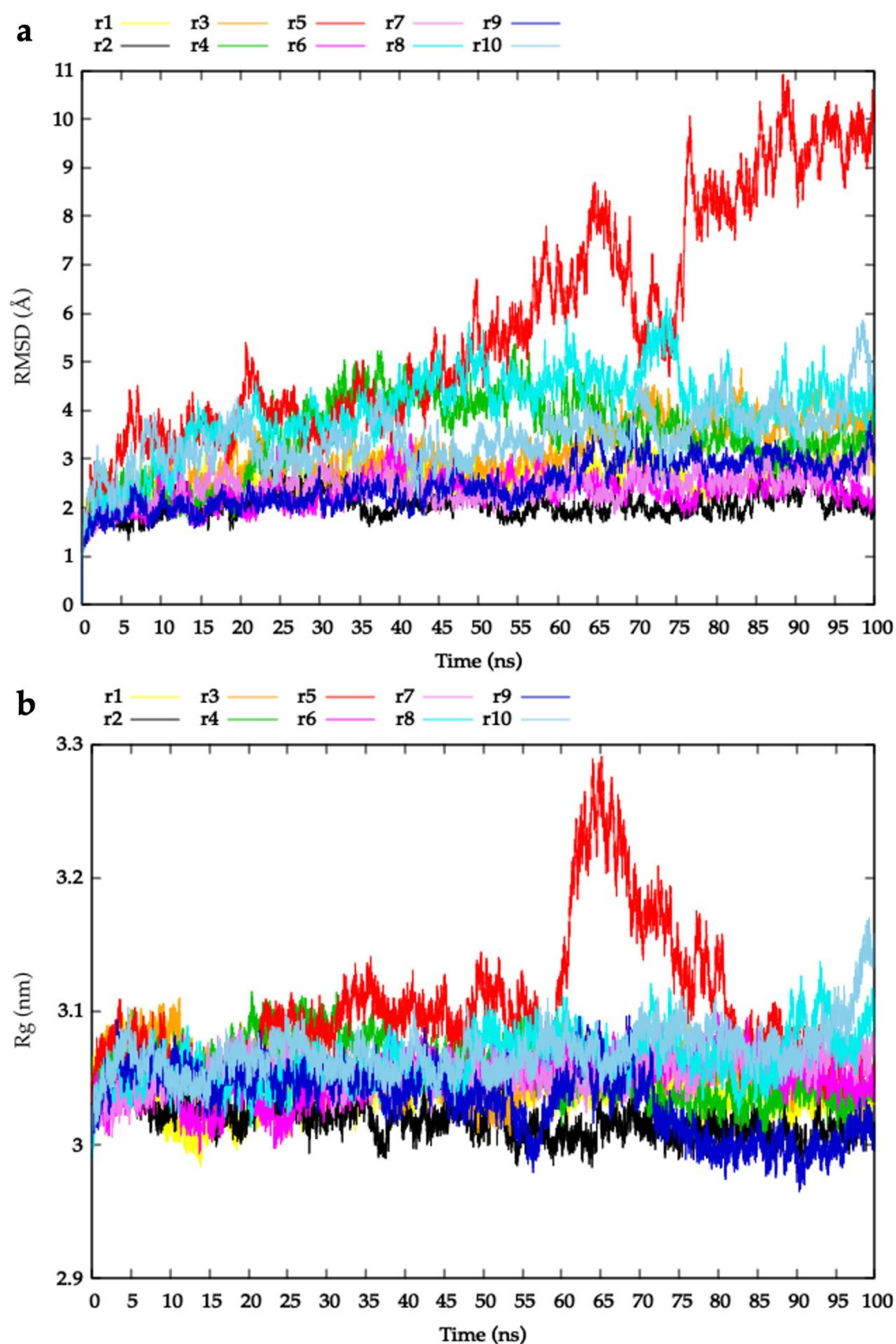


Figure S3. Molecular dynamic analysis of HblL1-HblB complexes. (a) Root-mean-square deviation of the position of backbone atoms in HblL1-HblB modelled complexes r1 - r10 throughout 100 ns MD simulations; (b) Radius of gyration of HblL1-HblB modelled complexes r1 - r10.

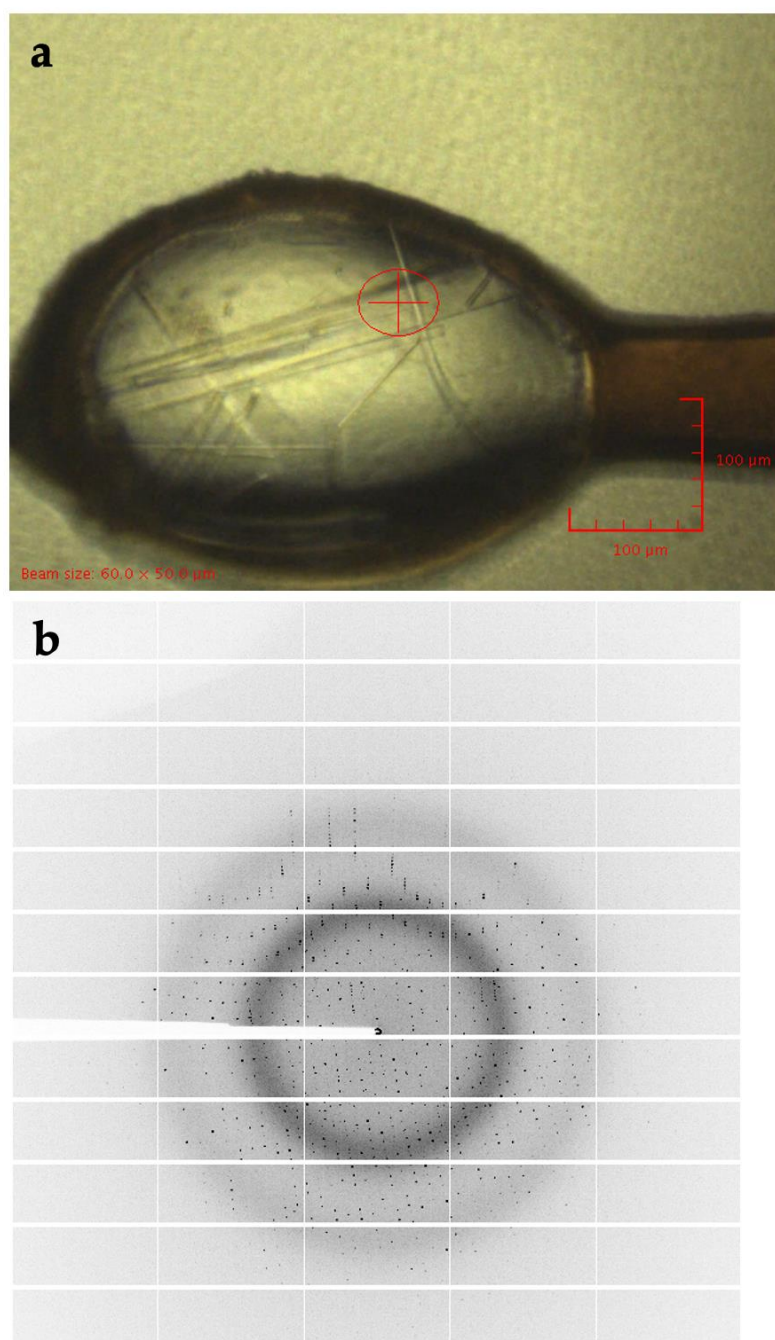


Figure S4. HbLL₁ crystal and diffraction image. (a) single crystal from which data were collected; (b) single frame from diffraction pattern derived from this crystal.