Structural Studies on $\it Cerebratulus \, lacteus \, Mini-Hb \, K(E10)W$ and $\it L(G12)A \, Mutants$

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A very short hemoglobin (CerHb; 109 amino acids) binds $\rm O_2$ cooperatively in the nerve tissue of the nemertean worm *Cerebratulus lacteus* to sustain neural activity during anoxia. The structure of oxygenated wild-type CerHb displays a substantial editing of the globin fold which makes CerHb unique among the known globin fold evolutionary variants [1].

Here we present the crystal structures of two CerHb mutants: Lys(E10)Trp (at 2.3 Å resolution) and Leu(G12)Ala (at 1.6 Å resolution) and its complex with xenon atoms (at 2.3 Å resolution).

The single mutation Lys(E10)Trp, intended to perturb the protein heme binding, has also a dramatic and unexpected effect on the H-bond network stabilizing the $\rm O_2$ ligand, and it makes the protein more susceptible to heme-iron oxidation.

The Leu(G12)Ala mutant and its complex with xenon atoms map a wide protein matrix tunnel connecting the distal site to a surface cleft between the E and H helices, thus suggesting a novel ligand access to heme.

[1] Pesce A., Nardini M., Dewilde S., Geuens E., Yamauchi K., Ascenzi P., Riggs A.F., Moens L., Bolognesi, M., *Structure*, 2002, **10**, 725.

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