

EXAFS Study of Myoglobin in Solution and in Crystalline Form

P. Eisenberger, B.M. Kincaid, A. Mayer, R.G. Shulman, and
B. Wyluda

Bell Laboratories, Murray Hill, New Jersey 07974, U.S.A. and
University of Bremen, Fachbereich Physik, 28 Bremen 33, FRG

The extended x-ray absorption fine structure (EXAFS) of myoglobin (mb) near the Fe K-edge was measured with synchrotron radiation (SLAC Stanford). A comparison was made between the aquo-met form of mb (sperm whale) in solution and in crystalline form. A solution of ca. 15 mM mb and single crystals of mb were used for this investigation. The analysis of the EXAFS data was performed as described in the references: The fluorescence data were Fourier-filtered to obtain the average distance of the first shell of ligands of the Fe. The unknown phase function of mb was determined empirically from the model system bis-imidazole-hemin. An average Fe-nearest neighbour distance of 2.045 ± 0.02 Å for mb in solution and of 2.035 ± 0.02 Å for mb in crystalline form was evaluated. Within the experimental accuracy these average distances are the same.

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