

What I did last

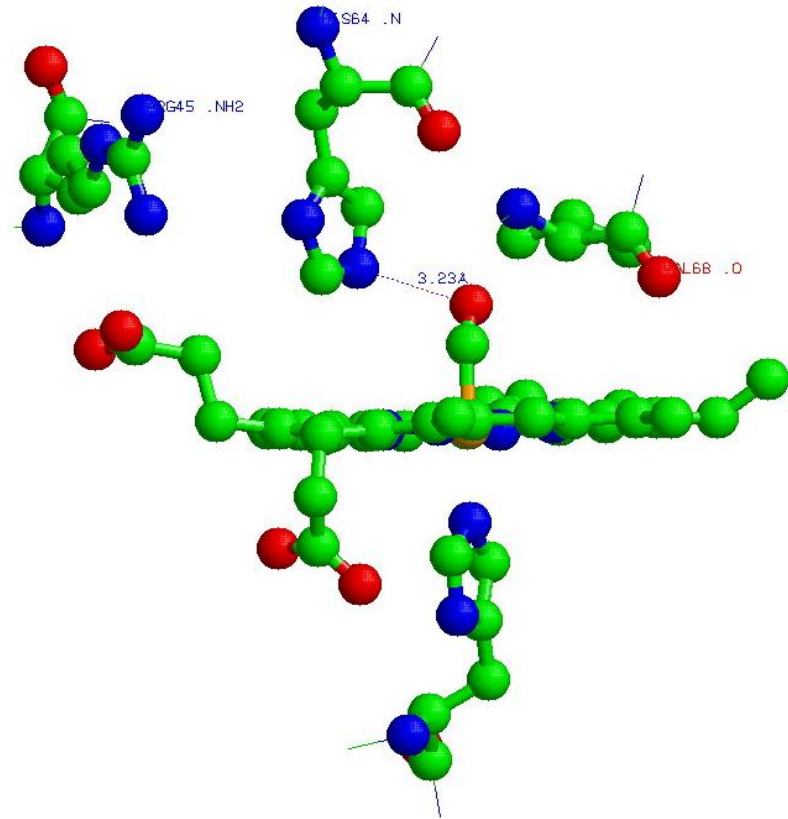
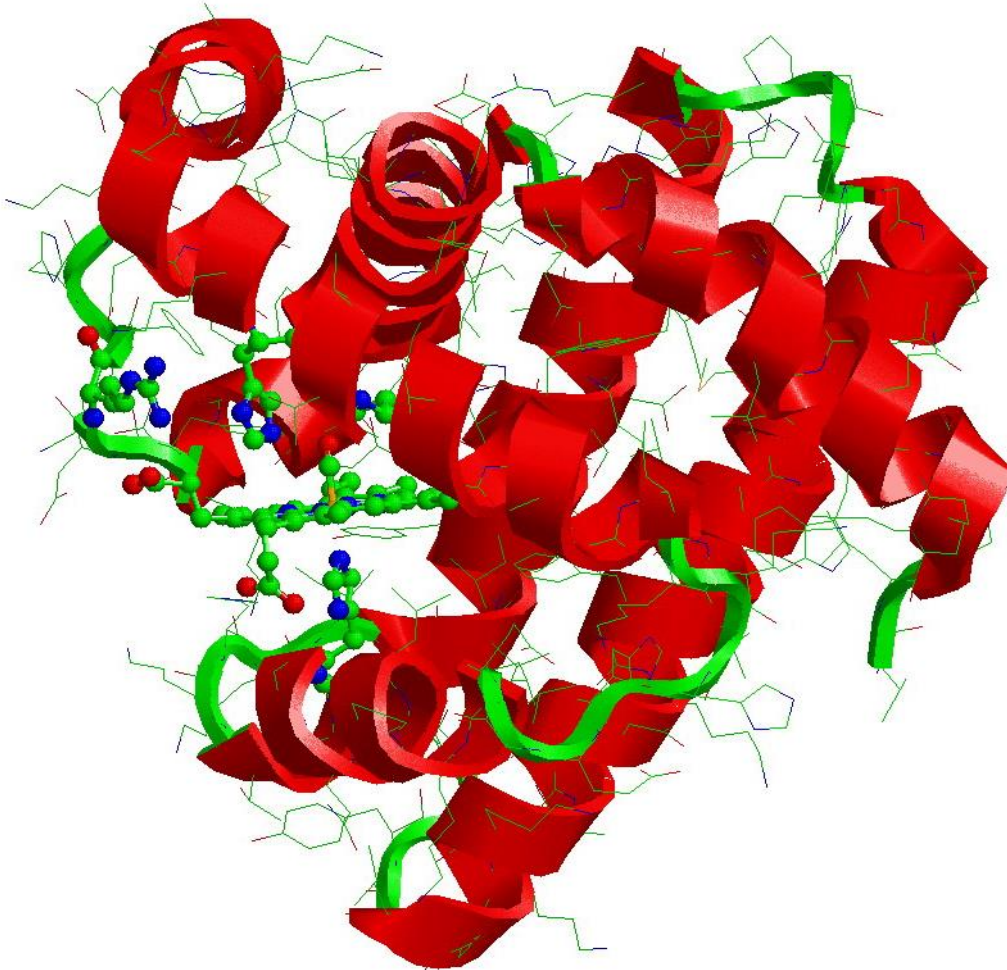
27 years

Solomon S. Stavrov

Sackler Institute of Molecular Medicine, Department of
Human Molecular Genetics and Biochemistry, Sackler
School of Medicine, Tel Aviv University, Israel
stavrov@post.tau.ac.il

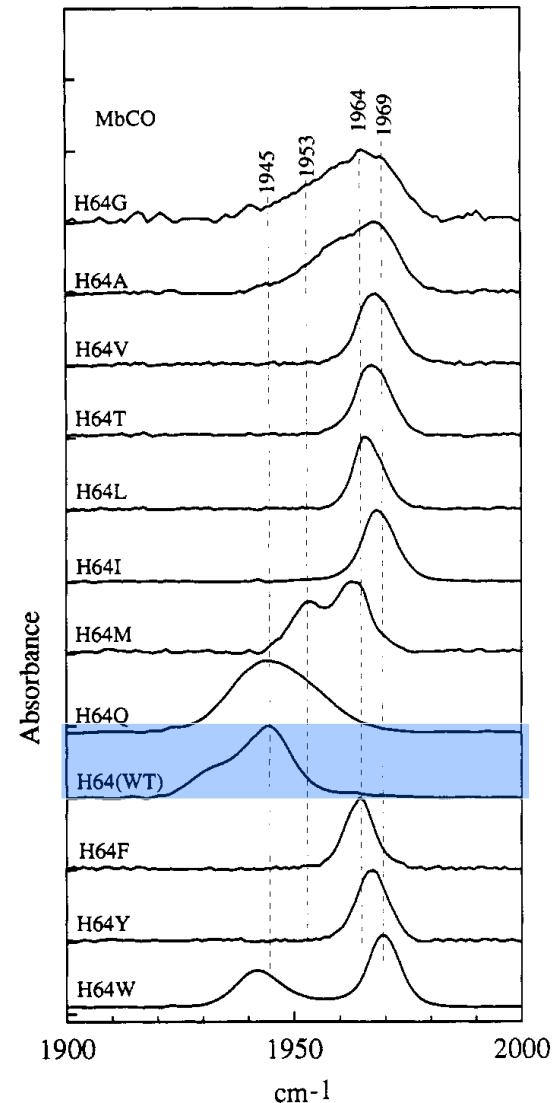
Objects

Carbonmonoxy myoglobin



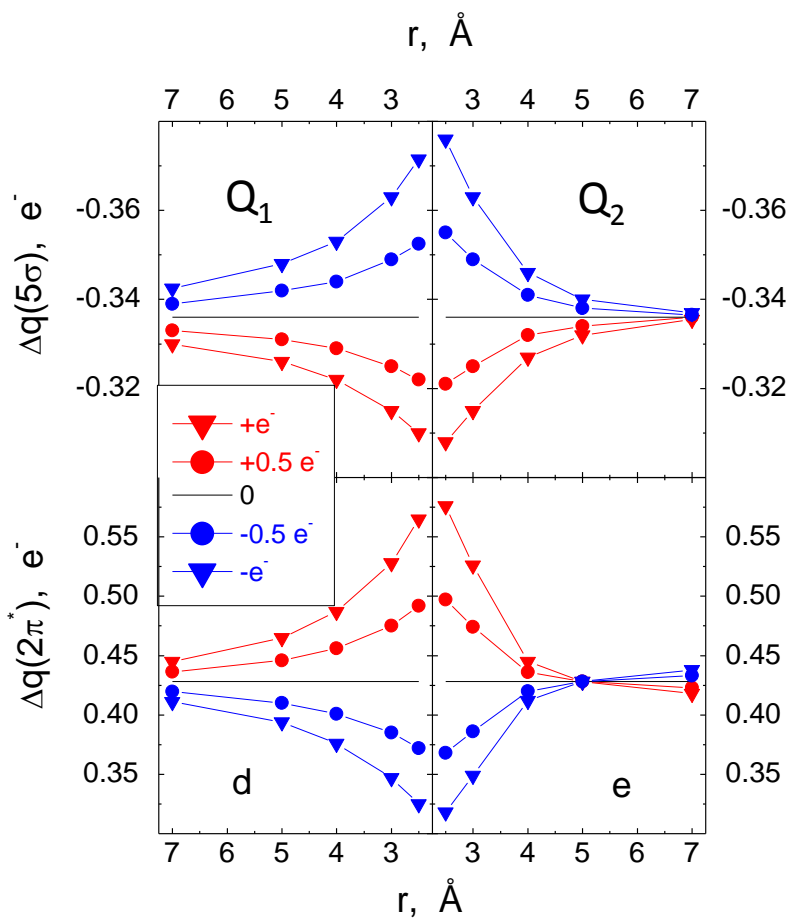
Effect of the His64 mutations on the CO infrared absorption bands of MbCO

(*Li et al., 1994*)

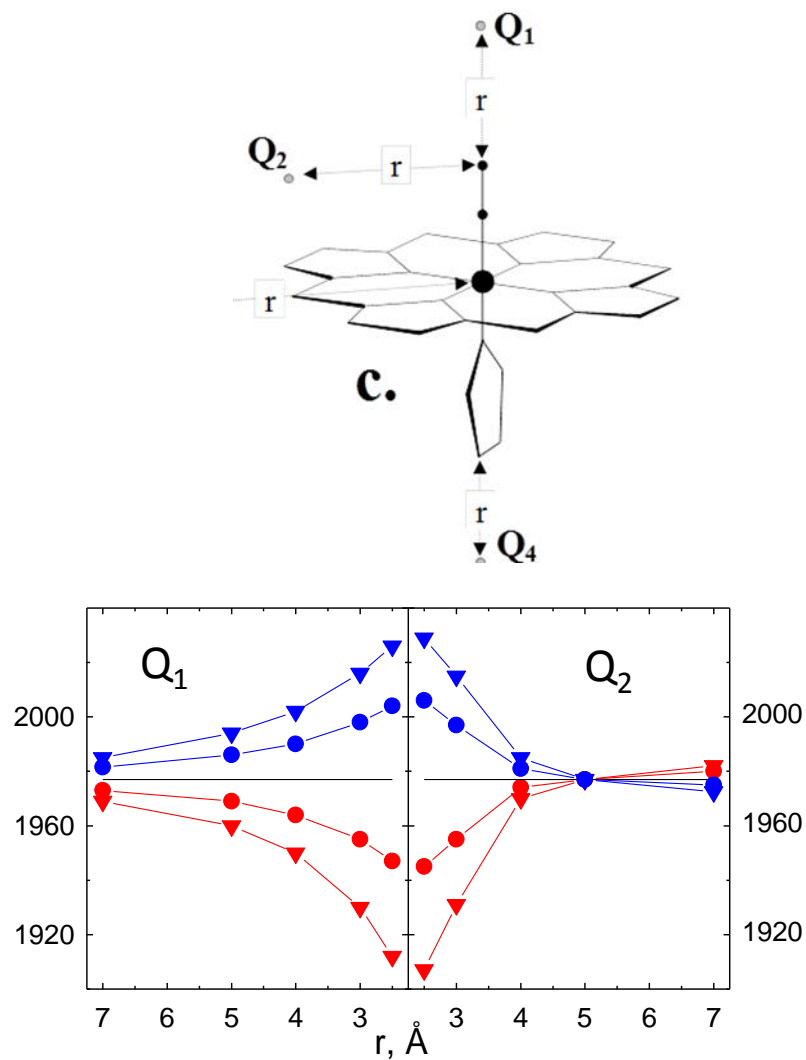


IR spectra of position 64 mutants of sperm whale CO-myoglobin measured at room temperature (~ 22 °C) in 0.1 M potassium phosphate/1 mM EDTA (pH 7.0). The mutant proteins are designated by a single letter abbreviation for the native residue, the position in the primary sequence, and the abbreviation for mutant amino acid, *viz.*, His⁶⁴→Gly is listed as H64G. WT stands for wild-type myoglobin. The spectra of the mutants are shifted to higher frequencies due to the loss of hydrogen-bonding interaction with His⁶⁴.

Effect of the electric field on the C-O vibrational frequency

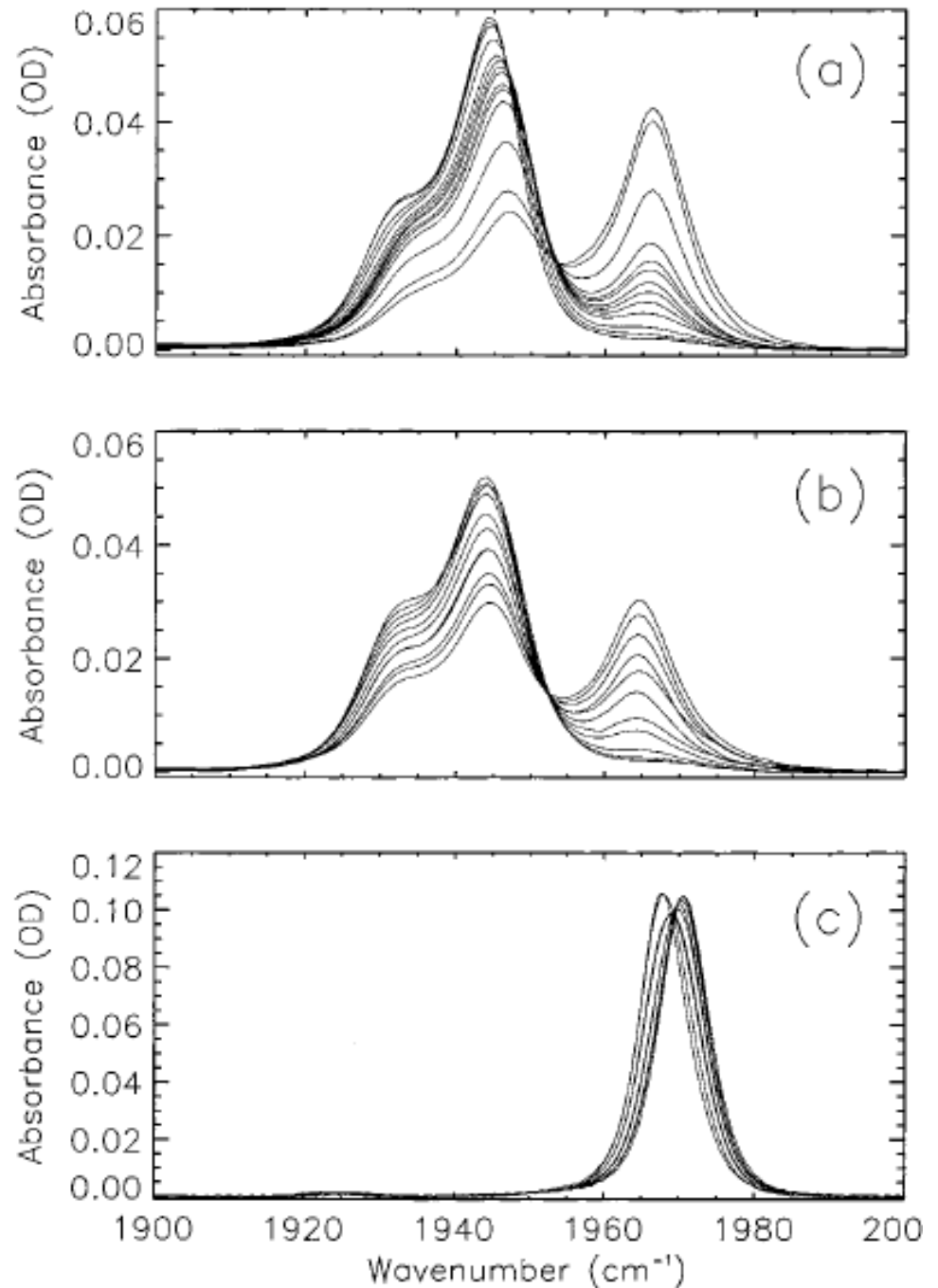


VTA \longrightarrow



Conformational Substates of MbCO Observed in the IR Absorption Spectra

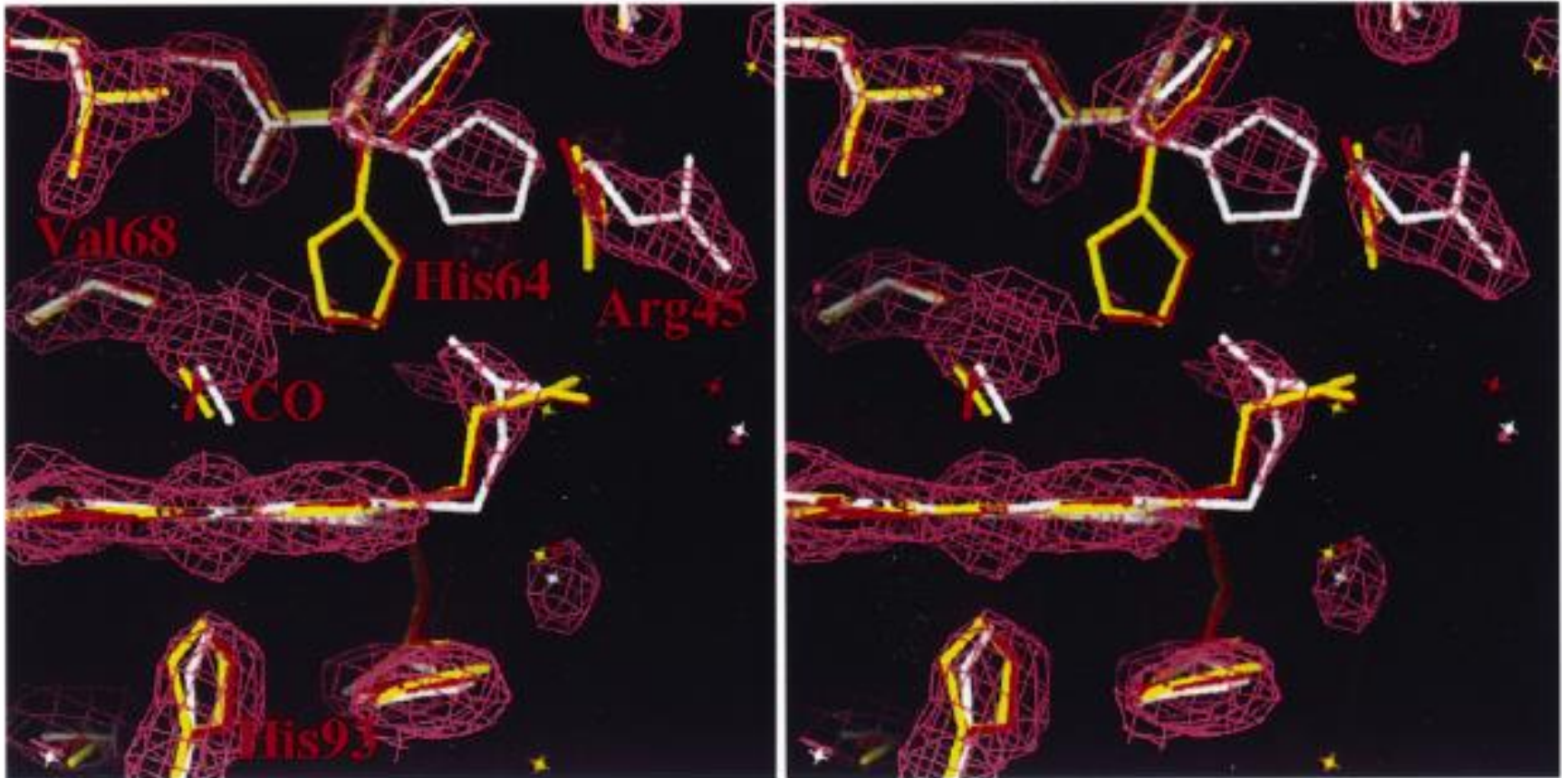
CO stretch infrared spectra in the pH range from 4.2 to 9.5 of (a) swMbCO, (b) proximal mutant H97F, and (c) distal mutant H64L (Muller et al. Biophysical Journal 77(2): 1036-1051)



Effect of pH on the Distal Histidine Position

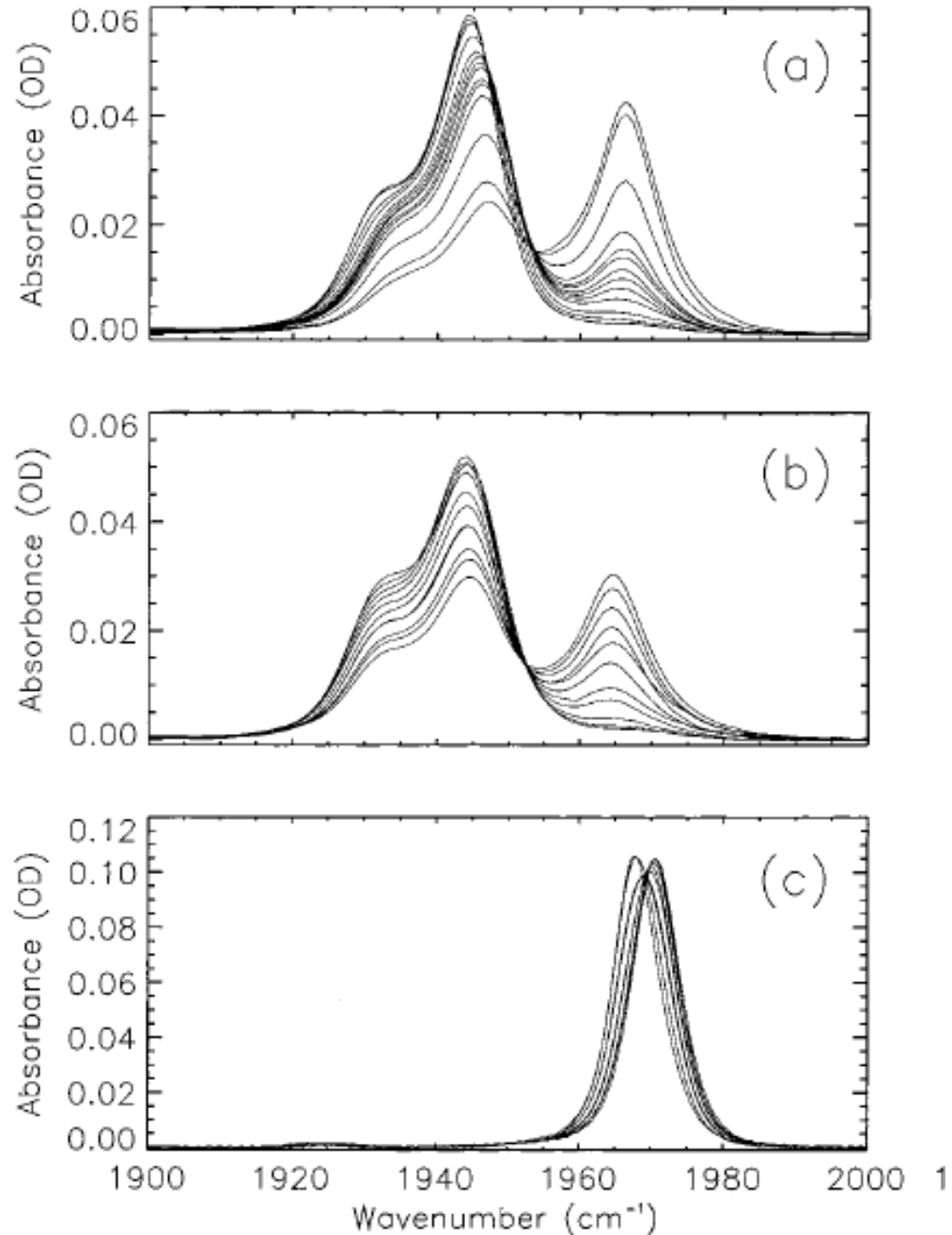
pH 4 (white), 5 (yellow) and 6 (red)

(Yang, F. and G. N. Phillips, 256(1996) 762)



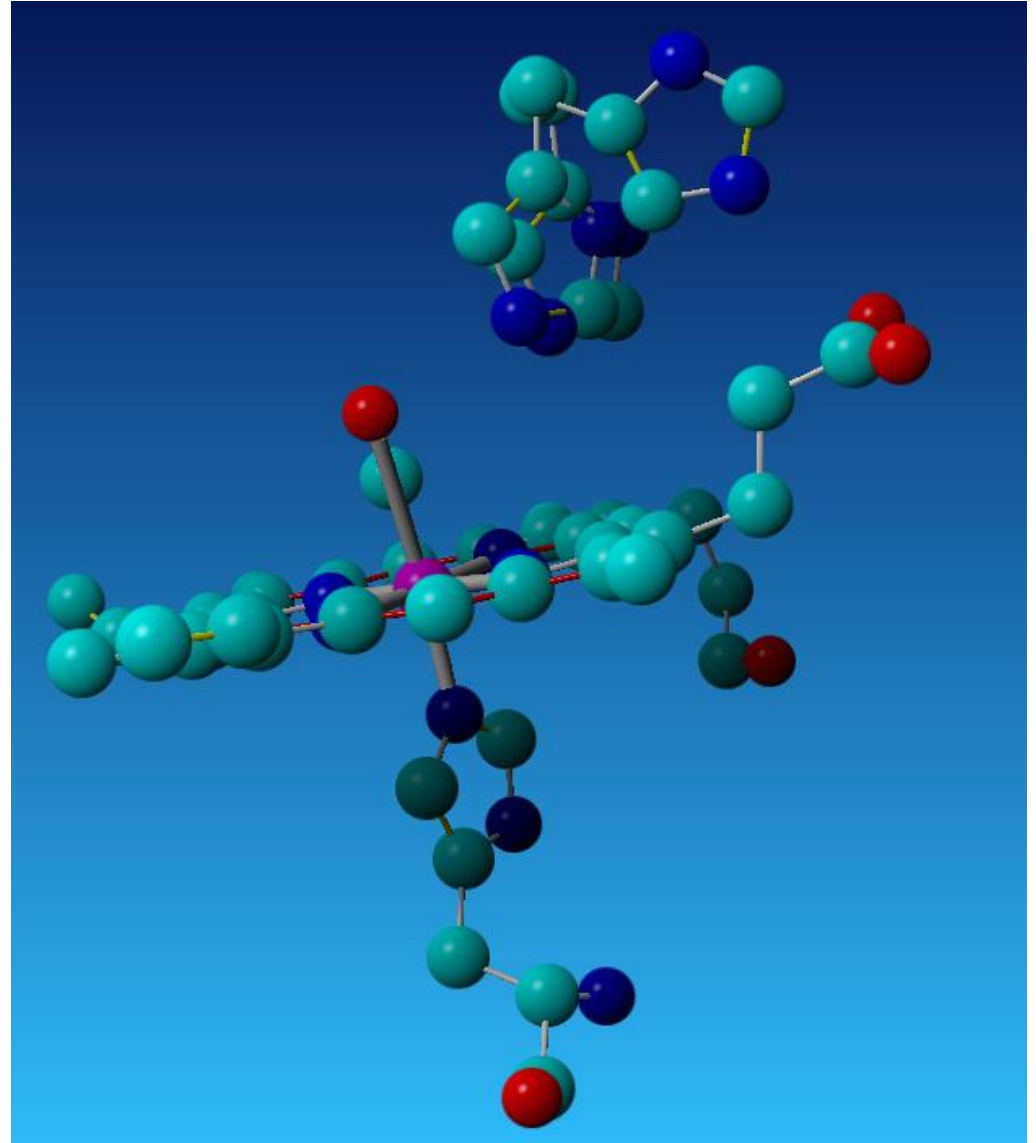
Conformational substates of MbCO observed in the IR absorption spectra

CO stretch infrared spectra in the pH range from 4.2 to 9.5 of (a) swMbCO, (b) proximal mutant H97F, and (c) distal mutant H64L (Muller et al. Biophysical Journal 77(2): 1036-1051)



The Possible Structure of the Conformational Substates

High-resolution (1.2 Å) X-ray study of MbCO showed presence of three different sub-structures of MbCO (*J. Vojtechovsky et al. Biophys. J., 77(1999, 2009) 2153-2174*)



Horseradish peroxidase

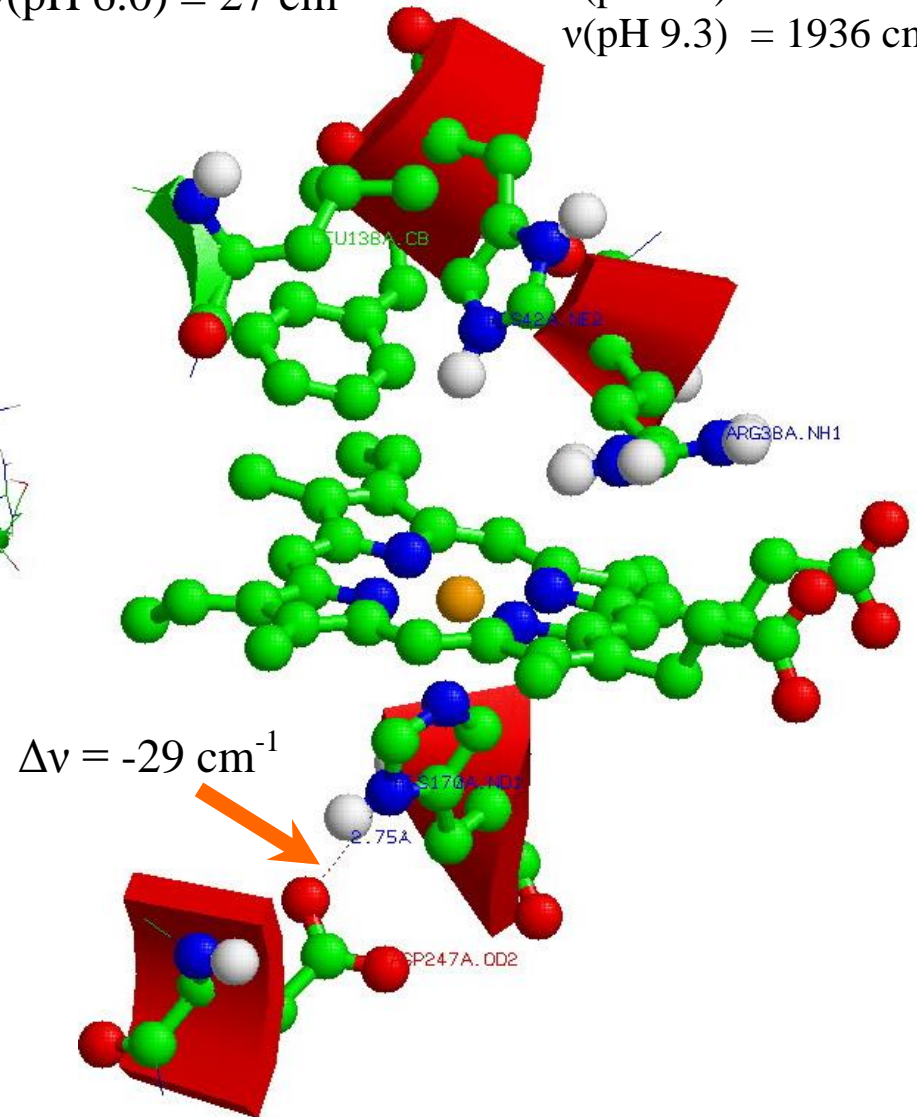
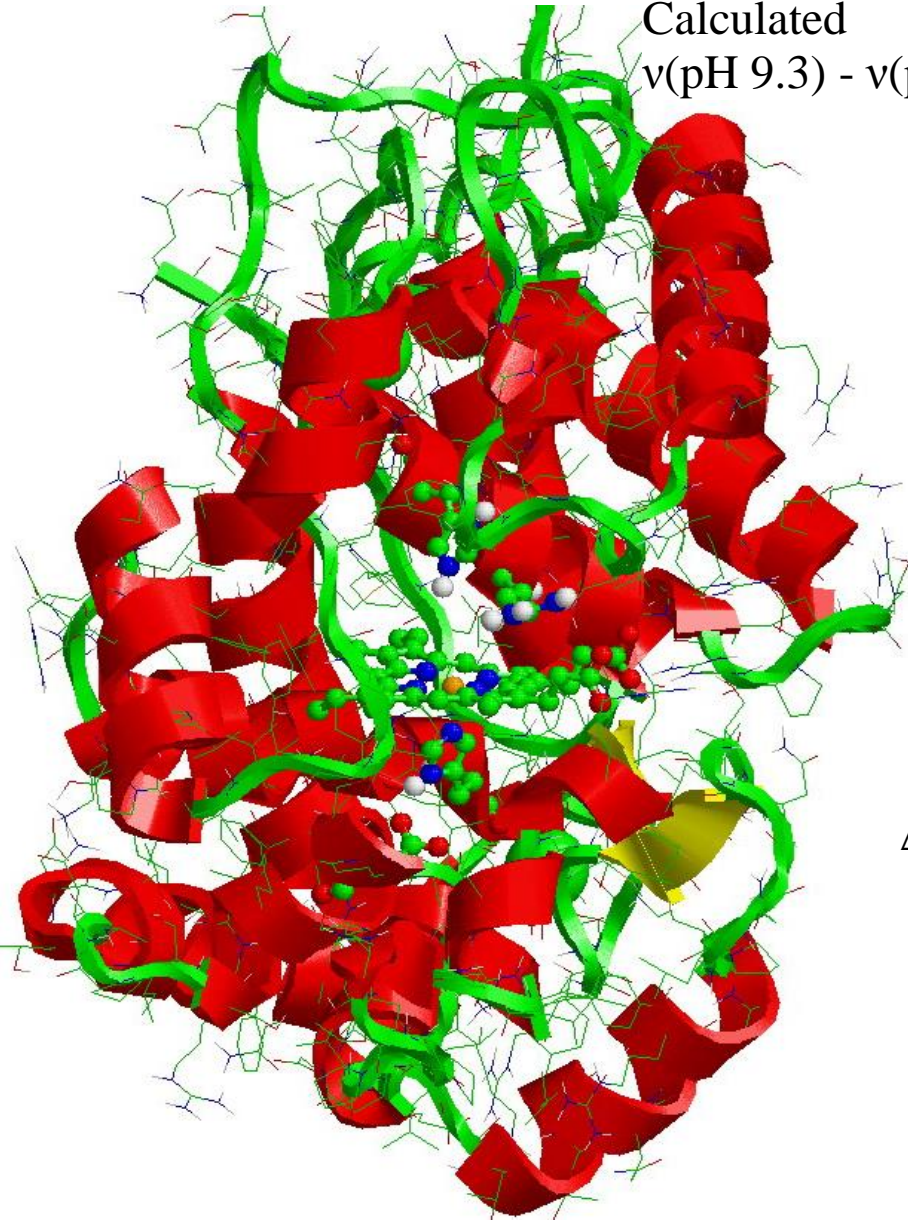
Calculated

$$v(\text{pH } 9.3) - v(\text{pH } 6.0) = 27 \text{ cm}^{-1}$$

Experimental

$$v(\text{pH } 6.0) = 1903 \text{ cm}^{-1}$$

$$v(\text{pH } 9.3) = 1936 \text{ cm}^{-1}$$



Horseradish peroxidase + BHA

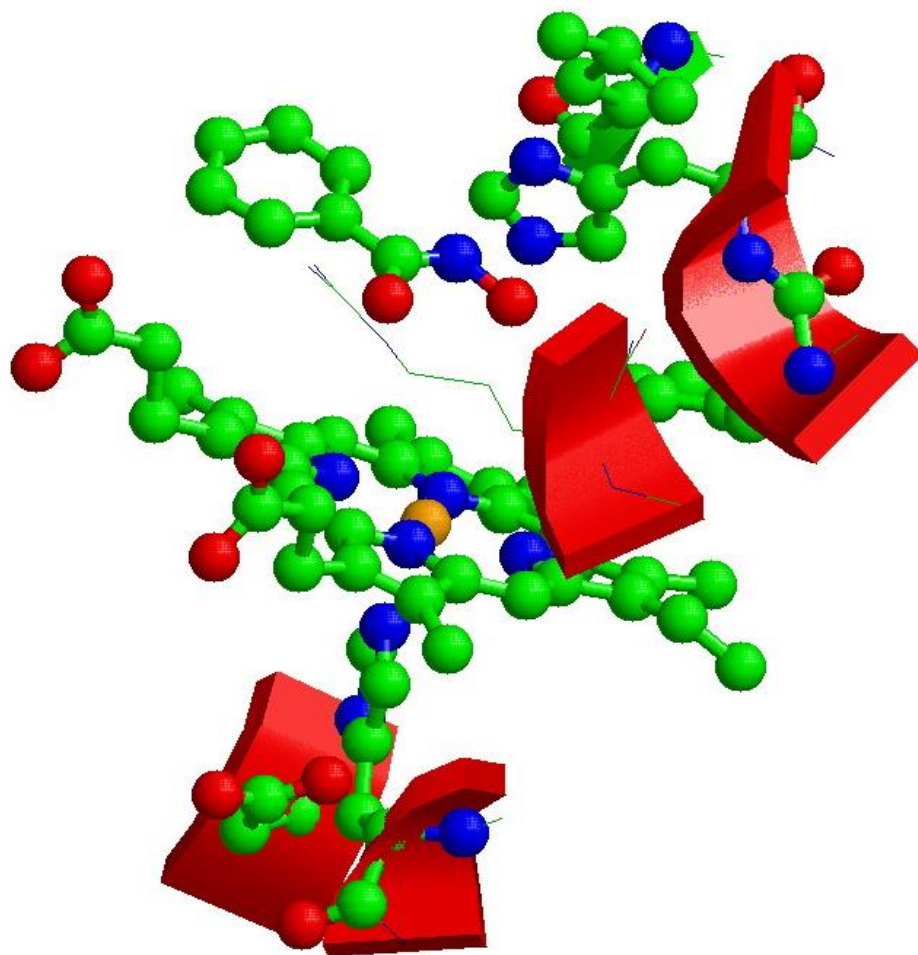
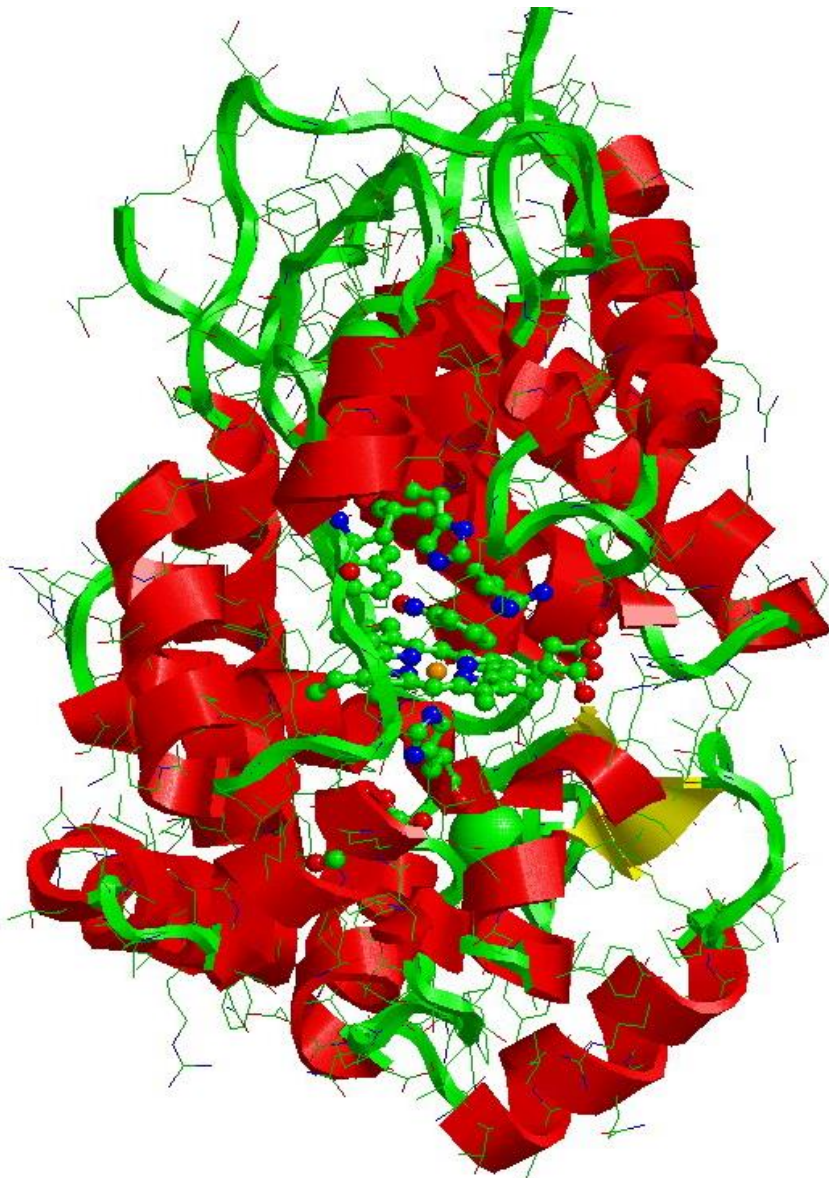
Calculated

$$\nu(\text{HRP+BHA}) - \nu(\text{HRP}) = 6 \text{ cm}^{-1}$$

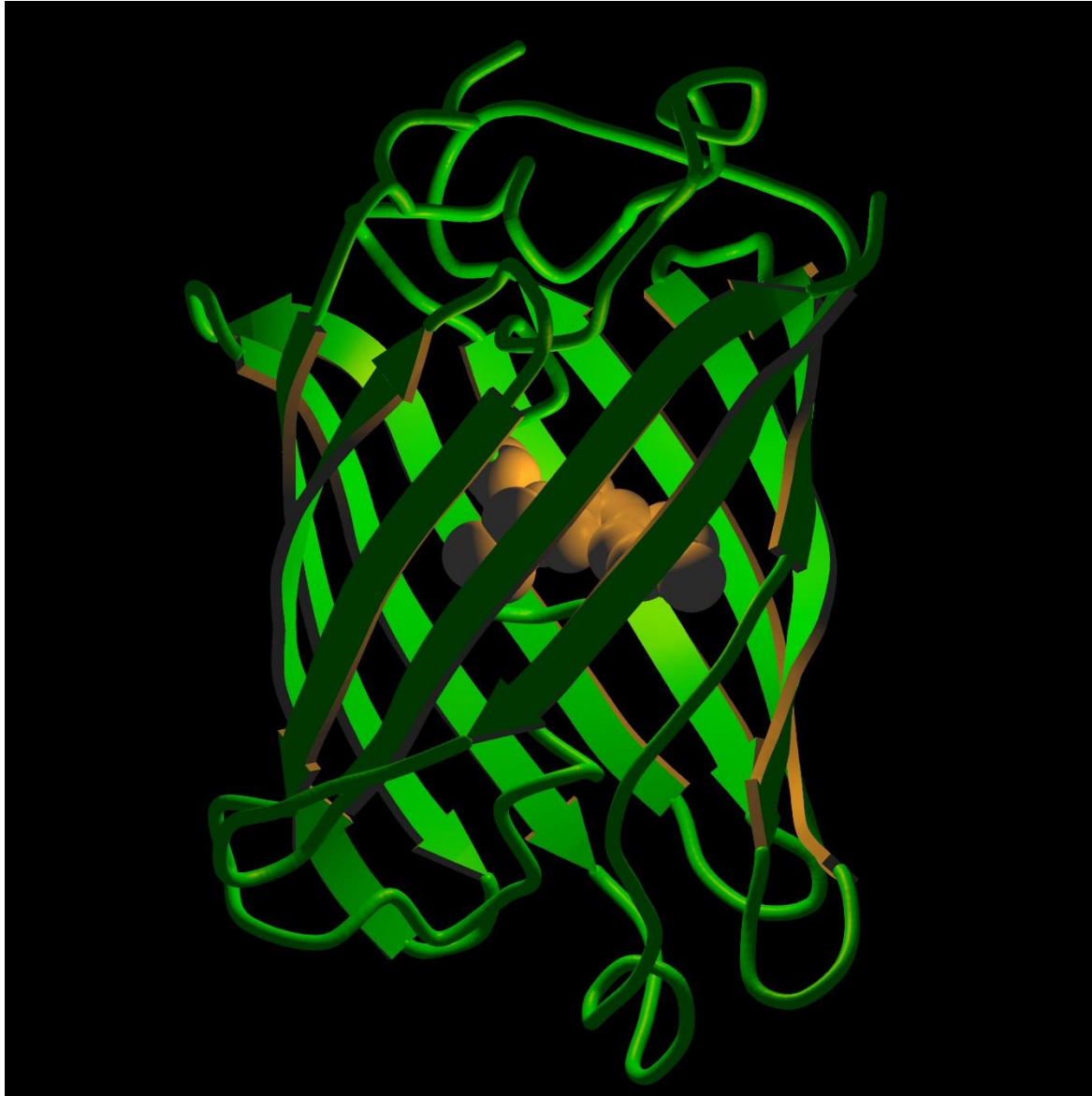
Experimental

$$\nu(\text{HRP}) = 1903 \text{ cm}^{-1}$$

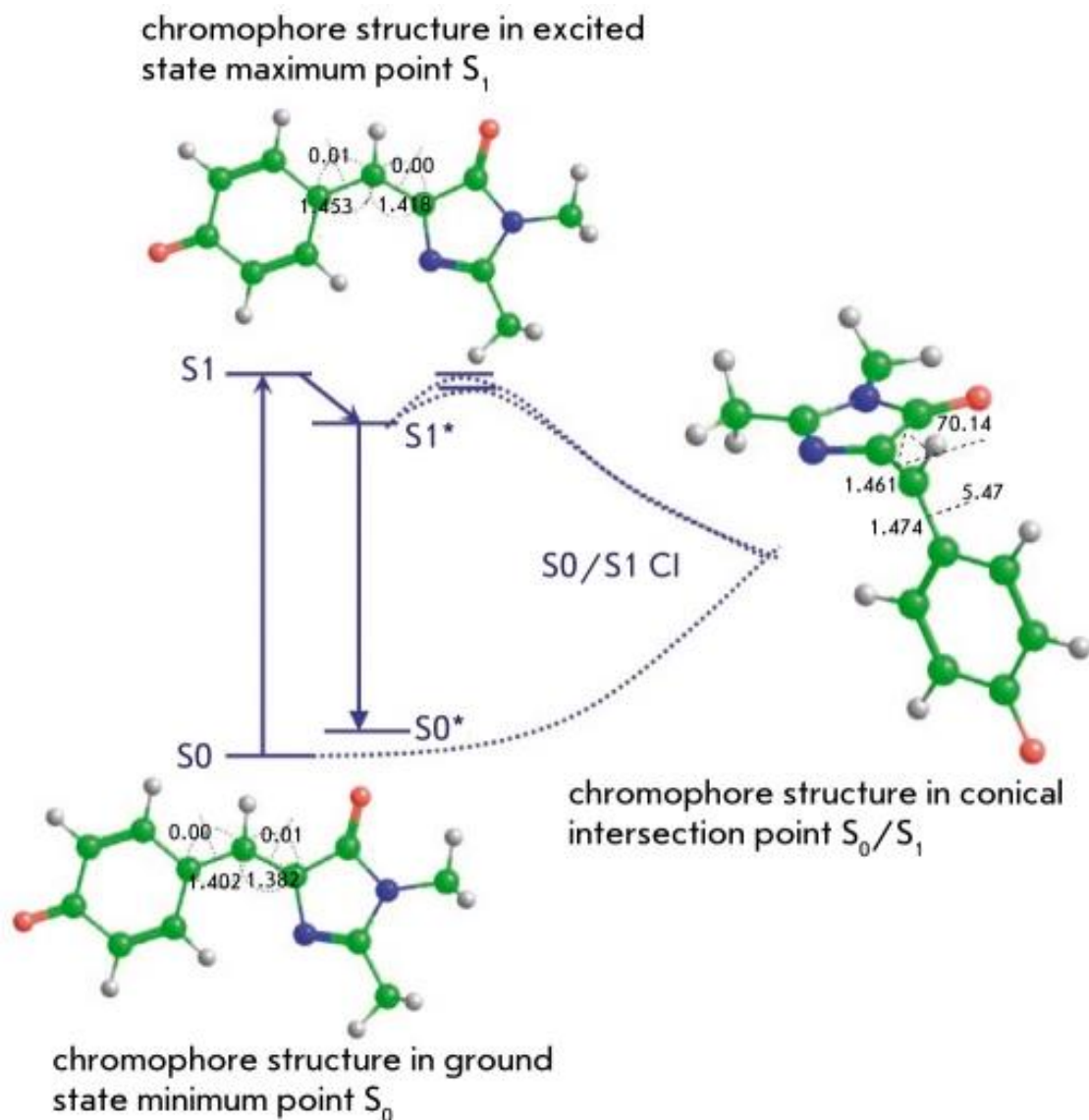
$$\nu(\text{HRP+BHA}) = 1911 \text{ cm}^{-1}$$



GFP



Ground state, excitation and decay of the excited state of the model chromophore



Relationship between the fluorescence spectra of the chromophore and protein

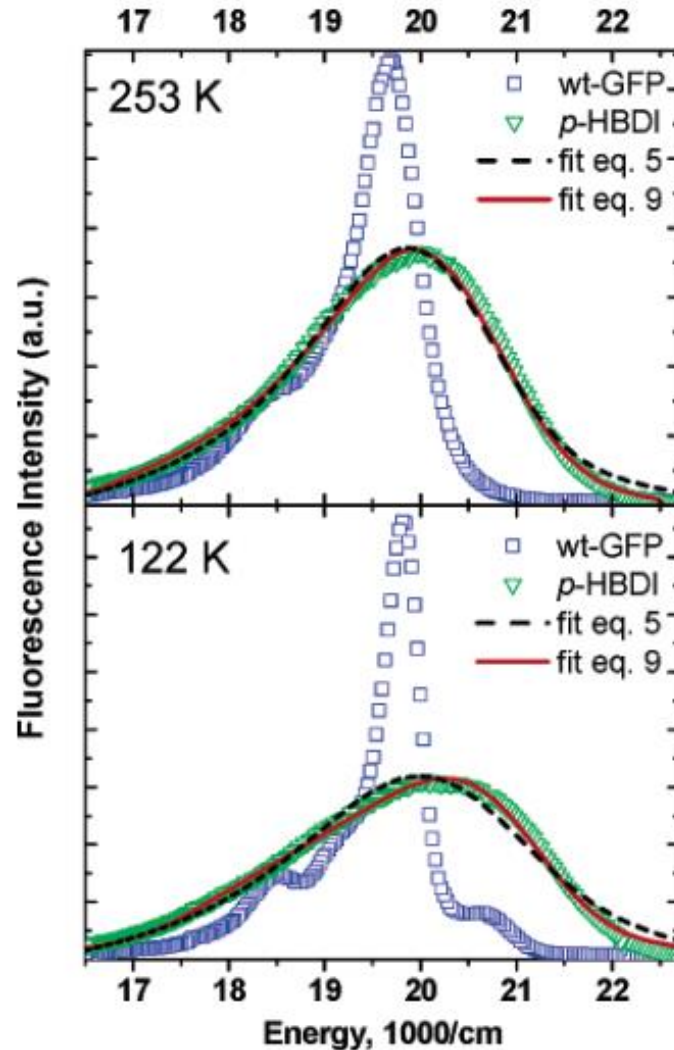
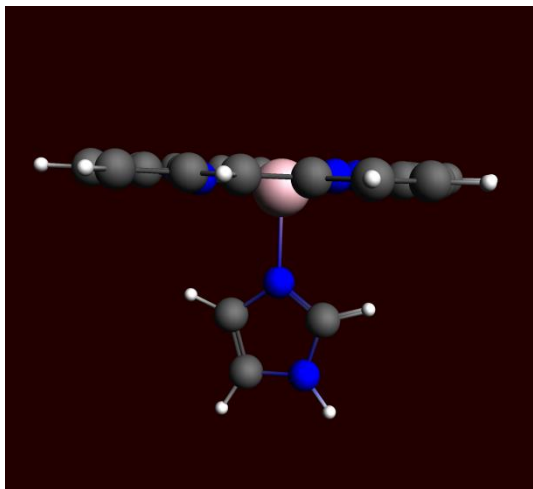


Figure 5. Reconstruction of the model spectra (∇) by utilizing the protein spectra |

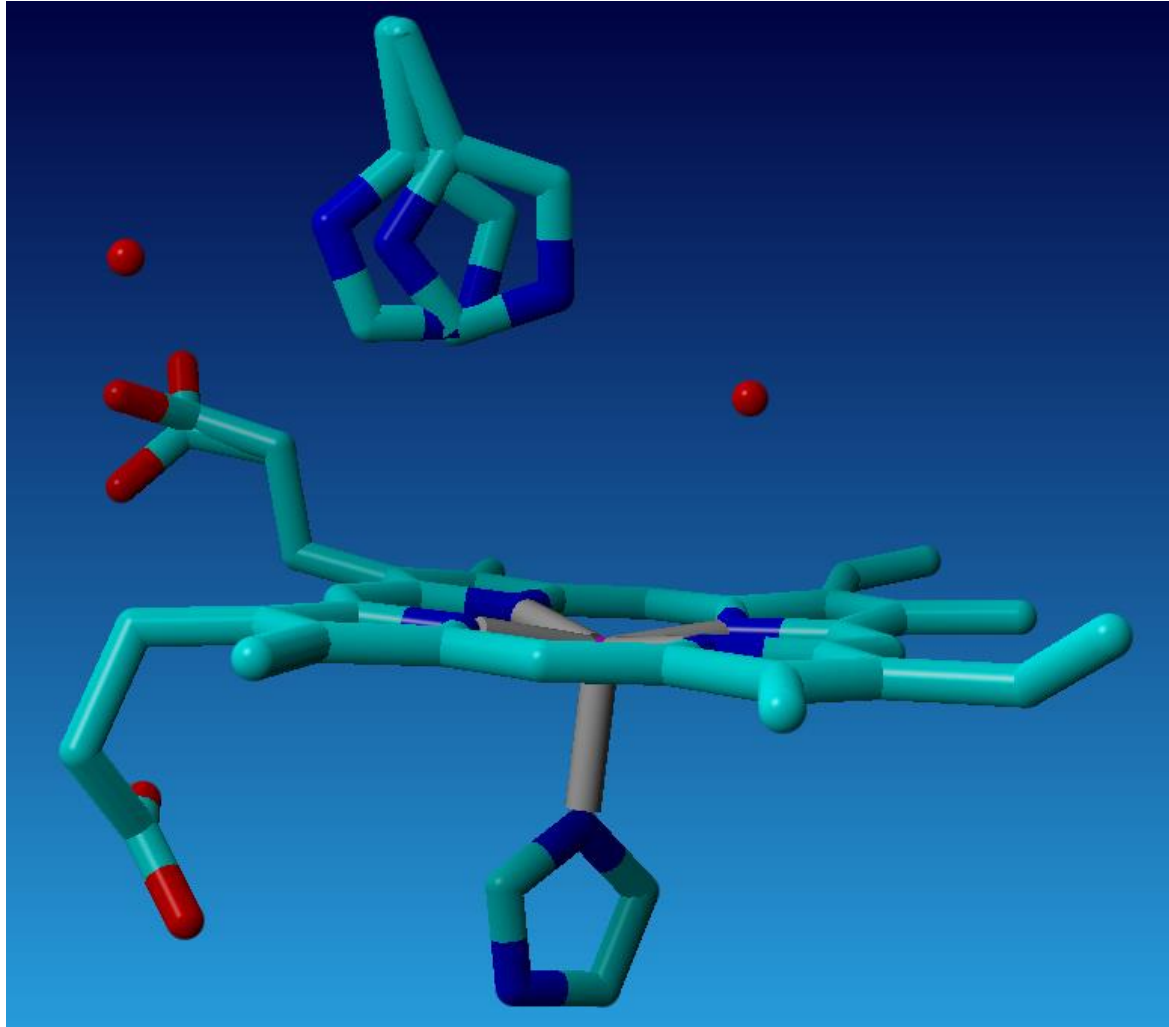
Mossbauer spectroscopy

Effect of the distortions and model electric fields



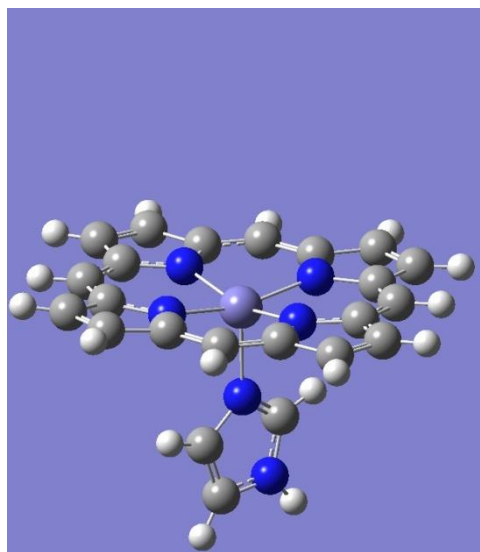
$R - \text{Fe}-\text{N}_{\text{Im}}; r - \text{Fe}-\text{Ct}$	$\Delta E_Q, \text{ mm/s}$	η
$r = 0.32 \text{ \AA}, R = 2.12 \text{ \AA}$ $E = 0$	-2.06	0.77
$r = 0.42 \text{ \AA}, R = 2.12 \text{ \AA}$ $E = 0$	-2.13	0.97
$r = 0.32 \text{ \AA}, R = 2.27 \text{ \AA}$ $E = 0$	-2.18	0.48
$r = 0.32 \text{ \AA}, R = 2.12 \text{ \AA}$ $E_{\parallel} = 0.01 \text{ a. u.}$	-2.06	0.76
$r = 0.32 \text{ \AA}, R = 2.12 \text{ \AA}$ $E_{\perp} = 0.01 \text{ a. u.}$	-1.94	0.91
$r = 0.32 \text{ \AA}, R = 2.12 \text{ \AA}$ $E_{\perp} = -0.01 \text{ a. u.}$	-2.11	0.64

Conformational substates of deoxymyoglobin
Resolution 1.15 Å at room temperature
(*Kachalova et al., Science 284, 1999, 473*)

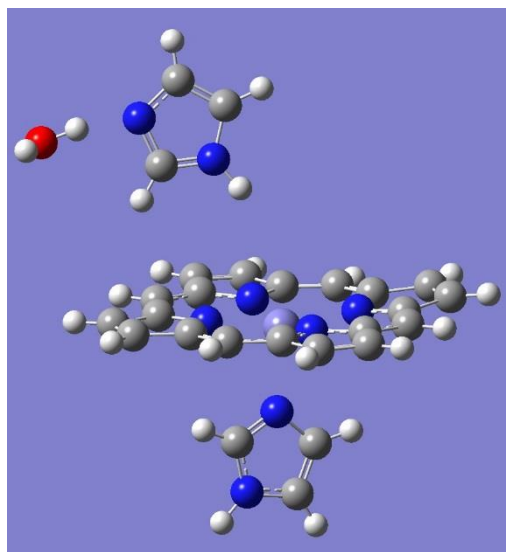


Effect of the distal imidazole electric field

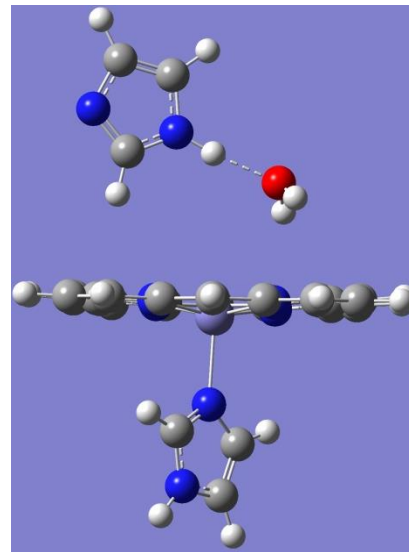
Resolution 1.15 \AA at room temperature (*Kachalova et al., Science 284, 1999, 473*)



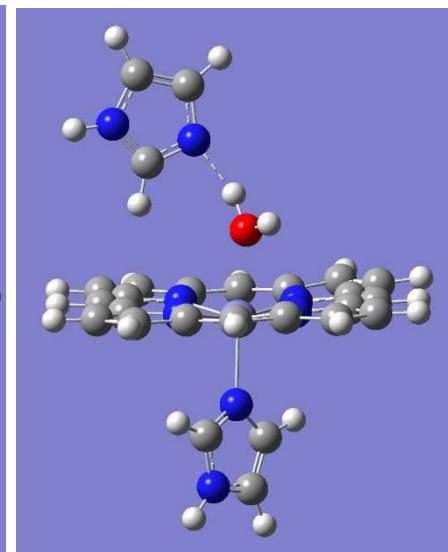
$\Delta E_Q \text{ (mm/s)} = -2.08$



-2.09



-2.18



-2.05

$\Delta E_{Q,\text{exp}}(\text{Mb}) = -2.16 - -2.22 \text{ (mm/s)}$
(Kent, T. A. et al. (1979), BBA, 580(2): 245-258.)

Professional activities

**Theoretical interpretation
of virtually any
spectroscopic studies of
molecules, including
biomolecules**