

# **Redox-linked protein dynamics of cytochrome c probed by time-resolved surface enhanced infrared absorption spectroscopy**

**Nattawadee Wisitruangsakul,<sup>1,3</sup> Ingo Zebger,<sup>1\*</sup> Khoa H. Ly,<sup>1</sup> Daniel H. Murgida,<sup>2</sup> Sanong Ekgasit,<sup>3</sup> and Peter Hildebrandt<sup>1\*</sup>**

<sup>1</sup> Technische Universität Berlin, Institut für Chemie, Sekr. PC 14, Straße des 17. Juni 135, D-10623 Berlin, Germany

<sup>2</sup> Departamento de Química Inorgánica, Analítica y Química Física / INQUIMAE-CONICET, Facultad de Ciencias Exactas y Naturales, Universidad de Buenos Aires. Ciudad Universitaria, Pab. 2, piso 1, C1428EHA-Buenos Aires, Argentina.

<sup>3</sup> Sensor Research Unit, Department of Chemistry, Faculty of Science, Chulalongkorn University, Bangkok 10330, Thailand

## **Content:**

Description of the model for approximating the electric field strength at the protein/SAM interface.

## Electric field strength at the protein/SAM interface

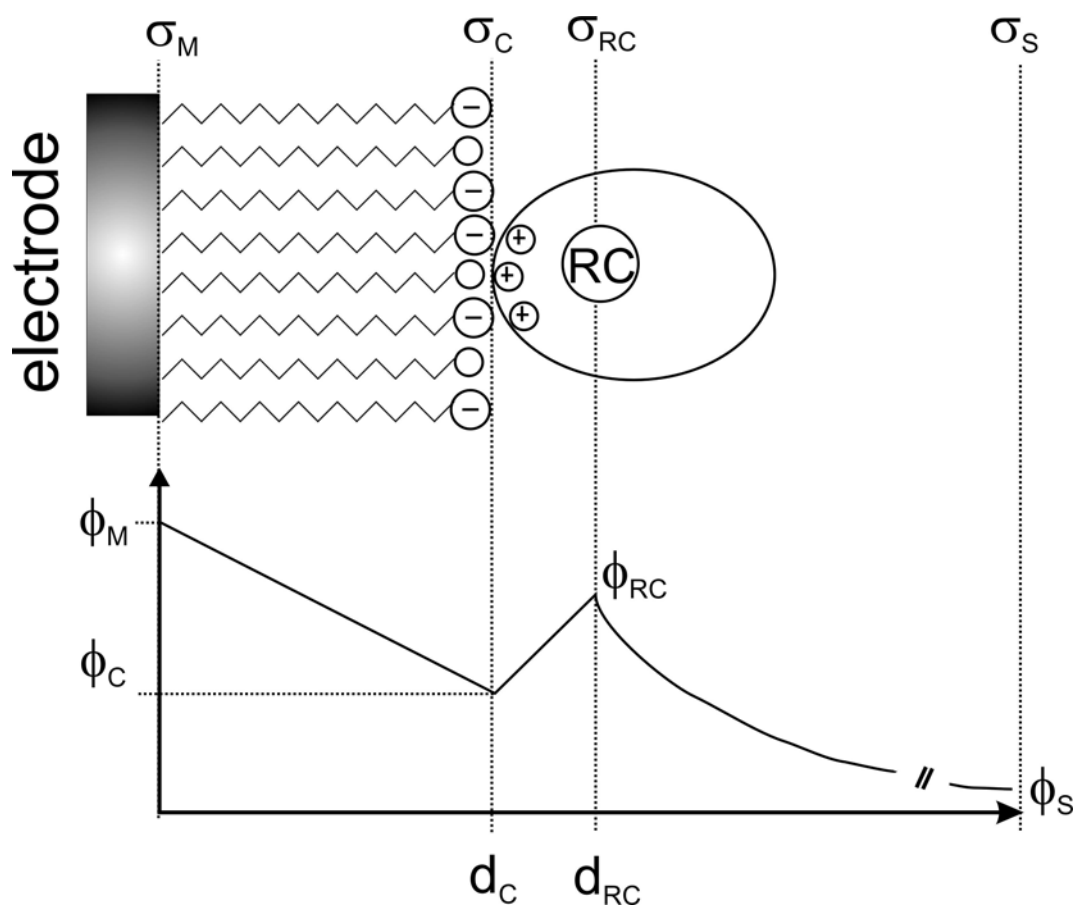


Fig. S1. Schematic representation of the potential distribution across the electrode/SAM/protein/solution interfaces.

Within the electrostatic model for the interfacial potential drops (Fig. S1), the electric field strength at the protein/SAM interface  $E_{EF}$  is given by

$$(S1) \quad E_{EF} = \frac{\sigma_M}{\epsilon_0 \epsilon_c}$$

or, with

$$(S2) \quad \sigma_M + \sigma_C + \sigma_{RC} + \sigma_S = 0$$

$$(S3) \quad E_{EF} = -\frac{\sigma_C + \sigma_{RC} + \sigma_S}{\epsilon_0 \epsilon_c}$$

where  $\sigma_M$ ,  $\sigma_C$ ,  $\sigma_{RC}$ , and  $\sigma_S$  are the charge densities on the metal, on the SAM, at the reaction site (heme), and in the solution, respectively (§ref). The quantities  $\epsilon_0$  and  $\epsilon_S$  refer to the permittivity and the dielectric constant in the SAM, respectively. Linearisation of the Gouy-Chapman expression for  $\sigma_S$  allows rewriting Eq. S3 to

$$(S4) \quad E_{EF} = -\frac{\sigma_C + \sigma_{RC} - \epsilon_0 \epsilon_S \kappa E_{RC}}{\epsilon_0 \epsilon_c}$$

where  $\epsilon_S$  and  $\kappa$  are the dielectric constant and the Debye length in solution.  $E_{RC}$  denote the potential drop at the redox site which can be expressed by

$$(S5) \quad E_{RC} = \frac{\sigma_C \epsilon_P d_C + \epsilon_o \epsilon_P \epsilon_C (E - E_{pzc}) + (d_C \epsilon_P + d_{RC} \epsilon_C) \sigma_{RC}}{\epsilon_o [\epsilon_C \epsilon_P + (d_C \epsilon_P + d_{RC} \epsilon_C) \epsilon_S \kappa]}$$

Here  $\epsilon_P$  is the dielectric constant in the protein and  $d_C$  and  $d_{RC}$  are the thickness of the SAM and the distance between the SAM/protein interface and the reaction site, as defined in Fig. S1. The quantities  $E$  and  $E_{pzc}$  refer to the electrode potential and the potential of zero charge, respectively. Inserting Eq. S5 into Eq. S4 one obtains

$$(S6) \quad E_{EF} = \frac{-\sigma_C [\epsilon_C \epsilon_P + d_{RC} \epsilon_C \epsilon_S \kappa] - \sigma_{RC} \epsilon_C \epsilon_P + \epsilon_0 \epsilon_S \epsilon_C \epsilon_P \kappa (E - E_{pzc})}{\epsilon_o \epsilon_C [\epsilon_C \epsilon_P + (d_C \epsilon_P + d_{RC} \epsilon_C) \epsilon_S \kappa]}$$

Since at pH values around 7,  $|\sigma_C| \gg |\sigma_{RC}|$  and furthermore  $\epsilon_C \epsilon_P \ll d_{RC} \epsilon_C \epsilon_S \kappa$ , Eq. S6 simplifies to

$$(S7) \quad E_{EF} = \frac{-\sigma_C d_{RC} + \epsilon_0 \epsilon_P (E - E_{pzc})}{\epsilon_o (d_C \epsilon_P + d_{RC} \epsilon_C)}$$