

# *In vitro* Observation of Direct Electron Transfer of Human Haemoglobin Molecules on glass/tin-doped Indium Oxide Electrodes

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**Abstract.** Electron transfer properties of the *haem* ( $\text{Fe}^{\text{III}}/\text{Fe}^{\text{II}}$ ) redox center, in human haemoglobin molecules, were investigated, *in vitro*, on glass/tin-doped indium oxide electrodes. The total surface concentration of electrochemically active haemoglobin molecules corresponded to a single protein monolayer. In addition, anodic and cathodic peak currents changed in direct proportion with the scan rate, suggesting a surface-controlled electrode process.

**Key words:** Cyclic voltammetry; direct-electron-transfer; human haemoglobin; tin-doped indium oxide electrode; surface electron transfer rate constant.

**Resumen.** Propiedades de transferencia electrónica, del centro redox ( $\text{Fe}^{\text{III}}/\text{Fe}^{\text{II}}$ ) del grupo *hemo*, en moléculas de hemoglobina humana, fueron investigadas, *in vitro*, sobre electrodos de óxido de indio dopado con estaño. La concentración superficial total de moléculas de hemoglobina electroquímicamente activa correspondió a una monocapa de proteína sencilla. Además, las corrientes de pico anódica y catódica cambiaron en proporción directa con la velocidad de barrido, sugiriendo un proceso de electrodo controlado superficialmente.

**Palabras clave:** Voltametría cíclica; transferencia electrónica directa; hemoglobina humana; electrodo de óxido de indio dopado con estaño; constante de velocidad de transferencia electrónica superficial.

## Introduction

The haemoglobin (Hb) molecule is an assembly of four globular protein subunits. Each subunit is composed of a protein chain tightly associated with a non-protein *haem* group. Each individual protein chain arranges in a set of  $\alpha$ -helix structural segments connected together in a “myoglobin fold” arrangement, so called because this arrangement is the same folding motif used in myoglobin proteins. This folding pattern contains a pocket that is suitable to strongly bind the *haem* group. A *haem* group consists of an iron atom held in a heterocyclic ring, known as a porphyrin. This iron atom is the site of oxygen binding. The iron atom is bonded equally to all four nitrogen atoms in the center of the ring, which lie in one plane. Two additional bonds perpendicular to the plane on each side can be formed with the iron to create the fifth and sixth positions, one of them being connected strongly to the protein through a histidine (Hys) residue, while the other one is available for the binding of oxygen. The iron atom can either be in the  $\text{Fe}^{\text{II}}$  or  $\text{Fe}^{\text{III}}$  state; however, ferrihemoglobin (methaemoglobin, metHb) ( $\text{Fe}^{\text{III}}$ ) cannot bind oxygen [1].

In adult humans, the most common Hb type is a tetramer (which contains 4 subunit proteins) called Hb A, consisting of two  $\alpha$  and two  $\beta$  subunits non-covalently bound. This is denoted as  $\alpha_2\beta_2$ . These subunits are structurally similar and about the same molecular size. Each subunit has a molecular weight of about 16,000 Da, for a total molecular weight of the tetramer of about 64,000 Da. The four polypeptide chains are bound to each other by salt bridges, hydrogen bonds, and hydrophobic

interaction. While Hb does not function physiologically as an electron transfer carrier, it does undergo oxidation and reduction at the *haem* center in certain cases *in vivo* [2]. Therefore the research on its electron transfer process might lead to a profound understanding of electron flow in biological systems.

To our knowledge, little work has been reported in the literature on electron transfer properties of human Hb in solution. For example, Topoglidis et al. [3] reported that titanium oxide and tin oxide allow the reduction of bovine metHb without the addition of any promoters and mediators. Recently, Ayato et al. [4, 5] reported that tin-doped indium oxide can induce the electron transfer of the *haem* ( $\text{Fe}^{\text{III}}/\text{Fe}^{\text{II}}$ ) redox center in bovine Hb molecules and that the protein directly adsorbed on the electrode surface of that material was not significantly denatured. Thus, semiconducting metal mixed oxides, e.g., tin-doped indium oxides,  $\text{In}_{2-x}\text{Sn}_x\text{O}_3$ , can act as both electron acceptors and electron donors, and can be considered a simple model system for mimicking a charge interface of the physiological binding domain. In order to begin a series of studies on this issue, the electron transfer properties of the *haem* ( $\text{Fe}^{\text{III}}/\text{Fe}^{\text{II}}$ ) redox center, in human Hb molecules, were investigated, *in vitro*, on commercial glass/tin-doped indium oxide (ITO) electrodes. Special emphasis is put in the Butler-Volmer model developed by Laviron for electron transfer between electrode and protein film, the morphological and structural properties of the electrode as well as the physiological milieu that was conditioned into the three-electrode cell system by means of a phosphate buffered saline (PBS) solution (0.01 mol L<sup>-1</sup> Na<sub>3</sub>PO<sub>4</sub>, 0.015 mol L<sup>-1</sup> NaCl, pH 7.2) and T = 25 °C.

## Results and Discussion

### Morphological, structural and electrochemical characterization of glass/ITO electrodes

#### Scanning electron microscopy

The surface morphology of a pretreated glass/ITO electrode was investigated using SEM (cf. Figure 1). The micrograph, taken at a 500  $\mu\text{m}$  scale, shows a well defined layer of ITO whose layer thickness was calculated to be ca. 90  $\mu\text{m}$ . Additionally, it is possible to observe a regular, uniform and flat electroodic surface. The superficial characterization of the electrode's roughness was carried out by means of a surface roughness tester HANDYSURF E-35A (TSK/Carl Zeiss®). The average roughness value obtained in this case was 0.017  $\mu\text{m}$  which is comparatively lower than the average roughness value obtained for a common glass slide (0.024  $\mu\text{m}$ ).

#### X-ray diffraction

The structural characterization of a pretreated glass/ITO electrode was investigated using XRD (cf. Figure 2). Figure 2 shows the XRD pattern of a glass/ITO substrate used like a working electrode. All of the distinct diffraction peaks corresponded to the (211), (222), (400), (440) and (622) reflections of the BCC structure of ITO ( $\text{In}_{2-x}\text{Sn}_x\text{O}_3$ ) (JCPDS Card File No. 89-4596). Almost all the peaks were very prominent and referred to the cubic rock salt structure of a very crystalline material. Moreover, strong (222) and (400) diffraction peaks are indicative of preferred orientations along the  $z$  and  $x$  directions, respectively [6]. An estimate of the mean crystallite or grain size for a given orientation was determined by using Scherrer's formula [7]:

$$D_{hkl} = \frac{K\lambda}{\beta_{hkl} \cos \theta} \quad (1)$$

where  $D_{hkl}$  is the crystallite size (nm),  $K$  is a constant (shape factor, about 0.90),  $\lambda$  is the X-ray wavelength (1.54  $\text{\AA}$  as

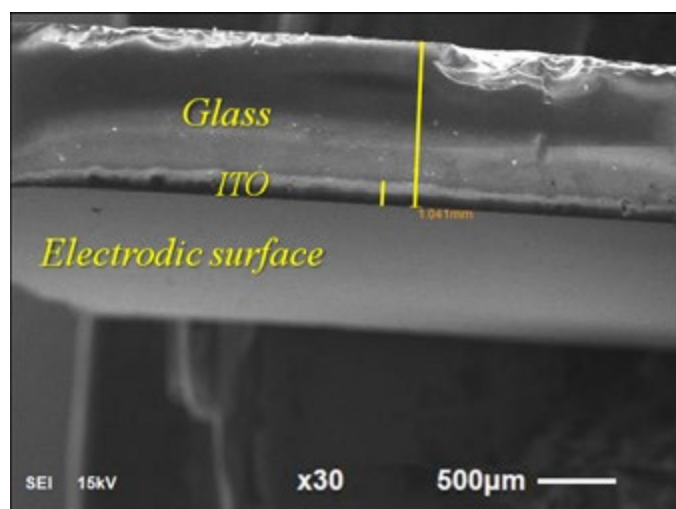


Figure 1. Cross-sectional SEM micrograph of a glass/ITO electrode.

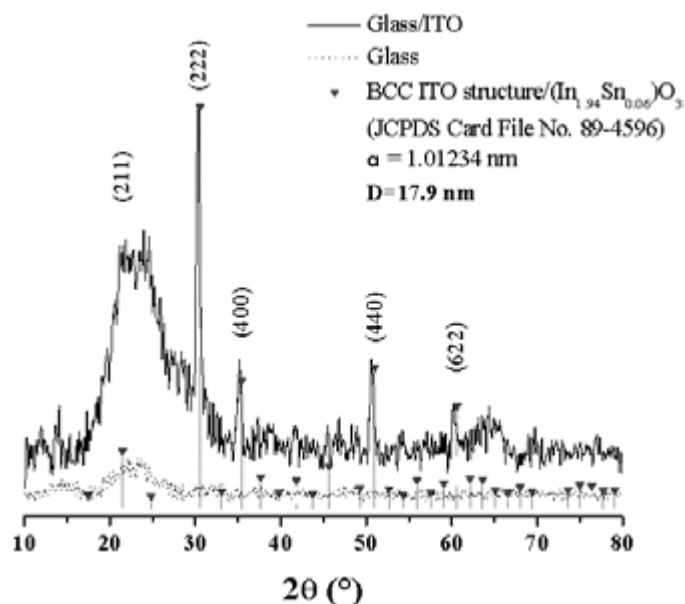


Figure 2. X-ray diffraction pattern of a glass/ITO electrode.

mentioned before),  $\beta_{hkl} = \Delta(2\theta)$  denotes the full width at half maximum (FWHM) or broadening of the diffraction peak (degree), and  $\theta$  is the diffraction angle (degree). The average  $D_{hkl}$  was estimated to be approximately  $D_{222} = 17.9 \text{ nm}$  for  $2\theta = 30.566^\circ$ . It is worth mentioning that the calculated lattice constant  $a$  for the glass/ITO substrate using Bragg's equation was  $a = 1.01234 \text{ nm}$ , which coincides with the reported value in the standard card.

#### Electroactive surface area determination

By measuring the peak current in cyclic voltammograms (CVs), the electroactive surface area of a pretreated glass/ITO electrode was determined according to the Randles-Ševčík equation for a reversible electrochemical process under diffusive control:

$$I_{pa} = 0.4463nFA_e C_{OX} \sqrt{\frac{nFvD}{RT}} \quad (2)$$

where  $I_{pa}$  is the anodic peak current (A),  $n$  is the number of electrons transferred in the redox reaction,  $F$  is Faraday's constant (96,485  $\text{C mol}^{-1}$  of electrons),  $A_e$  is the electroactive surface area of the electrode ( $\text{cm}^2$ ),  $C_{OX}$  the bulk concentration of an oxidant molecule in the solution ( $\text{mol cm}^{-3}$ ),  $v$  is the scan rate ( $\text{V s}^{-1}$ ),  $D$  the diffusion coefficient of the oxidant molecule in solution,  $(6.50 \pm 0.02) \times 10^{-6} \text{ cm}^2 \text{ s}^{-1}$ , for hexacyanoferrate (II) in 0.1 M KCl as supporting electrolyte at 25  $^\circ\text{C}$  [8],  $R$  is the gas universal constant (8.314  $\text{J K}^{-1} \text{ mol}^{-1}$ ), and  $T$  is the absolute temperature (K).

CVs for  $4.0 \times 10^{-3} \text{ M}$  hexacyanoferrate (II) in 0.1 M KCl were registered to different scan rates ( $v = 10, 20, 30, 40, 50, 60, 70, 80, 90$  and  $100 \text{ mV s}^{-1}$ ) with the glass/ITO electrode. The peak-to-peak potential separation was constant and linear relationships between the anodic and cathodic peak currents

and the square root of the scan rate:  $I_{pa} = 0.00101v^{1/2} - 5.8550 \times 10^{-7}$ ,  $R^2 = 0.9999$ ;  $-I_{pc} = 0.00101v^{1/2} + 3.2078 \times 10^{-7}$ ,  $R^2 = 0.9999$ , were achieved. From the slope of these equations,  $A_e$  was calculated to be  $1.36 \text{ cm}^2$ . The roughness factor ( $\rho$ ) of the GME, which is defined as the ratio ( $A_e/A_g$ ) [9], was estimated to be 1.18.

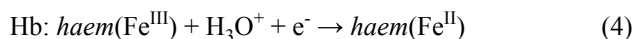
### Electrochemical behavior of human haemoglobin molecules

Figure 3 shows the CVs of glass/ITO electrodes in absence and presence of Hb molecules. Figure 3a, a CV recorded in PBS solution alone shows a non-Faradaic current behavior. The electrode had the largest background current in the non-electrolyte solution which reflected the properties of the electric double layer. The double layer capacitance ( $C_{dl}$ ) can be estimated by dividing the sum of the anodic and cathodic current with twice the scan rate [9]. So, the capacitances of the glass/ITO and glass/ITO/Hb electrodes were calculated from Figs. 3a and 3b as  $4.8$  and  $0.4 \mu\text{F cm}^{-2}$ , respectively. In Fig. 3b, a pair of redox peaks, at around  $-0.117 \text{ V}$  in the cathodic scan and at around  $-0.097 \text{ V}$  in the anodic scan, were found in that Hb-containing solution. This fact indicated that a Faradic current was generated over the glass/ITO electrode, which can be ascribed to the *haem* ( $\text{Fe}^{\text{III}}/\text{Fe}^{\text{II}}$ ) redox center in Hb molecules.

Once capacitive effects are counted out, the amount of electrochemically active Hb molecules could be estimated from integration of the charge  $Q$  (in C) under each peak in those CVs acquired at slow scan rates (i.e.,  $0.1$ ,  $0.3$  or  $0.5 \text{ V s}^{-1}$ ), given by Faraday's law:

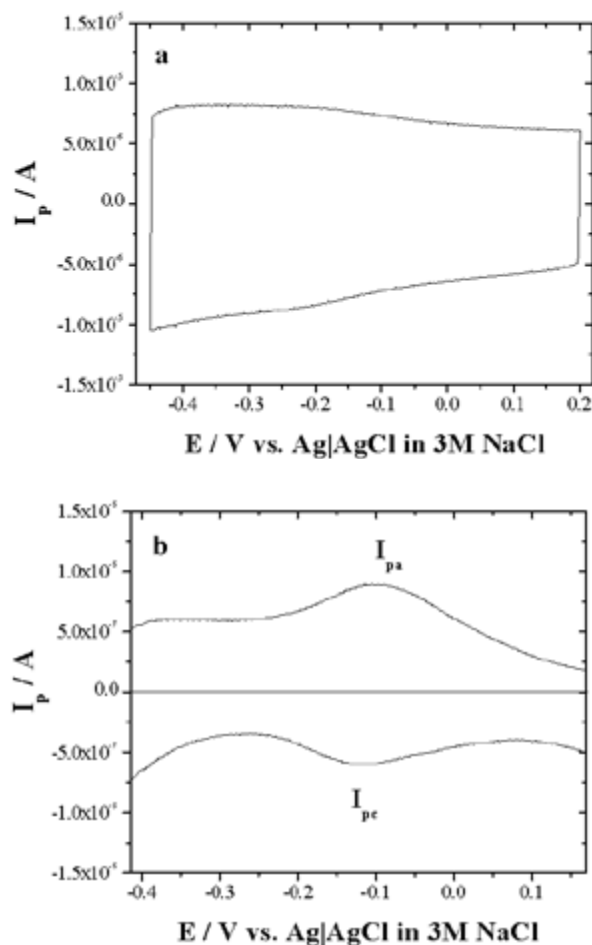
$$Q = nFA_e\Gamma_T \quad (3)$$

where  $\Gamma_T$  is total surface concentration of the protein molecule ( $\text{mol cm}^{-2}$ ),  $A_e$  is the electroactive surface area of electrode ( $\text{cm}^2$ ),  $F$  is Faraday's constant ( $96,485 \text{ C mol}^{-1}$  of electrons) and  $n$  is the number of electrons transferred in the redox reaction:



The total surface concentration of electroactive Hb molecules was estimated to be  $\Gamma_T = (4.69 \pm 0.52) \times 10^{-12} \text{ mol cm}^{-2}$ . On the other hand, the theoretical maximum coverage of a protein monolayer on the electrode surface was estimated as  $4.89 \times 10^{-12} \text{ mol cm}^{-2}$ , considering that one human Hb molecule in PBS solution has a Stokes radius of  $31.3 \text{ \AA}$  [10]. These data indicate that  $\theta = \Gamma_T/\Gamma_T^{\text{theo}} = 0.96$  of a protein monolayer was achieved.

Figure 4a shows CVs recorded with different scan rates, from  $0.1$  to  $3.5 \text{ V s}^{-1}$ . Nearly symmetric anodic and cathodic peaks were observed; in addition, they have roughly equal heights. The anodic to the cathodic peak potential difference ( $\Delta E_p$ ) was much greater than the ideal value of zero. At the lower scan rates, i.e.,  $0.1$ ,  $0.3$  or  $0.5 \text{ V s}^{-1}$ , the smaller redox peak currents were observed, while CVs with the largest redox peak currents corresponded to those acquired at the fastest scan rate, i.e.,  $3.5 \text{ V s}^{-1}$ . In such cases, the formal potential was taken

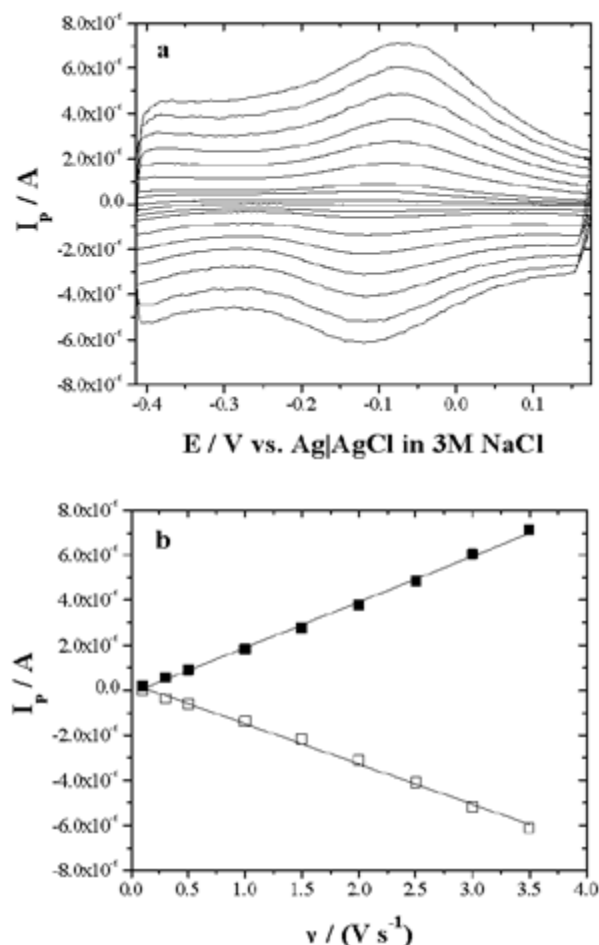


**Figure 3.** a) Cyclic voltammogram of a glass/ITO electrode in: a) PBS solution ( $0.01 \text{ mol L}^{-1} \text{ Na}_3\text{PO}_4$ ,  $0.015 \text{ mol L}^{-1} \text{ NaCl}$ , pH 7.2) and b) PBS solution ( $0.01 \text{ mol L}^{-1} \text{ Na}_3\text{PO}_4$ ,  $0.015 \text{ mol L}^{-1} \text{ NaCl}$ , pH 7.2) containing  $1.0 \times 10^{-4} \text{ mol human Hb L}^{-1}$  after subtraction of a), as a blank. The experiment was carried out at  $T = 25 \text{ }^\circ\text{C}$ . Scan rate:  $0.5 \text{ V s}^{-1}$ .

as the midpoint potential between the oxidation and reduction peaks if there is a small separation between them. Considering this criterium, an  $E^0 = -0.107 \text{ V}$  (vs.  $\text{Ag}|\text{AgCl}$  in  $3 \text{ M NaCl}$  solution) was determined at  $0.5 \text{ V s}^{-1}$ .

On the other hand, the anodic and cathodic peak currents,  $I_{pa}$  and  $I_{pc}$ , increased with increasing scan rates as observed in Figure 4b. These results are characteristic of quasi-reversible, surface confined electrochemical behavior, in which all electroactive proteins in their *haem* ( $\text{Fe}^{\text{III}}$ ) forms are reduced on the forward cathodic scan, and the reduced proteins in their *haem* ( $\text{Fe}^{\text{II}}$ ) forms are then fully oxidized to the *haem* ( $\text{Fe}^{\text{III}}$ ) forms on the reversed anodic scan. When the peak currents were plotted against the scan rate, direct linear relationships were obtained, indicating a surface-controlled electrode process. The origin of this process is indicative that the diffusion of  $\text{H}_3\text{O}^+$  ions towards the electrode surface was very fast. Therefore, the electron process can be expressed by Equation 4 as proposed above [11].

The linear regression equations for anodic and cathodic peak currents are as follows:  $I_{pa} = 2.0305 \times 10^{-6}v$ ,  $R^2 = 0.9987$ ;  $-I_{pc} = 1.7896v$ ,  $R^2 = 0.9985$ .



**Figure 4.** a) Corrected cyclic voltammograms of a glass/ITO electrode in PBS solution ( $0.01 \text{ mol L}^{-1} \text{ Na}_3\text{PO}_4$ ,  $0.015 \text{ mol L}^{-1} \text{ NaCl}$ , pH 7.2) containing  $1.0 \times 10^{-4} \text{ mol human Hb L}^{-1}$  ( $T = 25 \text{ }^\circ\text{C}$ ) as function of scan rate. Scan rates: 0.1, 0.3, 0.5, 1.0, 1.5, 2.0, 2.5, 3.0 and  $3.5 \text{ V s}^{-1}$ . b) Dependence of anodic (■) and cathodic (□) peak currents with scan rate:  $I_{pa} = 2.0305 \times 10^{-6}v$ ,  $R^2 = 0.9987$ ;  $-I_{pc} = 1.7896v$ ,  $R^2 = 0.9985$ .

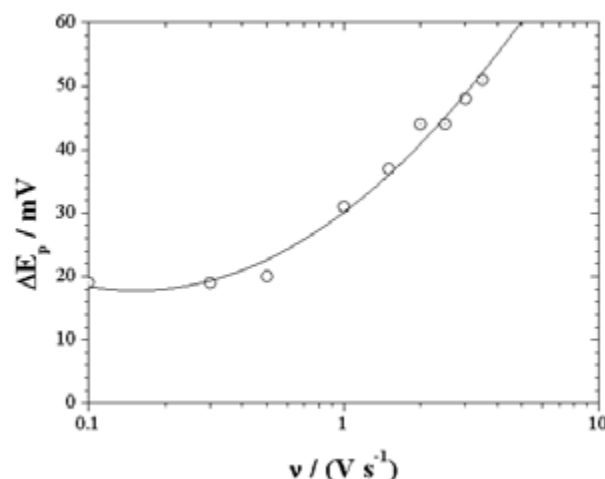
Linear plots of  $I_p$  vs.  $v$  were in good agreement with the following equation [12]:

$$I_p = \frac{n^2 F^2 A_c \Gamma_T v}{4RT} \quad (5)$$

However, their width at half height is nearly 200 mV, much larger than the ideal  $90.6/n \text{ mV}$  at  $25 \text{ }^\circ\text{C}$ .

Broadening or narrowing of CV peaks compared to the ideal  $90.6/n \text{ mV}$  at  $25 \text{ }^\circ\text{C}$  suggests a breakdown of the ideal model assumptions of no interactions between redox sites that all have the same  $E^{0'}$ . Some authors have modeled voltammograms of protein films utilizing the concept of distributions in  $E^{0'}$  and electron transfer rate constants to account for the peak broadening [13-15]. Other factors, including counterion transport efficiency, could also influence peak widths, but have not been investigated in detail for protein films.

At scan rates  $< 0.5 \text{ V s}^{-1}$ ,  $\Delta E_p$  was nearly constant in the films. As the scan rate increased, the peak potentials shifted



**Figure 5.** Influence of scan rate on the anodic to the cathodic peak potential difference ( $\Delta E_p$ ) for a glass/ITO electrode in PBS solution ( $0.01 \text{ mol L}^{-1} \text{ Na}_3\text{PO}_4$ ,  $0.015 \text{ mol L}^{-1} \text{ NaCl}$ , pH 7.2) containing  $1.0 \times 10^{-4} \text{ mol human Hb L}^{-1}$ . The experiment was carried out at  $T = 25 \text{ }^\circ\text{C}$ .

negatively (Figure 5). This is consistent with the onset of limiting kinetic effects as scan rates increase.

An increasing  $\Delta E_p$  as the scan rate is increased for an electroactive thin film suggests kinetic limitations of the electrochemistry [16] and is consistent with predictions of the Butler-Volmer model for electron transfer between an electrode and redox sites in a thin film on an electrode. Possible causes could be attributed to: (a) slow electron transfer between electrode and redox centers, (b) slow transport of charge within the film limited by electron or counterion transport, (c) uncompensated voltage drop within the film, and (d) structural reorganization of the protein accompanying the redox reactions.

When  $n\Delta E_p < 200 \text{ mV}$ , the surface electron transfer rate constant ( $k_s$ ) of the adsorbed Hb on the glass/ITO electrode can be estimated according to Laviron's equation for quasi-reversible thin-layer electrochemistry [17]:

$$\text{Log } k_s = \alpha \text{Log}(1-\alpha) + (1-\alpha)$$

$$\text{Log } \alpha - \text{Log } \frac{RT}{nFv} - \alpha(1-\alpha) \frac{nF\Delta E_p}{2.3RT} \quad (6)$$

Our experimental results showed that the scan rate in the range  $0.1$  to  $3.5 \text{ V s}^{-1}$  did not affect the  $k_s$  value, because  $n\Delta E_p < 200 \text{ mV}$ . Assuming a charge-transfer coefficient  $\alpha$  of 0.5, the  $k_s$  of the adsorbed Hb on the glass/ITO electrode was  $8.01 \text{ s}^{-1}$  at the onset of limiting kinetic effects ( $500 \text{ mV s}^{-1}$ ). This value is significantly higher than other previously reported values in the literature for different haemoglobin species and sort of electrodes. For comparison with data on bare or mediator-coated electrodes,  $k_s$  was converted to the standard heterogeneous rate constant ( $k^{0'}$ ) by using  $k^{0'} = k_s \cdot d$ , where  $d$  is the film thickness [17]. See for example [3-5, 18, 19] and references cited in [19].

This fact could be attributed to the morphological and structural properties shown by the electrode, i.e., surface

**Table 1.** Electrochemical parameters for different (superscripts: b-bovine, h-human) haemoglobin species at 25 °C.

pH	Sample   electrode	$E^0/\text{mV}$ (SHE)	$k^0/\text{cm s}^{-1}$	Reference
7.0	<sup>b</sup> Hb in solution   SnO <sub>2</sub>	-215	0.53 *	3
7.4	<sup>b</sup> Hb in solution   In <sub>2-x</sub> Sn <sub>x</sub> O <sub>3</sub>	-112	not determined	4, 5
7.2	<sup>h</sup> Hb in solution   In <sub>1.94</sub> Sn <sub>0.06</sub> O <sub>3</sub>	-62	8.01 *	this work
5.5	<sup>b</sup> Hb-DDAB-Nafion   edge-plane PG	80	5.7 *	18
5.5	<sup>h</sup> Hb-DDAB   edge-plane PG	84	2.7 *	19
5.5	<sup>h</sup> Hb in solution   edge-plane PG	not detected	not detected	op. cit. 19
7.0	<sup>b</sup> Hb in solution   Pt + MB	145	$2.0 \times 10^{-4}$	op. cit. 19
7.0	<sup>b</sup> Hb in solution   Pt + Azure A	180	$3.5 \times 10^{-6}$	op. cit. 19
7.0	<sup>b</sup> Hb in solution   Pt + BCG	184	$2.0 \times 10^{-7}$	op. cit. 19

DDAB = Didodecyldimethylammonium bromide, PG = pyrolytic graphite, MB = methylene blue, BCG = brilliant cresyl green

\* For comparison with CV data where diffusion control pertains,  $k_s$  was estimated from the standard heterogeneous rate constant ( $k^0$ ) by using  $k^0 = k_s \cdot d$ , where  $d$  is the film thickness [17].

roughness, crystallinity and hydrophilicity [20], and the physiological milieu of the three-electrode cell system that charged, negatively, to the working electrode [21] and protein [22]. All these factors played an important role in providing a more favorable microenvironment for the protein.

## Conclusions

In this study, we clearly demonstrated that human Hb molecules directly physisorbed on glass/tin-doped indium oxide substrates exhibited direct electron transfer (DET) in PBS (0.01 M Na<sub>3</sub>PO<sub>4</sub>, 0.015 M NaCl, pH = 7.2) solution and  $T = 25$  °C.

The experimental results suggest that acid-base equilibria and the water molecule coordinated to the *haem* group as the sixth ligand might play an important role in the electron transfer process between human haemoglobin and very crystalline and hydrophilic tin-doped indium oxide electrodes.

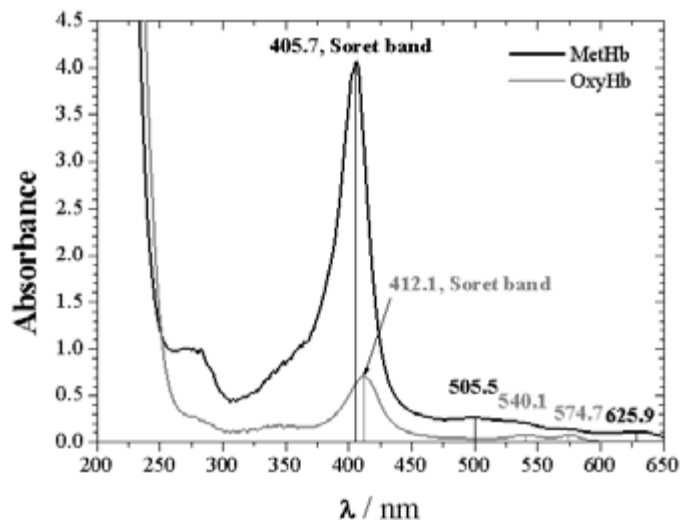
## Experimental

### Chemicals

Human Hb (Product No. H7379, MW = 64 500) and phosphate buffered saline (PBS) Packs (0.01 mol L<sup>-1</sup> Na<sub>3</sub>PO<sub>4</sub>, 0.015 mol L<sup>-1</sup> NaCl, pH 7.2), BupH™ were purchased from *Sigma-Aldrich*® and *Thermo Scientific*®, respectively, and used without further purification. The concentration of Hb was adjusted to  $1 \times 10^{-4}$  mol Hb L<sup>-1</sup> using the PBS solution.

### UV-Visible spectroscopy analysis

Absorption spectra of human Hb were measured at  $\lambda = 200 - 1000$  nm with an UV-Visible spectrophotometer GBC (Cintra 101), using the following parameters: step size = 0.16 nm, scan speed = 400 nm/min, slit width (SW) = 2 nm. The concentration



**Figure 6.** The UV-visible absorption spectra for metHb and oxyHb exhibiting the highly conjugated porphyrin macrocycle with intense features at 405.7 nm and 412.1 nm (the “Soret” bands), respectively, followed by several weaker absorptions (Q bands) at higher wavelengths (from 450 to 650 nm) [24-26].

of Hb referred above was estimated by this technique using the following absorptivity value:  $A_{540}^{1\%} = 5.97 \text{ cm}^{-1}$  [23].

### Scanning Electron Microscopy analysis

Scanning electron microscopy (SEM) micrographs were taken with a scanning electron microscope JSM-6510LV (*JEOL*) operated at an accelerating voltage of 15 KV.

### X-Ray Diffraction Analysis

The structural characterization was determined by X-ray powder diffraction (XRD) using a diffractometer D8 Advanced (*Bruker AXS*), using the following parameters: U = 40 kV, I =

35 mA, Ni-filter, and Cu-K $\alpha$  radiation,  $\lambda = 1.54 \text{ \AA}$ . A background diffractogram was subtracted using a glass/tin-doped indium oxide slide as blank. For qualitative analysis, XRD diffractograms were recorded in the interval  $10^\circ \leq 2\theta \leq 80^\circ$  at a scan speed of  $2^\circ/\text{min}$ .

### Electrochemical systems

Glass/tin-doped indium oxide (ITO) substrates were purchased from *TIRF Technologies, Inc.* The ITO film surface was cleaned according to the following standard procedure [27]: immersion for 15 min each in a series of ultrasonically agitated solvents (acetone, ethanol, water) then for 15 min each in ultrasonically agitated a) 2.0% (v/v) phosphate-free detergent solution Hellmanex (*Hellma*<sup>TM</sup>; sonification apparatus Super RK510, *Sonorex*), b) deionized water type I and c) ethanol at room temperature. In between the sonification steps the samples were rinsed in deionized water type I. Finally, the substrates were dried in a stream of nitrogen gas (*Praxair*, 99.999%) until further processing. The electrode potential was controlled with a potentiostat-galvanostat EW-4960 (*Epsilon*<sup>TM</sup> *BASi*) coupled to a PC/Processor Intel® Celeron, 3.06 GHz, using a three-electrode cell system. A glass/ITO substrate with a geometric area of  $1.15 \text{ cm}^2$  was used as working electrode. A platinum wire and a Ag|AgCl in 3 M NaCl solution electrode were used as counter and reference electrodes, respectively. This cell system was thermostated at  $25 \pm 0.1 \text{ }^\circ\text{C}$ . Prior to voltammetry, the Hb solution was purged with nitrogen gas (*Praxair*, 99.999%) for at least 30 minutes; then, a nitrogen atmosphere was maintained over the solution during experiments.

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