INTRODUCTION

Several epidemiological studies have reported possible effects on human health from exposure to extremely low frequency electromagnetic field (ELF-EMF).

In response to public concern over health effects of EMF exposure, in 1996, the International EMF Project was established by WHO and the Radiation and Environmental Health Unit, which coordinated studies on EMF relative to the Environmental Health Criteria (EHC).

In particular, the International Agency for Research on Cancer (IARC) formally evaluated the evidence for carcinogenesis from exposure to static and ELF-EMFs, concluding that ELF-EMFs are possibly carcinogenic to humans.\(^1\)

Milham and Ossiander have suggested that the appearance of the peak incidence at around age 3 in childhood acute lymphocytic leukemia is linked to electrification.\(^2\)

Further significant concern has been raised about the capacity of EMFs to cause DNA damage and chromosomes aberrations.\(^3-6\)
National radiation advisory authorities recommends measurements to minimize exposure to their citizens, as the exposure limits to electromagnetic fields recommended by the International Commission on Non-Ionizing Radiation Protection (I.C.N.I.R.P.)

However, we focused our attention on proteins. Indeed, proteins are fundamentals in organic metabolism of livings. In the cells each protein must fold into the specific conformational state in a complex and highly crowded environment, and the folding process is aided by a range of auxiliary proteins.

Otherwise it was largely demonstrated that several type of environmental stress agents can alter the secondary structure of proteins.

Recently it was proved that also ELF-EMFs and MWs can alter the secondary structure of proteins.

Haemoglobin is a heme-protein whose physiological importance is mainly related to its ability to bind molecular oxygen. The oxygen carried by heme-proteins is bound directly to the ferrous iron atom of the heme prosthetic group.

The heme portion of hemoglobin is extremely important because it aids in oxygen binding.

Haemoglobin is a tetrameric heme-protein found in erythrocytes where it is responsible for binding oxygen transporting the bound oxygen throughout the body to be used in aerobic metabolic functions.

The aim of this work was to investigate the alteration produced by exposure to EMFs on the secondary structure of this heme-protein that perform an important role in metabolism processes of organic systems, focusing the attention on aggregation mechanisms.

Fourier Transform Infrared (FTIR) Spectroscopy was used to investigate the effects of EMFs in the secondary structure of this protein in aqueous solution, in particular Amide I and II modes in the range 1700-1500 cm⁻¹. Indeed, it may be considered the most versatile spectroscopic technique for analyzing the secondary structure of a protein in diverse physiochemical environments.

**MATERIALS AND METHODS**

**Haemoglobin samples were obtained as previously reported**

The exposure system for haemoglobin consisted of a couple of Helmholtz coils, with pole pieces of round parallel polar faces, to produce a uniform magnetic field at the center of the coils distance. A magnetic flux density of 1 mT between the polar faces of the coils was generated by means of an AC voltage regulating up to 230 V. Samples were placed at the center of a uniform field area between the coils and the magnetic field was continuously monitored by a magnetic field probe GM07 of HIRST Magnetic Instruments Ltd, UK.

a. Analogue unexposed samples at the same room temperature were used as the control.

b. FTIR spectra were recorded by a spectrometer, Vertex 80v, from Bruker Optics.

c. The attenuated total reflection (ATR) method was chosen for spectrum collection.

d. For each spectrum, 64 interferograms were collected with a spectral resolution of 4 cm⁻¹ in the range from 4000 to 1200 cm⁻¹, using the techniques accurately described in 11.

e. Either exposed or control samples were located in the same room at a temperature of 20°C.

**RESULTS AND DISCUSSION**

**Exposure of Haemoglobin to 50 Hz EMF**

Samples of 250 µL of hemoglobin in bidistilled water aqueous solutions were exposed for 4 h to a uniform electromagnetic field of 1 mT at the frequency of 50 Hz at a room temperature of 20°C. Analogue unexposed samples at the same room temperature were used as the control.

Typical spectra from 1800 to 1400 cm⁻¹, obtained after exposure, are showed in Figure 1.

The spectra exhibited an intense amide I band centered at about 1654 cm⁻¹, corresponding mainly to an α-helix structure content due to C=O
stretching vibration and a N–H bending mode, a low intensity amide II, coupling of the N–H bending and C–N stretching modes. 

A significant decrease in intensity of the amide I and II modes was evidenced for exposed sample spectra, that can be due to a decrease of the α-helix component, a loss of α-helical and short-segment connecting α-helix segments in the amide I region. 

In addition, a perceptible increase of the β-sheet content in the region 1635-1610 cm$^{-1}$ can be observed in Figure 1, as well. 

Hence, Fourier self deconvolution (FSD) analysis was used to highlight the alterations in the amide I region. The concept of FSD is based on the assumption which a spectrum of single narrow bands is broadened in the liquid or solid state and cannot be

Fig. 1: Representative infrared spectra from 1800 to 1400 cm$^{-1}$ of hemoglobin in bidistilled water solution after 4 h of exposure to 50 Hz frequency EMF at 1 mT (red lines represent exposed samples spectra). The amide I and II regions are evidenced. The decrease in intensity after exposure was significant in particular in the amide II region.

Fig. 2: Representative FSD infrared spectra in the amide I region of hemoglobin in bidistilled water solution after 4 h of exposure to 50 Hz frequency EMF at 1 mT. The relative increases in intensity of β-sheet features (indicated by arrows) after exposure were evidenced by FSD analysis. Red lines represent exposed samples spectra.
distinguished in the amide envelope. A curve fitting procedure can be applied to estimate quantitatively the area of each component representing a type of secondary structure.

After FSD analysis on exposed and not-exposed spectra, vector normalization was used.

This analysis revealed the presence of five vibration bands centered at 1654, and around 1635, 1625, 1615 and 1685 cm\(^{-1}\) in the amide I region, as can be observed in Figure 2.

The band at 1654 cm\(^{-1}\) is due to \(\alpha\)-helix structures, and the other vibrations can be associated with \(\beta\)-sheet structures\(^{17-19}\).

This analysis relative to hemoglobin in bidistilled water aqueous solution revealed a significant increase in \(\beta\)-sheet bands after 4 h of exposure, indicated with arrows in Figure 2, comparing exposed and unexposed spectra. These features can be attributed to the formation of aggregates\(^{20-22}\).

In conclusion, an unfolding process of hemoglobin due to ELF-EMF exposure was enhanced and confirmed by FSD analysis.

CONCLUSION

The effects of exposure of 4 h to 50 Hz EMF at 1 mT on hemoglobin aqueous solutions were studied by means of FTIR techniques, showing that ELF-EMF can affect infrared vibration bands of hemoglobin.

In particular, a loss of \(\alpha\)-helical and short-segment connecting \(\alpha\)-helix segments was observed in amide I and amide II regions.

In addition, the use of FSD analysis evidenced a relative increase in intensity of the \(\beta\)-sheet content with respect to \(\alpha\)-helix component in the amide I region, that can be attributed to the formation of aggregates.

Previous literature have indicated that the propensity to cause the transition from \(\alpha\)-helix to \(\beta\)-sheet structure can be responsible for aggregation leading to the neurotoxicity and neurodegenerative disorders that can be considered as the first step to some pathologies. Hence, further research is needed to highlight bioprotective mechanisms against the effects of ELF-EMF.

REFERENCES


