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Effects of Extremely Low Frequency Magnetic Field on the Secondary Structures of β -Amyloid and Human Serum Albumin

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Abstract: Human serum albumin and β -amyloid were exposed to extremely low frequency (ELF) magnetic field of 1.5 mT intensity and 50 Hz frequency. The effects of exposure were investigated in the mid-infrared region by means of Fourier self-deconvolution spectroscopic analysis. The experimental results suggest that exposure to the ELF magnetic field has reversible effects on the out of phase combination of N-H in plane bending and C-N stretching vibrations of the secondary structures of the two proteins. The exposure of β -amyloid and human serum albumin to ELF magnetic field affected the absorption spectra of the vibration bands by changes in peak positions for the amide II bands and changes of intensities in most of the bands in the amide I and amide II regions. In the fingerprint region, the most sensitive vibrations to the magnetic field are found to be in the (720-600) cm^{-1} range. After removing the magnetic field, it took the vibration bands more than 10 minutes of a gradual change toward returning to their original spectra, obtained before the exposure. It is suggested that hydrogen bonds can alter the frequency of a stretching vibration depending on the increase or decrease of strain on the vibrations.

Keywords: FTIR-Spectroscopy, ELF-Magnetic Field, β -Amyloid, HAS, Protein Dynamics

1. Introduction

Infrared spectroscopy is one of the most important analytical techniques recognized in obtaining information on structural features of proteins. The excitation of molecular vibrations occurs when the frequency of the absorbed infrared radiation is equal to the difference between the corresponding vibrational energy levels. Fourier transform infrared spectroscopy (FTIR) spectroscopy is a valuable tool in measuring wavelength and intensity of the absorption of a protein sample by IR.

A considerable amount of scientific research had shown a strong correlation between band positions of the IR spectra and

the secondary structure of the studied proteins. The frequencies of the observed absorption bands in several proteins are linked to certain molecular vibrations in the secondary structures, such as α -helical or β -sheet structures [1-3]. Normal functionality of a protein depends on the three-dimensional structures of its constituents of peptides and polypeptide.

There are nine IR bands, named amide A, amide B and amides I-VII, in the order of decreasing frequency [3, 4]. The most intense and most useful for the analysis of the secondary structure of proteins was found to be the amide I band (1600-1700 cm^{-1}), which involves C=O stretching vibration [4, 5]. The amide II band (1480-1600) cm^{-1} and amide III band (1220 -1320) cm^{-1} contribute greatly to studies despite of the complex nature of their compositions

and can be used for secondary structure prediction [6-8]. The spectra in the region from about (1300 - 500) cm^{-1} arise due to complicated combinations of (C-C, C-O, C-N) stretching and bending vibration. This range, referred to as the fingerprint region, is important because each compound produces its unique pattern of peaks.

Within the cells, proteins must fold into the specific conformational state in a complex highly crowded environment aided by a range of auxiliary proteins [9, 10]. Changes in protein properties and enzyme reactions have been studied by analyzing secondary structures of proteins by IR spectroscopy. The regions of IR spectra corresponding to the amide I band, the amide II band, and amino acids side chain vibrations are especially suitable targets of such analysis.

β -amyloid (A β) which contains rich structured β -sheet, is insoluble in the physiological solutions and forms toxic oligomers within the human body in general leading to amyloidosis in different diseases [11-13]. Human serum albumin (HSA) is highly abundant protein in the blood plasma and usually binds and transports fatty acids, hormones, bilirubin, tryptophan, steroids, metal ions, and therapeutic agents. HSA serves as the major soluble protein constituent of the circulatory system, it contributes to colloid osmotic blood pressure, and it can bind and carry drugs which are poorly soluble in water [14].

Several studies have indicated a relationship between the exposure to extremely low frequency (ELF) electromagnetic fields and an increased risk of cancer [15, 16]. The world Health organization (WHO), had reported in 2007 that external ELF magnetic fields induce currents in the body and can lead to nerve and muscle stimulation in nerve cells. In 2002 The International Agency of Research on Cancer (IARC) had classified ELF magnetic fields as "possibly carcinogenic to humans" [17-21]. Although magnetic particles are found in the human body in various forms, not enough evidence appear to confer a sensitivity to environmental ELF magnetic fields. It seems that further scientific investigations are needed to explore any possible indirect effect under certain conditions.

The interest in the effect of ELF magnetic fields on living cells is significant for two reasons, first is the potential of its harmful implications on human health and the second is due to its vital role of sustaining biological functions for animals and humans. The effects on some biological processes, such as photosynthetic reaction of cryptochrome protein have demonstrated change in the protein orientations with respect to earth magnetic field causing it to act as a magnetic compass. In general, each of cryptochrome and photolyase can respond to magnetic field and transduce a physical signal into a chemical signal [22-28]. This magnetoreception in which an organism detects a magnetic field for navigating a location, is based on Zeeman's quantum effect where unpaired electron spins may be either antiparallel (\downarrow), corresponding to a singlet state, (S) or parallel (\uparrow) which correspond to a triplet state, (T) are created as a result of light absorption in the presence of applied magnetic field [29-35].

Magnetic field may affect some proteins through the

electrostatic interactions between electrons. It is possible to account for the anomalous Zeeman effect where weak magnetic field can yield a magnetic dipolar energy of ($E = g m_j \mu_B B$), where g is the Lande factor, B is the magnetic field, m_j is the quantum number for the total angular momentum and μ_B is Bohr magneton. If ($\mu_B B$) is less than the spin-orbit energy this will split the energy for the p and s states and spectral lines due to electric dipole transition will become allowed. If the Hamiltonian for two electron system commutes with the particle exchange operator then the degeneracy will be lifted creating $S=0$ state (singlet state) and $S=1$ (triplet state).

Charge motion and therefore magnetic fields are part of our environment at every level which leads to a natural assumption that magnetic field have a vital role in all biological systems and their functions. In this work, we intend to investigate the effects of exposing A β and HSA to extremely low frequency electromagnetic field.

2. Materials and Methods

A β (1-40) human, in powder form and human serum albumin HSA (fatty acid free) were purchased from Sigma Aldrich chemical company and used without any further purification.

2.1. Preparation of Stock Solutions

An amount of 0.5 mg of A β (1-40) was dissolved in 250 μL of triple distilled water to give a concentration of 0.462 mM as a stock solution.

HSA was dissolved in phosphate buffered saline (40 mg/ml). The concentration of HSA in the buffer solution was prepared using its list molecular weight of 66.5 kDa. The solutions of A β and HSA were incubated for 1 h (at 20°C) before spectroscopic measurements were taken.

2.2. FTIR Spectroscopic Measurements

The FTIR measurements were obtained on a Bruker IFS 66/S spectrophotometer equipped with a liquid nitrogen-cooled MCT detector and a KBr beam splitter. The spectrometer was continuously purged with dry air during the measurements. FTIR Samples were prepared after one hour of incubating the A β and HSA free solutions at room temperature. Four drops of each sample were placed on a certain area on a silicon window plate and left to dry at room temperature.

A spectrum was taken as an average of 60 scans to increase the signal to noise ratio, and the spectral resolution was at 4 cm^{-1} . The aperture used in this study was 8 mm, which was found to give the best signal to noise ratio. Baseline correction, normalization and peak areas calculations were performed for all the spectra by OPUS software. The peak positions were determined using the second derivative by OPUS software in addition to the Fourier self-deconvolution (FSD) procedure which is kept at 6 steps for all deconvolution process to be consisted for data

analysis. The concept of (FSD) assumes that a spectrum is composed of several single bands (each narrow band is characteristic of a secondary structure), which are broadened in the liquid or solid state.

The FTIR spectra of A β and HSA were obtained in the featured region of (4000 - 600) cm^{-1} . The FTIR spectra were acquired by subtracting the absorption spectrum of the buffer solution from the spectrum of the protein solution. For the net interaction effect, the difference spectra were generated using the featureless region of the A β and HSA protein solutions (1800 - 2200) cm^{-1} as an internal standard [5]. The accuracy of this subtraction method was tested using several control samples with the same protein concentrations, which resulted into a flat base line formation. The spectral differences were used to investigate the net effect of ELF magnetic field on A β .

2.3. ELF Magnetic Field Measurements

A magnetic field with a sinusoidal waveform was

generated by an electromagnet coil linked to an AC voltage regulating up to 220 volts, at the frequency of 50 Hz. This device was used to generate time-varying ELF electromagnetic fields. This set up is used to expose the samples to a fixed magnetic field of 1.5mT. The magnetic field was continuously monitored by a magnetic field axial probe PHYWE Gauss meter. In the last experimental part, each of A β and HSA samples were exposed to ELF separately for 10 minutes and after removing the magnetic field, the spectra were taken every 2minutes to monitor changes as a function of time.

3. Experimental Results

The FTIR absorption spectra are recorded in the range (4000-600) cm^{-1} including all the amide regions and the fingerprint region. The obtained spectra are taken for each of A β and HSA samples before and after exposure to ELF magnetic fields of 1.5 mT.

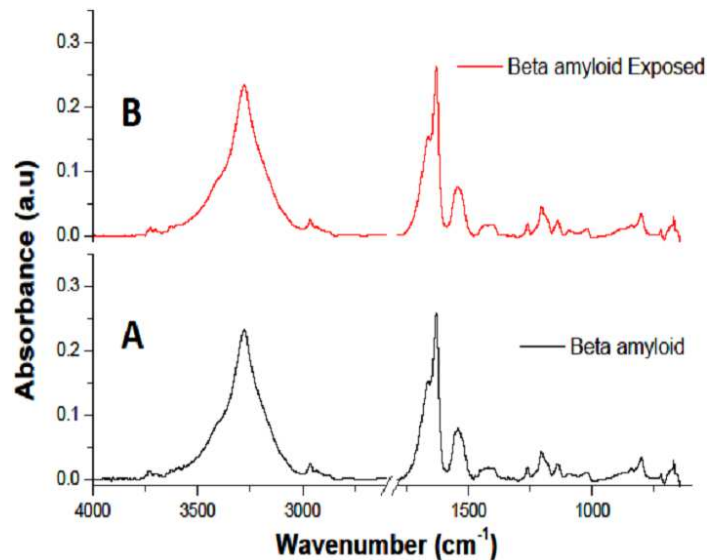


Figure 1. Absorption spectra of β -Amyloid (A) Sample is not exposed to ELF magnetic field, (B) Sample is exposed to ELF magnetic field 1.5 mT.

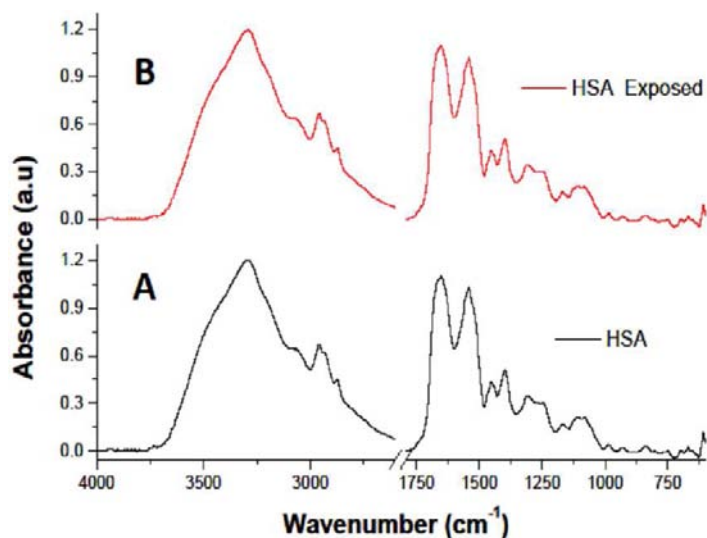


Figure 2. Absorption spectra of HSA (A) Sample is not exposed to ELF magnetic field, (B) Sample is exposed to ELF magnetic field 1.5 mT.

The absorption spectra of A β and HSA are shown in figures 1 and 2, respectively. The spectra is shown for each of A β and HSA before and after exposure to magnetic field. Each of the graphs shows the major absorption regions, including the amide bands and the fingerprint regions.

The second derivatives of the absorption spectra in the region of (1800 – 600) cm^{-1} are shown in figure 3 for both of HSA and A β . All major peaks have been identified based on the second derivative results and FSD spectra [36].

The analysis of the absorption spectra of A β in the amide I region (1600 -1700) cm^{-1} by FSD, yields several bands. The peaks of these bands correspond to the C=O stretching vibrations of the amide group, coupled to the C-N stretching and C-C-N deformation mode [37]. On the other hand, the FSD spectra shows the individual bands in the amide II (1600 -1480) cm^{-1} , which are mainly due to out of phase combination of N-H in plane bending and C-N stretching vibration modes [38]. As for amide III (1320-1220) cm^{-1} , the absorption bands are mainly due to in-phase combination of the N-H bending and the C-N stretching vibration with some contributions from the CO in plane bending and the CC stretching vibration [38, 39].

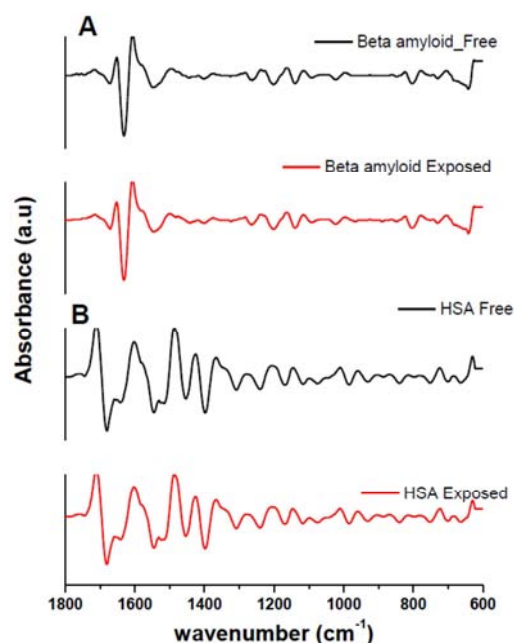


Figure 3. Second Derivative of Spectra of (A) Beta amyloid and (B) HSA.

Table 1. Bands assignment of the absorbance spectra (cm^{-1}) for β -amyloid in Amide (I, II, III) regions.

Region cm^{-1}	Band positions		Exposed to Field	Difference spectra
Amide I (1600–1700)	1609	β -sheet	1609	1612
	1629	β -sheet	1629	1628
	1648	Random	1645	1642
	1666	α -helix	1664	1660
	1682	β -turn	1680	1676
	1697	β -sheets	1695	1692
Amide II (1480–1600)	1490	β -sheets	1496	1499
	1508	Random	1520	1515
	1522	β -sheets	1532	1532
	1540	α -helix	1553	1550
	1556	α -helix	1569	1566
	1573	β -turn	1585	1584
Amide III (1220–1330)	1594	β -sheets	1596	1599
	1242	β -sheets	1242	1242
	1261	Random	1261	1263
	1279	β -turn	1280	1279
	1300	α -helix	1300	1295
	1318	α -helix	1319	1308

Table 2. Bands assignment of the absorbance spectra (cm^{-1}) for HSA in Amide (I, II, III) regions.

Region cm^{-1}	Peak positions		Exposed to Field	Difference spectra
Amide I (1600–1700)	---	β -sheet	1616	1614
	1633	β -sheet	1633	1630
	1651	α -helix	1651	1645
	1667	β -turn	1667	1660
	1682	β -turn	1682	1676
	1695	β -sheets	1695	1692
Amide II (1480–1600)	1494	β -sheets	1494	1499
	1511	Random	1511	1515
	1521	β -sheets	1521	1532
	1540	α -helix	1540	1549
	1556	β -turn	1556	1566
	1572	β -turn	1572	1584
Amide III (1220–1330)	1591	β -sheets	1590	1600
	1225	β -sheets	1225	1232
	1242	β -sheets	1242	1262

Region cm^{-1}	Peak positions		Exposed to Field	Difference spectra
	1269	Random	1267	1278
	1388	β -turn	1388	1293
	1298	β -turn	1298	1306
	1316	α -helix	1316	---
	---		---	1329

The amide I, amide II, and amide III bands are assigned in tables 1 and 2 based on several previous studies [40-44]. Furthermore, other bands in $(1300-900) \text{ cm}^{-1}$ range are assigned to C-O bending modes of saccharides (glucose,

lactose and glycerol), the peaks at $(1360-1430) \text{ cm}^{-1}$ attributed to vibrations of certain amino acids chains and $(1430-1480) \text{ cm}^{-1}$ is attributed to fatty acids, phospholipids and triglycerides [45, 46].

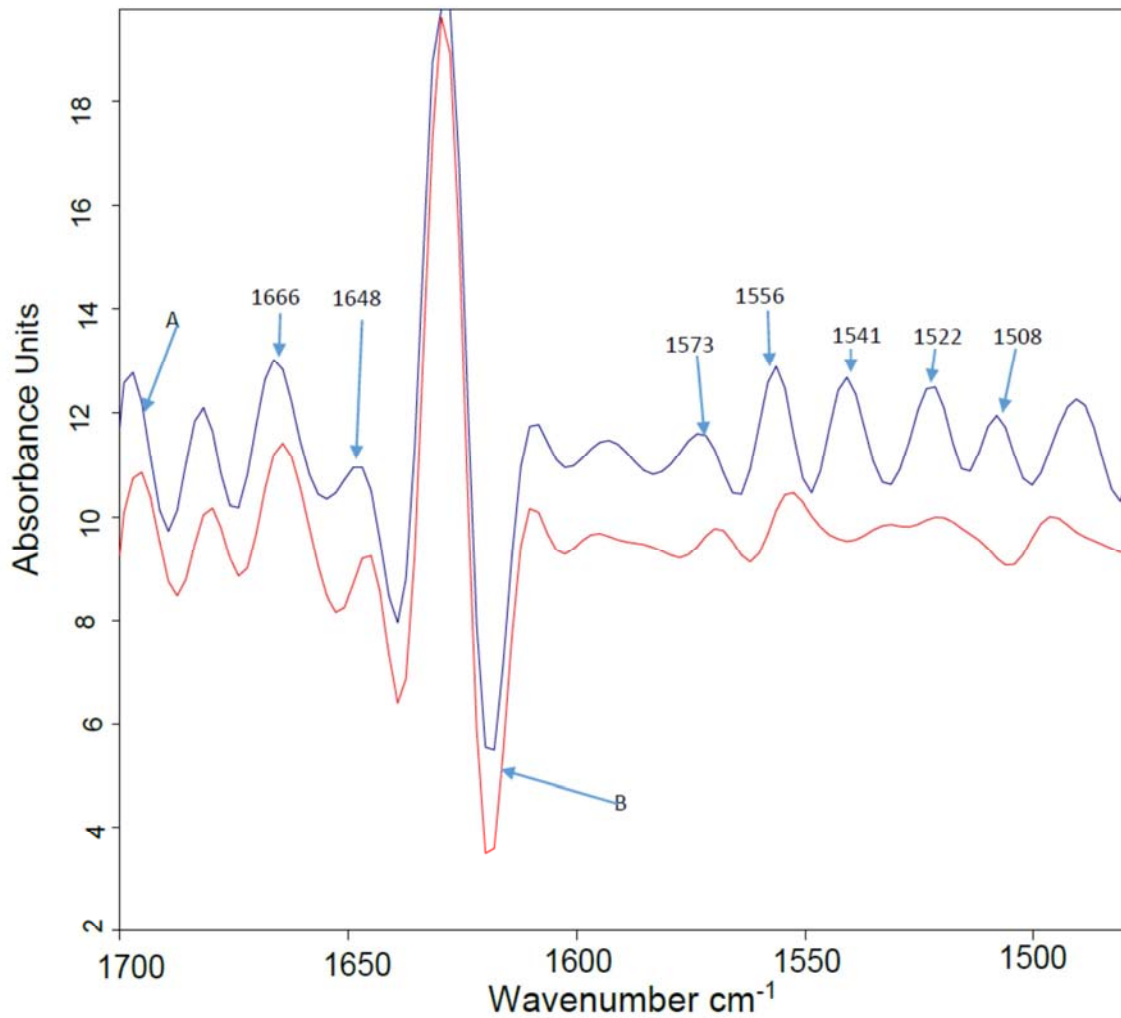


Figure 4. Deconvoluted absorption spectra for β -amyloid in the Amide I and Amide II regions.

A- β -amyloid free

B- β -amyloid exposed to ELF magnetic field 1.5 m T

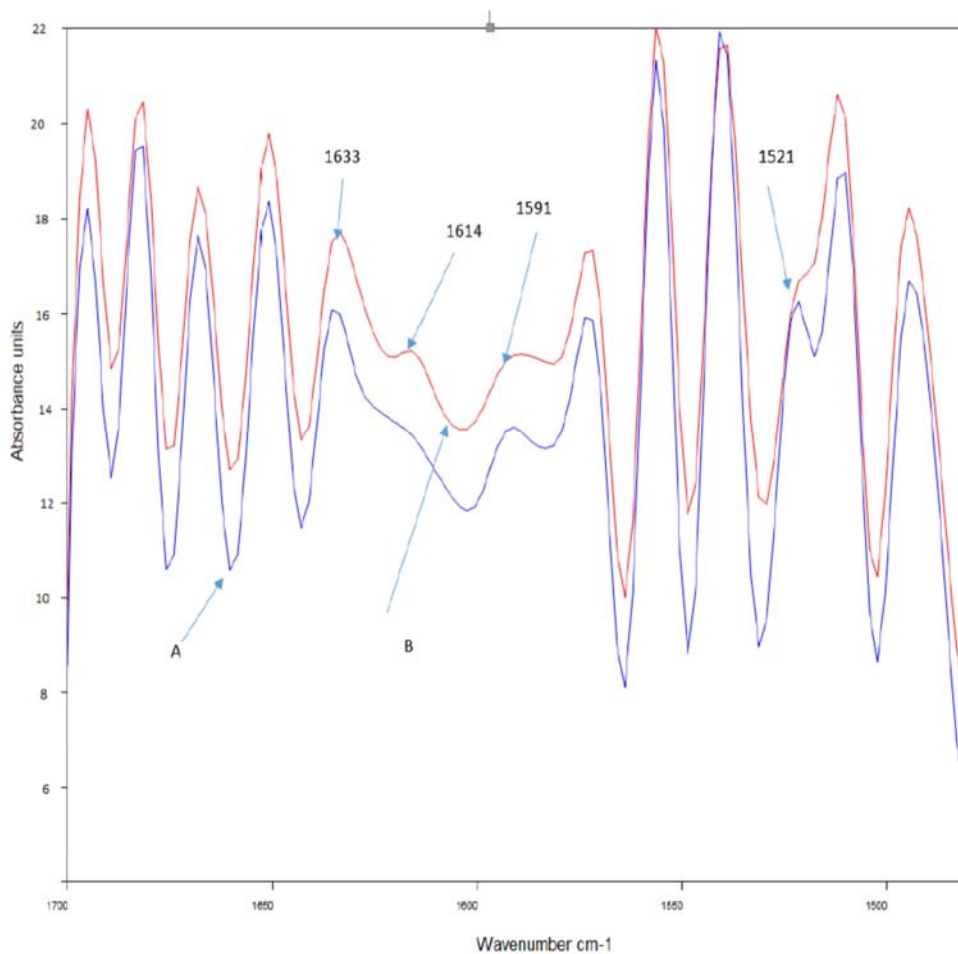
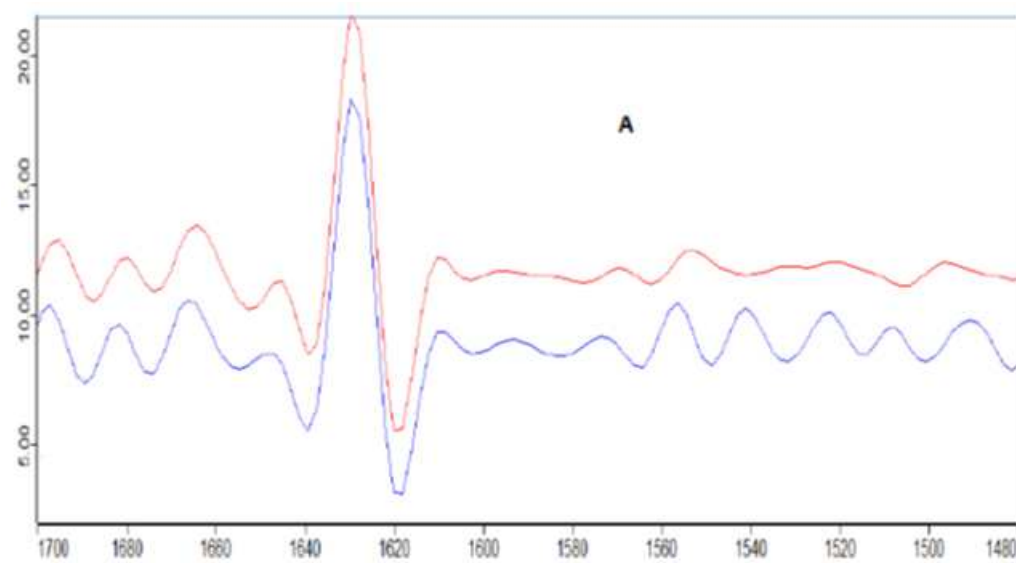


Figure 5. FSD spectra of HSA in the Amide I and Amide II regions.

A - Sample is not exposed to magnetic field
 B- Sample is exposed to 1.5 mT ELF magnetic field

The effect of exposing $A\beta$ and HSA to a 50 Hz oscillating magnetic field of 1.5 mT are shown in figures 4 and 5 for the amide I and amide II regions. The two figures show the FSD spectra for the two samples with all the main peaks that may reflect changes on the secondary structure of the two

proteins. In the case of $A\beta$, figure 4 shows the peaks' changes as listed in table 1, where peak positions and band intensities have been affected. On the other hand, figure 5 shows the changes for the HSA sample after exposure to the same magnetic field, and all peaks are listed in table 2.



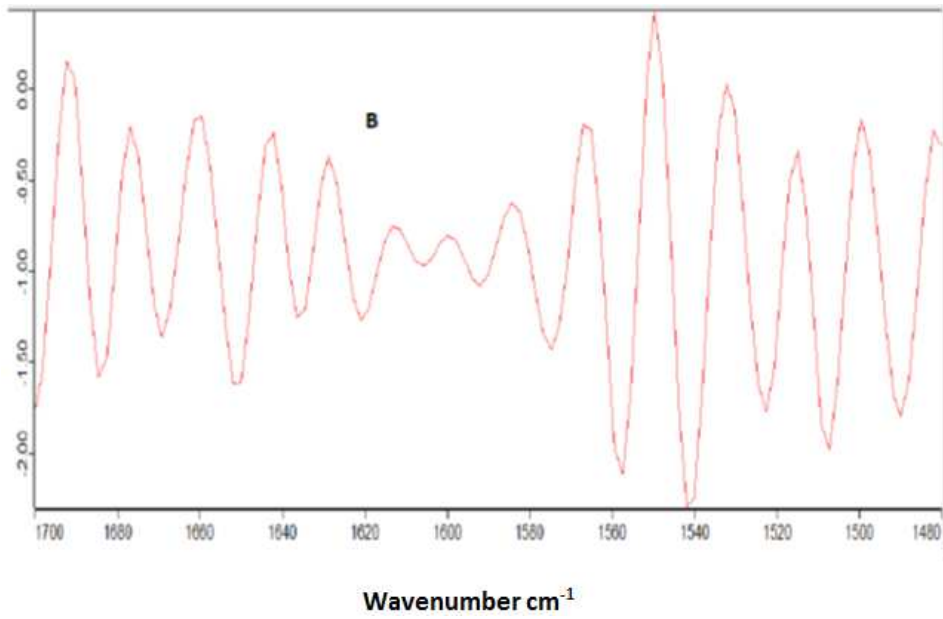


Figure 6. Difference of FSD spectra of β -Amyloid.

A- The two FSD spectra exposed to magnetic field (above) and no field (below)
 B- The difference of FSD spectra (β -Amyloid exposed to magnetic field – β -amyloid without magnetic field)

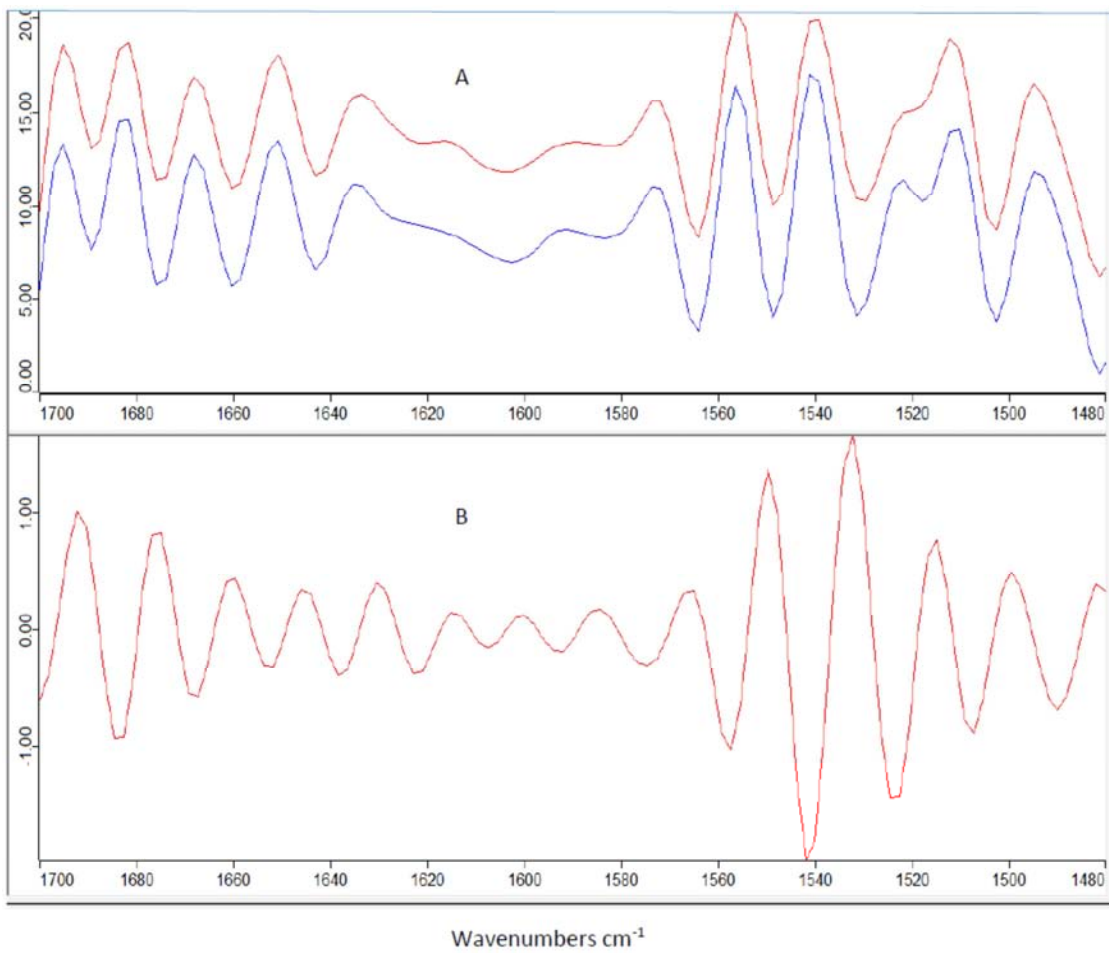


Figure 7. Difference of FSD spectra for HSA.

A- The two FSD spectra exposed to magnetic field (above) and no field (below)
 B- The difference of FSD spectra (HSA exposed to magnetic field – HSA without magnetic field)

The differences in the FSD spectra for each of $A\beta$ and HAS before and after exposure are shown in figures 6 and 7, respectively. Figure 6 shows the difference spectra which results from: {FSD spectra of $A\beta$ exposed to magnetic field — the FSD spectra of $A\beta$ }. The difference represents the net effect of the magnetic field on $A\beta$ as shown in figure 6-B, where it indicates

that the majority of the observed peaks have been affected in different ways: some show increase of intensity, others shifted in their peak positions. These variations in change pertaining to $A\beta$ have been listed in table 1. Similarly, changes for the HSA sample in the same range are shown in figure 6 and variations in changes are listed in table 2.

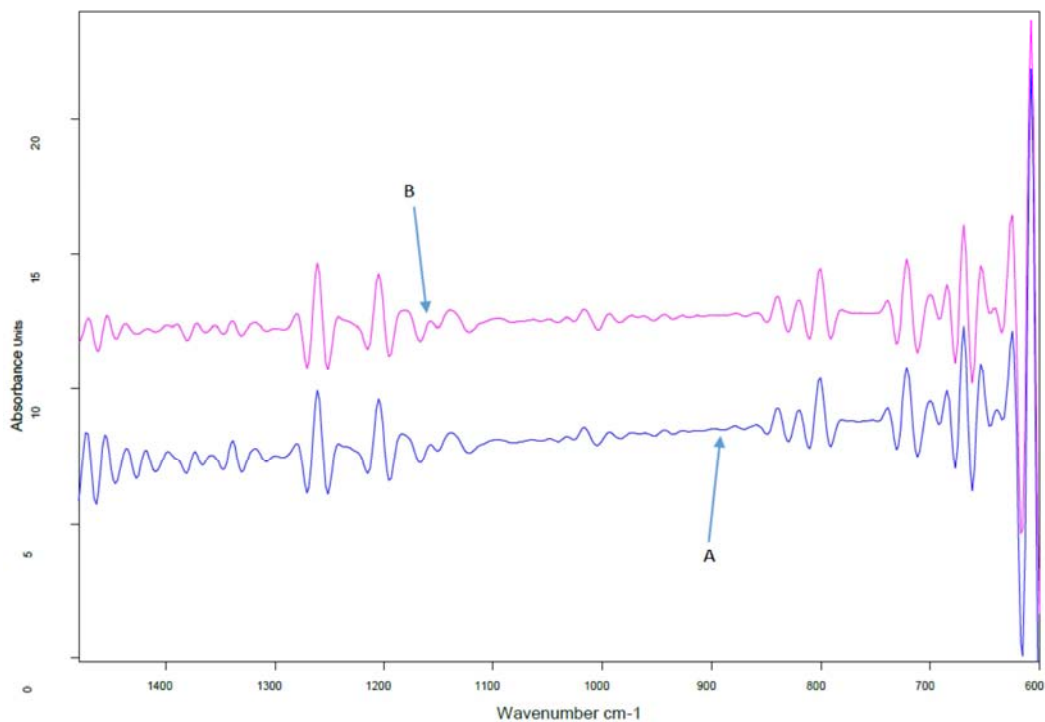


Figure 8. FSD spectra of $A\beta$ before and after exposure to ELF magnetic field.

A- Before exposure to magnetic field
B- After exposure to magnetic field

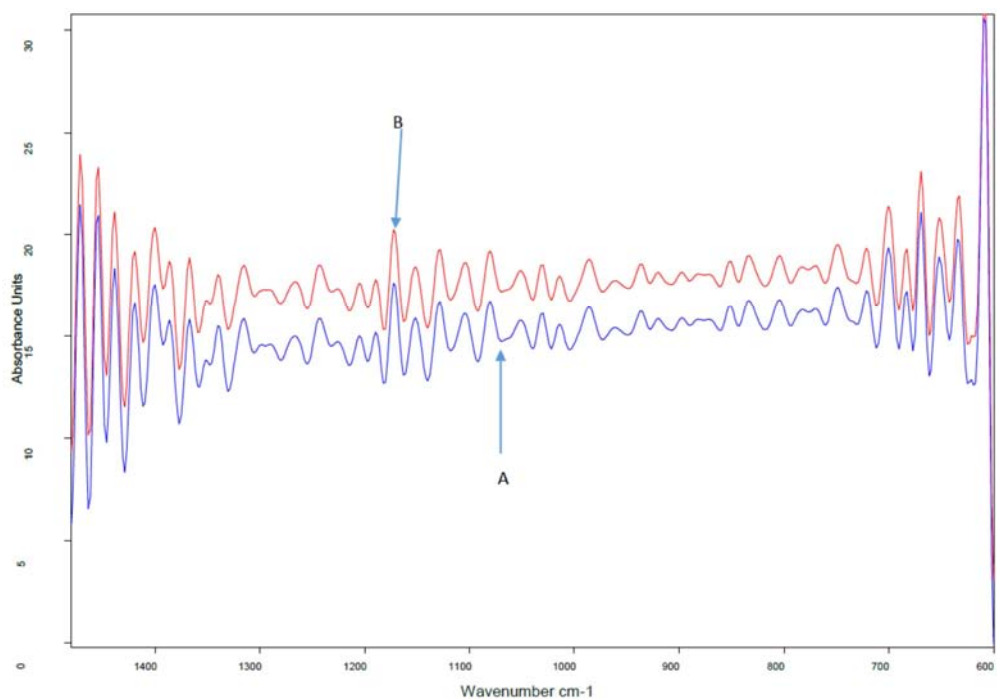


Figure 9. A- FSD spectra of HSA not exposed to magnetic field B- FSD spectra of HSA exposed to magnetic field.

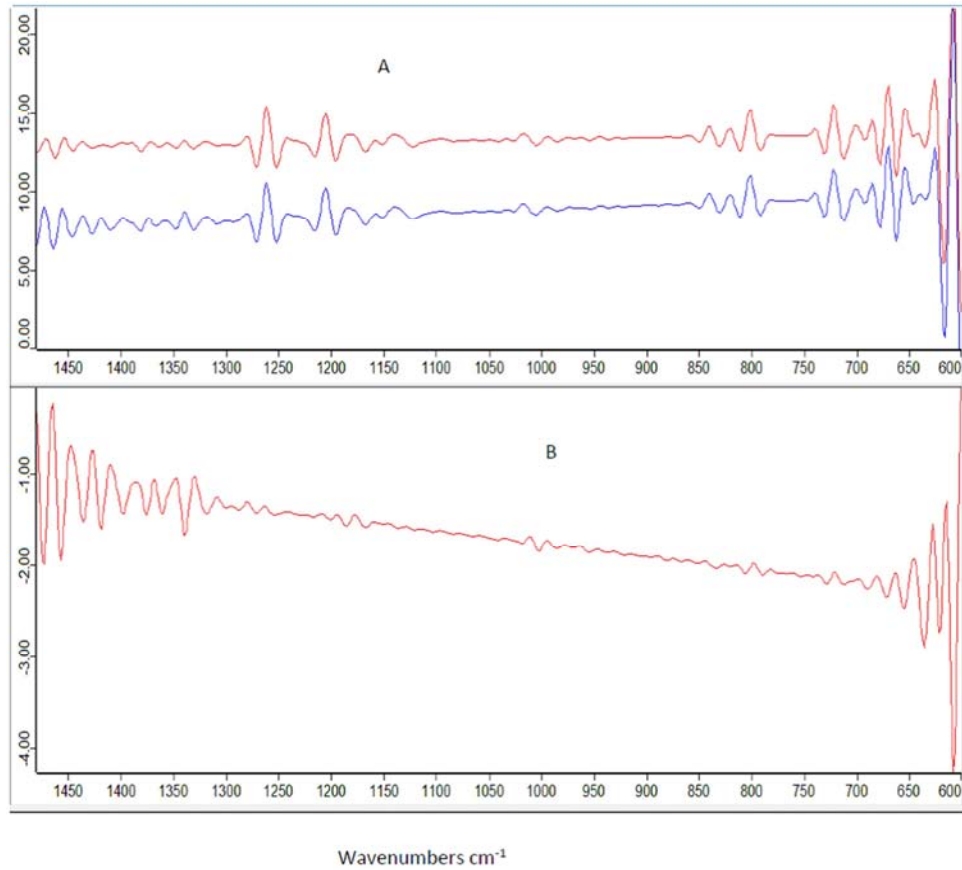


Figure 10. A: The FSD spectra for β -amyloid exposed to ELF 1.5 mT (above) and not exposed (below). B: The difference between $A\beta$ exposed and not exposed to magnetic field.

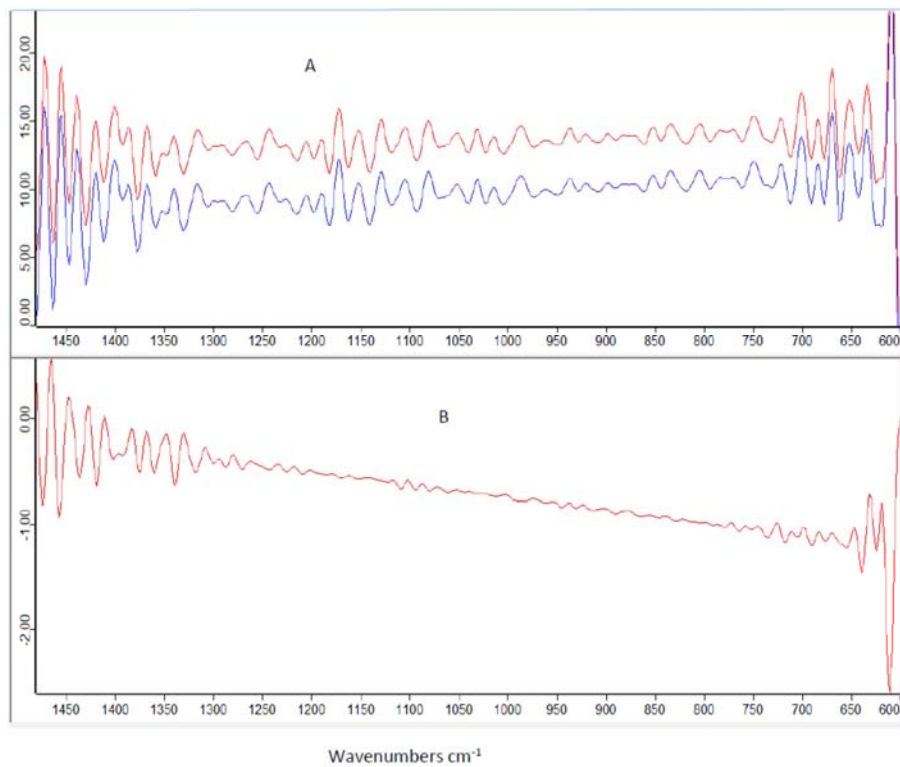


Figure 11. A- The FSD spectra of HSA exposed to ELF 1.5 mT (above) and not exposed (below) B- The difference between HSA exposed and not exposed to magnetic field.

Table 3. Peak positions of the absorbance spectra in the range (800-600) cm^{-1} .

A-Peak positions of the $A\beta$ absorption spectra		
$A\beta$ (not exposed)	Exposed to magnetic field	Difference spectra
608	608	---
---	---	613
625	625	626
640	641	644
653	653	661
669	669	680
685	685	--
700	700	--
721	721	--
739	739	--
782	782	--
800	800	--
B-Peak positions of the HSA absorption spectra in the range		
HSA (not exposed)	Exposed to magnetic field	Difference spectra
608	608	--
620	620	618
633	633	629
651	651	646
670	670	669
682	682	681
700	700	697
720	720	710
749	749	--
784	784	--
804	804	--

The amide III and the fingerprint region are included in the range (1480-600) cm^{-1} have been treated in a similar way and the results are shown in figures: 8, 9, 10 and 11, while the

observed changes are listed in tables 1 and 2 for the amide III bands, the observed changes in the fingerprint region are listed in table 3.

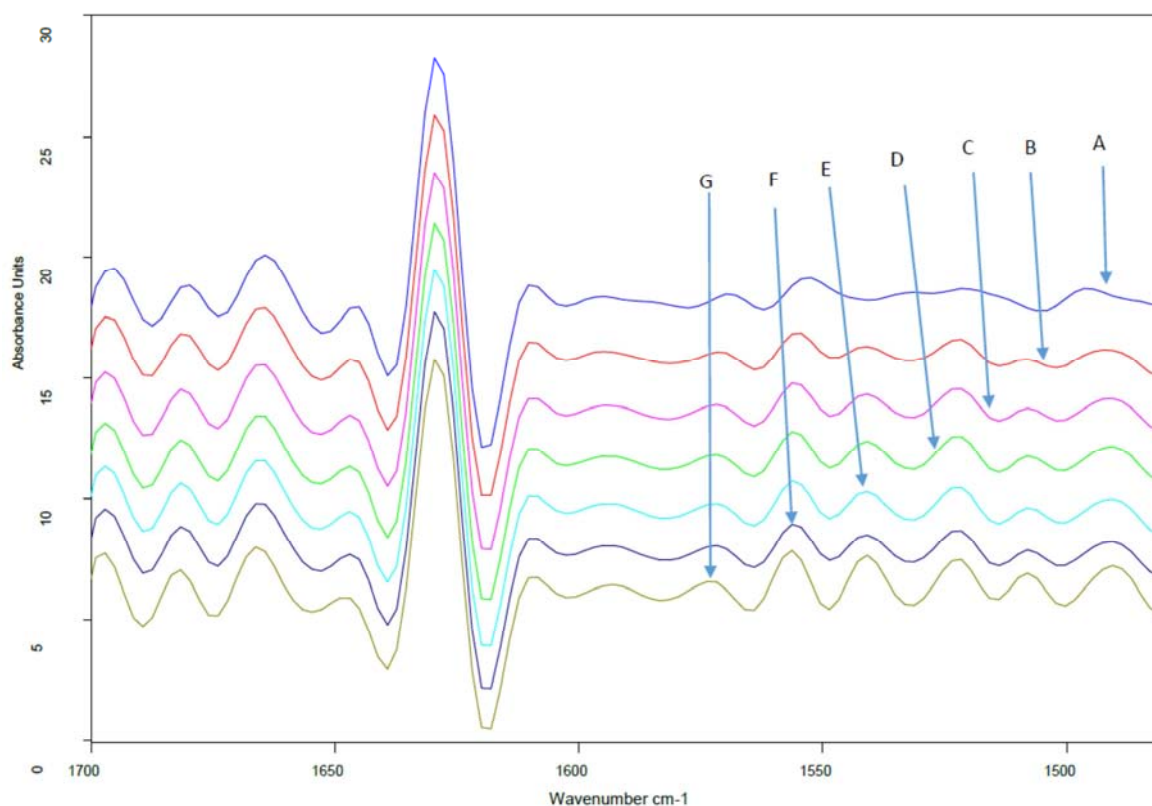


Figure 12. FSD spectra of $A\beta$ when (A- exposed to magnetic field of 1.5 mT), (B- 2 minutes after removal of field), (C- 4 minutes after removal), (D-6 minutes after removal), (E- 8 minutes after removal), (F- 10 minutes after removal), (G- No field).

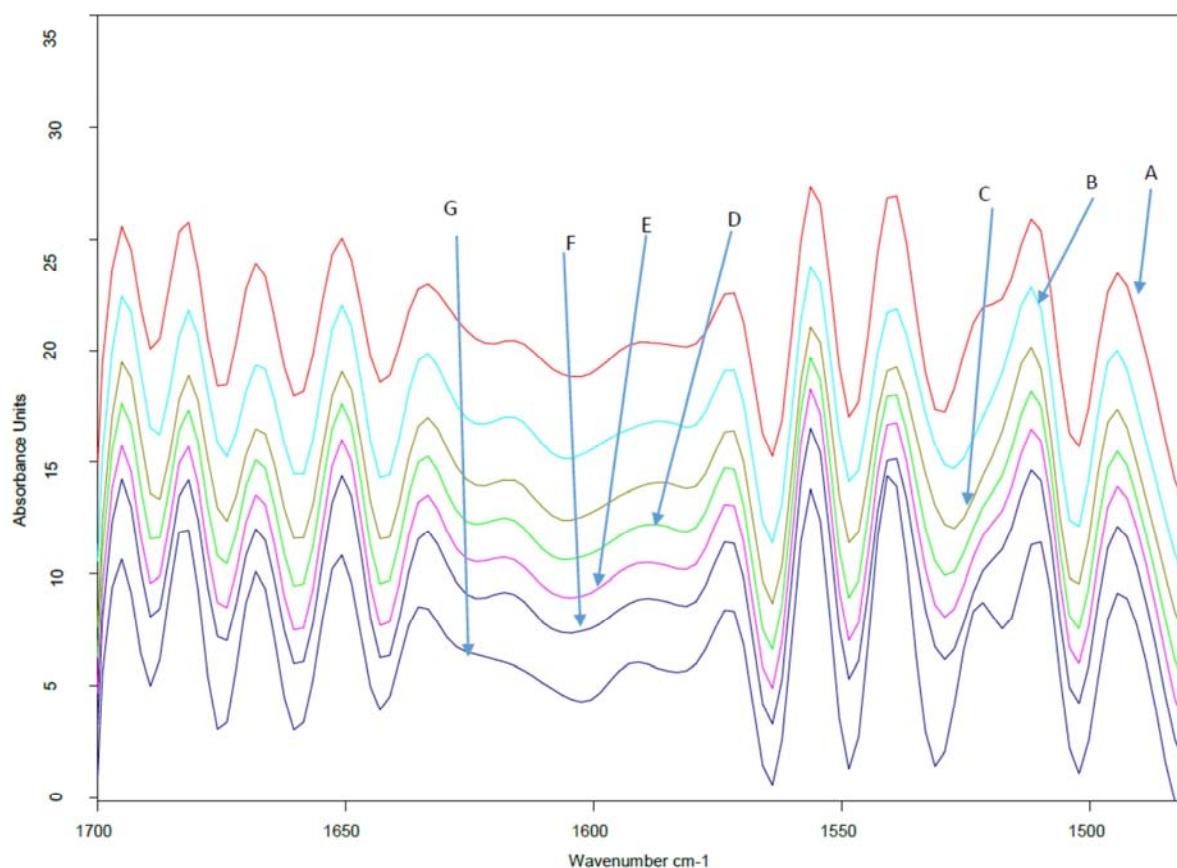


Figure 13. FSD spectra of HSA when (A-exposed to magnetic field), (B-2 minutes after removal of field), (C-4 minutes after removal), (D- 6 minutes after removal), (E- 8 minutes after removal), (F-10 minutes after removal), (G- before exposure to field).

The last part of the data is taken to investigate any possible remnant effect after removing the magnetic field on the samples. Each sample has been exposed to 1.5 mT for 10 minutes, then after turning off the magnetic field the spectra is taken every 2 minutes for five consecutive readings. The FSD spectra is shown in figures 12 and 13 for the A β and for the HSA samples respectively.

4. Discussions

In this work each of A β and HSA samples' FTIR spectra is obtained before and after exposure to the ELF magnetic field without moving or changing the sample position. Sample stationary position makes comparison of intensities and peak positions more accurate and minimizes any errors due to sample motion. Changes in the FSD convoluted graphs in figure 4 indicate secondary structure changes as a result of A β exposure to magnetic field. These changes can be summarized as follows: the bands at (1666, 1648, 1629, and 1609) cm^{-1} have shown little or no increase of intensity, while the rest of the bands in the figure have shown a sharp decrease of intensity. The following bands (1508, 1522, 1556, and 1573) cm^{-1} have shown large loss of intensity due to the effect of the magnetic field exposure. These remarks are also supported by figure 6-B, where the net effect of the magnetic field is shown to be strong near the above positions as listed in table 1.

The FSD convoluted spectra in figure 5 indicate secondary structure changes upon HSA exposure to magnetic field. The majority of the bands in amide I and amide II lost intensity due to the magnetic effect at different proportionality with the exception of two bands at (1633, and 1616) cm^{-1} . These two bands showed a little increase of their intensities. The shoulder at 1616 cm^{-1} has been enhanced strongly in intensity to become observable as peak by the effect of the magnetic field.

The peaks positions' changes for A β and HSA are listed in tables 1 and 2 respectively. The positions' changes include various range of changes and while some peaks have shifted positions by larger amounts, others retained their energy positions. In general, the peak position can be correlated to the exact frequency of the bond vibrations of the particular amide region, which also relies on the nature of the hydrogen bonding with the amide group, and by the particular secondary structure adopted by the protein. The difference of the FSD spectra, plotted in figures 6-B and 7-B for A β and HSA respectively, show that all the peaks have been affected by the magnetic field at different levels. The peaks of the difference spectra show the frequencies at which maximum effect by the magnetic field occurred.

In general a peak in the difference spectra represents a gain of intensity due to magnetic field effect, which is caused by a constructive interference at that frequency of the peak position. While an inverted peak position represents a loss of intensity due to the magnetic field caused by a destructive

interference at that frequency. If the difference spectra is a straight line, it indicates no contribution from the magnetic field implying no interference or energy exchange took place. A strong variations of the difference spectra in a particular region indicates a high degree of sensitivity to magnet effect in that region.

The quenching effect on the spectra of amide II region was stronger than the quenching effect on the spectra in the amide I region. [47] This difference indicates that the magnetic field has more damping influence on the out of phase combination of N-H, in-plane bending, and C-N stretching vibrations modes than on the in-phase combination of C=O stretching vibrations coupled to the C-N stretching.

The changes in the FSD spectra in the region (1480 - 1220) cm^{-1} which includes the amide III band are shown in figures 8 and 9 for A β and HSA respectively. The peak positions for this part of the spectra are also summarized in table 1 for A β bands and in table 2 for the HSA bands.

The changes in the FSD spectra of A β for the range of (1220 to 600) cm^{-1} are shown also in figures 8 and 10 however, the peak positions are summarized in table 3-A. It seems that most of the peaks show very little or no change in terms of their peak positions and in their intensities.

The difference spectra in figure 10-B, yields very little variations indicating little or no effect being imposed on most of the peak positions and their intensities by the magnetic field. The main changes are limited to the appearance of a new peak at 613 cm^{-1} , and few peaks changed positions as follows: (625 \rightarrow 626, 640 \rightarrow 644, and 653 \rightarrow 661) cm^{-1} .

The changes in the FSD spectra for the HSA in the range (1220-600) cm^{-1} are shown in figures 9 and 11 and the peak positions are listed in table 3-B. Most bands show little or no change in their peak positions and in their relative intensities due to the exposure to the magnetic field.

The difference spectra in figure 11-B shows peak positions shift as follows: (633 \rightarrow 629, 651 \rightarrow 646, 670 \rightarrow 669, 682 \rightarrow 681, 700 \rightarrow 697, and 720 \rightarrow 710) cm^{-1} . In addition to a strong magnetic field effect on band intensity occurring at (618, 629 and 646) cm^{-1} and very little effect on all other bands.

This implies that for both proteins the absorption bands in amide IV region (625-765) cm^{-1} , which are mainly caused by combinations of OCN bending vibrations mixed with out of phase N-H bending [48, 49] are highly sensitive to magnetic field effect. It is interesting to note that both regions amide II and amide IV show sensitivity to magnetic field exposure, and both bands caused by out of phase combinations of N-H bending mixed with OCN bending or stretching. One may speculate that out of phase combination of molecular vibrations leave molecules more susceptible to be twisted by the magnetic field.

The qualitative analysis of the secondary structure for A β and HSA proteins is based on the observed changes in peak position, band intensity and band width for the individual bands. It is factually accepted that general vibrational spectroscopy is highly sensitive and can detect any small change (0.02%) in bond strength [50]. Any changes in bond strength or the number of bonds is reflected in a change of

the band relative intensity.

Our experimental results show a large loss of intensity to most of the bands in the amide II region for both A β and HSA and a small decrease of intensity to some of the bands in amide I. The loss of intensity implies that exposure to magnetic field introduces a damping factor which alters molecular vibrations at the bonding sites. This may be explained by a destructive interference imposed by the magnetic field on the out of phase combinations of C-N stretching and N-H bending. On the other hand, the magnetic field did not interfere with the in-phase combinations of N-H bending and C-N stretching in the amide III region and minor changes were observed on its bands' intensities.

The interference of the magnetic field with molecular vibrations can be caused by the Hall Effect. Free charges tend to move by the magnetic force and the positive charges accumulate on one side while the negative charges accumulate at the opposite side creating region with potential difference. The accumulated charges on the opposing sides exert opposing forces on the ends of the vibrating molecule and produce a twisting effect on its vibration.

In the presence of a magnetic field in a medium several forces can influence the motion of charged particle. For example Lorentz force ($F=qv \times B$), and the magnetic gradient force ($F=(p_m \nabla) B$), where q and v are the charge and the velocity of the particle, B is the magnetic induction, and p_m is the dipole magnetic moment. These forces can have an impact but to a lesser degree than the impact of electric forces [51]. When the vibrations of the atoms are out of phase, these forces may produce a stronger torque on the molecules than the vibrations of the in-phase combinations case. This may explain why the peak positions of amide II bands are shifted to higher energies while at the same time lost some of their intensities. Physically these bonds are becoming stiffer with a larger spring force constant due to the twisting effect and losing intensity due to the damping effect by the Hall Effect. It is suggested that twisting of antiparallel beta sheets can also leads to a shift to higher energy by previous studies [52, 53].

These bands whose peaks shifted to a higher energy indicate a change in the energy of molecular vibration due to the increase in the spring restoring force, which is caused by additional hydrogen bonding. It is well known that hydrogen bonds help in stabilizing proteins which is needed for catalysis. Therefore, additional hydrogen bonds can lower the frequency of a stretching vibration. On the other hand, less hydrogen bonding decreases the strain on the molecules can shift the peak position to lower energy [54]. It is suggested that each of the two possible hydrogen bonds to the C-O group lowers the amide I frequency by 20 to 30 cm^{-1} and a hydrogen bond to the NH group by 10 to 20 cm^{-1} [55, 56].

The small increase in the intensity of β -sheets in amide I for A β spectra at 1629 cm^{-1} and for HSA spectra (1635 and 1616) cm^{-1} occurred after ELF magnetic field exposure can be attributed to disorder and unfolding processes in the structure of the protein [57-59].

The peak positions of the bands in amide I and amide III

are hardly affected by the number of amide groups in the strand of the sheet, but depends on the number of strands. Sheets with a larger number of strands have a lower spectral position of the main band [52, 60].

The effect of exposing A β and HSA to ELF magnetic field is examined further by monitoring spectra changes after removing the magnetic fields. Figures 12 and 13 show the spectra changes as a function of time. It seems that after removing the field, the affected absorbance intensities and the shift of the band positions takes on a gradual change slowly returning to their control levels, obtained before the exposure. The changes in the spectra are monitored every two minutes, but it is clearly evident that these two proteins need more than 10 minutes to revert to their initial control states.

This behaviour implies that the conformational changes on the secondary structure of the proteins are not momentarily, and the proteins' exposure to the magnetic field leads to internal energy changes in the proteins. Exposure to magnetic field can cause charge motion and induce Eddy currents within the protein. But these new currents induce a new magnetic field to counter the effect of the original magnetic field according to Lenz's law. Therefore, the protein may act as an inductor-capacitor circuit which causes energy oscillation and that leads to a time delay for a full recovery and the return to the original state.

5. Conclusion

The experimental results showed that A β and HSA are sensitive to ELF magnetic field exposure and the following changes have been observed: (1) Peak positions' shifts and intensity changes for amide II bands were highly noticeable. (2) A small decrease in band intensities of antiparallel β -sheets and β -turns in the amide I has been detected versus a small increase of intensities of α -helix and parallel β -sheets in that region. (3) No major changes in the amide III region have been observed (4) the region between (720-600) cm^{-1} has shown high sensitivity to ELF magnetic fields, and (5) Recovery of the magnetic effects are not instantaneous.

This study suggests that ELF magnetic fields can interact with proteins and may promote biological tasks or cause a dysfunction for the protein depending on whether the exchanged energy is appropriate for a particular task or not.

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