



Article

Application of FTIR and LA-ICPMS Spectroscopies as a Possible Approach for Biochemical Analyses of Different Rat Brain Regions

Mohamed H. M. Ali ^{1,*}, Fazle Rakib ², Volker Nischwitz ³, Ehsan Ullah ⁴, Raghvendra Mall ⁴, Amjad M. Shraim ², M. I. Ahmad ⁵, Zafar Khan Ghouri ⁶, Donald McNaughton ⁷, Stephan Küppers ³, Tariq Ahmed ¹ and Khalid Al-Saad ²

- Qatar Biomedical Research Institute (QBRI), Hamad Bin Khalifa University (HBKU), Doha 34110, Qatar; taahmed@qf.org.qa
- Department of Chemistry and Earth Sciences, Qatar University, Doha 2713, Qatar; fazle@qu.edu.qa (F.R.); amjad@qu.edu.qa (A.M.S.); kalsaad@qu.edu.qa (K.A.-S.)
- Forschungszentrum Jülich GmbH, Zentralinstitut für Engineering, 52425 Elektronik und Analytik, Germany; v.nischwitz@fz-juelich.de (V.N.); s.kueppers@fz-juelich.de (S.K.)
- Qatar Computing Research Institute (QCRI), Hamad Bin Khalifa University, Doha 34110, Qatar; eullah@hbku.edu.qa (E.U.); rmall@hbku.edu.qa (R.M.)
- ⁵ Central Laboratories Unit, Qatar University, Doha 2713, Qatar; mohammad.ibrahim@qu.edu.qa
- 6 Chemical Engineering Program, Texas A&M University at Qatar, Doha 23874, Qatar; zafarkhanghouri@hotmail.com
- Centre for Biospectroscopy and the School of Chemistry, Monash University, Clayton, Victoria 3800, Australia; donald.mcnaughton@monash.edu
- * Correspondence: mohamali@hbku.edu.qa

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Abstract: Fourier Transform Infrared Spectroscopy (FTIR) is a non-destructive analytical technique that has been employed in this research to characterize the biochemical make-up of various rat brain regions. The sensorimotor cortex, caudate putamen, thalamus, and the hippocampus were found to have higher olefinic content—an indicator of a higher degree of unsaturated fatty acids—rich in short-chain fatty acids, and low in ester and lipid contents. While the regions of the corpus callosum, internal, and external capsule were found to contain long-chained and higher-esterified saturated fatty acids. These molecular differences may reflect the roles of the specific regions in information processing and can provide a unique biochemical platform for future studies on the earlier detection of pathology development in the brain, as a consequence of disease or injury. Laser Ablation Inductively Coupled Plasma Mass Spectroscopy (LA-ICP-MS) is another vital analytical technique that was used in this work to analyze the elements' distribution patterns in various regions of the brain. The complementary data sets allowed the characterization of the brain regions, the chemical dominating groups, and the elemental composition. This set-up may be used for the investigation of changes in the brain caused by diseases and help create a deeper understanding of the interactions between the organic and elemental composition.

Keywords: FTIR imaging spectroscopy; biochemical analysis; brain; LA-ICP-MS

1. Introduction

The mammalian brain is sub-divided into numerous regions, each with a specific physiological function. Dissection down the midline identifies the two most visible features—an interior core of white matter (WM), composed of phospholipid rich myelinated axons, and an exterior layer of

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gray matter (GM), composed of arborized neurons with un-myelinated axons [1]. Due to the brain's complexity, there remains an important need for novel technologies that further delineate the (sub-) cellular components, in order to monitor and characterize the bio-chemical changes associated with neurological dysfunctions. The drawbacks of the current bio-diagnostic technologies are that they are too time-consuming, provide low resolution, and alter tissue integrity.

Current imaging techniques, such as positron emission tomography (PET), magnetic resonance imaging (MRI), and computerized tomography (CT), are used to identify disease pathology. CT imaging is rapid, highly reproducible and low in cost for clinical brain-injury screening. However, it is not equipped to detect all types of lesions in the brain, specifically, those related to the development of the secondary injury. MRI has long been the gold standard to study changes in gross brain morphology, and has been routinely used to examine the peri-infarct zone, following a stroke. Unfortunately, MRI does not have sufficient spatial resolution to study individual cells within the brain and reveals little information on the biochemical mechanisms of tissue insult [2].

Infrared (IR) spectra originate from the vibrational motions of atoms, in chemical bonds [3]. As the molecule absorbs the electromagnetic radiation, it produces a band in the IR spectrum, at specific frequencies [4]. The IR spectra vibrational frequencies provide information about chemical compounds, including bio-molecules, and produce characteristic fingerprint for proteins, lipids, cholesterols, phospholipids, carbohydrates, and nucleic acids, as well as providing a unique biochemical and biophysical fingerprint for the sample [5]. Fourier transform infrared (FTIR) spectroscopy is a simple, sensitive, accurate, highly-reproducible, non-destructive (with respect to sample preparation), and rapid, diagnostic technique [6]. In contrast to the standard histological staining methods, FTIR is the "state-of-the-art" bio-diagnostic and bio-imaging technology for fast diagnosis in medicine and biological studies. It has been used to screen and characterize a wide range of tissues and organs [7–9]. FTIR imaging spectroscopy has also been used to analyze the pathological changes associated with stroke, traumatic brain injury (TBI), and neurodegenerative diseases, such as Alzheimer's disease (AD), Parkinson's disease (PD), and multiple sclerosis (MS) [10–12]. We report here that FTIR technology is relatively straightforward and rapid. There is no requirement of prior treatment of brain samples with dyes, or reagents, in order to characterize the different brain structures, and the technique, in addition, is non-destructive, which allows the sample to be analyzed with additional techniques. Hence, FTIR can be a fast, reproducible technique, in order to characterize the biochemical make-up of various brain regions, to provide a good platform for investigating brain alterations at the molecular-level, such as in the cases of neurological disorders. Furthermore, a combined analysis of FTIR and Laser Ablation Inductively Coupled Plasma Mass Spectrometry (LA-ICP-MS) imaging can improve the brain tissue elemental characterization and classification, expediting the diagnostics and therapeutics. LA-ICP-MS has been shown to be a powerful technique for brain tissue imaging [13–17]. One disadvantage is that it is a destructive technique but, in combination with FTIR, the same sample can be analyzed to overlay the information of both techniques and can show changes, especially with respect to diseases, in the aging of brains. Specifically, the combination may be a tool to easily classify different diseases and may help to evaluate the biochemical impacts of different diseases.

2. Materials and Methods

2.1. Sample Preparation

All experimental protocols and animal handling procedures were approved by the University of Utrecht Animal Research Ethics Board. A total of 6 male (11-week-old Sprague-Dawley rats), were included in the study. The animals were sacrificed with an overdose of isofluorane, followed by transcardial perfusion. Then, the brains were extracted, embedded in paraffin, and the paraffin-embedded brain tissue blocks were serially cut into 10 µm sections, using a microtome (Leica RM 2155 semi-automated rotary microtome, Nussloch, Germany). All samples were treated in an identical manner, during sample processing and preparation. The areas of interest consists of

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seven brain regions including the somatosensory motor cortex (SSMC), the corpus callosum (CC), the caudate putamen (CP), from the Bregma 1.56 mm, and the external capsule (EC), the capsule interna (IC), the hippocampus (HC), and the thalamus (TH) from the Bregma 4.36 mm. From each rat, five brain sections of similar anatomy were collected and twenty FTIR spectra were recorded from each brain region. Serial and consecutive sections were collected for the LA-ICP-MS and a routine hematoxylin and eosin (H&E) staining analyses [2].

2.2. Fourier Transform Infrared Spectroscopy (FTIR)

FTIR spectra were recorded using the FTIR Cary 620 spectrometer in the reflection mode, within the range of 4000–700 cm $^{-1}$, with 64 scans per spectrum and a 4 cm $^{-1}$ spectral resolution. Background single beam spectra were measured on a substrate, without the biological tissue, by co-adding sixty-four scans. FTIR images were recorded using a 64 \times 64 focal plane array instrument with Mercury Cadmium Telluride (MCT, California, USA), Agilent Technology. FTIR imaging was performed with an objective of 15 \times magnification and 0.62 numerical aperture, yielding a pixel size of 5.5 \times 5.5 μ M.

2.3. FTIR Data Processing and Analysis

Spectra acquired from the border between a brain tissue and a substrate, affected by resonance Mie scattering (commonly present in biological samples) and spectra recorded from a substrate (outside the brain tissue area) were eliminated from further analysis [18]. FTIR spectra were vector normalized between 3500–900 cm⁻¹ and the baseline correction was performed on the full range of wavelengths for the spectra. The relative amount of the specific molecular content was quantified from the original absorbance spectra [2]. The approximate values were determined from the brain samples of different animals, based on their bio-chemical variations. Functional group images were obtained by calculating the ratios of the baseline corrected areas, under some peaks, based on Table 1 (see Figure 1). The total lipid content was represented by the C-H stretching region at 2994–2800 cm⁻¹ [19,20]. The protein concentration was represented by the amide II band at 1555-1535 [21,22]. For more insight and deep lipid analysis, the ratios of some specific lipid spectral bands ($v_{as}(CH_3)$ asymmetric, $v_s(CH_2)$ symmetric, $v_{as}(CH_2)$ asymmetric stretching, carbonyl ester v(C=O) ester, and olefinic =CH bands), to the total lipid (C-H stretching region), were calculated. For each animal, a mean and standard deviation was calculated and the average (n = 6) was generated. Mann Whitney-U test was performed on all data to test for significant differences (p < 0.05). All t-tests were two-tailed and the 95% confidence limit was used to test for significance.

Infrared Band	Spectral Rage (cm^{-1})		
Lipid components			
CH ₂ symmetric stretching	2852-2800		
CH ₂ asymmetric stretching	2915-2930		
CH ₃ asymmetric stretching	2950-2960		
* C-H stretching	2994-2800		
Olefin=CH	3000-3027		
Carbonyl ester (C=O) stretching	1745–1731		
Protein components			
Amide I	1700-1600		
Amide II	1555–1535		

Table 1. The spectral regions used for the analytical purposes.

Principal component analyses (PCA) were performed using "prcomp" function in the "stats" package in R. The PCA analysis was performed on the spectral second derivatives, in the range of 4000–700 cm⁻¹ (the range of 2500–2000 cm⁻¹ removed). PCA analysis was performed on a total of

^{*} Total lipid region.

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1270 wavenumbers, in the range of 4000–700 cm⁻¹, from the FTIR spectra obtained from the different brain regions, and PCA was performed, based on the covariance of these features. Furthermore, during PCA we randomly divided the spectral collection into 80% training and 20% test. The norm of the reconstruction error for each test spectra was 0.05, on average, while using the first three principal components. The 3D scores plot of these components enabled the spectra to be grouped according to the chemical information they contained. Hierarchical cluster analysis (HCA) was applied, using the "A2Rplot", to compare the seven brain regions, based on their PCA projections in the 3D score plot. HCA was used to group spectra that displayed the same degree of similarity. The result was visualized in a dendrogram and the grouping of the seven brain regions were presented in clusters, according to a heterogeneity scale [23,24].

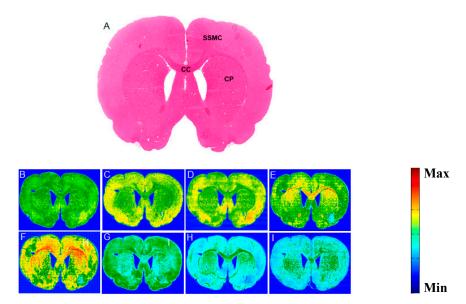


Figure 1. (**A**) H&E staining section of the rostral brain (**Bregma 1.56 mm**) showing the three regions of interest (ROI): SSMC, CC, and CP. (**B**) Corresponding FTIR image. (**C–I**) Representative images of the biochemical components—Amide I, Amide II, CH₃ asymmetric, CH₂ symmetric, CH₂ asymmetric, Olefin, and Ester, accordingly. The color bar represents a high concentration, in red, and low concentration, in blue.

2.4. Laser Ablation-ICP-MS

An Inductively Coupled Plasma Quadrupole Mass Spectroscopy ICP-QMS (Agilent 7900, Agilent Technologies, Hachioji, Japan) was coupled to a laser ablation system NWR 213 (New Wave Research, CA, USA) for the elemental imaging. The laser spot size was 60 µm with 30 µm between the lines, at a laser energy of about 29%. One measurement of each tissue sample was performed, with a scan speed of 70 μ m s⁻¹. The isotopes ¹³C, ³¹P, ³⁴S, ⁴³Ca, ⁴⁴Ca, ⁵⁴Fe, ⁵⁶Fe, ⁵⁷Fe, ⁶³Cu, ⁶⁵Cu, ⁶⁴Zn, ⁶⁶Zn, ²⁰⁶Pb, ²⁰⁷Pb, and ²⁰⁸Pb were monitored; in case of multiple available isotopes the most abundant was evaluated and the results were cross-checked with the additional isotopes of the same element, to check for possibly occurring spectral interferences. Quantification of the Fe, Cu, Zn, and Pb was performed, based on the rat brain-tissue standards. Blanks and five standards were prepared by spiking the homogenized rat brain tissue with elemental standard solutions. The resulting tissue, with the known concentrations (determined by the total elemental analysis), was frozen, cut in slices, and analyzed, in analogy to the samples. The calibration ranged from the blank tissue levels up to 19.8 mg/kg for Fe, up to 20 mg/kg for Cu, up to 30 mg/kg for Zn, and up to 2.5 mg/kg for Pb. The relative intensities of C, P, S, and Ca, were seen due to the lack of calibration standards. Image reconstruction was performed by a software developed in-house (Forschungszentrum Jülich), giving the scale in mg/kg, for the elements with calibration. Due to the complex process of ablating and transferring the aerosol into the ICP-MS, Appl. Sci. 2018, 8, 2436 5 of 13

there are several sources of uncertainty. Therefore in this work, the main focus was on identifying the elemental patterns within the tissue sections but not on an exact quantification.

3. Results

The formalin-fixed paraffin-embedded brain tissue samples (serial sections of 10 µm thickness), were collected from each rat for the FTIR, LA-ICP-MS, and the H&E analyses. The numerical comparisons of the lipid/protein ratio and the ratios of some specific lipid spectral of the regions of interest of this study, are listed in Table 1. Figure 1A shows a rostral section (Bregma 1.56), highlighting three other regions of interest—SSMC, CC, and CP; the corresponding FTIR image is shown in Figure 1B. Figure 1C-I shows the images of the biochemical components—Amide I, Amide II, $v_{as}(CH_3)$ asymmetric, $v_s(CH_2)$ symmetric, $v_{as}(CH_2)$ asymmetric stretching, olefin, and ester, respectively. These images were colored, according to the calculated biochemical content values, where red color corresponded to the highest bio-content and the blue color corresponded to the lowest bio-content, as shown on the color bars in the figure. The FTIR spectra of the SSMS, CC, and CP are shown in Figure 2A-C, respectively. Figure 2A represents the SSMC average FTIR spectrum. The spectrum revealed the ester absorption band that arose at ~1730 cm⁻¹ from the v(C=O) ester (carbonyl) group, had weak absorption. The FTIR spectrum revealed that a robust olefinic=CH band arose at about 3012 cm⁻¹. This was interpreted as that the SSMC structure contains lipids with a high degree of unsaturation. The protein absorption bands, such as amide II, amide I, and the N-H bands, arose at 1545, 1653, and 3290 cm⁻¹, respectively. The amide I and amide II have strong absorption.

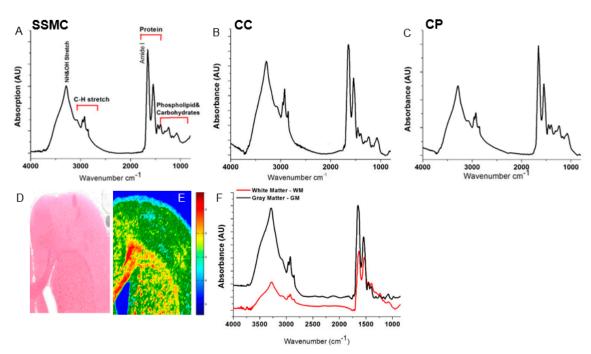


Figure 2. (A–C) Representative FTIR spectra of the SSMC, CC, and CP. (**D**) H&E images of the WM and the GM, and (**E**) FTIR image of the same region with a high-resolution, based on the lipid/protein ratio. (**F**) FTIR spectra of the WM and the GM. The color bar represents a high-concentration, in red, and a low-concentration, in blue.

The spectral data collected from the CC, in Figure 2B, revealed a strong peak which ascended from the ν (C=O) at 1470 cm⁻¹, which was attributed to the phospholipid. The FTIR spectrum revealed that the olefinic=CH band was weaker in the CC, than in the SSMC. Whereas, the FTIR spectrum identified that the band at 1464 cm⁻¹, which arose from the CH₂ bending vibration, was robust. The bands at 2852 and 2955 cm⁻¹ that arose from the ν_s (CH₂) and ν_{as} (CH₂), showed stronger absorption than

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the SSMC spectrum. The band at 2955 cm⁻¹ which arose from the $v_{as}(CH_3)$ of the lipids fatty acids, showed weaker absorption, in comparison to the SSMC spectrum.

Figure 2C is a representative FTIR spectrum of the CP. Here, the spectral band of the carbonyl group $\nu(C=O)$ of the ester component, had a low-absorption intensity, due to the low esterified-phospholipid content. The $\nu_s(CH_2)$ and $\nu_{as}(CH_2)$ spectral bands showed a weaker absorption, in comparison to the CC spectrum. The FTIR spectrum at the CP region exhibited strong olefinic=CH band, in comparison to the CC spectrum; indicating that the CP was rich in unsaturated lipids The FTIR spectrum was dominated by amide I and amide II spectral bands.

Figure 2D is a representative H&E image of the WM and GM, with the corresponding FTIR image obtained with a higher resolution, based on the lipid/protein ratio presented in Figure 2E. The data revealed here, that the white matter showed a higher intensity in the lipid/protein ratio signals, which correlated well with the presence of the myelin sheaths around the nerve axons located in this region. The spectra collected from these regions were dominated by the protein, lipid, and the ester bands (Figure 2F). The WM spectrum had two strong absorption bands, at 2850 and 2920 cm⁻¹, compared to the spectrum collected from the gray matter. These bands arose from the C-H stretching vibrations of the $\nu_s(\text{CH}_2)$ and $\nu_{as}(\text{CH}_2)$, respectively, from the fatty-acid content. We found from these results that there was a slightly stronger absorption in the ester region $\nu(\text{C=O})$, at 1740 cm⁻¹, for the WM, confirming that the WM is lipid rich.

Figure 3A is an example of the H&E image of a caudal brain section that represented three regions of interest: HC, TH, and EC. Figure 3B shows the corresponding FTIR image from these sub-fields. Figure 3C-I represent the biochemical components of this caudal brain section. The hippocampus H&E stain (Figure 4A), clearly shows the cyto-architecture of this organ. The figure revealed that the hippocampus has densely-packed neurons in the cornu ammonis (CA) sub-fields and the dentate gyrus (DG), and a smooth, less-dense upper region of the avleus (AV). The spectroscopic results (Figure 4B) indicate that there is a similarity between the CA and DG spectra, in contrast with the AV spectrum. The main differences were observed in the following spectral bands: (a) In the AV's FTIR spectrum, the intensity of the $v_s(CH_2)$, $v_{as}(CH_2)$, and the v(C=O) ester bands showed stronger absorption than the CA and DG spectra, indicating that the AV is enriched in lipid content. (b) The olefinic structure olefinic=CH band at 3012 was stronger in the CA and DG spectra than the AV spectrum. (c) The band at 2955 cm⁻¹ which arose from the $v_{as}(CH_3)$, was stronger in the CA and the dentate gyrus (DG) spectra than the AV spectrum. (d) The amide I and amid II bands exhibited a higher absorption in the in the CA and the DG spectra, in comparison to the AV spectrum. (e) The spectra revealed that the CA and DG regions yielded a low lipid-intensity, while the CH₂ spectral bands had a strong absorption, at the white matter location, which surrounded the entire hippocampus. These intense CH2 bands were due to the high lipid-bilayer concentration in the white matter.

The averaged spectral data, collected from the TH region (Figure 4C), revealed that the spectral bands of the $\nu_s(CH_2)$, $\nu_{as}(CH_2)$, and $\nu(C=O)$ ester, showed weaker absorptions; indicating that the esterified-phospholipid, in the TH, was low. The TH's FTIR spectrum exhibited strong amide I and amide II absorption bands. The FTIR spectrum revealed a high absorption in the spectral band around 3012 cm⁻¹, which arose from the olefinic=CH structure. These results indicated that the TH bio-chemical make-up contains lipids with a high degree of unsaturation. The FTIR spectrum of the TH yielded strong $\nu_{as}(CH_3)$ bands at 2955 cm⁻¹, which arose from the lipid structure, and a PO_4^{2-} band at 1240 cm⁻¹, which arose from the stretching vibrations of the nucleic acids.

The CE averaged spectrum (Figure 4D) identified that the $\nu_s(\text{CH}_2)$ and $\nu_{as}(\text{CH}_2)$ bands of the lipid content of the CE, showed a high absorption, due to the high lipid-content. The CE spectrum showed a high absorption peak at 1740 cm⁻¹ that arose from the $\nu(\text{C=O})$ ester band of the lipid structure. This $\nu(\text{C=O})$ band was found to be weaker in the CP region. These results indicated that the CP contains less lipid contents than the CE (see Tables 2 and 3). The spectrum revealed that the amide I and amide II bands had a high absorption. The FTIR spectrum for the capsule interna (IC) region had a large similarity between the FTIR spectrums collected from the capsule externa (data not shown).

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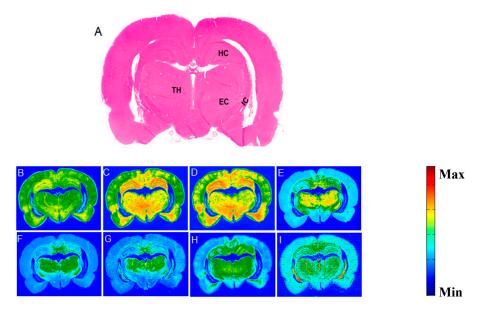


Figure 3. (**A**) H&E staining section of the caudal brain (**Bregma** −**4.36 mm**) showing four regions of interest (ROI): HC, TH, IC, and EC. (**B**) Corresponding FTIR image. (**C**−**I**) Representative images of the biochemical components—Amide I, Amide II, CH₃ asymmetric, CH₂ symmetric, CH₂ asymmetric, Olefin, and Ester, respectively.

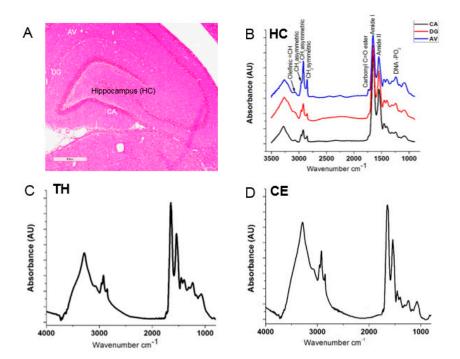


Figure 4. (A) H&E image of the Hippocampus showing AV, DG, and CA. **(B)** FTIR spectra of these three regions. **(C,D)** Representative FTIR spectra of TH and CE, respectively.

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Table 2. Differences in the band-area ratios of various functional groups, within the regions of interest (ROI), of a control healthy rat brain sections.

Ratio	SSMC	CC	CP	HP (CA)	TH	CE
Lipid/Protein	3.453 ± 0.170 *	6.902±0.204	3.143 ± 0.162	2.984 ± 0.144	3.251 ± 0.152 *	5.893 ± 0.215
CH ₂ /lipid	$0.2283 \pm 0.0019 *$	0.3213 ± 0.0028	0.2243 ± 0.0011 *	0.2015 ± 0.0012	$0.2146 \pm 0.0015 *$	0.2932 ± 0.0024
Carbonyl ester/lipid	0.0304 ± 0.0012	0.0382 ± 0.008	0.0291 ± 0.0010	0.0271 ± 0.0008	0.0264 ± 0.0014	0.0312 ± 0.0013
Olefinic=CH/lipid	0.0045 ± 0.0004	$0.0028 \pm 0.0002 *$	0.0043 ± 0.0003	0.0041 ± 0.0005	0.0044 ± 0.0010	$0.0023 \pm 0.0006 *$
CH ₃ /lipid	0.0741 ± 0.009	0.0582 ± 0.0007	0.0736 ± 0.013	0.0693 ± 0.008	0.0713 ± 0.007	0.0541 ± 0.0009

All the values are represented by the 'mean \pm SD' for each region of interest, since n = 6 and * p < 0.05.

Table 3. Identification and differentiation of the regions of interest (ROI), within the brain sections.

Characteristic Regions	Protein Region (Amide I, Amide II) at 1800–1480 cm ⁻¹	Lipid Region (C-H) Stretchii	ng 2994–2800 cm ⁻¹	ν(C=O) Carbonyl Ester at 1740 cm ⁻¹	Olefinic=CH at 3012 cm ⁻¹
		$\nu_s(CH_2)$ symmetric at 2852 and $\nu_{as}(CH_2)$ asymmetric at 2922 cm $^{-1}$	vas(CH3) asymmetric at 2955 cm ⁻¹		
SSMC	Strong	Moderate	Moderate	Weak	Moderate
СС	Strong	Strong	Weak	Moderate	Weak
СР	Strong	Moderate	Moderate	Weak	Moderate
HC (CA)	Strong	Moderate	Moderate	Weak	Weak
TH	Strong	Moderate	Moderate	Moderate	Moderate
EC	Strong	Strong	Weak	Moderate	Weak

Presented in Figure 5A, is a 3D image of the score plot of the PCA analysis for the seven brain regions, based on the spectral second derivatives, in the range of 4000–700 cm⁻¹ (the range of 2500–2000 cm⁻¹ removed). The score plot revealed the three main principal components that separate the seven brain regions and are accounted for by PC1 (90%), PC2 (2%), and PC3 (2%). These components form the PCA were based on the spectral variations between the FTIR spectra collected from each brain region. These variations were mainly located in the lipid, protein, ester structure, and the nucleic acids bio-contents. The dendrogram (Figure 5B) revealed that the spectral data were clustered into two major groups. The first two groups were sub-clustered under one main group, which was separated from the third group. The results indicated that the hippocampus structure was relatively close to the cortex, in the bio-chemical make-up, which indicated that the HC is a component part of the brain cortex, as evidenced by development studies—both are ectodermally-derived in the neural tube, during embryogenesis [25]. The results also indicated that the TH and CP are closely-related, due to the grey matter content. The IC, CE, and the CC are mainly related, due to the high content of the lipid axonal structure in the myelin sheaths of the white matter.

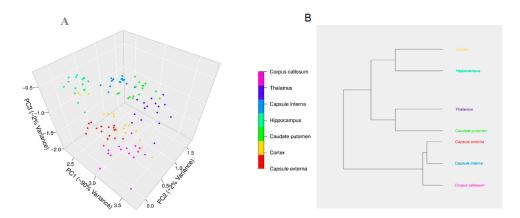


Figure 5. (**A**) PCA score plot for the seven brain regions. The three PCA values are given by PC1 (90%), PC2 (2%), and PC3 (2%). (**B**) Dendrogram representing the seven brain regions.

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3.1. Elemental analysis by Laser ablation (LA-ICP-MS)

LA-ICP-MS was used to assess the distributions of the important elements, in the rat brain sections, such as C, Fe, P, S, Cu, and Zn (Figure 6). From the rostral brain sections (Bregma 1.56), we detected high levels of C in the SSMC and in the CC (Figure 6). What was notable here was the high intensities of Ca, Mg, and Cu in the WM, and in particular in the CC. Two other elements, Fe and Zn, were evenly-distributed in the cortices and the basal ganglia, the former element was found to be enriched in the substantia nigra pars compacta—an area rich in dopamine neurons. Additionally, for the WM region, C, P, and S were also high in intensity—an indicator of high cell metabolism.

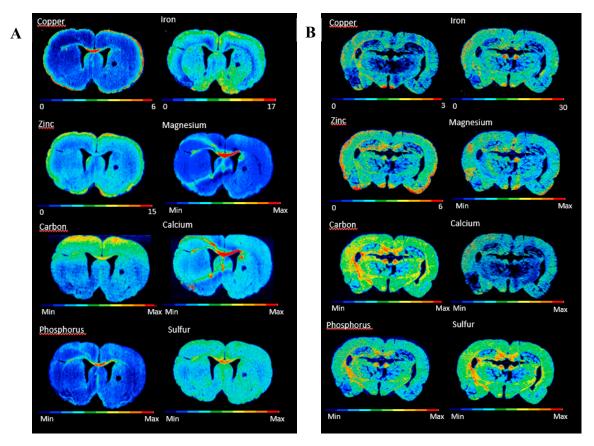


Figure 6. Distribution of the elements: (**A**) Cu, Fe, Zn, Mg, C, Ca, P, and S in the rostral, and (**B**) the caudal part of rat brain sections. The color bar represents a high-concentration, in red, and a low-concentration, in blue.

For the Caudal brain sections (Bregma -4.36), C was homogenously-distributed across the brain sections, with a high signal in the WM, the CC, and the Habenula. The divalent cations Ca, Mg, Cu, Fe, and Zn were fairly evenly-distributed, across the sections. As expected, Fe was also detected in the Habenula, the basal ganglia, and in the areas associated with the substantia nigra pars compacta. It should also be noted that the Mg, Cu, and Zn overlapped at specific regions and these hotspots were at brain sites closely-associated with active synaptic transmission. For the two inorganic elements, P and S, we detected a high intensity in the WM and the CC (Figure 6). Whereas, P and S are components of the nucleic acids, ATP, phospholipids, and proteins, which are all possible indicators of a high metabolism/turnover, in these regions.

4. Discussion

In this study the bio-chemical composition of seven different brain regions from healthy rats were investigated, using FTIR and LAICPMS spectroscopies. Application of these techniques to

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mammalian tissue, is not new, but in comparison with standard histopathological techniques, these approaches offer much more biochemical detail that may be important and pertinent to advance the understanding of pathologies. In combination with the PCA data, other brain regions that may prove to be susceptible to diseases, may be identified. The study revealed that FTIR spectroscopy allows for the investigation of molecular structures from biological samples, with minimal sample preparation; providing a unique information that cannot be gained by alternate methods. Importantly, the technique is a non-destructive, sensitive, and fast, bio-diagnostic tool [26].

FTIR results were in alignment with published work that the gray matter is made up of neural cell bodies, astro-glials, and capillaries, and contains very little lipid [27]. This data also revealed that the WM consists mostly of myelinated axons, with a high lipid-content, which showed strong absorption bands in the lipid region $(3000-2750 \text{ cm}^{-1})$ and the ester band (1740 cm^{-1}) .

In order to get a more detailed understanding of the lipid-structure configuration in the brain, the ratio between certain lipid spectral bands to the total lipid contents, were measured. This may be a good indicator and platform for detecting tissue alterations associated with pathologies. The lipid/protein ratios reveals information about the cell membrane function, since the changes in the lipid-protein content can affect the membrane symmetry and thickness, and leads to the confirmation of changes in the membrane receptors and ion-exchange channels [28]. For example, the ratio between the CH_2 spectral bands to the C–H stretching of the total lipid band revealed information about the lipids chain length. The ratio between the olefinic=CH band at 3000–3027 cm⁻¹ to the total lipid content at 2994–2800 cm⁻¹, provided information on the degree of unsaturation in the lipid biological system [29]. The changes of the band ratios of certain bio-chemical component can be used for characterization purposes, as well as to monitor tissue alterations.

The high CH₂/lipid and low CH₃/lipid ratios in the WM, CC, and CE indicated that these regions are characterized by long fatty-acid chains. These results are in a good agreement with a previous report [30], which stated that brain WM contains long-chained fatty acids. In another study, the authors showed that the sphingolipids from gray matter contained predominantly medium-chained fatty acids, while those from myelin or from white matter, contained predominantly long-chained fatty acids [31]. These regions also showed a high ratio of (C=O) carbonyl ester/lipid, indicating a high content of esterified-fatty acids. In contrast, the regions of SSMC, CP, TH, and the CA and DG of the hippocampus, exhibited low CH₂/lipid and high CH₃/lipid ratios, indicating short fatty-acid chains. These differences may reflect the functions of these regions. The low Olefinic=CH/lipid ratio in the WM, CC, and CE, in comparison to the GM regions of SSMC, CP, TH, and the CA and DG of the hippocampus, indicated that these WM, CC, and CE, were characterized by saturated and/or mono-saturated fatty acids, while the SSMC, CP, TH, and HP were predominantly characterized by polyunsaturated fatty acids. These results are in good agreement with the published data from Bazinet et al. [30]. The FTIR results revealed that in the HP, the CA and DG showed a lower lipid-content than in the AV region. The LA-ICP-MS chemical maps of carbon, phosphorus, and sulfur ions demonstrated a partial elevation of the concentration of those elements in the WM, which is a well-known difference between the WM and the GM, which is based on the abundant myelin in the WM, in comparison to the GM [32]. The MS chemical images related to the P, C, and S, provide a distinct contrast between the WM and the neighboring tissue types. Ca, Cu, and Fe play an important role in regulating brain excitability and neuronal plasticity, through their action in neurotransmission, or function as second messengers [33]. These metal ions are bound up in metal-protein complexes or metalloproteins, such as enzymes, transport, and storage proteins, in which metals ion are critical for proper protein folding, structural stability, or enzymatic catalysis [34]. Therefore, they have a central role in many biochemical pathways that are explained by the elevated distribution of these elements in specific brain regions [34]. For example, the higher level of the iron ions in the substantia nigra is due to the dopaminergic metabolism within these neurons. Thus, the LA-ICP-MS data complement the FTIR data by providing an extra insight into the total brain composition and a clue about the functions. Appl. Sci. 2018, 8, 2436

The chemometric analyses, such as principle component analysis and hierarchy dendrogram, showed that the seven regions of the brain were well-characterized and distinct, according to their spectral data. The score plot of the PCA delineated a clear separation between the brain regions, according to their lipid, ester, and protein content. Interpreting the dendrogram analyses we can argue that the hippocampus could be considered to be a distant part of the cerebral cortex. It was also concluded that all the regions that have a high-content of WM are clustered together, based on their lipid bio-chemical make-up. The eigenvalues graph showed that the control brain regions could be separated, based on the first three principal components, which accounted for more than 95% of the total variance in the spectral data. These variations were mainly located in the lipid, ester, and the protein spectral regions.

5. Conclusions

The FTIR spectroscopy is a good platform to reveal, examine, and investigate bio-chemical alterations associated with functional disorders in brain tissue. This makes the FTIR spectroscopy a valuable tool that can play an important role in the early detection of bio-chemical changes, at a sub-cellular level of neurological disorders, which would normally only be diagnosed when the pathology is at its most severe. The LA-ICP-MS elemental distribution maps provided information about what role and function these elements have in a particular brain region. This study showed that the FTIR and LA-ICP-MS techniques are powerful approaches to characterize the biochemical and elemental contents of brain tissue, providing complementary information to assess brain function. These techniques have been shown to be promising for tissue characterizations, as they offer unique opportunities for molecular and elemental assessments, for medical diagnoses.

Author Contributions: M.H.M.A. and F.R. performed all the FTIR and H&E measurements in this study. M.A., T.A., A.S., and K.A.-S. were involved in the data interpretation and in drafting the manuscript. V.N. and S.K. was involved in the elemental measurements and analysis studies. R.M. and E.U. were involved in processing the experimental data and performing the multivariate and statistical analysis. All authors discussed the results and commented on the manuscript and on the critical revisions of the manuscript.

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