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# Fourier-Transform Infrared Microspectroscopy (µFT-IR) Study on *Caput* and *Cauda* Mouse Spermatozoa<sup>†</sup>

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**Abstract:** Fourier-Transform Infrared micro-spectroscopy ( $\mu$ FT-IR) was used for an in vitro investigation on spermatozoa (SPZ) samples separately collected from *caput* and *cauda* of mouse epididymis. SPZ are characterized by deep biochemical changes during the transit along the epididymis and they can constitute ideal candidates for a  $\mu$ FT-IR investigation thanks to the ability of this technique in analyzing cells at a molecular level. Appreciable differences were reported in the infrared spectra from *caput* and *cauda* SPZ and biochemical changes in protein, nucleic acid, lipid, and carbohydrate content of cells were evidenced. The present investigation indicates that  $\mu$ FT-IR can constitute a valuable tool for monitoring in an easy and fast way the changes suffered by SPZ during the transit along the epididymis.

**Keywords:** *caput* and *cauda* spermatozoa; µFT-IR microspectroscopy; protein; nucleic acid; lipid; and carbohydrate components.

#### 1. Introduction

Fourier transform IR (FT-IR) spectroscopic technique is nowadays a very valuable tool for the nondestructive analysis of biological specimens. FT-IR results particularly useful in cytological and histological diagnosis also by using imaging procedures [1]. FT-IR is able to reveal bonds presenting a change of dipole moment during vibrational motion. These vibrational modes can be quantitatively characterized by IR spectroscopy, providing a very sensitive, label-free tool for studying molecular composition and dynamics without perturbing the sample. When FT-IR spectroscopy is performed on biological materials, the most important spectral regions are the fingerprint region (600–1450 cm<sup>-1</sup>) and the amide I and amide II (amide I/II) region (1500–1700 cm<sup>-1</sup>). The higher-wavenumber region (2800–3500 cm<sup>-1</sup>) is due to stretching vibrations such as S-H, C-H, N-H and O-H, whereas the lower-wavenumber regions typically correspond to bending and carbon skeleton fingerprint vibrations. These regions are able to give a complete biochemical fingerprint of the structure and function of investigated samples and have been adopted in a very large number of different samples of biological interest.

Recently, FT-IR spectroscopy has been adopted for investigating spermatozoa (SPZ) samples in order to principally take advantage of its non-invasivity. In particular, cyanide toxicity effects on rat sperms have been investigated at a molecular level by using FT-IR technique. Cyanide compounds are largely used in industries and are considered to be a ubiquitous pollutant in the environment. The use of FT-IR spectroscopy has also allowed an investigation on the secondary structure of protein

Proceedings **2019**, 2019 2 of 7

components. It is worth to note that a complementary vibrational technique, the Raman one, has been used for investigating oxidative DNA damage in human sperm [2].

In the present paper, FT-IR microspectroscopy (µFT-IR) has been adopted for characterizing SPZ separately collected from *caput* and *cauda* of the epididymis. Interestingly, SPZ are transcriptionally and translationally silent cells. During the epididymal transit, from *caput*-to-*cauda*, these cells undergo maturation via massive biochemical changes [3,4] that include: (i) decreases of cholesterol; (ii) qualitative composition of cytoplasmic proteins and RNA, (iii) protein degradation and phosphorylation; (iv) acquisition of GPI-anchored membrane proteins; (v) proteolytic remodelling of membrane proteins, (vi) fatty acid composition from saturated to polyunsaturated forms, (vii) DNA methylation, (viii) inter/intra protamine disulfide-bound formation. Because of their remarkable biochemical differences, *caput* and *cauda* SPZ can be considered ideal candidates for an FT-IR spectroscopy investigation.

In this report, we summarize the preliminary results of an investigation on mouse sperm samples from *caput* and *cauda* epididymis. To our knowledge, this is the first time that vibrational spectroscopies are adopted for investigating the changes occurring in SPZ during the epididymal transit

# 2. Materials and Methods

# 2.1. Collection of Sperm Samples from Caput and Cauda Epididymis

Adult male mice (CD1 background, from Charles River Laboratories, Lecco, Italy) were euthanized by asphyxia with CO2 and epididymis were dissected for sperm collection. Briefly, caput and cauda epididymis were separately immersed in PBS (pH 7.6) and slightly cut to endorse sperm extrusion from the epididymal duct, as previously reported [5]. Spermatozoa samples, separately collected from *caput* and *cauda* epididymis, were filtered throughout cheesecloth, washed by centrifugation (200xg for 5 min), in 0.9% NaCl solution and pellets were processed for FT-IR analysis.

All animal studies were carried out in accordance with the principles and procedures outlined in the National Institute of Health Guide for Care and Use of Laboratory Animals and approved by the Italian Ministry of Research and the Italian Ministry of Health.

### 2.2. FT-IR Spectral Analysis

For FT-IR spectral analysis SPZ pellets were resuspended in 0.9% NaCl solution and drops of few microliters were used for spectra acquisition. For this purpose, a Perkin Elmer Spectrum One FT-IR spectrometer equipped with a Perkin Elmer Multiscope system infrared microscope (Mercury Cadmium Telluride detector) was used. Spectral acquisitions were performed in specular-reflection mode with thin layers of samples (2  $\mu$ L) put on metallic IR-reflective surface. The background spectrum was collected from the metallic IR-reflective surface without sample. A sampling spot on the surface was selected through an objective (l0X optical or 15X infrared). All spectra were collected using 64 scans in the range from 4,000 to 600 cm<sup>-1</sup> with a 4 cm<sup>-1</sup> spectral resolution.

The spectra were preliminarily analyzed using the application routines provided by the software package ("Spectrum" User Guide, Perkin Elmer Inc. USA) controlling the whole data acquisition system. "Spectrum" (release 5.0.2, 2004) is the main Perkin Elmer software package for collecting, viewing and processing IR spectra.

## 3. Results and Discussion

The μFT-IR average spectra of mouse SPZ samples from *caput* and *cauda* epididymis (*caput* and *cauda* SPZ) were obtained in the 4000-600 cm<sup>-1</sup> range. The peak position and band assignments were done in agreement with ref. 6. In order to evidence the differences occurring between the two spectra, they were divided into four wavenumber ranges: 3600–3000 cm<sup>-1</sup> (Figure 1), 3050–2800 cm<sup>-1</sup> (Figure 2), 1800–1200 cm<sup>-1</sup> (Figure 3) and 1200–600 cm<sup>-1</sup> (Figure 4).

Figure 1 shows the average  $\mu$ FT-IR spectra of *caput* and *cauda* SPZ in the range of 3600–3000cm<sup>-1</sup>. In this range, there is the large contribution of amide A band (located at 3309 and 3290 for *caput* and

Proceedings **2019**, 2019 3 of 7

cauda SPZ, respectively) due to N-H and O-H stretching modes of proteins and intermolecular H bonding. It is worth to note that the shape of this band changes between caput and cauda samples. In particular, for caput SPZ the maximum is located at ~3400 cm<sup>-1</sup>, conversely, for cauda SPZ, the maximum is located 3290 cm<sup>-1</sup>. A very weak contribution ~3070 cm<sup>-1</sup> can be noticed only in the case of cauda samples and can be attributed to the N-H stretching of amide B. In order to analyze the changes in the lipid content of caput and cauda SPZ, the spectral region from 3050 to 2800 cm<sup>-1</sup> was considered (Figure 2). In this region, it is possible to evidence the contributions of CH<sub>3</sub> asymmetric stretching (located at 2957 and 2959 cm<sup>-1</sup> for caput and cauda samples, respectively), CH<sub>2</sub> asymmetric stretching (located at 2922 and 2924 cm<sup>-1</sup> for caput and cauda samples, respectively), CH<sub>3</sub> symmetric stretching (located at 2875 cm<sup>-1</sup> for both caput and cauda samples) and CH<sub>2</sub> symmetric stretching (located at 2851 and 2853 cm<sup>-1</sup> for caput and cauda samples, respectively). It's important to note that the two spectra are largely different in shape and relative intensity of the main peaks even though the position of the different CH2 and CH3 contribution do not change in a crucial way. In particular, the position of CH2 symmetric stretching doesn't show significant changes taking into account the spectral resolution of our equipment (4 cm<sup>-1</sup>). This can indicate that no important changes in membrane fluidity occur [7]. In this region of the spectra, it is also present a contribution at 3013 cm<sup>-1</sup> that is more evident for caput SPZ samples. This band is usually assigned to olefinic functional group in unsaturated lipids, mainly due to C-H stretching mode. This band can be used for further investigating lipid peroxidation in tissue [8].

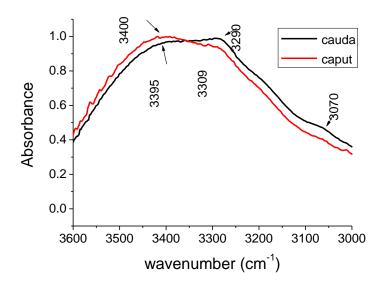
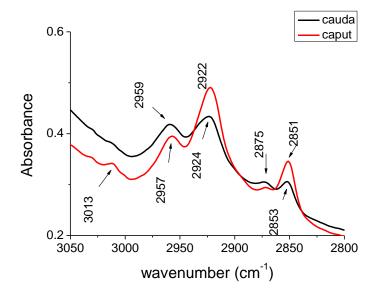


Figure 1. FT-IR spectra of caput and cauda sperm of WT mouse in 3600–3000 cm<sup>-1</sup> region.

Proceedings **2019**, 2019 4 of 7



**Figure 2.** FI-IR spectra of caput and cauda sperm of WT mouse in 3050–2800 cm<sup>-1</sup> region.

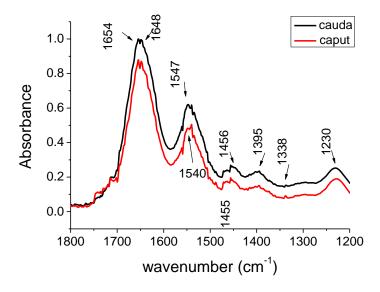


Figure 3. FI-IR spectra of caput and cauda sperm of WT mouse in 1800–1200 cm<sup>-1</sup> region.

Proceedings **2019**, 2019 5 of 7

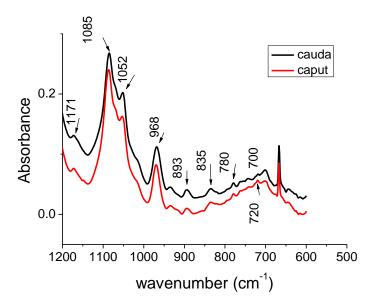


Figure 4. FI-IR spectra of caput and cauda sperm of WT mouse in 1200–600 cm<sup>-1</sup> region.

In Figure 3 the 1800-1200 cm<sup>-1</sup> region of μFT-IR spectra of *caput* and *cauda* SPZ is reported. In this region, the main contributions are due to protein content. In particular, the band due to the Amide I C=O stretching is positioned ~1650 cm<sup>-1</sup> for the two different samples, but differences can be noted in the band's shape. In Figure 3, the contributions of Amide II (at 1540 and 1547 cm<sup>-1</sup> for *caput* and *cauda* samples, respectively) and Amide III (at 1300 cm<sup>-1</sup> for both *caput* and *cauda* samples) are present. The small band positioned at 1455 and 1456 cm<sup>-1</sup> for *caput* and *cauda* samples, respectively can be due to CH<sub>3</sub> asymmetric bending and CH<sub>2</sub> scissoring mode that can be associated with lipid and protein contents. At 1395 cm<sup>-1</sup> a small feature can be noticed and it can be attributed to COO-symmetric stretching vibration mode related to fatty acids. The band located at 1230 cm<sup>-1</sup> is due to PO<sub>2</sub>- asymmetric stretching mode related to nucleic acids and phospholipids contributions.

In Figure 4 the region of short wavenumber (1200-600 cm<sup>-1</sup>) of μFT-IR spectra of *caput* and *cauda* SPZ is shown. This region is mainly related to DNA contributions. The most important ones are located at 1085 (PO<sub>2</sub><sup>-</sup> symmetric stretching) and 1052 cm<sup>-1</sup> (C–O stretching). In this region too, differences can be noted in the band shape. Also the other structures positioned at 968 cm<sup>-1</sup> (C–C stretching), 893 cm<sup>-1</sup> (deoxyribose ring), 835 cm<sup>-1</sup> (S- type sugar) are attributed to DNA. A complete list of the peak position and band assignments is reported in Table 1.

**Table 1.** Mode assignments for the FT-IR frequency values in caput and cauda sperm samples, abbreviation: as = asymmetric, v = vibration, v = vibration,

Peak caput cm <sup>-1</sup>	Peak cauda cm <sup>-1</sup>	Peak Assignments	Biochemical Molecules
3400	3395	Hydrogen bonded O-H v	protein water
		N-H v	
3309	3290	Amide A (N-H v)	protein
3013	3013	olefinic = $C-H v$	unsaturated lipid
2957	2959	CH <sub>3</sub> as v	lipid
2922	2924	CH <sub>2</sub> as v	lipid
2875	2875	CH <sub>3</sub> s v	lipid

Proceedings **2019**, 2019 6 of 7

2851	2853	CH <sub>2</sub> s v	lipid
1654	1654	Amide I α-helix	protein
1648	1648	Amide I	protein
1540	1547	Amide II	protein
1455	1456	CH <sub>3</sub> as b	lipid and protein
		CH <sub>2</sub> scis	
1395	1395	COO-s v	fatty acid
1300	1300	Amide III	protein
1230	1230	PO <sub>2</sub> - as v	nucleic acids and phospholipids
1171	1171	CO-O-C v	glycogen
1085	1085	PO <sub>2</sub> - s v	nucleic acids
1052	1052	C-O v	saccharides and DNA deoxyribose
968	968	C-C v	DNA backbone
893	893	Deoxyribose ring	DNA
835	835	DNA S-type sugar	DNA

#### 4. Conclusions

The results here reported indicate that µFT-IR spectroscopy can constitute a useful tool for investigated changes occurring in SPZ from *caput* and *cauda* epididymis of mouses in a rapid and easy way. Biochemical changes in protein, nucleic acid, lipid, and carbohydrate content of cell have been evidenced. Further work is in progress in order to quantitatively assess the observed differences.

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**Compliance with ethical standards:** All applicable international, national, and/or institutional guidelines for the care and use of animals were followed.

**Conflicts of Interest:** The authors declare that there is no conflict of interest.

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Proceedings **2019**, 2019 7 of 7

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