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Infrared spectroscopy and microscopy for biology and biochemistry

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IR fingerprints relevant to the biological applications of IR Microspectroscopy (IRMS)



Biology is the branch of natural science that studies life and living organisms.

The building blocks of life are the cells.

Cell

Tissue

Typical samples studied by means of IR microspectroscopy Organ

Living organism

The study of the chemical processes in living organisms is the subject of Biochemistry. It deals with the <u>structures</u> and <u>functions</u> of cellular components such as proteins, carbohydrates, lipids, nucleic acids and other biomolecules.

The lecture will illustrate recent achievements of SR InfraRed MicroSpectroscopy (IRMS) in studies of biochemical features of complex biological systems (tissues and cells)

Peptide



Adapted from: L. M. Miller, G.D. Smith and G. L. Carr, Journal of Biological Physics, 29 (2-3), 219-230, 2003



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Band intensity, position, width and shape (band components) are sensitive to subtle biochemical changes of bio-specimens.



Biological applications of IR microspectroscopy (IRMS)

Infrared spectroscopy and microspectroscopy of plant, animal or human cells, tissues and body fluids as well as of microbial species can be exploited for :

1- Characterizing and differentiating microbial cells and strains: rapid method for identifying micro-organisms responsible for infections.

2- Differentiation of healthy and diseased tissues and/or cells: new tool in medical diagnostic.

3-Charatherizing cell growth dependent phenomena and cell interaction with different agents such as drugs, pollutants, poisons, chemotactic agents: a complementary tool to conventional biochemical assays.

4- Analyzing blood and urine: fast and new clinical chemistry tool.

5- Great variety of other less conventional applications.

How can all this information be deduced from a series of infrared spectra?



Why fixed samples?

• *In-vivo* MIR imaging of internal organs (and cells) at the actual stage of the technique is still not possible

• Fiber-optic based ATR methods in the mid-infrared have been used for studying epithelial surfaces and surface skin contaminants *in-vivo* [Refs]. All the other tissues have to be analyzed *ex-vivo*.

• Conventional tissue slice thickness ranges from 5 to 20 microns, depending on the tissue type, in order to avoid signal saturation, that will make impossible the interpretation of IR data [Lambert-Beer Law].

•Single cell thickness is usually too small for inducing IR signal saturation.

The Lambert-Beer Law $A = -\log_{10} (I_1/I_0) = \varepsilon/c$ $\int_{I_0}^{\infty} c, \alpha = I_1$ $\int_{I_0}^{\infty} c, \alpha = I_1$ $\int_{I_0}^{\infty} c = mol \cdot L^{-1}$ $\int_{I_0}^{\infty} c = cm$

"Fiber-optic Probes for Mid-infrared Spectrometry", Peter J. Melling and Mary Thomson in *Handbook of Vibrational Spectroscopy*, John M. Chalmers and Peter R. Griffiths (Editors), John Wiley & Sons Ltd, Chichester, 2002
 L. Brancoleon, M.P. Bamberg and N. Kollias, *Appl. Spectrosc.*, 54, 1175 (2000).

Why fixed samples?

• *In-vitro* experiments under physiological conditions are limited by the strong water absorption bands in the Mid-IR regime.



In particular, the bending band of water centered at 1650cm⁻¹, completely hides the Amide I band of proteins, one of the most important for biologic application of IRMS.

New strategies are under development for allowing to perform *in-vitro* cellular experiments -> See "Mechanobiology of leucocytes"

Proper selection of fixation protocol

The aim of fixation is to preserve the structural and biochemical constituents of cells in as close to in vivo conditions as possible

•*Air-drying* can cause collapse of internal cellular structures and activation of cell autolysis (dramatic variation of osmotic pressure within the cells).

•*Flash-freezing* followed by cell lyophilisation (freeze-drying) can not be applied to most common FTIR materials, such as CaF_2 or BaF_2 , since too brittle and with poor thermal contact.

• Alcohol fixation results in water extraction and decrease of the cellular volume. Water is displaced from proteinaceous material, resulting in protein denaturation and organelle disruption. Alcohol extracts lipids from cells but has little effect on carbohydrates.

• *Formalin fixative* has bands potentially overlapping with cellular constituents bands (the most intense peak occurs at 1000 cm⁻¹); however it preserves most lipids and has little impact on carbohydrates. Formalin also appears to preserve protein secondary structure.

No definite solution since direct comparison with living cell IR spectra under physiological conditions has not been performed yet.

Which substrates?

Sampling technique	Material	Vis- Transparent	MIR- Transparent	MIR- Reflective	Biocompati bility	Other
Transflection	MirrIR-slides	No	No	Totally	Yes	Cheap
Both	Si, Ge	No	Partially	Partially	Yes	Cheap
Transmission	Diamond	Yes	Partially	No	Yes	Expensive +
	BaF ₂	Yes	Totally	No	No	Expensive
	CaF ₂	Yes	Partially	No	Possibly	Expensive
	ZnSe	Slightly	Partially	No	No	Expensive
	Si ₃ N ₄	Slightly	Partially	No	Yes	Fragile
	TEM grids	Yes	Totally	No	Yes	Fragile



The optimal IR substrate

- Biocompatible, for performing cell culture on it
- IR transparent/reflective in the MIR
- Vis transparent/reflective, to easily match Vis and IR data
- Cheap and/or recyclable
- Exploitable also for other investigation techniques



An infrared bio-experiment step by step Data collection_2

Schwarzschild objective





Acquisition parameters • Knife edge apertures, in order to match the desired experimental spatial resolution avoiding diffraction artifacts

• Number of scans per spectral point in order to enhance S/N ratio without dramatically increasing the acquisition time



Pre-processing of data



Pre-processing of data



Smoothed Spectrum Savitzky-Golay method, K+1 smoothing points

The method essentially performs a local polynomial regression on a series of values (k+1 points, equally spaced in the series) to determine the smoothed value for each point. This method is also provided for calculating the first up to the fifth derivatives of spectra.

First or second derivatives of spectra are often used for data analysis, in order to minimize baseline variations and maximize spectral resolution.



Pre-processing of data

Water vapor and carbon dioxide contributions to bio-specimen spectra as well as the spectral noise can be greatly reduced:

- Purging the interferometer with N2/dry air or operating in vacuum
- Purging the microscope stage environment with N2/dry air
- With a good conditioning of the laboratory environment



Too many people around the microscope!

- Enhancing the S/N spectral ratio
 - Increasing the number of scans
 - Increasing the signal -> Brighter Sources -> Synchrotron Radiation
 - Reducing any possible source of noise: vibrations, electronic noise,

Spectra comparison

Spectral analysis "by visual inspection"

For small data sets (few spectra), spectra can be compared "visually" in order to highlight spectral similarities and/or differences affecting:

• Band position (band shifts), width (band broadening) and shape (band components)

• Ratios of peak areas (different proportion of most fundamental tissue-cell constituents).

The reliability of biological conclusions drawn out from an experiment relies on the measurements of a statistically relevant number of samples.

Statistical analysis

Univariate and multivariate statistical analysis methods allow to compare a huge amount of spectra simultaneously, classifying them on the base of spectral similarities, affinities.

Univariate Methods: Average, standard deviation, regression techniques (PLS,....) Multivariate Methods: Cluster Analysis, Principal Component analysis (PCA),

1 - Cluster Analysis

1. Data pre-processing

Spectra are **normalized** in order to process an homogenous data set. Commonly employed normalization methods (alone or in combination) are: 1. Vector Normalization

 $a_{m} = \frac{\sum_{k} a(k)}{N} \text{ average value; } a'_{k} = a(k) - a_{m} \text{ subctracted spectrum; } a''_{k} = \frac{a'_{k}}{\sqrt{\sum_{k} (a'(k))^{2}}} \text{ normalized spectrum}$

2. (Extended) Multiplicative Scattering Correction (E)MSC



Single cell spectra often present slow oscillations of the baseline. The origin is that the interaction of electromagnetic radiations with dielectric spheres of dimensions close to that of the wavelength of light induces a strong scattering that can be modeled with the **Mie Scattering theory**. EMSC correct for this effects modeling the scattering object as a non-absorbing dielectric sphere. New algorithms are under development that consider the absorbing characteristics of the scattering object.

3. Derivative (1st or 2nd)

An infrared bio-experiment step by step Data Analysis_2 1- Cluster Analysis



2. Spectral distance calculation

Distance between spectra a and b can be calculated with many algorithms.

Euclidean spectral distance between a and b spectra is calculated over the all sampled k points.





There are many methods available to calculate spectral distances between a newly-created cluster and all the other spectra or clusters.



4. Spectra clustering





SR IRMS and Prion research

Protein structural information from Amide I band





Amide I band - 1700-1600 cm ⁻¹					
	Antiparallel ß-				
1692-1672	sheet/Aggregated strands				
1670-1660	3 ₁₀ - Helix				
1660-1648	a-helix				
1648-1640	Random coil				
1640-1625	β-sheet				
1628-1610	Aggregated strands				

... exploited for studying Transmissible spongiform encephalopathies (TSEs, also known as prion diseases)

M. Beekes et al., Analytical applications of Fourirer transform-infrared (FT-IR) spectroscopy in microbiology and Prion research, Veterinary microbiology (2007), 123:305-319



SR IRMS and Prion research



Cells were grown on silicon nitride 100nm membranes (GT1 cells do not preserve vitality when cultured on CaF_2 windows) and fixed in formalin 4%. An IR measurement chamber was then realized gluing a second window onto the first to guarantee the **biological safety**. Once concluded the experiment, samples are disposed (Si₃N₄ membranes are quite cheap).

- Data acquisition (TR mode)
- Individual whole-cell analysis
- Intra-cellular analysis
- Conventional biochemical assays

IR



Prion Laboratory , Neurobiology Sector, SISSA Prof. G. Legname, A. Didonna

Individual whole-cell analysis_1



Acquisition parameters

Spectra have been averaging 1024 scans. Knife edge apertures have been set at 30X30 μ m in order to match an entire cell

Data preprocessing

• Atmospheric compensation

• Cut :3800-1150 - Spectra have been cut at 1150 cm⁻¹ since (i) absorption band of silicon nitride is hardly compensate below 1200 cm⁻¹ (ii) the more intense band of formalin fixative is falling between 1100-1000 cm⁻¹. With this choice, both problems are avoided.

Is IR spectroscopy a quantitative analytical technique?

Yes, it is
$$\mathcal{A} = -\log_{10}\left(\mathcal{I}_{1}/\mathcal{I}_{0}\right) = \varepsilon/c$$

BUT the cell thickness is not homogenous and it differs from cell to cell, even within the same cell type

Nucleus Perinuclear region

Only relative variations in concentrations of most fundamental cellular macromolecules can be deduced (such as Protein to Lipid ratio) by rationing associated IR band areas (or heights) unless some additional information are available!

Individual whole-cell analysis_1

Semi-quantitative analysis

 GT1 and Sc-GT1 cells are indiscernible from a morphological point of view as known from the scientific literature and AFM (Atomic Force Microscopy) investigations.
 A conventional biochemical assay

(Bicichoninic acid) was performed in order to verify the protein cellular content variation upon infection, giving results comparable with IR analysis.





Individual whole-cell analysis_2 **Classification of spectra** GT1 1 Heterogeneity 0.8 2.5 0.5 2.0 1.0 1.5 0 0.6 0.4 Mr 0.2 GT1 Cluster Analysis 3500 3000 2500 2000 1500 Wavenumber / cm⁻¹ <u>אלה והו ואה הוא הדודה אוו חוז</u>ה Sc GT1 1 ScGT1 0.8 0.6 0.4 M 0.2 First derivative of spectra 3500 3000 2500 2000 1500 Wavenumber / cm⁻¹ Euclidean distances & Ward's algorithm



Intracellular analysis

Acquisition parameters

512 scans; Knife edge apertures set at 6X6

Amide I band ~ 6 μ m; Phospholipid band ~ 5.7 μ m; Lipid band 3.4 μ m

Data preprocessing



Adaptive immunity modulated by hBD2

Host Defense Peptides (HDPs): short introduction

 \checkmark Evolutionarily ancient component of INNATE IMMUNITY in multicellular organisms

✓ DIRECT KILLING of invading pathogens and/or MODULATION of IMMUNE and healing responses of the host.

✓ Huge DIVERSITY in sequence and structure

Human

defensin

hBD2

✓ Two major families: Cathelicidins and Defensins

TEMPLATE for development of antimicrobial compounds and/or immunomodulatory agents Human cathelicidin LL37

Anti-infective Laboratory Life Sciences Dept., University of Trieste Prof. A. Tossi, Prof. R. Gennaro and Prof. Sabrina Pacor PhD thesis of Francesca Morgera





Flow cytometry & single cell µ-FTIR



R1=completely differentiated, responsive iDC R2= non completely differentiated, unresponsive cells

Single cell µ-FTIR & Fluorescence microscopy





Plasma membrane Ch depletion affects hBD2 internalization

Living cell µ-FTIR measurements Mechanobiology of leucocytes





Fluidic cells with a carefully and reproducible path length (<9μm) for a precise and reliable water subtraction >Amide I band disclosure
Fluidic devices shaped with narrow channels (few microns wide and thick) in order to mimic microcapillaries and/or epithelium interstices

Microfabrication is needed

Prof. Sabrina Pacor

PhD activity of Giovanni Birarda

Sciences

Dept



The microfabrication issues

Calcium fluoride: new fabrication strategies

- + Vis-IR transparent
- + Lower water solubility (\rightarrow citotoxicity) than BaF₂
- High sensitivity to thermal shocks
- Low reactivity towards common reagents for wet etching



The spectroscopic approach



U937 monocytic (average diameter 8-10 microns) D- Gently deformed cells (★) [9 µm device] D Strongly deformed cells (★) [5 µm device] D+ Extremely deformed (▲) [3 µm device]





The synergic match between μ fabrication and μ -FTIR could extend the frontiers of FTIR microspectroscopy to almost unexplored fields such as mechanobiology.



Summary and Conclusions

- SR IR microspectroscopy: highly sensitive non-radiation damaging technique for the characterization of subtle biochemical changes in biological matter with sub-cellular spatial resolution.
- To unveil biological divergences associated to spectral differences, a good experimental design has to be done: sample preparation, data collection and data analysis.
- The present limitations for studying living systems in physiological environment can be overcome developing microfabrication approaches suitable for IR-transparent materials.
- The complexity of biological word can be addressed only by a multidisciplinary/multi-technique approach

Selected literature

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Biomedical Vibrational Spectroscopy Peter Lasch PhD (Author), Janina Kneipp (Author) Wiley-Interscience, 2008