Fourier-Transform Infrared Spectroscopy of Epiretinal Membranes and Internal Limiting Membranes after Pars Plana Vitrectomy with Membrane Peeling

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Keywords
Fourier-transform infrared spectroscopy · Epiretinal membrane · Internal limiting membrane · Vitrectomy with membrane peeling

Abstract

Introduction: Fourier-transform infrared imaging (FTIRI) enables examination of protein secondary structure in the analyzed tissues. The aim of our study was to examine the distribution of secondary structures in epiretinal membranes (ERMs) and internal limiting membranes (ILMs), and to explore possible associations to other diagnostic variables.

Methods: This prospective pilot study included patients scheduled for pars plana vitrectomy with membrane peeling. ERMs and ILMs were harvested during surgery and placed on a BaF\textsubscript{2} window for postsurgical FTIRI analysis. Infrared hyperspectral images were subjected to second and fourth derivative analysis to obtain information of the protein secondary structures present in the tissues.

Results: Samples of 43 patients were analyzed, with the triple helical domain showing the highest prevalence in the examined tissues. The other secondary structures (beta-sheet, random coil, and beta-turn) showed a heterogeneous distribution in the examined samples, without specific associations to indication of surgery, comorbidities, outcomes from optical coherence tomography, and intraoperative findings.

Conclusions: FTIRI enables analysis of the spatial distribution of protein secondary structures in the examined tissues; thus, it is a useful analytical technique for the analysis of ERMs and ILMs.
The aim of our study was to use FTIRI to examine the distribution of specific secondary structures, in ERMs and ILM, and to explore possible associations with other diagnostic outcomes.

**Materials and Methods**

This prospective study included patients scheduled for pars plana vitrectomy with membrane peeling between August 2017 and December 2018 at the ophthalmic department of the Hanusch Hospital in Vienna, Austria. The inclusion criteria were age above 18 years, presence of an ERM with indication for surgery (defined as significant loss of vision and/or metamorphopsia due to the ERM), full thickness macular hole, and lamellar macular hole. All patients, samples could not be harvested during surgery, 43 (86%) patients could be analyzed by FTIRI, while in 4 patients, samples were not evaluable. The mean age of patients was 72.5 (SD 7.8) years. An overview about the number and exact position of the underlying peaks that can be assigned to specific secondary structures. In instances where it was obvious that the calculated second derivative peaks were not single but still composite ones (presence of discernible shoulders), 4th derivative spectra were calculated as well.

Continuous data are described using mean and standard deviation, and for categorical data, absolute frequencies and percentages are presented. To examine associations of FTIRI outcomes to diagnosis and general factors, χ² test and Fisher’s exact test were performed. In case of a significant finding, we planned to apply Benjamini and Hochberg correction for multiple testing in order to control for the false discovery rate; however, no significant relationship was found.

**Results**

From 50 patients enrolled in the study, samples from 43 (86%) patients could be analyzed by FTIRI, while in 4 patients, samples could not be harvested during surgery, and in 3 cases, samples were not evaluable. The mean age of patients was 72.5 (SD 7.8) years. An overview about indications for surgery is provided in Table 1.

**Table 1. Distribution of secondary structures of proteins in ERM and ILM**

| (n)          | Beta-sheet domain, % (n) | Random coil domain, % (n) | Alpha-helical domain, % (n) | Triple helical domain, % (n) | Beta-turn domain, % (n) |
|--------------|--------------------------|---------------------------|-----------------------------|-----------------------------|------------------------|
| iERM (17)    | 47 (8)                   | 18 (3)                    | 24 (4)                      | 88 (15)                     | 71 (12)                |
| dERM (9)     | 67 (6)                   | 11 (1)                    | 22 (2)                      | 100 (9)                     | 78 (7)                 |
| FTMH (7)     | 29 (2)                   | 14 (1)                    | 14 (1)                      | 86 (6)                      | 43 (3)                 |
| Other ILM samples (4) | 50 (2)            | 0 (0)                     | 0 (0)                       | 100 (4)                     | 50 (2)                 |
| Atrophic LMH (3) | 67 (2)            | 33 (1)                    | 33 (1)                      | 100 (3)                     | 100 (3)                |
| ERM foveoschisis (4) | 50 (2)            | 0 (0)                     | 25 (1)                      | 75 (3)                      | 50 (2)                 |
| Pseudomacular hole (2) | 0 (0)            | 0 (0)                     | 0 (0)                       | 100 (2)                     | 0 (0)                  |

ERM, epiretinal membrane; iERM, idiopathic epiretinal membrane; dERM, diabetic epiretinal membrane; FTMH, full thickness macular hole; LMH, lamellar macular hole; ILM, internal limiting membrane.
from 75 to 100%), whereas other protein conformations did not reach such high levels of presence in the samples, with the exception of beta-turn domain in samples from atrophic lamellar macular holes (Table 1; Fig. 1).

Statistical analysis of results showed no significant associations between conformations found in FTIRI and indications for surgery (idiopathic ERM, diabetic ERM, full-thickness macular hole, atrophic lamellar macular hole, ERM foveoschisis, and pseudomacular hole), comorbidities (such as arterial hypertension, hypercholesterinemia, and diabetes mellitus), retinal properties (such as adherence of membranes [fully or partly adherent] and intraretinal cystoid changes, both diagnosed with optical coherence tomography), and intrasurgical findings (such as good or poor staining of membranes with chromovitrectomy dye, and whether only ERM was peeled, ERM and ILM were peeled separately, or ERM and ILM were peeled en bloc) (Table 2).

Fig. 1. FTIRI analysis of an ERM sample of a patient with an idiopathic ERM: native tissue sample (upper panel), organic matrix distribution based on FTIRI (middle panel), and second derivative analysis with peaks at wavenumber 1661, indicative for triple helix domain (lower panel). FTIRI, Fourier-transform infrared imaging; ERM, epiretinal membrane.
Discussion

In this prospective pilot study, we observed that all specific protein conformations, indicated by FTIRI, could be found in ERM and ILM with triple helical domain showing the highest prevalence. Nevertheless, the distribution of specific ones was heterogeneous, with no typical distribution patterns among different types of ERMs (idiopathic ERM, diabetic ERM, and different types of lamellar macular holes) and ILM. The results of our study, therefore, indicate a heterogeneous composition of proteins in these membranes, even in case of the same indications for surgery. This finding is in accordance with the fact that results concerning the presence of types of collagen in epiretinal tissue examined with immunohistochemical analysis were reported with differences between authors [3–6].

In general, collagen types I, II, III, IV, and VI were found in ERMs [3–6]. Kritzenberger et al. [3] found a high amount of collagen type VI and a relative absence of collagen types I and II in cellophane maculopathy compared to preretinal fibrosis. Bu et al. [4, 5] found collagen types I, II, IV, and VI in ERM samples and collagen types I, III, and V in membrane samples from macular hole surgery, whereas Compera et al. [6] found collagen types I and III in samples from lamellar macular holes. Apart from collagen, also laminin, fibronectin, and α-SMA were found in ERMs [3, 4, 6]. Collagen types I, II, III, and V are fibril-forming collagens, with characteristic patterns of bands in electron microscopy, whereas collagen type IV is a sheet-forming collagen, typical for all basal laminas, and collagen type VI is a fibril-associated collagen, associated with collagen type I [7].

All of the abovementioned types of collagen show a triple helical configuration in the central part (for collagen types IV and VI, the central triple helices are interrupted, enabling more lateral flexibility than the fibril-forming types of collagen I, II, III, and V) with telopeptides at both ends (the C and N terminals). Triple helical domain in FTIRI is, therefore, indicative for the presence of collagen molecules. We have no explanation why the triple helical domain could not be shown in all samples. Nevertheless, ERM and ILM also consist of other proteins with possible influence on FTIRI outcomes.

As the FTIRI results resemble information of spatial characteristics of proteins and their neighborhoods in the examined tissue, other proteins in ERM and ILM either from telopeptides or the extracellular matrix have the potential to influence the results. Up to now the presence of specific conformations of proteins in epiretinal tissues, such as the beta-sheet domain, random coil domain, alpha-helical domain, and beta-turn domain, has not been examined enough to give conclusive answers with respect to that question. Due to the fact that this study was planned as a pilot study, investigating FTIRI for epiretinal tissues in general, future research will be needed to improve knowledge on protein origins and their impact on FTIRI outcomes.

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| Table 2. Associations between diagnosis and general factors to secondary structures of proteins |
|-------------------------------------------------|-----------------|-----------------|-----------------|-----------------|-----------------|
|                                                  | Beta-sheet domain, p | Random coil domain, p | Alpha-helical domain, p | Triple helical domain, p | Beta-turn domain, p |
| iERM                                             | 0.961             | 0.682            | 1.02             | 1.02             | 0.531           |
| dERM                                             | 0.312             | 1.02             | 1.02             | 0.562            | 0.52            |
| FTMH                                             | 0.422             | 1.02             | 1.02             | 0.522            | 0.42            |
| Atrophic LMH                                      | 0.622             | 0.352            | 0.492            | 1.02             | 0.292           |
| ERM foveoschisis                                  | 1.02             | 1.02             | 1.02             | 0.312            | 0.623           |
| Pseudomacular hole                               | 0.492             | 1.02             | 1.02             | 1.02             | 0.133           |
| Arterial hypertension                            | 0.812             | 0.382            | 0.472            | 1.02             | 0.392           |
| Hypercholesterinemia                             | 0.342             | 1.02             | 1.02             | 1.02             | 0.282           |
| Diabetes mellitus                                | 0.182             | 1.02             | 1.02             | 1.02             | 0.492           |
| Adherence of membranes                           | 0.783             | 0.672            | 0.462            | 1.02             | 0.942           |
| Intraretinal cystoid changes                     | 0.861             | 1.02             | 0.682            | 1.02             | 1.02            |
| Staining properties                              | 0.881             | 0.622            | 1.02             | 0.562            | 0.722           |
| ERM, ILM, ERM + ILM en bloc                      | 0.512             | 1.02             | 0.672            | 0.822            | 0.512           |

ERM, epiretinal membrane; iERM, idiopathic epiretinal membrane; dERM, diabetic epiretinal membrane; FTMH, full thickness macular hole; LMH, lamellar macular hole; ILM, internal limiting membrane. 1χ2 test. 2Fisher’s exact test.
tion with aldehydes, as this procedure has potential for creation of artifacts in FTIRI results.

We did not find any significant associations of specific conformations, identified by FTIRI, to the indications for surgery, comorbidities of patients, retinal properties, and intrasurgical findings most likely due to low patient numbers, as this study was designed as an exploratory one (Table 2). FTIRI results, however, provide a completely new mode of examination of excised tissues in ophthalmology, and further research is needed to better understand the clinical impact of the spectroscopic findings.

A limitation of the study is the fact that the study was planned and performed as a pilot study with the former classification of lamellar macular holes, and, therefore, numbers of samples from patients with atrophic lamellar macular holes, ERM foveoschisis, and pseudomacular holes are low. Nevertheless, 100% presence of beta-turn domain and triple helical domain in atrophic lamellar holes, and the fact that in pseudomacular holes only triple helical domain could be found are novel results with the need to be validated with a higher number of samples. In conclusion, FTIRI is a new method of analysis of ERMs and ILM and their composition of proteins, with spatial information including specific conformation of molecules.

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