

# Measurement of permittivity of biological tissues from 300 MHz to 3 GHz using an open-ended coaxial line

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## **ABSTRACT / RESUMEN**

A comparative numerical study of three probe types with the same radial dimensions, three probe models, and five reference materials is presented for six biological tissues: liver, muscle, kidney, heart, blood and fat. The Probe-Model-Reference (PMR) combinations that lead to the most accurate results of the permittivity at frequencies from 300 MHz to 3 GHz are investigated for a coaxial probe without flange, a coaxial probe with flange, and a coaxial probe open into a propagation circular wave guide. The probe models considered are capacitive model, antenna model, and virtual line model. The reference materials are distilled water, 0.02 M NaCl(aq) solution, ethanol, methanol and 0.5 M NaCl(aq) solution. The results corroborates that when measuring the RF permittivity of biological materials using an open-ended coaxial line the proper selection of the probe type, probe model and reference material is crucial if good accuracy of measurements is expected. The presented methodology could be useful not only for the design of experiments, but also for the development of improved coaxial probes.

Key words: coaxial probe, complex permittivity measurement, vector reflection coefficient measurement

*Para seis tejidos biológicos –hígado, músculo, riñón, corazón, sangre y grasa– se presenta un estudio numérico comparativo de tres tipos de sonda con las mismas dimensiones radiales, tres modelos de sonda y cinco materiales de referencia. Se investigan las combinaciones Sonda-Modelo-Referencia (SMR) que conducen a los resultados más precisos de la permitividad a frecuencias desde 300 MHz hasta 3 GHz para una sonda coaxial sin pestaña, una sonda coaxial con pestaña, y una sonda coaxial abierta hacia una guía de onda circular. Los modelos de sonda considerados son el capacitivo, el de antena y el de la línea virtual. Los materiales de referencia son agua destilada, solución de NaCl(ac) 0.02 M, etanol, metanol y solución de NaCl(ac) 0.5 M. Los resultados corroboran que cuando se mide la permitividad de RF de materiales biológicos usando una línea coaxial abierta en un extremo la selección apropiada del tipo de sonda, del modelo de sonda y del material de referencia es crucial si se espera una buena exactitud de las mediciones. La metodología presentada pudiera ser útil no solamente para el diseño de experimentos, sino también para el desarrollo de sondas coaxiales mejoradas.*

*Palabras claves: sonda coaxial, medición de la permitividad compleja, medición del coeficiente de reflexión vectorial*

*Medición de la permitividad de los tejidos biológicos desde 300 MHz hasta 3 GHz usando una línea coaxial abierta en un extremo*

## 1. - INTRODUCTION

Measurement of dielectric properties of biological tissues at radio frequencies has been the subject of many researches in the last four decades [1-26]. For frequencies below 20 GHz the commonly used technique to obtain the complex dielectric permittivity ( $\epsilon$ ) consists of measuring the complex reflection coefficient ( $S_{11}$ ) using a vector network analyzer. Afterwards,  $\epsilon$  is obtained through a model that gives the complex admittance of the probe/material interface as a function of the dielectric permittivity of the material under test (MUT), which is considered nonmagnetic, isotropic, homogeneous and semi-infinite [7]. Since the applied fields are weak, it is assumed that they elicit linear responses [27].

Various admittance models of an open-ended coaxial sensor have been proposed to determine complex permittivity of materials [2, 3, 28-30]. These models include the so-called capacitive [2, 3], antenna [28] and virtual line [29] models. A common procedure is to assume that their parameters do not depend upon  $\epsilon$  or frequency, and determine them by measuring the admittances of substances with known permittivities (reference materials) and then use the calibrated probe to measure substances with unknown  $\epsilon$ . However, discrepancies of 10 % or more may occur between the measured and the expected values [31, 32]. Fortunately, this can be attenuated if the reference material used during the calibration procedure had very similar properties to the specimen [32-34]. However it is unlikely that one would obtain such references for all possible specimens at all frequencies.

There is a large family of coaxial probe types designed for dielectric measurements. These include the coaxial probe without flange, the coaxial probe with flange, and the coaxial probe open into a propagation circular wave guide, among others [1, 4, 35]. Commonly, the appropriate probe is selected taking into account only the suitability for in vivo measurements, ease for the control of the sample temperature, sample size, compatibility with liquid, semiliquid and solid samples, as well as range of operating frequencies [1, 2]. However, when making this selection the biological material, the probe admittance model and the reference material are not considered.

Although the limitations of the capacitive, the antenna and the virtual line models have been recognized [31, 32-34, 36, 37], no comparative studies are available in the open literature to assess the precision of these models, when measurements are made on biological tissues, and the influence of both, the reference material and the open-ended coaxial probe physical characteristics, are considered. In this work a numerical such comparative study is made, and the Probe-Model-Reference (PMR) combinations that lead to the most accurate results of the permittivity of six biological tissues (liver, muscle, kidney, heart, blood and fat) at frequencies from 300 MHz to 3 GHz are investigated for three coaxial probe types (without flange, with flange, and open into a propagation circular wave guide) with the same radial dimensions, three probe models (capacitive, antenna, virtual line), and five reference materials (distilled water, 0.02 M NaCl(aq) solution, ethanol, methanol and 0.5 M NaCl(aq) solution).

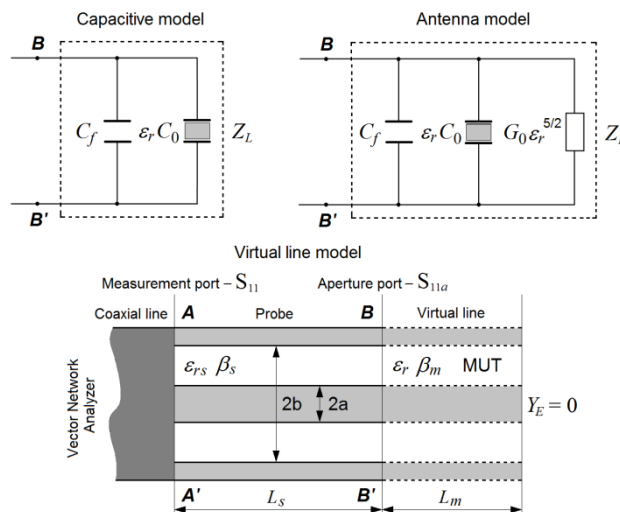


Figure 1

Schema of the three models used in this study.

## 2. - MODELING OF THE OPEN-ENDED COAXIAL PROBE

The schema of the capacitive, the antenna and the virtual line models is shown in Figure 1. The following sections present a brief description of these models:

## 2.1. - CAPACITIVE MODEL

The equivalent circuit for this model consists of two capacitances connected in parallel, as shown in Figure 1.

The reflection coefficient  $S_{11a}$  at the probe/material interface (the aperture port) of the open-ended sensor is obtained by considering the complex admittance of the equivalent circuit

$$S_{11a} \equiv |S_{11a}|e^{j\varphi} = \frac{Z_L - Z_C}{Z_L + Z_C} = \frac{1 - j\omega Z_C(\varepsilon_r C_0 + C_f)}{1 + j\omega Z_C(\varepsilon_r C_0 + C_f)} \quad (1)$$

where  $\omega = 2\pi f$  is the angular frequency,  $Z_C$  is the characteristic impedance of the transmission line, and  $\varepsilon_r = \varepsilon'_r - j\varepsilon''_r$  is the complex relative permittivity of the MUT. From (1)  $\varepsilon'_r$  and  $\varepsilon''_r$  are calculated as

$$\varepsilon'_r = \frac{-2|S_{11a}|\sin\varphi}{\omega Z_C C_0(1 + 2|S_{11a}|\cos\varphi + |S_{11a}|^2)} - \frac{C_f}{C_0} \quad (2)$$

$$\varepsilon''_r = \frac{1 - |S_{11a}|^2}{\omega Z_C C_0(1 + 2|S_{11a}|\cos\varphi + |S_{11a}|^2)} \quad (3)$$

In order to calculate the complex relative permittivity from the complex reflection coefficient we should know the values of  $C_f$  and  $C_0$ . These two parameters are usually obtained by calibrating the open-ended coaxial probe with a reference sample of known permittivity, for example deionized or distilled water, saline solution, ethanol or methanol. The two unknown parameters are then given by the following equations:

$$C_0 = \frac{1 - |S_{11a}|^2}{\omega Z_C \varepsilon''_r(1 + 2|S_{11a}|\cos\varphi + |S_{11a}|^2)} \quad (4)$$

$$C_f = \frac{-2|S_{11a}|\sin\varphi}{\omega Z_C(1 + 2|S_{11a}|\cos\varphi + |S_{11a}|^2)} - \varepsilon'_r C_0 \quad (5)$$

## 2.2. - ANTENNA MODEL

In this case the coaxial probe is considered as an antenna in a lossy dielectric [28]. The equivalent circuit consists of two capacitances ( $C_f$  and  $\varepsilon_r C_0$ ) and a radiation conductance ( $G_0 \varepsilon_r^{5/2}$ ), all connected in parallel, as shown in Figure 1. The admittance of the circuit is given by

$$Y = \frac{1}{Z_L} = j\omega C_f + j\omega \varepsilon_r C_0 + G_0 \varepsilon_r^{5/2} \quad (6)$$

Commonly,  $C_f \ll \varepsilon'_r C_0$  [17]. Neglecting  $C_f$  in comparison with  $\varepsilon'_r C_0$  in (6)

$$Y = j\omega \varepsilon_r C_0 + G_0 \varepsilon_r^{5/2} \quad (7)$$

Admittance  $Y$  is related to the reflection coefficient  $S_{11a}$  at the probe aperture as

$$Y = Y_C \frac{1 - S_{11a}}{1 + S_{11a}} \quad (8)$$

where  $Y_C = 1/Z_C$  is the characteristic admittance of the coaxial line.

The method of solving the complex equation (7) consists in splitting it into real and imaginary parts, obtaining thus a set of two real equations for the two real unknowns, which are either  $C_0$  and  $G_0$  (when calibrating the probe with a reference material) or the relative permittivity  $\varepsilon'_r$  and the loss factor  $\varepsilon''_r$  of MUT [6].

The splitting of equation (7) into real and imaginary parts gives a set of two real equations like this:

$$\begin{cases} Y' = a_{11}C_0 + a_{12}G_0 \\ Y'' = a_{21}C_0 + a_{22}G_0 \end{cases} \quad (9)$$

where the coefficients  $a_{11}$ ,  $a_{12}$ ,  $a_{21}$  and  $a_{22}$  depend on  $\varepsilon_r$ .

For the unknown  $C_0$  and  $G_0$  the system of equations (9) is linear and thus can be easily solved, but for the unknowns  $\varepsilon'_r$  and  $\varepsilon''_r$  it is nonlinear and must be solved using an iterative procedure. For this, an objective function is defined as  $F(\varepsilon_r) = Y(\varepsilon_r) - Y_m$ , where  $Y(\varepsilon_r)$  is the admittance of the aperture calculated using the system of equations (9), and  $Y_m$  is the admittance of the aperture obtained by measurement or simulation of  $S_{11a}$  and then calculated using equation (8). The permittivity of the sample can be calculated by finding the zero of the objective function.

## 2.3. - VIRTUAL LINE MODEL

The fringing field at the extremity of the open-ended coaxial probe terminated by a dielectric sample is modeled as a segment of equivalent lossy transmission line which has the same radial dimensions as the physical line, and a virtual length  $L_m$  (Figure 1) [29, 39].

There are two variants for the virtual line model: one in which  $L_s$  is the true length of the probe, and another in which  $L_s$  is a virtual length. In the first variant, one must consider the existence of a coupling admittance  $Y$  between the virtual line and the physical line, so that the load admittance of the physical line was equal to the input admittance of the virtual line. In this case the input admittance of the virtual line can be expressed as

$$Y_{BB'} = Y_{0m} \frac{Y_E + jY_{0m} \tan(\beta_m L_m)}{Y_{0m} + jY_E \tan(\beta_m L_m)} + Y \quad (10)$$

where

$$Y_{0m} = \frac{1}{Z_{0m}} = \frac{\sqrt{\varepsilon_r}}{60 \ln(b/a)} \quad (11)$$

is the characteristic admittance of the virtual line,  $Y_E$  is the load admittance of the virtual line,  $\varepsilon_r$  is the complex relative permittivity of the MUT, and  $\beta_m$  is the propagation constant in the MUT:

$$\beta_m = \omega \sqrt{\mu \varepsilon} = \omega \sqrt{\mu_0 \varepsilon_0 (\varepsilon'_r - j \varepsilon''_r)} = \frac{2\pi f \sqrt{\varepsilon_r}}{c} \quad (12)$$

Since the virtual line is terminated by an open circuit,  $Y_E = 0$ , (10) becomes

$$Y_{BB'} = jY_{0m} \tan(\beta_m L_m) + Y \quad (13)$$

The load admittance of the physical line can be expressed as:

$$Y_{BB'} = Y_{0s} \frac{1 - S_{11} e^{j2\beta_s L_s}}{1 + S_{11} e^{j2\beta_s L_s}} = Y_{0s} \frac{1 - S_{11a}}{1 + S_{11a}} \quad (14)$$

where

$$Y_{0s} = \frac{1}{Z_{0s}} = \frac{\sqrt{\varepsilon_{rs}}}{60 \ln(b/a)} \quad (15)$$

$$\beta_s = \omega \sqrt{\mu \varepsilon_s} = \omega \sqrt{\mu_0 \varepsilon_0 \varepsilon_{rs}} = \frac{2\pi f \sqrt{\varepsilon_{rs}}}{c} \quad (16)$$

$S_{11}$  is the reflection coefficient at the measurement port,  $Y_{0s} = Y_C$  is the characteristic admittance of the physical line, and  $\varepsilon_{rs}$  is the relative permittivity of the dielectric inside the probe. Equating the expressions (13) and (14) and substituting (11) and (15) in the result:

$$\frac{\sqrt{\varepsilon_{rs}}}{60 \ln(b/a)} \frac{1 - S_{11a}}{1 + S_{11a}} = j \frac{\sqrt{\varepsilon_r}}{60 \ln(b/a)} \tan(\beta_m L_m) + Y \quad (17)$$

or

$$\sqrt{\varepsilon_{rs}} \frac{1 - S_{11a}}{1 + S_{11a}} = j \sqrt{\varepsilon_r} \tan(\beta_m L_m) + Y 60 \ln(b/a) \quad (18)$$

Therefore, the relation between the reflection coefficient at the aperture port and the complex permittivity can be formulated into the following equations as:

$$y = \operatorname{Re} \left\{ \sqrt{\varepsilon_{rs}} \frac{1 - S_{11a}}{1 + S_{11a}} - j \sqrt{\varepsilon_r} \tan(\beta_m L_m) \right\} \quad (19)$$

$$\varepsilon_r = - \left[ \left( \frac{1 - S_{11a}}{1 + S_{11a}} \sqrt{\varepsilon_{rs}} - y \right) \cot(\beta_m L_m) \right]^2 \quad (20)$$

where  $y \equiv Y 60 \ln(b/a)$

The two unknown parameters  $L_m$  and  $y$  are calculated from (19) within an iterative procedure developed here that requires only one reference material, contrary to the iterative procedures used in [31] and [35] that requires two. In essence, for each value of the frequency  $f$  the value of the length  $L_m$  is searched that when being substituted in (19) gives the value of  $y$  that, when being substituted in (20), gives the permittivity of the reference material.

Once  $L_m$  and  $y$  are known, the complex relative permittivity of an unknown material can be found using the equations (12) and (20) within a second iterative procedure that minimizes the difference between the proposed value for the permittivity in the equation (12) and the calculated value for the permittivity using the equation (20).

### 3. - NUMERICAL SIMULATION OF THE REFLECTION COEFFICIENT

A numerical simulation based on the Finite Element Method (FEM) was used to calculate the reflection coefficient at the interface probe/sample in a range of frequency that goes from 300 MHz to 3 GHz with a step of 100 MHz. The wave equation for the electric field  $\mathbf{E}$  in the frequency domain

$$\nabla \times (\mu^{-1} \nabla \times \mathbf{E}) - \omega^2 \varepsilon \mathbf{E} = 0 \quad (21)$$

was solved in a 2D axisymmetric geometry. The domains of the model and the boundary conditions for the three types of open-ended coaxial probe and sample considered are shown in Figure 2.

An adaptive mesh of triangular elements was generated. The number of elements was 5 425 for the (a)-model, 5 014 for the (b)-model, and 2 274 for the (c)-model. In all cases the mesh maximum element size was 0.2 mm. For solving the generated system of linear equations the MUMPS code [40, 41] was used with a memory assignation of 1.2 and a pivot threshold of 0.1. The reflection coefficient at the probe's aperture port was calculated using

$$S_{11a} = S_{11} e^{j2\beta_s L_s} = \frac{\int_{port1} (\mathbf{E}_c - \mathbf{E}_1) \cdot \mathbf{E}_1^* dS_1}{\int_{port1} \mathbf{E}_1 \cdot \mathbf{E}_1^* dS_1} e^{j2\beta_s L_s} \quad (22)$$

where  $\mathbf{E}_c$  is the computed total electric field on the excitation port (port 1), and  $\mathbf{E}_1$  is the excitation analytical electric field.

The materials considered as sample were distilled water, 0.02 M NaCl(aq) solution, 0.5 M NaCl(aq) solution, ethanol, methanol, liver tissue, fat tissue, muscle tissue, kidney tissue, heart tissue, and blood. All these materials were modeled by Cole-Cole or Debye relaxation models [5]. Probes with  $a = 0.65$  mm,  $b = 2.05$  mm,  $L_s = 10$  mm, and  $\varepsilon_{rs} = 1.9$  were considered. A radius of 5.38 mm was assigned to the probe flange.

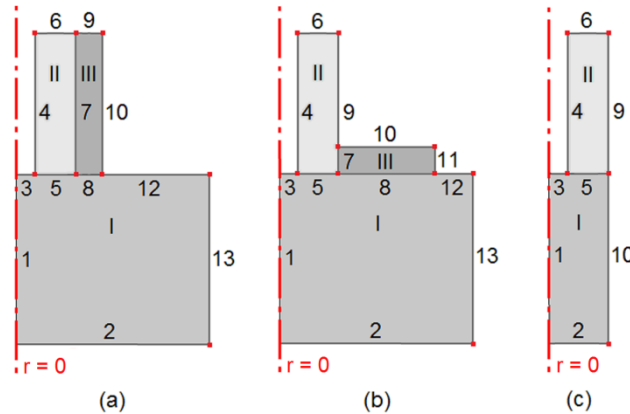


Figure 2

**Domains and boundary conditions for the three types of probe and sample. Probe types: (a)- without flange, (b)- with flange, (c)- open into a propagation circular wave guide. Domains: (I)- Sample; (II)- Dielectric inside the probe; (III)- Copper. Boundaries and conditions: (1)- Axial Symmetry, (3, 4, 9, 10, 11)- Perfect Electric Conductor, (6)- Excitation Coaxial Port, (2, 12, 13)- Absorbing, (5, 7, 8)- Continuity.**

## 4. - RESULTS

Using the results of the simulations and the permittivity extraction procedures described before, the relative permittivity  $\varepsilon'_r$  and the loss factor  $\varepsilon''_r$  as well as their errors  $\Delta\varepsilon'_r/\varepsilon'_r$  and  $\Delta\varepsilon''_r/\varepsilon''_r$  where determined for the six biological tissues, considering the all possible PMR combinations that can be made up from the three probe types, three probe models and the five reference liquids.

The maximum errors on  $\varepsilon'_r$  and  $\varepsilon''_r$  obtained for the liver tissue, the muscle tissue, the kidney tissue, the heart tissue, the blood and the fat tissue, using the 45 possible PMR combinations for each tissue are shown in Table 1. Since we are interested in good accuracy of measurements, we have highlighted the values that don't surpass, for example, 6% for both, the permittivity  $\varepsilon'_r$  and the loss factor  $\varepsilon''_r$ , when using the same PMR combination. Considering only the highlighted values we can make the following statements:

- 1) The coaxial probe without flange can be used for the determination of the complex permittivity of the liver tissue, the muscle tissue, the kidney tissue, the heart tissue, the blood and the fat tissue, only when using the antenna model.
- 2) The coaxial probe with flange can be used for the determination of the complex permittivity of the liver tissue, the muscle tissue, the kidney tissue, the heart tissue, the blood and the fat tissue, using the antenna model. Using the capacitive model this probe can be used for the determination of the complex permittivity of the blood and the fat tissue.
- 3) The coaxial probe open into a propagation circular wave guide can be used for the determination of the complex permittivity of the liver tissue, the muscle tissue, the kidney tissue, the heart tissue and the blood, using anyone of the three models.
- 4) The antenna model, with the distilled water or the saline 0.02 M NaCl(aq) solution as reference liquid, can be used for the determination of the complex permittivity of the liver tissue, the muscle tissue, the kidney tissue, the heart tissue and the blood, using anyone of the three considered types of coaxial probe.
- 5) The virtual line model is not appropriate for the determination of the complex permittivity of the fat tissue, using anyone of the three considered types of coaxial probe.
- 6) The capacitive model and the virtual line model are more appropriate for the coaxial probe without flange and for the coaxial probe open into a propagation circular wave guide.
- 7) The complex permittivity of the fat tissue is determined with greater accuracy using the coaxial probe open into a propagation circular wave guide, the capacitive model and the ethanol as reference liquid.

Table 1

Maximum errors on  $\epsilon'_r$  and  $\epsilon''_r$  for the six studied biological tissues, using the 45 possible PMR combinations for each tissue. The values that don't surpass 6% for both,  $\epsilon'_r$  and  $\epsilon''_r$ , when using the same PMR combination are highlighted. Legend: P1- probe without flange, P2- probe with flange, P3- probe open into a propagation circular wave guide, M1- capacitive model, M2- antenna model, M3- virtual line model, R1- distilled water, R2- 0.02 M NaCl(aq) solution, R3- ethanol, R4- methanol, R5 -0.5 M NaCl(aq) solution, No sol- no solution for  $\epsilon'_r$  or  $\epsilon''_r$  was found.

|    |    | Liver                              |      |      |                                     |      |      | Muscle                             |        |      |                                     |        |      |
|----|----|------------------------------------|------|------|-------------------------------------|------|------|------------------------------------|--------|------|-------------------------------------|--------|------|
|    |    | Maximum error on $\epsilon'_r$ (%) |      |      | Maximum error on $\epsilon''_r$ (%) |      |      | Maximum error on $\epsilon'_r$ (%) |        |      | Maximum error on $\epsilon''_r$ (%) |        |      |
|    |    | P1                                 | P2   | P3   | P1                                  | P2   | P3   | P1                                 | P2     | P3   | P1                                  | P2     | P3   |
| M1 | R1 | 21.2                               | 24.0 | 3.0  | 24.0                                | 28.1 | 8.0  | 11.5                               | 13.0   | 1.1  | 21.3                                | 23.8   | 5.8  |
|    | R2 | 11.5                               | 13.0 | 3.6  | 7.1                                 | 9.0  | 6.5  | 6.5                                | 7.6    | 1.1  | 7.1                                 | 4.7    | 5.8  |
|    | R3 | 0.8                                | 1.6  | 2.0  | 15.0                                | 16.5 | 6.9  | 1.3                                | 2.9    | 3.1  | 21.2                                | 24.5   | 9.5  |
|    | R4 | 1.3                                | 0.8  | 1.4  | 12.2                                | 13.5 | 5.5  | 1.5                                | 1.7    | 2.4  | 18.5                                | 20.9   | 7.9  |
|    | R5 | 13.4                               | 15.2 | 6.0  | 3.0                                 | 4.3  | 6.4  | 8.7                                | 10.2   | 3.8  | 3.0                                 | 2.1    | 4.1  |
| M2 | R1 | 5.2                                | 5.5  | 4.6  | 1.8                                 | 1.2  | 1.9  | 3.7                                | 4.1    | 3.6  | 2.7                                 | 1.5    | 1.5  |
|    | R2 | 5.6                                | 6.0  | 4.6  | 2.3                                 | 3.8  | 5.0  | 4.4                                | 4.4    | 3.6  | 3.1                                 | 4.5    | 5.4  |
|    | R3 | 6.8                                | 12.0 | 29.0 | 25.2                                | 40.0 | 40.0 | 7.9                                | 15.0   | 33.0 | 40.0                                | 60.0   | 55.0 |
|    | R4 | 2.8                                | 3.6  | 1.7  | 23.0                                | 18.1 | 2.0  | 3.0                                | 4.4    | 2.5  | 42.0                                | 34.5   | 3.3  |
|    | R5 | 7.0                                | 8.0  | 6.1  | 10.0                                | 7.5  | 9.0  | 6.2                                | 6.2    | 5.0  | 8.0                                 | 10.0   | 11.0 |
| M3 | R1 | 2.5                                | 2.9  | 1.8  | 22.0                                | 28.0 | 3.7  | 1.9                                | 2.2    | 1.4  | 15.4                                | 21.0   | 2.3  |
|    | R2 | 0.5                                | 0.6  | 0.9  | 25.0                                | 31.0 | 9.1  | 0.8                                | 1.3    | 0.5  | 18.6                                | 23.0   | 7.4  |
|    | R3 | 1.6                                | 1.3  | 3.4  | 11.0                                | 11.8 | 1.4  | 1.5                                | 1.2    | 3.2  | 16.0                                | 18.0   | 1.5  |
|    | R4 | 1.6                                | 2.1  | 0.5  | 10.0                                | 11.0 | 2.9  | 2.3                                | 2.7    | 0.8  | 15.5                                | 17.5   | 4.1  |
|    | R5 | 3.0                                | 3.6  | 0.9  | 36.0                                | 43.0 | 16.0 | 3.3                                | 4.4    | 0.8  | 28.5                                | 34.0   | 14.0 |
|    |    | Kidney                             |      |      |                                     |      |      | Heart                              |        |      |                                     |        |      |
|    |    | Maximum error on $\epsilon'_r$ (%) |      |      | Maximum error on $\epsilon''_r$ (%) |      |      | Maximum error on $\epsilon'_r$ (%) |        |      | Maximum error on $\epsilon''_r$ (%) |        |      |
|    |    | P1                                 | P2   | P3   | P1                                  | P2   | P3   | P1                                 | P2     | P3   | P1                                  | P2     | P3   |
| M1 | R1 | 11.4                               | 12.9 | 0.9  | 22.5                                | 26.1 | 6.0  | 9.8                                | 11.1   | 0.7  | 21.1                                | 25.0   | 5.4  |
|    | R2 | 6.0                                | 7.0  | 1.6  | 7.3                                 | 6.7  | 4.5  | 5.4                                | 6.4    | 1.5  | 7.1                                 | 5.4    | 4.0  |
|    | R3 | 1.0                                | 1.7  | 2.7  | 17.5                                | 20.0 | 9.3  | 0.9                                | 2.3    | 3.0  | 19.2                                | 22.8   | 9.9  |
|    | R4 | 1.8                                | 0.8  | 2.0  | 14.8                                | 17.0 | 7.7  | 1.7                                | 1.0    | 2.4  | 16.5                                | 19.0   | 8.4  |
|    | R5 | 8.5                                | 9.5  | 3.6  | 1.0                                 | 1.4  | 4.4  | 7.6                                | 8.9    | 3.4  | 2.0                                 | 0.7    | 3.7  |
| M2 | R1 | 4.0                                | 4.4  | 3.8  | 2.9                                 | 1.8  | 1.4  | 3.7                                | 4.1    | 3.6  | 3.4                                 | 2.0    | 1.5  |
|    | R2 | 4.3                                | 4.3  | 3.6  | 1.1                                 | 2.0  | 3.3  | 3.9                                | 4.1    | 3.3  | 1.8                                 | 2.6    | 3.9  |
|    | R3 | 10.0                               | 17.0 | 35.0 | 27.0                                | 37.0 | 26.0 | 9.6                                | 17.5   | 35.5 | 32.5                                | 45.1   | 34.0 |
|    | R4 | 5.8                                | 7.2  | 2.8  | 30.0                                | 23.0 | 3.0  | 5.0                                | 6.1    | 3.0  | 36.0                                | 28.0   | 3.2  |
|    | R5 | 6.0                                | 6.2  | 4.8  | 6.0                                 | 6.5  | 7.0  | 6.0                                | 6.0    | 4.6  | 7.0                                 | 8.0    | 8.5  |
| M3 | R1 | 2.7                                | 3.1  | 1.6  | 11.3                                | 15.2 | 1.5  | 2.4                                | 2.8    | 1.4  | 11.0                                | 15.0   | 1.3  |
|    | R2 | 0.7                                | 0.4  | 0.7  | 13.0                                | 16.0 | 5.0  | 0.5                                | 0.8    | 0.5  | 13.0                                | 15.8   | 5.0  |
|    | R3 | 1.8                                | 1.5  | 3.1  | 12.0                                | 13.9 | 1.5  | 1.7                                | 1.3    | 3.2  | 13.8                                | 15.8   | 1.5  |
|    | R4 | 1.5                                | 1.9  | 0.7  | 12.8                                | 14.5 | 4.0  | 1.8                                | 2.3    | 0.8  | 14.0                                | 16.1   | 4.2  |
|    | R5 | 2.7                                | 3.4  | 0.8  | 19.8                                | 23.0 | 9.5  | 2.8                                | 3.8    | 0.8  | 20.0                                | 23.2   | 10.0 |
|    |    | Blood                              |      |      |                                     |      |      | Fat                                |        |      |                                     |        |      |
|    |    | Maximum error on $\epsilon'_r$ (%) |      |      | Maximum error on $\epsilon''_r$ (%) |      |      | Maximum error on $\epsilon'_r$ (%) |        |      | Maximum error on $\epsilon''_r$ (%) |        |      |
|    |    | P1                                 | P2   | P3   | P1                                  | P2   | P3   | P1                                 | P2     | P3   | P1                                  | P2     | P3   |
| M1 | R1 | 7.1                                | 8.0  | 0.2  | 21.4                                | 24.0 | 4.6  | 411                                | 452    | 110  | 29.0                                | 37.0   | 13.0 |
|    | R2 | 3.9                                | 4.6  | 1.0  | 7.5                                 | 4.8  | 3.2  | 228                                | 250    | 105  | 14.0                                | 19.5   | 12.0 |
|    | R3 | 1.2                                | 2.4  | 3.4  | 20.0                                | 24.0 | 10.8 | 1.1                                | 0.6    | 1.5  | 6.5                                 | 3.2    | 1.8  |
|    | R4 | 1.9                                | 1.0  | 2.5  | 17.5                                | 20.0 | 9.3  | 16.0                               | 14.0   | 5.4  | 4.0                                 | 0.8    | 1.8  |
|    | R5 | 6.3                                | 7.4  | 2.8  | 1.9                                 | 1.6  | 2.9  | 220                                | 240    | 120  | 10.0                                | 15.2   | 11.8 |
| M2 | R1 | 3.3                                | 3.5  | 3.3  | 3.8                                 | 3.0  | 1.5  | 5.5                                | 7.2    | 3.6  | 2.5                                 | 6.3    | 6.4  |
|    | R2 | 3.4                                | 3.4  | 2.8  | 1.4                                 | 2.3  | 3.2  | 6.6                                | 8.0    | 3.9  | 2.5                                 | 7.8    | 7.0  |
|    | R3 | 10.9                               | 19.2 | 38.0 | 33.0                                | 44.0 | 29.0 | 1.6                                | 2.2    | 5.3  | 8.0                                 | 8.5    | 19.0 |
|    | R4 | 6.0                                | 7.1  | 3.6  | 39.0                                | 31.0 | 3.5  | 2.0                                | 2.0    | 3.1  | 2.5                                 | 3.3    | 1.7  |
|    | R5 | 5.0                                | 4.6  | 4.0  | 6.0                                 | 8.0  | 8.4  | 9.0                                | 10.5   | 6.0  | 6.0                                 | 10.0   | 9.5  |
| M3 | R1 | 2.5                                | 2.8  | 1.3  | 8.0                                 | 11.0 | 0.6  | 3.1                                | 3.2    | 4.7  | 85                                  | 85.0   | 63.0 |
|    | R2 | 0.6                                | 0.8  | 0.5  | 9.2                                 | 11.2 | 3.7  | 3.2                                | 3.2    | 4.8  | 90                                  | 85.0   | 79.0 |
|    | R3 | 1.8                                | 1.4  | 3.2  | 14.0                                | 16.5 | 1.6  | 2.3                                | 2.3    | 3.5  | 13.8                                | 17.8   | 35.0 |
|    | R4 | 1.7                                | 2.2  | 0.9  | 15.0                                | 17.5 | 4.8  | 4.4                                | 3.7    | 4.1  | 60.0                                | 52.0   | 16.0 |
|    | R5 | 2.8                                | 3.6  | 0.7  | 15.2                                | 17.5 | 8.0  | No sol                             | No sol | 5.5  | No sol                              | No sol | 87.0 |

Taking into account their feasible practical application, we considered also the PMR combinations made up from the same probes and reference liquids but different admittance models. Again, we focus our attention in the combinations such as both maximum errors, on  $\varepsilon_r'$  and on  $\varepsilon_r''$ , do not surpass 6%. From the analysis of the data tabulated in Table 1 we found 19 such combinations for the liver tissue, 25 for the muscle tissue, 28 for the kidney tissue, 25 for the heart tissue, 32 for the blood, and 16 for the fat tissue. Next we focus our attention in the PMR combinations that lead to the most accurate results for the permittivity of the studied tissues in the both cases, using the same admittance models and using different admittance models. These better combinations are shown in Table 2.

Table 2

Better PMR combinations for the measurement of  $\varepsilon_r'$  and  $\varepsilon_r''$  obtained for the studied tissues.

| Tissue | Using different admittance models |                   |                   |                   | Using the same admittance model |                   |                   |                   |
|--------|-----------------------------------|-------------------|-------------------|-------------------|---------------------------------|-------------------|-------------------|-------------------|
|        | $\varepsilon_r'$                  |                   | $\varepsilon_r''$ |                   | $\varepsilon_r'$                |                   | $\varepsilon_r''$ |                   |
|        | Comb.                             | Maximum error (%) | Comb.             | Maximum error (%) | Comb.                           | Maximum error (%) | Comb.             | Maximum error (%) |
| Liver  | P3M3R4                            | 0.5               | P3M2R4            | 2.0               | P3M3R4                          | 0.5               | P3M3R4            | 2.9               |
| Muscle | P3M1R1                            | 1.1               | P3M2R1            | 1.5               | P3M3R1                          | 1.4               | P3M3R1            | 2.3               |
| Kidney | P1M3R2                            | 0.7               | P1M2R2            | 1.1               | P3M3R1                          | 1.6               | P3M3R1            | 1.5               |
| Heart  | P3M1R1                            | 0.7               | P3M3R1            | 1.3               | P3M3R1                          | 1.4               | P3M3R1            | 1.3               |
| Blood  | P3M1R1                            | 0.2               | P3M3R1            | 0.6               | P3M3R1                          | 1.3               | P3M3R1            | 0.6               |
| Fat    | P2M2R4                            | 2.0               | P2M1R4            | 0.8               | P3M1R3                          | 1.5               | P3M1R3            | 1.8               |

Figure 3 shows the dielectric properties of liver tissue, muscle tissue, kidney tissue, heart tissue and blood obtained using the better PMR combinations with different admittance models. The corresponding theoretical curves are plotted for comparison.

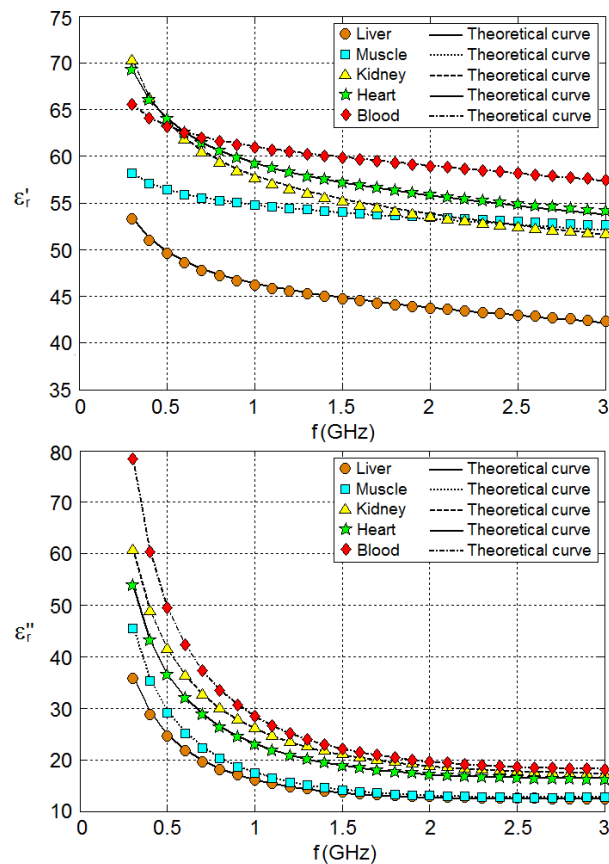


Figure 3

Dielectric properties of liver tissue, muscle tissue, kidney tissue, heart tissue and blood, obtained using the better PMR combinations with different admittance models.



## 5. - DISCUSSION

The results showed in Table 1 corroborates that when measuring the RF permittivity of biological materials using an open-ended coaxial line the proper selection of the probe type, probe model and reference material is crucial if good accuracy of measurements is expected. These reveal that overall the accuracy of measurements is better for the biological tissues of high water content (liver, muscle, kidney, heart, blood) than for the biological tissues of low water content (fat). According to obtained results we can state that when determining the permittivity of biological tissues of high water content using the PMR combinations with the same probes, reference liquids and admittance models, the probe which overall gives the best results is the coaxial probe open into a propagation circular wave guide, the models which overall gives the best results are the virtual line model and the antenna model, and the reference materials which overall gives the best results are the distilled water and 0.02 M NaCl(aq) solution. However, a disadvantage of the coaxial probe open into a propagation circular wave guide is that it cannot be used for the in vivo measurements, and the sample preparation may be difficult, especially when the probe is too small, except when the sample is liquid.

Ours results indicate that the PMR combinations which overall lead to the most accurate results of the permittivity for the studied biological tissues are those with the same probe and reference liquids, but with different admittance models (left part of the Table 2). They indicate also that the 0.5 M NaCl(aq) solution is not adequate as a reference material for dielectric property measurements on the considered biological tissues.

An advantage of the iterative procedure described here for the permittivity extraction when using the virtual line model is that, contrary to the iterative procedures used in [31] and [35] which requires two reference materials, only one reference material is needed for the determination of the two model parameters. Moreover, the execution of the permittivity extraction algorithm was about 10 times faster for the virtual line model than for the antenna model.

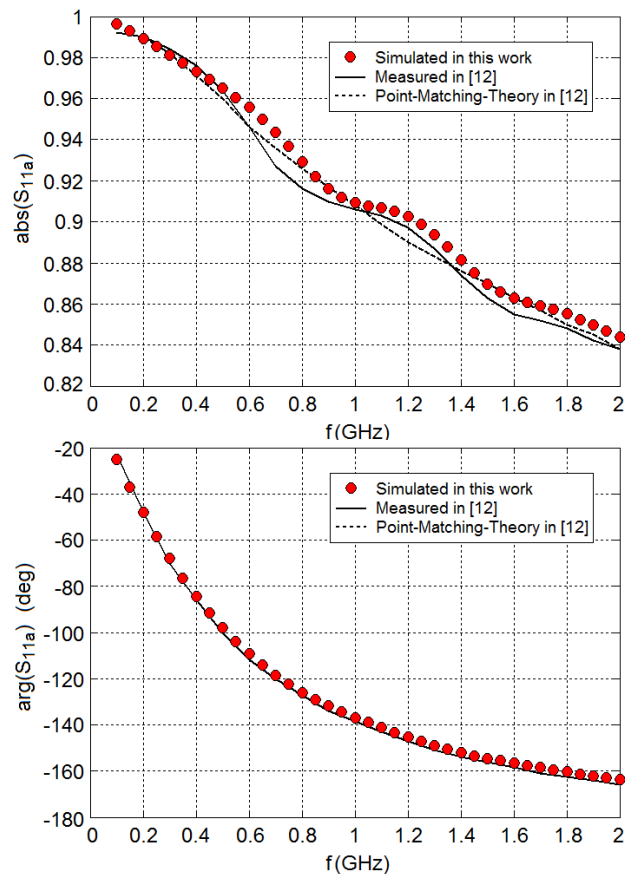


Figure 4

Modulus and phase of distilled water reflection coefficient for a coaxial probe similar to that studied in [32].

In [31] Bérubé et al. investigated the accuracy of four models in measuring 0.5 M NaCl(aq) and 1.0 M NaCl(aq) solution and their robustness as a function of the calibration materials in the frequency range from 50 MHz to 20 GHz. They used a coaxial probe without flange twice smaller than ours, but with the same characteristic impedance, and deionized water and methanol as reference materials. Three of their conclusions are connected with our work: (1)- The antenna model can give accurate values on  $\varepsilon''_r$ , however, its accuracy for  $\varepsilon'_r$  is poor, (2)- The best results are obtained by the virtual line model and the antenna model, being the obtained results for  $\varepsilon''_r$  accurate while those obtained for  $\varepsilon'_r$  acceptable, (3)- The capacitive model does not give accurate results for low frequencies, especially for  $\varepsilon''_r$ .

In this respect we can say that our results support the conclusion (1), but only when the distilled water or 0.02 M NaCl(aq) solution are used as reference material, since for the other reference materials the behavior is the opposite. Regarding conclusion (2), we have confirmed that the best results are obtained by the virtual line model and the antenna model, but, for the virtual line model, on the contrary, we obtained results for  $\varepsilon'_r$  more accurate than those obtained for  $\varepsilon''_r$ . Our results support also the conclusion (3), but not when the fat tissue is used as MUT and ethanol or methanol as reference material. It should be noticed that despite the fact that we investigated biological tissues instead of saline solutions, overall our results are consistent with those reported by Bérubé et al. in [31], nevertheless, they showed that the dielectric behavior of the saline solutions not always can be extrapolated to biological tissues.

The presented methodology constitutes a low cost tool that could be useful not only for the design of experiments, but also for the development of improved coaxial probes. Based on this methodology a computer program was developed and used here for data processing. Finally, we considered important to validate the numerical procedure used in this work for calculating the reflection coefficient at the interface probe/sample. With this aim, we present in Figure 4 the results obtained for the Finite Element model developed here of a coaxial probe with flange, similar to that studied in [32]. As we can see, our results agree quite well with those reported in that work.

## 6. - CONCLUSIONS AND FUTURE WORK

A numerical comparative study was made to assess the precision of three coaxial probe models (capacitive, antenna and virtual line), when measurements are made on liver, muscle, kidney, heart, blood and fat tissues, and the influence of both, the reference material (distilled water, 0.02 M NaCl(aq) solution, ethanol, methanol and 0.5 M NaCl(aq) solution) and the open-ended coaxial probe type (without flange, with flange, and open into a propagation circular wave guide) are considered. The Probe-Model-Reference combinations that lead to the most accurate results of the permittivity of the considered tissues at frequencies from 300 MHz to 3 GHz were investigated. It was found that the combinations which overall lead to the most accurate results of the permittivity for the studied biological tissues are those with the same probe and reference liquids, but with different admittance models.

The results corroborated that when measuring the RF permittivity of biological materials using an open-ended coaxial line the proper selection of the probe type, probe model and reference material is crucial if good accuracy of measurements is expected. They showed also that the dielectric behavior of the saline solutions not always can be extrapolated to biological tissues.

The presented methodology constitutes a low cost tool that could be useful for the design of experiments, for example, those for obtaining the temperature dependences of dielectric properties of biological tissues during their heating processes, which is fundamental to suitable model medical procedures such as thermotherapy and RF thermal ablation. This tool could be useful also for the development of improved coaxial probes.

In a future study we will consider other biological tissues and investigate the effect of the probe dimensions and the frequency range on the obtained results.

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