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K-space data processing for Magnetic Resonance Elastography (MRE)

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Short title: MRE using raw MR data

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Abstract

Object: Magnetic Resonance Elastography (MRE) requires substantial data processing based on phase image reconstruction, wave enhancement and inverse problem solving. The objective of this study is to propose a new, fast MRE method based on MR raw data processing, particularly adapted to applications requiring fast MRE measurement or high elastogram update rate.

Material and Methods: The proposed method allows measuring tissue elasticity directly from raw data without prior phase image reconstruction and without phase unwrapping. Experimental feasibility is assessed both in a gelatin phantom and in the liver of a porcine model in vivo. Elastograms are reconstructed with the raw MRE method and compared to those obtained using conventional MRE. In a third experiment, changes in elasticity are monitored in real-time in a gelatin phantom during its solidification by using both conventional MRE and raw MRE.

Results: The raw MRE method shows promising results by providing similar elasticity values to the ones obtained with conventional MRE methods while decreasing the number of processing steps and circumventing the delicate step of phase unwrapping. Limitations of the proposed method are the influence of the magnitude on the elastogram and the requirement for a minimum number of phase offsets.

Conclusion: This study demonstrates the feasibility of directly reconstructing elastograms from raw data.

Keywords: Elasticity Imaging Techniques; Magnetic Resonance Imaging; Radiology, Interventional

Introduction

Magnetic Resonance Elastography (MRE) is a non-invasive technique for measuring the mechanical properties of soft tissues. MRE allows to detect elasticity changes resulting from the presence of a pathology. For example, hepatic fibrosis [1, 2] has been shown to be associated with increased values of shear modulus. Higher modulus values may also reflect the presence of cancerous tissue as it has been shown in liver and breast [3–5]. On the contrary, neurodegenerative disorders may be related to lower stiffness values [6–8]. MRE requires a mechanical exciter that generates a wave propagating through the tissue. The displacement resulting from the propagation of the shear wave is encoded on phase images by using a specific MR-pulse sequence. An inverse problem solver allows reconstructing an elastogram based on the fact that the shear wave velocity is directly related to the underlying mechanical properties [9].

The duration of the overall MRE exam is highly variable as it depends on the nature of the information that is searched for. Highly quantitative MRE methods aiming at measuring advanced mechanical properties such as viscoelasticity or poroelasticity [5, 10–14] may require highly resolved images, 3D motion encoding in several slices, advanced post-processing and inverse problem solving techniques, that will make the whole process last from several minutes up to one hour. On the other extreme, MRE for interventional radiology [15, 16] aims at providing the physician with an updated elastogram as fast as possible. The whole acquisition and online reconstruction process has been shown to be achievable down to a couple of seconds in interventional MRE [16]. Such fast methods cannot be considered as quantitative as biomechanically-oriented MRE methods in terms of tissue mechanical properties that are calculated. However, they allow to provide the user with a stiffness parameter at a very high refresh rate, which is highly important in specific applications such as interventional radiology. As an example of how elastography may help to interventional procedures, tissue elasticity

has been reported as a valuable biomarker for the monitoring of thermal ablations in real-time [17, 18].

Fast MRE usually relies on the use of fractional encoding [19], i.e. the frequency of the motion-sensitizing gradient is higher than the MRE excitation frequency, and on the use of fast MR pulse sequences such as echo-planar imaging, fast gradient echo sequence or optimal k-space filling strategies [20–22]. Despite these acceleration strategies, such methods are based on classical MRE protocol, namely, the encoding of shear wave on phase images, the reconstruction of the phase images, filtering and processing of such phase images typically with temporal Fourier transform and phase unwrapping, and the resolution of the inverse problem on the resulting data. In this paper, we propose an alternative to this process aiming at accelerating and simplifying the reconstruction of the elastogram, for applications that require fast MRE information. The key feature of the proposed method is that spatial frequencies, and hence wavelengths, are estimated directly from the raw MR data, without any need for phase image reconstruction. This allows circumventing several steps such as spatial inverse and forward Fourier transforms and unwrapping algorithms.

Material and Methods

Theoretical background

As mentioned above, conventional MRE uses MR phase data for wave image and elastogram reconstruction. The proposed method uses directly the raw complex MRI data. A schematized comparison between the two methods is provided in **Fig. 1**. This section introduces the theoretical background of the proposed method.

Conventional process

New process

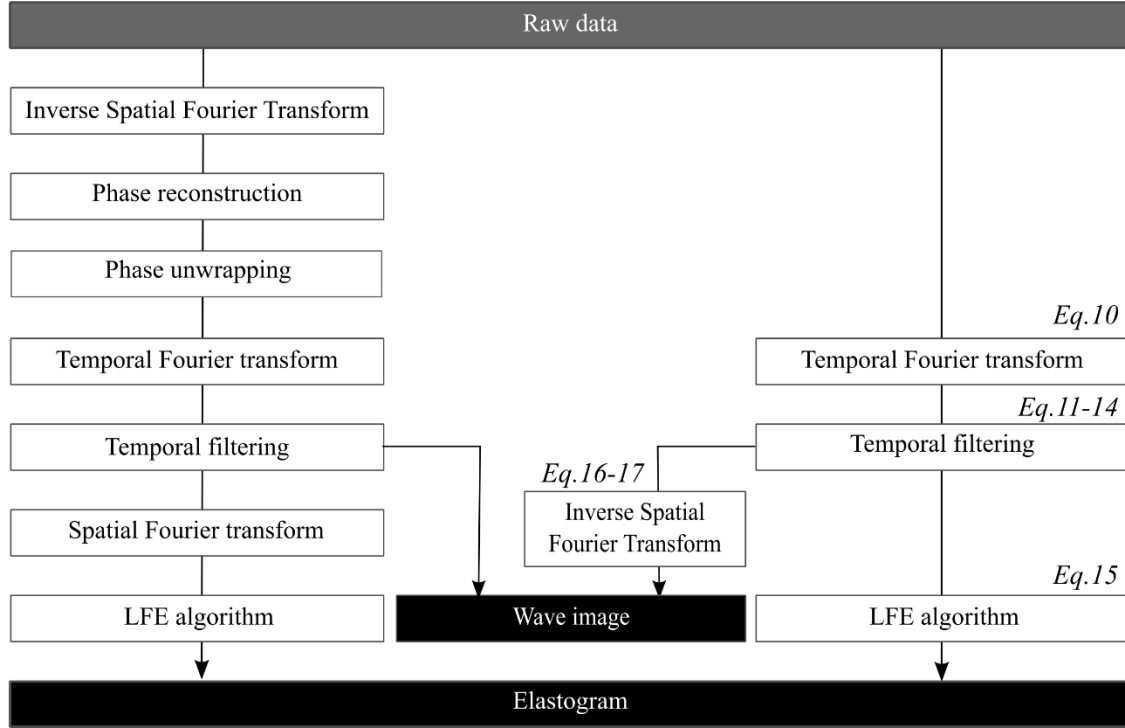


Fig. 1 Schematic description of the conventional MRE method and the proposed, raw data-based method

Conventional MRE

MRE relies on the encoding of tissue motion in the phase Φ of the complex MRI signal. Let us consider a harmonic oscillation of one isochromat with maximum amplitude u_0 at a given temporal frequency f_e , where \vec{u} is the position vector:

$$\vec{u}(t) = \vec{u}_m + \vec{u}_0 \sin(2\pi f_e t - \vec{k} \cdot \vec{u} + \alpha) \quad [1]$$

where \vec{k} is the wave number, \vec{u}_m is the mean position and α is the initial phase offset.

A number N_p of periods of a sinusoidal motion encoding gradient (MEG), called G with maximum amplitude G_0 is implemented in the MR-pulse sequence and its polarity is switched with respect to the mechanical excitation frequency f_e .

$$G(t) = \begin{cases} G_0 \sin(2\pi f_e t) & t \in [0, N_p/f_e] \\ 0 & \text{otherwise} \end{cases} \quad [2]$$

The observed MRI phase shift Φ_s acquired with a gradient echo sequence including the described MEG is proportional to the scalar product of the position vector and the gradient vector [9]:

$$\Phi_s = \gamma \frac{N_p (\vec{G}_0 \cdot \vec{u}_0)}{2f_e} \cos(\vec{k} \cdot \vec{u} - \alpha) \quad [3]$$

A similar equation can be written when the MEG frequency f_g and the mechanical excitation frequency f_e are not equal [19]:

$$\Phi_s = \gamma \frac{f_g \vec{G}_0 \cdot \vec{u}_0}{\pi(f_g^2 - f_e^2)} \sin\left(\frac{\pi f_e}{f_g}\right) \cos\left(\frac{\pi f_e}{f_g} - \vec{k} \cdot \vec{u} + \alpha\right) \quad [4]$$

More generally, the observed MRE phase shift can simply be written as:

$$\Phi_s = C_1 \cos(-\vec{k} \cdot \vec{u} + C_2) \quad [5]$$

The same expression can be obtained for spin echo sequences with different expressions for C_1 and C_2 . MRE relies on estimating the wavenumber k from the phase of the acquired MRI signal. The conventional protocol involves the reconstruction of the phase image within the range $[-\pi, \pi]$, the use of a phase unwrapping algorithm, the harmonic analysis on a set of images with different initial phase offsets α , and finally, the resolution of the inverse problem.

The objective of this study is to propose a new approach that circumvents several of the aforementioned steps while being more adapted to fast MRE protocols.

MRE using raw data

The proposed theoretical framework will be developed in 1D for clarity. It can be easily extended to 2D images. The raw MRI signal S acquired in the k-space can be defined with respect to the transverse magnetization $I(x)$ as:

$$S(v_x) = \sum_{x=0}^{N_x-1} I(x) e^{-j2\pi \frac{v_x x}{N_x}} \quad [6]$$

where x is the arbitrary discrete spatial dimension considered in this example, N_x is the total number of points, v_x is the discrete spatial frequency and $I(x)$ is the transverse magnetization $I(x) = M(x)e^{j\Phi(x)}$, M being its magnitude and Φ being its phase. In the specific case of MRE, Φ is given by:

$$\Phi(x) = \Phi_0(x) + \Phi_s(x) \quad [7]$$

where Φ_0 is the background phase, i.e. the constant phase obtained without motion encoding, and Φ_s is the phase shift related to the encoded displacement as described in Eq.[5].

The k-space acquired in MRE experiments can therefore be written as follows:

$$S(v_x) = \sum_{x=0}^{N_x-1} M(x) e^{j\Phi_0(x) + jC_1(x) \cos\left(-\frac{k_x x}{N_x} + C_2\right)} e^{-\frac{j2\pi v_x x}{N_x}} \quad [8]$$

where $\frac{k_x}{N_x}$ corresponds to the normalized discrete wave number.

Eq.8 includes an exponential of a cosine function that can be rewritten using the Jacobi-Anger expansion [23]

$$S(v_x) = \sum_{x=0}^{N_x-1} M(x) e^{j\Phi_0(x)} e^{-\frac{j2\pi v_x x}{N_x}} \sum_{n=-\infty}^{\infty} j^n J_n(C_1(x)) e^{-\frac{nk_x x}{N_x} + n C_2} \quad [9]$$

where J_n is the n^{th} Bessel function. This equation highlights the presence of an infinite number of harmonics of the spatial frequency of interest in the k-space due to the presence of a cosine pattern in the phase signal. A similar characteristic has already been described in the framework of motion compensation in MRI [24].

In order to illustrate this phenomenon, a simulated 1D MRE signal is represented in **Fig. 2** showing the simulated spectrum of the resulting MRE 1D-signal. In this figure and the following, only the modulus of complex signals are represented in the frequency domain.

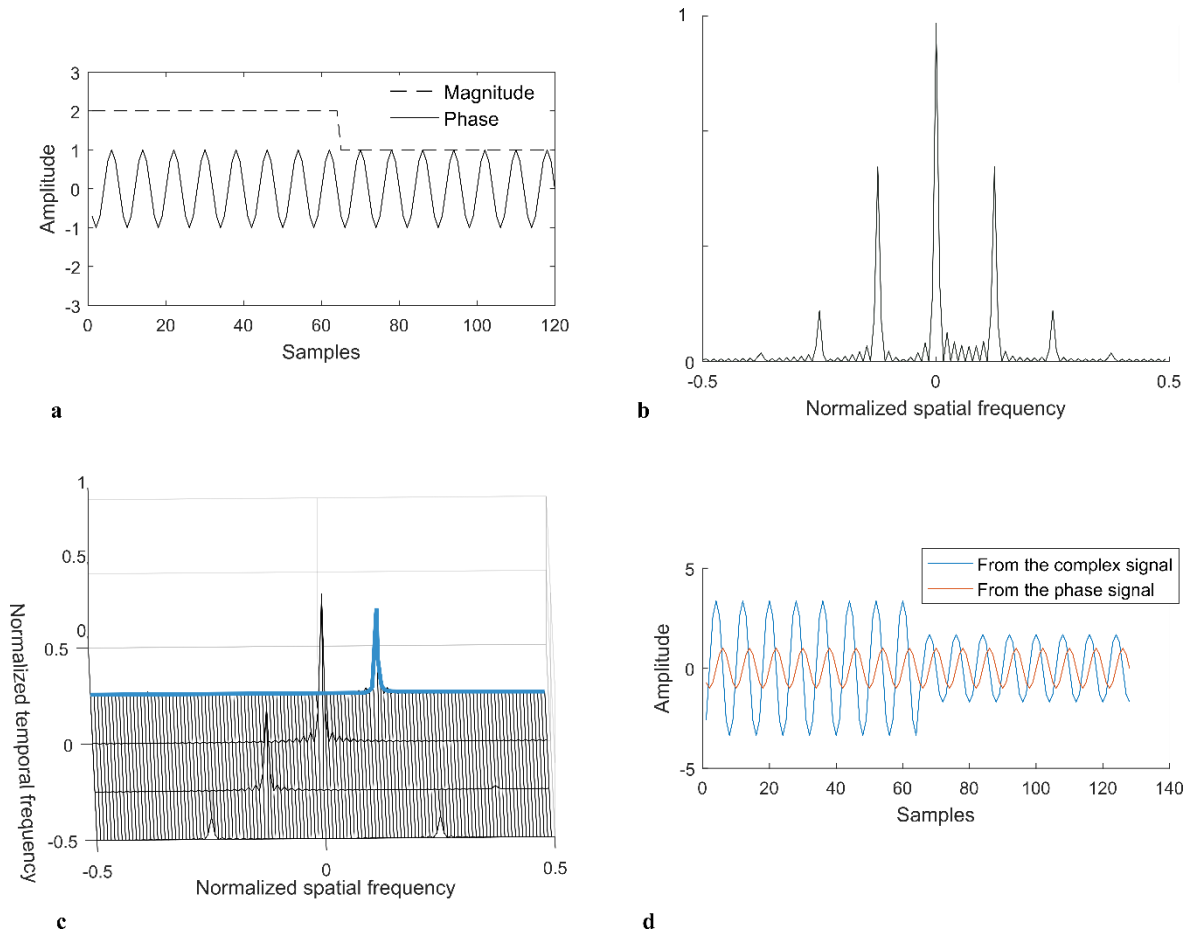


Fig. 2 a) Simulated 1D MR signal with heterogeneous magnitude and sinusoidal phase. b) Raw data corresponding to the simulated 1D MR signal. c) Plot of the temporal Fourier transform of a simulated data set

with 4 varying delays between excitation and MEG. Selection of the component of interest of the spatial frequency in blue. d) Wave signal obtained from the phase signal in red (conventional process) and wave signal obtained from the complex signal in blue (new proposed method). All ordinates are in arbitrary units

The spectrum is composed of an infinite number of harmonics as predicted by Eq.[9]. In order to remove all harmonics and select only the component of interest, a simple low pass filter could be used. However, the intrinsic heterogeneity of real tissue results in the presence of several spatial frequencies of interest (**Fig. 3**). Hence, distinguishing the component of interest of one frequency from the harmonic of another one is impossible.

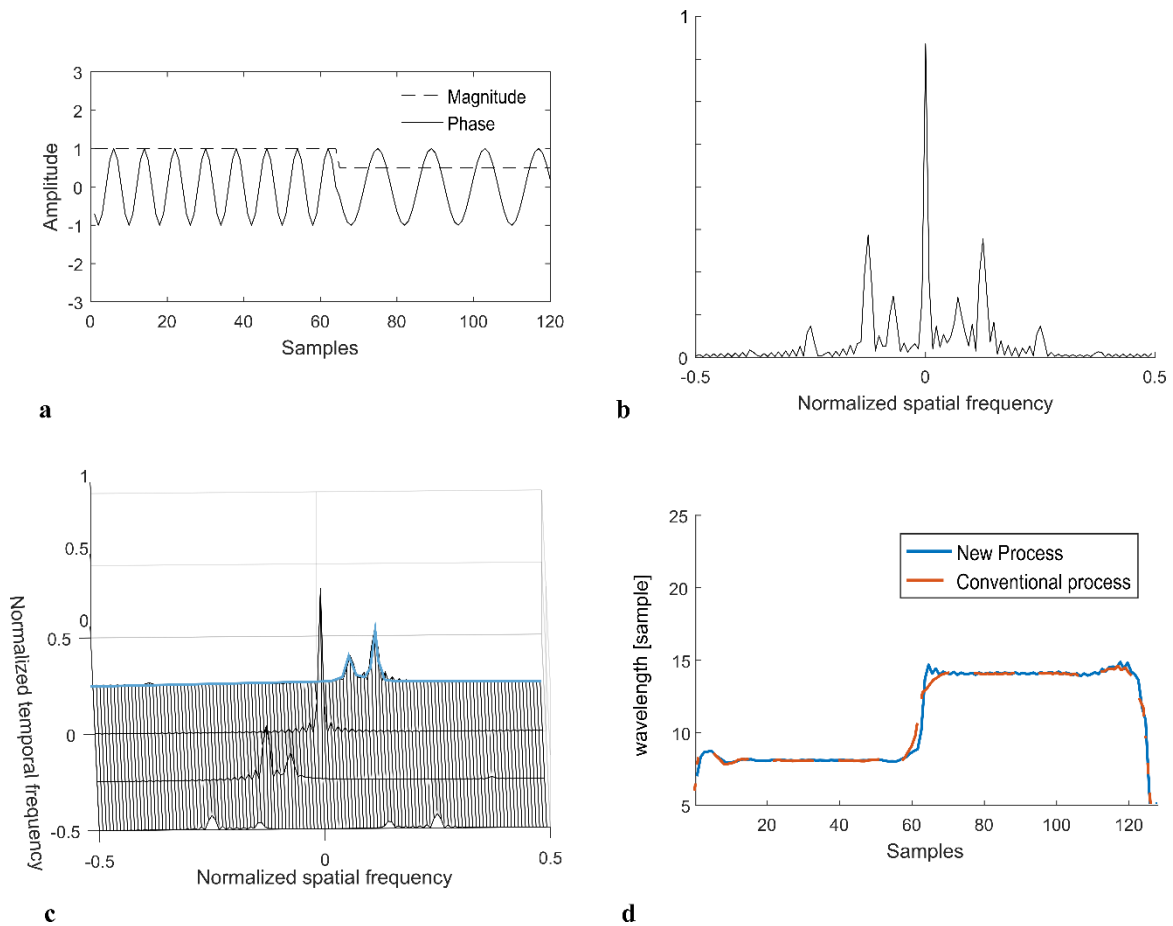


Fig. 3 a) Simulated 1D MR signal with heterogeneous magnitude and a signal composed of two frequencies. b) Raw data corresponding to the simulated 1D MR signal. c) Plot of the temporal Fourier transform of a simulated data set with 4 varying delays between excitation and MEG. Selection of the component of interest of the spatial

frequency in blue. d) Local frequencies returned by the LFE-algorithm performed on the filtered k-space (blue) compared to the conventional result obtained with the same filters (red) applied to phase signal

The proposed solution is inspired from the harmonic analysis used in conventional MRE. As previously mentioned, several images are acquired with varying phase shifts between mechanical excitation and motion encoding gradients. A temporal Fourier transform is subsequently performed on phase images in order to select only the displacement induced by the generated mechanical wave at a specific excitation frequency. The following paragraph aims at demonstrating that a similar analysis directly performed on the raw data allows identifying and extracting the components of interest.

Let us consider N_t acquisitions with varying phase offsets evenly spaced around the mechanical excitation cycle (i.e. sampling frequency f_s equal to $N_t f_e$). The temporal discrete Fourier transform of the series, including the Jacobi-Anger expansion yields:

$$S(v_x, v_t) \quad [10]$$

$$= \sum_{x=0}^{N_x-1} M(x) e^{j\Phi_0(x)} e^{-\frac{j2\pi(v_x x)}{N_x}} \sum_{n=-\infty}^{\infty} j^n J_n(C_1(x)) e^{-jn \frac{(k_x x)}{N_x} + jn C_2} \sum_{t=0}^{N_t-1} e^{-\frac{j2\pi(v_t t)}{N_t}} e^{\frac{jn(2\pi t)}{N_t}}$$

Figure 2.c illustrates the effects of the temporal Fourier transform on the previously simulated signal with 4 varying phase-offsets evenly spaced around the mechanical excitation. Spatial harmonics are thus separated over the temporal frequency range. The spatial component of interest is now distinguishable since it is associated to the temporal frequency of interest.

Let us consider the previous expression Eq.[10] at the temporal excitation frequency f_e (with $f_e = \frac{f_s}{N_t}$ which corresponds to $v_t=1$):

$$S(v_x, 1) \quad [11]$$

$$= \sum_{x=0}^{N_x-1} M(x) e^{j\phi_0(x)} e^{-\frac{j2\pi(v_x x)}{N_x}} \sum_{n=-\infty}^{\infty} j^n J_n(C_1(x)) e^{-jn\frac{(k_x x)}{N_x} + jn C_2} \sum_{t=0}^{N_t-1} e^{-j2\pi\frac{t}{N_t}} e^{jn2\pi\frac{t}{N_t}}$$

The sum over N_t is non null and equal to N_t only if $n = 1 + kN_t$ ($k \in Z^*$), therefore the Eq.[11] reduces to:

$$S(v_x, 1) = \sum_{n \in U} \sum_{x=0}^{N_x-1} M(x) e^{j\phi_0(x)} e^{-\frac{j2\pi(v_x x)}{N_x}} j^n N_t J_n(C_1(x)) e^{jn\left(-\frac{(k_x x)}{N_x} + C_2\right)} \quad [12]$$

$$U = \{n \mid n = kN_t + 1, k \in Z^*\}$$

Let us assume that the number of phase-offsets is sufficient to satisfy the following condition:

$$\log\left(\frac{|J_1(C_1(x))|}{|J_{kN_t+1}(C_1(x))|}\right) > 1, k \in Z^* \quad [13]$$

Details of this condition are discussed in Appendix A. Eq.[12] then simplifies to:

$$S(v_x, 1) = \sum_{x=0}^{N_x-1} M(x) e^{j\phi_0(x)} e^{-\frac{j2\pi(v_x x)}{N_x}} j N_t J_1(C_1(x)) e^{j\left(-\frac{k_x x}{N_x} + C_2\right)} \quad [14]$$

The component of interest is separated from the harmonics and selected (blue line in Fig. 2). The data associated to Eq.[14] will be referred to as the filtered k-space raw data, since only the spatial frequencies associated to the temporal frequency of interest are selected.

At this point, one of the interesting features of the proposed method is that it offers the possibility of estimating the wavelength distribution using the Local Frequency Estimation (LFE) algorithm. The LFE algorithm is a commonly used method for the resolution of the inverse problem in MRE. In conventional MRE with LFE processing, post-processed phase images are spatially Fourier transformed and filtered in order to find spatial distribution of spatial frequencies [25]. With our method, the LFE

calculations can be directly performed on the filtered k-space data (Eq. [14]), circumventing the need for an additional spatial Fourier transform step. For this purpose, classic image filters of the LFE algorithm [26] conventionally used in the frequency domain of the phase signal, are applied to the filtered k-space raw data. The local frequency of a real signal is defined as the derivative of the phase of its analytic signal. A pair of two lognormal quadrature filters R_1 and R_2 are classically applied in the frequency domain of the enhanced wave image and the local frequency f_l is given by:

$$f_l = \sqrt{f_1 f_2} \operatorname{Re} \left(\frac{F^{-1}(R_1 S)}{F^{-1}(R_2 S)} \right) \quad [15]$$

where f_1 and f_2 are the central frequencies of R_1 and R_2 , respectively. F^{-1} is the inverse Fourier transform. Usually, S is the signal in the spatial frequency domain of the enhanced wave image resulting from phase reconstruction, phase unwrapping, temporal frequency filtering and finally spatial Fourier transform. In this work, it is proposed to apply the same algorithm on k-space data obtained after temporal frequency filtering. In this case, S is described by Eq.[16]. This complete process is illustrated in **Fig. 3** on a 1D simulated MRI signal, yielding the local frequency for each sample. This method allows to reconstruct LFE-derived elastograms without phase image reconstruction and without spatial Fourier transform since LFE filtering is directly performed in the k-space. It is important to note that this method circumvents the problem of the phase wrapping, since the k-space of a signal with a phase wrapped in the range $[-\pi; \pi]$ and the same theoretical signal with a continuous phase out of the range $[-\pi; \pi]$ are exactly the same. The proposed method processes directly data in the k-space (Eq. [14]) and is therefore not affected by phase wrapping as would conventional phase images be. It must be noticed that the filtered k-space raw data is not the same as the k-space of the phase signal alone. Firstly, the filtered k-space raw data does not include high spatial frequencies related to the steep variations occurring in a wrapped phase. Secondly, the filtered k-space is convolved by the magnitude spectrum.

A qualitative visualization of the shear wave pattern is possible by performing an inverse spatial Fourier transform of the filtered k-space data, yielding the following equation:

$$s(x) = \frac{N_t}{N_x} M(x) e^{j\phi_0} J_1(C_1(x)) e^{j\left(-\frac{k_x x}{N_x} + C_2\right)} \quad [16]$$

The imaginary part of Eq. [16] is equal to:

$$\text{Im}(s(x)) = \frac{N_t}{N_x} M(x) J_1(C_1(x)) \cos\left(-\frac{k_x x}{N_x} + C_2 + \Phi_0\right) \quad [17]$$

It must be highlighted that the wave pattern described by Eq.[17] is weighted by the magnitude $M(x)$ and Bessel coefficients that depend on the encoded wave amplitude $C_1(x)$. Besides, a phase shift related to the background phase is introduced. The relationship between this signal and the actual displacement is therefore highly complex. This signal cannot be directly compared to the displacement image usually obtained in conventional MRE, and it is not directly exploitable for further processing. However, Eq. [17] allows obtaining a qualitative visualisation of the shear wave propagation without any need for phase image reconstruction.

Experimental Methods

Phantom experiments

The proposed method was evaluated on a gelatin phantom (17 cm in diameter) made of two parts with different concentrations (4% and 8%). A commercial pneumatic surface exciter (Resoundant, Rochester, MN) was placed on the top of the phantom in order to generate shear waves. The experiment was performed in a 1.5 T MRI scanner (MAGNETOM Aera, Siemens, Germany) and bipolar motion encoding gradients were implemented in a spoiled gradient echo sequence. The body antenna was used to obtain a single channel data set. The excitation and the encoding frequencies were set to 120 Hz and 220 Hz, respectively. The motion encoding direction was set through slice and the slice

was parallel to the surface of the pneumatic exciter. 4 phase-offsets evenly spaced across a mechanical period were acquired. Main imaging parameters are the following: FOV 340 mm x 340 mm, acquisition matrix 128 x 128, slice thickness 10 mm, MEG amplitude 20 mT/m, TE/TR 6.13/8.33 ms, flip angle 15° and bandwidth frequency 795 Hz/pixel. The proposed raw MRE method is used to reconstruct the wave pattern image and the elastogram, both directly from the raw data without reconstructing nor the phase nor the amplitude image. These results are compared to those obtained with conventional MRE involving phase image reconstruction, followed by the use of an unwrapping algorithm [27] and temporal Fourier transform. For this experiment and the two following ones, the LFE algorithm is applied with 10 pairs of filters in 4 orthogonal directions. The smaller central frequency of the filters is 0.07 cycle/sample.

In vivo experiments

In vivo experiments were performed on swine liver and were approved by the local ethics committee (ICOMETH C2EA - 38). The animal was anesthetized (propofol 3 mg kg⁻¹, pancuronium 0.2 mg kg⁻¹ and isoflurane 2 %) and placed in the large bore 1.5 T scanner. The body antenna was used. A surface pneumatic exciter was strapped around the chest and one slice perpendicular to the surface exciter was acquired. Relevant parameters of acquisition include: excitation/encoding frequencies 40/90 Hz, encoding direction: through slice, slice thickness 10 mm, FOV 320 mm x 320 mm, acquisition matrix 128 x 128, MEG amplitude 20 mT/m, TE/TR 12.7/25 ms, flip angle 15° and bandwidth frequency 795 Hz/pixel. 4 images with different phase-offsets evenly spaced across a mechanical period were acquired. The raw MRE method was compared to the conventional MRE in the same manner as in the phantom experiment. Images were acquired in a single end expiration breath-hold.

Real-time monitoring of elasticity changes in phantom

A third experiment was performed on a phantom in order to evaluate the possibility of monitoring elasticity changes without reconstructing any phase image. This experiment is similar to the experiment described in [16]. Shortly, a hollow gelatine phantom (10 %) was filled with freshly

prepared (still liquid) gelatine (15%) (**Fig. 4**). The elasticity changes occurring during the solidification of the gelatine were observed over time thanks to continuous acquisition of MRE data. MRE experiments were performed during the first 50 minutes of the cross-linking process of gelatin. During these first 50 minutes, the inclusion is still softer than the matrix even though its concentration in gelatin is higher.

A needle MRE driver was used to generate the mechanical wave. Relevant parameters of MRE acquisition include: excitation/encoding frequency: 120/290 Hz, MEG amplitude 20 mT/m, encoding direction: through slice, one slice orthogonal to the needle, slice thickness 10 mm, FOV 300 mm × 300 mm, acquisition matrix 128 × 128, TR/TE 8.34/5.62 ms, flip angle 15° and bandwidth frequency 780 Hz/pixel. 3 phase offsets were acquired. The 12 cm loop coil was placed around the needle MRE driver and used for data acquisition.

The proposed method and the conventional method were compared for estimating the variation of elasticity in the region of interest. In the conventional MRE method, phase images are reconstructed, unwrapped, and submitted to a temporal Fourier transform. Finally, only one frequency is selected and the LFE is applied to the resulting images (Fig. 4.b). In the proposed method, elastograms are reconstructed as described in the theoretical part, without reconstructing any phase image. The temporal Fourier transform is directly applied on the k-space raw data (Fig. 4.c). The LFE is applied on the resulting filtered data (Fig. 4.d). The elasticity in the same region of interest is calculated as a function of time.

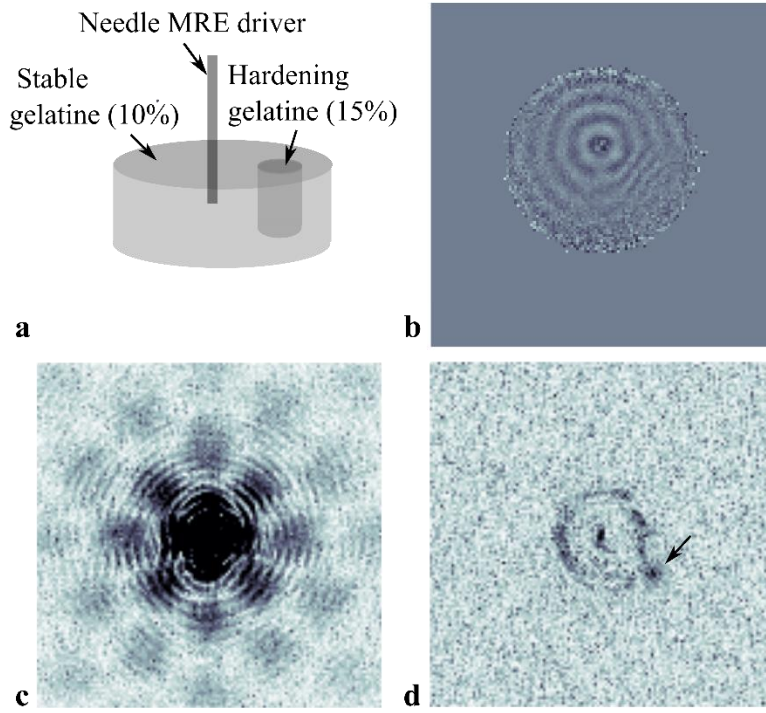


Fig. 4 a) The needle MRE driver is inserted in a stable gelatine phantom (10 %). A small part of the phantom is composed of gelatine (15 %) still under solidification process. b) Conventional phase image after phase unwrapping and temporal filtering on resulting phase images. c) Magnitude of the k-space raw data. d) Magnitude of the filtered k-space raw data at $t = 1280$ s. The ring corresponds to the frequency of the wave propagating in the stable gelatin, the small region indicated by the arrow corresponds to the wave propagating in the hardening gelatin.

Results

Fig. 5 illustrates the results obtained from a conventional MRE processing including phase unwrapping (left column) and the proposed method without phase image reconstruction (right column). A qualitative shear wave pattern obtained using Eq. [17] is displayed (Fig. 5.b) for visual comparison with a displacement image obtained by conventional MRE (Fig. 5.a). LFE-derived elastograms obtained by both methods are very similar (Fig. 5.c-d). Mean values and standard deviations of the local wavelength estimation obtained with the raw MRE method and the conventional method are

respectively 20.9 ± 10.9 mm and 19.81 ± 10.7 mm in the left softer half, and 30.7 ± 4.8 mm and 32.7 ± 3.8 mm in the right stiffer half.

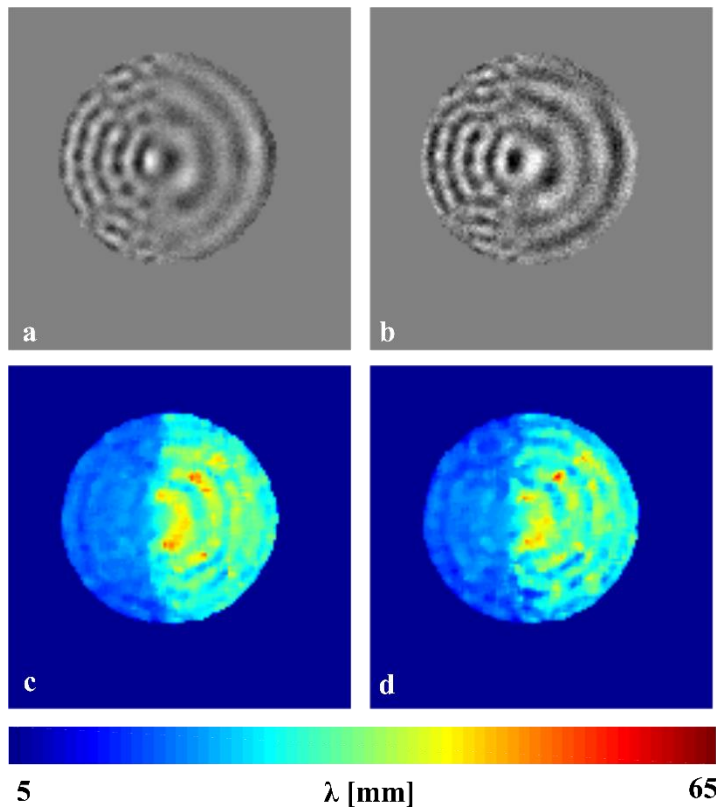


Fig. 5 (At the top) (a) Wave image obtained by unwrapping the phase image (conventional method) compared to the wave pattern image returned by the raw MRE method (b). (At the bottom) (c) Elastogram reconstructed by the conventional method compared to the one returned by the raw MRE method (d). A median filter (kernel 4x4) was applied on the 2 resulting elastograms

Wave pattern images and LFE-derived elastograms obtained in vivo by the two methods are displayed in **Fig. 6**. The local wavelength obtained by the raw MRE method and conventional MRE in the liver is equal to 28.45 ± 5.44 mm and 30.50 ± 7.9 mm, respectively. In both phantom and in vivo experiments, mean values and standard deviations of the local wavelength obtained with the proposed method were found to be very close to those obtained with the conventional one.

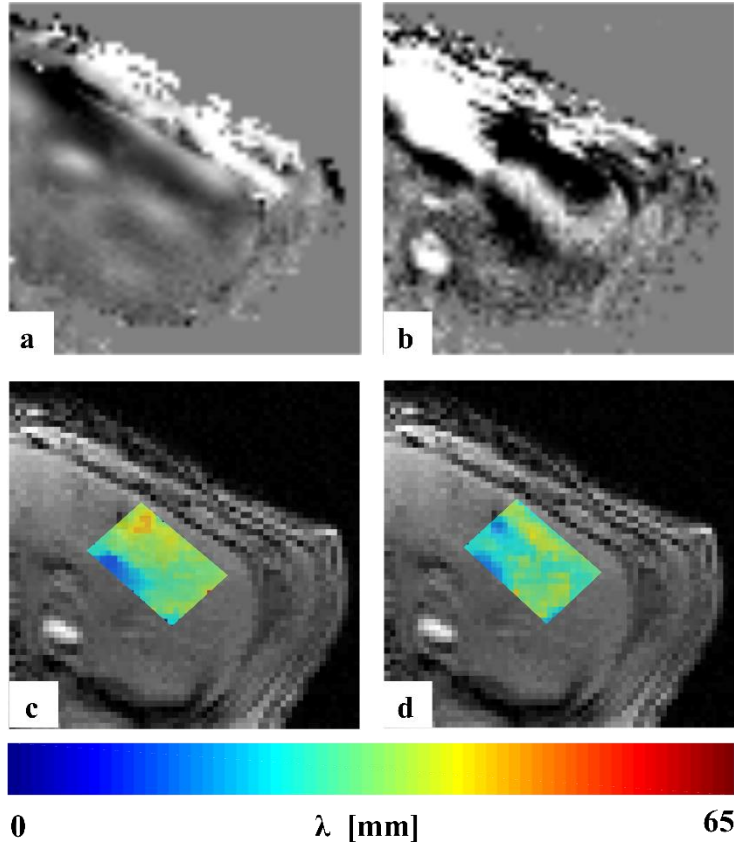


Fig. 6 (At the top) (a) Wave image obtained by unwrapping the phase image (conventional method) compared to the wave pattern image returned by the raw MRE method (b). (At the bottom) (c) Elastogram reconstructed by the conventional method compared to elastogram returned by the raw MRE method (d). A median filter (kernel 4x4) was applied on the two resulting elastograms

The variation of elasticity observed during the solidification of the gelatin is plotted in **Fig. 7**. Elasticity is estimated in real-time without phase image reconstruction and compared to the elasticity measured by the conventional MRE method. The evolution of the shear modulus over time obtained with the proposed method (red curve) is very similar to the one obtained with the conventional MRE method (blue curve).

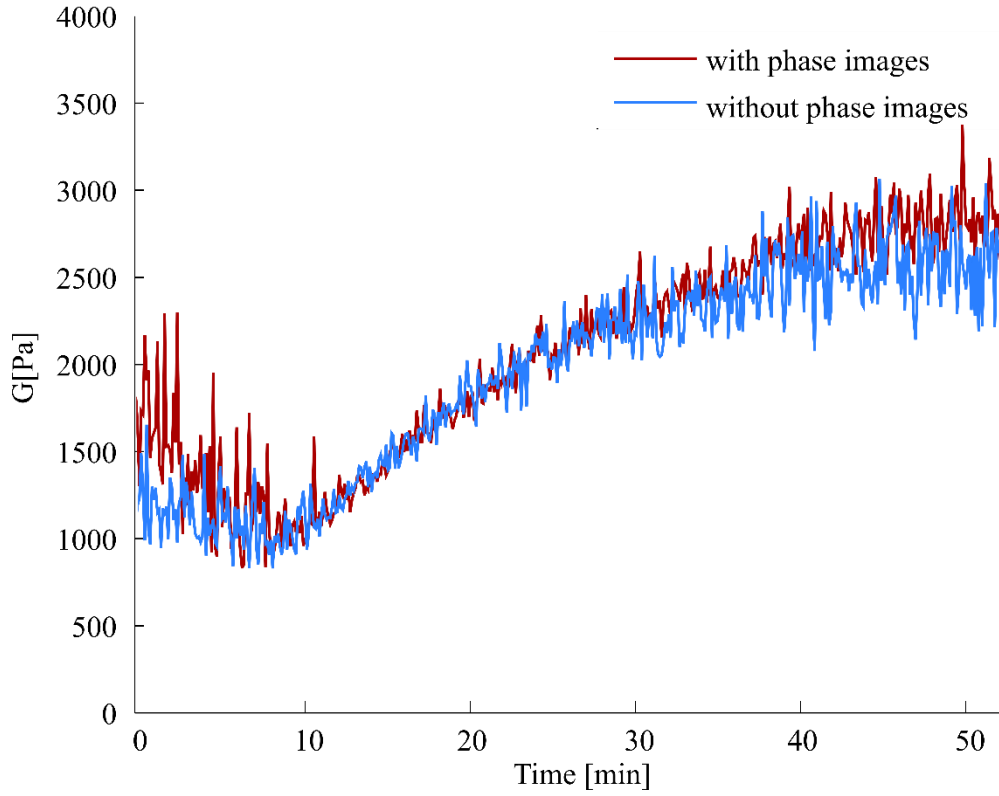


Fig. 7 Evolution of shear elastic modulus of the gelatin during its solidification measured on elastograms reconstructed by the conventional MRE (red), and the proposed raw data-based method (blue)

Discussion

This study introduces an alternative to the conventional processing protocol used in MRE. Elastograms can be reconstructed using directly raw MRI data, without phase reconstruction, phase unwrapping and spatial 2D transforms, as summarized in **Fig. 1**. This simplified and accelerated process is particularly interesting for applications that require very fast elastogram reconstruction or those requiring high elastogram update rate, such as in interventional radiology. The raw data obtained during an MRE acquisition is not easily readable because of the infinite number of harmonics induced by the presence of one sinusoidal pattern in the phase signal. The solution proposed here to overcome this difficulty is to perform the harmonic analysis directly on raw data. The method was assessed experimentally in gelatin phantoms and in vivo, and results were found to be very similar to those obtained with the conventional MRE process with phase unwrapping. The potential of the proposed

method for providing elasticity changes at high temporal sampling rate was demonstrated in a third experiment where elasticity changes were monitored in gelatin during its solidification.

Fundamental aspects of the proposed method need to be discussed. As previously highlighted, this method is sensitive to the choice of the number of phase-offsets. This number must be high enough to allow the selection of only one frequency component. However, this method is intended to applications requiring fast MRE information, which assumes the use of particularly fast MRE sequences. In most of the cases, such sequences involve lower amplitude encoding than conventional MRE sequences due to acceleration strategies such as fractional encoding. This makes the condition outlined in Appendix A easier to reach. The second point that deserves to be mentioned is the assumption concerning the phase ϕ_0 accumulated without the motion encoding gradient. The present theory relies on the assumption that ϕ_0 is locally uniform. Should this assumption not be valid, the phase offset could be removed similarly to what is commonly done in conventional MRE, i.e. by performing a difference between two images acquired with opposite MEG polarities. However, in this case, the phase difference could then be performed through a direct product in the spatial complex domain or a convolution in the frequency complex domain, the former avoiding the need for an additional direct/inverse spatial Fourier transform. One of the main advantages of the proposed method is that it is inherently immune to phase wrapping, a known challenge in MRE [28, 29] because its framework remains in the complex domain. Indeed, a complex signal with a theoretical phase signal out of the range $[-\pi; \pi]$ and the same signal with a phase wrapped in the range $[-\pi; \pi]$ have exactly the same spectrum.

The major limitation of the reconstruction of elastograms without phase images is that it is limited to the use of a unique inverse problem solving approach, namely, the LFE method. This algorithm is particularly well adapted for the proposed method, as it operates on k-space data usually obtained through spatial Fourier transform of filtered phase images. With the proposed method, the LFE can be directly applied on the filtered k-space raw data, without any need for phase image reconstruction and subsequent spatial Fourier transform. One must also remain aware that the filtered k-space is convolved by the magnitude spectrum, as seen in Eq. [14]. Although this influence is deemed to be

negligible in most cases, high frequencies corresponding to strong magnitude variations may add noise to the filtered k-space and influence the elastogram. *Further work is needed to investigate such artifacts on the resulting elastogram. This may be particularly important in situations with significant physiological motion, resulting in residual, potentially periodic patterns that are not due to MRE shear wave motion. In such cases, motion correction methods may be necessary.*

As we have already mentioned, a qualitative wave pattern image can be obtained without prior phase image reconstruction (Eq. [17]). However, the wave pattern image returned by the proposed method is weighted by different parameters, such as the magnitude image, the mechanical wave amplitude and the parameters of motion encoding. Therefore, such signal is not directly exploitable at this point beyond the sole qualitative visualization of the wave field, and it cannot be compared to the usual displacement image in conventional MRE used for further processing and other inverse solving methods. The relationship between the signal described by Eq. [17] and the actual displacement, as well as its sensitivity to noise compared to the one in conventional MRE need to be thoroughly studied to evaluate whether this method could be coupled with other inverse problem solving methods and post-processing techniques.

Substantial developments are still needed to make this method compatible with multi-channel arrays and undersampled k-space raw data. However, this approach could lead to new applications and alternative ways of estimating biomechanical properties in the future. Ongoing work is focused on in vivo monitoring of thermal ablations using this method for elastogram reconstruction in real time.

Conclusion

An alternative method for MRE data processing is proposed here. Elastograms can be independently computed without prior phase image reconstruction. Despite its limitations regarding the minimum number of phase offsets and the influence of the magnitude on the elastogram, this method is particularly well suited for applications requiring fast update rate of MRE information and it provides an interesting alternative to the conventional MRE process.

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Compliance with Ethical standards

Conflict of Interest: The authors declare no conflict of interest.

Research involving Animals: All applicable international, national, and/or institutional guidelines for the care and use of animals were followed. In vivo experiments of this study were performed according a protocol approved by a local ethics committee (ICOMETH C2EA - 38).

Informed consent: Not applicable to this study.

Author's contribution

Corbin: Protocol & Project development; Data collection or management; Data Analysis

Breton: Protocol & Project development; Data collection or management

De Mathelin: Protocol & Project development;

Vappou: Protocol & Project development; Data collection or management

Appendix

A temporal Fourier transform is applied on a set of k-space raw data acquired with varying delays between mechanical excitation and motion encoding gradients. By selecting only the temporal frequency of interest, most of the harmonics of the spatial frequency are removed.

The result is described by Eq. [12]:

$$S(v_x, 1) = \sum_{n \in U} \sum_{x=0}^{N_x-1} M(x) e^{j\phi_0(x)} e^{-\frac{j2\pi(v_x x)}{N_x}} j^n N_t J_n(C_1(x)) e^{jn\left(-\frac{k_x x}{N_x} + C_2\right)}$$

$$U = \{n \mid n = kN_t + 1, k \in \mathbb{Z}\}$$

As described by the set U , the component of interest ($n = 1$) is not the only one component selected by the filtering because of aliasing. In most cases, the other components are negligible compared to the first one (blue signal in **Fig. 8.b**), nevertheless the selection of other non-negligible components remains possible when the number of phase-offsets is poorly chosen as illustrated in **Fig. 8.c-d** (red signal). It is therefore relevant to assess conditions that favor an optimal ratio between the amplitude of the 1st and the $(kN_t + 1)^{\text{th}}$, $k \in \mathbb{Z}^*$ components of the spectrum that are susceptible to being associated to the same temporal frequency. The ratio R is deduced from the previous equation:

$$R = \log\left(\frac{J_1(C_1)}{J_{kN_t+1}(C_1)}\right)$$

Let us consider that a component is negligible compared to the first one when its amplitude is at least 10 times inferior, i.e. when $R > 1$. The number of phase-offsets N_t required to satisfy this condition is plotted in **Fig. 9** with respect to the C_1 parameter. Under the assumption that $R > 1$, Eq.[12] simplifies to Eq.[14]

$$S(v_x, 1) = \sum_{x=0}^{N_x-1} M(x) e^{j\phi_0(x)} e^{-\frac{j2\pi(v_x x)}{N_x}} j N_t J_1(C_1(x)) e^{j\left(-\frac{k_x x}{N_x} + C_2\right)}$$

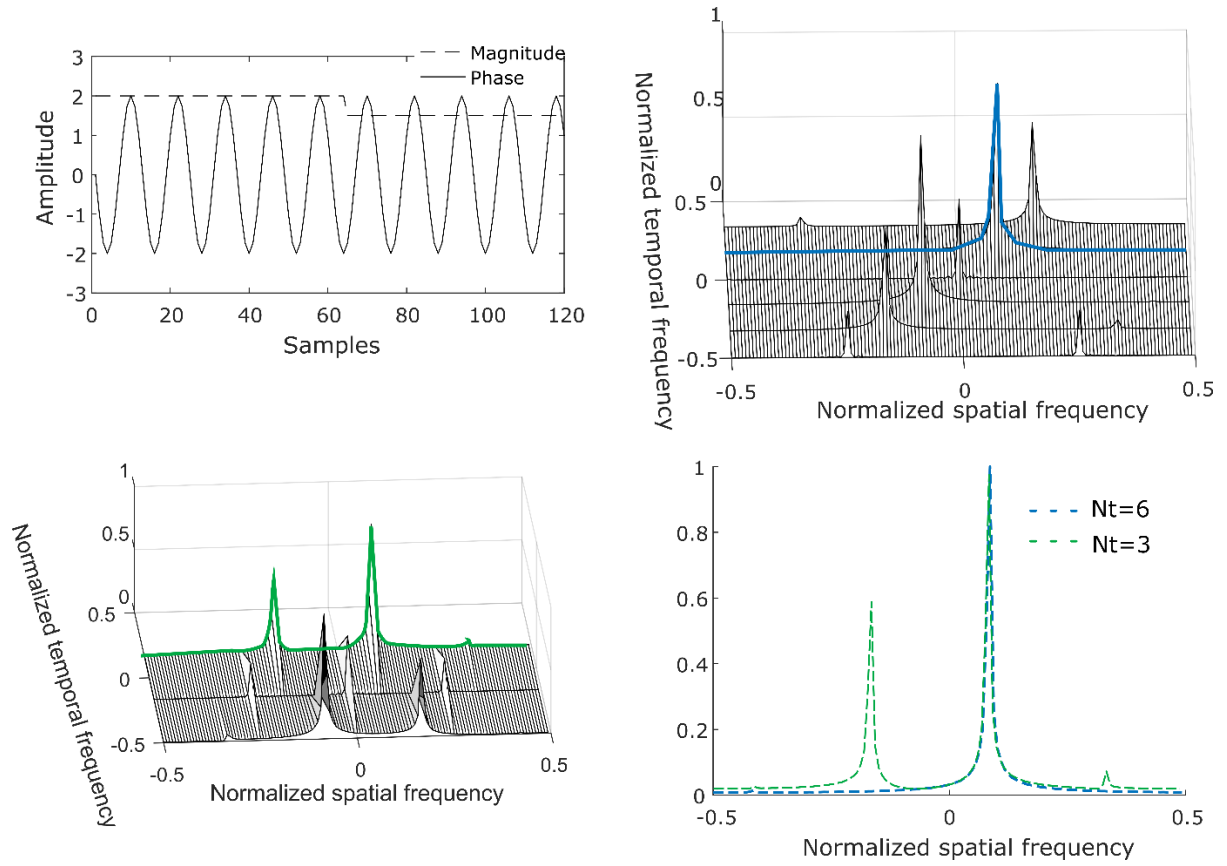


Fig. 8 a) Simulated 1D MR signal with heterogeneous magnitude and sinusoidal phase. b) Plot of the temporal Fourier transform of the previous k-space performed from the data set simulating 6 varying delays between motion encoding gradients and mechanical excitation. c) Plot of the temporal Fourier transform performed from the data set simulating 3 varying delays between motion encoding gradients and mechanical excitation. d) Plot of the filtering at the temporal frequency of interest in blue for the case with 6 phase-offsets and in red for the case with 3 phase-offsets

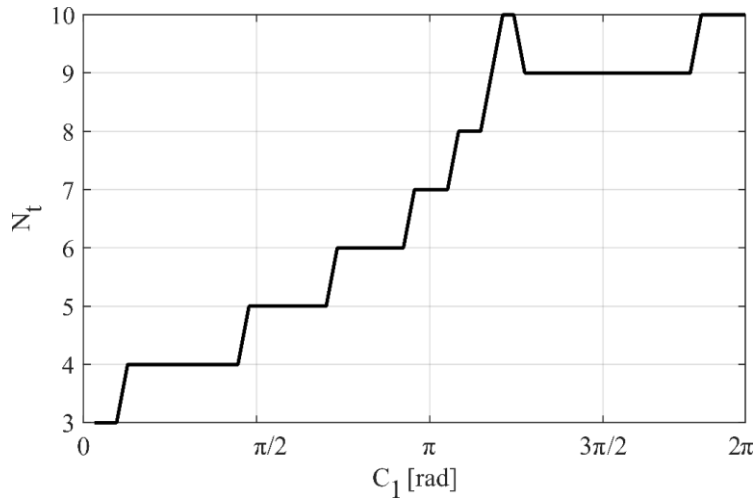


Fig. 9 Assessment of the minimal number of phase-offsets N_t required to satisfy the condition $R > 1$ with respect to the amplitude of the encoded wave C_1

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