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# RESTORATION OF ULTRASONIC IMAGES USING NON-LINEAR SYSTEM IDENTIFICATION AND DECONVOLUTION

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## ABSTRACT

This paper studies a new ultrasound image restoration method based on a non-linear forward model. A Hammerstein polynomial-based non-linear image formation model is considered to identify the system impulse response in baseband and around the second harmonic. The identification process is followed by a joint deconvolution technique minimizing an appropriate cost function. This cost function is constructed from two data fidelity terms exploiting the linear and non-linear model components, penalized by an additive  $\ell_1$ -norm regularization enforcing sparsity of the solution. An alternating optimization approach is considered to minimize the penalized cost function, allowing the tissue reflectivity function to be estimated. Results on synthetic ultrasound images are finally presented to evaluate the algorithm performance.

**Index Terms**— non-linear model, system identification, harmonic ultrasonic imaging, optimization, ADMM, polynomial Hammerstein model, deconvolution.

## 1. INTRODUCTION

Ultrasound (US) imaging is an effective, low cost, harmless and non-invasive medical imaging modality [1]. It is therefore widely used for clinical diagnosis (mainly for soft tissue applications related to cardiovascular and various cancers), blood flow velocity assessment and obstetrics. However, because of instrumentation constraints, US images suffer from a relatively low contrast, reduced spatial resolution and signal-to-noise ratio. Specifically, US images are contaminated by an intrinsic noise called speckle, which appears as a granular texture in the image. It is well known that speckle deeply reduces the general image quality in terms of contrast and resolution and increases the margin of inaccuracy in diagnostic analysis. Improving US image quality through post-processing image restoration techniques is therefore an important research area. Note that instrumentation approaches or pre-processing methods such as non-conventional beamformers have been also explored in the context of US [2]. In addition to the aforementioned limitations of ultrasound imaging, the propagation of ultrasound in many tissues is non-linear. Consequently, the resulting signals are combination of linear and non-linear components. This paper explores a deconvolution approach based on a new non-linear image formation model. This model is able to express different degrees of nonlinearity generated by the tissues themselves or by ultrasound contrast agents [3]. Most of the existing US image restoration methods use the first order Born approximation and consequently assume that the tissue reflectivity function (TRF) is related to the US radiofrequency (RF) image through a 2D convolution with the system point spread func-

tion (PSF) [4–7]. However, this forward model widely explored in the literature does not exploit the non-linear relationships that can exist between the unknown TRF and the observed image. We propose herein a new image restoration method originating from a study of non-linear ultrasonic wave propagation in a medium [1]. More specifically, we investigate a different perspective for US image deconvolution, *i.e.*, we estimate the TRF by eliminating the contribution of the system excitation and the PSF using a non-linear image formation model. The non-linearity considered in this work results from a polynomial Hammerstein model allowing PSF identification. More specifically, we focus in this paper on identification methods for a generalized polynomial Hammerstein model based on an exponential swept-sine input signal, allowing one-path estimation of the unknown linear filters. This identification method, originally proposed by Farina *et al.* [8] and further explored by Novak *et al.* [9] and Rebillat *et al.* [10], allows the TRF of the explored medium to be estimated. The proposed estimation method optimizes a joint cost function based on both the linear and non-linear components of the model using the alternating direction method of multipliers (ADMM). The remainder of the paper is organized as follows. Section 2 describes the non-linear model considered in this study and the corresponding parameter identification approach. Section 3 introduces the proposed image restoration framework and the corresponding ADMM algorithm. Simulation results are reported in Section 4 before concluding the paper.

## 2. PROBLEM STATEMENT

### 2.1. Nonlinear Model

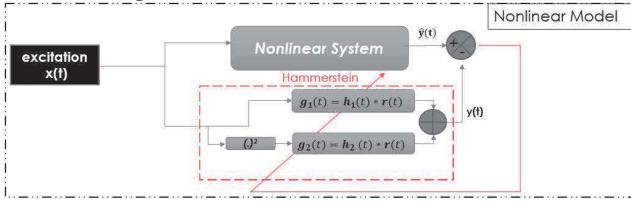
The use of non-linear models in US imaging is mainly motivated by the strong non-linearities that are caused by the interaction between the US waves and some particular tissues. Therefore, several non-linear models can be found in the literature for non-linear US simulation [11] or for the analysis of US images, such as Volterra series [12], neural networks [13] or Hammerstein models [14]. These models depend on unknown parameters or kernels to be estimated from the data. In this paper, we use the non-parametric generalized polynomial Hammerstein model due to its generic structure: this non-linear system can be represented by  $N$  parallel branches, each branch being composed of a non-linear static polynomial element followed by a linear impulse response block. All branches have a common excitation  $\mathbf{x}(t)$ . The relationship between the input  $\mathbf{x}(t)$  and the output  $\mathbf{y}(t)$  is given hereafter

$$\mathbf{y}(t) = \sum_{n=1}^N \mathbf{g}_n(t) * \mathbf{x}^n(t) \quad (1)$$

where  $*$  denotes convolution and  $g_n(t)$  is the impulse response of the  $n^{\text{th}}$  branch. In this work,  $g_n(t)$  includes both the system PSF at the corresponding order  $n$  and the TRF representing the true imaged field topography. In this work, the number of system branches is limited to two, *i.e.*, a linear component resulting from the linear image formation model and a second one associated with the first harmonic resulting from the non-linear interactions in the field. For  $N = 2$  and taking into account the transducer's limited bandwidth (both in reception and emission), the model in (1) can be reformulated as

$$\mathbf{y}(t) = \mathbf{x}(t) * \mathbf{g}_1(t) + \mathbf{x}^2(t) * \mathbf{g}_2(t) + n(t) \quad (2)$$

where  $g_1(t)$  and  $g_2(t)$  are the system impulse response in baseband and around the first harmonic and  $n(t)$  is an additive white Gaussian noise (AWGN). A schematic representation of the model in (2) is shown in Fig. 1.



**Fig. 1.** Schematic representation of the non-linear generalized polynomial Hammerstein model considered in this work.

## 2.2. System identification

This work assumes that the excitation  $\mathbf{x}(t)$  is known such that the identification process only concerns the estimation of  $g_1(t)$  and  $g_2(t)$  from  $\mathbf{y}(t)$  and  $\mathbf{x}(t)$ . The estimation of the model impulse responses in (2) has been already explored in the literature and depends on the nature of the excitation  $\mathbf{x}(t)$  (see, *e.g.*, [15] for a sine wave excitation or [16] for a random excitation). Assuming the model input  $\mathbf{x}(t)$  is known and observing the output signal  $\mathbf{y}(t)$ , the identification approach used herein is based on the convolution between the output of the non-linear system and the inverse filter associated with the input. This work focuses on a swept-sine signal  $\mathbf{x}(t)$ . In this case, the identification process has an analytical solution obtained from the instantaneous frequency step parametrization during one time step [9, 10, 17]. The analytical inverse filter, denoted by  $\hat{\mathbf{x}}(t)$ , is a matched filter, *i.e.*, the time-reversed signal associated with the input  $\mathbf{x}(t)$ . As explained in [9], the convolution between the system output  $\mathbf{y}(t)$  and the estimated input  $\hat{\mathbf{x}}(t)$  allows the excitation  $\mathbf{x}(t)$  to be removed from (2). Indeed, the result of this so-called "non-linear convolution" can be written

$$\mathbf{z}(t) = \mathbf{y}(t) * \hat{\mathbf{x}}(t) = \sum_{n=1}^2 \mathbf{g}_n(t + \Delta t_n) \quad (3)$$

where  $\Delta t_n$  is the temporal gap between the non-linear impulse response (IR) ( $n = 2$ ) and the fundamental IR ( $n = 1$ ). One of the main advantages of this convolution is that it allows the system IRs to be estimated in one path, *i.e.*, the result of the convolution is a set of IRs separated in time with a delay  $\Delta t_n$ . This separation allows the IRs of the two branches  $g_1(t)$  and  $g_2(t)$  to be estimated after time windowing. The two estimated IRs are then used in the next section for estimating the TRF of the US image.

## 3. ULTRASOUND IMAGE DECONVOLUTION

The non-linear model and its identification presented in Section 2 have been included in a US image deconvolution framework allow-

ing the system excitation and the PSF to be separated from the TRF. The proposed TRF restoration method follows two steps. First, the excitation is eliminated through the identification presented in Section 2.2. In a second step, a deconvolution procedure cancelling the effect of the PSF is processed. The proposed deconvolution method combines two data fidelity terms, related to the two branches (fundamental and first harmonic) of the non-linear propagation model. A time windowing operation is used to separate the two system IRs and form the two RF signals. The envelopes of these two signals are computed for each column of the image and are concatenated into two vectors  $\mathbf{g}_1 \in \mathbb{R}^{N \times 1}$  and  $\mathbf{g}_2 \in \mathbb{R}^{N \times 1}$  representing the linear and non-linear image components (where  $N$  is the number of image pixels). The proposed deconvolution method assumes that the two vectors  $\mathbf{g}_1$  and  $\mathbf{g}_2$  are related with the unknown TRF  $\mathbf{r} \in \mathbb{R}^{N \times 1}$  by convolutions with  $\mathbf{h}_1$  and  $\mathbf{h}_2$  associated with the system 2D PSF in baseband (linear propagation) and around the first harmonic (non-linear propagation). The optimization problem related to image deconvolution can then be formulated as the minimization of the sum of three terms: two data fidelity terms associated with  $\mathbf{g}_1$  and  $\mathbf{g}_2$  and an  $\ell_1$ -norm regularization. Note that the  $\ell_1$ -norm is a common choice in order to promote a sparse solution and was used in several applications including US image restoration [18, 19]. We finally have to solve the following problem

$$\min_{\mathbf{r}} \frac{1}{2} \|\mathbf{g}_1 - \mathbf{H}_1 \mathbf{r}\|_2^2 + \frac{1}{2} \|\mathbf{g}_2 - \mathbf{H}_2 \mathbf{r}\|_2^2 + \mu \|\mathbf{r}\|_1 \quad (4)$$

where  $\mathbf{H}_1, \mathbf{H}_2 \in \mathbb{R}^{N \times N}$  are two 2D convolution matrices corresponding to the convolutions between the unknown TRF  $\mathbf{r}$  with  $\mathbf{h}_1$  and  $\mathbf{h}_2$ . In this paper, the classical assumption of cyclic boundary condition for 2D convolution is considered, *i.e.*,  $\mathbf{H}_1$  and  $\mathbf{H}_2$  are block circulant matrices with circulant blocks. We also assume that the TRFs associated with the linear and non-linear components  $\mathbf{g}_1$  and  $\mathbf{g}_2$  are the same as in [20]. To solve (4), we use the alternating direction method of multipliers (ADMM) whose principles can be found in [21, 22]. The ADMM is able to solve the following problem

$$\begin{aligned} \min_{\mathbf{u}, \mathbf{v}} \quad & f_1(\mathbf{u}) + f_2(\mathbf{v}) \\ \text{s.t.} \quad & \mathbf{A}\mathbf{u} + \mathbf{B}\mathbf{v} = \mathbf{c} \end{aligned} \quad (5)$$

where  $f_1$  and  $f_2$  are closed convex functions and  $\mathbf{A}, \mathbf{B}$  and  $\mathbf{u}, \mathbf{v}, \mathbf{c}$  are matrices and vectors of correct sizes. The ADMM is iteratively alternating minimizations over the variables  $\mathbf{u}$  and  $\mathbf{v}$  [22]

$$\begin{aligned} \text{For } k = 0, \dots \\ \left[ \begin{array}{l} \mathbf{u}^{k+1} \in \operatorname{argmin}_{\mathbf{u}} \mathcal{L}_A(\mathbf{u}, \mathbf{v}^{(k)}, \boldsymbol{\lambda}^{(k)}) \\ \mathbf{v}^{k+1} \in \operatorname{argmin}_{\mathbf{v}} \mathcal{L}_A(\mathbf{u}^{(k+1)}, \mathbf{v}, \boldsymbol{\lambda}^{(k)}) \\ \boldsymbol{\lambda}^{(k+1)} = \boldsymbol{\lambda}^{(k)} + \beta(\mathbf{A}\mathbf{u}^{(k+1)} + \mathbf{B}\mathbf{v}^{(k+1)} - \mathbf{c}) \end{array} \right. \end{aligned} \quad (6)$$

where  $\mathcal{L}_A(\mathbf{u}, \mathbf{v}, \boldsymbol{\lambda})$  is the augmented Lagrangian function related to the optimization problem in (5),  $\beta$  is the penalty parameter for the linear constraint, and  $\boldsymbol{\lambda}$  is the Lagrangian multiplier attached to the linear constraints. The following parametrization is chosen in order to transform (4) into (5)

$$\begin{cases} f_1(\mathbf{u}) = \frac{1}{2} \sum_{n=1}^2 \|\mathbf{g}_n - \mathbf{H}_n \mathbf{u}\|_2^2, & f_2(\mathbf{v}) = \mu \|\mathbf{v}\|_1 \\ \mathbf{A} = \mathbf{I}_N, \mathbf{B} = -\mathbf{I}_N, \mathbf{c} = \mathbf{0}_N \end{cases} \quad (7)$$

with  $\mathbf{I}_N$  the identity matrix of size  $N \times N$  and  $\mathbf{0}_N$  a vector containing  $N$  zeroes. The augmented Lagrangian associated with (4) is

$$\mathcal{L}_A(\mathbf{u}, \mathbf{v}, \boldsymbol{\lambda}) = f_1(\mathbf{u}) + f_2(\mathbf{v}) + \frac{\beta}{2} \|\mathbf{u} - \mathbf{v} - \boldsymbol{\lambda}/\beta\|_2^2. \quad (8)$$

The solution of this problem can be iteratively reformulated using the following three steps.

Step 1: Update  $\mathbf{u}$

$$\mathbf{u}^{k+1} \in \operatorname{argmin}_{\mathbf{u}} \frac{1}{2} \sum_{n=1}^2 \|\mathbf{g}_n - \mathbf{H}_n \mathbf{u}\|_2^2 + \frac{\beta}{2} \|\mathbf{u} - \mathbf{v}^k - \boldsymbol{\lambda}^k / \beta\|_2^2. \quad (9)$$

This step admits an analytical solution implemented in practice in the Fourier domain (denoted by  $\mathcal{F}$ ), as shown below

$$\mathbf{u}^{k+1} = \mathcal{F}^{-1} \left\{ \frac{\mathcal{F} \{ \mathbf{H}_1^T \mathbf{g}_1 + \mathbf{H}_2^T \mathbf{g}_2 + \beta \mathbf{v}^k - \boldsymbol{\lambda}^k \}}{\mathcal{F}^* \{ \mathbf{h}_1 \} \mathcal{F} \{ \mathbf{h}_1 \} + \mathcal{F}^* \{ \mathbf{h}_2 \} \mathcal{F} \{ \mathbf{h}_2 \} + \beta \mathbf{I}_N} \right\}. \quad (10)$$

Step 2: Update  $\mathbf{v}$

$$\mathbf{v}^{k+1} \in \operatorname{argmin}_{\mathbf{v}} \mu \|\mathbf{v}\|_1 + \frac{\beta}{2} \|\mathbf{u}^{k+1} - \mathbf{v} - \boldsymbol{\lambda}^k / \beta\|_2^2. \quad (11)$$

The solution for (11) can be obtained using the classical soft-thresholding operator (denoted as  $\operatorname{soft}$  [5]), leading to

$$\mathbf{v}^{k+1} = \operatorname{soft}_{\frac{\mu \|\mathbf{v}\|_1}{\beta}} (\mathbf{v} + \boldsymbol{\lambda}^k / \beta). \quad (12)$$

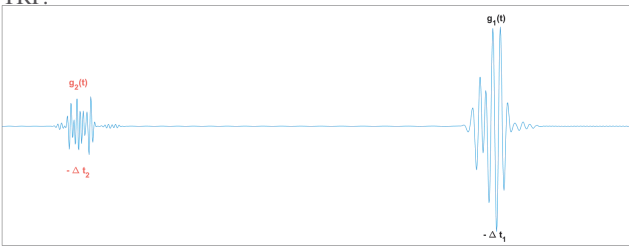
Step 3: Update of the Lagrangian operator  $\boldsymbol{\lambda}$

$$\boldsymbol{\lambda}^{k+1} = \boldsymbol{\lambda}^k + \beta (\mathbf{u}^{k+1} + \mathbf{v}^{k+1}). \quad (13)$$

## 4. SIMULATION RESULTS

### 4.1. System identification

We analyze in this subsection the capability of separating two impulses responses  $\mathbf{g}_1(t)$  and  $\mathbf{g}_2(t)$  following (3). For illustration purpose, we considered an exponential swept-sine excitation, *i.e.*, an exponential chirp with starting frequency  $f_1 = 1$  MHz and stopping frequency  $f_2 = 10$  MHz and two impulse responses  $\mathbf{g}_1(t)$  and  $\mathbf{g}_2(t)$  to be estimated. The output of the non-linear system has been generated by convolution between a column of the TRF image in Fig. 3 (a) and two PSFs in baseband and around the first harmonic. The central frequency was fixed at  $f_0 = 3.5$  MHz. The identification process discussed in Section 2.2 allows the two time-shifted IRs to be estimated in one path, as illustrated in Fig. 2 showing a typical example of non-linear convolution  $z(t)$ . Note that the time shift  $\Delta t_2 - \Delta t_1$  represents a controllable transition parameter from  $f_1$  to  $2f_1$  for the instantaneous frequency of the exponential chirp. After identification of  $\mathbf{g}_1(t)$  and  $\mathbf{g}_2(t)$ , these functions are used to form the cost function  $f_1(u)$  in (7) in order to estimate the unknown TRF.



**Fig. 2.** Example of non-linear convolution  $z(t)$ , where the two IRs  $\mathbf{g}_1(t)$  and  $\mathbf{g}_2(t)$  are well separated (with a separation of  $\Delta t_2 - \Delta t_1$ ) allowing the linear and non-linear components to be estimated.

### 4.2. Image restoration

The proposed non-linear model-based deconvolution method was tested on simulated data with a controlled ground truth TRF. US

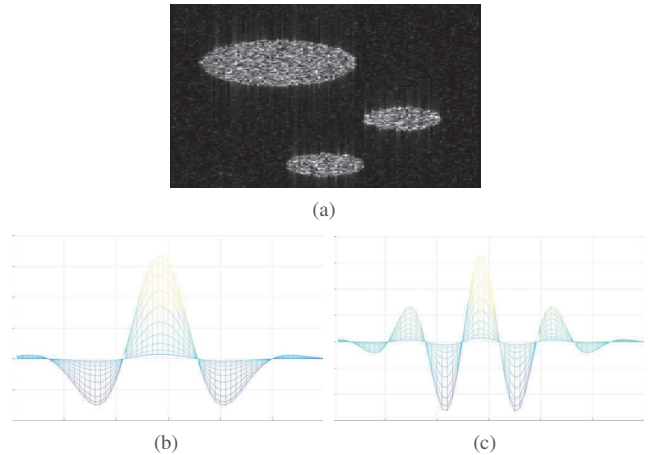
images,  $\mathbf{g}_1$  and  $\mathbf{g}_2$ , were obtained by 2D convolution between two spatially invariant US PSF (in baseband and around the first harmonic, with baseband and harmonic frequencies  $f_0 = 3.5$  MHz and  $2f_0 = 7$  MHz) and the TRF. The TRF corresponds here to a simple medium representing three round hypoechoic inclusions into a homogeneous medium. The pixels located inside and outside the inclusion were generated independently according generalized Gaussian distributions (GGDs) with different parameters. The images were finally contaminated by an AWGN corresponding to a blurred-signal-to-noise ratio of 40 dB. The results obtained using the proposed method are compared to those obtained by a classical linear deconvolution method, having as input a non-linear forward model controlled by a parameter  $\alpha$  that determines the degree of non-linearity of the input. Specifically,  $\alpha$  is defined by

$$\alpha = \frac{\|\mathbf{g}_2\|_2^2}{\|\mathbf{g}_1\|_2^2}. \quad (14)$$

In this case, the optimization problem becomes

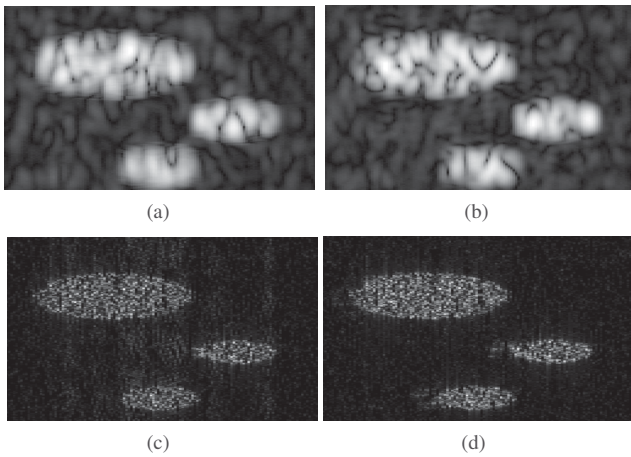
$$\min_{\mathbf{r}} \frac{1}{2} \|\mathbf{s} - \mathbf{H}_1 \mathbf{r}\|_2^2 + \mu \|\mathbf{r}\|_1 \quad (15)$$

where  $\mathbf{s} \in \mathbb{R}^{N \times 1}$  is a weighted sum (depending on  $\alpha$ ) between the linear and the non-linear images  $\mathbf{g}_1$  and  $\mathbf{g}_2$ . The TRF, baseband PSF and first harmonic PSF are shown in Figs. 3 (a, b, c) respectively. Figs. 4 (a, b) show the simulated observed B-mode image (log-compressed envelope image of the corresponding beamformed RF image commonly used for visualization purpose in US imaging) obtained from the TRF convolved with the two PSFs in baseband and around the first harmonic respectively. The estimated TRF using the classical US image restoration approach following (15) is shown in Fig. 4 (c) (for  $\alpha = 0.01$ ). Fig. 4 (d) displays the estimated TRF using the proposed method. The quality of the deconvolution can be appreciated by comparing the estimated TRFs obtained with the two methods. The density of the estimated TRF inside the inclusions is higher with the proposed method. Visual results are complemented by quantitative performance measures, *i.e.*, by the structural similarity (SSIM) [23] and the NRMSE (see Table 1). The proposed method provides estimates of the imaged medium's reflectivity that are closer to the reference image while taking advantage of the blurred data corresponding to the linear and the non-linear branches of the considered model.



**Fig. 3.** (a) TRF phantom, (b, c) US PSF in baseband frequency and around the first harmonic (with baseband frequency  $f_0 = 3.5$  MHz).





**Fig. 4.** (a, b) Blurred image with baseband and first harmonic PSF, (c) Estimated TRF using the classical approach for  $\beta = 0.0012$ ,  $\mu = \sigma^2$  and  $\alpha = 0.01$ , (d) Estimated TRF using the proposed method.

Index	$\alpha$	SSIM(%)	NRMSE(%)
Proposed method	-	97.10	0.68
	0	97.16	0.69
Classical method [19]	0.01	97.05	0.69
	0.05	93.22	0.87
	0.1	83.14	1.38
	0.5	32.55	7.01

**Table 1.** Performance measures for different values of the degree of non-linearity  $\alpha$

## 5. CONCLUSION

In presence of non-linearities affecting a medium of interest, standard linear image processing techniques can have serious limitations leading to image anomalies. The origin of these anomalies is due to the fact that linear models cannot explain the non-linear interactions between the imaged medium and the observed image. This paper showed the interest of considering non-linear models and dedicated image processing strategies to estimate the tissue reflectivity from an ultrasound image. Our results are encouraging and open up new prospects. Future works will be devoted to apply the proposed approach to *in vivo* data taking into consideration that the reflectivity depends on the type of the interaction between the tissues and the beamformed waves. It would be also interesting to consider blind deconvolution methods allowing a spatially variant PSF to be estimated.

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