# Data-Driven Sensor Array Subsampling for Plane-Wave Ultrasound Imaging

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Abstract—Ultrafast plane-wave ultrasound imaging offers very high frame rates but entails large volumes of backscattered data collected by sensor array over multiple plane-wave emissions at different angles. We propose a simple method for reducing the total amount of sampled data (subject to subsequent beamforming and coherent compounding). First, we acquire the zeroangle data in full, and then perform deterministic subsampling of the remaining nonzero-angle data. Our subsampling patterns are angle-specific and derived based on the zero-angle data.

Index Terms—Plane-wave ultrasound imaging, coherent compounding, reduced data acquisition

#### I. INTRODUCTION

Ultrafast plane-wave (PW) ultrasound imaging [1] typically involves acquiring a 3D raw dataset  $RF(t, x, \theta)$  by a transducer array, where  $\theta$  denotes the steering angle of an emitted PW pulse (giving rise to the backscattered signals to be sampled), xrefers to the individual sensor positions within the array, and t represents the sampling time instance. After beamforming and coherent compounding, we get a 2D compounded image dataset D(z, x), where z = ct/2 is the imaging depth, and cis the speed of sound in the insonified medium.

Our objective is to reduce the amount of sensor array data recorded in  $RF(t, x, \theta)$ , whose subsequent processing would still produce acceptable-quality D(z, x). One way to achieve this goal is to use powerful techniques from the compressed sensing theory [2]. Briefly, to obtain an unknown data vector u, we first acquire a smaller-sized vector of its measurements  $\mathbf{v} = \mathbf{A}\mathbf{u} = \mathbf{A}\mathbf{\Phi}\mathbf{w}$ , where (typically)  $\mathbf{A}$  is a sub-Gaussian random matrix, and  $\Phi$  is some fitting orthonormal basis, such that  $\mathbf{u} = \mathbf{\Phi} \mathbf{w}$  with  $\mathbf{w}$  being sparse. We can then recover  $\mathbf{u}$ by solving a suitable constrained optimization problem, e.g.,  $\arg \min \|\mathbf{w}\|_1$  subject to  $\|\mathbf{A} \mathbf{\Phi} \mathbf{w} - \mathbf{v}\|_2 \le \epsilon$  for some  $\epsilon > 0$ . In this work, however, we explore a different approach that does not involve using  $A, \Phi$ , or iterative optimization. Instead, we first acquire zero-angle  $RF(t, x, \theta = 0)$  in full, from which we derive deterministic subsampling patterns for the remaining nonzero-angle raw data. Next, we subsample  $RF(t, x, \theta \neq 0)$ accordingly, then beamform the acquired samples, and finally, compound the resulting beamformed data to produce D(z, x).

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Among various methods for computing D(z, x), we focus on Fourier-domain image reconstruction [3]–[7]. Specifically, we shall make use of the zero-offset depth migration technique from our previous work [7], which is briefly outlined below. In the sequel, the *f*-axis,  $k_x$ -axis, and  $k_z$ -axis data refer to the outputs of the Fourier transform applied to the *t*-axis, *x*-axis, and *z*-axis data, respectively.

For each angle  $\theta$ , we first transform the corresponding 2D raw data  $P_{\theta}(t, x)$  into its spectrum  $F_{\theta}(f, k_x)$ .<sup>1</sup> Next, we remap the *f*-axis data into the  $k_z$ -axis data via linear interpolation, which yields the migrated spectrum  $K_{\theta}(k_z, k_x)$ . Then, we transform the latter into its  $(k_z, x)$ -domain representation (via the inverse Fourier transform along the  $k_x$ -axis), and multiply the result element-wise by  $\exp(j\pi k_z x \tan(\theta))$ , thus producing the phase-shifted spectrum  $S_{\theta}(k_z, x)$ . Finally, we compound all of such  $\theta$ -specific spectra  $S_{\theta}(k_z, x)$  by adding them together (i.e., summing over a given set of  $\theta$  values), and then apply the inverse Fourier transform along the  $k_z$ -axis to obtain D(z, x). The spectral remapping step mentioned above is performed using the following formulas [7]:

$$\mathsf{K}_{\theta}(k_z, k_x) = A(k_z, k_x, \theta) \cdot \mathsf{F}_{\theta}\left(f_{\mathsf{mig}}(k_z, k_x, \theta), k_x\right), \quad (1)$$

$$f_{\rm mig}(k_z, k_x, \theta) = \frac{ck_z}{1 + \cos(\theta)} \left[ 1 + (k_x/k_z)^2 \right], \qquad (2)$$

$$A(k_z, k_x, \theta) = \frac{c}{1 + \cos(\theta)} \left[ 1 - (k_x/k_z)^2 \right].$$
(3)

In the sequel, we shall focus on H(z, x), the complex-valued Hilbert transform of D(z, x) (along the z-axis), which will be our desired output instead of D(z, x).

## II. PROPOSED METHOD

Let  $N_t$ ,  $N_x$ , and  $N_a$  denote the number of sampling time instances, the number of sensors (channels), and the number of PW emission angles, respectively. The full acquisition of  $RF(t, x, \theta)$  would entail collecting as many as  $N_t \times N_x \times N_a$ samples. We want to reduce the total number of acquired raw data samples, while still aiming to produce acceptable-quality compounded images.

<sup>1</sup>Note that  $P_{\theta}(t, x)$  is simply a 2D "slice" of  $RF(t, x, \theta)$  for a particular value of  $\theta$ .

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We begin the acquisition process by fully sampling the zeroangle 2D raw RF data  $P_0(t, x)$ , i.e., our initial cost is  $N_t \times N_x$ samples. Then, for each  $\theta \neq 0$ , we are to acquire some fraction p < 1 of the corresponding 2D raw RF data  $P_{\theta}(t, x)$ . The total cost will become equal to  $N_t \times N_x + p \times (N_t \times N_x) \times (N_a - 1)$ samples, which implies the savings of  $\frac{1+p \times (N_a - 1)}{N_a} \times 100\%$ .

Let  $\mathsf{P}^*_{\theta}(t, x)$  denote a subsampled representation of  $\mathsf{P}_{\theta}(t, x)$ , and let  $\mathsf{M}_{\theta}(t, x)$  be a binary sampling matrix for  $\mathsf{P}_{\theta}(t, x)$ . If we have  $\mathsf{M}_{\theta}(t_l, x_m) = 1$  for some specific location  $(t_l, x_m)$  within our sampling grid, then  $\mathsf{P}^*_{\theta}(t_m, x_l) = \mathsf{P}_{\theta}(t_m, x_l)$ ; otherwise,  $\mathsf{P}^*_{\theta}(t_m, x_l) = 0$ . We express such subsampling compactly as  $\mathsf{P}^*_{\theta}(t, x) = \mathsf{M}_{\theta}(t, x) \odot \mathsf{P}_{\theta}(t, x)$ , where  $\odot$  denotes the elementwise multiplication. The total number of ones in  $\mathsf{M}_{\theta}(t, x)$  must be equal to  $N_p = p \times (N_t \times N_x)$ .

Given multiple  $\theta$ -specific  $\mathsf{P}^*_{\theta}(t, x)$  as inputs, we apply our Fourier-domain reconstruction sequence from [7] to all of them individually, which is then followed by compounding (over all nonzero  $\theta$  values) and the Hilbert transform computations:

$$\begin{split} \mathsf{P}^*_\theta(t,x) &\to \mathsf{F}^*_\theta(f,k_x) \to \mathsf{K}^*_\theta(k_z,k_x) \to \mathsf{S}^*_\theta(k_z,x),\\ &\sum_{\theta \neq 0} \mathsf{S}^*_\theta(k_z,x) \to \mathsf{H}^*(z,x). \end{split}$$

The zero-angle data  $P_0(t, x)$ , acquired in full, yields

$$\mathsf{P}_0(t,x) \to \mathsf{F}_0(f,k_x) \to \mathsf{K}_0(k_z,k_x) \to \mathsf{S}_0(k_z,x) \to \mathsf{H}_0(z,x).$$

Finally, our target output H(z, x) is estimated as follows:

$$\widehat{\mathsf{H}}(z,x) = \mathsf{H}^*(z,x) + \mathsf{H}_0(z,x) + \widetilde{\mathsf{H}}(z,x), \tag{4}$$

where  $\widetilde{H}(z, x)$  serves as an auxiliary "data filler" to mitigate the negative impact of subsampling. In the worst case,  $\widetilde{H}(z, x)$ is simply all zeros. In the best case, it provides an adequate approximation of unknown  $\overline{H}^*(z, x)$  that represents what we would have obtained from the missing data:

$$\overline{\mathsf{P}}_{\theta}^{*}(t,x) \to \overline{\mathsf{F}}_{\theta}^{*}(f,k_{x}) \to \overline{\mathsf{K}}_{\theta}^{*}(k_{z},k_{x}) \to \overline{\mathsf{S}}_{\theta}^{*}(k_{z},x),$$
$$\sum_{\theta \neq 0} \overline{\mathsf{S}}_{\theta}^{*}(k_{z},x) \to \overline{\mathsf{H}}^{*}(z,x),$$

where each "hypothetical"  $\overline{\mathsf{P}}^*_{\theta}(t,x)$  contains the data samples from  $\mathsf{P}_{\theta}(t,x)$  that are not in  $\mathsf{P}^*_{\theta}(t,x)$ . Equivalently,  $\overline{\mathsf{P}}^*_{\theta}(t,x) = \overline{\mathsf{M}}_{\theta}(t,x) \odot \mathsf{P}_{\theta}(t,x)$ , where  $\overline{\mathsf{M}}_{\theta}(t,x)$  is the complemented version of binary  $\mathsf{M}_{\theta}(t,x)$ . Since  $\overline{\mathsf{P}}^*_{\theta}(t,x) = \mathsf{P}_{\theta}(t,x) - \mathsf{P}^*_{\theta}(t,x)$ , we get  $\mathsf{H}(z,x) - \widehat{\mathsf{H}}(z,x) = \overline{\mathsf{H}}^*(z,x) - \widetilde{\mathsf{H}}(z,x)$ , which stresses the importance of  $\widetilde{\mathsf{H}}(z,x)$  being close to unknown  $\overline{\mathsf{H}}^*(z,x)$ .

The main idea of this work pertains to the derivation of the  $\theta$ -specific sampling pattern matrices  $M_{\theta}(t, x)$  from the zeroangle information stored in  $P_0(t, x)$ . The first step is to identify which  $N_p$  data samples in  $P_0(t, x)$  are deemed most useful: for example, we may select those having the largest absolute values. Next, we record their corresponding locations as ones in our initial zero-angle matrix  $M_0(t, x)$ , and then generate its  $(k_z, x)$ -domain representation denoted by  $S_0^M(k_z, x)$ :

$$\mathsf{M}_0(t,x) \to \mathsf{F}_0^\mathsf{M}(f,k_x) \to \mathsf{K}_0^\mathsf{M}(k_z,k_x) \to \mathsf{S}_0^\mathsf{M}(k_z,x).$$
(5)

For any given  $\theta \neq 0$ , the binary sampling matrix  $M_{\theta}(t, x)$ can now be obtained from  $S_0^{\mathsf{M}}(k_z, x)$  as follows. We multiply  $S_0^{\mathsf{M}}(k_z, x)$  element-wise by  $\exp(-j\pi k_z x \tan(\theta))$ , transform it into its  $(k_z, k_x)$ -domain version, and perform demigration (i.e., we remap from  $f_{\mathsf{mig}}$  back to f), thus producing a particular  $\theta$ specific  $(f, k_x)$  spectrum. Then, we apply the inverse Fourier transform to the said spectrum, which yields the desired (t, x)domain matrix  $\widetilde{\mathsf{M}}_{\theta}(t, x) \approx \mathsf{M}_{\theta}(t, x)$ . To get our final  $\mathsf{M}_{\theta}(t, x)$ , we simply binarize  $\widetilde{\mathsf{M}}_{\theta}(t, x)$  via thresholding: for example, the first  $N_p$  largest-magnitude elements of  $\widetilde{\mathsf{M}}_{\theta}(t, x)$  become 1, while the rest are set to 0. Essentially,  $\mathsf{S}_0^{\mathsf{M}}(k_z, x)$  goes through the reconstruction sequence (5) in reverse, except that  $\theta \neq 0$  is now incorporated into the corresponding remapping equations.

## **III. EVALUATION RESULTS**

To evaluate our proposed subsampling method, we have used two experimental datasets from PICMUS-2017 [8], corresponding to  $N_a = 5$  PW emissions (5.208-MHz frequency, 67% bandwidth, 2.5-cycle excitation) at  $-16^{\circ}$ ,  $-8^{\circ}$ ,  $0^{\circ}$ ,  $+8^{\circ}$ , and  $+16^{\circ}$ . These datasets, referred to as TYPE-1 and TYPE-2, were recorded by a 128-element 38.4-mm linear probe, whose sampling frequency was set to 20.832 MHz [8]. For each  $\theta$ , the  $N_t$ -by- $N_x$  size of raw RF data frames was  $1536 \times 128$ . After reconstruction, we generated the B-mode images by log-compressing their respective normalized envelope sections covering the z-axis range [5, 50] mm, as shown in Fig. 1.



Fig. 1. Reference TYPE-1 (left) and TYPE-2 (right) compounded images.

Fig. 2 illustrates the effect of subsampling applied to the 16°-angle TYPE-2 raw RF data frame. The top part of Fig. 2 shows the grayscale image of the absolute values of  $P_{\theta}(t, x)$  elements (100% RF data), while the bottom part shows the corresponding image of  $P_{\theta}^{*}(t, x)$  for p = 0.03 (3% RF data). Note that we were able to capture many large-magnitude samples, whose locations were specified in advance by  $M_{\theta}(t, x)$  derived from  $S_0^{M}(k_z, x)$  for  $\theta = +16^{\circ}$ . Recall that  $S_0^{M}(k_z, x)$  itself is generated from  $M_0(t, x)$ . In this work, we obtain  $M_0(t, x)$  from the initial zero-angle RF data  $P_0(t, x)$  acquired in full: we simply identify the locations of the first  $N_{p=0.03}$  largest-magnitude samples in  $P_0(t, x)$ , and then set the corresponding elements in  $M_0(t, x)$  to 1 (with the others made equal to 0).

Table I provides the quantitative quality indicators for the TYPE-1 and TYPE-2 images under consideration. These metrics have been computed using the original PICMUS-provided



Fig. 2. Subsampling illustration: TYPE-2 RF data,  $\theta = +16^{\circ}$ , p = 0.03.

evaluation routines [8]. The TYPE-1 images are assessed based on the contrast-to-noise ratios (CNR, dB) for the top/bottom cyst phantoms (two anechoic cylinder targets), as well as the axial/lateral full widths at half-maximum (FWHM, mm) for the bottom-right point phantom (a wire target). The TYPE-2 images are assessed based on the average axial/lateral FWHM, computed over all seven point phantoms.

In this work, we report the following cases of subsampling  $N_a - 1 = 4$  nonzero-angle raw RF data frames:

- A) p = 0.01, resulting in 79.2% overall savings;
- B) p = 0.03, resulting in 77.6% overall savings;
- C) p = 0.09, resulting in 72.8% overall savings.

Note that our initial (full) acquisition of the zero-angle data incurs the immediate cost of  $N_t \times N_x$  samples, which amounts to 20% with respect to their total number  $(N_t \times N_x) \times N_a$ . In all cases, we let

$$\widetilde{\mathsf{H}}(z,x) = \frac{\|\mathsf{H}^*(z,x) + \mathsf{H}_0(z,x)\|_2}{\|\mathsf{H}_0(z,x)\|_2} \cdot \overline{\mathsf{H}}_0^*(z,x), \qquad (6)$$

where  $\overline{\mathsf{H}}_0^*(z,x)$  denotes the reconstruction result for the input  $\overline{\mathsf{P}}_0^*(t,x) = \overline{\mathsf{M}}_0(t,x) \odot \mathsf{P}_0(t,x)$  that simulates the "missing" portion of the initial zero-angle raw RF data frame (i.e., as if it underwent  $\mathsf{M}_0(t,x)$ -based subsampling). The multiplication factor in (6) accounts for the energy content difference arising due to compounding.

TABLE I Acquisition Cost and Resulting Image Quality Indicators.

	Acq.	Type-1		Type-2
Case	Cost	Top/Bottom	Axial/Lateral	Axial/Lateral
	(%)	CNR (dB)	FWHM (mm)	FWHM (mm)
5PW	100	8.0/9.5	0.49/0.48	0.47/0.44
1PW	20.0	8.6/6.0	0.48/0.85	0.48/0.80
A	20.8	8.4/6.0	0.46/0.79	0.46/0.68
В	22.4	6.1/6.0	0.46/0.64	0.47/0.60
C	27.2	3.5/6.2	0.47/0.60	0.46/0.55

Fig. 3 shows the images available after the initial acquisition of the zero-angle RF data frame. Acquiring additional four nonzero-angle RF data frames in full would yield the compounded images seen in Fig. 1. Comparing the 1-PW and 5-PW entries in Table I, it is reasonable to anticipate that compounding zero-angle full-input reconstruction data with nonzero-angle partial-input reconstruction data would produce the top and bottom CNR values within [8.0, 8.6] and [6.0, 9.5] dB, respectively; we would also expect the lateral FWHM values to decrease from 0.80-0.85 towards 0.44-0.48 mm.

Table I and Fig. 4, 5, and 6 confirm the expected FWHM improvement (down to 0.55-0.60 mm), but the overall contrast performance is actually much worse than anticipated.<sup>2</sup> This discrepancy can be attributed to the fact that our current approach favors large-magnitude samples, thus aiming to capture signals from strong reflectors, such as wire targets. Formation of the image portions containing weak reflectors and anechoic targets is done mostly by  $H_0(z, x)$  and  $\tilde{H}(z, x)$ . Our current choice of  $\tilde{H}(z, x)$  appears to be inadequate for the TYPE-2 images (in terms of the CNR values obtained), meaning that further investigations into enhanced  $\tilde{H}(z, x)$  are needed.

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<sup>2</sup>We should also mention that in all cases the resulting images did pass the Kolmogorov-Smirnov tests for speckle preservation [8].



Fig. 3. Initial zero-angle images: single PW emission at  $\theta = 0$ .



Fig. 4. Compounded images: nonzero-angle RF data subsampling, p = 0.01 (case A).



Fig. 5. Compounded images: nonzero-angle RF data subsampling, p = 0.03 (case B).



Fig. 6. Compounded images: nonzero-angle RF data subsampling, p = 0.09 (case C).

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