Random noise attenuation using $f$-$x$ regularized nonstationary autoregression$^a$

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**ABSTRACT**

We propose a novel method for random noise attenuation in seismic data by applying regularized nonstationary autoregression (RNA) in frequency-space ($f$-$x$) domain. The method adaptively predicts the signal with spatial changes in dip or amplitude using $f$-$x$ RNA. The key idea is to overcome the assumption of linearity and stationarity of the signal in conventional $f$-$x$ domain prediction technique. The conventional $f$-$x$ domain prediction technique uses short temporal and spatial analysis windows to cope with the nonstationary of the seismic data. The proposed method does not require windowing strategies in spatial direction. We implement the algorithm by iterated scheme using conjugate gradient method. We constrain the coefficients of nonstationary autoregression (NA) to be smooth along space and frequency in $f$-$x$ domain. The shaping regularization in least square inversion controls the smoothness of the coefficients of $f$-$x$ RNA. There are two key parameters in the proposed method: filter length and radius of shaping operator. Synthetic and field data examples demonstrate that, compared with $f$-$x$ domain and time-space ($t$-$x$) domain prediction methods, $f$-$x$ RNA can be more effective in suppressing random noise and preserving the signals, especially for complex geological structure.

**INTRODUCTION**

Random noise attenuation in seismic data can be implemented in the frequency-space ($f$-$x$) and time-space ($t$-$x$) domain using prediction filters (Abma and Claerbout, 1995). Linear prediction filtering assumes that the signal can be described by an autoregressive (AR) model. When the data are contaminated by random noise, the signal is considered to be predicted by the AR filter and the noise is the residual (Bekara and van der Baan, 2009). A number of approaches in $f$-$x$ domain have been proposed and been used for attenuating random noise. The $f$-$x$ prediction technique was introduced by Canales (1984) and further developed by Gulunay (1986). The $f$-$x$ domain prediction technique is also referred as $f$-$x$ deconvolution by Gulunay (1986). Sacchi and Kuehl (2001) utilized the autoregressive-moving average (ARMA) structure of the signal to estimate a prediction error filter (PEF) and the noise sequence is estimated by self-deconvolving the PEF from the filtered data. Hodgson et al. (2002)
presented a novel method of noise attenuation for 3D seismic data, which applies a smoothing filter (e.g. 2D median filter) to each targeted frequency slice and allows targeted filtering of selected parts of the frequency spectrum. The conventional $f$-x domain prediction uses windowing strategies to avoid that the seismic events are not linear. The data are assumed to be piecewise linear and stationary in an analysis temporal and spatial window. To overcome the potentially low performance of $f$-x deconvolution that arises with processing structural complex data, Bekara and van der Baan (2009) proposed a new filtering technique for random and coherent noise attenuation in seismic data by applying empirical mode decomposition (EMD) (Huang et al., 1998) on constant-frequency slices in the $f$-x domain and removing the first intrinsic mode function. In addition, in the research field of seismic data interpolation, Naghizadeh and Sacchi (2009) proposed an adaptive $f$-x prediction filter, which was used to interpolate waveforms that have spatially variant dips. The $f$-x domain prediction technique can be implemented in the frequency slice and also in pyramid domain (Sun and Ronen, 1996). The implemented in pyramid domain makes the operators more efficient because one only needs to estimate one prediction filter from many different frequencies (Sun and Ronen, 1996; Hung et al., 2004; Guitton and Claerbout, 2010).

The prediction process can be also achieved in $t$-x domain (Claerbout, 1992). Abma and Claerbout (1995) discussed $f$-x and $t$-x approaches to predict linear events and concluded that $f$-x prediction is equivalent to $t$-x prediction with a long time length. Crawley et al. (1999) proposed smooth nonstationary PEFs with micropatches and radial smoothing in the application of seismic interpolation, which typically produces better results than the rectangular patching approach. Izquierdo et al. (2006) proposed a technique for structural noise reduction in ultrasonic nondestructive examination using time-varying prediction filter. Sacchi and Naghizadeh (2009) proposed an algorithm to compute time and space variant prediction filters for noise attenuation, which is implemented by a recursive scheme where the filter is continuously adapted to predict the signal.

Fomel (2009) developed a general method of nonstationary regression with shaping regularization (Fomel, 2007). Shaping regularization has an advantage of a fast iterative convergence. Regularized nonstationary regression (RNA) has been used in multiple subtraction Fomel (2009), time-frequency analysis (Liu et al., 2011b), and nonstationary polynomial fitting (Liu et al., 2011a). Liu and Fomel (2010) introduced an adaptive PEFs using RNA in $t$-x domain which has been used for trace interpolation.

In this paper, we investigate the $f$-x domain prediction technique and propose $f$-x domain RNA to attenuate random noise in seismic data. Firstly, we review the theory of $f$-x stationary autoregression. Then, we describe the $f$-x RNA and extend to complex number domain. Next we provide the methodology of random noise attenuation using $f$-x RNA. Finally, we use synthetic and real data examples to evaluate and compare the proposed method with other noise attenuation techniques, such as $f$-x domain and $t$-x domain prediction techniques.
We first consider a seismic section \( S(t, x) \) that consists of a single linear event with the slope \( p \) and constant amplitude. The frequency domain representation of \( S(t, x) \) is given by

\[
S(f, x) = A(f) e^{j2\pi fp},
\]

where \( A(f) \) is the wavelet spectrum, \( f \) is the temporal frequency, and \( x \) is the spatial variable. We assume \( x = n\Delta x \), where \( n = 1, 2, ..., N \), \( N \) is the number of traces in the whole section. The relationship between the \( n \)-th trace and (\( n-1 \))-th trace can be easily shown as

\[
S_n(f) = a_1(f) S_{n-1}(f),
\]

where \( a_1 = \exp(j2\pi fp\Delta x) \). This recursion is a first-order differential equation also known as an AR model of order 1 and represents a single complex-valued harmonic (Bekara and van der Baan, 2009). If there are \( M \) linear events in \( x-t \) domain, we can have a difference equation of order \( M \) (Sacchi and Kuehl, 2001)

\[
S_n(f) = \sum_{i=1}^{M} a_i(f) S_{n-i}(f).
\]

The recursive filter \( \{a_i(f)\} \) can be found for predicting a noise-free superposition of complex harmonics. Considering seismic data with additive random noise and non-causal prediction with order \( 2M \) which includes both forward and backward prediction equations (Spitz, 1991; Naghizadeh and Sacchi, 2009), we can obtain

\[
\varepsilon_n(f) = S_n(f) - \sum_{i=1}^{M} a_i S_{n-i}(f) - \sum_{i=-1}^{M} a_i S_{n-i}(f),
\]

where \( \varepsilon_n(f) \) is a complex noise sequence. Canales (1984) argues a causal estimate of signal \( \sum_{i=1}^{M} a_i(f) S_{n-i}(f) \) is the predictable part of data obtained by an AR model. This operation is usually called \( f-x \) deconvolution (Gulunay, 1986). Noise-free events that are linear in the \( t-x \) domain manifest as a superposition of harmonics in the \( f-x \) domain and these harmonics can be perfectly predicted using AR filter. If seismic events are not linear, or the amplitudes of wavelet are varying from trace to trace, they no longer follow Canales assumptions (Canales, 1984). One needs to perform \( f-x \) deconvolution over a short sliding window in time and space. This leaves the choice of window parameters (window size and length of overlapping between adjacent windows). Bekara and van der Baan (2009) discuss some limitations of conventional \( f-x \) deconvolution in detail.
F-X DOMAIN REGULARIZED NONSTATIONARY AUTOREGRESSION

Nonstationary autoregression has been developed and used in signal processing (Bakrim et al., 1994; AllenAboutajdine et al., 1996; Izquierdo et al., 2006) and seismic data processing (Sacchi and Naghizadeh, 2009). Fomel (2009) developed a general method of nonstationary autoregression using shaping regularization technology and applied it to multiples subtraction. In this paper, we extend the RNA method to f-x domain for complex numbers and apply it to seismic random noise attenuation.

Consider two adjacent seismic traces $S_n(f)$ and $S_{n-1}(f)$ in f-x domain of a seismic section that consists of a single nonlinear event with the slope $p_n$ and varying amplitude $B_n(f)$. Similar to equation 1, we can write

$$
S_n(f) = B_n(f) e^{j2\pi f \Delta x p_n} S_{n-1}(f) = a_1(f) S_{n-1}(f).
$$

(5)

From equation 5 we can find that the coefficients of AR is the function of space index $n$ and frequency index $f$. Therefore, we can use nonstationary autoregression to describe this problem. Equation 5 describes the relation between two traces. If we consider multiple traces, the nonstationary autoregression can be defined as (Fomel, 2009)

$$
\varepsilon_n(f) = S_n(f) - \sum_{i=1}^{M} a_{n,i}(f) S_{n-i}(f),
$$

(6)

where $n$ and $f$ are the coordinate of space and frequency, respectively. If considering the situation of non-causal nonstationary autoregression, we can rewrite the Nonstationary autoregression models (equation 6) as

$$
\varepsilon_n(f) = S_n(f) - \sum_{i=1}^{M} a_{n,i}(f) S_{n-i}(f) - \sum_{i=-1}^{-M} a_{n,i}(f) S_{n-i}(f).
$$

(7)

Equation 7 indicates that one trace noise-free in f-x domain can be estimated by weighted stacking adjacent traces with the weights $a_{n,i}(f)$, which is varying along the space and frequency. Note that the difference between equations 4 and 6 is that the coefficients are varying with space coordinate $n$ in equation 7. To obtain the coefficients $a_{n,i}(f)$ from equation 7, we can transform equation 7 to the following least square problem:

$$
\min_{a_{n,i}(f)} ||S_n(f) - \sum_{i=1}^{M} a_{n,i}(f) S_{n-i}(f) - \sum_{i=-1}^{-M} a_{n,i}(f) S_{n-i}(f)||_2^2,
$$

(8)

where $\|\|_2^2$ denotes the squared L-2 norm. Note that both the data $S_n(f)$ and the coefficients $a_{n,i}(f)$ are in complex numbers domain.

The problem of the minimization in equation 7 is ill-posed because it has more unknown variables than constraints. To obtain the spatial-varying coefficients in nonstationary autoregression, several methods can be employed (AllenAboutajdine et al.,
1996). Some of these are related to the expansion of the spatial-varying coefficients in terms of a given sets of orthogonal basis functions and estimation of the coefficients of the expansion by the least-square method (Izquierdo et al., 2006). Naghizadeh and Sacchi (2009) used exponentially weighted recursive least square (EWRLS) to solve the adaptive problem and applied it to seismic trace interpolation. Fomel (2009) proposed regularized nonstationary autoregression in which shaping regularization technology (Fomel, 2007) is used to constrain the smoothness of the coefficients of nonstationary autoregression. In this paper, we also adopt shaping regularization to solve the under-constrained problem equation 8.

With the addition of a regularization term, equation 8 can be written as

$$\min_{a_{n,i}} \|S_n(f) - \sum_{i=-M,i\neq 0}^{M} a_{n,i}(f)S_{n-i}(f)\|^2_2 + R[a_{n,i}(f)],$$

where \(R\) denotes the shaping regularization operator. Fomel (2009) compared classic Tikhonov’s regularization with shaping regularization in RNA problem. Shaping regularization Fomel (2007) provides a particularly convenient method of enforcing smoothness in iterative optimization schemes. Shaping regularization has clear advantages of a more intuitive selection of regularization parameters and a faster iterative convergence (Fomel, 2009). In the shaping regularization, we assume the initial value for the estimated model \(a_{n,i}(f)\) is zero. If we choose a more appropriate initial value, we can have a fast iterative convergence of the conjugate-gradient iteration in shaping regularization.

Note that the RNA equation in this paper (equation 9) is in complex number domain while the RNA used in multiples subtraction (Fomel, 2009) is in real number domain. Analogous to RNA in real number domain, we force the complex coefficients \(a_{n,i}(f)\) in equation 9 to have a desired behavior, such as smoothness. In shaping regularization technology, we need to choose a shaping operator \(S\) (Fomel, 2007). In this paper, we choose shaping operator \(S\) as Gaussian smoothing with adjustable radius \(r\). Fomel (2007) indicated Gaussian smoothing can be implemented by repeated triangle smoothing operator. Both the data \(S_n(f)\) and the coefficients \(a_{n,i}(f)\) are complex numbers, but the shaping operator \(S\) is real. Therefore, shaping operator \(S\) is operated in real and imaginary parts of the complex coefficients respectively and the L-2 norm in equation 9 is the norm of complex numbers. When using the algorithm of conjugate-gradient iterative inversion with shaping regularization proposed by Fomel (2007) to solve the complex RNA, we only need to replace transpose of real number by conjugate transpose of complex number.

Once we obtain the complex coefficients of RNA, we can achieve an estimation of signal

$$\tilde{S}_n(f) = \sum_{i=-M,i\neq 0}^{M} a_{n,i}(f)S_{n-i}(f).$$

When we use \(f-x\) RNA to noise attenuation, we first select a time window in \(t-x\) domain and transform the data to \(f-x\) domain. The usage of time window is to
guarantee that the data is approximately stationary in time. The $f$-$x$ NAR can deal with spatial nonstationary data in $f$-$x$ domain. Then we use equation 9 with shaping regularization to compute the coefficients $a_{n,i}(f)$ and use equation 10 to estimate the signal in $f$-$x$ domain. Finally, we transform data back to the $t$-$x$ domain and repeat for the next time window.

The computational cost of the proposed method is $O(N_dN_fN_{iter})$, where $N_d$ is data size, $N_f$ is filter size, $N_{iter}$ is the number of conjugate-gradient iterations. If $N_{iter}$ and $N_f$ are small, this is a fast method. In practical implementation, we can choose the range of computed frequency to reduce the computation cost. In addition, if we can simplify the coefficients not to be frequency-dependent, we can apply RNA in frequency slice. In that case the different-frequency slices can be processed in parallel. And the computation cost and memory requirements can be further reduced.

Consider a simple section (Figure 1a), which includes one event, 501 traces. The event is obtained by convolution with Ricker wavelet. Both the dip and amplitude of the event are space-varying. The travel time is a sine function and the amplitude of the Ricker wavelet is multiplied by $B(x) = 0.2(x - 2.5)^2 + 0.5$. This event is obviously nonstationary in space. We add some random noise to it (Figure 1b). The event is greatly contaminated by random noise, especially in the middle part, because the signal of the middle part is poorer than the sideward and the noise levels are the same in the whole section. We use three methods, $f$-$x$ domain prediction, $t$-$x$ domain prediction, and $f$-$x$ domain RNA, to suppress random noise. The $f$-$x$ domain and $t$-$x$ domain prediction methods we used in this paper are discussed by (Abma and Claerbout, 1995). The $f$-$x$ domain prediction is implemented over a sliding window of 20 traces width with 50% overlap and the filter length is 4, $M = 2$ in equation 4 and the $t$-$x$ domain prediction is implemented over the same sliding window and the filter length in space and time are 4 and 5 respectively. The $f$-$x$ RNA is implemented with the parameters: the filter length is 4, $M = 2$ in equation 8; the smoothing radiuses in space and frequency axes of shaping regularization are respectively 20 and 3, $r_x = 20$, $r_f = 3$. Comparing the results of $f$-$x$ RNA (Figure 1f) with other prediction methods (Figures 4c and 4d, we find that the $f$-$x$ RNA is more effective in random noise attenuation for this simple nonstationary data. From the difference sections (Figure 2), we find that all the three methods remove some effective signals. However, the proposed method removes fewest signals than other two methods. Note that the removed signals by $t$-$x$ and $f$-$x$ prediction methods are bigger in the sideward part than middle part, because the filters are the same in a sliding time window while the amplitude and slope of the event are different.

In order to quantitatively evaluate the effect of denoising between different methods, we use signal-to-noise ratio (SNR) to compare these three methods. The SNR is estimated by

$$SNR = 10\log_{10}\left(\frac{\sum_{t,n}[d_n(t)]^2}{\sum_{t,n}[d_n(t) - r_n(t)]^2}\right),$$

where $r_n(t)$ is the noisy section and $d_n(t)$ is the noisy-free section or the section after
Figure 1: (a) Synthetic data. (b) Noisy data (SNR=1.53). (c) Result of $f$-$x$ domain prediction (SNR=2.53). (d) Result of $t$-$x$ domain prediction (SNR=3.12). (e) Result of $f$-$x$ RNA without constraint of smoothness along the frequency axis (SNR=4.87). (f) Result of $f$-$x$ RNA with $r_f = 3$ (SNR=5.06). We decimate the data for display purpose.
Figure 2: The real part of coefficients at a given shift $a_{n,i}(f)$.

Figure 3: Difference sections of $f$-$x$ domain prediction (a), $t$-$x$ domain prediction (b), and $f$-$x$ RNA (c).
noise attenuation. In this simple example, the SNR of the input data without processing is 1.53. After noise attenuation, the SNR of $f$-x RNA is 5.06 dB, and the SNRs of $f$-x domain and $t$-x domain prediction are respectively 2.53 dB and 3.12 dB. The $f$-x RNA can improve SNR more greatly.

To test the sensitivity to the constraint of smoothness along the frequency axis, we give the result of $f$-x RNA without constraint of smoothness along the frequency axis in this simple example (Figure 1e). The result without constraint along frequency axis (Figure 1e) is a little worse than that with constraint along frequency. But both of them are better than the results of conventional $f$-x domain and $t$-x domain prediction. Therefore, to reduce the computation cost, we can simplify the coefficients not to be frequency-dependent when the input seismic data is huge. It is a tradeoff between computation cost and effect of noise attenuation.

Because the dip and amplitude of the event are varying smoothly, $f$-x RNA can predict the signal with smooth coefficients $a_{n,i}(f)$. To specify the coefficients, we display the real parts of the complex coefficients at a given shift $a_{n,i=1}(f)$ (Figure 2). The reason of displaying real parts not imaginary parts is that the signs of real parts of coefficients are the same for forward and backward prediction. We find that the real parts of the complex coefficients are smooth. The smoothing radius controls the smoothness of the coefficients. If we only use one adjacent trace to predict the trace, the coefficients should have the expression

$$a_{n,i=1}(f) = \frac{B_n(f)}{B_{n-1}(f)} e^{j2\pi f \Delta x p_n}. \quad (12)$$

From equation 12 we can note that if the dip and amplitude are smoothly varying, the coefficients are smooth. Therefore, we use Gaussian shaping regularization to constrain the coefficients when solving the least squares equation 9.

**EXAMPLES**

We demonstrate the effectiveness of the proposed $f$-x RNA on a synthetic shot gather and a field poststack dataset.

**Synthetic shot gather**

Figure 4a shows a synthetic shot gather with four hyperbolic events, 501 traces. Some random noise is added to this gather. We do not use windows in time for this example. For $f$-x RNA, the length of filter is $M = 4$ and the smoothing radiuses in space and frequency axes are respectively 20 and 3, $r_x = 20$, $r_f = 3$. The $f$-x domain prediction is implemented over a sliding window of 20 traces width with 50% overlap and the filter length is 6, $M = 3$ and the $t$-x domain prediction is implemented over the same sliding window and the filter length in space and time are 6 and 5 respectively. The estimated nonstationary coefficients by the proposed $f$-x RNA are shown in Figure
Figure 4: (a) Synthetic shot gather. (b) Noisy gather. (c) Result of $f$-$x$ domain prediction (SNR=0.98). (d) Result of $t$-$x$ domain prediction (SNR=1.25). (e) Result of $f$-$x$ RNA (SNR=3.12). (f) The real part of coefficients at a given shift $a_{n,i}(f)$. We decimate the data in (a)-(e) for display purpose.

Figure 5: Difference sections of $f$-$x$ domain prediction (a), $t$-$x$ domain prediction (b), and $f$-$x$ RNA (c).
4f. Note that the middle coefficient is bigger than the sideward, which is because the dip of the middle is smaller than the sideward. The results of three methods are shown in Figures 4d-6d, respectively. The $f$-$x$ RNA achieves a similar result to $f$-$x$ domain and $t$-$x$ domain prediction methods. However, we use equation 11 to compute the SNRs of the results of three methods. The SNRs of three methods are 0.98 dB, 1.25 dB, 1.67 dB, respectively. The $f$-$x$ RNA can improve SNR more greatly. The $f$-$x$ RNA solves the nonstationary case by allowing the coefficients smoothly varying, while $f$-$x$ domain or $t$-$x$ domain prediction method uses windowing strategies. From the difference sections (Figure 7a-5c), we find that $f$-$x$ domain and $t$-$x$ domain prediction methods damage more signals than $f$-$x$ RNA. If we use windows in time for this example, we can obtain better results. This example shows that $f$-$x$ RNA can be used for random noise attenuation in shot gather.

Field poststack dataset

Figure 6: (a) A field marine data set. (b) The result of $f$-$x$ domain prediction. (c) The result of $t$-$x$ domain prediction. (d) The result of $f$-$x$ RNA.

Figure 6a is a seismic image from marine data after time migration. The preprocessing, such as bandpass filtering and migration, has removed some noise. However, some noise still exists in this image (indicated by arrows). This dataset is not structurally too complex and the noise seems random. Therefore, we can use $t$-$x$ domain
Figure 7: Difference sections of $f$-$x$ domain prediction (a), $t$-$x$ domain prediction (b), and $f$-$x$ RNA (c).
Figure 8: Zoomed sections. (a) Original data. (b) The result of $f-x$ domain prediction. (c) The result of $t-x$ domain prediction. (d) The result of $f-x$ RNA.
Figure 9: Comparison on spectra of one trace at 6000 m in field marine data (Figure 6a-6d). (a)-(d) are the amplitude spectra of one trace at 6000 m in Figures 6a-6d, respectively.
or \( f \)-\( x \) domain prediction methods to attenuate the random noise. Figures 6b-6d show the results of random noise attenuation using \( f \)-\( x \) domain prediction, \( t \)-\( x \) domain prediction and \( f \)-\( x \) RNA, respectively. The length of time window is 512 ms in all the three methods. For \( f \)-\( x \) RNA, the filter length is \( M = 4 \), and the smoothing radiuses in space and frequency axes are respectively 20 and 3, \( r_x = 20 \), \( r_f = 3 \). The \( f \)-\( x \) domain prediction is implemented over a sliding window of 20 traces width with 50\% overlap and the filter length is \( M = 4 \) and the \( t \)-\( x \) domain prediction is implemented over the same sliding window and the filter length in space and time are 6 and 5 respectively. The \( f \)-\( x \) domain and \( t \)-\( x \) domain prediction methods removes random noise well in the case that the events are approximately linear. In the area of complex structure, however, both of the \( f \)-\( x \) domain and \( t \)-\( x \) domain prediction methods can not obtain a good result. Compared to \( f \)-\( x \) domain and \( t \)-\( x \) domain prediction methods, \( f \)-\( x \) RNA removes more noise and preserves signals (Figure 7a-7c). Note that \( f \)-\( x \) domain and \( t \)-\( x \) domain prediction methods remove some signals, especially for complex structure (indicated by arrows in Figures 7a and 7b). We display the zoomed section in Figure 8a-8d. The zoomed sections show the proposed method is more effective than other methods, especially in the area of complex structure indicated by arrow. The \( f \)-\( x \) RNA gives a good result not only for linear events but also for curving events (indicated by arrows in Figure 8a-8d). From the comparison on the spectra of a trace randomly chosen as shown in Figure 9a-9d, we can see that \( f \)-\( x \) domain and \( t \)-\( x \) domain prediction methods greatly attenuate frequency components in [10,30] Hz, which includes effective signals. Thus, the difference sections of \( f \)-\( x \) domain and \( t \)-\( x \) domain prediction methods (Figures 7a and 7b) include more signals than \( f \)-\( x \) RNA (Figure 7c). For high frequency random noise, all the three methods can achieve a similar result (Figure 9a-9d).

**CONCLUSIONS**

We have proposed a novel method for random noise attenuation using \( f \)-\( x \) domain regularized nonstationary autoregression. \( f \)-\( x \) RNA uses shaping regularization to constrain the complex nonstationary coefficients to be smooth along space and frequency axes. Contrary to conventional noise-reduction technology, \( f \)-\( x \) domain and \( t \)-\( x \) domain prediction, \( f \)-\( x \) RNA invokes no piecewise-stationary assumption. The parameters used in \( f \)-\( x \) RNA are intuitive because the parameters directly control the smoothness of complex coefficients. The proposed method has two key parameters: filter length and smoothing radius of shaping operator. Filter length is related to the number of events and smoothing radius is related to the smoothness of desired RNA complex coefficients. As the smoothing radius increases, the result of RNA approaches the result of stationary autoregression. This approach does not require breaking the input data into local windows along space axis, although it is conceptually analogous to sliding spatial windows with maximum overlap. Both synthetic and field data examples confirm that the proposed approach can be significantly more effective than other noise-reduction methods in improving signal-to-noise ratio and preserving the signals. A comparison with the recently published \( t \)-\( x \) RNA method
Noise attenuation using $f$-$x$ RNA has not been attempted, but remains of interest for further investigation. The proposed method is easy to extend to the 3D case ($f$-$x$-$y$ domain). One only needs to add a space dimension in the equation 9 when applied in 3D case. Besides random noise attenuation, $f$-$x$ RNA may have other applications in seismic data processing, such as seismic trace interpolation.

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