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SPECIALTY SECTION

This article was submitted to
Ocean Observation,
a section of the journal
Frontiers in Marine Science

RECEIVED 21 September 2022

ACCEPTED 23 November 2022

PUBLISHED 16 December 2022

CITATION

Huang P and Yang P (2022)
Synthetic aperture imagery for
high-resolution imaging sonar.
Front. Mar. Sci. 9:1049761.
doi: 10.3389/fmars.2022.1049761

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Synthetic aperture imagery for high-resolution imaging sonar

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Synthetic aperture sonar (SAS) can provide high-resolution underwater images. Traditional fast imaging algorithms designed for multi-receiver synthetic aperture sonar (MSAS) are complex because the point target reference spectrum (PTRS) deduction and imaging algorithm development are complicated. This paper proposes an imaging algorithm for the MSAS system to solve this issue. The proposed method first approximates the two-round slant range based on the phase center approximation method. The PTRS, including the quasi-monostatic and bistatic deformation terms, can be easily deduced. After compensating for the bistatic deformation term based on the interpolation and complex multiplication with the preprocessing step, the MSAS imagery can be simplified to the focus of the traditional monostatic SAS. Therefore, the conventional imaging algorithms designed for traditional monostatic SAS can be used directly. The proposed method providing high-resolution imaging results is more efficient than the traditional methods.

KEYWORDS

synthetic aperture sonar, high-resolution, imaging algorithm, phase center approximation, imaging performance

1 Introduction

The complex ocean environment (Hunter, 2006; Davis et al., 2009; Xia et al., 2016; Zhang et al., 2020) makes its observation challenging. Therefore, underwater imaging is essential for scientists and engineers working in ocean engineering. Currently, the high-resolution imaging equipment named synthetic aperture sonar (SAS) (Gough and Hawkins, 1996; Gough and Hawkins, 1997; Gough and Hawkins, 1998; Zhang and Ying, 2022; Dillon and Steele, 2022; Yang and Liu, 2022) can generate high-resolution images. Compared with traditional imaging sonar, such as side scan and multibeam sonar (Cutrona, 1975), SAS can provide high resolution in the along-track dimension independent of the frequency and detection range. Therefore, the SAS image can be used for seabed mapping (Barclay et al., 2005; Tan et al., 2019) and target detection (Dobeck, 1999; Rankin et al., 1999; Cong et al., 2000; Gough et al., 2006), such as searching for

shipwrecks, airplane wrecks (LeHardy and Larsen, 2015), and pipelines (Sæbø et al., 2010).

Synthetic aperture image formation is a crucial technique for the SAS system. The monostatic SAS transmitter is still used as the receiver. The point target reference spectrum (PTRS) can be easily deduced using the stationary phase method. Based on the PTRS of the monostatic SAS, fast imaging algorithms such as the range-Doppler (RD) algorithm (Zhong and Liu, 2006; Neo et al., 2008; Wu et al., 2014; Zhang et al., 2017b), chirp scaling (CS) algorithm (Zhong and Liu, 2009; Zhong and Liu, 2010; Zhang et al., 2013), and range migration algorithm (RMA) (Liu et al., 2009; Shin and Lim, 2009; Zhang et al., 2021c) can be easily designed. For pulsed sonar, a longer detection range in the cross-track dimension requires a larger pulse repetition interval (PRI). However, higher resolution in the along-track dimension requires a lower PRI. Therefore, there is a significant contradiction between the cross-track dimension mapping swath and the along-track dimension resolution.

A multi-receiver synthetic aperture sonar (MSAS) (Rolt and Schmidt, 1992; Gough and Miller, 2003; Zhang et al., 2014a; Zhang et al., 2021c), which includes a single transmitter and many receivers in the along-track dimension, is proposed to solve this problem. MSAS can provide a wider mapping swath at a given resolution than the monostatic SAS system. Unfortunately, the use of the MSAS system involves two major challenges. Firstly, the two-round slant range of MSAS is characterized by double square root terms, which makes it difficult to deduce the PTRS of MSAS based on the stationary phase method (Wang et al., 2008; Wang et al., 2011). Secondly, all of the imaging algorithms have been designed for the monostatic SAS system and cannot be directly used for MSAS systems. To deduce the PTRS based on the stationary phase method, several approximations, including the Taylor approximation of the two-round slant range and Loffeld's bistatic formula (LBF) (Loffeld et al., 2004; Zhang et al., 2021b), are used.

In addition, the imaging algorithms used for monostatic SAS are further extended to the MSAS system. In general, these extensions can be grouped into two classes. The first involves developing a new algorithm according to the RD, CS, and RMA. This operation often employs the series approximation of PTRS, which is exceedingly tedious. The other involves directly using the monostatic SAS system imaging algorithms after converting the MSAS data into a monostatic SAS-equivalent signal. The second approach is much easier to achieve when comparing the advantages and disadvantages of both methods. The PTRS of MSAS must be decomposed into a quasi-monostatic term and a bistatic deformation term in order to directly use the imaging algorithms designed for monostatic SAS. After compensating for the bistatic deformation term, traditional imaging algorithms can be used directly.

Based on the assumption that the transmitter and the receiver contribute equally to the Doppler phase in the along-

track dimension, Loffeld et al. (2004) developed a PTRS (often called LBF) that can be decomposed into a quasi-monostatic term and a bistatic deformation term. The bistatic deformation term of LBF depends significantly on the range, instantaneous frequency, and Doppler frequency. The sub-block processing method (Zhang et al., 2019) in the cross-track dimension is usually exploited to cancel this term; therefore, the sub-block width determines the imaging performance. In other words, a wider sub-block width reduces performance, while a narrower width increases performance. Unfortunately, the use of a narrow sub-block width is time-consuming.

In general, there is a significant contradiction between imaging performance and efficiency when the LBF is used. The phase center approximation (PCA) method (Gough et al., 2000; Gough and Hayes, 2005) supposes an element located at the midpoint between the transmitter and the receiver. This element is equivalent to a monostatic transducer used for signal transmission and reception. However, this operation results in an approximation error, called the PCA error (Zhang et al., 2021a). Traditional methods do not compensate for this approximation error, and image distortion is introduced at a close range. Some methods neglect the stop-and-hop error (Zhang et al., 2017a; Zhang et al., 2018), seriously affecting long-range SAS imaging results.

This paper presents an imaging method based on the PCA method. Firstly, the two-round slant range was approximated by the second-order Taylor approximation and inverse Taylor approximation. Then, the slant range equivalent to the traditional monostatic SAS was obtained. Because the PTRS of the monostatic SAS can be easily obtained, the stationary phase method was exploited to deduce the PTRS of MSAS, including the quasi-monostatic and bistatic deformation terms. After compensating for the bistatic deformation term, the MSAS data were converted into monostatic SAS-equivalent data, considered the input to the RD algorithm in this paper. Lastly, simulations were performed to validate the proposed method. The remainder of this paper is organized as follows. *Section 2* discusses the imaging geometry. *Section 3* describes the proposed method. In *Section 4*, the simulations are described in detail. Finally, the last section summarizes the conclusions of this study.

2 MSAS imaging geometry

Figure 1 shows the MSAS imaging geometry. The sonar system is towed in the along-track direction, and the horizontal axis is in the cross-track direction. The fast time is denoted by τ and the slow time by t . An ideal target was assumed in this imaging scenario. The coordinate in the cross-track dimension is r , while that in the along-track dimension is 0. The transmitter and receiver arrays were deployed in the along-track dimension, where the black rectangle is the transmitter and the remaining

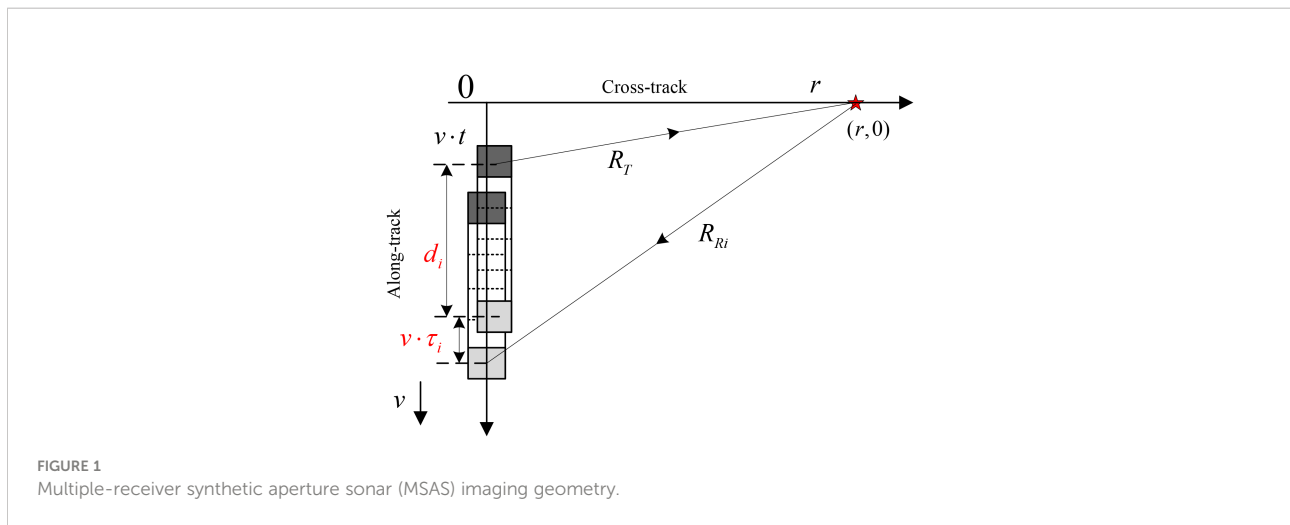


FIGURE 1 Multiple-receiver synthetic aperture sonar (MSAS) imaging geometry.

rectangles are the receivers, which were uniformly spaced at equal distances. The distance between the transmitter and the i th receiver is d_i , where $i (i \in [1, M])$ is the receiver index. M is the total number of receivers.

Based on Figure 1, the two-round slant range between the transmitter to the target and back to the i th receiver is

$$R_i(t; r) = R_T(t) + R_{Ri}(t) = \sqrt{r^2 + (vt)^2} + \sqrt{r^2 + (vt + d_i + 2v \cdot r/c)^2}. \quad (1)$$

In Equation 1, $R_T(t) = \sqrt{r^2 + (vt)^2}$ is the distance between the transmitter and the target. The distance between the target and the i th receiver is denoted by $R_{Ri}(t) = \sqrt{r^2 + (vt + d_i + 2v \cdot r/c)^2}$, where v is the velocity of the sonar platform and c is the sound velocity in water. Traditional methods neglect the stop-and-hop approximation (Zhang et al., 2017a) on the SAS system because traditional SAS systems work with low resolution and narrow mapping swath. A lot of high-resolution SAS systems with wide mapping swaths are currently being developed. If the stop-and-hop error is neglected, distortions such as higher side lobes and along-track shifting are introduced into the SAS image. Compared with the traditional signal model, Equation 1 considers the influence of the stop-and-hop approximation (Zhang et al., 2017a) on the SAS system.

The SAS system often uses a chirp signal. The received signal of the i th receiver is given by

$$ss_i(\tau, t) = p\left(\tau - \frac{R_i(t; r)}{c}\right) \omega_a(t) \cdot \exp\left\{-j2\pi f_c \frac{R_i(t; r)}{c}\right\} \quad (2)$$

In Equation 2, the transmitted chirp is denoted by $p(\tau)$. The beam pattern of the SAS system is denoted by $\omega_a(t)$. Because $\omega_a(t)$ does not affect the SAS focusing, it is suppressed in the following discussions. The carrier frequency is f_c . Because the

two-round slant range in Equation 1 has two square root terms, it is challenging to deduce the PTRS of MSAS using the stationary phase method.

3 MSAS imaging algorithm

3.1 MSAS PTRS

The PTRS (Wu et al., 2011) should be obtained first to develop the MSAS imaging algorithm. Based on Equation 1, the second-order Taylor expansion is used, and the result is

$$R_i(t; r) \approx \sqrt{r^2 + (vt)^2} + \sqrt{r^2 + (vt + d_i + v \frac{2r}{c})^2} = 2r + \frac{v^2}{r} \left[t + \left(\frac{r}{c} + \frac{d_i}{2v} \right) \right]^2 + \frac{1}{4r} (v \frac{2r}{c} + d_i)^2. \quad (3)$$

With the inverse Taylor expansion, Equation 3 can be further reformulated as

$$R_i(t; r) \approx 2r + \frac{v^2}{r} \left[t + \left(\frac{r}{c} + \frac{d_i}{2v} \right) \right]^2 + \frac{1}{4r} (v \frac{2r}{c} + d_i)^2 \approx 2\sqrt{r^2 + (vt + v \frac{r}{c} + \frac{d_i}{2})^2} + \frac{(v \frac{2r}{c} + d_i)^2}{4r} \quad (4)$$

The square root term in Equation 4 is similar to the slant range of the monostatic SAS system. The second term in Equation 4 denotes the error introduced by PCA and the stop-and-hop approximation. With the second term in Equation 4 compensated by the preprocessing, the following work focuses on the MSAS data using the traditional imaging algorithms of a monostatic SAS system.

The current work deduces the PTRS. Starting with Fourier transformation (FT) in the cross-track dimension for the echo signal (Equation 2),

$$SS_i(f_\tau, t) = P(f_\tau) \exp \left\{ -j2\pi(f_c + f_\tau) \frac{R_i(t; r)}{c} \right\} \quad (5)$$

In Equation 5, $P(f_\tau)$ is the spectrum of the transmitted chirp signal. The instantaneous frequency is denoted by f_τ . Performing an FT in the along-track dimension for Equation 5,

$$SS_i(f_\tau, f_t) = P(f_\tau) \cdot \int_{-T_s/2}^{T_s/2} \exp \left\{ -j2\pi(f_c + f_\tau) \frac{R_i(t; r)}{c} - j2\pi f_t t \right\} dt \quad (6)$$

In Equation 6, the Doppler frequency is denoted by f_t , and the synthetic aperture time is represented by T_s . The exponential phase in Equation 6 is further denoted by

$$\varphi_i(f_\tau, f_t; r) = -2\pi(f_c + f_\tau) \frac{R_i(t; r)}{c} - 2\pi f_t t. \quad (7)$$

Performing the stationary phase method (Wang et al., 2009) on Equation 7 yields

$$\frac{\partial R_i(t_i; r)}{\partial t} + \frac{c f_t}{f_c + f_\tau} = 0. \quad (8)$$

In Equation 8, the stationary point is denoted by t_i . Based on Equation 4, Equation 8 is then reformulated as

$$\frac{2v \left(vt_i + v \frac{r}{c} + \frac{d_i}{2} \right)}{\sqrt{r^2 + \left(vt_i + v \frac{r}{c} + \frac{d_i}{2} \right)^2}} + \frac{c f_t}{f_c + f_\tau} = 0. \quad (9)$$

The solution to Equation 8 is expressed as

$$t_i = -\frac{c f_t}{v(f_c + f_\tau)} \cdot \frac{1}{\sqrt{4v^2 - \left[\frac{c f_t}{(f_c + f_\tau)} \right]^2}} - \frac{r}{c} - \frac{d_i}{2v} \quad (10)$$

Substituting Equation 10 into Equation 7, we obtain the PTRS phase given by

$$\begin{aligned} \varphi_i(f_\tau, f_t; r) &= -2\pi(f_c + f_\tau) \frac{R_i(t_i; r)}{c} - 2\pi f_t t_i \\ &= -2\pi(f_c + f_\tau) \frac{2\sqrt{r^2 + \left(vt_i + v \frac{r}{c} + \frac{d_i}{2} \right)^2} + \frac{(v \frac{r}{c} + \frac{d_i}{2})^2}}{c} - 2\pi f_t t_i \\ &= -4\pi \frac{f_c}{c} r \sqrt{\left(1 + \frac{f_\tau}{f_c} \right)^2 - \frac{c^2 f_\tau^2}{4v^2 f_c^2}} + \pi f_t \frac{d_i}{v} + 2\pi f_t \frac{r}{c} - \pi(f_c + f_\tau) \frac{(2 \frac{v}{c} r + d_i)^2}{2rc} \end{aligned} \quad (11)$$

Inspection of Equation 11 reveals that the first term is similar to the PTRS of traditional monostatic SAS. The spatial sampling of each receiver causes the second term. The third term denotes the azimuthal shifting caused by the stop-and-hop approximation. The last term is the Doppler error and micro-range migration caused by the PCA and stop-and-hop errors. Based on the idea of the monostatic conversion-based method, the phase error related to the displaced distance, d_i , is the bistatic deformation term, which is

$$\begin{aligned} \Psi_i(f_\tau, f_t; r; d_i) &= -\pi f_c \frac{\left(2 \frac{v}{c} r + d_i \right)^2}{2rc} - \pi f_\tau \frac{\left(2 \frac{v}{c} r + d_i \right)^2}{2rc} \\ &\quad + \pi f_t \frac{d_i}{v} \end{aligned} \quad (12)$$

The remaining phase denotes the quasi-monostatic term given by

$$\Phi(f_\tau, f_t; r) = -4\pi \frac{f_c}{c} r \sqrt{\left(1 + \frac{f_\tau}{f_c} \right)^2 - \frac{c^2 f_\tau^2}{4v^2 f_c^2}} + 2\pi f_t \frac{r}{c} \quad (13)$$

Subsequently, Equation 11 can be further reformulated as

$$\varphi_i(f_\tau, f_t; r) = \Phi(f_\tau, f_t; r) + \Psi_i(f_\tau, f_t; r; d_i) \quad (14)$$

It is apparent from Equation 14 that the MSAS imagery can be converted into monostatic SAS focusing after compensating for the bistatic deformation term, $\Psi_i(f_\tau, f_t; r; d_i)$.

3.2 MSAS imagery

The MSAS imagery includes two primary operations. The first operation compensates for the bistatic deformation term, $\Psi_i(f_\tau, f_t; r; d_i)$, and is called preprocessing. The second operation conducts the MSAS data focusing based on the quasi-monostatic term.

The compensation of the bistatic deformation term is now discussed. The first step is the compression in the cross-track dimension. The matched filtering function is written as

$$H_{\text{cross}} = P^*(f_\tau) \quad (15)$$

In Equation 15, the superscript asterisk denotes the complex conjugate operation.

The second step is the compensation of the Doppler phase, the first term in Equation 12, in the two-dimensional (2D) time domain. The filtering function is expressed as

$$\Gamma_i(\tau, d_i) = \exp \left\{ \pi f_c \frac{\left(2 \frac{v}{c} r + d_i \right)^2}{2rc} \right\} \quad (16)$$

The second step corrects the micro-range migration, the second term in Equation 12, which is caused by the PCA and stop-and-hop errors. From Equation 12, the micro-range migration in the 2D time domain corresponding to the second term in the 2D frequency domain is given by

$$\Delta_i(\tau, d_i) = \frac{\left(2 \frac{v}{c} r + d_i \right)^2}{4rc} \quad (17)$$

The term shown in Equation 17 is range-variant. Interpolation is exploited to correct the micro-range migration.

The third step of the preprocessing operation compensates for the spatial sampling error, the last term in Equation 12. This can be achieved by a complex multiplication in the cross-track

time, along-track Doppler domain. The filtering function is expressed as

$$\Lambda_i(\tau, d_i) = \exp\left\{-\pi f_t \frac{d_i}{v}\right\} \quad (18)$$

The last step is the recovery of the chirp characteristic. Filtering is conducted in the cross-track frequency domain, and the filtering function directly uses the spectrum of the transmitted signal.

A block diagram of the preprocessing operation is depicted in Figure 2 based on the steps discussed above.

After conducting the preprocessing operations, the remaining work involves MSAS focusing using the RD algorithm based on the quasi-monostatic term in Equation 13. The use of the RD algorithm is enabled by approximating the first term in Equation 13 by the second-order Taylor expansion. After the series approximation, Equation 13 is reformulated as

$$\begin{aligned} \Phi(f_r, f_t; r) \approx & -\frac{4\pi r \beta}{\lambda} - \frac{4\pi r}{c\beta} f_r + 2\pi \frac{r\lambda}{c^2} \left(\frac{1}{\beta^3} - \frac{1}{\beta}\right) f_r^2 \\ & + 2\pi f_t \frac{r}{c} \end{aligned} \quad (19)$$

Where

$$\beta = \sqrt{1 - \frac{c^2 f_t^2}{4v^2 f_c^2}} \quad (20)$$

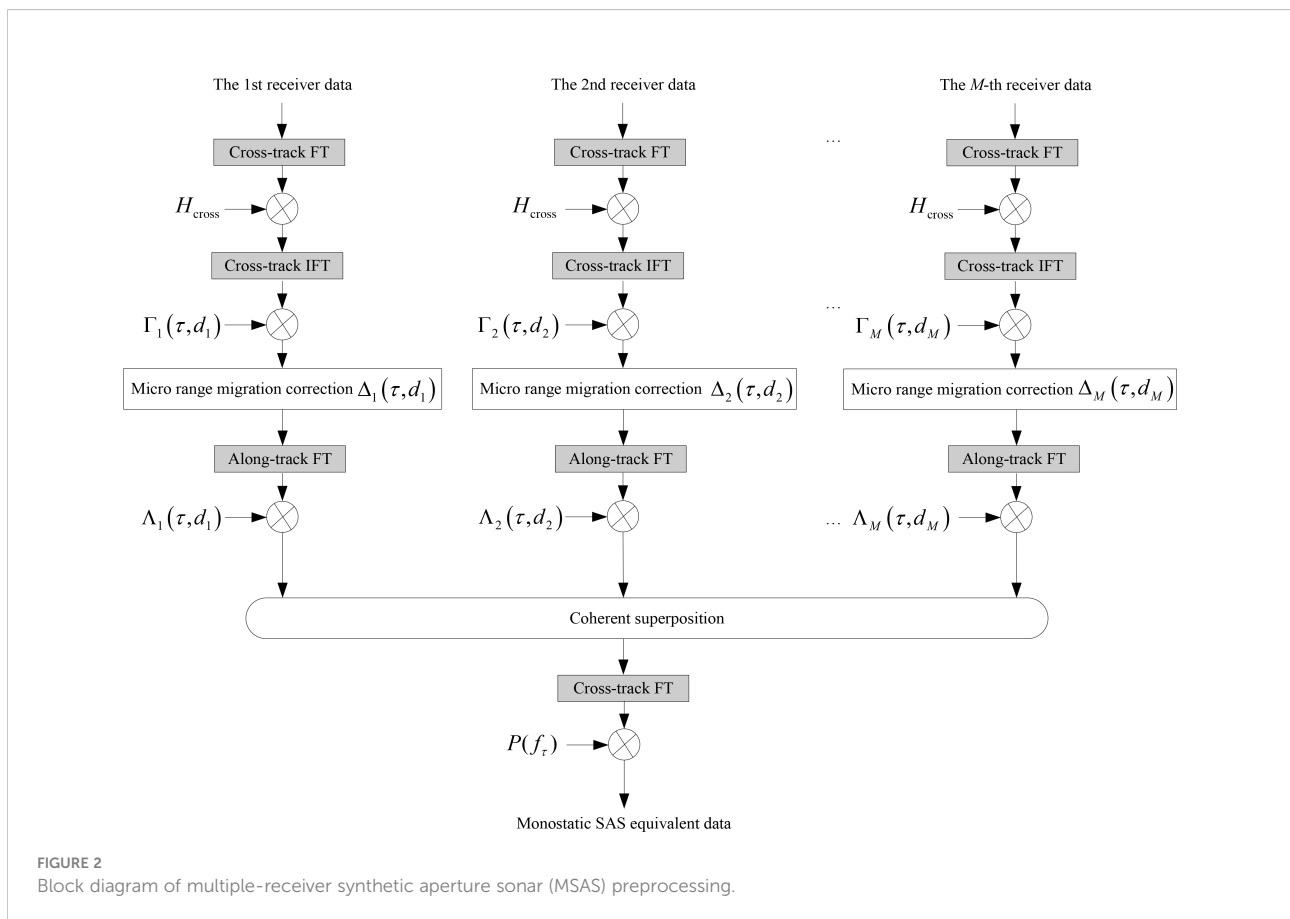
The first term in Equation 19 denotes the along-track modulation, the second term is the range migration, and the third term denotes the cross-track modulation. The last term is the along-track shifting.

The first step in the RD algorithm requires performing the cross-track compression in the 2D frequency domain based on the third term in Equation 19. Because the third term in Equation 19 is slightly range-variant, this term can be compensated using the sub-block processing method (Zhang et al., 2019). For the n th sub-block, the filtering function is expressed as

$$H_n(f_r, f_t; r_n) = \exp\left\{j\pi \frac{f_r^2}{\gamma_e(f_t; r_n)}\right\} \quad (21)$$

Where

$$\gamma_e(f_t; r_n) = \frac{1}{\gamma} - 2\lambda r_n \frac{1 - \beta^2}{c^2 \beta^3} \quad (22)$$



In Equation 21, the subscript n ($n \in [1, N]$) is the sub-block index and N is the total number of sub-blocks. In Equation 22, γ is the chirp rate of the transmitted signal and r_n is the center range of the n^{th} sub-block. In practice, Equation 21 is also called the secondary range compression (SRC), which is range-variant. To some degree, the phase error within each sub-block width should not exceed $\pi/16$, and this can be expressed as $|\gamma_c(f_i; r_n) - \gamma_c(f_i; r_n \pm 0.5\Delta r)| < \pi/16$, where Δr is the sub-block width. The sub-block width can be easily determined with this method. In general, the SRC depends on the range, to some extent. The data can now be coarsely segmented into several sub-blocks in the cross-track dimension.

The second term in Equation 19 denotes the range migration, which can be corrected by interpolation in the range-Doppler domain. Based on the second term in Equation 19, the range migration in the cross-track time, along-track Doppler domain is denoted by

$$\Delta R(f_i; r) = \frac{2r}{\sqrt{1 - \left(\frac{\lambda f_i}{2v}\right)^2}} - 2r \quad (23)$$

After this step, the coupling between the cross-track and along-track dimensions can be decoupled. Subsequently, the along-track compression can be performed based on the first term in Equation 19. The matched filtering function in the along-track dimension is denoted by

$$H_{\text{along}}(f_i; r) = \exp\left\{j4\pi \frac{\beta}{\lambda} r\right\} \quad (24)$$

The last step involves correcting the along-track shifting based on the last term in Equation 19. The filtering function is given by

$$H_{\text{shifting}}(f_i; r) = \exp\left\{-j2\pi f_i \frac{r}{c}\right\} \quad (25)$$

Correction of the along-track shifting can also be carried out with the along-track compression. A block diagram based on the RD algorithm discussed above is shown in Figure 3.

4 Simulation results

The simulations presented in this section validate the proposed method. The MSAS parameters are listed in Table 1, which indicates that there are 32 receivers.

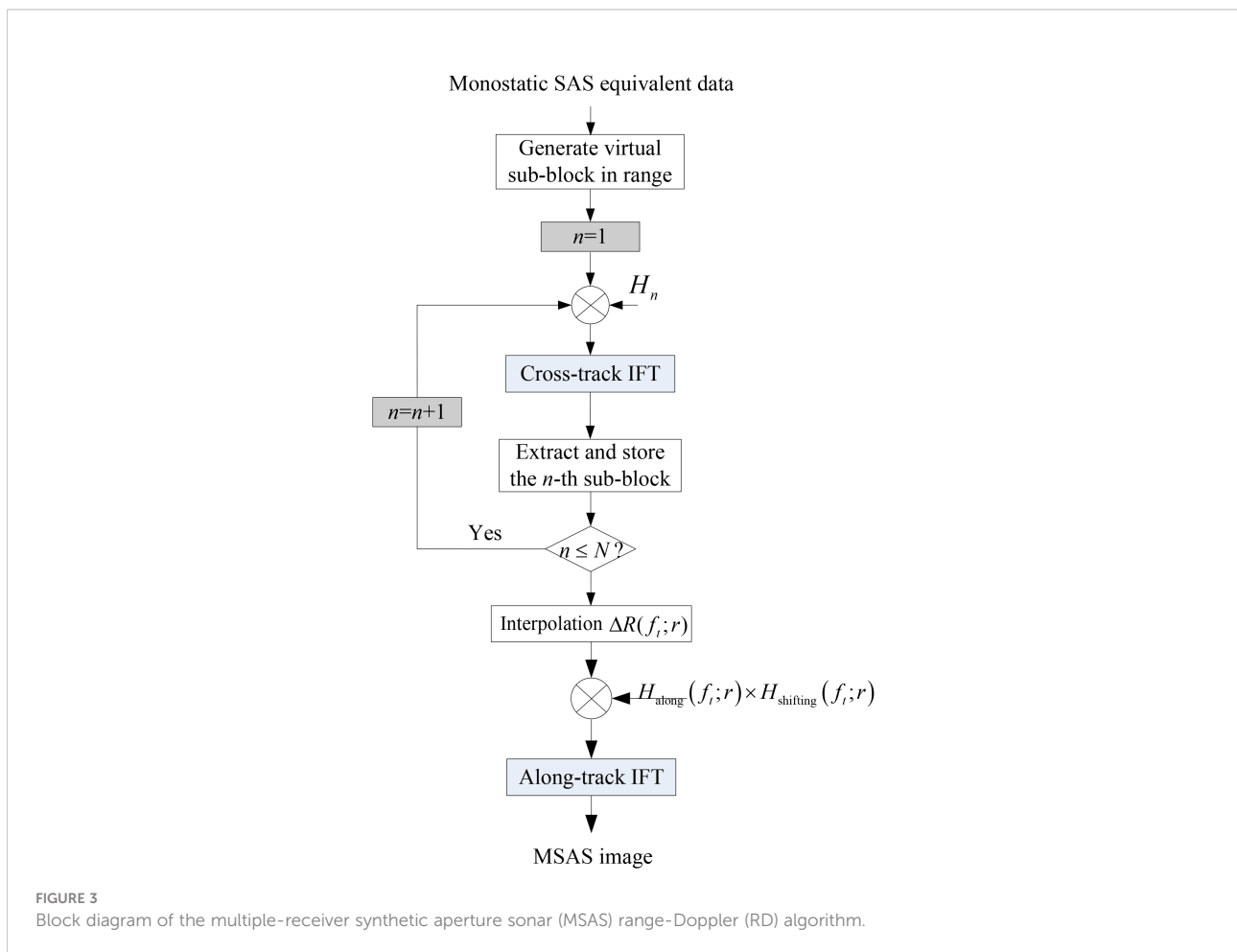


FIGURE 3 Block diagram of the multiple-receiver synthetic aperture sonar (MSAS) range-Doppler (RD) algorithm.

TABLE 1 Synthetic aperture sonar (SAS) parameters.

Parameter	Value	Unit
Platform velocity	2	m/s
Pulse repetition interval	0.32	s
Signal bandwidth	20	kHz
Carrier frequency	150	kHz
Receiver array length	1.28	m
Receiver width	0.04	m
Transmitter width	0.08	m

There are six ideal point targets in the imaging scenario depicted in Figure 4. These targets are numbered T1, T2, ..., and T6 to simplify the description.

The back-projection (BP) algorithm (Zhang et al., 2014b; Wang et al., 2015) can provide high-performance results based on the interpolation theory and is therefore used as the benchmark. After focusing the simulated MSAS data, the imaging results from the proposed method are shown in Figures 5A, B shows the results from the BP algorithm. Comparing the imaging results of the proposed method to those of the BP algorithm (Zhang et al., 2014b), it can be seen that the performance of the proposed method is very similar to that of the BP algorithm, indicating that the proposed method can accurately reconstruct the targets.

The along-track slices are shown in Figure 6. It is readily apparent that the along-track slices of the present method agree well with the slices from the BP algorithm. These results further confirm our conclusion that the present method can accurately recover the targets.

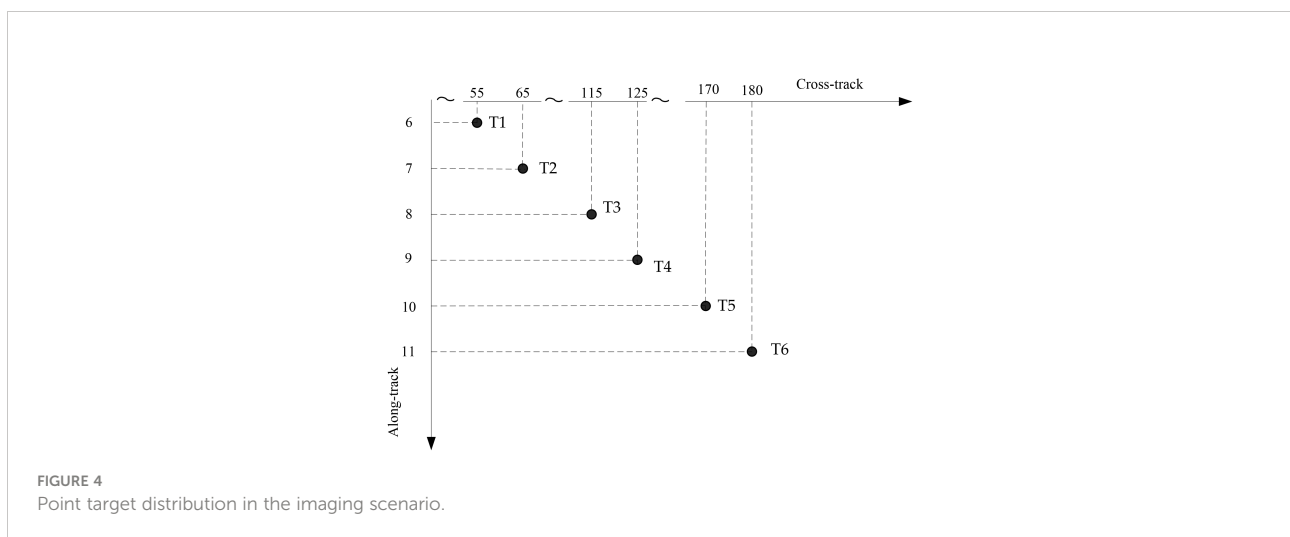
Figure 7 shows the along-track resolution and peak side lobe ratio (PSLR) of the focused targets. These results show that the proposed method can obtain high-resolution imaging results consistent with those of the BP algorithm.

The PSLR, integrated side lobe ratio (ISLR), and the along-track resolution (AR) were calculated to further evaluate the imaging performance of the proposed method. The quality parameters of the proposed method are listed in Table 2, indicating that the difference in the PSLRs between the proposed method and the BP algorithm was about 0.19 dB. The difference in the ISLRs between the proposed method and the BP algorithm was about 0.08 dB. The degradation in the quality parameters was very modest, and the performance loss can be neglected. Both methods obtained essentially the same along-track resolution. Therefore, it can be concluded that the proposed method can focus the targets well. Furthermore, the proposed method can obtain high-resolution imaging results similar to those of the BP algorithm.

5 Processing results for real data

In this section, real data were used to verify the proposed method. The real data included 800 sampling points in the range dimension. Because the BP algorithm is considered precise, the BP results were again used as the benchmark. Figure 8A depicts the results of the proposed method, while Figure 8B shows the results of the BP algorithm. It can be observed that the proposed method can obtain nearly the same results as the BP algorithm.

Table 3 lists the processing times of the proposed method and the BP algorithm, which showed that the BP algorithm is extremely time-consuming. However, the proposed method can significantly improve the imaging efficiency by a factor of 62 in terms of the processing speed. Consequently, it is concluded that the proposed method can accurately focus the targets without loss of imaging performance. In addition, the imaging efficiency can be improved using the proposed method.



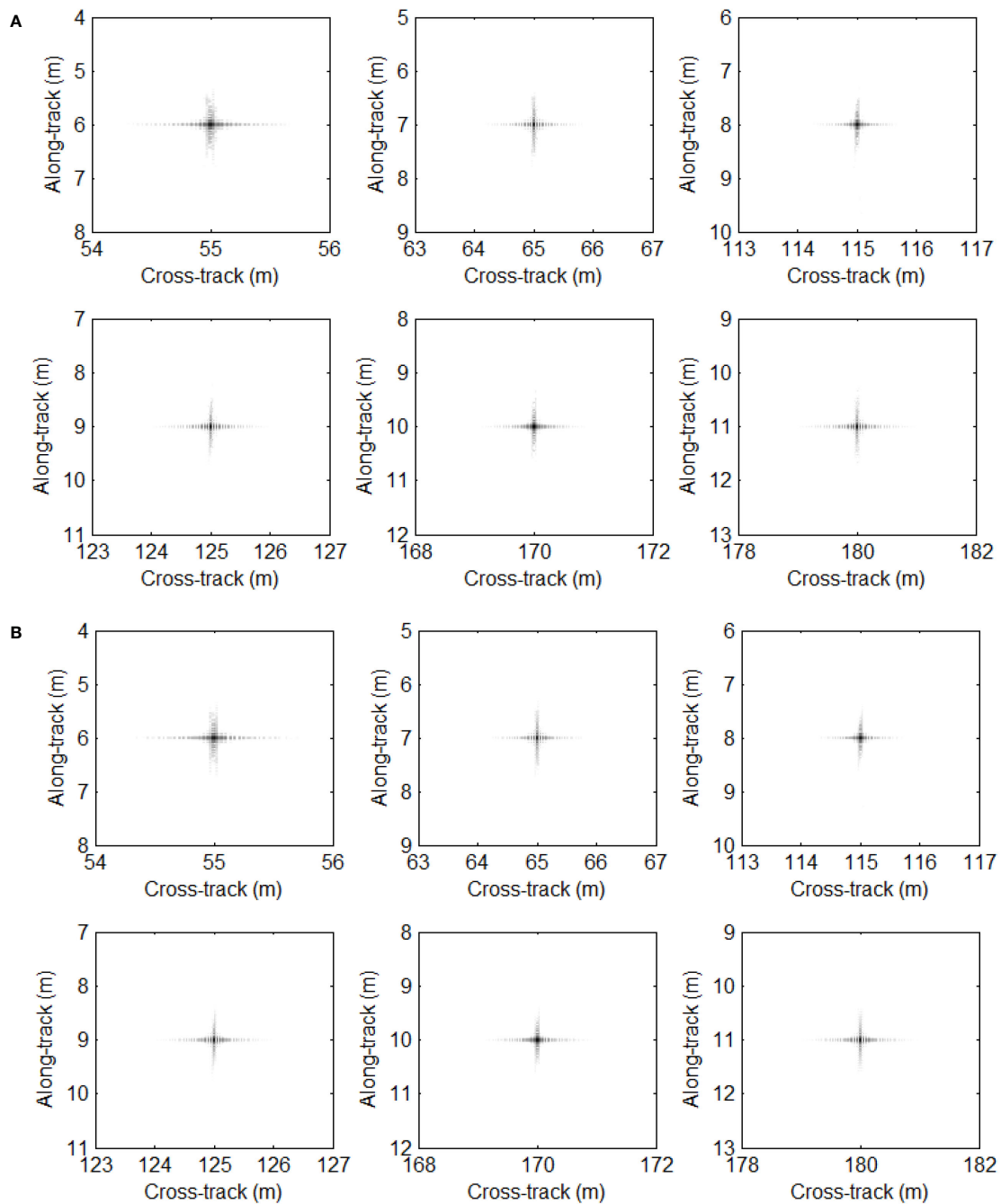


FIGURE 5 Imaging results using simulated data. (A) Proposed method. (B) Back-projection (BP) algorithm.

6 Conclusions

MSAS imagery is challenging because the two-round slant range is characterized by double square root terms. The method proposed in this paper transforms the two-round

slant range into a single square root term based on the second-order Taylor expansion and inverse Taylor expansion to solve this problem. Subsequently, the PTRS can be decomposed into quasi-monostatic and bistatic deformation terms. The bistatic deformation term can be

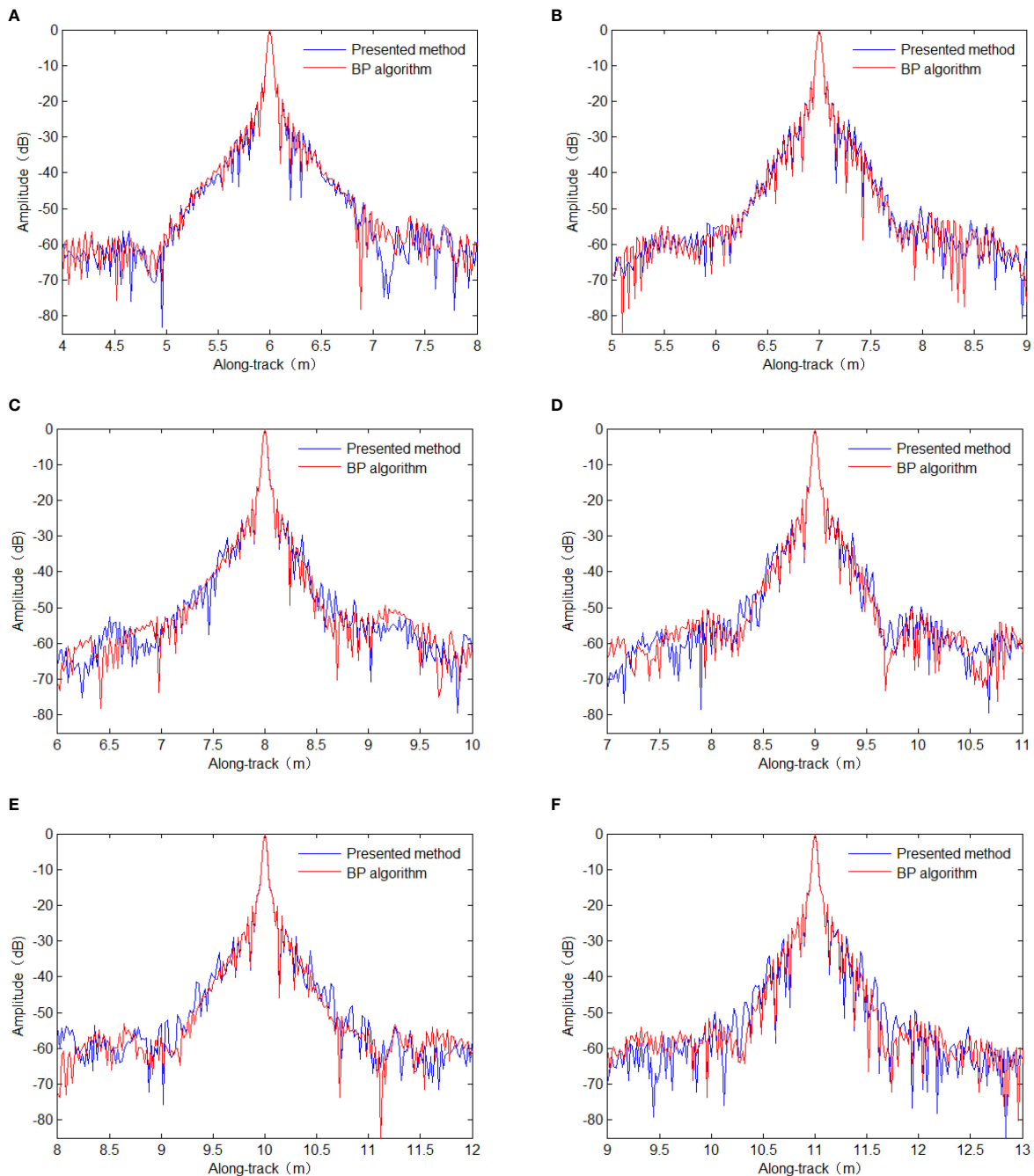


FIGURE 6
 Along-track slices of the focused targets. (A) T1. (B) T2. (C) T3. (D) T4. (E) T5. (F) T6. BP, back-projection.

anceled with the preprocessing step based on complex multiplication, interpolation, coherent superposition, and dechirping operations. Consequently, the MSAS data can be considered a monostatic SAS-equivalent signal, which is used

as the input to an RD algorithm. Considering the cross-track variance of the SRC, this paper exploits the sub-block processing method in the cross-track dimension. The proposed method can generate high-resolution results very

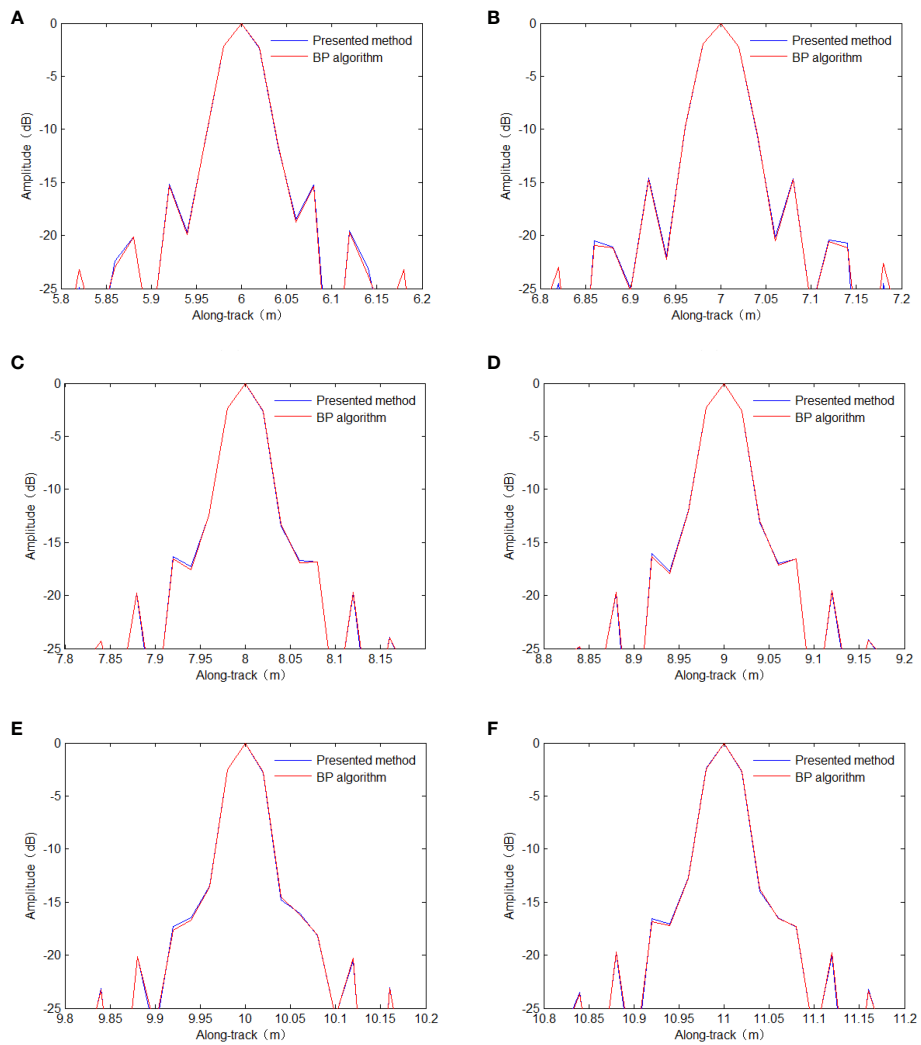


FIGURE 7
 Along-track resolution. (A) T1. (B) T2. (C) T3. (D) T4. (E) T5. (F) T6. BP, back-projection.

TABLE 2 Quality parameters of the focused targets.

Method	PSLR (dB)	ISLR (dB)	AR (m)	Target
Proposed method	-14.44	-8.61	0.04	T1
BP	-14.57	-8.69	0.04	
Proposed method	-14.34	-8.09	0.04	T2
BP	-14.47	-8.11	0.04	
Proposed method	-14.64	-9.51	0.04	T3
BP	-14.77	-9.56	0.04	
Proposed method	-14.63	-9.36	0.04	T4
BP	-14.74	-9.41	0.04	
Proposed method	-14.65	-9.95	0.04	T5
BP	-14.84	-10.00	0.04	
Proposed method	-14.63	-9.65	0.04	T6
BP	-14.76	-9.69	0.04	

PSLR, peak side lobe ratio; ISLR, integrated side lobe ratio; AR, along-track resolution; BP, back-projection.

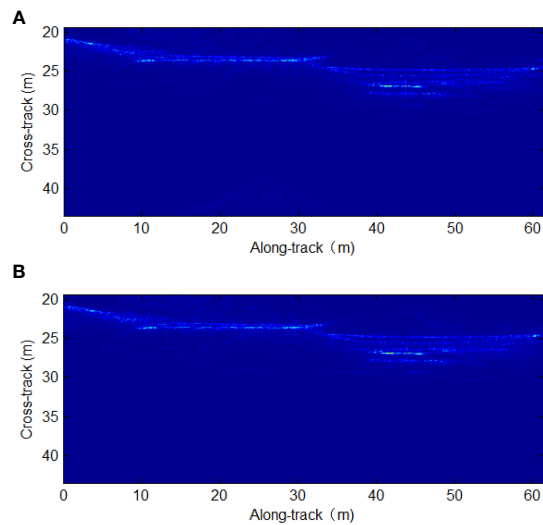


FIGURE 8
Processing results for real data. (A) Proposed method. (B) Back-projection (BP) algorithm.

TABLE 3 Processing times of the imaging algorithms.

	Proposed method	BP algorithm
Processing time (s)	28	1,761

BP, back-projection

similar to those of the BP algorithm benchmark in a significantly shorter processing time. Simulations using a simulated imaging scenario and real sonar data further validated the proposed method.

Data availability statement

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

Author contributions

PH and PY designed the study, wrote the manuscript, managed communication between the authors, and analyzed the data. All authors contributed to the article and approved the submitted version.

Funding

This work was supported by the Natural Science Foundation of Shandong Province (grant no.

ZR2019MA022) and the Enterprise Marine Project (grant no. 20210701).

Acknowledgments

The authors thank the reviewers and editors for their valuable comments that helped improve this paper.

Conflict of interest

PY is employed by the Acoustic Signal and Electronics Science and Technology Corporation.

The remaining author declares that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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