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**Li, Y, Li, H, Zhang, C, Zhao, Y and Yang, Z (2024) Incorporation of adaptive compression into a GPU parallel computing framework for analyzing largescale vessel trajectories. Transportation Research Part C: Emerging Technologies, 163. ISSN 0968-090X** 

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# Transportation Research Part C



journal homepage: [www.elsevier.com/locate/trc](https://www.elsevier.com/locate/trc) 

# Incorporation of adaptive compression into a GPU parallel computing framework for analyzing large-scale vessel trajectories

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# ARTICLE INFO

*Keywords:*  Trajectory compression Parallel computing Graphics processing unit (GPU) Unmanned ships Maritime safety

#### ABSTRACT

Automatic Identification System (AIS) offers a wealth of vessel navigation data, which underpins research in maritime data mining, situational awareness, and knowledge discovery within the realm of intelligent transportation systems. The flourishing marine industry has prompted AIS satellites and base stations to generate massive amounts of vessel trajectory data, escalating both data storage and calculation costs. The conventional Douglas-Peucker (DP) algorithm used for trajectory compression sets a uniform threshold, which hampers effective compression. Additionally, compressing and accelerating the computation of large datasets poses a significant challenge in real-world applications. To address these limitations, this paper aims to develop a new Graphics Processing Unit (GPU) parallel computing and compression framework that enables the acceleration of the optimal threshold calculation for each trajectory automatically in maritime big data mining. It achieves this by incorporating a new Adaptive DP with Speed and Course (ADPSC) algorithm, which utilizes the dynamic navigation characteristics of different vessels. It can effectively solve the associated computational time cost concern when using the ADPSC algorithm to compress vast trajectory datasets in the real world. Additionally, this paper proposes a novel evaluation metric for assessing compression efficacy based on the Dynamic Time Warping (DTW) method. Comprehensive experiments encompass vessel trajectory datasets from three representative research areas: Tianjin Port, Chengshan Jiao Promontory, and Caofeidian Port. The experimental results demonstrate that 1) the newly developed ADPSC method outperforms in terms of compression, and 2) the designed GPU parallel computing framework can significantly shorten the compression time for extensive datasets. The GPU-accelerated compression methodology not only minimizes storage and transmission costs for data from both manned and unmanned vessels but also enhances data processing speed, supporting real-time decision-making. From a theoretical perspective, it provides the key to the puzzle of realizing the real-time anticollision of manned and unmanned ships, particularly in complex waters. It hence makes significant contributions to maritime safety in the autonomous shipping era.

# **1. Introduction**

The rise of economic globalization has catalyzed frequent import and export trade among countries, spurring the continuous growth in the shipping industry [\(Li et al., 2023a; Tagiltseva et al., 2022](#page-38-0)). Among various transport sectors enabling international trade,

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<https://doi.org/10.1016/j.trc.2024.104648>

Received 27 September 2023; Received in revised form 12 April 2024; Accepted 30 April 2024

Available online 9 May 2024

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# **Nomenclature**



the maritime industry has emerged as the preferred one for global trade transportation. Its dominance is attributed to cost-effective shipping and the ability to accommodate a diverse range of goods [\(Li et al., 2022; Li and Lam, 2017\)](#page-38-0). Vessels are pivotal in the maritime industry and can generate massive dynamic data during navigation [\(Li et al., 2024\)](#page-38-0). This data typically includes details like time stamp, longitude, latitude, Speed Over Ground (SOG), and Course Over Ground (COG). Integrated into an Automatic Identification System (AIS) ([Li et al., 2023b; Yang et al., 2019](#page-38-0)), this data is transmitted to designated servers through base stations and satellites, as illustrated in [Fig. 1](#page-3-0). Once received, AIS data undergoes processing through algorithms such as mathematical statistics, visual analysis, and anomaly recognition on the server side, yielding valuable information (e.g., traffic intensity, traffic flow distribution characteristics, and collision risk) ([Li et al., 2023; Xin et al., 2023](#page-38-0)). Such information not only aids in monitoring vessel navigation dynamics but also provides critical insights for developing intelligent maritime transportation systems.

The robust growth of the global maritime industry has led to a steady rise in the number of vessels, resulting in a surge of vessel navigation (or AIS) data transmitted to servers. This data expansion strains storage capacity, often containing redundant information [\(Sun et al., 2020; Zheng et al., 2020](#page-39-0)). It is crucial to remove such redundancies for accurate data analysis. For instance, in a vessel's straight-line navigation, merely noting the starting and ending points suffices to understand its trajectory, rendering intermediate points unnecessary.

Trajectory compression technology emerges as an effective solution to these challenges in accurate data analysis. Its core idea is to pinpoint feature points in a trajectory to replace the original data, eliminating redundancy and thus reducing storage costs [\(Liu et al.,](#page-38-0) [2019b; Tang et al., 2021b](#page-38-0)). Furthermore, trajectory compression offers computational cost savings ([Chen et al., 2020b, 2020a](#page-37-0)). Since the compressed vessel trajectory efficiently captures sailing characteristics while reducing computational time for other research algorithms, numerous scholars employ it for future studies, including trajectory clustering [\(Bai et al., 2023; Tang et al., 2021a](#page-37-0)), route extraction (Karataş et al., 2021; Yan et al., 2020), path planning [\(Gu et al., 2023; Liu et al., 2019a\)](#page-38-0), trajectory anomaly recognition [\(Dogancay et al., 2021; Liang et al., 2022; Rong et al., 2020](#page-37-0)), and collision avoidance [\(Wang et al., 2023, 2024b,a; Xin et al., 2023\)](#page-39-0). It is evident that trajectory compression is a pivotal data preprocessing technique and stands as an essential research component in data mining [\(Li et al., 2022](#page-38-0)).

The trajectory compression research predominantly encompasses two approaches: batched compression [\(Li et al., 2019\)](#page-38-0) and online compression [\(Liu et al., 2016](#page-38-0)). Their essential difference is that online compression methods utilize local points for calculation when trajectory data is incomplete, making them suitable for application scenarios that involve compressing during transmission. In contrast, batched compression methods leverage the overall distribution of trajectories to identify feature points. Given that the vessel

<span id="page-3-0"></span>

**Fig. 1.** Visual illustration of satellite and terrestrial AIS networks.

trajectory data acquired in this paper are complete, the focus will be primarily on batched compression techniques. The Douglas-Peucker (DP) algorithm ([Saalfeld, 1999](#page-39-0)) stands out as a renowned, widely adopted, and effective batched compression method, capable of identifying feature points based on the vessel trajectory distribution, subsequently supplanting the original trajectory data. However, the original DP algorithm confronts two notable limitations in handling vessel trajectory compression tasks. First, it mandates the predetermination of a compression threshold, which is conventionally set manually. An excessively high threshold risks omitting vital feature points, considering them redundant, whereas an unduly low threshold may fail to eradicate superfluous data. Additionally, it leads to suboptimal compression results for the majority when applying a uniform compression threshold across all trajectories. Secondly, the original DP algorithm faces challenges in efficiently compressing large-scale data, diminishing its practical applicability.

To address the above challenges, this paper aims to develop a new Graphics Processing Unit (GPU) parallel computing and compression framework, embedding with a novel Adaptive Douglas-Peucker with Speed and Course (ADPSC) algorithm to realize massive trajectory data preprocessing and compression. It can automatically calculate compression thresholds rooted in individual vessel navigation details, including time stamps, longitude and latitude coordinates, SOG, and COG. Moreover, the newly developed ADPSC algorithm has been refined into GPU parallel algorithms ([Cheng and Gen, 2019; Kallioras et al., 2015; Swirydowicz](#page-37-0) et al., 2022; [You et al., 2022](#page-37-0)), which greatly enhances its capability to handle large-scale trajectory data, leading to a significant reduction in algorithm execution time. Finally, a new evaluation index based on the Dynamic Time Warping (DTW) approach is proposed to measure trajectory compression performance. From a theoretical standpoint, this paper provides a crucial piece of the puzzle in achieving real-time anti-collision capabilities for manned and unmanned ships, especially in complex water environments. Consequently, it makes significant contributions to enhancing maritime safety in the era of autonomous shipping.

This paper proposes an adaptive and accelerated compression framework, which can not only accurately simplify vessel trajectories but also process large-scale data quickly to adapt to practical application scenarios in maritime industries. [Section 2](#page-4-0) provides an overview of vessel trajectory compression methods and GPU parallel computing frameworks, revealing the relevant gaps and contributions. [Section 3](#page-6-0) serves as the preparation phase for the study, covering two contents. Firstly, it elucidates essential definitions that offer a theoretical overview of this paper. Secondly, it details the preprocessing of experimental data, encompassing tasks such as denoising vessel trajectory data and converting geographical coordinates. [Section 4](#page-11-0) delves into an in-depth exploration of the relevant theories behind the original DP, optimized ADPSC, and GPU-based parallel acceleration ADPSC algorithms. In [Section 5,](#page-23-0) a comparative experiment is conducted, evaluating the new ADPSC algorithm both qualitatively and quantitatively. The evaluation underscores its superior performance over the original DP algorithm in addressing vessel trajectory data compression challenges. Meanwhile, the

<span id="page-4-0"></span>speedup ratio is employed to analyze the acceleration benefits of the optimized ADPSC algorithm when compressing large-scale vessel trajectory data using the GPU parallel computing framework, as opposed to traditional Central Processing Unit (CPU) serial approaches. [Section 6](#page-35-0) sheds light on the relevant research conclusions and outlines potential avenues for future sustainable research.

#### **2. Literature review**

Many scholars have recently investigated trajectory compression to reduce data storage costs. Concurrently, there is a growing inclination towards implementing algorithms on GPU parallel computing frameworks to decrease algorithmic computation time. Advanced methods specific to vessel trajectory compression will be presented first in Section 2.1. Subsequently, [Section 2.2](#page-5-0) delves into the evolutionary development of GPU parallel computing. [Section 2.3](#page-6-0) summarizes the research contributions of this paper.

# *2.1. Review of studies on maritime trajectory compression methods*

Vessel trajectory refers to how the position of a vessel in space changes over time, with its change function being continuous. Unlike simple function curves, trajectory data contains not only position information but also dynamic details such as time, direction, and velocity. At its core, trajectory compression seeks to identify an approximate trajectory with fewer data points to substitute for the original trajectory. The development of the batched compression methods is elaborated upon in Section 2.1.1, while the evolution of the online compression techniques is detailed in Section 2.1.2.

#### *2.1.1. Batched compression methods*

It is necessary to collect the completed trajectory when performing vessel trajectory compression tasks using batched methods to further identify feature points in the data and eliminate redundant information [\(Arslan et al., 2018; Liu et al., 2019b\)](#page-37-0). This entire trajectory is taken into account in the batched methods, making it easier to achieve global optimization during compression.

The Uniform Sampling (US) and DP algorithms are the fundamental batched compression techniques frequently utilized in trajectory data compression. These methods have seen extensive refinement over the years, with the DP algorithm garnering particular attention for enhancements. The US algorithm is straightforward, with the core idea of retaining one point out of every *W* point ([Lv](#page-38-0) [et al., 2015; Sun et al., 2016](#page-38-0)). For instance, given a trajectory with 16 points and deploying the US method to keep one point out of every 5, the resultant trajectory would be formed by the first, sixth, eleventh, and sixteenth points. While the US approach boasts efficiency and reduced computational demand, its shortcoming lies in its inability to effectively conserve crucial trajectory features, leading to disparities between the compressed and the original trajectories.

In contrast, the DP algorithm [\(Douglas and Peucker, 1973](#page-37-0)) emerged to address the limitations of the US algorithm. This algorithm, essentially recursive, iteratively identifies feature points based on trajectory distributions and point deviations [\(Huang et al., 2020](#page-38-0)). Over time, researchers have improved the DP algorithm to optimize its compression capabilities. As trajectories encapsulate both spatial (longitude and latitude) and temporal (time) aspects, [Meratnia and de By \(2004\)](#page-38-0) proposed the Top-Down Time-Ratio (TD-TR) algorithm. This approach integrates spatial distance with time ratios, generating more accurate compressed trajectories. To expedite the compression and boost the compression rate, [Hansuddhisuntorn and Horanont \(2019\)](#page-38-0) put forward an improved version of the TD-TR algorithm called TD-TR Reduce. [Liu et al. \(2015\)](#page-38-0) utilized data structures like convex hulls to optimize the spatial and temporal complexity of the DP algorithm. Their algorithm could achieve the best compression effect for trajectory data with a space complexity of *O*(1) and a time complexity of *O*(*n*). [Zhao and Shi \(2018\)](#page-39-0) merged the directional changes of vessel trajectories with the DP algorithm, yielding impressive results under high compression. [Zhou et al. \(2023\)](#page-39-0) pointed out the shortcomings of the DP algorithm in compressing vessel trajectory data, including three specific points: (1) there is a continuous turning phenomenon in the trajectory, and its compression effect is poor; (2) it does not take into account the impact of vessel speed and course; (3) there's a possibility of errors, such as the compressed vessel trajectory intersecting obstacles. To address the above issues, they introduced a Multi-objective Peak DP algorithm (MPDP), which incorporated a peak sampling strategy. In recent years, scholars identified that universally applying the same manually set compression threshold across all trajectories compromises compression accuracy. Therefore, some adaptive compression algorithms have been proposed by scholars. [Liu et al., \(2019b\)](#page-38-0) proposed the Adaptive Douglas-Peucker (ADP) algorithm for compressing vessel trajectory data, which employs the average distance from the trajectory point to the baseline (a line linking the starting and ending points) to automatically calculate the compression threshold. [Tang et al., \(2021b\)](#page-39-0) utilized the threshold change rate to determine the feature points of each trajectory. [Li et al. \(2022\)](#page-38-0) developed the compression algorithm proposed by [Liu et al., \(2019b\),](#page-38-0) incorporating not just the distance from the point to the baseline, but also the velocity change rate of each trajectory point when determining the compression threshold. Their method, termed Adaptive Douglas-Peucker with Speed (ADPS), efficiently extracts the featured points for knowledge discovery.

The adaptive compression algorithm autonomously determines the compression threshold for each trajectory, guaranteeing trajectory accuracy even at elevated compression rates. Unlike some existing adaptive compression algorithms, the newly developed ADPSC algorithm takes into account variations in speed and direction for each trajectory point, addressing a longstanding research gap. Furthermore, the ADPSC algorithm is redesigned for a parallelized version tailored for GPU computing frameworks, enabling rapid compression of extensive vessel trajectory datasets for real-world applications.

#### *2.1.2. Online compression methods*

The online compression methods employ local features of trajectories to identify critical points ([Gao et al., 2019](#page-37-0)). It is suitable for application scenarios where compression occurs simultaneously with transmission. [Keogh et al. \(2001\)](#page-38-0) introduced two algorithms,

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**Fig. 2.** Overview of differences between CPU and GPU computing frameworks. (a) CPU architecture, (b) GPU architecture, and (c) conceptual architecture of the CUDA parallel computing model.

namely Sliding Window (SW) and Open Window (OW). The core idea of SW employs an initialized sliding window to gradually search for feature points in the trajectory. If no feature points are found in the trajectory segment inside the sliding window, it expands, encompassing new trajectory points until the final trajectory point is reached. While OW and SW share similar compression fundamentals, they differ in their approach to assessing feature points. [Gao and Shi \(2019\)](#page-37-0) extract key feature points from vessel trajectories based on vessel heading angle deviation, position deviation, and spatiotemporal features of AIS data, and then optimize the SW algorithm. This new method is called the vessel spatiotemporal key feature point online extraction algorithm. [Sun et al. \(2020\)](#page-39-0) proposed a Scan-Pick-Move (SPM) trajectory data compression algorithm based on SW to address the high compression rate and long processing time in existing online trajectory compression algorithms. They used the maximum offset distance reference trajectory point to determine whether the current trajectory point can be compressed, aiming to reduce the storage space. [Zhu and Ma \(2021\)](#page-39-0) used the trajectory change rate and velocity change rate in SW as the criteria for simplifying trajectory points. Compared with the DP algorithm, SW algorithm, and Opening Window Time Ratio (OPW-TR) algorithm, their method effectively considered vessel behavior patterns and compressed vessel trajectory data. [Liu and Yang \(2023\)](#page-38-0) proposed an improved opening window trajectory simplification algorithm, analyzing the impact of distance and velocity thresholds on algorithm performance to determine appropriate simplification thresholds. It is worth noting that this algorithm better preserves the position information and spatial features of the original trajectory. [Potamias et al. \(2006\)](#page-39-0) put forward the threshold-guided sampling algorithm, which employs speed and direction to construct secure regions and then searches for feature points. [Muckell et al. \(2011\)](#page-38-0) developed an online compression algorithm called the Spatial QUalIty Simplification Heuristic (SQUISH). To compress a trajectory, SQUISH starts by initializing a priority queue, and then progressively adds trajectory points. Once the queue hits its capacity, it removes the point, resulting in the least error. After this, the priority of each trajectory point is updated. SQUISH not only maintains the accuracy of compressed trajectories at high compression rates, but also boasts a relatively low time complexity. Consequently, it is a popular choice for online trajectory data compression research, with several scholars suggesting enhanced versions. For example, [Muckell et al. \(2014\)](#page-38-0) designed the SQUISH-Extended (SQUISH-E) algorithm, which can achieve the best compression rate within a given error threshold. [Han et al. \(2018\)](#page-38-0) adopted a multi-core computing framework to accelerate the SQUISH-E algorithm, introducing a Parallel version of the SQUISH-E algorithm called PSQUISH-E. Additionally, they developed the GPU-assisted PSQUISH-E algorithm called G-PSQUISH-E.

Both batched and online compression methods compress trajectory data in an Euclidean space. However, their primary distinction lies in their approach to traversing trajectory points. While batched compression methods extract global feature points from trajectory data, online compression techniques rely on local trajectory information to identify these points, which can result in a higher compression error. Meanwhile, the average time complexity of batched compression technology is generally higher than that of online compression technology.

#### *2.2. Review of studies on GPU computation*

Given the increased computational complexity of the optimized ADPSC algorithm compared to the DP algorithm and the vast scale of trajectory datasets that need compression, there is a significant computational demand for trajectory compression. Traditional CPU serial computing frameworks struggle to process optimization algorithms for large-scale dataset compression within a reasonable time frame. It would significantly increase computational costs for engineers to upgrade CPU specifications to meet these demands continuously. New solutions to these challenges are demanded with urgency. GPU offers robust computational capabilities, positioning them as formidable platforms for large-scale data mining ([Chen et al., 2018; Heywood et al., 2019; Jurczuk et al., 2021\)](#page-37-0). Currently, <span id="page-6-0"></span>while mainstream CPUs typically include several cores, each corresponding to two threads for simultaneous computing, conventional GPUs can run thousands of threads concurrently [\(Huang et al., 2020; Jeong et al., 2022; Qu and Zhou, 2017\)](#page-38-0). The contrasting structures of CPUs and GPUs are illustrated in [Figs. 2](#page-5-0) (a) and (b).

The GPU's capacity to support thousands of threads underscores its superior computing prowess compared to the CPU, as depicted in [Fig. 2](#page-5-0) (c). In the GPU parallel computing architecture, a thread is the foundational calculation unit. One block consists of up to 1,024 threads, and several blocks combine to form a grid [\(Lin et al., 2023\)](#page-38-0). Within a block, threads can communicate via shared memory. However, inter-block thread communication relies on global memory. Notably, data transfer in shared memory is more efficient than that in global memory (Cagigas-Muñiz [et al., 2022; Shanbhag et al., 2022\)](#page-37-0). It is challenging to apply the intricate GPU directly in conventional parallel computing. To simplify the GPU calculation process, the General-Purpose GPU (GPGPU) was proposed, mapping general computing tasks to graphic hardware [\(Owens et al., 2007](#page-38-0)). Although GPGPU offers a more accessible entry point than GPU, only professional engineers well-versed in graphic APIs can proficiently master it. As a result, Nvidia launched Compute Unified Device Architecture (CUDA) in 2006, overcoming the drawbacks of both GPU and GPGPU. CUDA provides a versatile programming interface, facilitating data processing and analysis tasks on NVIDIA GPUs [\(Basnet et al., 2022](#page-37-0)).

Parallel algorithms can map data across GPU threads, enabling swift execution of tasks like processing large-scale datasets [\(Manduhu and Jones, 2019; Roberge and Tarbouchi, 2021\)](#page-38-0). Since vessel trajectories consist of distinct points, each being logically independent, it is evident that designing parallel algorithms within the GPU computing structure is suitable for processing extensive vessel trajectory data in the maritime sector. However, the current literature reveals that very few studies relating to the use of GPU for vessel trajectory analysis, and to the authors' best knowledge, no studies have been undertaken to address the design of parallel algorithms in a GPU computing structure in the maritime sector, disclosing the theoretical novelty of this work.

# *2.3. Contributions of our study*

To address the aforementioned limitations and research gaps, this paper proposed an ADPSC method, an innovative DP-based trajectory compression algorithm, harnessing vessel trajectories' distribution characteristics and additional dynamic navigation data such as time stamps, SOG, and COG to determine the compression threshold for each trajectory autonomously. Consequently, it proficiently prunes unnecessary data, addressing the first limitation of the DP algorithm. Furthermore, this paper refines the ADPSC algorithm to expedite the compression of massive vessel trajectories within the GPU parallel computing environment, effectively tackling the DP algorithm's second limitation and research gaps in GPU computation. The paper's core contributions are delineated as follows.

(1) *Propose an adaptive compression algorithm based on multiple factors.* 

The new ADPSC algorithm leverages the dynamic navigation information of ships, including location, SOG, and COG. This ensures a unique compression threshold for each trajectory. The proposed approach primarily resolves the problem of employing a universal threshold for all trajectories, which previously led to suboptimal compression outcomes.

(2) *Develop a GPU parallel computing framework that integrates an ADPSC algorithm.* 

Given the vast datasets requiring compression in real-world applications, this paper further fine-tunes the ADPSC algorithm, constructing parallel computing strategies to facilitate trajectory data compression on GPUs. This enhancement notably boosts computational speed, making it apt for real-time data processing in practical settings.

(3) *Design a new index for evaluating compression effectiveness.* 

Evaluation metrics offer a quantitative assessment of algorithmic compression performance. The DTW approach serves as a metric to gauge the similarity between trajectories pre and post-compression, thereby appraising the algorithm's compression quality. This paper deviates from conventional DTW measurement methods. Instead, it first computes the distance between trajectory points and the baseline both before and after compression, generating two vectors. Subsequently, the DTW method measures the similarity between these vectors, using the resultant value to evaluate compression efficacy. This improved method proves more apt for assessing the compression outcomes of trajectory data.

(4) *Conduct comparative experiments using three substantial and representative datasets.* 

To demonstrate the universality of the experiment, this paper collected actual AIS data from three representative regions: Tianjin Port, Chengshan Jiao Promontory, and Caofeidian Port. In the comparative analysis of compression performance, a mix of qualitative and quantitative methods is employed to evaluate the superior performance of the ADPSC algorithm over the original DP algorithm. Additionally, this paper uses the acceleration ratio as a metric to quantitatively gauge the speed-up efficiency of the ADPSC algorithm when compressing trajectory data within the GPU parallel computing framework.

# **3. Preliminary**

This section provides clear definitions essential for understanding and implementing trajectory compression, as detailed in [Section](#page-7-0) [3.1](#page-7-0). Additionally, vessel trajectory data should undergo processes such as denoising and coordinate conversion before compression. The denoising step focuses on identifying and rectifying noisy data, with its specifics elaborated in [Section 3.2.](#page-8-0) Meanwhile, coordinate conversion translates trajectory data from the World Geodetic System − 1984 Coordinate System to the Mercator Projection Coordinate System, a procedure that will be described in [Section 3.3](#page-9-0).

### <span id="page-7-0"></span>**Table 1**  List of the notations.



# *3.1. Definitions*

This list of notations is presented in Table 1 to enhance clarity throughout the content. Various essential definitions are provided below.

**Definition 1. (***The original vessel trajectory.***)** A vessel trajectory  $T_{ori}$  with the length *O* includes a series of points collected by the AIS base station. The mathematical expressions are as follows,

$$
T_{ori} = \left\{ P_{ori}^{1}, P_{ori}^{2}, ..., P_{ori}^{n}, P_{ori}^{n+1}, ..., P_{ori}^{O} \right\},\tag{1}
$$

$$
P_{\text{ori}}^{n} = \left\{ t_{\text{ori}}^{n}, \text{lon}_{\text{ori}}^{n}, \text{lat}_{\text{ori}}^{n}, \text{sog}_{\text{ori}}^{n}, \text{cog}_{\text{ori}}^{n} \right\}, n = 1, 2, ..., O
$$
\n(2)

where  $P_{ori}^n$  denotes the *n*-th point in the original vessel trajectory  $T_{ori}$ ,  $t_{ori}^n$ ,  $lon_{ori}^n$ ,  $lat_{ori}^n$ ,  $sog_{ori}^n$ , and  $cog_{ori}^n$  represent the timestamp, longitude, latitude, SOG, and COG data in the *n*th trajectory point, respectively.

**Definition 2**. (*Compressed vessel trajectory.*) A sequence of data *Tcom* with a length *C* is compressed from the original vessel trajectory, whose mathematical expressions are represented in Eqs. (3) and (4),

$$
T_{com} = \{P_{com}^1, P_{com}^2, ..., P_{com}^{m-1}, P_{com}^m, ..., P_{com}^C\},\tag{3}
$$

$$
P_{com}^{m} = \{t_{com}^{m}, \text{lon}_{com}^{m}, \text{lat}_{com}^{m}, \text{sog}_{com}^{m}, \text{cog}_{com}^{m}\}, m = 1, 2, ..., C.
$$
\n
$$
(4)
$$

where  $P_{com}^m$  is the m-th point in the compressed vessel trajectory.  $t_{com}^m, lon_{com}^m, lat_{com}^m, sog_{com}^m$ , and  $co_{com}^{m}$  represent the timestamp, longitude, latitude, SOG, and COG data in the *m*th trajectory point, respectively. Specifically, the disparity between *O* and *C* indicates the volume of redundant data removed by the compression algorithm.

**Definition 3. (AIS data matrix.)** A matrix of  $5 \times O$  contains a primary dynamic AIS information of a vessel, defined as follows,



where *O* denotes the total number of points in all trajectory data, and 5 counts the number of the involved parameters, including *t*, *lon*, *lat, sog, and cog. AISM* is used as the input data of the GPU-accelerated ADPSC algorithm to reduce frequent data transmission in this paper.

<span id="page-8-0"></span>

Fig. 3. Visual illustration of vessel trajectory data denoising, (a) trajectory data with noise, and (b) trajectory data after noise removal. In particular, the green dots represent noisy data.



**Fig. 4.** Visualisation of noisy trajectory data in three research areas: (a) Tianjin Port, (b) Chengshan Jiao Promontory, and (c) Caofeidian Port. In particular, the orange boxes indicate noisy data.

#### *3.2. Trajectory data denoising*

During the transmission process of vessel trajectory data between AIS base stations and satellites, there may be noise data, as illustrated in Fig. 3 (a). Unprocessed noise data can impede further research. For instance, when analyzing the compressed content of trajectory data, algorithms may mistakenly assume that noisy data are feature points and preserve them. Therefore, as a preparatory step before conducting specific research in this paper, it is essential to denoise the original trajectory and obtain high-quality trajectory data, as shown in Fig. 3 (b).

The denoising process of original trajectory data mainly includes two aspects: one is to identify noisy data, and the other is to repair noisy data. There are many methods for identifying noisy data, and the most commonly used is to separate noisy data based on clustering methods. This paper uses Density-Based Spatial Clustering of Applications with Noise (DBSCAN) [\(Bai et al., 2023; Li et al.,](#page-37-0) [2021\)](#page-37-0), the most representative clustering method, to identify noisy data in the original vessel trajectory. This method has two important parameters. One is the domain radius when defining density, abbreviated as *Eps*. The other refers to the threshold for defining core points, which represents the number of minimum point sets within the cluster, abbreviated as *Minpt*. These two parameters determine which cluster each data in the dataset belongs to. In practical clustering calculations, the DBSCAN method divides the trajectory points in the dataset into three categories: core points, boundary points, and noise points. The core point indicates that the number of data points within its radius *Eps* exceeds *Minpt*. The boundary point means that the number of data points contained in its radius *Eps* is less than *Minpt*, and the point falls within the area of the core point. The trajectory points in the dataset that are neither core nor boundary points are classified as noise points. DBSCAN method can accurately identify the noise data in the trajectory. The values of *Eps* and *Minpt* are configured as 0.01 and 3, respectively, as demonstrated in the parameter optimization process detailed in [Appendix A.](#page-36-0) These values represent the optimal parameters for the DBSCAN algorithm to identify outliers across all vessel trajectories within the three study areas.

The next step is to remove these noise data and repair them with the linear interpolation method ([Blu et al., 2004\)](#page-37-0). This paper assumes that the trajectory points (*lont*, *latt*) at time *t* are noise data. The detailed calculation process of using the trajectory coordinates at  $t - 1$  and  $t + 1$  to repair the longitude and latitude data of the noise point is shown in Eqs. [\(6\) and \(7\)](#page-9-0), respectively,

<span id="page-9-0"></span>

**Fig. 5.** The distribution of vessel trajectories after denoising in three research areas: (a) Tianjin Port, (b) Chengshan Jiao Promontory, and (c) Caofeidian Port.

An example of vessel trajectory points (i.e., longitude and latitude) in two coordinate systems.



$$
lon_{t} = lon_{t-1} + (t - (t - 1)) \times \left(\frac{lon_{t+1} - lon_{t-1}}{(t+1) - (t-1)}\right)
$$
\n(6)

$$
lat_{t} = lat_{t-1} + (t - (t - 1)) \times \left(\frac{lat_{t+1} - lat_{t-1}}{(t+1) - (t-1)}\right)
$$
\n
$$
(7)
$$

where  $lon_{t-1}$  and  $lon_{t+1}$  represent the longitude of trajectory points at  $t-1$  and  $t+1$ , respectively. *lat<sub>t-1</sub>* and  $lat_{t+1}$  denote the latitude of trajectory points at  $t - 1$  and  $t + 1$ , respectively.

Eqs. (6) and (7) take the time information of the noise point and adjacent points as the baseline, and use the coordinates of adjacent points to interpolate the longitude and latitude data of the noise point separately. [Figs. 4 and 5](#page-8-0) show the distribution of noisy trajectories and preprocessed trajectories in the three study areas on the map, respectively. In summary, DBSCAN and linear interpolation methods can accurately detect and repair noise data, thereby obtaining high-quality trajectory data for subsequent compression research.

#### *3.3. Conversion of geographical coordinates*

Calculating the spherical distance between two consecutive points in a trajectory based on the World Geodetic System − 1984 Coordinate System is challenging, potentially leading to significant errors [\(Huang et al., 2020](#page-38-0)). Hence, it is not advisable to analyze raw vessel trajectory data directly with compression algorithms. To address the issue, this paper proposes converting the trajectory data coordinates using the Mercator Projection Coordinate System instead of retaining the original system. Meanwhile, the World Geodetic System − 1984 Coordinate System exhibits significant deformation in high latitude areas in two-dimensional space, unlike the Mercator Projection Coordinate System, which does not have this issue. Suppose ( $\langle \textit{lon}_W, \textit{lat}_W \rangle$  and  $(\textit{lon}_M, \textit{lat}_M)$  denote the longitude and latitude data of the vessel trajectory points in the original and transformed coordinate systems, respectively. The calculation process of coordinate conversion is as follows:

$$
R = \frac{d}{\sqrt{1 - e^2 \sin^2 g}} \times \cos g,\tag{8}
$$

$$
S = \text{Intan}\left(\frac{\pi}{4} + \frac{lat_W}{2}\right) + \frac{e}{2}\ln\frac{1 - e\sinlat_W}{1 + e\sinlat_W},\tag{9}
$$

$$
lon_M = lon_W \times R,
$$
\n(10)

$$
lat_M = S \times R. \tag{11}
$$

where *R* denotes the radius of a parallel circle at the standard latitude, and *d* represents the long radius of the Earth's ellipsoid. *g*  denotes the standard latitude in the Mercator projection, while *e* represents the first eccentricity in the Earth's ellipsoid. *S* denotes isometric latitude.

Table 2 displays the representation of a single point's information from vessel trajectory data in two different coordinate systems.

<span id="page-10-0"></span>

**Fig. 6.** The whole framework.

<span id="page-11-0"></span>List of the notations.





**Fig. 7.** The schematic of the original DP compression algorithm, (a) an original vessel trajectory, (b) feature points selection based on the threshold, (c) trajectory segmentation and feature points searching within different trajectory segments, (d) identify new feature points iteratively, and (e) the compressed vessel trajectory.

The unit of latitude and longitude coordinates has changed from degrees to meters.

# **4. Methodology**

# *4.1. A GPU-accelerated compression framework*

The entire framework of this paper is illustrated in [Fig. 6](#page-10-0), encompassing AIS data collection, data preprocessing, GPU-accelerated adaptive compression technique, compression evaluation criteria, and experimental analysis. Embedded within this methodology are the innovative ADPSC approach and its GPU-accelerated counterpart. In particular, AIS data collection and preprocessing are the

<span id="page-12-0"></span>

**Fig. 8.** Visual illustration of trajectory points' offset and SOG decomposition relative to the baseline: (a) and (b) reflect the distance distribution between the midpoint and the baseline for two trajectories, respectively, and (c) displays the SOG decomposition process of trajectory points based on the coordinate system with baseline as the *x*-axis.

preparatory work of the new methodology, aimed at obtaining high-quality vessel trajectory data to avoid the compression algorithm capturing feature points incorrectly. These contents are elaborated in [Section 3.2](#page-8-0). Sections 4.2 and 4.3 provide comprehensive explanations of the novel ADPSC algorithm and the GPU-accelerated ADPSC method, respectively. They constitute the core content and main contributions of this paper. To verify the effectiveness of the newly proposed algorithms, [Sections 5.2, 5.3, and 5.4](#page-24-0) introduce evaluation metrics, compression effect, and acceleration performance, respectively. This section introduces new notations for describing the methodology listed in [Table 3](#page-11-0).

#### *4.2. The proposed ADPSC-based vessel trajectory compression method*

The essence of the DP algorithm is to find the feature points in the vessel trajectory data by setting a threshold, illustrated in [Fig. 7](#page-11-0). These feature points can accurately reflect the features of the original trajectory. A compilation of these features then takes the place of the original trajectory data, achieving the compression objective [\(Li et al., 2016\)](#page-38-0). Owing to its straightforwardness and efficacy in maritime transportation contexts, it has garnered considerable interest.

The specific steps of the DP algorithm in executing trajectory data compression tasks are as follows:

(1) The starting and ending points of the trajectory serve as feature points and are generated as the baseline, such as points  $P_1$ (P<sub>Begin</sub>) and P<sub>16</sub> (P<sub>End</sub>) in [Fig. 7](#page-11-0) (b). Using the DP algorithm, the distance from intermediate points (i.e., P<sub>2</sub> to P<sub>15</sub>) to the baseline is calculated and compared with the threshold. If the distance exceeds the threshold, the point becomes a new feature point, like point  $P_{10}$ ( $P_{Feature}$ ) in [Fig. 7](#page-11-0) (b).

(2) In [Fig. 7](#page-11-0) (c), the new feature point P<sub>10</sub> divides the original trajectory into two segments. Subsequently, the feature points P<sub>5</sub> and  $P_{12}$  in these segments can be obtained by repeating the calculation process in step (1). Notably, the distance from point  $P_{12}$  to the baseline of the second trajectory segment is below the threshold, resulting in only point P5 becoming a new feature point and splitting the first segment trajectory into two subsets, as depicted in [Fig 7](#page-11-0) (d).

(3) Although  $P_3$  and  $P_8$  are the farthest from the baseline in new segments, their distances remain below the threshold.

In summary, only points  $P_1$ ,  $P_5$ ,  $P_{10}$ , and  $P_{16}$  are retained as feature points, replacing the original trajectory in [Fig. 7](#page-11-0) (e). The DP algorithm continues to search for feature points until the maximum distance from any segment point to the baseline is less than the threshold.

The traditional DP algorithm relies on manually establishing a threshold for all trajectory data, typically based on experience rather than theoretical principles. However, this approach results in varied compression thresholds across various vessel trajectories, posing challenges. Utilizing a uniform threshold for all trajectories leads to difficulties in handling redundant data and the potential loss of crucial trajectory feature points. Consequently, the development of adaptive compression algorithms emerges as a crucial research avenue.

Determining a compression threshold is quantifying the extent of deviation (essentially the distance) between trajectory points and the baseline, progressively pinpointing feature points. A trajectory point with a substantial offset from the baseline is more likely to be a feature point. In line with this approach, some researchers compute the average distance from all intermediary points in the tra-jectory to the baseline, forming a threshold ([Li et al., 2022; Liu et al., 2019b](#page-38-0)). The functional expressions for this threshold are as follows,

$$
threshold = \frac{1}{lt - 2} \sum_{i=2}^{lt-1} dis_i
$$
\n
$$
(12)
$$

$$
dis_i = \frac{|(lon_S - lon_i) \times (lat_E - lat_i) - (lon_E - lon_i) \times (lat_S - lat_i)|}{\sqrt{(lon_S - lon_E)^2 + (lat_S - lat_E)^2}}
$$
\n(13)

where *lt* represents the total number of trajectory points, and *disi* denotes the distance from the *i-*th trajectory point to the baseline. (*lon<sub>S</sub>*, *lat<sub>S</sub>*) and (*lon<sub>E</sub>*, *lat<sub>E</sub>*) are the coordinates of the starting and ending points of the trajectory point, respectively. (*lon<sub>i</sub>*, *lat<sub>i</sub>*) represents the longitude and latitude coordinates of the *i-*th trajectory point. In particular, the denominator in Eq. (13) can not be 0 due to the distinct positions of the collected vessel trajectory data's starting and ending points. Furthermore, the initialization feature points exclude the trajectory's starting and ending points from the calculation, so the value of *i* is between 2 and *lt*-1.

<span id="page-13-0"></span>

**Fig. 9.** The scheme of compression threshold calculation: (a) offset weight, (b) SOG variation weight, and (c) weight fusion.

The average calculation method has a limitation, potentially obscuring critical information. As depicted in [Fig. 8](#page-12-0) (a), the distance from  $P_6$  to the baseline is the largest, implying that its contribution to the threshold calculation is more significant than distances from other trajectory points. Hence, a weighted average calculation method is proposed.

Let a vector  $DV = \{dis_2, dis_3, ..., dis_{lt-1}\}$  is used to store the distances from all intermediate trajectory points to the baseline. Another vector  $RV = \{rate_2, rate_3, ..., rate_{l-1}\}$  denotes the proportion of each distance to the total. The calculation process for each proportional value is outlined as follows,

$$
Dis = \sum_{i=2}^{h-1} dis_i,
$$
  
\n
$$
rate_i = \frac{dis_i}{Dis}, i = 2, 3, ..., h-1
$$
\n(14)

where *Dis* represents the cumulative distance from the *i*-th trajectory point to the baseline. *disi* and *ratei* represent the distance and proportional weight of the *i-*th trajectory point to the baseline, respectively.

The threshold calculation based on the weighted average can be expressed by,

$$
threshold = \sum_{i=2}^{h-1} dis_i \times rate_i, i = 2, 3, ..., h-1
$$
\n(15)

$$
(15)
$$

The weighted average calculation method effectively addresses the issue of direct average calculation. However, it also exposes another limitation. This arises when the distance from consecutive  $n (n \geq 3)$  points to the baseline is identical, and the proportion is the largest. This scenario is exemplified in [Fig. 8](#page-12-0) (b) by points  $P_6$ ,  $P_7$ , and  $P_8$ . These points form a line segment, and according to the compression principle of the DP algorithm, point  $P<sub>7</sub>$  becomes redundant and can be removed. If the weighted average calculation method is employed, the distance from  $P<sub>7</sub>$  to the baseline also needs to be included, carrying a substantial weight. Consequently, this could result in significant errors in the calculated threshold. To effectively address these issues, this paper incorporates SOG and COG in threshold calculation.

When a vessel navigates in a straight line, even if the SOG changes, retaining the starting and ending points is sufficient to accurately capture the distribution characteristics of the trajectory during compression tasks. Conversely, if a trajectory point quickly deviates from the baseline in a short period, it is highly likely to be a feature point. To address this, the approach taken in this paper involves establishing a coordinate system using the trajectory baseline as the *x*-axis. This system decomposes the velocity of each trajectory point, as depicted visually in [Fig. 8](#page-12-0) (c). A substantial change rate of the SOG component perpendicular to the *x*-axis between two points suggests a rapid deviation from the baseline. It is worth noting that COG assists SOG in its decomposition within this novel coordinate system.

According to the above analysis, the deviation of each intermediate trajectory point from the baseline and the change rate of SOG along the *y*-axis in the new coordinate system jointly determine whether it qualifies as a feature point. Thus, this paper combines these two aspects of information to calculate the compression threshold, as illustrated in [Fig. 9](#page-13-0). The core concept involves determining the distance ratio from each intermediate point to the baseline concerning the sum of all distances. Concurrently, the proportion of the change rate of SOG along the *y*-axis component for each intermediate point is calculated based on the overall change rate. The contribution value of the point to the compression threshold can be obtained by fusing the above two proportional weights and multiplying them with the distance to the baseline. The specific steps of the compression threshold computation are outlined as follows,

Step 1. *Offset weight calculation*. As shown in [Fig. 9](#page-13-0) (a), this step utilizes Eq. [\(13\)](#page-12-0) to calculate the distance between each intermediate point and the baseline, which is then stored in vector *DV*. Furthermore, Eq. [\(14\)](#page-13-0) is used to obtain a vector *RV* containing the proportion of each intermediate point's distance to the baseline in relation to the total.

Step 2. *SOG variation weight calculation*. Following [Fig. 9](#page-13-0) (b), the SOG of each trajectory point is decomposed in a new coordinate system, which is established with the trajectory baseline as the *x*-axis. In particular, this paper exclusively selects the SOG component along the *y*-axis to compute the compression threshold because the speed at which a trajectory point deviates from the baseline to some extent determines whether the point is a feature point. The decomposition of SOG requires the assistance of COG. However, it is unfeasible to employ the original COG directly because COG needs to be converted into the new coordinate system. The new coordinate system can be obtained by rotating the old coordinate system clockwise or counterclockwise by a certain degree  $\alpha$  ( $0 \le \alpha \le 180$ ) around the trajectory point. The original COG add or subtract *α* can get the heading value in the new coordinate system. Hence, it is a pivotal step to calculate the angle between the trajectory baseline and the *x*-axis of the original coordinate system. The details of the calculation process are thoroughly outlined in Algorithm 1.

#### **Algorithm 1:** Calculation of included angle



For ease of calculation, this paper uniformly establishes that the new coordinate system is achieved by rotating the existing coordinate system counterclockwise by *α* degrees. The components of SOG along the *y*-axis in the new coordinate system can be derived based on Algorithm 2.

**Algorithm 2:** Calculation of SOG components along the *y*-axis

**Input:** *SOG*, *COG*, *α* // The definition of *COG* in the original coordinate system is the angle between it and the positive half-axis of the *y*-axis, and its value range is [0*,* 360).

**Output:** *SOGY* 

<sup>1.</sup> **if** 0⩽*α*⩽90 **then** 

<sup>2.</sup> **if**  $COG = 0$  **then**  $SOGY = SOG \times \sin(|90 - \alpha| \times \frac{\pi}{180})$ ) ;

**Algorithm 2:** Calculation of SOG components along the *y*-axis

3. **else if**  $0 < COG \le 90$  then  $SOGY = SOG \times \sin\left(\left(\frac{90 - COG}{9}\right) - \alpha\right) \times \frac{\pi}{180}$ ) ; 4. **else if**  $90 < COG \le 180$  **then**  $SOGY = -SOG \times \sin(((COG - 90) + \alpha) \times \frac{\pi}{1800})$ ) ; 5. **else if** 180 *< COG <* 270 **then**  6. *cogConvert* = 270 – *COG*; 7. **if**  $\cos$ Convert ==  $\alpha$  **then**  $\text{SOGY} = 0$ ; 8. **else if**  $\cos$ *Convert*  $>$  *a* **then** 9.  $\text{SGGY} = -\text{SGS} \times \sin((\text{cogConvert} - \alpha) \times \frac{\pi}{180})$ ) ; 10. **else if** *cogConvert < α* **then**  11.  $\text{SGGY} = \text{SGG} \times \sin\left((\alpha - \text{cogConvert}) \times \frac{\pi}{180}\right)$ ) ; 12. **end if**  13. **else if**  $COG = 270$  **then**  $SOGY = SOG \times \sin\left(\alpha \times \frac{\pi}{180}\right)$ ) ; 14. **else if**  $270 < \text{COG} < 360$  then  $\text{SOGY} = \text{SOG} \times \sin\left(\left(\frac{(\text{COG} - 270)}{180}\right) + \alpha\right) \times \frac{\pi}{180}$ 15. **end if**  16. **else if**  $90 < a < 180$  **then** 17. **if**  $COG == 0$  **then**  $SGGY = SOG \times \sin((90 - (180 - \alpha)) \times \frac{\pi}{180})$ ) ; 18. **else if**  $0 < COG \leq 90$  **then** 19.  $SOGY = SOG \times \sin\left(\left(\frac{90 - COG}{+ (180 - \alpha)}\right) \times \frac{\pi}{180}\right)$ ) ; 20. **else if** 90 *< COG* ≤ 180 **then**  21.  $coeConvert = COG - 90$ 22. **if**  $\cos$ Convert == (180 –  $\alpha$ ) **then** 23.  $SOGY = 0;$ 24. **else if**  $\cos$ Convert  $\lt$  (180 –  $\alpha$ ) **then** 25.  $\text{SOGY} = \text{SOG} \times \sin\left(\left(\frac{180 - \alpha}{\text{O}}\right) - \text{cogConvert}\right) \times \frac{\pi}{180}$ ) ; 26. **else if**  $\cos$ Convert > (180 –  $\alpha$ ) **then** 27.  $\text{SOGY} = -\text{SOG} \times \sin((\text{cogConvert} - (180 - \alpha)) \times \frac{\pi}{180})$ ) ; 28. **end if**  29. **else if** 180 *< COG <* 270 **then**  30.  $SOGY = -SOG \times \sin\left(\left(\frac{180 - a}{+90 - (270 - COG)}\right)\right) \times \frac{\pi}{180}$ ) ; 31. **else if**  $COG = 270$  **then**  $SOGY = -SOG \times \sin((180 - \alpha) \times \frac{\pi}{180})$ );<br>); 32. **else if** 270 *< COG <* 360 **then**  33. *cogConvert* = *COG* − 270; 34. **if**  $\cos \theta$ *convert* ==  $(180 - \alpha)$  **then** 35.  $SOGY = 0$ ; 36. **else if**  $\cos$ Convert  $\lt$  (180 –  $\alpha$ ) **then** 37.  $\text{SGGY} = -\text{SG} \times \sin\left(\left(\frac{180 - \alpha}{\text{Cov}}\right) - \text{cogConvert}\right) \times \frac{\pi}{180}$ ) ; 38. **else if** *cogConvert >* (180 − *α*) **then**  39.  $\text{SOGY} = \text{SOG} \times \sin\left((\text{cogConvert} - (180 - \alpha)) \times \frac{\pi}{180}\right)$ ) ; 40. **end if**  41. **end if**  42. **end if** 

According to Algorithm 2, the SOG component along the *y*-axis for each trajectory point can be computed and stored in the vector  $SGGYV = \{SOGY_1, SOGY_2, \ldots, SOGY_{lt}\}$ . To determine the SOG change rate of each trajectory point along the *y*-axis, the timestamp of each trajectory point is retained in vector  $TV = \{t_1, t_2, ..., t_t\}$ . In particular, the time information in AIS data is expressed in hours, minutes, and seconds. To facilitate calculation, this paper adopts Eq. (16) to convert time into seconds uniformly as follows,

) ;

$$
t = 3600 \times dataH + 60 \times dataM + dataS \tag{16}
$$

where *dataH*, *dataM*, and *dataS* represent the hour, minute, and second in the timestamp, respectively. The velocity change rate along the *y*-axis at each intermediate point can be obtained based on vectors *SOGYV* and *TV* below.

$$
sogyC_{i+1} = \frac{SOGY_{i+1} - SOGY_i}{t_{i+1} - t_i}, i = 1, 2, ..., lt - 2
$$
\n
$$
(17)
$$

where *i* represents the *i-*th value in two vectors. *lt* denotes the total number of points in the trajectory. The velocity change rate component at each intermediate point is stored in the vector  $SGYCV = \{sogyC_2, sogyC_3, ..., sogyC_{lt-1}\}.$ 

In addition, the proportion of the change rate of SOG along the *y*-axis component for each intermediate point to the total change rate is determined by Eq. [\(18\)](#page-17-0) and expressed as a vector  $SOGYCRV = {sogyCR_2, sogyCR_3, ..., sogyCR_{l-1}}$ .

<span id="page-16-0"></span>

Kernel<sub>7</sub>



trajectory has feature points

Determine whether the trajectory has completed compression calculation

<span id="page-17-0"></span>

$$
Sum = \sum_{2}^{h-1} sogyC_i,
$$
  
\n
$$
sogyCR_i = \frac{sogyC_i}{Sum}, i = 2, 3, ..., h-1
$$
\n(18)

Step 3. *Weight fusion calculation*. The objective is to integrate the offset weights calculated in Steps 1 and 2 with the SOG change weights. The contribution value of each intermediate point to the threshold, essentially a weighted result, is determined by multiplying its fusion weight with the distance from the baseline. In particular, if a certain intermediate point significantly contributes to the threshold, the calculated fusion weight value will be higher. The final threshold is achieved by summing the contribution values of all intermediate points, as expressed by the following function.

$$
threshold = \sum_{i=2}^{l-1} dis_i \times sogyCR_i \times rate_i
$$
\n(19)

where *disi* denotes the distance from the *i-*th trajectory point to the baseline. *sogyCRi* is the change weight of the *i-*th trajectory point along the *y*-axis SOG component. *rate<sub>i</sub>* represents the offset weight of the *i*-th trajectory point relative to the baseline.

The purpose of multiplying two weights is to exclude the contribution of a factor if it evaluates to zero during threshold calculation. In such cases, the distance from that particular point to the baseline does not play a role in the threshold calculation. The ADPSC algorithm employs this calculated threshold in iterative searching for feature points, thereby realizing the trajectory compression task.

<span id="page-18-0"></span>

**Fig. 11.** Coalesced global memory access of AIS data.



**Fig. 12.** Visual comparison of trajectory data (or vector) and images (or matrix) in parallel computing framework design. (a) and (b) depict the thread distribution architecture for parallel computing in image and trajectory data, respectively.

#### *4.3. GPU-accelerated ADPSC compression method*

The ADPSC algorithm proposed in this paper entails an extended compression threshold calculation process compared to the original DP algorithm. It means that compressing massive vessel trajectory data demands more time. To align with real-world applications, this paper enhances the ADPSC algorithm for large-scale trajectory data compression within the GPU parallel computing framework, substantially diminishing execution time. The ADPSC algorithm predominantly encompasses two calculation processes during trajectory data compression: threshold and compression calculation. Accordingly, the designed parallel algorithm is structured around these two calculation components, as illustrated in [Fig. 10](#page-16-0). The threshold and compression calculation in the parallel ADPSC algorithm consists of multiple different functional functions, respectively. In the programming framework of CUDA, each function is encapsulated into a kernel, constituting a CUDA parallel computing function executed on the GPU. Each thread within the GPU conducts the kernel function in parallel to process the trajectory points to accomplish compression. Specifically, the operational logic of the kernel functions numbered 1 to 7 in [Fig. 10](#page-16-0) follows a progressive rather than a random execution order. The necessary notations are listed in [Table 4.](#page-17-0)

When executing parallel algorithms on a GPU, it cannot directly obtain data from memory and needs to transfer the data from memory to video memory. This transfer process is time-consuming. If vessel trajectories in the dataset are compressed one by one in sequence, although it might simplify the design of parallel algorithms, it would still demand substantial computational time. To tackle this challenge, this paper uniformly stores all trajectory data (i.e., time stamp, longitude, latitude, SOG, and COG) in a matrix as represented in Eq. [\(5\)](#page-7-0). This approach allows copying all necessary data from memory to video memory in a single operation, as displayed in Fig. 11. This helps avoid frequent data copying between memory and video memory, thereby reducing computational costs. In practical parallel computing, it is necessary to set a label vector to determine whether a trajectory point belongs to a specific vessel. [Sections 4.3.1 and 4.3.2](#page-19-0) explain the threshold and compression parallel computing processes within the ADPSC algorithm, respectively.

#### <span id="page-19-0"></span>*4.3.1. Parallelization of the threshold calculation*

Each point in the vessel trajectory is discretely distributed, and they exhibit independence from one another. This feature provides convenience for processing each trajectory point by using parallel compression algorithms. Each trajectory point can be systematically linked to threads within the GPU, with each thread executing the compression task by activating the kernel function. GPUs are renowned for their potent computational capabilities, often employed in image processing (or matrix calculation). Parallel algorithms devised for image processing adapt the distribution of threads in GPUs based on the matrix size, as depicted in [Fig. 12](#page-18-0) (a). The thread arrangement in the parallel algorithm of trajectory data differs from that of image processing, assuming a vector-oriented layout illustrated in [Fig. 12](#page-18-0) (b). In particular, the total thread count equals the sum of all points in the vessel trajectory dataset.

According to [Fig. 10](#page-16-0) (a), the parallel calculation of the threshold mainly consists of five functions, which are encapsulated within five kernels. Each thread executes these five kernel functions in turn to obtain the compression threshold of each vessel trajectory. The parallel computing process of these five kernel functions in each thread is as follows:

(1) *Kernel1*. Its purpose is to calculate the SOG components decomposed along the *y*-axis for each trajectory point in the new coordinate system. While all trajectory points are distributed in each thread of the GPU for parallel computing tasks, this presents two challenges. One is the inability to discern the relationship between trajectories and their respective points. Another issue is the inability to match the starting and ending points of trajectories.

To address these two challenges, this paper sets up two label vectors, *Ltpr* and *Lntp*. They are designed to determine the relationship between points and trajectories and identify the starting and ending points in trajectories. For illustration, suppose there are three vessel trajectories in a dataset with track point counts of 4, 5, and 6, respectively. The values of vectors *Ltpr* and *Lntp* would be [1, 1, 1, 1, 2, 2, 2, 2, 2, 3, 3, 3, 3, 3, 3] and [0, 4, 9, 15], respectively. The length of vector *Ltpr* is the total number of all trajectory points in the dataset, with each value indicating the specific trajectory to which a point belongs. The value in vector *Lntp* is the accumulation of the number of points in each trajectory. According to the above example analysis, the starting and ending points of the first trajectory are in the 0th and (4–1)th threads.

Based on Algorithm 3, the parallel calculation results of SOG decomposition along the *y*-axis for each trajectory point can be obtained. In particular, the function 'cauSogDecompose' integrates the abilities of Algorithm 1 and Algorithm 2.

#### **Algorithm 3:** Parallel computing of SOG decomposition

**Input:** *AISM*, *Ltpr*, *Lntp, sogDecL* // According to Eq. [\(5\)](#page-7-0), the first to fifth rows of the *AISM* matrix represent time, longitude, latitude, SOG, and COG, respectively. *sogDecL* is the initialization vector, whose length equals the total number of trajectory points in the dataset. All values in the vector *sogDecL* are 0. **Output:** *sogDecL* // A vector stores the SOG decomposition results of each trajectory point.

1. *thd* = *blockIdx.x*  $\times$  *blockDim.x* + *threadIdx.x*; // thread number.

2. *posS* = *Lntp*[*Ltpr*[*thd*] – 1]; // The starting position of the trajectory where the current point being processed by the current thread is located.

3. *posE* = *Lntp*[*Ltpr*[*thd*]] – 1; // The ending position of the trajectory where the current point being processed by the current thread is located.

4. *lonS* = *AISM*[1,:][*posS*]; // Longitude of starting point.

5. *latS* = *AISM*[2,:][*posS*]; // Latitude of starting point.

6. *lonE* = *AISM*[1,:][*posE*]; // Longitude of ending point.

7. *latE* = *AISM*[2,:][*posE*]; // Latitude of ending point.

8. *sogData* = *AISM*[3,:][*thd*];

9. *cogData* = *AISM*[4,:][*thd*];

10. *sogDecL*[*thd*] = cauSogDecompose(*lonS*, *latS*, *lonE*, *latE*, *sogData*, *cogData*);

(2) *Kernel2*. This kernel function is used to calculate the distance between each trajectory point and the baseline. During the execution of parallel computing, vectors *Ltpr* and *Lntp* remain essential to determine the relationship between points and trajectories, as well as to identify the starting and ending points. The parallel calculation process is described in Algorithm 4.

**Algorithm 4:** Parallel computing of distance from trajectory point to baseline

**Input:** *AISM*, *Ltpr*, *Lntp*, *DisL* // *DisL* is the initialization vector whose all values are 0.

**Output:** *DisL* // A vector stores the distance from each trajectory point to the baseline.

4. **if** *thd*!= *posS* && *thd*!= *posE* **then** // The starting and ending points in each trajectory do not participate in the calculation.

5. *lonS* = *AISM*[1,:][*posS*];

6. *latS* =  $AISM[2,1]$ [*posS*];

7. *lonE* = *AISM*[1,:][*posE*];

8. *latE* = *AISM*[2,:][*posE*];

9. *lonData = AISM*[1,:][*thd*]; // Longitude of current trajectory point.

- 10. *latData* = *AISM*[2,:][*thd*]; // Latitude of current trajectory point.
- 11. *disData* = cauDistance(*lonS*, *latS*, *lonE*, *latE*, *lonData*, *latData*);<br>12. *posData* = *thd* (*Ltpr*[*thd*]  $\times$  2 1); // Calculate where the da
- 12. *posData* = *thd*  (*Ltpr*[*thd*] × 2 1); // Calculate where the data (or distance) is stored in the vector *DisL*.

13. *DisL*[*posData*] = *disData*;

14. **end if** 

Algorithm 4 reflects the parallel calculation process of the point-to-baseline distance for each trajectory. The function 'cauDistance' integrates the ability of [Eq. \(13\).](#page-12-0) The length of the vector *DisL* is not equal to the total number of all trajectory points in the dataset. This

<sup>1.</sup>  $thd = blockIdx.x \times blockDim.x + threadIdx.x;$ 

<sup>2.</sup> *posS* = *Lntp*[*Ltpr*[*thd*] – 1];

<sup>3.</sup>  $posE = Lntp[Ltnr[thd]] - 1$ :

discrepancy arises primarily because the distance calculation from the trajectory point to the baseline excludes both the starting and ending points. The formula for calculating the length of the vector *DisL* is,

# *length* = *poNum* − 2 × *traNum* (20)

where *poNum* represents the total number of trajectory points in the dataset, and *traNum* denotes the number of trajectories.

(3) *Kernel3*. Its core idea is to calculate the SOG change rate of the intermediate point of each trajectory in the *y*-axis direction. The results of the first kernel function (*sogDecL*) are utilized in the calculation process. A detailed description of the parallel computing method is shown in Algorithm 5.

**Algorithm 5:** SOG change rate along the *y*-axis component



Based on Algorithm 5, the SOG change rate along the *y*-axis at all intermediate points of each trajectory can be quickly calculated. Each trajectory's starting and ending points do not participate in calculating the SOG change rate, so the length of vector *sogyCL* is the same as that of vector *DisL*.

(4) *Kernel4*. This kernel serves two primary purposes. Firstly, it calculates the cumulative distance of all intermediate points of each trajectory from the baseline. Secondly, it determines the collective SOG change rates along the *y*-axis for all intermediate points within each trajectory. Unlike previous kernel functions, where the focus was individual points, this kernel targets entire trajectories, with each trajectory in the dataset being sequentially mapped to the GPU threads. The parallel computation methodology is detailed in Algorithm 6.

### **Algorithm 6:** Calculate the sum of distance and SOG component change rate separately

**Input:** *DisL*, *sogyCL*, *Lntp*, *sumDisL*, *sumSogyCL* // *sumDisL* and *sumSogyCL* are the initialization vectors whose all values are 0.

**Output:** *sumDisL, sumSogyCL* // Two vectors store the sum of the distances from all intermediate points of each trajectory to the baseline and the sum of the SOG change rates of all intermediate points along the *y*-axis, respectively.

1. *thd* = *blockIdx.x*  $\times$  *blockDim.x* + *threadIdx.x*;

2.  $posDataS = Lntp[thd] - 2 \times thd;$ 

- 3. *posDataE* = *Lntp*[*thd* + 1] (2 × *thd* + 2); // Steps 2 and 3 calculate the boundary positions of two vectors (i,e., *DisL* and *sogyCL*) to match the relationship between the data and the trajectory.
- 4. **for** *i* = *posDataS*: *posDataE* **do**
- 5.  $sumDist[thd] = sumDist[thd] + Dist[i];$
- 6. *sumSogyCL*[*thd*] = *sumSogyCL*[*thd*] + *sogyCL*[*i*];
- 7. **end for**

Algorithm 6 performs parallel computing tasks based on trajectories. Therefore, the lengths of vectors *sumDisL* and *sumSogyCL* equal the total number of vessel trajectories in the dataset.

(5) Kernel<sub>5</sub>. This kernel function calculates the offset weight and SOG component variation rate weight of trajectory points according to Eqs. [\(14\) and \(18\)](#page-13-0), respectively. It then integrates these two weights, using Eq. [\(19\),](#page-17-0) to determine the weighted distance of each trajectory point from the baseline, yielding the threshold. This kernel is the same as the fourth kernel because the object processed by each thread is the vessel trajectory. The detailed calculation process is presented in Algorithm 7.

**Algorithm 7:** Parallel computing of compression threshold

**Input:** *DisL*, *sogyCL*, *Lntp*, *sumDisL*, *sumSogyCL*, *thrL* // *thrL* is the initialization vector whose all values are 0.

**Output:** *thrL* // A vector stores the compression threshold for each trajectory.

<sup>1.</sup> *thd* = *blockIdx.x*  $\times$  *blockDim.x* + *threadIdx.x*;

<sup>2.</sup>  $posDataS = Lntp[thd] - 2 \times thd;$ 

<sup>3.</sup> *posDataE* = *Lntp*[*thd* + 1] – (2  $\times$  *thd* + 2); // The functions of steps 2 and 3 are consistent with those in Algorithm 6.

<sup>4.</sup> **for** i = posDataS: posDataE **do** 



Algorithm 7, similar to Algorithm 6, executes parallel computing tasks based on vessel trajectories. As a result, the size of vector *thrL*  corresponds to the total number of trajectories in the dataset. Following the parallel computing process mentioned above, the compression threshold for each vessel trajectory is determined. This underpins the parallel compression task for vessel trajectory data discussed in Section 4.3.2.

# *4.3.2. Parallelization of compression calculation*

The ADPSC algorithm utilizes the calculated compression threshold to iteratively search for trajectory feature points. It uses these feature points to form a new data sequence to replace the original trajectory, thereby completing the compression task. According to the execution principle of the compression algorithm, the parallel process of compressed computing mainly consists of two parts, shown in [Fig. 10](#page-16-0) (b). On the one hand, it calculates the distance from each trajectory point to the baseline of the matching trajectory segment. This calculation process is dynamic, as the newly generated feature points will repartition the trajectory. On the other hand, it compares the threshold with the distance from the trajectory point to the baseline to determine the feature points. Meanwhile, a condition must be set to terminate the calculation process to represent that the trajectory has been compressed. The reason is that the compression algorithm is a recursive process requiring a constraint to end the operation. According to [Fig. 10](#page-16-0) (b), the parallel calculation of the compression mainly consists of two kernel functions. Each thread executes these two kernel functions in sequence to obtain compressed trajectories. The parallel computing process of these two kernel functions in each thread is as follows,

(1) *Kernel6*. Its purpose is to calculate the distance from each trajectory point to the baseline of the matching trajectory segment and store the results in the vector *DisL*. The length of this vector is equal to the total number of all trajectory points in the dataset, whose all values are 0 as the initialization state. During each iteration of searching for new feature points, a portion of the values in *DisL* is constantly changing. The reason is that each trajectory generates new feature points that divide it into multiple different trajectory segments.

In the specific parallel computing process, four problems must be addressed to complete parallel computing tasks successfully. The initial challenge concerns matching the relationship between points and trajectories. In the parallel algorithm, each trajectory point is assigned to a separate thread for operation, yet the GPU cannot identify which trajectory a point belongs to. This paper solves this issue by setting a label vector *Lfp*, where each value in the vector can match the logical relationship between each point and the trajectory. For example, suppose there are three trajectories in the dataset. Each trajectory has 4, 5, and 6 points, respectively. The values of vector *Lfp* are  $[1, 1, 1, 1, 2, 2, 2, 2, 2, 2, 3, 3, 3, 3, 3]$ . The second issue is to determine which points are the feature points calculated by the algorithm. The vector *Lfp* is utilized to solve this issue. If a point is a feature point of the corresponding trajectory, then its corresponding value in vector *Lfp* is negative. Before the compression algorithm executes each trajectory, the starting and ending points are their initialization feature points. Hence, the values of label vector *Lfp* are [-1, 1, 1, -1, -2, 2, 2, -2, -3, 3, 3, 3, 3, 3, 3, -3] as an initial state. The third issue is to pinpoint the position of each trajectory's starting and ending points in *AISM*. To address the problem effectively, this paper sets up another label vector *Lntp*, whose value accumulates the number of points in each trajectory. Based on the above example, the values of vector *Lntp* is [0, 4, 9, 15]. The fourth issue is to identify whether a certain trajectory has completed the compression task and does not need to execute this kernel function. The vector *Lntp* will also assist in solving this problem. For instance, if the second trajectory mentioned above has already completed the compression task, and the other two trajectories still need to continue searching for new feature points. The values of vector *Lntp* is [0, 4, -9, 15]. The detailed calculation process will be introduced in the seventh kernel function. In summary, these four vectors will aid parallel algorithms in calculating the distance from the trajectory point to the baseline and then comparing it with the threshold to search for feature points iteratively.

Each trajectory will generate new feature points during continuous iteration, which will repartition the trajectory and form different trajectory segments. Different trajectory segments in each trajectory have their corresponding starting and ending points, which cannot be identified using the vector *Lntp*. The reason is that the vector *Lntp* can only recognize the positions of the starting and ending points in a trajectory and assist in determining whether the trajectory has completed the compression task. To calculate the baseline distance from each point to the corresponding trajectory segment, this paper uses double pointers to find the starting and ending points of the trajectory segment corresponding to each point, as shown in Algorithm 8.

**Algorithm 8:** Double-pointer search for starting and ending points

**Output:** *leftPos, rightPos* // *leftPos* and *rightPos*, respectively, represent the positions of the starting and ending points of a certain trajectory segment in *AISM*.

1. *leftPos* = *thd*  $-1$ :

**Input:** *Lfp*, *thd* // *thd* denotes the thread number and also represents the *thd-*th trajectory point in the dataset that the current thread is processing.

<sup>2.</sup> *rightPos = thd + 1; // leftPos and rightPos initialization.* 

<sup>3.</sup> *labelData* = *Lfp*[*thd*]; // Calculate the value corresponding to the trajectory point in the vector *Lfp*.

while Lfp[leftPos]!=  $-$ labelData || Lfp[rightPos]!=  $-$ labelData do // When labelData is negative, it indicates that the position of the pointer is at the boundary of the trajectory segment (starting or ending point).

<sup>5.</sup> **if** *Lfp*[*leftPos*]!= − *labelData* **then** *leftPos* = *leftPos* – 1;



**Algorithm 8:** Double-pointer search for starting and ending points

6. **end if** 

7. **if** *Lfp*[*rightPos*]!= − *labelData* **then** *rightPos* = *rightPos* – 1; 8. **end if**  9. **end while** 

According to the results of Algorithm 8, this paper can continue to calculate the distance from each trajectory point to the corresponding trajectory segment baseline in parallel, as shown in Algorithm 9.

**Algorithm 9:** Computing of distance from points to the trajectory segment baseline

**Input:** *AISM*, *Lfp*, *Lntp*, *DisL* 

**Output:** *DisL* 

1. *thd* = *blockIdx.x*  $\times$  *blockDim.x* + *threadIdx.x*;

2.  $lntpData = Lntp[abs(Lfp[thd])];$ 

3. **if** *lntpData >* 0 **then** // When *lntpData* is a positive number, it indicates that there are still feature points in the current trajectory, and it is necessary to continue calculating the distance between the intermediate point and the corresponding trajectory segment baseline.

4.  $labelData = Lfp[thd];$ 

- 5. **if** *labelData >* 0 **then**
- 6. *leftPos*, *rightPos* = cauStartEndPos(*Lfp*, *thd*);
- 7. *lonS* = *AISM*[1,:][*leftPos*]; // Longitude of starting point.
- 8. *latS* = *AISM*[2,:][*leftPos*]; // Latitude of starting point.
- 9. *lonE* = *AISM*[1,:][*rightPos*]; // Longitude of ending point.
- 10.  $latE = AISM[2, :] [rightPos]; // Latitude of ending point.$
- 11. *lonData* = *AISM*[1,:][*thd*]; // Longitude of current trajectory point.
- 12. *latData* = *AISM*[2,:][*thd*]; // Latitude of current trajectory point.
- 13. *DisL*[*thd*] = cauDistance(*lonS*, *latS*, *lonE*, *latE*, *lonData*, *latData*);
- 14. **else if** *labelData <* 0 **then**
- 15. *DisL*[*thd*] = 0; // When *labelData* is negative, it indicates that the current point is a feature point and the distance from the baseline is zero.<br>16. **end if**
- 16. **end if**
- 17. **end if**

Algorithm 9 reflects the detailed process of computing the distance between each trajectory point and the baseline of the corresponding trajectory segment in parallel. In particular, the functions 'cauStartEndPos' and 'cauDistance' respectively integrate the capabilities of [Eq. \(13\)](#page-12-0) and Algorithm 8.

(2) Kernel7. This kernel function plays two essential roles in parallel computing. One is to judge whether each trajectory segment has new feature points according to the results of the sixth kernel function. If a new feature point is generated, the value matching the point with the vector *Lfp* is changed from a positive number to a negative number. Another function is determining whether a certain trajectory has completed the compression task. When a certain trajectory has completed the compression task, the corresponding value in vector *Lntp* changes from positive to negative. Since the object of this kernel function is a trajectory, the threads in the GPU are mapped to each trajectory in the dataset one by one, which is different from the sixth kernel function. The parallel calculation process is shown in Algorithm 10.

#### **Algorithm 10:** Iterative search for feature points

**Input:** *Lfp*, *Lntp*, *DisL*, *thrL* // Vector *thrL* stores the compression threshold for each vessel trajectory.

**Output:** *Lfp*, *Lntp* 

1. *thd* = *blockIdx.x*  $\times$  *blockDim.x* + *threadIdx.x*;

2.  $$ 

- 3. **if** *lntpData >* 0 **then** // When *lntpData* is a positive number, it indicates that there are still feature points in the current trajectory which require further iterative search.
- 4. *ifFeature* = 0; // A variable used to determine whether new feature points are generated in iterative search. Its initial value is zero. When it becomes one during the calculation process, it indicates that new feature points have been generated in the iterative search.
- 5. *threshold = thrL*[*thd*]; // The compression threshold of the current trajectory.
- 6. *num* = 0; // It assists in searching for feature points in trajectory segments with an initial value of zero.
- 7.  $begin = 0;$
- 8.  $end = 0$ ; // begin and end are used to define the positions of the starting and ending points for trajectory segments, respectively. Their initial values are zero.
- 9. *maxDisData* = 0; // This variable defines the maximum value from a point to the baseline of a trajectory segment. Its initial value is zero.

10. **for**  $(i = Lntp[thd + 1] - 1; i > abs(Lntp[thd]) - 1; i-)$  **do** // Traverse search for feature points for each trajectory segment.

11.  $disData = Dist$ .

<sup>12.</sup> **if**  $\frac{d}{dx}$  *disData* == 0  $\& \& num == 0$  **then** // Locate the position of the ending point for a trajectory segment.<br>13.  $num = num + 1$ :

 $num = num + 1$ ;

<sup>14.</sup>  $end = i$ ;

<sup>15.</sup> **else if** *disData*!= 0 && *num* == 1 **then** // Calculate the maximum distance from all intermediate points to the baseline of the trajectory segment.

<sup>16.</sup> **if** *disData > maxDisData* || *disData* == *maxDisData* **then** 

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<span id="page-23-0"></span>

In executing parallel algorithms, each thread performs the sixth and seventh kernel functions repeatedly, which reflects that the algorithm is essentially a recursive process. The condition for recursive termination is that all values in the vector *Lntp* are negative. *DisL*, *Lfp*, and *Lntp* are three important vectors that assist parallel algorithms in finding feature points of trajectories through continuous iterations. The vectors *DisL* and *Lfp* length equal the total number of trajectory points. The size of vector *Lntp* is the total number of trajectories plus one. Before executing parallel compression tasks, three vectors need to be assigned initialization values, as illustrated in [Fig. 13](#page-24-0) (a). The sixth kernel function calculates the distance from all intermediate points to the baseline in parallel and updates the value in the vector *DisL* to store the distance, as shown in [Fig. 13](#page-24-0) (b). [Figs. 13](#page-24-0) (c), (d), and (e) reflect the process of two kernel functions iteratively searching for feature points and the changes of all values in the three vectors. When the vessel trajectories meet the termination conditions of the iteration, the final compression result is obtained, as illustrated in [Fig. 13](#page-24-0) (f). In particular, the trajectory points corresponding to negative values in vector *Lfp* are feature points. The changes of all values in the two label vectors before and after trajectories compression are shown in [Figs. 14](#page-25-0) (a) and (b), respectively.

#### **5. Experimental results and analysis**

To verify the effectiveness of the proposed adaptive and accelerated compression framework, this paper conducts experiments from two perspectives. First, it showcases the superior performance of the ADPSC algorithm in addressing vessel trajectory compression issues compared to the original DP algorithm, considering both qualitative and quantitative perspectives. Second, the acceleration ratio is employed to quantitatively evaluate the enhanced speed of the optimized parallel compression algorithm compared to the original serial algorithm. Concurrently, a GPU parallel computing framework is established as a requisite experimental environment. The required notations are offered in [Table 5. Table 6](#page-26-0) provides a detailed overview of the hardware and software environments.

#### *5.1. Datasets description*

This paper collects AIS data from three different research areas: Tianjin Port, Chengshan Jiao Promontory, and Caofeidian Port. The datasets are used to verify the efficacy of the proposed ADPSC compression algorithm and cover the period from 1 July 2020 to 30 September 2022. The distribution and density visualization effects of vessel trajectories in the three regions are illustrated in [Figs. 15](#page-25-0) (a), (b), and (c), respectively. Meanwhile, the statistical information related to the above research areas is shown in [Table 7](#page-26-0). It is well known that vessel trajectory data can be affected during transmission through base stations and satellites, resulting in noisy data. Thus, high-quality trajectory data can be obtained from the newly collected original data after preprocessing in [Section 3.3,](#page-9-0) which can be used for subsequent comparative experiments.

<span id="page-24-0"></span>

**Fig. 13.** Procedures of the proposed GPU-based ADPSC parallel compression of vessel trajectory. (a) original vessel trajectory marked with labels *DisL*, *Lfp*, and *Lntp*, (b) parallel computation of *dis*, (c) searching for the maximal *dis* to count the feature trajectory points, (d) segmenting trajectory and updating the new feature points, (e) iteratively finding the feature trajectory points, and (f) obtaining the final compressed vessel trajectory.

#### *5.2. Compression evaluation metrics*

To provide a quantitative assessment of the proposed ADPSC algorithm in performing vessel trajectory data compression tasks compared to the original DP algorithm, this paper selects Compression Ratio (CR), Rate of Length Loss (RLL), and DTW as key performance evaluation metrics. In particular, when the CR is high, and both the RLL and DTW are low, it signifies optimal compression performance. Concurrently, the Speedup Ratio (SR) is employed to quantitatively evaluate the acceleration efficiency of the ADPSC algorithm when compressing extensive vessel trajectory data in the GPU parallel computing framework.

# *5.2.1. Compression ratio*

CR is a universal and standard indicator to measure compression efficacy, particularly in illustrating variations in the number of

<span id="page-25-0"></span>

**Fig. 14.** Visual illustration of changes in two critical labels (i.e., *Lfp* and *Lntp*) before and after performing compression tasks. (a) initialization status of labels *Lfp* and *Lntp* before compression, and (b) data changes in labels *Lfp* and *Lntp* after completing the compression task.



**Fig. 15.** Density visualisation of realistic vessel trajectory datasets: (a) Tianjin Port, (b) Chengshan Jiao Promontory, and (c) Caofeidian Port.

trajectory points. A higher CR value signifies removing a larger number of trajectory points. Its formula is defined as,

$$
CR = \left(1 - \frac{N_{com}}{N_{ori}}\right) \times 100\%
$$
\n(21)

where *N<sub>com</sub>* and *N<sub>ori</sub>* represent the number of points in all compressed and original trajectories, respectively.

#### *5.2.2. Rate of length loss*

RLL can reflect the length loss of the vessel's trajectory before and after compression. A lower RLL value suggests minimal distortion in the compressed trajectory, ensuring optimal compression results. The formula for RLL is given by

$$
RLL = \frac{\sum_{i=1}^{I} |T_{ori}^{i}| - \sum_{i=1}^{I} |T_{com}^{i}|}{\sum_{i=1}^{I} |T_{ori}^{i}|}
$$
(22)

where *I* represents the total number of vessel trajectories in the dataset.  $|T_{ori}^i|$  $|$  and  $|T^i_{com}$ ⃒ ⃒ are the lengths of the *i-*th original and compressed trajectories, respectively.

<span id="page-26-0"></span>

#### **Table 6**

Hardware and software environments.

Hardware	Model	Software	Version
<b>CPU</b>	i7-12700KF	Python	3.8.3
	Dodeca Core		
<b>Host Memory</b>	32GB	<b>CUDA</b>	11.7
GPU	GTX 3080	MySQL	8.0
Global Memory	12GB	$\sim$	

#### **Table 7**

Statistical information related to three different regions.



# *5.2.3. Dynamic time warping*

DTW is a prevalent measurement technique adept at calculating the similarity (inversely proportional to distance) between two sequential data [\(Lahreche and Boucheham, 2021\)](#page-38-0). Its core idea is to find the minimum distance between two sequence data using the dynamic programming method. In this paper, DTW serves as a quantitative evaluation method to assess the similarity between original and compressed trajectory data [\(Li et al., 2020](#page-38-0)). [Huang et al. \(2020\)](#page-38-0) also used DTW as an evaluation indicator when investigating compression algorithms for vessel trajectories. A smaller DTW value between two trajectories suggests that the compression algorithm introduces minimal distortion post-processing, indicating commendable compression efficacy.

A matrix *M* of size *O* × *C* is created, where each element *Ma,b* along the warping path refers to the cumulative distance between the *a*-th point  $P_{ori}^a$  in the original trajectory and the *b*-th point  $P_{com}^b$  in the compressed trajectory. Meanwhile, let *Q* represent the warping path between  $T_{ori}$  and  $T_{com}$ , essentially a sequence  $Q = \{q_1, q_2, q_3, \dots, q_L\}$  with  $q_l = (0,0] \in [1:0] \times [1:0]$ . The set of all potential warping paths can be represented by  $W_{OC}$ . The warping cost  $d_Q(T_{ori}, T_{com})$  of the *Q* can be defined as follows,

<span id="page-27-0"></span>

**Fig. 16.** The display of a new index for evaluating compression effectiveness, (a) shows the morphology of the original (T<sub>ori</sub>) and compressed (T<sub>com</sub>) trajectories, respectively; (b) reflects the process of using the optimized measurement method to calculate the similarity values of two trajectories before and after compression.

$$
d_Q(T_{\text{ori}}, T_{\text{com}}) = \sum_{l=1}^{L} d(P_{\text{ori}}^{Ol}, P_{\text{com}}^{Cl})
$$
\n(23)

where *L* represents the length of sequence *Q*, and its range is  $\max(O, C) \le L \le O + C$ .  $d(\cdot, \cdot)$  is the squared Euclidean distance. The DTW metric between  $T_{\text{ori}}$  and  $T_{\text{com}}$  related to the minimum warping cost is given as follows:

$$
DTW(T_{ori}, T_{com}) = \sqrt{d_{Q^*}(T_{ori}, T_{com})} = \min\left\{\sqrt{d_Q(T_{ori}, T_{com})} | Q \in W_{OC}\right\}
$$
\n(24)

where *Q*∗ is the minimum warping cost. Meanwhile, each element in *M* can be calculated as

$$
M_{a,b} = d\left(P_{ori}^a, P_{com}^b\right) + \min\{M_{a-1,b-1}, M_{a-1,b}, M_{a,b-1}\}\tag{25}
$$

The original DTW method attempts to match all points in the original trajectory with those in the compressed trajectory to determine the best warping path and calculate the similarity value. However, this approach introduces significant errors and inaccuracies in assessing the similarity between the original and compressed trajectories due to the considerable difference in the number of points between them. For example, if a vessel's trajectory forms a straight line and the compressed result only retains the starting and ending points, as illustrated in Fig. 16 (a), the compressed trajectory effectively reflects the original trajectory's distribution characteristics. Consequently, one might expect the similarity measure between these two trajectories to be zero. However, due to the reduction in the number of compressed trajectory points compared to the original trajectory, the similarity measure is not zero, as indicated by the DTW calculation process described earlier.

To address this issue, further optimization of the DTW measurement approach is proposed. The essence of this optimization lies in gauging the compression quality by examining the similarity in distances of each point in the original and compressed trajectories to the baseline. This baseline represents a line connecting the trajectory's starting and ending points, as shown in Fig. 16 (b).

To further elaborate on the computational process of the newly proposed measurement method, this paper breaks it down into three steps.

(1) *Distance calculation*. Eq. [\(13\)](#page-12-0) is employed to calculate the distance from each point in the original and compressed trajectories to the baseline, respectively. This process yields two distinct distance vectors: *Vori* for the original and *Vcom* for the compressed trajectory.

(2) *Distance symbol judgment*. The core of this step lies in discerning the sign (positive or negative) of values within the two distance vectors, *Vori* and *Vcom*. As illustrated in Fig. 16 (b), this paper characterizes the positional relationship of trajectory points relative to the baseline by allocating a positive value to the distance of points above the baseline and a negative value to those below. Concurrently, if a trajectory point coincides with the baseline, it is assigned a value of zero within the vector. In specific judgments, the coordinates of the starting and ending points of the trajectory are used to generate the equation of the baseline as  $y = kx + b$ . The calculation method for *k* (slope) and *b* (intercept) is shown in Eqs. (26) and (27),

$$
k = \frac{lat_E - lat_S}{lon_E - lon_S} \tag{26}
$$

$$
b = lat_E - \frac{lat_E - lat_S}{lon_E - lon_E} \times lon_E
$$
\n(27)

Subsequently, the intermediate trajectory point coordinates (*lon*, *lat*) are brought into the linear equation. If inequality *lat* − *k* × *lon* − *b* > 0 is satisfied, the distance between the trajectory point and the baseline is denoted as positive. In contrast, if inequality  $lat - k \times lon - b < 0$  prevails, the distance value is considered negative. An exception arises when *lon<sub>E</sub>* and *lon<sub>S</sub>* are equal,

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**Fig. 17.** Visual illustration of the compression threshold distribution calculated by the ADPSC algorithm in three research areas: (a) Tianjin Port, (b) Chengshan Jiao Promontory, and (c) Caofeidian Port.

# **Table 8**

The average and standard deviation results of the ADPSC algorithm for calculating compression threshold in three water areas.





**Fig. 18.** Visualisation of original and compressed (using ADPSC algorithm) vessel trajectories in three different water areas: (a) Tianjin Port, (b) Chengshan Jiao Promontory, and (c) Caofeidian Port.



<span id="page-29-0"></span>The evaluation results of all trajectories data compressed based on original DP (i.e., 0.1m, 0.5m, 1.0m, 5.0m, 10.0m, 20.0m, 50.0m, and 100.0m), other adaptive DP (i.e., ADP and ADPS) and our proposed ADPSC algorithms in Tianjin Port.

Note: ↑ indicates that a larger CR value corresponds to better compression performance. ↓ denotes that smaller values of RLL, DTW, and TP-DTW result in better compression performance.

# **Table 10**

The evaluation results of all trajectories data compressed based on original DP (i.e., 0.1m, 0.5m, 1.0m, 5.0m, 10.0m, 20.0m, 50.0m, and 100.0m), other adaptive DP (i.e., ADP and ADPS) and our proposed ADPSC algorithms in Chengshan Jiao Promontory.



implying the non-existence of *k*. For instance, the distance value is categorized as positive when the inequality  $\log \frac{1}{n}$  /  $\log \frac{1}{n}$  holds true. Conversely, it is designated as negative.

(3) *Measurement results*. Based on the calculation results in step (2), the DTW method is used to compute the similarity between the newly obtained vectors,  $V_{ori}$  and  $V_{com}$ , resulting in the final measurement result.

This new measurement method is called Trajectory Points-based DTW (TP-DTW). The conventional similarity calculation is rooted in direct measurements between two trajectories. This original method is designated as DTW herein. [Section 5.3](#page-30-0) delves deeper into the distinctions between DTW and TP-DTW in assessing compression efficacy.

# *5.2.4. Speedup ratio*

SR quantifies the acceleration capabilities of the proposed ADPSC algorithm in performing compression tasks under the GPU parallel computing framework compared to traditional computing methods. A notably higher SR value suggests a more enhanced acceleration yielded by this new computational framework. The underlying equation for SR is as follows,

$$
SR = \frac{T_{CPU}}{T_{GPU}}\tag{28}
$$

where *T<sub>CPU</sub>* and *T<sub>GPU</sub>* represent the execution time of the ADPSC algorithm when performing the same task under the CPU serial and GPU parallel computing framework, respectively.

In this paper, an essential aspect to take into account is that the computational expense of the CPU serial framework solely comprises the runtime of the ADPSC algorithm when compressing vessel trajectory data. On the other hand, the computing cost of the GPU parallel framework includes not only the running time of the algorithm but also the time of original data transmission from memory to video memory and the transmission of compressed data from video memory back to memory. If parallel algorithms lack optimal design, they will consume more computational time, especially when dealing with large-scale datasets. Therefore, in designing the GPU parallel algorithms, this paper accounts for the implications of data transfer on computation time, ensuring a more effective integration with the real-world application environment.

<span id="page-30-0"></span>The evaluation results of all trajectories data compressed based on original DP (i.e., 0.1m, 0.5m, 1.0m, 5.0m, 10.0m, 20.0m, 50.0m, and 100.0m), other adaptive DP (i.e., ADP and ADPS) and our proposed ADPSC algorithms in Caofeidian Port.





**Fig. 19.** Visual comparison of compression performance of four representative trajectories based on original DP (i.e., 0.1m and 100m), other adaptive DP (i.e., ADP and ADPS), and the proposed ADPSC algorithms in Tianjin Port.

#### *5.3. Compression performance of the ADPSC method*

The primary advantage of the proposed ADPSC algorithm in this paper lies in its ability to calculate compression thresholds based on the characteristics of different trajectory data automatically. [Fig. 17](#page-28-0) reflects the distribution of compression thresholds calculated based on the ADPSC algorithm for all vessel trajectories in the three research areas. The threshold distribution of the three regions mainly falls within the 1 m to 50 m range. In particular, the trajectory threshold distribution of Tianjin Port is between 1 m and 5 m, which accounts for the most significant proportion. According to the results in [Figs. 17](#page-28-0) (b) and (c), the threshold distribution of Chengshan Jiao Promontory and Caofeidian Port appear closely assigned, with the most common range being 10 m to 20 m. Concurrently, [Table 8](#page-28-0) shows the average and standard deviation of all trajectory compression thresholds in three regions. The standard deviations across the regions are consistent. Among them, Chengshan Jiao Promontory has the highest average, Tianjin Port is the lowest, and Caofeidian Port's average is close to that of Chengshan Jiao Promontory.

To provide a clear visual representation of the ADPSC algorithm, this paper projected the original and compressed trajectories of



**Fig. 20.** Visual comparison of compression performance of four representative trajectories based on original DP (i.e., 0.1m and 100m), other adaptive DP (i.e., ADP and ADPS), and the proposed ADPSC algorithms in Chengshan Jiao Promontory.

three regions onto the map, as shown in [Fig. 18.](#page-28-0) At a macro level, this visual comparison effectively illustrates that the distribution of compressed trajectories closely mirrors that of the original ones.

To further quantitatively evaluate the overall effectiveness of the new ADPSC compression algorithm on all trajectory data in each study area, this paper conducts comparative experiments with the original DP and two other classic adaptive compression algorithms, namely ADP ([Liu et al., 2019b\)](#page-38-0) and ADPS [\(Li et al., 2022](#page-38-0)). Notably, the ADP and ADPS algorithms, along with the ADPSC algorithm proposed in this paper, are essentially optimized based on the original DP algorithm. The advantage of the ADPSC algorithm compared to ADP and ADPS algorithms lies in simultaneous consideration of the SOG and COG of trajectory points, enabling more accurate preservation of feature points. Therefore, the comparative experiment between ADPSC and these two adaptive compression algorithms is highly valuable. This experiment accumulates the CR and RLL values of each trajectory to obtain the consolidated results in [Tables 9,](#page-29-0) [10, and 11.](#page-29-0) Additionally, the DTW and TP-DTW values of each trajectory are calculated to determine their average value and standard deviation. According to the results in [Tables 9, 10, and 11,](#page-29-0) setting a small threshold (e.g., 0.1 m and 0.5 m) in the DP algorithm yields high similarity between the compressed and original trajectories; however, the CR remains low, retaining redundant data. Conversely, an excessively large threshold (e.g., 50.0 m and 100.0 m) might result in a high CR, but the compressed trajectory will deviate significantly from the original distribution. Conclusively, the three adaptive compression algorithms overcome the conventional DP algorithm's limitations by effectively eliminating redundant data and mirroring the original trajectory's distribution characteristic.

Further examination of the compression effects between ADP, ADPS, and ADPSC reveals that while ADP and ADPS can compress more trajectory points compared to ADPSC, the trade-off is the destruction of the original trajectory's data structure. According to the

<span id="page-32-0"></span>

**Fig. 21.** Visual comparison of compression performance of four representative trajectories based on original DP (i.e., 0.1m and 100m), other adaptive DP (i.e., ADP and ADPS) and the proposed ADPSC algorithms in Caofeidian Port.





results in [Tables 9, 10, and 11,](#page-29-0) compared to ADPSC, ADP compresses trajectory data from three water areas, increasing CR by 13.7595 %, 6.9819 %, and 7.2871 %, respectively. Meanwhile, the growth rates of indicator RLL in three different water areas are 348.8745 %, 55.5300 %, and 114.2494 %, respectively. The growth rates of average DTW in three different water areas are 372.3583 %, 112.4785 %, and 80.0441 %, respectively. The growth rates of average TP-DTW in three different water areas are 478.5035 %, 102.2880 %, and 63.3042 %, respectively. Additionally, compared with ADPSC, ADPS compresses vessel trajectory data from three water areas,

The comparative evaluation of compression results of four representative trajectories based on original DP (i.e., 0.1m and 100m), other adaptive DP (i.e., ADP and ADPS), and the proposed ADPSC algorithms in Chengshan Jiao Promontory.



#### **Table 14**

The comparative evaluation of compression results of four representative trajectories based on original DP (i.e., 0.1m and 100m), other adaptive DP (i.e., ADP and ADPS), and the proposed ADPSC algorithms in Caofeidian Port.



increasing CR by 11.7251 %, 5.2574 %, and 6.1618 %, respectively. Regarding the three indicators of evaluating trajectory quality, the growth rates of indicator RLL in three different water areas are 267.2026 %, 37.7880 %, and 101.5267 %, respectively. The growth rates of average DTW in three different water areas are 275.5338 %, 97.5731 %, and 43.4611 %, respectively. The growth rates of average TP-DTW in three different water areas are 365.8212 %, 84.1827 %, and 56.7656 %, respectively.

Based on the comparison of the growth rates of the above four indicators in different compression algorithms, although ADP can compress more trajectory points than ADPS, it deletes more key feature points, thus altering the trajectory structure. This result is because ADP represents the compression threshold by calculating the average distance from each trajectory point to the baseline, which may blur key information, as analyzed in [Section 4.2](#page-12-0). Furthermore, the ADPSC algorithm proposed in this paper has a slightly lower CR value compared to the ADPS algorithm, but it offers the highest compressed data quality. In practical application scenarios, the compression quality of vessel trajectory data is paramount. Hence, the ADPSC algorithm outperforms other classic adaptive compression algorithms, such as DP, ADP, and ADPS, in vessel trajectory data compression tasks.

Further deepening the exploration, four representative trajectories with multiple feature points from each region are selected. These trajectories undergo compression using the original DP, two other adaptive compression algorithms (i.e., ADP and ADPS), and the optimized ADPSC algorithm. This allows for a direct comparison of their efficiency in handling trajectory compression. The chosen trajectories are specifically selected for their multiple feature points, providing a comprehensive representation of the predominant

<span id="page-34-0"></span>

**Fig. 22.** Comparison of two similarity measurement methods (i.e., DTW and TP-DTW) for evaluating compression performance: (a) Tianjin Port, (b) Chengshan Jiao Promontory, and (c) Caofeidian Port.

Comparison of storage space occupied before and after data compression. In particular, the stored data mainly includes timestamps, longitude and latitude coordinates, SOG, and COG.

Water areas	Storage size without compression (MB)	Storage size with compression (MB)
Tianjin Port	938.3889	154.0128
Chengshan Jiao Promontory	1735.2518	175.6007
Caofeidian Port	2673.6407	329.6135

distribution characteristics found in most trajectories. The compression effects of deploying the DP, ADP, ADPS, and proposed ADPSC algorithms on these select trajectories within the three areas are illustrated in [Figs. 19, 20, and 21.](#page-30-0) Unlike the broader overview presented in [Fig. 18](#page-28-0), these figures offer a more detailed, micro-level assessment of compression performance.

When the four representative trajectories of each research area are compressed using the DP algorithm, thresholds of 0.1 m and 100 m are selected and used for comparative experiments. As depicted in [Figs. 19, 20, and 21](#page-30-0), a larger threshold set by the DP algorithm for trajectory compression leads to a more distorted trajectory compared to its original version, failing to accurately reflect its inherent distribution characteristics. At a compression threshold of 0.1 m with the DP method, the trajectory retains features almost identical to the original, but as shown in [Tables 12, 13, and 14](#page-32-0), the CR value is notably low, indicating the presence of redundant data. In summary, developing adaptive compression algorithms to automatically calculate the compression threshold for each trajectory has practical application and research value.

On the contrary, the ADPSC algorithm is designed to automatically determine suitable thresholds based on the unique distribution attributes of each trajectory and its dynamic navigation details. This approach ensures that significant features from the original trajectory are retained while eliminating redundant data. Although ADP and ADPS algorithms can also automatically calculate the compression threshold for each trajectory and compress more trajectory points compared to the ADPSC algorithm, they incur the cost of losing the original structure of the trajectory, which is not advisable in practical applications. The visual representations in [Figs 19,](#page-30-0) [20, and 21](#page-30-0) qualitatively indicate that the ADP and ADPS algorithms sometimes fail to accurately preserve the feature points of vessels turning slightly, while the new ADPSC algorithm can achieve this. This implies that the trajectory compressed using the ADPSC algorithm is consistent with the original spatial distribution on the map. Additionally, [Tables 12, 13, and 14](#page-32-0) show that all the trajectories compressed via the ADPSC method achieve lower RLL and similarity metrics (either DTW or TP-DTW) while maintaining a superior CR value.

[Tables 12, 13, and 14](#page-32-0) preliminarily reflect the disparities between DTW and TP-DTW in evaluating compression performance. To

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**Fig. 23.** Accelerated effect of the proposed ADPSC compression algorithm based on GPU parallel computing framework, (a) the CPU (serial) execution times (average value + standard deviation), (b) GPU (parallel) execution times (average value + standard deviation), and (c) the speedup ratio between CPU and GPU. Specifically, the calculation time across ten iterations for each experimental scenario shows minimal variation, resulting in a very low standard deviation.

further explore the differences in assessing the compression performance of each trajectory using these two measurement methods, this experiment randomly selected 2000 trajectories from each study area. At the same time, the DTW and TP-DTW metrics of each trajectory after compression are drawn into a line chart, depicted in [Fig. 22.](#page-34-0)

According to the results displayed in the line chart, it becomes apparent that the novel TP-DTW metric consistently produces lower values than its DTW counterpart when calculating the similarity between the original and compressed trajectories. This is largely because traditional DTW when assessing the similarity between the two trajectories, tends to record enhanced measurements even with minor trajectory alterations (such as a reduction in trajectory points). In contrast, the TP-DTW approach focuses on assessing similarity by evaluating the deviations of points in both trajectories from a baseline. The advantage of the TP-DTW method over DTW is that it measures compression quality by checking the similarity between the distance from each point in the original trajectory and the baseline in the compressed trajectory. Specifically, the baseline represents a line connecting the trajectory's starting and ending points. In summary, if a compression algorithm retains all pivotal points in the compressed trajectory, the value derived from TP-DTW will be considerably lower than that from DTW. For instance, the trajectory depicted in [Fig. 16](#page-27-0) (a) retains all key points after compression, mirroring the original distribution characteristics. Theoretically, their similarity metric should stand at zero. While TP-DTW aligns with this theoretical estimate, DTW offers a different result. Conclusively, the newly proposed TP-DTW metric emerges as a more fitting tool for assessing compression efficacy.

The above analysis demonstrates the effectiveness of the ADPSC algorithm in compressing vessel trajectory data and its ability to eliminate redundant information in the dataset. To further quantitatively illustrate how these compression algorithms can economize a storage space, this experiment counted the storage size of three regional datasets before and after compression, with the specifics displayed in [Table 15.](#page-34-0) The trajectory data from Tianjin Port, Chengshan Jiao Promontory, and Caofeidian Port, when compressed using the ADPSC algorithm, led to savings of 784.3761 MB, 1559.6511 MB, and 2344.0272 MB, respectively. In summary, the newly proposed ADPSC algorithm in this paper excels in removing redundant data from trajectories, thereby substantially curtailing storage costs.

#### *5.4. Acceleration performance of GPU-based ADPSC method*

To verify the enhanced efficiency of the accelerated ADPSC compression algorithm in processing vast vessel trajectory data, this paper tests the ADPSC algorithm on trajectory data from three experimental areas by both CPU serial and GPU parallel computing frameworks. The execution time results are displayed in Figs. 23 (a) and (b), respectively. Moreover, each experimental scenario is executed ten times, and the average is calculated to minimize random variations. Fig. 23 (c) employs the indicator SR to visually demonstrate the significant speed advantage of the ADPSC algorithm under the GPU parallel computing framework compared with traditional CPUs. According to the statistical results in [Table 7,](#page-26-0) Caofeidian Port has 35,967 more vessel trajectories than Chengshan Jiao Promontory, with a discrepancy of 8,431,912 in the count of trajectory points. On the other hand, Tianjin Port records the least data compared to the other two zones, trailing Chengshan Jiao Promontory by 12,018 trajectories and 8,445,629 trajectory points. The SR of the accelerated ADPSC algorithm, when processing data in the Caofeidian Port region, stands at 51.4988. This is 11.4 and 2.2 points higher than the rates for Chengshan Jiao Promontory and Tianjin Port, respectively. Conclusively, the ADPSC algorithm's acceleration under the GPU parallel framework becomes more evident as the scale of the vessel trajectory dataset increases.

#### **6. Conclusion and future perspectives**

This paper enhances the conventional DP algorithm by developing an adaptive ADPSC method. This innovative compression approach autonomously determines compression thresholds for individual trajectories, relying on the distribution characteristics of vessel trajectories and navigational data, encompassing time stamps, SOG, and COG. Unlike applying a universal compression threshold to all trajectories, this method ensures optimal compression outcomes. Additionally, a GPU parallel computing framework is <span id="page-36-0"></span>proposed to expedite the ADPSC algorithm's data compression processes. This optimization framework divides the algorithm into two segments: parallel threshold calculation and parallel compression. This customized methodology is better suited for managing largescale datasets than conventional CPU-based systems.

Furthermore, this paper introduces the TP-DTW method, which calculates the similarity between trajectory point distributions on the baseline before and after compression to evaluate the compression effectiveness. Compared with the traditional DTW method, the novel TP-DTW method is based on the different distributions of the trajectory points, enhancing the compression analysis capability of the algorithm. Experimental datasets from three research areas, Tianjin Port, Chengshan Jiao Promontory, and Caofeidian Port, demonstrate that the ADPSC algorithm outperforms the DP counterpart by delivering better compression rates while minimizing data distortion. Simultaneously, leveraging the GPU parallel computing framework accelerates the ADPSC algorithm, especially when handling massive trajectory data. In essence, the proposed framework offers efficient and rapid trajectory compression that eliminates redundant data, making it crucial for intelligent maritime transportation systems. The GPU-accelerated adaptive compression methodology serves a multifaceted function by reducing data size while preserving accuracy and essential information. This enables efficient data storage, faster transmission over networks, cost savings in storage and bandwidth, improved data analysis, and enhanced safety and reliability in applications in maritime operations. Overall, it optimizes data management, enhancing cost-effectiveness and efficiency without compromising critical data integrity.

Nevertheless, future research directions could use multiple GPUs to construct new parallel computing frameworks, enhancing accelerated ADPSC compression algorithms. While the single GPU parallel compression algorithm highlighted in this paper already showcases promising acceleration outcomes, subsequent investigations could leverage multiple GPUs to either shorten execution times even further or process even larger data volumes more efficiently. It remains crucial to determine different thresholds for each vessel trajectory during compression tasks. Yet, trajectories exhibiting significant similarities could potentially be compressed using a shared threshold without compromising compression quality. Hence, an intriguing future research direction lies in accurately clustering vessel trajectories, followed by tailored threshold calculations for different trajectory clusters, which holds the promise of substantial computational savings.

# **CRediT authorship contribution statement**

**Yan Li:** Conceptualization, Data curation, Formal analysis, Investigation, Methodology, Resources, Software, Validation, Visualization, Writing – original draft, Writing – review & editing. **Huanhuan Li:** Conceptualization, Data curation, Formal analysis, Investigation, Methodology, Project administration, Resources, Software, Supervision, Visualization, Writing – original draft, Writing – review & editing. **Chao Zhang:** Formal analysis, Investigation, Validation, Visualization, Writing – original draft. **Yunfeng Zhao:**  Formal analysis, Investigation, Validation, Visualization, Writing – original draft. **Zaili Yang:** Funding acquisition, Methodology, Project administration, Resources, Validation, Writing – original draft, Writing – review  $\&$  editing.

#### **Declaration of competing interest**

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

#### **Data availability**

Data will be made available on request.

# **Acknowledgements**

This work is supported by the European Research Council (ERC) under the European Union's Horizon 2020 research and innovation program (Grant Agreement No. 864724) and Royal Society International Exchanges 2021 Cost Share (NSFC) (IEC\NSFC \211211).

#### **Appendix A**

Clustering evaluation metrics, including the Silhouette Coefficient (SC), Calinski-Harabasz Score (CHS), and Davies-Bouldin Index (DBI), assess clustering effectiveness by examining point density within clusters and separation between clusters ([Li et al., 2022; Li and](#page-38-0) [Yang, 2023\)](#page-38-0). After performing the DBSCAN clustering task, each trajectory obtains SC, CHS, and DBI values. Additionally, the average values of these metrics across all trajectories in the three study areas are calculated. The experimental results showing the average SC, CHS, and DBI values for varying *Eps* and *Minpt* parameters are displayed in [Fig. 24](#page-37-0). According to the experimental findings, when *Eps* is 0.01 and *Minpt* is 3, the three study areas achieve the maximum SC, CHS, and minimum DBI. Therefore, these values are optimal for the DBSCAN algorithm to identify outliers in all vessel trajectories within these study areas.

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**Fig. 24.** Visual illustration of the average matrices for SC, CHS, and DBI for different parameter values (i.e., *Eps* and *Minpt*) in three water areas. Specifically, the vertical axis of each matrix represents the value of the parameter *Minpt*, while the horizontal axis represents the value of the parameter *Eps*.

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