



Vertical Fine-Scale Distribution of *Calanus sinicus* in the Yellow Sea Cold Water Mass During the Over-Summering Process

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Pan J, Cheng F, Yu F, Shi Y, Sun F, Si G, Wei C, Diao X and Zhao Y (2021) Vertical Fine-Scale Distribution of Calanus sinicus in the Yellow Sea Cold Water Mass During the Over-Summering Process. Front. Mar. Sci. 8:644043. doi: 10.3389/fmars.2021.644043 Calanus sinicus, a temperate copepod with a lethal temperature >27°C, is one of the key species in Chinese coastal marine ecosystems. The C. sinicus population increases in spring and declines in early summer annually due to increasing water temperature. Numerous C. sinicus individuals then congregate in the Yellow Sea Cold Water Mass (YSCWM) and remain under the thermocline from early summer to early autumn. Development and reproduction is halted in this cold and foodless bottom water and they avoid ascending to the hot surface water, which is regarded as an oversummering strategy. Based on discrete water sampling approaches, previous studies demonstrated that higher chlorophyll a (Chl a) levels appeared in the mixed hot surface water layer; however, the subsurface chlorophyll a maximum layer (SCML) has seldom been described. In the present study, various probes and a visual plankton recorder (VPR) were used to determine the fine vertical distributions of environmental factors and C. sinicus. VPR observations showed the ecological responses in fine scale and indicated that few C. sinicus individuals ascend at night, the main population preferred to remain below the SCML all day long. The results demonstrated that a constant thin SCML existed in the YSCWM area, and that the SCML location coincided with or was beneath the thermocline and halocline layers, where the temperature was suitable for C. sinicus. The relationship between abundance and Chl a, showed the diel vertical migration trend of C. sinicus to feed at night in the YSCWM area. In addition to temperature as a main influencing factor, dissolved oxygen concentrations and column depth were also influencing factors. Therefore, in addition to avoiding high surface temperature, energy supplement may be an important driving force confining the diel vertical migration of *C. sinicus* in the Yellow Sea in summer.

Keywords: copepod, *Calanus sinicus*, Yellow Sea Cold Water Mass, diel vertical migration, subsurface chlorophyll a maximum layer

INTRODUCTION

Calanus sinicus is a warm-temperate copepod species widely distributed in the shelf ecosystems of the Northwest Pacific Ocean (Chen, 1964; Li et al., 2016), and plays an important role in the marine food web (Uye, 2000; Sun et al., 2010). It is a key species in China's marine pelagic ecosystems, contributing 73-99% of the biomass of the large copepod functional group all year round in the Yellow Sea (Sun et al., 2010; Kim and Kang, 2018). The C. sinicus population usually disappears during summer in most parts of southern China seas due to increased water temperature (Li et al., 2016). However, the population remains in the Yellow Sea all year round (Chen, 1964; Li et al., 2016). Numerous C. sinicus individuals reside in the Yellow Sea Cold Water Mass (YSCWM) to survive through the hot summer, which is regarded as an over-summering strategy (Zhang et al., 2005). The fifth copepodite stage (C5) and female individuals dominate the population during the over-summering process, exhibiting low ingestion and metabolic rates, suspended development, confined diel vertical migration (DVM), a well-developed oil sac, and a low RNA:DNA ratio (Li et al., 2004; Zhou and Sun, 2016; Wang et al., 2017). In autumn, C. sinicus in the YSCWM serve as the parental individuals for population development and can be transported southward by monsoon-driven currents (Hwang and Wong, 2005; Yin et al., 2011). These physiological and ecological phenomena are similar to the over-wintering strategy of the dormant copepod species Calanus finmarchicus in the North Atlantic Ocean (Jónasdóttir, 2010).

The *C. sinicus* population dynamics has mainly been examined using water samplers or nets in the Yellow Sea (Li et al., 2004; Zhang et al., 2005). However, these traditional approaches might obscure the behavioral details and ecological responses in fine scale, such as vertical migration, feeding rhythm, predator avoidance, and the vertical distribution pattern of *C. sinicus* (Pan et al., 2018). The refined vertical distribution of *C. sinicus* in the YSCWM with corresponding environmental factors needs to be studied during the over-summering period, to enhance our understanding of relationships between the ecological strategies of *C. sinicus* and environmental elements (Ross, 2014; Stanley et al., 2017), and to improve zooplankton models (Everett et al., 2017).

The Visual Plankton Recorder (VPR) has been used to examine real-time abundance and distribution of zooplankton individuals for over two decades (Davis et al., 1992; Ashjian et al., 2001). By imaging a given water volume and various fitting sensors, the VPR can continuously provide images of plankton species and simultaneous environmental elements, such as temperature, salinity, nitrate, and fluorescent chlorophyll a (Chl a) concentration (Benfield et al., 1996; Davis et al., 2005). Previous studies have shown that the VPR can provide comparable information to net samples on plankton vertical and horizontal distributions with accurate species components, and moreover fine scale distribution information (Broughton and Lough, 2006; Takahashi et al., 2015).

The vertical position of copepod individuals might be a tradeoff of many factors, such as temperature, light intensity, food concentration, and predator abundance, producing different

fitness that appears as mortality, growth rate and birth rate. The VPR can help to illustrate the copepod vertical distribution, improve comprehension of the profound consequences of vertical distribution and shed light on the behavioral feedback to environmental factors (Bo et al., 2006; Sainmont et al., 2014). In this study, we used the VPR to 1) describe the horizontal and vertical distributions of *C. sinicus*, and 2) represent the transformation of hydrographical conditions across the YSCWM, to determine the influences of the thermocline and fluorescence on the vertical and horizontal distributions of *C. sinicus*.

MATERIALS AND METHODS

Sampling Area and Time in the Yellow Sea

The Yellow Sea is a shallow and semi-closed marginal sea between the Chinese mainland and Korean Peninsula. The YSCWM is a seasonal water mass that is characterized by low temperature $(<9^{\circ}C)$ and high salinity (>32.5), occupying the third deepest part of the Yellow Sea in summer (Wang et al., 2003; Park et al., 2011). The formation of the YSCWM in summer has a significant impact on hydrographic features, biogeochemical cycles, marine planktonic biomass, and fishery resources in the Yellow Sea (Huo et al., 2012; Su et al., 2013).

The present investigation was conducted during a cruise in the Yellow Sea from 25 August to 7 September 2017 (**Figure 1**). A zooplankton net (mesh size: 505 mm) was hauled vertically from a depth of 2–4 m above the bottom to the surface at a speed of 1 m s⁻¹. Zooplankton samples were collected by the vertical net, which were then preserved in 5% formalin seawater solution after collection, identification and counting for morphological verification in the laboratory by ZooScan.

Temperature and salinity were recorded by a CTD instrument (Seabird 911) at all stations. The vertical profiles of turbidity, fluorescence, concentration of dissolved oxygen (DO), and nitrate were generated in each station simultaneously by equipment such as FLNTU (Wetlabs ECO Puck), DO (SBE-43), and Submersible Ultraviolet Nitrate Analyzer (SUNA) sensors. The CTD and additional sensors were calibrated before the cruise.

Survey of Visual Plankton Recorder and Environmental Elements Using Sensors

We employed the VPR for vertical tows from the bottom to the surface with the running speed at approximately 0.5–1 m s⁻¹ (Norrbin et al., 2009), and the sampling stations are shown in **Table 1**. The VPR used in the study consisted of two pressure cases, one which contained a Uniq UC-1830CL camera equipped with lenses that used four stepper motors (S0–S3) for accurate and repeatable positioning of the camera-lens extension, and the other contained a high-powered strobe in a ring configuration. The camera could generate compressed pictures with 1 megapixel resolution (1,024 × 1,024), and 10-bit color depth in the frame rate of around 15 s⁻¹ via an RS232 interface.



The imaged volume was calibrated before the survey using calibration files and calibration software (VPR_Cal) (Sainmont et al., 2014). Each VPR tow produced a file consisting of compressed images that were captured and recorded by the embedded computer stack. The pictures were obtained by the camera using the highest magnification with a field view of 42×42 mm (Ashjian et al., 2001; Sainmont et al., 2014). The ancillary CTD data were transmitted to a processing computer by two methods (USB adapter or Ethernet cable) after every VPR tow.

Data Analysis

The images were as determined for the gradient in grayscale and segmented at a user-selected brightness threshold (Davis et al., 1992; Ross, 2014). Regions of Interest (ROIs) were named by sampling time in milliseconds and stored in a special data directory structure. They were then time-stamped for subsequent spatial mapping (time in milliseconds within the day) and sampling depth via the time stamp using Mathworks (MATLAB software, 2015a), in order to allow them to be merged with the data from the CTD which were written in separate data files (Davis et al., 2005; Gislason et al., 2016).

Regions of interests were sorted automatically into different taxa using classifiers after training with a set of images of taxa assigned by morphological identifying specialists. All classified ROIs were subsequently checked manually to confirm classification and counted (Hu and Davis, 2006; Gislason et al., 2016). The hydrographic parameters could also be related to the ROIs by time stamping. Images of different taxa were listed by observation times and binned into 1s bins of sensor data, and then counted. The image volume and counts of manually sorted images were used to calculate abundance and then plotted with vertical profiles of physical parameters (Davis et al., 2005; Sainmont et al., 2014; Gislason et al., 2016).

The formula used to calculate abundance was as follows:

$$Abu = \frac{N_{ind}}{N_{frame} \times Volume} \times 10^3 \tag{1}$$

Where, Abu is the number of individuals per cubic meter, N_{ind} is the number of individuals observed in a second, N_{frame} is the

number of frames per second, and Volume is the image volume of a single frame (e.g., S3 setting, the volume is 300 ml).

For S3 stepper, the camera was focused such that the field of view of each image was 42 mm × 42 mm, the lens was adjusted to give a field of view of 1.7 cm and the calibrated image volume was 300 ml (Davis et al., 2005). The horizontal distribution of *C. sinicus* was related to water depth, number of detected individuals e.g. the cross section is w = 42 mm, l = 170 mm, h = depth (m), the volume is w × l × h (m³); and the horizontal abundance is the number of individuals detected/volume (ind/m³).

As C. sinicus is the dominant species in the Yellow Sea with a main body size of 2.4-3.5 mm (Chen, 1964), VPR S3 setting was chosen as the main stepper in this study. The range of zooplankton length that appeared in the obtained images was generally 1.8-16 mm. We did not discriminate the developmental stages of C. sinicus due to some ambiguous images. However, they did not hamper identification of C. sinicus, as it highly dominated at most stations according to the net samples and its obvious exterior helped identification. Based on the Savitzky-Golay method, the profiles of C. sinicus abundance are shown using MATLAB (R2015a). According to the DIVA gridding method, vertical profiles of temperature, salinity, fluorescence and nitrate were visualized using Ocean Data View software version (4.7.6) (Schlitzer, 2002). The depth of the subsurface Chl a maximum layer (SCML) was determined by the downward profile of fluorescence.

RESULTS

Environmental Conditions

All VPR stations were under highly stratified conditions, and temperature, salinity, and concentrations of dissolved oxygen (DO) generally exhibited steep changes between 10 and 40 m depth (Figure 2). The thickness of the thermocline (>0.5°C change in 1 m depth interval) ranged from 5 to 20 m (Figure 2). The boundary of the YSCWM was defined by the 10°C bottom isotherm (Fei et al., 2006; Figure 1). Most of the VPR stations in our survey (13 stations in all) were inside the YSCWM, and three stations (3400-06, 3825-05, and 3875-02) were outside and located at the edge of the YSCWM frontal area (bottom temperatures were <11.4°C). In addition, two stations (3500-05 and 3875-05) were located outside the YSCWM with bottom temperatures > 14.1°C (**Figures 1, 2**). The spatial coverage of the YSCWM was observed from 122°E to 124°E and from 34°N to 38°N in this study.

The range of salinity was 28.5–32.9, the salinity coverage in the YSCWM was more than 32. The salinity ranged from 31.16 to 32.93 among the 18 VPR stations, generally showing inverse changes compared with the temperature, and the haloclines mirrored the thermoclines at most stations (**Figure 2**). Above the thermocline, there was a mixed layer of warm water (>23°C) with salinity < 31.9. The bottom water showed relatively high salinity (>32) and low

TABLE 1 Time and depths of VPR stations used during August	to
September, 2017.	

Cast	Location	Data (Day/Month/Year)	Local time	Sampling Depth (m)	Bottom Depth (m)
		,		,	• • • •
3400-06	34°N, 123°E	07/September/2017	15:00	0–65	67
3400-07	34°N, 123.5°E	07/September/2017	19:00	0–67	69
3500-05	35°N, 121.5°E	02/September/2017	10:00	0–35	39
3500-08	35°N, 123°E	02/September/2017	1:00	0–68	72
3500-09	35°N, 123.5°E	01/September/2017	23:00	0–72	77
3600-05	36°N, 122.5°E	25/August/2017	7:00	0–55	59
3600-06	36°N, 123°E	25/August/2017	11:00	0–67	70
3600-08	36°N, 124°E	25/August/2017	17:00	0–75	77
3650-05	36.5°N, 123°E	26/August/2017	2:00	0–59	61
3650-07	36.5°N, 124°E	25/August/2017	21:00	0–74	76
3700-02	37°N, 123.5°E	27/August/2017	10:00	0–66	72
3700-03	37°N, 124°E	27/August/2017	13:00	0–74	75
3775-04	37.75°N, 123°E	31/August/2017	17:00	0–57	63
3775-05	37.75°N, 123.5°E	01/September/2017	5:00	0–69	70
3825-03	38.25°N, 122°E	31/August/2017	7:00	0–50	54
3825-05	38.25°N, 123°E	31/August/2017	18:00	0–54	55
3875-02	38.75°N, 122.5°E	28/August/2017	6:00	0–48	52
3875-05	38.75°N, 124°E	27/August/2017	22:00	0–55	60

temperature, ranging from 6.6°C (38.25°N, 122°E) to 14.2°C (35°N, 121.5°E) with the exception of station 3875-05 (bottom temperature 19.2°C, bottom salinity 31.94). The maximum temperature difference between the surface and bottom layers was up to 20.0°C in the center of the YSCWM (**Figure 2**). The concentration of DO in the water column of the stations assessed using a calibrated sensor varied from 2.60 to 7.66 mg. L⁻¹. In addition, the concentration of DO increased from the surface layer to the thermocline and was always higher near the SCML, then decreased with increased depth (**Figures 2, 3**).

A pronounced SCML formed around 15–30 m inside or below the thermocline at most stations except station 3875-05 (**Figure 3**). The highest fluorescence values of the SCML at all stations were from 1.3 to 3.65 mg. m^{-3} according to the fluorescence probe. The SCML may contribute substantial phytoplankton biomass in the water column. Our results also showed that the depth of the SCML was shallower (about 20 m) in the middle of the YSCWM region and the northern Yellow Sea, and was deeper (30 to 40 m) in the eastern part of the YSCWM region (**Figure 3**).We statistically analyzed the mean values and standard deviation of depth, Chl a, temperature, salinity and oxygen which are shown in **Table 2**. Pearson correlation analysis was performed to examine the relationships between *C. sinicus* abundance and environmental factors (**Table 3**).



Distribution of *Calanus sinicus* in the Yellow Sea

Calanus sinicus was detected by the VPR at all 18 stations and verified using the monograph 'Atlas of Common Zooplankton of Chinese Coastal Seas' (Sun et al., 2016). C. sinicus abundance was high at temperatures ranging from 6 to 10° C in summer. The abundance of C. sinicus was high at the optimal salinity from 32 to 32.9 during summer. In general, the results from most

stations indicated that the majority of the *C. sinicus* population remained below the thermocline and the SCML all day long and rarely appeared above the thermocline (**Figures 2**, **3**). The only exception was found at station 3825-05, located between Yantai and Dalian. Only a few *C. sinicus* individuals were detected in the SCML by the VPR at stations 3600-05, 3650-05, 3775-04, 3775-05, 3825-03, 3825-05, and 3875-02 (**Figure 3**). However, at some night stations, such as 3400-07, 3500-08, 3500-09, 3600-08, and



individuals captured by the VPR; h: survey time (24 h).

3650-07, they were not detected in or above the SCML and the thermocline (**Figure 3**). The peak of *C. sinicus* abundance at each station ranged from 222.2 to 888.9 ind/m³ (**Figure 3**). The highest

abundance of *C. sinicus* appeared at station 3775-04 (in the SCML) and station 3775-05 (in the layer below the thermocline). Both stations are located in the northern Yellow Sea.

We also used a plankton net (505 μ m) to collect samples at 17 stations, but not at station 3400-07, as the net sample was unavailable. We calculated the abundance of C. sinicus at each station to show the horizontal distribution. The VPR range of abundance was 15.3-668.4 ind/m³. To validate the copepod abundance results obtained by ZooScan, the net sample results are also shown. A similar pattern was revealed by both the VPR and the plankton net (Figure 4). The plankton net results showed that the area with high abundance was similar to the area calculated by the VPR, and the order of magnitude was the same, which proved that the VPR had high sampling accuracy. Net sample results usually reflect the overall distribution of C. sinicus at each station. Shi et al. (2019) showed the distribution results using a steel sampler at different depths (0, 5, 10, 20, 35, 50, 75, and 2-5 m from the bottom) in order to improve the vertical distribution of C. sinicus abundance.

DISCUSSION

Potential Roles of Chlorophyll A, Subsurface Chlorophyll A Maximum Layer, and Dissolved Oxygen

The survey stations were located in the YSCWM-influenced area, where the hydrodynamic environment is relatively stable in summer (Fei et al., 2006; Xia et al., 2006). Strong stratification, which suspended physical mixing of the upper seawater layer and the layer below the thermocline, led to limited nutrients for phytoplankton growth and generated low fluorescence in the surface waters of the YSCWM area (Wei et al., 2016). The rhythm of both the nutricline and SCML depths were coincident with the pycnocline depth due to stratification in summer, which has been reported by in situ studies in the Yellow Sea (Zhou and Sun, 2016). The SCML was found to be only several meters thick in the YSCWM area in summer using a submersible fluorometer probe in situ (Li et al., 2011). Consequently, few studies have described the detailed vertical distribution pattern of chlorophyll fluorescence values and the thin phytoplankton layer. Photosynthetic pigment analysis indicated a steep peak of fluorescence which coincided with or beneath the depth of the SCML (Fu et al., 2009). The SCML has been attributed to the contribution of brown algae and green algae, and is controlled by the mechanism of vertical nutrients transport and the depth of the pycnocline. The present study showed that the SCML appeared at 17 of 18 stations (Figure 3). The profiles of Chl a were closely related to those of nitrate, which was in accordance with the results of previous studies (Gong et al., 2017). In the present study, the fluorescence probe provided the fine scale vertical profiles of primary production and reliable food concentration data in the Yellow Sea.

Sun et al. (2020) showed that decreasing DO had potential negative effects on early recruitment of *C. sinicus* in the northern Yellow Sea, China. However, low DO (4-6 mg/L) may not affect the survival rate of *C. sinicus* (Wang et al., 2013). In the present study, the abundance of detected *C. sinicus* and the environmental factors at a typical YSCWM station (3500-09) were used in the Pearson correlation analysis (**Table 3**).

TABLE 2 Average values of environmental factors	s at	18
stations by VPR and CTD.		

Factor	Mean values	Standard deviation	Median	
	Weall values		weatan	
Temperature (°C)	9.162	2.347	8.526	
Salinity	32.356	0.229	32.264	
Depth (m)	53.002	12.619	55.386	
Oxygen (mg/L)	5.31	0.402	5.303	
Chl a (mg/m ³)	0.279	0.482	0.207	

TABLE 3 Results of correlation analysis between environmental variables and abundance of *Calanus sinicus* at YSCWM station (3500-09) at night.

Factor	Values of correlation analysis	P-value	
Salinity	-0.283	>0.05	
Depth	-0.317*	< 0.05	
DO	0.433*	< 0.05	
Chla	0.312*	< 0.05	

"*" indicate significantly higher with P < 0.05, respectively.

Our results showed that all of the individuals detected by the VPR remained in the 9°C layers with DO (4-7 mg/L) at 40-70 m. Chl a and DO had a positive relationship, and both were positively correlated with *C. sinicus* abundance. These results suggested that *C. sinicus* may use diel vertical migration to feed at station 3500-09 at night, in addition, avoiding hypoxia may be another purpose but not the main driving factor. Furthermore, field surveys with more stations in different seasons should be conducted in the future to verify the purpose of this strategy.

Over-Summering Hypothesis of Calanus sinicus

Wang et al. (2003) focused on the life history strategies of *C. sinicus* in summer, and the majority of *C. sinicus* (C5) remained in the YSCWM. Based on samples collected from different depths, high *C. sinicus* abundance, mainly composed of C5 and female adults, were resident in the bottom layer ($<12^{\circ}$ C) of the YSCWM in summer. *C. sinicus* was found to have a normal DVM in the Yellow Sea but was confined to a certain extent, as the majority of the population ascended to the thermocline but not to the mixed layer at night, and then descended after several hours. Chl a concentration was regarded as the main proxy for potential phytoplankton food of *C. sinicus* in the YSCWM area (Wang et al., 2009). However, simultaneous Chl a produced by the discrete water sampling approach always showed the highest values in mixed surface water above the thermocline with no description of the SCML (Li et al., 2004).

Previous studies proposed that the upper thermal limit of *C. sinicus* might be about 23° C in the field (Uye, 1988) or 27° C after the individuals have acclimated (Zhang et al., 2007). High surface temperature was harmful for breeding, egg hatching and developmental processes. As the *C. sinicus* population could not enter the food-rich hot surface layer (higher than 25° C), researchers presumed that *C. sinicus* had to adopt a confined DVM and use the bottom layer of the YSCWM as a refuge (Zhang et al., 2007). As a result, low temperature and scarce food in the



YSCWM restrained physiological processes such as reproduction and development of *C. sinicus* (Zhou et al., 2016). Previous studies suggested that clearance of *C. sinicus* in the YSCWM was high on board, but gut pigment contents were relatively low in situ (Li et al., 2004). This situation of low ingestion rates and relatively high oxygen consumption rates induced a net loss of body carbon in *C. sinicus*. Consequently, female individuals could not spawn due to lack of energy, and their oil sac decreased gradually from late spring to early autumn (Sun et al., 2011). Both respiration rate and RNA expression suggested that C5 individuals stopped developing, but were not in diapause (Li et al., 2004; Zhou et al., 2016).

The main observation data in this study were on vertical distribution and analysis. The data collected using the plankton net showed that the range of abundance in the surveyed area was 0-700 ind/m³. Shi et al. (2019) obtained similar results with a range of abundance of 16.9-423.7 ind/m³, and the data on the layered network of the YSCWM has a certain reference significance for analysis. Liu et al. (2003) focused on the frontal area in the Yellow Sea, and the abundance was less than 200 ind/m³ in the frontal zone. The results of Yin et al. (2011) showed the seasonal change, where the spring abundance was $23 \pm (77.78)$ ind/m³ and the summer abundance was 13.74 \pm (45.1) ind/m³ in the South China Sea. Previous research showed the results of using a traditional net to collect C. sinicus in the whole water layer and average them. While the local high value was not truly reflected, the results of this study better describes the local high value in order to reflect the actual C. sinicus vertical distribution using the VPR.

The Yellow Sea Warm Current, a branch of the Tsushima Warm Current originating from the Kuroshio region in the East China Sea, carries warmer and saltier water northward into the Yellow Sea. This current, together with the southward flowing coastal currents, plays an important role in the water exchange within the Yellow Sea (Shi et al., 2019). In winter, zooplankton communities in the Yellow Sea were dominated by temperate and warm-temperate species, such as *C. sinicus* (Chen et al., 2020). However, we have no relevant data at present on advection and

require further data in future. *C. sinicus* inside the YSCWM is probably quiescent in summer (Zhou et al., 2016), their activity and energy supply are insufficient to achieve migration, and we believe that most of the species observed using the VPR are local species.

Vertical Migration and Its Driving Force in Calanus sinicus

Investigations into the relationships between vertical distributions of zooplankton and the SCML require finescale sampling approaches and datasets on contemporary circumstances. The VPR provided more details on both vertical distribution profiles of C. sinicus and simultaneous Chl a levels in the YSCWM area in summer (Figure 3). The SCML was only several meters higher than the location of the C. sinicus population, which potentially induced DVM. However, only a small portion of C. sinicus ascended to the SCML at seven of 18 stations (3600-05, 3650-05, 3775-04, 3775-05, 3825-02, 3825-05, and 3875-02), while the main population of C. sinicus remained in the foodless YSCWM regardless of the survey time (Figure 3). Pu et al. (2004) showed that the vertical migration ability of C. sinicus was weak and only a few migrated to the thermocline; in general, the vertical distribution of C. sinicus revealed the confined range of classical DVM, which was similar to the results of previous studies (Zhang et al., 2007). Kang et al. (2013) observed that the major C. sinicus population did not migrate into the surface water even at stations where the surface temperature was comfortable (less than 20°C), indicating that there were other driving forces in addition to surface temperature. Thus, predation was speculated to be the main reason for the confined DVM.

Irrespective of simulated or in situ studies in either fresh water or marine ecosystems, numerous reports proposed that the DVM is a zooplankton behavioral feedback determined by trade-off between growth and evasion of predation risk (Pinti and Visser, 2019). Prey often use cold and dark deep water as a refuge to avoid predation from visual predators, and ascend to feed in the surface water at night (Mehner, 2012; Thygesen and Patterson, 2019; Moller et al., 2020). Due to the disadvantages of a cold and foodless environment in the deep layer, migrants often reduce their rate of development and reproductive fitness. Due to their energy intake requirements, herbivorous copepods may ascend to ingest food when circumstances are appropriate. Consequently, remaining in the SCML might be an optimal choice for them at night. In the English Channel, the majority of the C. helgolandicus population was found to remain in the SCML during the night in summer, where the seawater column was also highly stratified (Harris, 1988). Xu et al. (2016) reported that the sound scattering layer, which was attributed to copepods (mainly C. sinicus and Euchaeta spp.), migrated upward and remained in the SCML beneath the thermocline at night in the East China Sea. Particles 100-150 µm in size decreased simultaneously in the layer, and were considered to be ingested by copepods (Xu et al., 2016).

Previous studies suggested that predation could induce the descent of marine copepods such as Acartia hudsonica and Pseudodiaptomus spp. (Frost and Bollens, 2011), and reduce their feeding activation regardless of illumination (Bollens, 1996). The over-wintering strategy of Calanus spp., which descend to the deep, cold, and dark domain of the Atlantic Ocean seasonally in a dormant state, was recognized to be partially driven by predation from planktivorous fish (Bagoien et al., 2001). In the Yellow Sea, the C. sinicus population suffered significant predation pressure from the anchovy Engraulis japonicus (Kang et al., 2013). Predation by sardines and anchovies was thought to be the primary reason affecting the DVM of C. sinicus in the Japan inland sea (Uye, 1988). The hot areas of anchovy distribution were mainly located in and at the edge of the YSCWM area in summer (Niu et al., 2013) and E. japonicus often aggregated in and above the thermocline layer (Wan et al., 2008). The VPR and probes demonstrated confined vertical migration of C. sinicus in the Yellow Sea in the present study, although the distribution of the SCML was suitable. Therefore, avoidance of predators was presumed to be the most probable driving force regulating the DVM of C. sinicus in the Yellow Sea in summer. The simultaneous diel vertical distributions of Chl a, copepod C. sinicus and anchovy E. japonicus should be further studied.

Lipid accumulation is deemed to be an important factor affecting the DVM of copepods. In the Arctic Ocean, Calanus spp. with a high lipid load were found to remain in the deep water, while individuals with a lower lipid load and all Metridia longa developmental stages continued normal DVM. Substantial lipid reserves of *Calanus* spp. are considered to be the main energy resource after diapause, while M. longa adopted a combined strategy of active feeding and lipid accumulation to overcome the Arctic conditions in winter (Schmid et al., 2018). The average value of the relative oil sac volume (expressed as % of body volume) of C. sinicus (about 30-35%) in the YSCWM (Sun et al., 2011) was lower than that of Calanus spp. (about 50-80%) in the Arctic Ocean (Schmid et al., 2018). In addition, part of the C. sinicus population developed and the oil sac shrank gradually during summer (Wang et al., 2009) and evidence from the grazing impact of C. sinicus (Li et al., 2004), oxygen consumption rate and gene expression (Zhou et al., 2016), all support that the oversummering strategy of C. sinicus in the YSCWM is quiescent and not diapause, which is similar to the combined strategy of *M. longa.* Therefore, lipid accumulation may not be the factor that ceases DVM of *C. sinicus.* The detailed relationships between lipid fullness and DVM of *C. sinicus* remain to be explored in future studies.

CONCLUSION

In this study, the VPR was used to determine the refined vertical distribution characteristics of *Calanus sinicus* in the Yellow Sea. Environmental factors affecting the vertical distribution of *C. sinicus* are discussed and statistically analyzed in this study. The influence of the SCML on *C. sinicus* vertical distribution has seldom been described. The relationship between abundance and Chl a, showed that the diel vertical migration trend in *C. sinicus* is to feed at night in the YSCWM. In addition to temperature, DO concentrations, and the column depth are also influencing factors.

DATA AVAILABILITY STATEMENT

The datasets generated for this study are available on request to the corresponding author.

AUTHOR CONTRIBUTIONS

JP, FC, FY, and YS contributed to the conception of the study. JP, XD, CW, and FS contributed to sample collection and data collation. JP, GS, FS, and YZ analyzed the samples and conducted the data analyses. JP, FC, and YS wrote the manuscript and contributed to the definition of aims. All authors contributed to improvement of the manuscript before approving it for submission.

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