

Copepod communities related to water masses in the southwest East China Sea

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Abstract The East China Sea is characterized by a complex hydrographic regime and high biological productivity and diversity. This environmental setting in particular challenged a case study on the use of mesozooplankton community parameters as indicators of water masses. In order to reveal spatial patterns of zooplankton communities during summer, a large scale oceanic transect study was conducted. Two transects were taken in the southwest East China Sea region, covering for the first time the China shelf, slope, and the estuaries of the Yangtze river and of the Minjiang river, the northern Taiwan Strait, and the Kuroshio Current region. A total of 77 copepod species were quantified. Copepod abundance was significantly higher in the estuary of the Yangtze River runoff mixture waters and lowest at the Kuroshio Current Region. The calanoid *Parvocalanus crassirostris* was the most frequently occurring and abundant species retrieved from 27 samples of a total of 39 samples. The use of multivariate cluster analysis separated the Mainland China Shelf from the northern Taiwan Strait and the Kuroshio Current Region at

the first hierarchical level. The use of an indicator value method (IndVal) associated with each cluster of stations revealed characteristic species assemblages. Two hierarchical levels defined 4 assemblages within geographical sectors representing copepod assemblages of the Kuroshio Current Region, of the northern Taiwan Strait and the southern China Shelf near the estuary of the Minjiang River and northern stations near the estuary of the Yangtze River. Overall, there was a strong correspondence between the distribution of certain copepod species and water masses. Differences between the Mainland China shelf, the northern Taiwan Strait and the Kuroshio Current Region were characterized by differences in species composition and abundance. Water mass boundaries in the study area were exclusively indicated by distinct differences in species composition, emphasizing a correlation between copepod communities and water masses of the southwest East China Sea in summer.

Keywords Copepod composition · Water mass · East China Sea · Taiwan Strait · Kuroshio Current

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Introduction

The present study was carried out within a long-term project “Long-term Observation and Research of the East China Sea (ECS) (I)—Factors Controlling Carbon Fluxes (LORECS)”. The west border of the East China Sea to the mainland of China links with the Yellow Sea in the north and with the Taiwan Strait (TS) in the south. Water masses passing the Kuroshio Current region (KCR) along the east coast of Taiwan flow northward (Ashjian et al. 2005; Shahidul Islam et al. 2006). There are many rivers carrying the largest runoff of low-saline water to the East China Sea

from the mainland of West China (Liu et al. 2000), e.g. the Yangtze River, Qiantang River, Minjiang River and Wujiang River. The annual seawater temperature is primarily influenced by the monsoons and mediated by ocean currents. In summer, the sea surface water temperature is nearly homogeneous at about 27–29°C and is influenced by the Yangtze River, where a low-saline high-thermal water tongue is sometimes entering the estuarine regions (Chen 1992).

Zooplankton communities support higher trophic levels and copepods are an important determinant as for their abundance and ecological roles (Turner 1998; Hwang et al. 1998; Wong et al. 1998). In the upwelling of the Kuroshio Current and in the East China Sea near to Taiwan copepods are the most abundant constituents of mesozooplankton communities (Chen and Chen 1992). Investigating copepod community structure and distribution patterns appears therefore as an essential step towards the understanding of the trophic ecology of pelagic systems (Chen 1992).

Compared to other oceanographic regions, there have been only a few studies on copepod ecology in the East China Sea (Chen and Zhang 1965; Tan 1967; Chen et al. 1974; Chen and Chen 1992; Shih and Young 1995; Shih and Chiu 1998; Lan et al. 2004; Hwang and Wong 2005; Hwang et al. 2006). Planktonic copepods in upwelling regions of the southern East China Sea have been studied previously (Shih et al. 2000; Lo et al. 2004). Several studies have been published on the biological oceanography of the East China Sea area, including zooplankton composition (Chen 1992), copepod transport by monsoon driven currents (Hwang and Wong 2005; Hwang et al. 2006), chlorophyll *a* (Chang et al. 2003a; Gong et al. 2000; Gong and Liu 2003), primary productivity in correlation with chemical hydrography (Gong et al. 1996, 2000), photosynthetic-irradiance models (Gong et al. 2001), primary production (Gong and Liu 2003), cyanobacteria (Chiang et al. 2002; Chang et al. 2000, 2003b), nutrient fluxes (Liu et al. 2000), carbon cycling (Chang et al. 2003a; Shiah et al. 2003), and potential anthropogenic impacts (Hwang et al. 2004). Previous publications suggest a correlation between temporal and spatial distribution patterns of zooplankton that affect the dynamics of biological and chemical interactions (Berasategui et al. 2006). In the present paper, the spatial distribution of copepods in the southwest ECS and in summer was examined for the first time. The present study provides additional and new information on the composition, structure and dynamics of copepod communities in relation to water masses of estuaries and rivers in general and those of the ECS, the shelf of mainland China, the northern Taiwan Strait (NTS) and the Kuroshio Current water masses in particular.

Materials and methods

Field sampling

The area of investigation in the southwest ECS (Fig. 1) is a border region of mainland China adjacent to the NTS and Taiwan. We set up 13 stations along two perpendicular transects and sampled mesozooplankton during cruise 618 of Ocean Research Vessel I of the National Science Council of Taiwan (Republic of China) from 15 to 29 July 2001. A transect line with 8 stations was chosen north of the estuary of Yangtze River (EYR) towards the southern estuary of the estuary of Minjiang River (EMR). A second transect line with five stations was located near the western EMR through the NTS east of KCR. The locations of stations are shown in Fig. 1 and details are shown in Table 1. Copepods were collected from the seabed to the surface at all stations in oblique hauls by 100 µm mesh nets. Tow speed was set at 33–50 cm s⁻¹. Seabed depths of stations 12 and 13 were 577 m and 1,760 m, respectively—but samples were taken only above 150 m water depth. Since the copepod composition is known to change diurnally, samples were taken every 6 h at four stations with the following specificities: EYR (six samples, 20 m vertical net tow), EMR (eight samples, 10 m vertical net tow), NTS (eight samples, 60 m vertical net tow), and KCR (eight samples, 150 m vertical net tow). The water volume filtered was estimated from flow

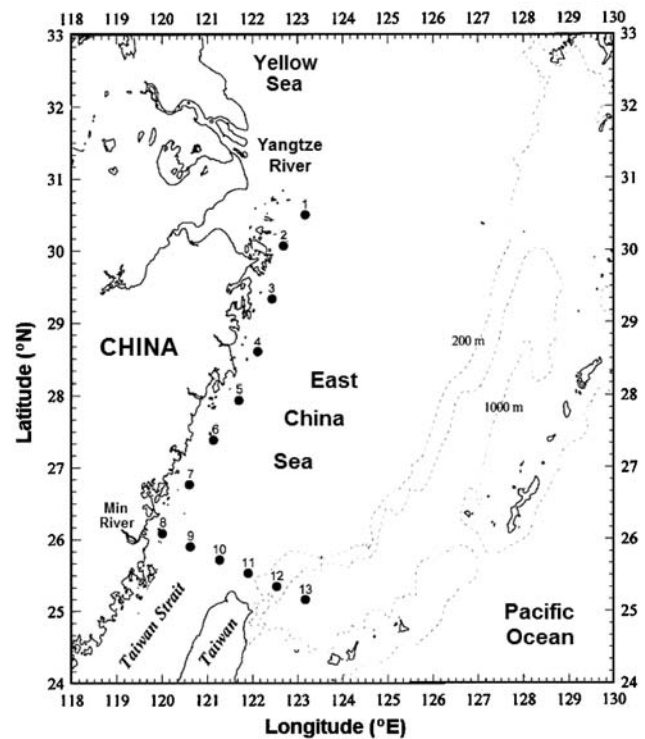


Fig. 1 Map of the sampling stations of LORECS OR1 618, 15–29 July 2001

Table 1 Locations, dates, times, sampling depths, and CTD data collected at LORECS OR1 CR-618 stations

Station	Latitude (N)	Longitude (E)	Date (-July 2001)/Time	Sampling depth (m)	CTD data depth (m)
1	30°30′	123°10′	16/22:00, 17/04:00, 17/10:00, 17/16:00, 17/22:00, 18/04:00	30	30
2	30°30′	122°52′	18/12:50	45	45
3	29°19′	122°25′	18/18:00	25	25
4	28°36′	122°07′	18/23:30	25	25
5	27°56′	121°42′	19/05:00	25	25
6	27°22′	121°08′	19/10:50	25	25
7	26°46′	120°36′	19/16:50	30	30
8	26°01′	119°54′	22/16:00, 22/22:00, 23/04:00, 23/10:00, 23/16:00, 23/22:00, 24/04:00, 24/10:00	10	20
9	25°54′	120°37′	24/20:45	60	60
10	25°43′	121°16′	20/11:00, 20/17:00, 20/23:00, 21/05:00, 21/11:00, 21/17:00, 21/23:00, 22/05:00	70	70
11	25°32′	121°54′	25/07:00	115	115
12	25°20′	122°32′	25/11:30	150	150
13	25°10′	123°10′	25/17:00, 25/23:00, 26/05:00, 26/11:00, 26/17:00, 26/23:00, 27/05:00, 27/11:00	150	150

meter readings mounted at the net openings. Samples were immediately preserved in buffered formalin after retrieval. We used seawater diluted formalin at final concentrations of 5%.

Identification and enumeration

In the laboratory, each sample was split with a Folsom splitter for taxonomic identification and enumeration. Species composition and abundances were determined by counting the adults. General references for identification were Chen and Zhang (1965), Chen et al. (1974), Zheng et al. (1965, 1982) and Huys and Boxshall (1992).

Statistical analyses

For statistical treatment a matrix was made of species as variables and samples or stations as objects. A double square root transformation of the original data of species densities was applied first, before computing a similarity coefficient between samples. Then, a hierarchical classification of samples was realized using the Bray-Curtis similarity coefficient and clustering approach of flexible links as has been demonstrated by Souissi et al. (2001) and Anneville et al. (2002). The advantage of using dendrograms is that different “resolutions” can be obtained from the dataset depending on the choice of cut-off levels, i.e. the first cut-off level produces two groups, the next level produces three groups etc. For the present analysis, the first three cut-off levels were retained and analysed. In order to characterize species assemblages for each individual group and across all cut-off levels, indicator species were calculated from an

indicator value index (IndVal) proposed by Dufrene and Legendre (1997) as in Souissi et al. (2001) and Anneville et al. (2002). This index is obtained by multiplying two-independent information: specificity measures (measure of affiliation) and fidelity (measure of occurrence) of a species for a group. The species characterizing a group best is indicated by a high IndVal. For the present study, only species having an indicator value greater than the arbitrary threshold level of 25% were retained in the assemblage. Furthermore, the characteristic species for each hierarchical level, showing the maximal value of their IndVal in that category are pointed out (Dufrene and Legendre 1997). Different steps of the cluster analysis and the graphical representation were programmed using Matlab Software (ver. 6.1).

Results

Faunistic peculiarities

From 39 samples we found 77 copepod species belonging to 43 genera and 25 families: 54 species of Calanoida, 6 species of Cyclopoida, 4 species of Harpacticoida, and 13 species of Poecilostomatoida (Table 2). There were 24 species present in ten or more samples but 13 species occurred only once. The ten most abundant and common species were *Parvocalanus crassirostris* (Dahl, 1893), *Oithona nana* Giesbrecht, 1892, *Paracalanus parvus* (Claus, 1863), *Euterpina acutifrons* (Dana, 1847), *Paracalanus nanus* Sars, 1907, *Corycaeus (Ditrichocorycaeus) affinis* McMurrich, 1916, *Paracalanus aculeatus* Giesbrecht, 1888, *Oithona rigida* Giesbrecht, 1896, *Acartia (Odontacartia)*

Table 2 Planktonic copepod species recorded during summer sampling in the East China Sea, 15–29 July 2001

Copepod taxa	Abbrev.	OR	RA	Mean	SD
CALANOIDA					
ACARTIIDAE					
<i>Acartia (Odontacartia) spinicauda</i> Giesbrecht, 1889	<i>Acart. spin.</i>	41.03	2.70	100.09	184.62
<i>Acartia negligens</i> Dana, 1849	<i>Acart. negl.</i>	35.90	0.55	20.35	38.21
AETIDEIDAE					
<i>Undeuchaeta plumosa</i> (Lubbock, 1856)	<i>Undeu. plum.</i>	2.56	0.02	0.69	4.29
AUGAPTILIDAE					
<i>Haloptilus longicornis</i> (Claus, 1865)	<i>Halop. long.</i>	2.56	0.01	0.29	1.81
CALANIDAE					
<i>Calanus sinicus</i> Brodsky, 1965	<i>Calan. sini.</i>	20.51	1.83	67.89	189.61
<i>Canthocalanus pauper</i> (Giesbrecht, 1888)	<i>Canth. paup.</i>	61.54	1.42	52.45	63.23
<i>Cosmocalanus darwini</i> (Lubbock, 1860)	<i>Cosmo. darw.</i>	20.51	0.13	4.92	11.07
<i>Mesocalanus tenuicornis</i> (Dana, 1863)	<i>Mesoc. tenu.</i>	2.56	0.05	2.04	12.71
<i>Nannocalanus minor</i> (Claus, 1863)	<i>Nanno. mino.</i>	12.82	0.07	2.62	8.18
<i>Neocalanus gracilis</i> (Dana, 1849)	<i>Neoca. grac.</i>	2.56	0.02	0.66	4.10
<i>Undinula vulgaris</i> (Dana, 1849)	<i>Undin. vulg.</i>	38.46	1.44	53.51	103.54
CANDACIIDAE					
<i>Candacia bradyi</i> A. Scott, 1902	<i>Canda. brad.</i>	17.95	0.19	7.21	20.55
<i>Candacia catula</i> (Giesbrecht, 1889)	<i>Canda. catu.</i>	15.38	0.11	4.02	12.08
<i>Paracandacia truncata</i> (Dana, 1849)	<i>Pcand. trun.</i>	2.56	0.04	1.58	9.85
CENTROPAGIDAE					
<i>Centropages calaninus</i> (Dana, 1849)	<i>Centr. cala.</i>	7.69	0.03	1.06	3.82
<i>Centropages dorsispinatus</i> Thompson & Scott, 1903	<i>Centr. dors.</i>	5.13	0.17	6.20	27.23
<i>Centropages furcatus</i> (Dana, 1849)	<i>Centr. furc.</i>	10.26	0.06	2.15	7.11
CLAUSOCALANIDAE					
<i>Clausocalanus arcuicornis</i> (Dana, 1849)	<i>Claus. arcu.</i>	7.69	0.03	0.95	3.68
<i>Clausocalanus furcatus</i> (Brady, 1883)	<i>Claus. furc.</i>	46.15	1.77	65.62	119.77
EUCALANIDAE					
<i>Mecynocera clausi</i> Thompson, 1888	<i>Mecyn. clau.</i>	5.13	0.03	1.01	5.60
<i>Pareucalanus attenuatus</i> (Dana, 1849)	<i>Peuca. atte.</i>	12.82	0.07	2.53	8.12
<i>Rhincalanus rostrifrons</i> Dana 1849	<i>Rhcal. rost.</i>	5.13	0.03	1.02	4.54
<i>Subeucalanus crassus</i> (Giesbrecht, 1888)	<i>Seuca. cras.</i>	7.69	0.06	2.21	8.82
<i>Subeucalanus subcrassus</i> (Giesbrecht, 1888)	<i>Seuca. subc.</i>	66.67	2.59	96.07	136.75
<i>Subeucalanus subtenis</i> (Giesbrecht, 1888)	<i>Seuca. subt.</i>	17.95	0.15	5.67	14.03
EUCHAETIDAE					
<i>Euchaeta concinna</i> (Dana, 1849)	<i>Eucha. conc.</i>	64.10	2.15	79.62	179.29
<i>Euchaeta plana</i> Mori, 1937	<i>Eucha. plan.</i>	10.26	0.20	7.49	28.94
<i>Euchaeta rimana</i> Bradford, 1973	<i>Eucha. rima.</i>	5.13	0.13	4.71	20.82
HETERORHABDIDAE					
<i>Heterorhabdus papilliger</i> (Claus, 1863)	<i>Heter. papi.</i>	2.56	0.02	0.62	3.86
LUCICUTIIDAE					
<i>Lucicutia flavicornis</i> (Claus, 1863)	<i>Lucic. flav.</i>	15.38	0.17	6.42	17.96
METRIDIIDAE					
<i>Pleuromamma abdominalis</i> (Lubbock, 1856)	<i>Plmam. abdo.</i>	5.13	0.03	1.09	4.91
<i>Pleuromamma gracilis</i> (Claus, 1863)	<i>Plmam. grac.</i>	10.26	0.08	2.86	9.88

Table 2 continued

Copepod taxa	Abbrev.	OR	RA	Mean	SD
PARACALANIDAE					
<i>Acrocalanus gibber</i> Giesbrecht, 1888	<i>Acala. gibb.</i>	12.82	0.23	8.48	25.50
<i>Acrocalanus gracilis</i> Giesbrecht, 1888	<i>Acala. grac.</i>	30.77	0.54	20.10	44.06
<i>Acrocalanus monachus</i> Giesbrecht, 1888	<i>Acala. mona.</i>	15.38	0.15	5.67	15.27
<i>Calocalanus pavo</i> (Dana, 1849)	<i>Caloc. pavo.</i>	20.51	0.30	11.06	33.68
<i>Calocalanus pavoninus</i> Farran, 1936	<i>Caloc. pvon.</i>	20.51	0.50	18.48	54.16
<i>Calocalauns plumulosus</i> (Claus, 1863)	<i>Caloc. plum.</i>	25.64	0.52	19.38	55.27
<i>Paracalanus aculeatus</i> Giesbrecht, 1888	<i>Pcala. acul.</i>	56.41	4.55	168.67	288.14
<i>Paracalanus nanus</i> Sars, 1907	<i>Pcala. nanu.</i>	64.10	6.07	224.87	365.03
<i>Paracalanus parvus</i> (Claus, 1863)	<i>Pcala. parv.</i>	46.15	10.69	395.93	863.85
<i>Parvocalanus crassirostris</i> (Dahl, 1893)	<i>Parvc. cras.</i>	69.23	16.92	626.88	1158.89
PONTELLIDAE					
<i>Calanopia elliptica</i> (Dana, 1849)	<i>Capia. elli.</i>	7.69	0.05	1.98	7.38
<i>Calanopia minor</i> A. Scott, 1902	<i>Capia. mino.</i>	10.26	0.08	2.96	11.05
<i>Labidocera acuta</i> (Dana, 1849)		5.13	0.03	1.18	5.56
<i>Labidocera bipinnata</i> Tanaka, 1936	<i>Labid. bipi.</i>	5.13	0.07	2.48	10.88
<i>Labidocera minuta</i> Giesbrecht, 1889	<i>Labid. minu.</i>	2.56	0.04	1.58	9.85
<i>Pontellina plumata</i> (Dana, 1849)	<i>Ponte. plum.</i>	5.13	0.07	2.48	11.18
SCOLECITHRICIDAE					
<i>Scaphocalanus echinatus</i> Farran, 1905	<i>Scaph. echi.</i>	2.56	0.03	1.09	6.84
<i>Scolecithricella longispinosa</i> Chen & Zhang, 1965	<i>Scoll. long.</i>	35.90	2.37	87.73	289.11
<i>Scolecithrix danae</i> (Lubbock, 1856)	<i>Scolx. dana.</i>	12.82	0.19	7.09	24.30
TEMORIDAE					
<i>Temora discaudata</i> (Giesbrecht, 1889)	<i>Temor. disc.</i>	43.59	0.75	27.80	49.16
<i>Temora turbinata</i> (Dana, 1849)	<i>Temor. turb.</i>	17.95	0.33	12.30	31.78
TORTANIDAE					
<i>Tortanus gracilis</i> (Brady, 1883)	<i>Tortan. grac.</i>	2.56	0.02	0.65	4.04
CYCLOPOIDA					
OITHONIDAE					
<i>Oithona attenuata</i> Farran, 1913	<i>Oitho. atte.</i>	10.26	0.14	5.22	21.08
<i>Oithona fallax</i> Farran, 1913	<i>Oitho. fall.</i>	7.69	0.10	3.71	14.19
<i>Oithona nana</i> Giesbrecht, 1892	<i>Oitho. nana</i>	48.72	14.32	530.75	1238.39
<i>Oithona rigida</i> Giesbrecht, 1896	<i>Oitho. rigid.</i>	56.41	4.49	166.23	277.06
<i>Oithona setigera</i> (Dana, 1849)	<i>Oitho. seti.</i>	23.08	0.35	13.08	32.16
<i>Oithona similis</i> Claus, 1866	<i>Oitho. simi.</i>	20.51	0.29	10.78	42.86
HARPACTICOIDA					
CLYTEMNESTRIDAE					
<i>Clytemnestra scutellata</i> Dana, 1847	<i>Clyte. scut.</i>	20.51	0.38	13.97	33.76
ECTINOSOMIDAE					
<i>Microsetella norvegica</i> (Boeck, 1846)	<i>Micro. norv.</i>	61.54	1.23	45.43	59.04
EUTERPINIDAE					
<i>Euterpina acutifrons</i> (Dana, 1847)	<i>Euter. acut.</i>	66.67	9.66	357.85	746.05
MACROSETELLIDAE					
<i>Macrosetella gracilis</i> (Dana, 1847)	<i>Macro. grac.</i>	30.77	0.60	22.08	64.79
POECILOSTOMATOIDA					
CORYCAEIDAE					
<i>Corycaeus (Corycaeus) crassiusculus</i> Dana, 1849	<i>Coryc. cras.</i>	7.69	0.08	3.08	13.37
<i>Corycaeus (Corycaeus) speciosus</i> Dana, 1849	<i>Coryc. spec.</i>	5.13	0.03	1.18	5.56

Table 2 continued

Copepod taxa	Abbrev.	OR	RA	Mean	SD
<i>Corycaeus (Ditrichocorycaeus) affinis</i> McMurrich, 1916	<i>Coryc. affi.</i>	64.10	5.33	197.39	368.02
<i>Corycaeus (Onychocorycaeus) pumilus</i> M. Dahl, 1912	<i>Coryc. pumi.</i>	7.69	0.05	1.77	6.91
<i>Corycaeus (Urocorycaeus) lautus</i> Dana, 1849	<i>Coryc. laut.</i>	20.51	0.28	10.45	33.12
<i>Farranula concinnus</i> (Dana, 1847)	<i>Farra. conc.</i>	2.56	0.04	1.65	10.33
<i>Farranula gibbula</i> (Giesbrecht, 1891)	<i>Farra. gibb.</i>	28.21	0.37	13.88	31.06
ONCAEIDAE					
<i>Oncaea conifera</i> Giesbrecht, 1891	<i>Oncae. conif.</i>	25.64	0.79	29.13	76.47
<i>Oncaea media</i> Giesbrecht, 1891	<i>Oncae. medi.</i>	17.95	0.20	7.50	22.35
<i>Oncaea venusta</i> Philippi, 1843	<i>Oncae. venu.</i>	30.77	0.33	12.41	32.57
SAPPHIRINIDAE					
<i>Copilia mirabilis</i> Dana, 1849	<i>Copil. mira.</i>	2.56	0.03	1.09	6.84
<i>Sapphirina intestinata</i> Giesbrecht, 1891	<i>Sapph. inte.</i>	2.56	0.03	1.13	7.03
<i>Sapphirina nigromaculata</i> Claus, 1863	<i>Sapph. nigr.</i>	2.56	0.03	0.98	6.10

OR occurrence rate (%); RA relative abundance (%), Mean average abundance (in d m^{-3}); SD standard deviation (in d m^{-3})
Taxa in bold-face represent the most abundant taxa retrieved during the investigation

spinicauda Giesbrecht, 1889 and *Subeucalanus subcrassus*, (Giesbrecht, 1888). *Parvocalanus crassirostris* did not only occur in 27 of 39 samples, but represented the most abundant species with an average density of 626.88 (individuals m^{-3}).

Hydrological structure

Monthly-averaged information derived from advanced very high resolution radiometer (AVHRR) recordings for sea surface temperature and seawater chlorophyll *a* values for July 2002 are shown in Fig. 2. The image for sea surface temperatures (Fig. 2a) shows the estuary of the Yangtze River with a high temperature fresh water output, providing water temperatures above 30°C. Surrounding waters of northern Taiwan and coastal mainland China were above

29°C. Sea surface concentration of chlorophyll *a* (Fig. 2b) shows the highest regional distribution along the coast of the mainland of China, especially in the EYR, reaching levels above 5.0 mg m^{-3} . A warmer fresh-water mass tongue (>30°C, <30 practical salinity unit—PSU) was found in the EYR, providing high concentrations of chlorophyll *a* in the Yellow Sea and the north-western ECS. The chlorophyll *a* concentration of areas from the EMR towards the NTS and the KCR were lower than along the mainland China coast, ranging between 0.5 and 1.0 mg m^{-3} .

CTD profiles

The vertical profiles of temperature and salinity at each sampling station are shown in Fig. 3. The hydrography of

Fig. 2 Monthly-averaged information derived from AVHRR for sea surface temperature (a) and SeaWiFS chlorophyll *a* (b) of July 2001

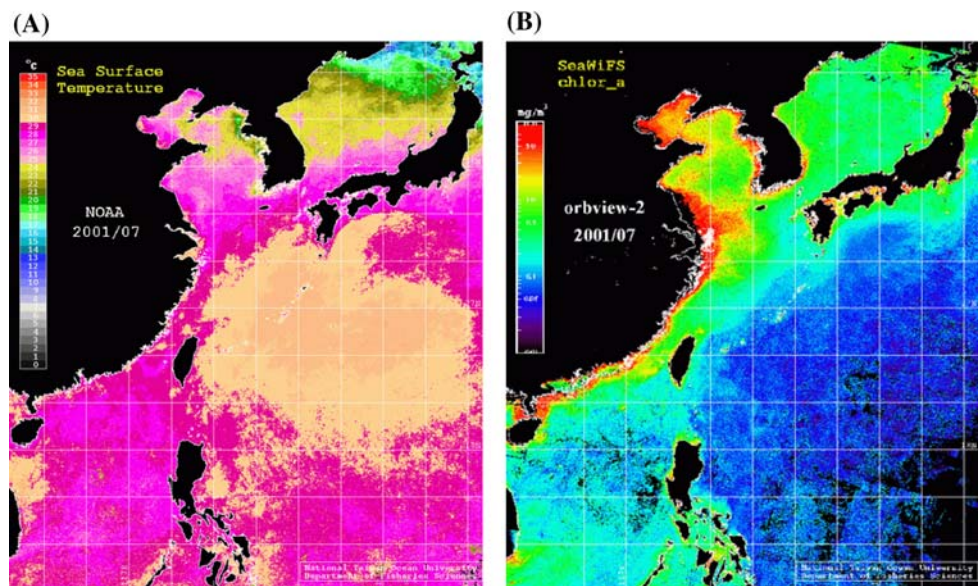
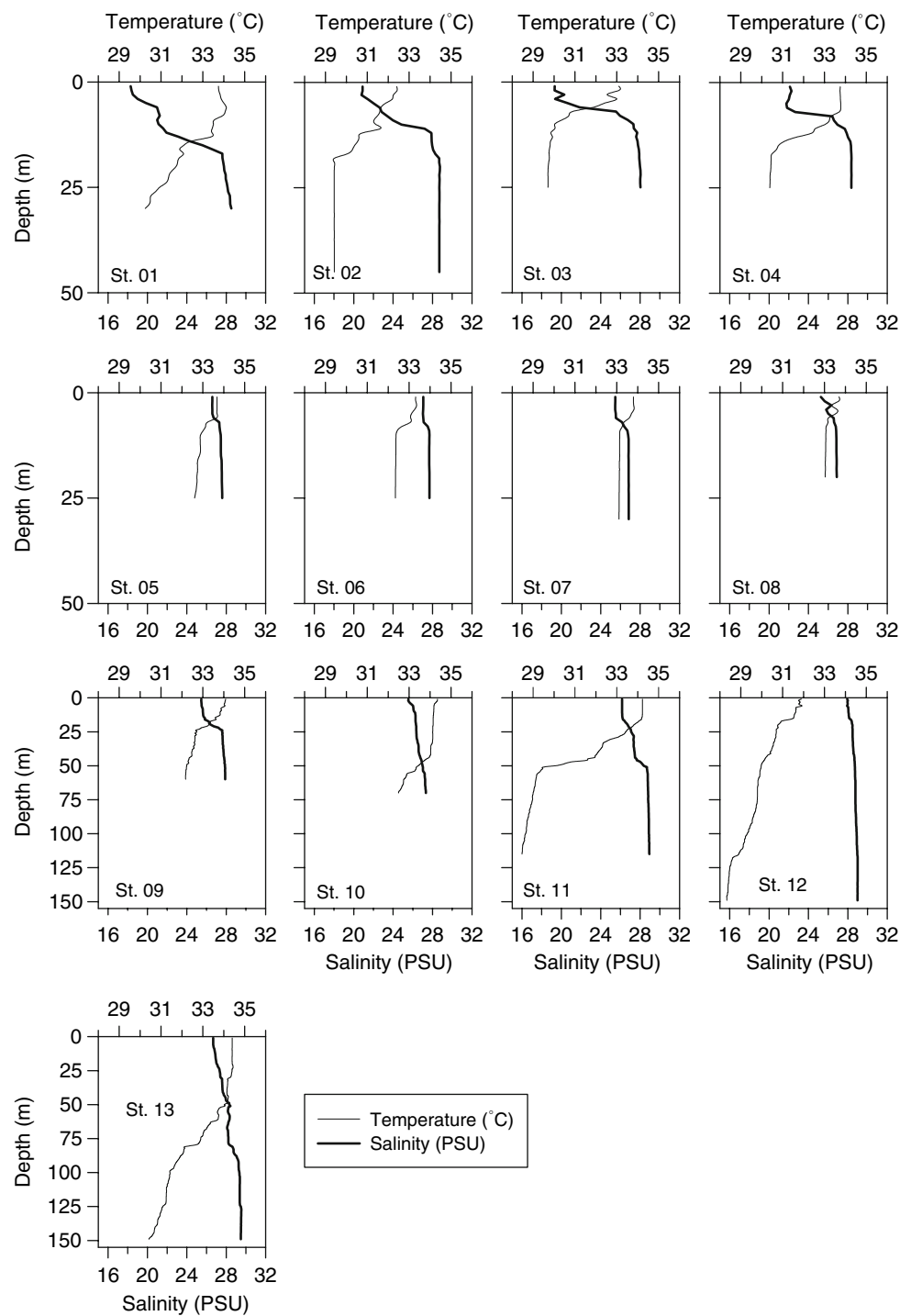


Fig. 3 Vertical variation of temperature and salinity at the sampling stations 1–13



sampling stations can be grouped within five categories. First, station 9 (EMR) belongs to estuarine water. The runoff from the Minjiang River shows low salinity and higher temperatures. Here, the salinity can be lower than 29.0. Second, station 1 (EYR) grouped as the continental coastal water, is influenced by the Yangtze River with water resulting from a mixing of river runoff into the sea with a low-saline and high-temperature water mass. A third

group, with stations 11 (NTS), 12 and 13 belongs to the ECS with mixed water from the KC and the Taiwan Strait Kuroshio branch water with the continental coastal water. Hence, salinity values range in the middle between these two. A fourth group with station 13 (KCR) being influenced by the Kuroshio surface layer water mass is characterized by high-temperature and high-salinity; the salinity can be higher than 34.2. A fifth group with low-saline and

low-temperature water belongs to stations 3 and 4. This water is formed by the Yangtze River mixed with continental coastal waters forming a long strip along the coast.

Hierarchical classification and identification of copepod assemblages

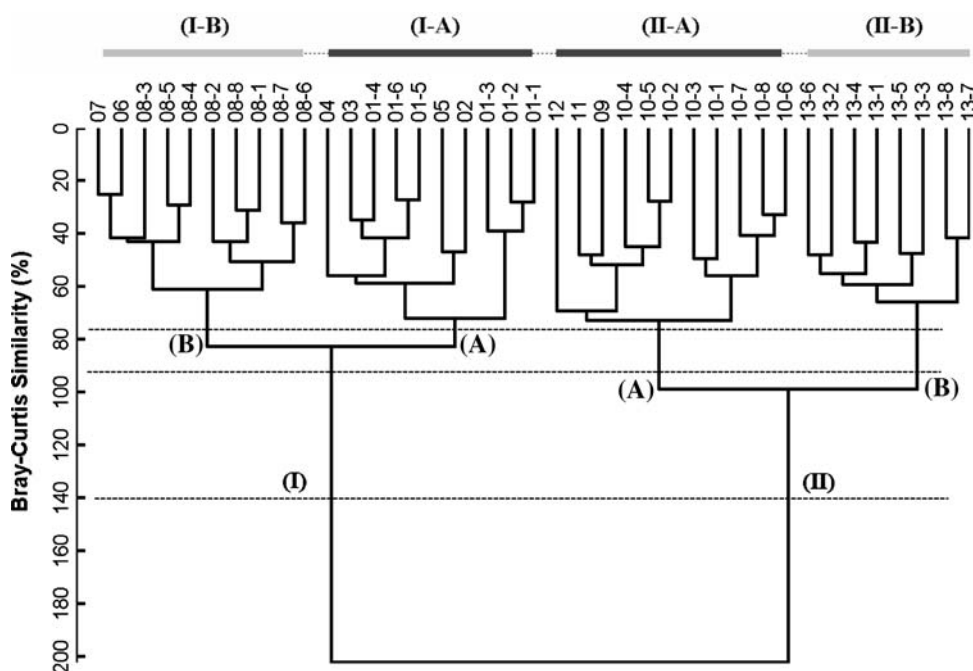
The different regional copepod communities in the East China Sea were distinguished using the first three cut-off levels of the dendrogram (Fig. 4): The first cut-off level separated the mainland China Shelf stations from the other stations in the NTS and the KCR. In the next hierarchical level, the NTS stations were separated from station 13 that was sampled for eight times during day and night in the KCR (Fig. 4). The last hierarchical level considered here allowed to split the Mainland China Shelf stations into two clusters: (1) the northern stations near the EYR, and (2) the southern stations near the EMR. The cluster analysis showed that the geographical variability in copepod composition is greater than the variability at the same stations over day and night periods. In fact, the four stations (1, 8, 10 and 13) that were sampled several times during day and night belong to four different clusters at the four-group hierarchical level (Fig. 4).

The different assemblages of copepod species corresponding to the different hierarchical levels discussed above are selected and shown in Fig. 5. For the first hierarchical subdivision in two groups, 40 species (52% of the total set of species) were retained in these assemblages. The Mainland China shelf assemblage was characterised by the harpacticoid species *Euterpina acutifrons*, two calanoid species (*Parvocalanus crassirostris* and *Acartia (Odont-*

cartia) spinicauda), the poecilostomatoid species *Corycaeus (Ditrichocorycaeus) affinis* and two cyclopoid species (*Oithona nana* and *Oithona rigida*). The indicator values of all these species were larger than 70% and at maximum for this assemblage (Fig. 5). A second assemblage of NTS and the KCR was characterised by 27 species (Fig. 5), but for only three calanoid species (*Acartia negligens*, *Clausocalanus furcatus* and *Paracalanus nanus*) the indicator values were maximal and larger than 70%. Additional 12 copepod species reached their maximal IndVal for this group (Fig. 5). In the next hierarchical level, the NTS stations were separated from the KCR stations and the IndVal of several species increased significantly and reached maximal values at this cut-off levels. The IndVals of both calanoid species *Undinula vulgaris* and *Temora discaudata* increased between the first and the second cut-off level from 67.21 and 76.24% to 89.48 and 84.69%, respectively. These species are most characteristic for this assemblage. Even if the IndVal of *Paracalanus nanus* slightly decreased at the second hierarchical level, it remains high (86.73) to consider this species as being characteristic for the NTS region. The maximal values of IndVal in this group were reached for six additional copepod species, namely *Pareucalanus attenuatus*, *Acrocalanus gracilis*, *Acrocalanus monachus*, *Paracalanus aculeatus*, *Calanopia minor* and *Corycaeus (Onychocorycaeus) pumilus* (Fig. 5). The harpacticoid *Macrosetella gracilis* and the calanoid *Lucicutia flavicornis* showed the highest individual values for the KCR assemblage (Fig. 5).

The third cut-off level allowed separating the EYR assemblage indicated by nine species from the EMR assem-

Fig. 4 Hierarchical classification of the sampling sites for planktonic copepods during 15–29 July 2001 in the East China Sea. The characteristics of the sampling sites (depth, date and time) are shown in Table 1. The first three cut-off levels are indicated by dotted lines



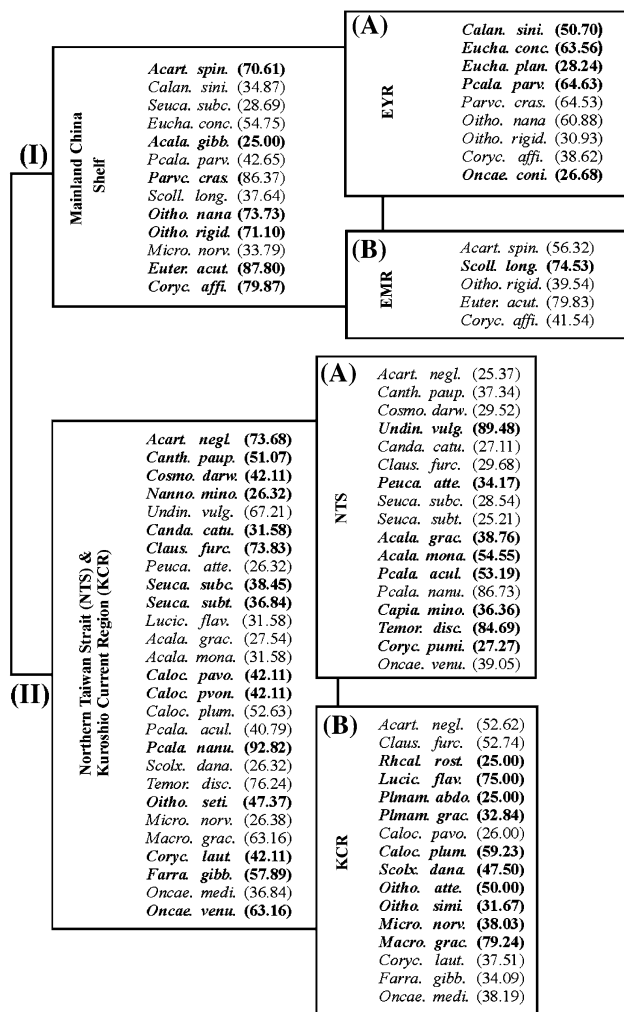


Fig. 5 Representation of the summer planktonic copepod assemblages in different regions in the East China Sea obtained from Fig. 4 (three cut-off levels). Species abbreviations shown in Table 2 are used to identify the different assemblages. The indicator values are shown between parentheses; maximal values in bold

blage indicated by five species (Fig. 5). From the indicator species of the EYR assemblage, we distinguish the northern species, i.e. *Calanus sinicus* and *Euchaeta concinna*. The species *Scolecithricella longispinosa* and *Euterpina acutifrons* are most characteristic of the EMR assemblage (Fig. 5). The spatial pattern of most indicator species for each copepod assemblage is provided in Fig. 6. The small harpacticoid *Euterpina acutifrons* (Fig. 6a) is characteristic for neritic assemblages occurring at all shallow stations along the China shelf. Its pattern of distribution is opposed to the one of *P. nanus* (Fig. 6b) at the deeper stations 9–13. However, the highest abundance of *P. nanus* was observed in the Taiwan Strait. For the higher resolution assemblages, the northern distribution of *Paracalanus parvus* (Fig. 6c) is clearly opposed to the spatial pattern of *Scolecithricella longispinosa* (Fig. 6d), being highly abundant at station 8. *Undinula vulgaris* (Fig. 6e) shows a low abundance in

northern Taiwan at this time of the year, whereas *Macrosetella gracilis* (Fig. 6f) is more abundant at the Kuroshio Current station. The high variability in the abundance of some species shown in Fig. 6 can be explained by a day–night difference in their abundance. It is worth noting that this shows a typical spatial distribution though. The indicator species of NTS and KCR (see Fig. 5) are characterised by higher abundances at stations 9 and 13, respectively.

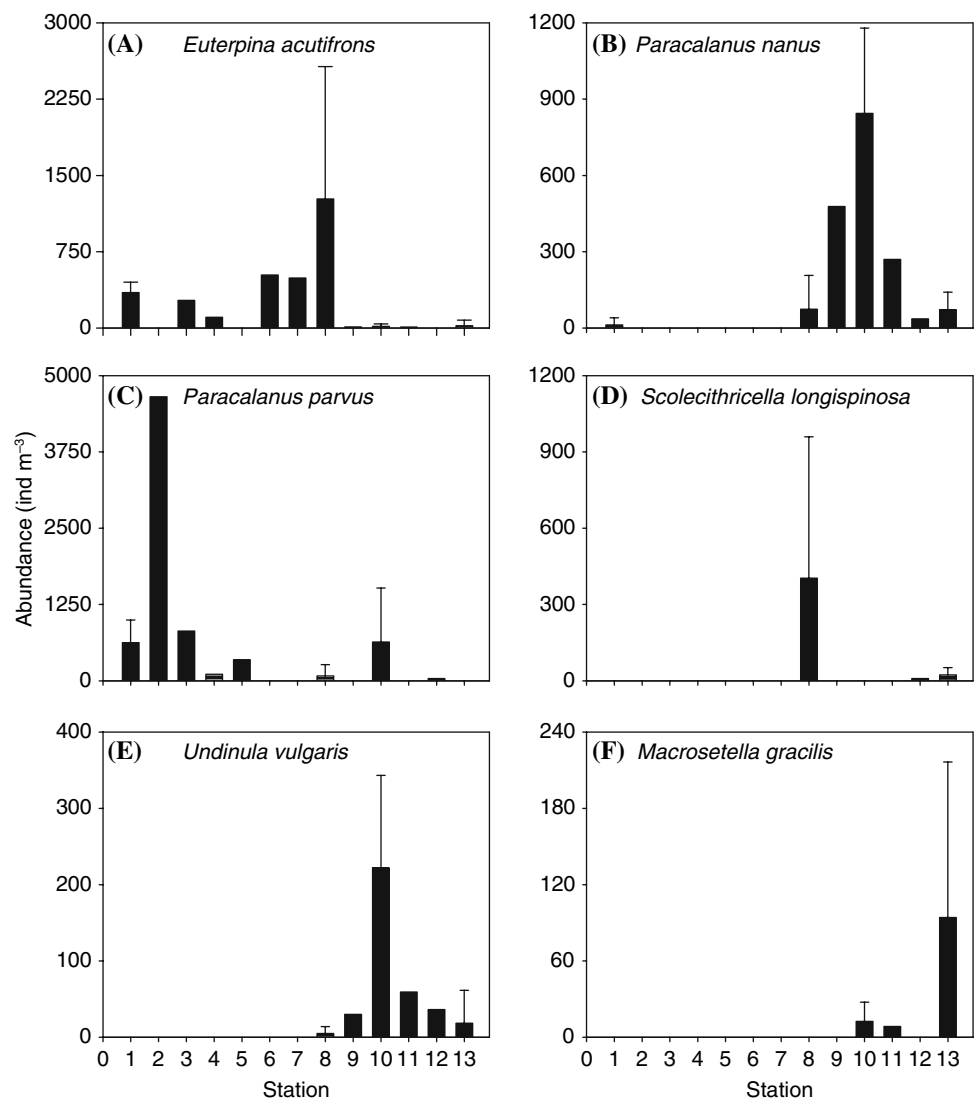
The patterns of abundance and diversity of copepods in the area studied are provided in Fig. 7. Copepod numbers were highest at the northern China shelf station 3 (12,901.20 individuals m⁻³) and lowest at station 12 (680.70 individuals m⁻³), whereas the Shannon–Wiener diversity index remained similar along the coastline of China. Species richness and diversity indices increase from the NTR to the KCR. The richness index values varied from a lowest value of 0.64 at station 2 to the highest values at station 10 (2.88 ± 0.33). It can thus be concluded that copepod species diversity is highly influenced by the complex hydrographic regime in this region. The Shannon–Wiener diversity index varied from 1.37 at station 2 to 2.78 ± 0.13 at station 13, diversity showing highest values at KCR. The evenness index varied from 0.63 at station 9 to 0.92 ± 0.03 at station 13. No other significant difference was found between stations.

Discussion

The present study provides a correlation between copepod abundances and sampling sites. Cluster results indicate that copepod species distribution patterns were correlated with water masses in the southwest ECS area. A similar study by Shih and Chiu (1998) found that the abundance of copepods was higher in the north of Taiwan than in coastal waters or areas influenced by the KC. The present results confirm that different water masses are characterized by different copepod compositions from the KC area to the north of Taiwan. Our data support the notion that copepod distribution patterns correspond to water masses reported by Shih and Chiu (1998), Shih et al. (2000), Hwang et al. (2004), Hwang and Wong (2005), and Hwang et al. (2006).

The present study found copepod densities several times higher than reported by Shih and Chiu (1998) from the southern East China Sea and by Shih et al. (2000) in upwelling regions of the western North Pacific. We explain this with the smaller mesh sized zooplankton nets and different sampling depths applied in the present study. A smaller mesh size (100 µm) clearly caused an increase of retrieved copepod abundances in the present study. Shih and Chiu (1998) and Shih et al. (2000) used 500 and 333 µm mesh size plankton nets for their studies, respectively. Such differences in sampling design could be

Fig. 6 Spatial patterns of indicator species for the sampling stations 1–13

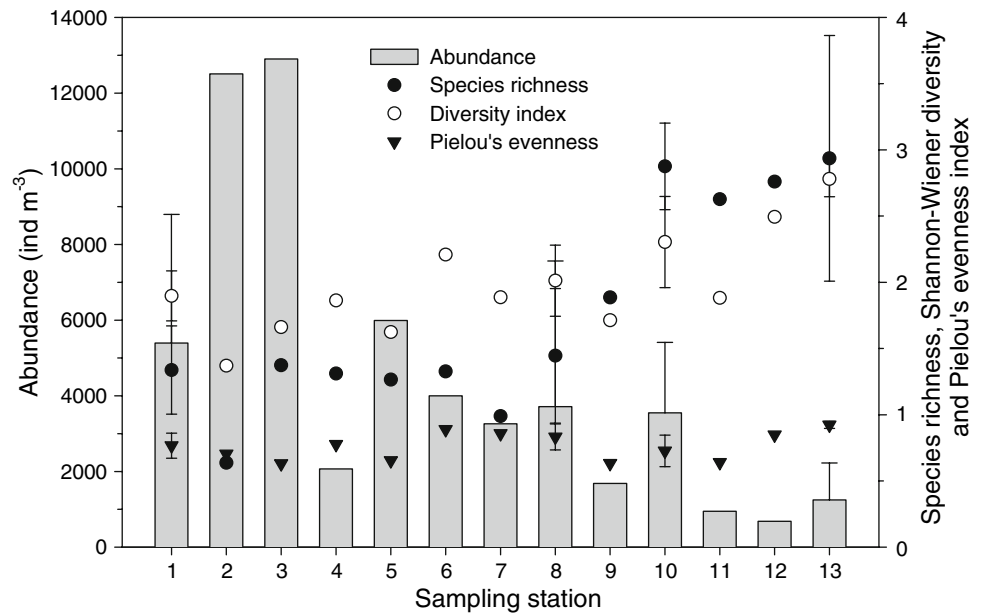


responsible for the variations in the densities obtained, and for the size composition of copepods. The use of a 100 μm mesh allowed the retrieval of several smaller mesozooplanktonic copepod species (i.e. species belonging to: *Paracalanus*, *Clausocalanus*, *Acartia*, *Oithona*, *Oncaea*, *Corycaeus*) that are known to be abundant, particularly in coastal waters (Turner 2004). Accordingly, copepod abundances in the present study were much higher than in several long-term studies (Hwang et al. 2006). Different sampling depths and sampling times affect copepod densities and species occurrences (Shih et al. 2000). In addition, change copepod abundance and composition with the diurnal migrations of plankton (Lo and Hwang 2000; Shih et al. 2000). Our samples are from shallower waters than those of Shih and Chiu (1998) and Shih et al. (2000), resulting in higher copepod abundances.

The present results of hierarchical classification of the ECS samples show that group similarity is positively correlated with the geographic distance of stations to the main-

land of China. Shih and Chiu (1998) observed that relative similarity is higher at stations nearby the mainland of China and lower at coastal stations compared to the KC (Hsieh et al. 2004). Hsieh and Chiu (1997) studied copepod assemblages in estuarine and coastal waters and found the species composition to be correlated with the geographic location of sampling stations. The properties of our stations 9–12 may be related to the fact that these stations are located close to upwelling areas. This could explain the different copepod composition relative to that of stations north of the Taiwan transect line, and the separation into the same group (group II-A, Fig. 4) by the cluster approach. Stations 1–8 belong to coastal waters from the EYR to the EMR area (Fig. 1). A similar cluster analysis for copepod composition confirms that these stations belong to the same group (group I, Fig. 4). Two different groups can be separated: a northern coastal group of stations (stations 1–5, group I-A), and a southern coastal group of stations (stations 6–8, group I-B). The stations at the northern coast close to EYR

Fig. 7 Copepod abundance, species richness, Shannon-Wiener diversity index and Pielou's evenness index during 15–29 July 2001 in the East China Sea. Only those stations sampled for several times during the day and at night are provided with error bars (=standard deviations)



and the stations of the southern coast group close to EMR belong to different water masses, which are indicated by different copepod assemblages. We conclude from AVHRR temperature images (Fig. 2a) that the EYR intrusion waters cannot affect the EMR area. At the first hierarchical level, NTS and KCR stations constituted a single assemblage composed of 77 different copepod species (Fig. 4). For these areas close to Taiwan it is possible to compare our findings with other studies from the northern coast of Taiwan. In the present study, the number of species observed in the southwest East China Sea area is higher than that recorded by Hwang et al. (1998) from the coastal waters off the northern tip of Taiwan in summer where these authors identified 25 species of calanoid copepods. Dominant species that were found at all stations were *Acrocalanus gracilis*, *Canthocalanus pauper*, *Temora discaudata*, *Temora turbinata* and *Undinula vulgaris*. Wong et al. (1998) identified 37 species of calanoid copepods near a nuclear power plant area in northern Taiwan and found *A. gracilis* as being dominant there (see also Hwang et al. 2004). The main copepod species found by Hwang et al. (1998) and Wong et al. (1998) in the summer season in northern Taiwan were also found in the present study. Other reports from the northern Taiwan Strait (Hsieh and Chiu 2002) and the upwelling area of northern Taiwan (Lo et al. 2004) confirmed higher species numbers than in the present study. A 5-year study by Hwang et al. (2006) showed similar results. Here, a total of 110 copepod species were reported and their distribution patterns were related to the NE and SW monsoons (Hwang et al. 2006). Diversity estimates of copepod species in other studies of this geographical sector are very variable due to different seasons or years making a comparison with previous studies difficult. The present

study provides species numbers higher than those in coastal waters of southwest Taiwan (67 species by Lo et al. 2001), but lower than those in the northern South China Sea (78 species by Hwang et al. 2000).

Reports of species composition show a remarkable variability at temporal scales, such as seasonal or monthly differences (Hsieh and Chiu 1997). Comparing the species composition found during the present study with those of Shih and Chiu (1998), Chen and Zhang (1965) and Chen et al. (1974), all copepod species found in previous studies were claimed to be characteristic for the southern East China Sea.

All members of the copepod communities identified in the present study were found in reports of other studies from the ECS (Chen and Zhang 1965; Chen et al. 1974; Shih and Chiu 1998). In addition, copepod communities from the ECS were dominated by calanoid species that were reported from several localities around Taiwan (Hsieh and Chiu 1997, 2002; Hwang et al. 1998, 2000; Lo and Hwang 2000; Lo et al. 2001; Shih and Chiu 1998; Shih et al. 2000; Tseng 1975, 1976; Wong et al. 1998; Hwang 2004; Hwang et al. 2006; Dur et al. 2007). According to the classical copepod survey by Shih and Young (1995), almost all species reported from adjacent seas of Taiwan are wide spread in the East and the South China Sea. However, the majority of recorded copepod species were not widely distributed, with 13 species found in only one sampling area. No species occurred at all stations and in more than 70% of the samples. The calanoid *P. crassirostris* with a 69.23% occurrence ratio and an average abundance of 626.88 individuals m⁻³ was the most frequently occurring and most abundant species in all samples. Lo et al. (2004) surveyed only the upwelling area of northern Taiwan and

found that *P. crassirostris* belonged to one of the three most abundant species. Our data confirms this species to be dominant in northern Taiwan. Its dominance on the coastal shelf extends to northern Taiwan where it still shows high abundances. Through station 13 in the upwelling area, the density of *P. crassirostris* decreases and disappears completely at KCR. We suppose that this species prefers low salinities and is wide spread along the coast of the mainland of China and in the southwest ECS. The calanoid *P. crassirostris* can be an important indicator-species in the ECS. The relative occurrence of copepod species in the present study was similar to that found during April in the southern ECS north of Taiwan (Shih and Chiu 1998). Three species, *C. sinicus*, *E. rimana*, and *P. aculeatus* were the most abundant species among the top 20, corresponding to the results of the present study. This demonstrates the dominance of these three copepod species from April to July in this area. It is reasonable to assume that the composition of copepod assemblages follows a seasonal shift, primarily affected by the TS Current and annual monsoonal changes (Chen 1992; Chen et al. 1998). The dominant species of the NTS station were similar in their temporal occurrence and locations to these previous reports.

There are some results of the present study that deserve to be emphasized here, e.g., the highest copepod abundance during the present survey occurred at stations 3 and 4 along the coast. Water mass characteristics at these two stations were: low salinity and low temperature (Fig. 2). Major contributing species were *P. parvus*, *P. crassirostris* and *O. nana*. We speculate these species to tolerate low salinity and low temperature conditions (Chen and Zhang 1965). However, these species were also dominant in coastal areas and occurred there at high temperature and low salinities (Chen and Zhang 1965). In the present study, *C. sinicus* appeared at all stations except the KCR station. This species is characteristic of the East China Sea as reported in previous papers (Chen 1992; Shih and Chiu 1998; Shih et al. 2000; Souissi et al. 2004; Hwang and Wong 2005). Chen (1992) argues that *C. sinicus* is one of the seven most dominant copepods in the ECS continental waters. Shih and Chiu (1998) found it to be dominant in the ECS and few individuals could be found in the Kuroshio region, which confirmed the observation of Chen (1992). When Shih et al. (2000) surveyed an upwelling area (near our station 13), they rarely found *C. sinicus* in their samples. Our result confirms that *C. sinicus* is abundant in the ECS, rare in mid-way stations and absent near the Kuroshio Current (Hwang and Wong 2005; Hwang et al. 2004, 2006). Considering that the Kuroshio Current flows along the east coast towards the north to Japan, we suppose that water masses begin to mix and exchange with the ECS water mass edge in the northeast of the Taiwan upwelling regions. That might be the reason why Shih and Chiu (1998) found

C. sinicus distributed in the Kuroshio Current area of the warm water mass, at a station north of our KCR station.

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