

## Copepod Diversity and Composition as Indicators of Intrusion of the Kuroshio Branch Current into the Northern Taiwan Strait in Spring 2000

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**Chih-Hao Hsieh, Tai-Sheng Chiu and Chang-tai Shih (2004)** Copepod diversity and composition as indicators of the intrusion of the Kuroshio Branch Current into the northern Taiwan Strait in spring 2000. *Zoological Studies* 43(2): 393-403. This study was carried out during the waning of the northeastern monsoon, at which time the Kuroshio Branch Current intrudes into the northern Taiwan Strait. The Kuroshio Branch Current was characterized by high temperature and salinity in contrast to the low temperature and salinity of the China Coastal Current. A front was induced south of the Haitan Island, which resulted in a phytoplankton bloom when the China Coastal Current and Kuroshio Branch Current met. The front is marked by a temperature of  $18 \pm 2$  °C and a salinity of  $33 \pm 0.5$  psu. The intrusion of the Kuroshio Branch Current apparently influenced the species diversity of copepods. Species richness and diversity were higher in the area affected by the Kuroshio Branch Current than that affected by the China Coastal Current. No significant differences in the copepod abundances between the two water masses were observed. Species assemblages were distinct in the two water masses. The China Coastal Current contained only coastal/neritic cold-water species, while the Kuroshio Branch Current consisted of coastal, neritic, and oceanic warm-water species. Indicator species of each water mass were identified and are discussed. <http://www.sinica.edu.tw/zool/zoolstud/43.2/393.pdf>

**Key words:** China Coastal Current, Kuroshio Branch Current, Indicator, Species diversity, Species composition.

The Taiwan Strait is a shallow channel connecting the East China Sea and South China Sea and serves as an important pathway for faunal exchange between these two waters in the margin of the western North Pacific. It is also a main route for migratory fishes and thus an important fishing ground. Three main currents, the China Coastal Current (CCC), Kuroshio Branch Current (KBC), and South China Sea Surface Current (SCSSC), all driven by the monsoon system, affect the hydrographic conditions of the Strait (Jan et al. 2002). When the northeastern monsoon prevails during winter (Nov. to Mar.), the CCC moves southward into the Strait; meanwhile, the monsoon also forces the KBC to penetrate into the Strait through the Penghu Channel (Wang and

Chern 1988). Stagnated by the CCC, the KBC is blocked south of the Changyun Ridge when the northeastern monsoon is strong. Only when the northeastern monsoon wanes in the spring, does the KBC flow over the Changyun Ridge or move northwestward along the local isobaths into the northern Strait. A front south of the Haitan Is. is induced when the northwardly flowing KBC encounters the retreating CCC. During summer (June to Aug.), the southwestern monsoon causes the SCSSC to move northward into the Strait in place of the KBC. The seasonal variation in circulation of the Taiwan Strait was described by Jan et al. (2002). The physical oceanography of the intrusion of the KBC into the Strait during the waning of northeastern monsoon has been well stud-

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ied (Jan et al. 1998); however, the biological effects of this circulation are not known.

The CCC is a neritic water mass characterized by low salinity, low temperature, and high nutrient due to river runoff from the mainland China (Liu et al. 2000). The KBC, which originates from the oceanic Kuroshio Current, has high temperature and high salinity, but counter-intuitively contains a nutrient level comparable to that of the CCC (Chung et al. 2001). It is known that zooplankton composition and abundance are affected by the properties of water masses (Boucher et al. 1987, Kouwenberg 1994, Lopes et al. 1999). Since our samples were collected across both the CCC and KBC, this survey allowed us to test the following hypotheses: (i) distinct species compositions occur in the CCC and KBC; (ii) low species diversity occurs in the CCC (a neritic water mass subject to frequent disturbances) and high diversity occurs in the KBC (an oceanic water mass with greater stability); and (iii) equal abundances of copepods occur in the CCC and KBC according to their similar nutrient contents.

Previous faunal surveys in the Taiwan Strait have been either confined to a small area or conducted without attention to hydrographic effects. Tzeng and Wang (1992 1993) studied the composition and seasonal variation of larval fish fauna in the Tanshui River estuary off northwestern Taiwan; Chiu and Chang (1995) reported diurnal cycling in the vertical distribution of larval fishes at a fixed station in the northeastern Strait; Chiu and Chang (1994) compared the larval fish compositions during winter and spring in the northeastern Strait; Chiu (1992) and Chiu and Chen (1997) examined the larval fish composition in relation to the hydrography in the southern Strait. The copepod fauna was reported by Tan (1967) around the Tanshui River estuary; Hsieh and Chiu (1998) compared the species assemblages and abundances of copepods in the Tanshui River estuary and adjacent area; Zheng et al. (1982) and Chen (1992) reported copepod species in the Strait. A relatively large-scale survey including benthic, planktonic, and larval fish faunas in the Strait was conducted by Zhu et al. (1988), but was limited to the western Strait and hydrographic effects on biological variation were not examined. The present study is thus a preliminary survey of copepods covering the northern Taiwan Strait and analyzes hydrographic effects on copepod assemblages. Copepods are a major component of marine zooplankton and play an important role in marine food webs (Runge 1988, Poulet and Williams 1991, Williams et al.

1994). Studies on copepods are crucial for understanding the dynamics of marine ecosystems and are necessary for purposes of management and conservation of marine resources. The objectives of our study are to provide basic information about the copepod fauna and to determine the effects of the Kuroshio intrusion in the Taiwan.

## MATERIALS AND METHODS

### Sampling

A cruise was carried out on board the *Ocean Research I* on 20~25 Mar. 2000 in the Taiwan Strait (Fig. 1). Copepods were sampled using a standard North Pacific zooplankton net with a mouth diameter of 45 cm and mesh size of 150  $\mu\text{m}$ , and samples were preserved in seawater with 5% formaldehyde. A flowmeter was mounted at the center of the net mouth to record the volume of water filtered. The net was towed obliquely from near the bottom to the surface. Vertical profiles of temperature, salinity, fluorescence, and dissolved oxygen were recorded at each station from the bottom to the surface with a conductivity-temperature-depth (CTD) profiler (Sea-Bird Electronics, Bellevue, USA) equipped with a fluorometer. The fluorescence value was not calibrated but used as a reference index of the relative phytoplankton biomass. During the cruise, plankton samples were collected at only 10 of 36 stations, because the ship time allocated for our study was limited.

In the laboratory, copepods were sub-sampled using a Folsom splitter until the sample size was reduced to 300~500 specimens. Copepods (including copepodid stages) were identified to species whenever possible. The abundance of copepods was expressed as the number of individuals/ $\text{m}^3$ .

### Data analysis

Shannon's diversity index was used to measure the species diversity of each sampling station. A correspondence analysis based on the relative abundance of the 20 most abundant species was applied to simultaneously examine the station associations and the corresponding species (Pielou 1984). For reducing the patchiness effect of zooplankton, data were square-root transformed (Krebs 1989). K-means classification based on the station scores of the correspondence analysis was applied to distinguish station groups

(Wilkinson 1990). By using K-means classification, the number of groups was decided in advance (in the current case, the number of groups was two), and an iterative algorithm was used to assign stations to the groups by maximizing the variance (of input variables) between groups and minimizing the variance within groups. This procedure results in an objective classification of stations. Indicator species of each station groups were extracted by an indicator species analysis (Dufrene and Legendre 1997). The indicator value (Ind Val) of each species was computed as follows:

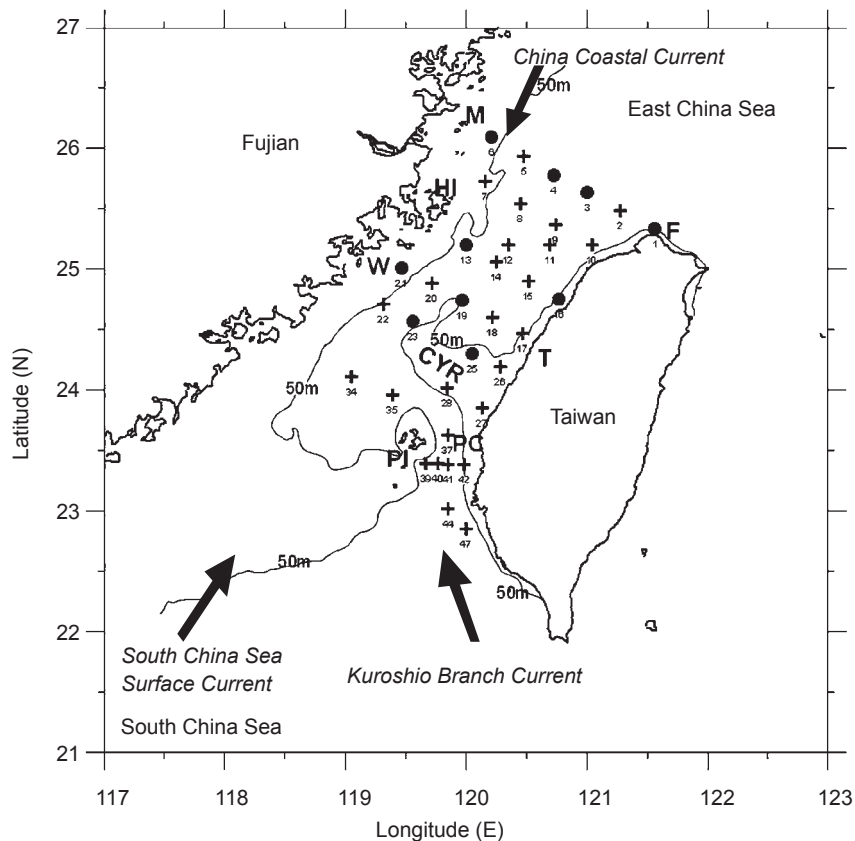
$$\text{Ind Val} = \text{RA}_{kj} \times \text{RF}_{kj} \times 100;$$

where  $\text{RA}_{kj}$  is the relative abundance of species  $j$  in group  $k$ , and  $\text{RF}_{kj}$  is the relative frequency (presence/absence) of species  $j$  in group  $k$ . The highest indicator value for a given species across groups was saved as a summary of the overall indicator

value of that species. A Monte Carlo method was used to test the significance of their indicator values by randomly permuting the data and recalculating Ind Val. One thousand permutations were made to compare the observed Ind Val and recalculated Ind Val. The result was deemed significant when the observed Ind Val fell in the 5% upper tail. Only individuals identified to species were included in the above analyses. A Mann-Whitney U test was used to test for differences in copepod abundance, species richness, and diversity between the station groups affected by the China Coastal Current and Kuroshio Branch Current.

## RESULTS

The two water masses have distinctive temperature-salinity (T-S) properties (Fig. 2). Stations 1, 3, 4, 6, and 21 characterized by low temperature and salinity were influenced by the CCC, while sta-



**Fig. 1.** Map showing sampling stations and isobaths in the Taiwan Strait. Crosses indicate stations with CTD data only; filled circles indicate stations with both CTD and copepod samples; arrows indicate the main currents during the cruise; HI, Haitan Island; PI, Penghu Island; PC, Penghu Channel; CYR, Changyun ridge; F, Cape Fuguei; M, Matsu; W, Wuchiu; T, Taichung.

tions 16, 19, 23, and 25 characterized by high temperature and salinity indicated intrusion of the KBC. Station 13 was at the transitional location with a broad range of temperature and salinity. The T-S curve of station 41 (no plankton sample, c.f. Fig. 1) was used to highlight the intrusion of the KBC into the Penghu Channel. Typical T-S curves<sup>1</sup> of the CCC, KBC, and SCSSC were also plotted in figure 2 for comparison. The SCSSC is distinct from the KBC by having higher temperature and lower salinity in the surface layer. This distinction provides supportive evidence that our samples were taken during the intrusion of the KBC. Since the purpose of this work was to compare the CCC and KBC, distinguishing the KBC and SCSSC is not discussed.

<sup>1</sup>The typical T-S curves represent characteristics of the three water masses (Jan, in prep.). The KBC was taken on 01/05/1991 in the central Penghu Channel, the SCSSC was taken on 08/16/1999 off the southwestern Taiwan, and the CCC was taken on 03/21/2000 in the northwestern Taiwan Strait. Note that coincidentally the chosen typical CCC is the station 6 of this study.

The distinctive properties of the CCC and KBC were clarified by observing surface water temperature and salinity. From contour plots of surface water temperature and salinity (Fig. 3a, b), one can observe the high-temperature and -salinity water of the KBC flowing over the Changyun Ridge (see Fig. 1) and penetrating into the northern Taiwan Strait. Dense contour lines of temperature and salinity south of the Haitan Is. indicate that a front had formed, and that the front resulted in a phytoplankton bloom represented by high fluorescence values (Fig. 3c). Furthermore, profiles of temperature and salinity of the northern transect (Matsu-Cape Fuguei) showed a high-temperature and -salinity core at a distance 60~80 km from the western end, which indicated that the influence of the KBC had extended to the northern Strait (Fig.

4a, c). The extreme low-temperature and -salinity water along the China coast was the southward extension of the CCC and river runoff from Fujian Province (Fig. 4).

In our samples, 70 species of copepods were identified, belonging to 30 genera, 17 families, and 4 orders (only individuals identified to species were counted) (Table 1), and some clarification on copepod taxonomy is provided in the footnotes. Species richness and diversity were higher in the area influenced by the KBC than by the CCC (Fig. 5, Table 1). The correspondence analysis resulted in two station clusters through K-means classification, namely the CCC and KBC (Fig. 6), and the first 2 axes explained 78.11% of the variance. The CCC included stations 1, 3, 4, 6, and 21 with corresponding species of *Calanus sinicus*, *Corycaeus (Ditrichocorycaeus) affinis*, *Oithona similis*, and *Paracalanus parvus*, while the KBC included stations 13, 16, 19, 23, and 25 with corresponding species of *Acartia negligens*, *Acrocalanus gracilis*, *Calocalanus gracilis*, *Cal. plumulosus*, *Cosmocalanus darwini*, *Clausocalanus furcatus*, *Cl. minor*, *Euterpina acutifrons*, *Lucicutia flavicornis*, *Oithona plumifera*, *Oncaea venusta*, *Paracalanus aculeatus*, *Parvocalanus crassirostris*, *Subeucalanus mucronatus*, *Temora stylifera*, and *T. turbinata*. The indicator species of the two water masses with significant indicator values are shown in Table 2. Only one species, *Corycaeus affinis*, belonged to the CCC, and 16 species of the KBC were recognized as indicator species. Species with higher indicator values are more representative of the water mass, and a value of 100 indicates a perfect indicator. Two species, *Calanus sinicus* and *Oithona similis*, belonging to the CCC and one species, *Clausocalanus arcuicornis*, belonging to the KBC with marginal significance, are also listed in table 2. The average copepod abundance was 1368 individuals/m<sup>3</sup> in the CCC and 1296 individuals/m<sup>3</sup>

**Table 1.** Copepod species composition (%), species richness, diversity, and abundance, and environmental information at each sampling station. (only individuals identified to species were included in the measurement of species richness and diversity)

Family Species	1	3	4	6	13	16	19	21	23	25
ORDER: CALANOIDA										
ACARTIIDAE										
<i>Acartia negligens</i>	0	0	0	0	0	0.22	2.34	0	2.51	1.62
CALANIDAE										
<i>Calanus sinicus</i>	14.26	7.01	2.38	9.04	5.56	1.76	1.82	17.15	3.23	8.12

Table 1. (Cont.)

Family Species	1	3	4	6	13	16	19	21	23	25
<i>Canthocalanus pauper</i>	0	0	0	0	0.38	0.22	1.04	0	0.72	0.32
<i>Cosmocalanus darwinii</i>	0	0	0	0	0	0	0	0	2.87	0.65
<i>Mesocalanus tenuicornis</i>	0	0	0	0	0	0	0.26	0	0	0
<i>Nannocalanus minor</i>	0	0.19	0	0	0	0.22	0.26	0	0	0.32
<i>Undinula vulgaris</i>	0	0	0	0	0	0	0.26	0	0.72	0
CANDACIIDAE										
Candaciidae copepodid	0	0	0	0	0	0	0.26	0	0	0.32
CLAUSOCALANIDAE										
<i>Clausocalanus arcuicornis</i>	0	0.19	0	0	0	0.44	0.52	0	0.36	0.32
<i>Clausocalanus furcatus</i>	0	0	0	0	3.26	0	1.56	0	2.87	2.27
<i>Clausocalanus mastigophorus</i>	0	0	0	0	0	0.44	0	0	0.36	0
<i>Clausocalanus minor</i>	0	0	0	0	1.34	1.32	2.34	0	1.08	2.92
EUCALANIDAE										
<i>Pareucalanus attenuatus</i>	0	0	0	0	0	0	0	0	0.36	0.32
<i>Rhincalanus rostrifrons</i> <sup>1</sup>	0	0	0	0	0	0	0	0	0.72	0
<i>Subeucalanus crassus</i>	0	0	0	0	0	0	0	0	0.36	0.32
<i>Subeucalanus mucronatus</i>	0	0	0	0	0	0.88	1.04	0	1.79	0.65
Eucalanidae copepodid	0	0	0	0	0.38	0	0	0	0.72	3.57
EUCHAETIDAE										
<i>Euchaeta concinna</i>	0	0	0	0	0	0	0.26	0	0	0
<i>Euchaeta plana</i>	0	0.19	0	0.6	0	0.22	0	0	0	0
<i>Euchaeta rimana</i> <sup>2</sup>	0	0	0	0	0.19	0	1.04	0	1.08	0
<i>Euchaeta spinosa</i>	0	0	0	0	0	0	0	0	0	0.32
Euchaetidae copepodid	0.4	0.76	0.95	0.6	1.53	4.18	4.95	0	5.73	0.65
LUCICUTIIDAE										
<i>Lucicutia flavicornis</i>	0	0	0	0	0	1.1	0.78	0	0.72	0.32
<i>Lucicutia gaussae</i>	0	0	0	0	0	0	0	0	0.36	0
METRINIDINIDAE										
<i>Pleuromamma abdominalis</i>	0	0	0	0	0	0	0	0	0	0.32
<i>Pleuromamma gracilis</i>	0	0	0	0	0	0.22	0	0	0	0
PARACALANIDAE										
<i>Acrocalanus gibber</i>	0	0	0	0	0.77	1.54	2.6	0	0.72	2.6
<i>Acrocalanus gracilis</i>	0	0	0	0	0	0	0	0	0	0.32
<i>Calocalanus contractus</i>	0	0	0	0	0	0	0.78	0	0	0
<i>Calocalanus gracilis</i>	0	0	0	0	0	0.22	1.04	0	2.87	0
<i>Calocalanus monospinus</i>	0	0	0	0	0	0	0	0	0.72	0
<i>Calocalanus pavoninus</i>	0	0	0	0	0	0	0	0	1.08	0
<i>Calocalanus plumulosus</i>	0	0	0	0	0	0.66	0.26	0	0.36	1.62
<i>Calocalanus styliremis</i>	0	0	0	0	0	0	0.26	0	1.08	0
<i>Paracalanus aculeatus</i>	0	0.19	0	0	3.45	6.81	5.99	7.56	7.17	7.47
<i>Paracalanus parvus s.l.</i>	29.32	55.3	59.62	64.76	42.53	70.33	40.89	32.56	24.73	41.56
<i>Parvocalanus crassirostris</i>	0	0	0	0	0.38	1.1	2.6	0	2.15	2.6
<i>Parvocalanus sp.</i>	0	0	0	0	0	0	0	0	0	0.32
PONTELLIDAE										
Pontellidae copepodid	0	0	0	0	0.19	0	0.26	0	0	0.65
SCOLECITRICHIDAE										
<i>Scolecithricella longispinosa</i>	0	0	0	0	0	0.22	0.26	0	0	0
<i>Scolecithricella sp.</i>	0	0	0	0.3	0	0	0	0	0	0
<i>Scolecithrix danae</i>	0	0	0	0	0	0.22	0.26	0	1.08	0.32
TEMORIDAE										
<i>Temora discaudata</i>	0	0	0	0	0	0	0	0	0.36	0
<i>Temora stylifera</i> <sup>3</sup>	0	0	0	0	1.34	0.44	1.3	0	1.08	4.55
<i>Temora turbinata</i>	0	0	0	0	0.57	0.22	0.78	0.87	0	0.32
ORDER: CYCLOPOIDA										
OITHONIDAE										
<i>Oithona atlantica</i>	0	0	0	0	0	0	0.26	0	0.36	0
<i>Oithona attenuata</i>	0	0	0	0	0.38	0	0	0	0	0



Table 1. (Cont.)

Family Species	1	3	4	6	13	16	19	21	23	25
<i>Oithona brevicornis</i>	0	0	0	0	0.19	0	0	0	0	0
<i>Oithona decipiens</i>	0	0	0	0	0	0	0.26	0	0.36	0.65
<i>Oithona fallax</i>	0	0	0	0	0	0	0	0	0	1.3
<i>Oithona fragilis</i>	0	0	0	0	0	0	0	0	0.36	0
<i>Oithona plumifera</i>	0	0	0	0	0.19	0.22	6.77	0	12.9	2.92
<i>Oithona similis</i>	1.61	18.75	8.08	18.37	14.94	1.32	1.82	24.71	0	1.62
<i>Oithona simplex</i>	0	0	0	0	0.19	0	0	0	0	0
ORDER: HARPACTICOIDA										
CLYTEMNESTRIDAE										
<i>Clytemnestra rostrata</i>	0	0	0.24	0.3	0	0	0.26	0	0	0.32
EUTERPINIDAE										
<i>Euterpina acutifrons</i>	0	0	0	0	0.57	0	2.6	0	1.08	2.27
MIRACIIDAE										
<i>Macrosetella gracilis</i>	0	0	0.24	0	0.57	0	0	0	0	0
ORDER: POECILOSTOMATOIDA										
CORYCAEIDAE										
<i>Corycaeus (Agetus) flaccus</i>	0	0	0	0	0	0	0.26	0	0	0
<i>Corycaeus (Corycaeus) speciosus</i>	0	0	0	0	0	0	0	0	1.08	0
<i>Corycaeus (Ditrichocorycaeus) affinis</i>	54.02	16.67	28.27	3.92	2.3	0	1.3	16.86	0	0
<i>Corycaeus (Ditrichocorycaeus) dahli</i>	0	0	0	0	0.77	0	0	0	0.36	0.32
<i>Corycaeus (Ditrichocorycaeus) erythraeus</i>	0	0	0	0	0	0	0.52	0	0	0
<i>Corycaeus (Ditrichocorycaeus) subtilis</i>	0	0	0	0	0	0	0	0	0.72	0
<i>Corycaeus (Onychocorycaeus) pacificus</i>	0	0	0	0	0.96	0.22	0	0	0	0.32
<i>Corycaeus (Urochocorycaeus) furcifer</i>	0	0	0	0	0	0	0	0	0.36	0
<i>Farranula gibbula</i>	0	0	0	0	0.38	0	0	0	2.15	0
<i>Farranula rostrata</i>	0	0	0	0	0.19	0	0	0	0	0
<i>Corycaeidae copepodid</i>	0	0	0	0	8.81	2.2	1.56	0	1.79	1.62
ONCAEIDAE										
<i>Oncaea media</i>	0	0.19	0	0	0.19	0	0	0	1.08	0
<i>Oncaea mediterranea</i>	0	0	0	0	0.96	0.22	0.78	0	0.72	0
<i>Oncaea venusta</i>	0.4	0.38	0	2.11	6.32	2.64	5.73	0.29	6.09	2.6
<i>Oncaea</i> sp1	0	0	0	0	0.19	0	0	0	0	0
<i>Oncaea</i> sp2	0	0	0.24	0	0	0	0	0	0	0
<i>Triconia conifera</i> <sup>4</sup>	0	0.19	0	0	0	0	0.78	0	0	0
<i>Triconia minuta</i>	0	0	0	0	0	0.22	0	0	0	0
<i>Triconia similis</i>	0	0	0	0	0	0	0.78	0	0	0
SAPPHIRINIDAE										
<i>Copilia mirabilis</i>	0	0	0	0	0	0	0	0	0	0.32
<i>Sapphirina gemma</i>	0	0	0	0	0	0	0	0	0.36	0
<i>Sapphirina ovatolanceolata</i>	0	0	0	0	0	0	0	0	0.36	0
No. of species	5	11	6	7	26	27	39	7	43	33
Shannon's diversity	1.059	1.22	0.986	1.067	1.933	1.212	2.443	1.567	2.878	2.295
Density (ind./m <sup>3</sup> )	1119	1153	1719	855	1246	2123	1159	1497	1351	960
Depth (m)	70	80	72	41	56	50	46	35	60	31
Sea surface temperature (°C)	14.71	13.56	13.66	10.55	13.67	18.70	21.31	13.08	22.52	21.84
Sea surface salinity (psu)	32.63	32.11	32.27	29.56	31.22	33.43	34.35	30.42	34.42	34.40

1. Bowman (1971) recognized the two forms of *Rhincalanus cornutus*, f. *typica* from the Indian and Pacific oceans, and f. *atlantica* from the Atlantic Ocean as two species, based on the difference in the structure of the 5th legs as well as the relative length of distal segments of the 1st antenna of female. He retained the species name for the Atlantic form and reinstalled the Indopacific form as *Rhincalanus rostrifrons*, which was already recognized by Dana (1853).

2. Bradford (1974) found the so-called *Euchaeta marina*.

3. According to Fleminger and Hulsemann (1973), *Temora stylifera* is endemic to the Atlantic Ocean. However, the specimens of *T. stylifera* in the current study are consistent with those described in Chen and Zhang (1965). Whether *T. stylifera* also distribute in the Pacific Ocean or the species described in Chen and Zhang (1965) is another unknown new species is a question that required further study.

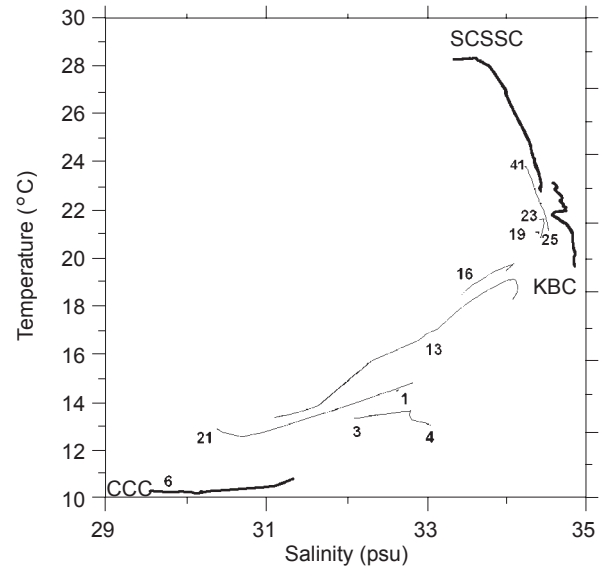
4. Bottger-Schnack (1999) established the new genus *Triconia*, and separated it from *Oncaea*.

in the KBC. There was no significant difference in abundance between these two water masses ( $U = 10.000$ ,  $p = 0.602$ ). The species richness and diversity were higher in the KBC than those in the CCC ( $U = 0.000$ ,  $p = 0.009$  and  $U = 2.000$ ,  $p = 0.028$  respectively).

## DISCUSSION

Intrusions of the Kuroshio Branch Current (KBC) into the northern Taiwan Strait during the waning of the northeastern monsoon have been described by Jan et al. (1998 2002). According to their observations from 1985 to 1992 and simulation studies, they found a recurring pattern of the KBC intruding into the retracting China Coastal Current (CCC) during spring, which was generally defined as occurring in Apr. to May. Although the present study was conducted in late Mar., the observed hydrography is consistent with the results of Jan et al. (2002), i.e., the presence of the high-temperature and -salinity KBC intruding into the low-temperature and -salinity CCC (Figs. 3, 4). The front induced by the interaction of the KBC and CCC is marked by the temperature of around  $18 \pm 2^\circ\text{C}$  and salinity of around  $33 \pm 0.5$  psu (Fig. 3a, b). The temperature and salinity properties showed a strong contrast between the CCC

and KBC (Figs. 2, 3a, b). The hydrographic pattern was clear, but the temperature and salinity values should be treated with caution since long-term variations in oceanographic properties in the Taiwan Strait are still under examination (Jan,



**Fig. 2.** Temperature-salinity diagram of copepod sampling stations. Station 41 represents the temperature-salinity curve in the Penghu Channel; KBC, Kuroshio Branch Current; SCSSC, South China Sea Surface Current; CCC, China Coastal Current.

**Table 2.** Indicator species of China Coastal Water and Kuroshio Branch Water. (Ind Val, indicator value; p-value, probability of Type-I error through Monte-Carlo test; \*, species with only marginal significance)

China Coastal water			Kuroshio Branch Water		
Species	Ind Val	p-value	Species	Ind Val	p-value
<i>Corycaeus (Ditrichocorycaeus) affinis</i>	97	0.004	<i>Acrocalanus gibber</i>	100	0.004
* <i>Calanus sinicus</i>	70.4	0.073	<i>Canthocalanus paupur</i>	100	0.004
* <i>Oithona similis</i>	74.9	0.082	<i>Clausocalanus minor</i>	100	0.004
			<i>Oithona plumifera</i>	100	0.004
			<i>Parvocalanus crassirostris</i>	100	0.004
			<i>Temora stylifera</i>	100	0.004
			<i>Oncaea venusta</i>	90	0.004
			<i>Paracalanus aculeatus</i>	83	0.032
			<i>Acartia negligens</i>	80	0.038
			<i>Calocalanus plumulosus</i>	80	0.038
			<i>Clausocalanus furcatus</i>	80	0.045
			<i>Euterpina acutifrons</i>	80	0.045
			<i>Lucicutia flavicornis</i>	80	0.038
			<i>Oncaea mediterranea</i>	80	0.044
			<i>Scolecithrix danae</i>	80	0.038
			<i>Subeucalanus mucronatus</i>	80	0.038
			* <i>Clausocalanus arcuicornis</i>	70	0.081

pers. comm.).

No significant differences in copepod abundance between the areas affected by the CCC and KBC were detected, which is consistent with the studies of nutrient fluxes (Liu et al. 2000, Chung et al. 2001). The front induced south of the Haitan Island resulted in a phytoplankton bloom (Fig. 3c), but no corresponding spatial pattern of zooplankton was observed. However, we have a limited sample size and do not have in situ rate measurements of phytoplankton and zooplankton. Therefore, no attempt was made to explain the trophic links among nutrients, phytoplankton, and zooplankton.

The species richness and diversity of copepods were higher in the area affected by the KBC than that by the CCC (Fig. 5). The Kuroshio is an oceanic current with a more-stable environment for plankton compared with China coastal water, which is subject to disturbance and nutrient enrichment from the land. The fluctuating coastal environment might be the reason why we only saw a few coastal and neritic species in the CCC (Table 1). The study by Shih and Chiu (1998) in the water north of Taiwan also showed a similar trend that copepod diversity decreases from the Kuroshio area toward the China coastal area. A similar coastal-oceanic gradient of copepod diversity was also observed in the waters off Rio de Janeiro, southeast Brazil (Lopes et al. 1999). On the other hand, the intrusion of the KBC brought warm-water oceanic species into the Taiwan Strait,

added to the coastal species that can tolerate high temperature and salinity, and resulted in a combined assemblage with a higher diversity. Enrichment in zooplankton diversity caused by mixing the Kuroshio assemblage with the coastal assemblage was also observed in the coastal area south of Japan (Noda et al. 1998). A long-term study is required to check if the lower diversity in the CCC and higher diversity in the KBC are invariant biological properties of these two water masses. It will be interesting to observe the long-term dynamics of the copepod diversity in the frontal area where the CCC meets the KBC.

The presence of the two station clusters, namely the CCC and KBC, based on the copepod compositions is consistent with the results derived from hydrographic data. According to the records of Chen and Zhang (1965) and Chen et al. (1974), among the four corresponding species of the CCC, *Calanus sinicus* is abundant in the coastal and neritic waters of the Yellow Sea and East China Sea, *Corycaeus (Ditrichocorycaeus) affinis* and *Oithona similis* are abundant in the coastal waters of the Yellow Sea, and *Paracalanus parvus* is abundant in coastal waters off China. Of the 16 corresponding species of the KBC, *Acartia negligens*, *Calocalanus gracilis*, *Calocalanus plumulosus*, *Cosmocalanus darwinii*, *Lucicutia flavicornis*, and *Subeucalanus mucronatus* are oceanic species; *Acrocalanus gibber*, *Clausocalanus furcatus*, *Clausocalanus minor*, *Oithona plumifera*, *Oncaea venusta*, *Paracalanus aculeatus*, *T. stylif-*

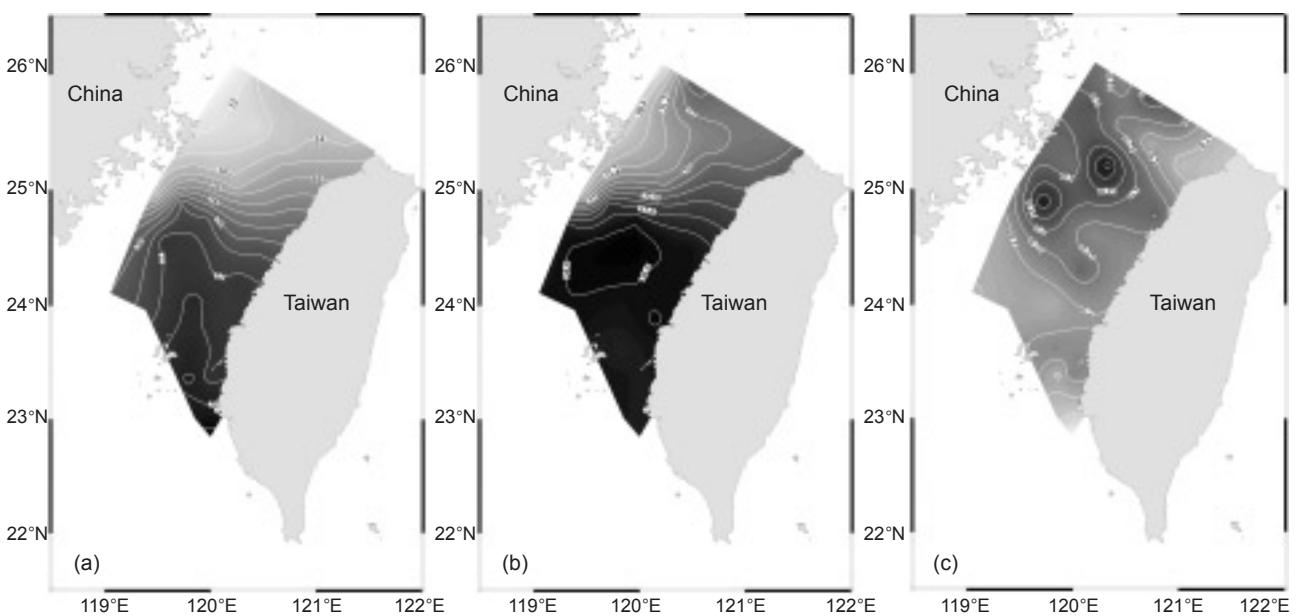


Fig. 3. Contour plots of surface water (a) temperature, (b) salinity, and (c) fluorescence.

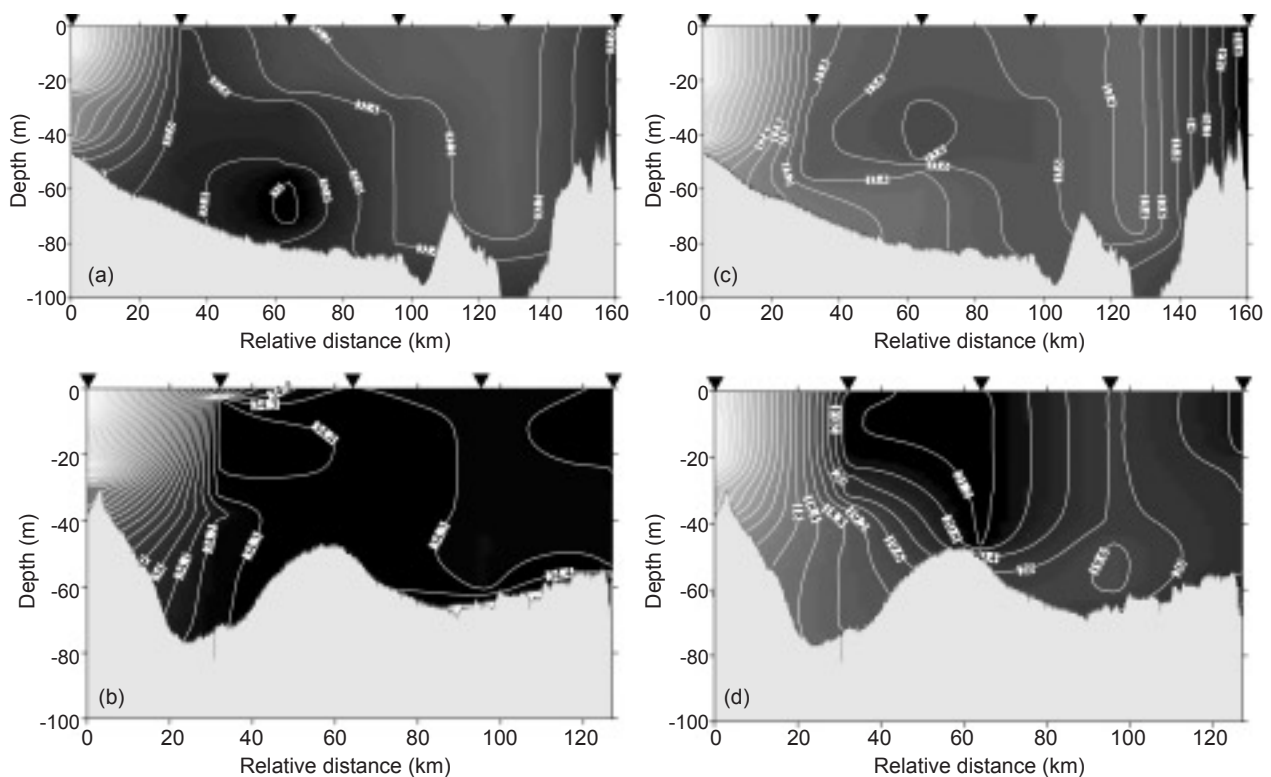


*era*, and *T. turbinata* are neritic species; and *Euterpina acutifrons* and *Parvocalanus crassirostris* are coastal species. They are all confined to warm waters (Chen and Zhang 1965, Chen et al. 1974, Chen 1992). From our study, we believe that *Calanus sinicus*, *Corycaeus affinis*, and *Oithona similis* followed the CCC into the Taiwan Strait from the north, while warm-water oceanic species accompanied the KBC into the Strait from the south and mixed with coastal and neritic species in the northern Strait. *Paracalanus parvus* was the most abundant species and had a broad distribution in this study.

In conclusion, the CCC contained only coastal/neritic cold-water species, but the KBC consisted of coastal, neritic, and oceanic warm-water species. The copepod community in the KBC is a mixture of the Kuroshio assemblage and coastal/neritic assemblage in the Taiwan Strait, which reflects the high diversity mentioned above. Using plankton assemblages to characterize the hydrography is important for understanding marine ecosystems. For example, Boucher et al. (1987) showed the spatial distribution of zooplankton in relation to the hydrography in the Ligurian Sea

front; Kouwenberg (1994) examined the copepod distribution in relation to the spatial structure in the northwestern Mediterranean; Gowen et al. (1998) used plankton distribution to characterize the frontal structure in the Irish sea; and Lopes et al. (1999) studied the distribution pattern of copepods related to the advection of South Atlantic Central Water southeast of Brazil.

Besides showing distinctive community structures in the CCC and KBC, we provide indicator species of each water mass. Among the 16 good indicators for the KBC, three are not included in the 20 most abundant species. These three, *Canthocalanus pauper*, *Oncaea mediterranea*, and *Scolecithrix danae*, are also warm-water species according to the records of Chen and Zhang (1965) and Chen et al. (1974). We have to emphasize that the copepod community in the KBC differs from that of the main Kuroshio Current by having a mixture of Kuroshio oceanic species and coastal/neritic warm-water species. By extracting the indicator species, we hope to provide useful information for monitoring Taiwan Strait ecosystems. Some species that are not abundant but appear frequently could be important targets

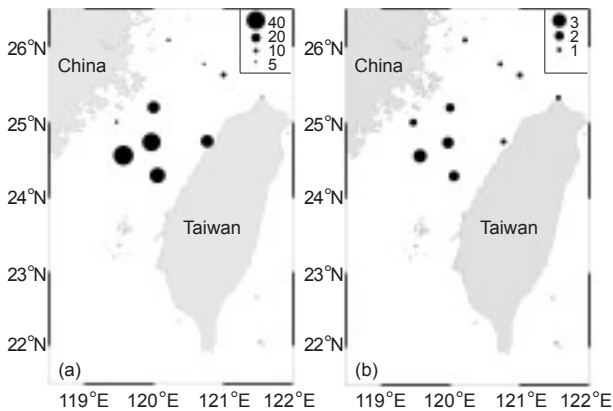


**Fig. 4.** Vertical sections of temperature of the (a) northern (Matsu-Cape Fuguei) and (b) central (Wuchiu-Taichung) transects and of salinity of the (c) northern (Matsu-Cape Fuguei) and (d) central (Wuchiu-Taichung) transects. Triangles indicate sampling stations.

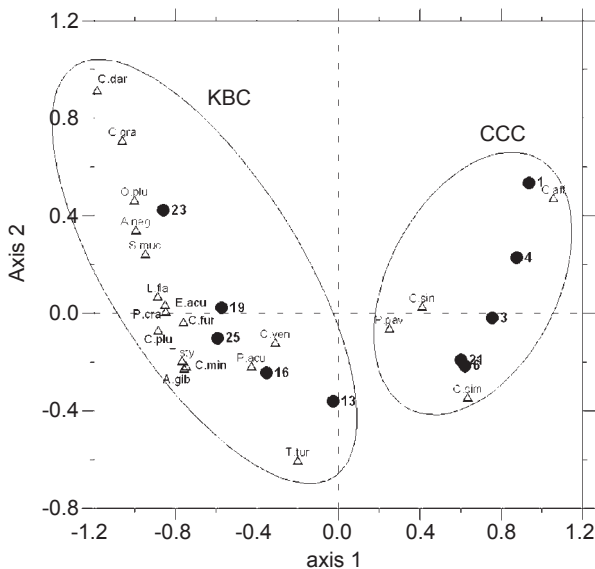
for long-term observation. The use of biological indicators to label water masses is as effective as physical or chemical methods in many cases, and it also provides useful information for observing ecosystem dynamics, especially when plankters are sensitive to environmental changes (Haury and Pieper 1988). Also, variations in zooplankton composition and abundance are widely used as proxies to observe long-term changes in marine ecosystems (Rebstock 2001 2002, Beaugrand et

al. 2002, Lindley and Reid 2002). Our future plan is to continually monitor this ecosystem in an attempt to understand seasonal and inter-annual variations.

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**Fig. 5.** Scatter plots of copepod species (a) richness (number of species) and (b) diversity (Shannon's index) at each sampling station.



**Fig. 6.** Ordination diagram derived from correspondence analysis based on the relative abundance of the 20 most abundant species of copepods, showing station associations and corresponding species. The ellipses indicate two station clusters determined by K-means classification. The first letter of the genus and the first three letters of the species name were used as abbreviations in the plot.

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