INTRODUCTION

Understanding the recruitment dynamics of exploitable fish is a central issue in fisheries science. The early life stage is the most important stage for determining annual recruitment of fishes supporting various major commercial fisheries, and many studies have focused on ecological aspects of the larval stage, such as distribution, growth and survival (Chambers & Trippel 1997, Fuiman & Werner 2002). However, marine fish exhibit high species diversity, and not all larval-stage fishes, including various commercially important species, can readily be morphologically identified to the species level (e.g. Okiyama 1988, Moser 1996). Analysis of mtDNA has been used to identify fish species at all life stages (e.g. Graves et al. 1990, Chow et al. 2003, Karaiskou et al. 2005, Neira & Keane 2008), and this method can help to examine the ecology of larval-stage fishes for which species identification cannot be readily distinguished based solely on morphological characteristics.

The chub mackerel *Scomber japonicus* and spotted mackerel *S. australasicus* occur widely in temperate and subtropical waters of the Indo-Pacific Ocean, and display antitropical distributions (Collette & Nauen 1983, Scales et al. 1998, Collette 1999). In the East China Sea (ECS), which is one of the largest marginal seas of the western Pacific Ocean, both species occur abundantly and are commercially fished, mainly with purse seines, by the countries adjacent to the ECS. These 2 species form a shared stock that migrates across the boundaries within the ECS.
across the boundaries of adjacent Exclusive Economic Zones (EEZs) of 2 or more coastal countries (FAO 2006). The catches of *S. japonicus* and *S. australasicus* by Japanese and Korean fisheries during 2000–2008 ranged between 190 000 and 307 000 and between 37 000 and 91 000 t, respectively for the two species, in the ECS and the adjacent sea areas (Fisheries Agency & Fisheries Research Agency of Japan 2010). Based on their gonadal development, these 2 species spawn in the ECS during February to June (Yukami et al. 2009).

Information on the processes associated with survival during the vulnerable larval stage of *Scomber japonicus* and *S. australasicus* in the ECS is important for understanding mechanisms of year-to-year variation in recruitment. Survival during the larval stage, i.e. during the first weeks of planktonic life, is considered critical for recruitment in the congeneric Atlantic mackerel *S. scombrus* in the northwestern Atlantic Ocean (Robert et al. 2007, Castonguay et al. 2008). In the northeastern Atlantic Ocean off the coast of Europe where considerable information has been accumulated on the egg and larval biology of *S. scombrus*, an individual-based model was developed for the prediction of year-to-year variations in transport, growth and survival of the early life stages (Bartsch & Coombs 2004, Bartsch et al. 2004). However, information on the recruitment processes of *S. japonicus* and *S. australasicus* is limited, and little is known on the distribution and growth of the larvae in the ECS. The Fisheries Research Agency of Japan began large scale larval sampling survey from the southern to northern ECS in 2001, and extremely high abundances of *Scomber* spp. larvae were found in the shelf break region of the southern ECS south of 28°N during February to March (our Fig. 1, Sassa et al. 2006). This suggests that their primary spawning ground is formed there. However, few characteristics are available to distinguish between *S. japonicus* and *S. australasicus* larvae based on their morphology and pigmentation patterns (Watanabe 1970). Although Ozawa (1984) suggested that *S. australasicus* possesses distinctive melanophores on the surface of the hindbrain throughout the postlarval stage, which are not seen in *S. japonicus*, it remains to be verified how valid this feature is as a diagnostic characteristic (C. Sassa unpubl. data).

Recently, a species identification method for *S. japonicus* and *S. australasicus* has been established based on PCR-restriction fragment length polymorphism (PCR-RFLP) analysis of mtDNA (Sezaki et al. 2001).

In this study, we examined the distribution and abundance of *Scomber japonicus* and *S. australasicus* larvae in the southern ECS during February to March in 2004 and 2005 based on PCR-RFLP analysis of mtDNA (Sezaki et al. 2001). The distribution patterns are discussed in relation to the physical oceanography to infer the larval transport as well as between-species and between-year differences in distribution. We also examined the larval growth and the between-year differences of the 2 mackerels based on otolith increments, since early growth is one of the most important factors determining recruitment success of fishes (Takasuka et al. 2003, Takahashi & Watanabe 2004, Robert et al.

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**Fig. 1. Scomber japonicus and S. australasicus.** Horizontal distributions of mackerel larvae in the East China Sea from February to March in 2004 and 2005. Circles represent the abundance as a continuous range of values. Crosses indicate no catch. N: total number of sampling stations; KBCNT: Kuroshio Branch Current north of Taiwan; TSWC: Tsushima Warm Current
Sassa & Tsukamoto: Distribution and growth of *Scomber* larvae (2007). The larval growth is discussed in response to habitat conditions, such as water temperature and food availability experienced by the larvae. Our study provides fundamental information on which to base future studies, including modeling, to allow prediction of the spawning and recruitment variability of the 2 mackerel species of these shared stocks in the ECS.

**MATERIALS AND METHODS**

**Study area.** The southern ECS between 25 and 28° N is an extremely dynamic oceanic region, as indicated by the prominent frontal area between the Kuroshio Current and shelf waters and the between-year difference in water mass distribution (Tang et al. 2000, Sassa et al. 2008b). Two current systems are active in the southern ECS during winter: the Kuroshio Current and the Kuroshio Branch Current north of Taiwan (KBCNT) (Fig. 2, lower right panel; Ichikawa & Beardsley 2002). The Kuroshio Current flows along the 200 m isobath at the shelf break. A branch of the Kuroshio Current intrudes onto the shelf (<200 m depth) northeast of Taiwan between ~122 and ~125° E, i.e. the KBCNT (Ichikawa & Beardsley 2002). The KBCNT flows first northward and then northeastward at a relatively slow speed. The intrusion of the Kuroshio waters is related to the northeast monsoon; thus, the KBCNT is expected to be prevalent in winter (Gong et al. 1997, Ichikawa & Beardsley 2002). These 2 currents can significantly affect transport processes of larval carangid fishes (Kasai et al. 2008, Sassa et al. 2008b).

**Sample collection.** Larvae were collected along 5 transects in the shelf break region of the southern ECS between 25° 45’ N and 28° 15’ N in 2004 and 2005 (Fig. 2). From 27 February to 9 March 2004 and between 1 and 11 March 2005, samples were collected at 36 and 37 stations, respectively, from the RV ‘Yokomaru’ (Seikai National Fisheries Research Institute, Fisheries Research Agency; 499 t) (Fig. 2). A paired bongo net (60 cm mouth diameter, 0.33 mm mesh) was adopted for larval sampling, with a flowmeter and depth recorder attached to the net for quantitative sampling. A double-oblique tow was conducted at each station from the surface down to 150 m depth or close to the bottom at shallow stations. The towing speed was approximately 1 m s\(^{-1}\) (2 knots). A CTD (Alec Electronics) with a submersible fluorometer (Clorotec, ACL220-PDK, Alec Electronics) was used at each sampling station to 200 m depth or close to the bottom at shallower stations. Plankton samples were preserved in 99.5% ethanol onboard immediately after capture.

In the laboratory, all *Scomber* spp. larvae were sorted from the samples and counted. Body length (BL) was measured to the nearest 0.1 mm with an ocular micrometer of a stereomicroscope. Notochord length (NL) was measured for preflexion larvae and standard length (SL) for flexion and postflexion larvae (Moser 1996). No correction was made for larval shrinkage due to net capture or preservation by the ethanol. Each larva was numbered and individually preserved in a separate vial for further analysis. In total, we collected 368 and 274 *Scomber* spp. larvae in 2004 and 2005, respectively, of which, 306 and 254 larvae, respectively, in good condition were used for DNA and otolith extraction.

**Species identification.** Total DNA from the muscle tissues of *Scomber* spp. larvae was extracted using the DNeasy Tissue Kit (QIAGEN). DNA fragments encoding the mitochondrial cytochrome \(b\) gene were
amplified by the PCR using the primer pair SACB-7L (5'-AGT CCC ATA CGT CGG TAC TA-3') and SACB-8H (5'-CAT TCA GGC TTA ATA TGA GG-3') (Sezaki et al. 2001). The reaction mixtures were preheated at 98°C for 30 s followed by 30 cycles of amplification (at 98°C for 10 s in denaturation, 55°C for 30 s in annealing, and 72°C for 60 s in extension) with a final polymerization step at 72°C for 120 s. Amplified products were purified with the GFX™ PCR DNA and Gel Band Purification Kit (Amersham Biosciences).

The full length of the cytochrome b gene consisted of 1140 nucleotides irrespective of *Scomber japonicus* and *S. australasicus*, in which 16 species-specific variations were observed (Sezaki et al. 2001). Among the various restriction enzymes that recognize the cytochrome b genes from the 2 fish species, MvaI is specific to the gene of *S. japonicus* and MboI to that of *S. australasicus*, with both enzymes producing 2 fragments (Sezaki et al. 2001). In this study, the 10 µl of amplified products were directly digested by 2 restriction enzymes (MvaI and MboI) for 1 h. The digested samples were electrophoresed through 1.5% agarose gel (Agarose X, Nippon Gene) for ~25 min. The DNA bands were visualized and photographed after electrophoresis and staining with ethidium bromide.

In the present study, 98.9% of the larvae showed restriction fragment length polymorphism (RFLP) profiles consistent with those of Sezaki et al. (2001) (see ‘Results—Species identification’). For the other 6 larvae having inconsistent RFLP profiles, the nucleotide sequences of the cytochrome b were determined. Then, the species identifications were carried out by comparing the sequences between the larvae and morphologically well-identified adults (DDBJ/EMBL/GenBank under accession numbers AB032515–AB032520).

**Hydrographic analysis.** To clarify the position of the Kuroshio axis along the shelf break, currents were measured with an acoustic Doppler current profiler (ADCP) and were routinely monitored at 10, 25 and 50 m depths throughout the cruise. However, it is not possible to accurately assess the position and flow strength of the KBCNT, since the flow direction and velocity on the shelf region of the ECS fluctuates with the diurnal and semidiurnal tidal flows (Katoh et al. 2000). Thus, we defined the KBCNT based on water mass distributions and sea surface temperature (SST) fields.

A cluster analysis using water temperature and salinity at 20 m depth was conducted to distinguish the water properties at each sampling station. Since (1) the water column is mixed vertically in winter due to the strong northeast monsoon and the mixed layer depth is usually deeper than 30 m (Ichikawa & Beardsley 2002, Sassa et al. 2006), and (2) *Scomber* spp. larvae are concentrated in the upper 30 m layer with peak densities at ~20 m depth in the southern ECS (C. Sassa unpubl. data), the temperature and salinity at 20 m directly represent the water properties in the epipelagic layer where the larvae occur. The standardised Euclidean distance was used to evaluate the dissimilarity between water properties of each sampling station. Clustering by the farthest neighbour strategy was used to construct dissimilarity matrices for data of the total 73 sampling stations (Wilks 1995, Sassa et al. 2008b). Cluster analysis was performed with the PRIMER v.6 software package (Clarke & Gorley 2006). We used similarity profile permutation tests (SIMPROF, PRIMER v.6) to identify significant clusters of locations at the 95% significance level.

**Analysis of larval distribution.** The description of the geographic distribution of larvae was based on their abundance (number of larvae per 10 m² sea surface). If the catch was too large to identify all larvae using mtDNA (only 4 stations), the larvae randomly picked out from the sample at the station were analysed and the total abundance of *Scomber* spp. was multiplied by the ratio of *S. japonicus* to *S. australasicus*. To examine the larval transport processes, horizontal distributions were analyzed using the following 2 body size classes: <5 mm and ≥5 mm BL (hereafter referred to as ‘small larvae’ and ‘large larvae’, respectively). Notochord flexion begins to occur between 5 and 6 mm BL (Ozawa 1984); thus, the small larvae were at the preflexion stage, while the large larvae included flexion and postflexion stages.

Spatial overlap between *Scomber japonicus* and *S. australasicus* larvae was measured in each size class during 2004 and 2005 using Schoener’s index (Schoener 1970) which, in this case, compared the relative proportions of the square root-transformed abundance of each species. This index ranges from 0% (no overlap) to 100% (complete overlap).

Since species-specific temperature optima during the early life stages are key to understanding multispecies population dynamics (Takasuka et al. 2008), quotient analysis (Ibaibarriaga et al. 2007, Neira & Keane 2008) was performed on larval abundance data across 2 surveys combined to describe characteristics of the habitat temperature of each species. We used temperatures at 20 m depth of each station. For this analysis, square root-transformed abundance of the larvae within 1°C temperature classes was expressed as a percentage of total abundance, divided by the percentage frequency of stations under each temperature. Quotients > 1 indicate positive habitat selection, i.e. range of optimum temperature.

**Chlorophyll a concentration and copepod nauplii density.** The chlorophyll a (chl a) fluorescence mea-
sured by the Clorotec was calibrated based on standard procedures using extracted chl a from 250 ml water samples at 20 m depth (Sassa & Konishi 2006). The chl a concentration (mg m⁻²) was integrated for the 0 to 50 m water column.

Copepod nauplii are an important prey item for Scomber spp. larvae in the study area (Sassa et al. 2008a). Data on nauplii density were taken from Okazaki et al. (2008). The nauplii were collected from 1 l of surface water and concentrated using a plankton net with 0.05 mm mesh and fixed in 5% buffered formalin seawater. Since there was no significant difference in the nauplii density between the sea surface and 20 m depth (C. Sassa unpubl. data), we used data at the sea surface as a proxy for the food available to the larvae.

The weighted mean values of temperature (WMT), chl a concentration (WMC) and copepod nauplii density (WMN) of Scomber japonicus and S. australasicus habitat for each of the 2 body size class were calculated using the following equations for each year:

\[
\begin{align*}
WMT &= \frac{\sum_{i=1}^{n} (s_i \times t_i)}{\sum_{i=1}^{n} s_i} \\
WMC &= \frac{\sum_{i=1}^{n} (s_i \times c_i)}{\sum_{i=1}^{n} s_i} \\
WMN &= \frac{\sum_{i=1}^{n} (s_i \times n_i)}{\sum_{i=1}^{n} s_i}
\end{align*}
\]

where \(s_i\) is the abundance of S. japonicus or S. australasicus larvae in the 1th sampling station (no. larvae 10 m⁻²), and \(t_i\), \(c_i\) and \(n_i\) are the water temperature at 20 m depth, the chl a concentration and the copepod nauplii density at the 1th sampling station, respectively. Before the analysis, the larval abundance was square root-transformed to reduce the bias caused by sampling stations with extremely large catches.

**Otolith analysis and growth rates.** Sagittal otoliths were extracted from all Scomber spp. larvae identified to the species level, and increments were counted under a microscope to determine growth. The first increment was observed at approximately 5 µm from the otolith core in both larvae. Since the otolith radius of newly hatched larvae of S. australasicus was approximately 5 µm in a rearing experiment (M. Saito, Japan NUS Co. Ltd. pers. comm.), we defined the first increment as the hatch check, and the total increment number on the outside of the hatch check as age. We postulated that the increments were deposited daily, as determined in studies for S. japonicus and S. scombrus (Mendiola & Álvarez 2008, M. Takahashi unpubl. data).

Instantaneous growth rate (\(G\)) and relative growth rate (\(K\)) were estimated as follows (Yamashita & Bailey 1989, Mendiola et al. 2009):

\[
L_t = L_0 e^{G t}
\]

and

\[
K = e^G - 1
\]

where \(L_0\) is the initial body length (mm) and \(L_t\) is the body length at time \(t\) (d). The daily specific growth rate was defined as \(K \times 100\%\) (Mendiola et al. 2009). To test the suitability of the exponential model, the body length-at-age data were also fitted by linear regression.

The differences in the instantaneous growth rates (G) by years and by species were evaluated using analysis of covariance (ANCOVA), which was performed for linearised exponential models.

**RESULTS**

**Water mass distributions and current features**

Based on a cluster analysis, 5 water masses were recognized (SIMPROF, \(p < 0.05\)) (Fig. 2). Based on their temperature and salinity (T-S) properties and distributions, they were respectively named the Kuroshio waters, Kuroshio branch current waters-I and -II, coastal waters, and mixed waters (hereafter KUR, KB-I, KB-II, COA, and MIX, respectively) (Figs. 2 & 3a). Temperature gradually decreased from the highest values in the KUR (22.9 to 24.4°C) to the lowest in the COA (13.8 to 15.8°C) (Fig. 3a). Salinity in the KUR, KB-I, and KB-II was high (34.5 to 34.8) with no significant difference (Kruskal-Wallis, \(p > 0.05\)), while salinity in both the MIX and COA was lower and broader in range (34.1 to 34.8 and 33.4 to 34.6, respectively) compared with the above 3 subareas (Mann-Whitney U-test, \(p < 0.05\)) (Fig. 3a).

Between ~26 and ~29°N, the Kuroshio front was observed along the 200 m isobath at the shelf break based on the ADCP observations during our cruises. This corresponds with the boundary between the 2 water masses of the KUR and KB-I and the SST fields (~21 to 23°C isotherms) in both years (Fig. 2).

Water mass distribution was significantly different between 2004 and 2005 (Fig. 2). An intrusion of the Kuroshio branch current north of Taiwan (KBCNT) was much more evident in 2004 than in 2005, since the distribution of the KB-I and KB-II in 2004 extended more northward to ~28°N (Fig. 2). In 2005, on the other hand, the KBCNT was weak based on the water mass distribution and the SST fields. Instead, it was notable that the cold COA extended southward to ~27°N (Fig. 2). SST was significantly higher in 2004 than in 2005 (20.8 ± 2.0 versus 18.8 ± 3.0°C, mean ± SD) (Mann-Whitney U-test, \(p < 0.05\)).
Chl \(a\) concentration and copepod nauplii density

In 2004, high concentrations of chl \(a\) (>40 mg \(m^{-2}\)) were observed over a wide area in the northwest edge and central part of the study area (Fig. 4), corresponding with the KBCNT area (Fig. 2). In 2005, on the other hand, the high chl \(a\) concentrations were restricted in a narrower area of the northeast edge of the study area (Fig. 4). In both years, the chl \(a\) concentrations were low in the KUR, with a mean value of 22 mg \(m^{-2}\) (Fig. 4). Mean (±SD) chl \(a\) concentrations in 2004 and 2005 were 37.8 ± 14.4 and 30.3 ± 9.8 mg \(m^{-2}\), respectively, and chl \(a\) was significantly higher in 2004 (Mann-Whitney \(U\)-test, \(p < 0.05\)).

The nauplii density showed positive correlations with chl \(a\) concentration (Pearson’s correlation coefficient: \(r = 0.433\), \(n = 73\), \(p < 0.05\)). In 2004, high densities of nauplii (>15 ind. \(l^{-1}\)) were observed mainly in the KBCNT area (Figs. 2 & 4). In 2005, the nauplii densities were lower than those in 2004, although the maximum density of 41 ind. \(l^{-1}\) was observed at the northernmost station (Fig. 4). In both years, the nauplii densities were low in the KUR, with mean density of 5 ind. \(l^{-1}\) (Fig. 4).

Species identification

A total of 544 \textit{Scomber} spp. larvae ranging from 2.6 to 11.7 mm BL were identified to species level based on the PCR-RFLP analysis. Only 6 larvae showed inconsistent RFLP profiles with those described by Sezaki et al. (2001), and all of them were assigned to \textit{S. australasicus} based on the nucleotide sequence analysis. Ten

![Fig. 3](image-url) (a) Temperature-salinity (T-S) diagram for each of the 73 sampling stations at 20 m depth. Each water mass resulting from the cluster analysis is shown by symbols. (b,c) Numerical abundance of \textit{Scomber japonicus} and \textit{S. australasicus} larvae in relation to the temperature and salinity of their habitats. Circles represent the abundance as a continuous range of values > 0. Crosses indicate no catch. KUR: Kuroshio waters; KB-I: Kuroshio branch current waters-I; KB-II: Kuroshio branch current waters-II; COA: coastal waters; MIX: mixed waters

![Fig. 4](image-url) Horizontal distributions of the chl \(a\) concentration (mg \(m^{-2}\)) in the upper 50 m of the water column and copepod nauplii density (ind. \(l^{-1}\)) at the sea surface in the southern East China Sea in 2004 and 2005. Contours indicate the chl \(a\) concentration. Circles represent the nauplii density as a continuous range of values > 0. Cross indicates no catch. The 100 and 200 m isobaths are shown with dashed lines.
larvae were unidentifiable since the DNA extracted was not sufficient to analyse, possibly due to problems related to sample preservation. In the present study, *S. japonicus* and *S. australasicus* accounted for 57.6 and 42.4% of the total larvae identified, respectively.

**Larval abundance and body length**

Mean abundances of *Scomber japonicus* larvae in 2004 and 2005 were comparable, with values of 27.4 and 32.5 larvae 10 m$^{-2}$, respectively (Mann-Whitney U-test, $p > 0.05$) (Table 1). Mean abundance of *S. australasicus* larvae in 2004 was 36.3 larvae 10 m$^{-2}$, which was approximately twice that of 2005, although no significant difference was detected (Mann-Whitney U-test, $p > 0.05$). In both years, there were no significant differences in abundances between species (Mann-Whitney U-test, $p > 0.05$) (Table 1).

Larval *Scomber japonicus* and *S. australasicus* larvae were significantly larger in 2004 (5.7 and 5.4 mm BL, respectively) than in 2005 (4.9 and 5.0 mm BL, respectively) (Mann-Whitney U-test, $p < 0.05$) (Fig. 5).

**Larval distribution and transport**

Small larvae of *Scomber japonicus* (<5 mm BL) were collected abundantly to the west of 124°E in 2004, with mean abundance of 70.1 larvae 10 m$^{-2}$ at positive stations (Fig. 6a). Distribution of the large larvae (≥5 mm BL) tended to extend northeastward

<table>
<thead>
<tr>
<th>Sampling period</th>
<th>Species</th>
<th>n</th>
<th>Positive tows (%)</th>
<th>Abundance (ind. 10 m$^{-2}$)</th>
</tr>
</thead>
<tbody>
<tr>
<td>27 Feb–9 Mar 2004</td>
<td><em>S. japonicus</em></td>
<td>147</td>
<td>50.0</td>
<td>27.4 ± 12.0</td>
</tr>
<tr>
<td></td>
<td><em>S. australasicus</em></td>
<td>149</td>
<td>30.6</td>
<td>36.3 ± 14.2</td>
</tr>
<tr>
<td>1–11 Mar 2005</td>
<td><em>S. japonicus</em></td>
<td>170</td>
<td>48.6</td>
<td>32.5 ± 12.0</td>
</tr>
<tr>
<td></td>
<td><em>S. australasicus</em></td>
<td>84</td>
<td>29.7</td>
<td>14.7 ± 8.1</td>
</tr>
</tbody>
</table>

Table 1. *Scomber japonicus* and *S. australasicus*. Percentage of positive tows and mean ± SE abundance (ind. 10 m$^{-2}$) of the larvae in the shelf break region of the southern East China Sea in 2004 and 2005. n: total number of fish collected

Fig. 5. *Scomber japonicus* and *S. australasicus*. Length frequency distributions of larvae collected by a bongo net in the southern East China Sea in 2004 and 2005. n: total number of fish collected
to 125° 30' E (Fig. 6b), corresponding with the intrusion of the KBCNT (Fig. 2). In 2005, the small larvae of *S. japonicus* occurred abundantly in the westernmost transect and the Kuroshio Current frontal area at ~124° E (Fig. 6c). In addition, they were collected in the northeastern area between 124° 30' E and 125° 30' E, but with a relatively low abundance (Fig. 6c). In 2005, distribution of the large larvae shifted slightly northeastward in the area west of 123° 30' E (Fig. 6d), although this pattern was not so clear compared with that in the previous year. The large larvae also occurred along the Kuroshio front in the area east of 124° E, with a relatively low abundance (Fig. 6d).

The small larvae of *Scomber australasicus* were concentrated to the west of 124° E between 26 and 27° N in 2004, with mean abundance of 76.5 larvae 10 m$^{-2}$ at positive stations (Fig. 7a). As larvae grew, their center of distribution shifted northward and northeastward (Fig. 7b), corresponding closely with the direction of the KBCNT (Fig. 2). In 2005, the abundance and distribution pattern of the small larvae of *S. australasicus* was similar to that in the previous year (Fig. 7c), while that of the large larvae was considerably different from that in the previous year, i.e. they showed a dispersed pattern along the Kuroshio front (Fig. 7d). The change in larval transport conditions corresponded with the between-year difference in the pattern of the currents (Fig. 2).

**Difference in distribution and habitat temperature between species**

In both years, the percentage of positive tows of *Scomber japonicus* larvae showed higher values than that of *S. australasicus* (48.6 to 50.0 versus 29.7 to 30.6%) (Table 1), and the distributional area was broader in *S. japonicus* larvae (Figs. 6 & 7). Although there were sampling stations where both *S. japonicus* and *S. australasicus* larvae were abundantly collected, the center of distribution tended to differ between spe-
cies in both size classes (Figs. 6 & 7). That is, the distribution center of *S. australasicus* was in a slightly more southern area than that of *S. japonicus*. In 2004, spatial overlap of these 2 species, based on Schoener's index, was relatively high at 54.9% in small larvae, but it declined to 31.4% in large larvae. The spatial overlap in 2005 was low in both small and large larvae (29.9 and 31.2%, respectively).

High abundances of *Scomber japonicus* larvae were observed broadly in the 5 water masses (Fig. 3b), while those of *S. australasicus* larvae were restricted mainly in the KUR and KB-I (Fig. 3c). The abundance quotients of *S. japonicus* larvae across 2 surveys showed high values (>1) between 15 and 22°C, except between 18 and 19°C (Fig. 8). On the other hand, the abundance quotients of *S. australasicus* larvae peaked sharply
between 20 and 23°C (Fig. 8), showing a higher and narrower range of habitat temperature than that of *S. japonicus*, although there was overlap between 20 and 22°C.

### Larval growth and its between-year variations

The exponential model fitted the body length (BL) and age data of both species better than the linear model (Fig. 9, Table 2), except for data of *Scomber japonicus* in 2005 when the linear model fitted better than the exponential one ($r^2 = 0.858$ versus 0.837). In both species, there were significant differences in the instantaneous growth rate ($G$) between the 2 yr. That is, $G$ of both species in 2004 was significantly higher than values in 2005 (ANCOVA: df = 1, $F = 30.88$, $p < 0.05$ for *S. japonicus*; df = 1, $F = 8.95$, $p < 0.05$ for *S. australasicus*) (Fig. 9, Table 2).

The daily specific growth rate ($K$) of *Scomber japonicus* larvae was higher in 2004 (8.2% BL d$^{-1}$) than in 2005 (6.2% BL d$^{-1}$) (Table 2). Similarly, $K$ of *S. australasicus* larvae was higher in 2004 (9.3% BL d$^{-1}$) than in 2005 (7.7% BL d$^{-1}$), although the between-year difference was smaller compared with that of *S. japonicus* (Table 2).

<table>
<thead>
<tr>
<th>Species</th>
<th>Year</th>
<th>$L_0$</th>
<th>$G$</th>
<th>$K$ (%)</th>
<th>$n$</th>
<th>$r^2$</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>S. japonicus</em></td>
<td>2004</td>
<td>2.516</td>
<td>0.079</td>
<td>8.2</td>
<td>147</td>
<td>0.838</td>
</tr>
<tr>
<td></td>
<td>2005</td>
<td>2.947</td>
<td>0.060</td>
<td>6.2</td>
<td>170</td>
<td>0.837</td>
</tr>
<tr>
<td><em>S. australasicus</em></td>
<td>2004</td>
<td>2.300</td>
<td>0.089</td>
<td>9.3</td>
<td>146</td>
<td>0.799</td>
</tr>
<tr>
<td></td>
<td>2005</td>
<td>2.622</td>
<td>0.074</td>
<td>7.7</td>
<td>84</td>
<td>0.859</td>
</tr>
</tbody>
</table>

Predicted absolute growth rates of *Scomber japonicus* between 5 and 15 d were 0.28 to 0.63 and 0.23 to 0.42 mm d$^{-1}$ in 2004 and 2005, respectively, and those of *S. australasicus* were 0.31 to 0.74 and 0.27 to 0.57 mm d$^{-1}$ in 2004 and 2005, respectively (Table 3). The predicted absolute growth rates of both species of larvae increased with age (Table 3).

### Between-year variations in habitat conditions

The WMT of *Scomber japonicus* habitat was ~2 to 3°C higher in 2004 than in 2005 in both size classes,
while there was no remarkable between-year difference in the WMT of *S. australasicus* habitat, i.e. only 0.1 to 0.5°C higher in 2004 (Table 4).

In 2004, the occurrence of *Scomber japonicus* and *S. australasicus* larvae (Figs. 6 & 7) closely corresponded with the area of high chl a concentration (>40 mg m⁻³) (Fig. 4). In 2005, on the other hand, they occurred in the area where the chl a concentration was low (15 to 30 mg m⁻³). The weighted mean chl a concentration (WMC) of both *S. japonicus* and *S. australasicus* habitat was ~1.5 times higher in 2004 than in 2005 (Table 4).

The WMN of *Scomber japonicus* habitat was much higher in 2004 than in 2005 in both size classes (15.5 to 16.5 versus 9.4 to 9.7 ind. ¹⁻¹) (Table 4). The WMN of *S. australasicus* habitat was slightly higher in 2004 than in 2005 (9.3 to 10.5 versus 8.1 ind. ¹⁻¹) (Table 4).

### DISCUSSION

#### Limitations of the sampling and analysis

In this study, dense distributions of *Scomber japonicus* and *S. australasicus* larvae were observed in the southern ECS in both 2004 and 2005; however, there are 3 potential limitations in our sampling and analysis. Firstly, the abundance of larvae of both species in the southern ECS should be considered as an underestimate in this study since a large number of larvae of both species occurred at the westernmost stations. In the west of our study area, there is usually a northeastward flow with a maximum speed sometimes reaching ~0.6 to 0.8 knots, although inter-seasonal differences are known; i.e. the main route changes off northeastern Taiwan and a low velocity of <0.1 to 0.2 knots is observed outside of the flow (Tang et al. 2000). This indicates that their distribution extends to the west of our study area, i.e. in the coastal and offshore waters around Taiwan. Although the occurrence of *Scomber* spp. larvae in the inshore and offshore waters of northeastern Taiwan has been reported, differences in distribution of each species were not differentiated based on biochemical techniques (Chiu 1999).

Secondly, the present study might not clarify the northern border of the distribution of *Scomber japonicus*. The lower limit of the habitat temperature of *S. japonicus* larvae was ~15°C. Based on monthly mean SST fields in the ECS during the study periods (Japan Meteorological Agency, available at www.jma.go.jp/jma/; accessed 20 Mar 2010), the 15°C isotherm extended to ~29° N in 2004; thus, the distribution of *S. japonicus* larvae would have been underestimated. In 2005, on the other hand, our survey covered the 15°C isotherm in the study area.

Finally, in the present study, species identification was based on the mtDNA, that is maternal inheritance; thus, our method could not detect hybrid individuals if they occur. Although the Schoener’s spatial overlap index between *Scomber japonicus* and *S. australasicus* showed relatively low values, both species were collected abundantly at the few sampling stations where hybrid individuals might potentially be produced. Although occurrence of morphologically intermediate adult individuals between the 2 mackerels has been reported, Kijima et al. (1986), using isozyme markers, could not find genetic evidence of hybridization among them. Species identification using nuclear DNA analysis is needed to clarify the occurrence of hybrid larvae in the field.

#### Species identification

The reliability of PCR-RFLP analysis depends on the magnitude of intraspecific variation, which requires a large number of wild samples (Chow & Inoue 1993, Chow et al. 2003). In the present study, 98.9% of larvae were identified to species based on the PCR-RFLP analysis. Only 6 larvae had inconsistent RFLP profiles and all were identified as *Scomber australasicus* by the nucleotide sequence analysis. This indicated that there were intraspecific variations in the cytochrome b gene sequences in *S. australasicus* from those reported by Sezaki et al. (2001), but the frequency of occurrence was low (<3%).

### Table 3. *Scomber japonicus* and *S. australasicus*. Predicted absolute growth rates (mm d⁻¹) of the larvae at various ages from hatching in 2004 and 2005

<table>
<thead>
<tr>
<th>Age (d)</th>
<th><em>S. japonicus</em></th>
<th><em>S. australasicus</em></th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>2004</td>
<td>2005</td>
</tr>
<tr>
<td></td>
<td>2004</td>
<td>2005</td>
</tr>
<tr>
<td>5</td>
<td>0.28</td>
<td>0.23</td>
</tr>
<tr>
<td>10</td>
<td>0.42</td>
<td>0.31</td>
</tr>
<tr>
<td>15</td>
<td>0.63</td>
<td>0.42</td>
</tr>
</tbody>
</table>

#### Table 4. *Scomber japonicus* and *S. australasicus*. Weighted mean temperature (WMT, °C), chl a concentration (WMC, mg m⁻³) and copepod nauplii density (WMN, ind. ¹⁻¹) associated with larvae of each size class during 2004 and 2005

<table>
<thead>
<tr>
<th>Species</th>
<th>Size class (mm)</th>
<th>WMT</th>
<th>WMC</th>
<th>WMN</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>2004</td>
<td>2005</td>
<td>2004</td>
<td>2005</td>
</tr>
<tr>
<td><em>S. japonicus</em></td>
<td>Small larvae (&lt;5)</td>
<td>20.4</td>
<td>18.7</td>
<td>50.0</td>
</tr>
<tr>
<td></td>
<td>Large larvae (≥5)</td>
<td>20.3</td>
<td>17.7</td>
<td>46.2</td>
</tr>
<tr>
<td><em>S. australasicus</em></td>
<td>Small larvae (&lt;5)</td>
<td>21.5</td>
<td>21.0</td>
<td>43.7</td>
</tr>
<tr>
<td></td>
<td>Large larvae (≥5)</td>
<td>20.9</td>
<td>20.8</td>
<td>57.1</td>
</tr>
</tbody>
</table>
**Spawning ground**

High abundances of the small larvae of *Scomber japonicus* and *S. australasicus* (<5 mm BL) were observed in the shelf break region of the southern ECS south of 27°N during February to March. Since larvae of this size are less than 10 d old after fertilization (Fig. 9), the distributional area of the small larvae would represent the approximate location of the spawning ground. Based on the analysis of seasonal gonad development of adults, both *S. japonicus* and *S. australasicus* spawn primarily during February to March off the coast of northeastern Taiwan (Ku & Tzeng 1985a,b), which corresponds with the larval occurrence of the both species in the present study. Based on catch statistics and biometric data, the potential spawning ground of both species is estimated to be in the broad area of the shelf break region of the ECS between 26 and 31°N during February to March (Yukami et al. 2009). From our study, the estimated spawning ground was determined to be much more specific.

The distribution center of larvae tended to differ between species, and *Scomber australasicus* occurred in a slightly more southern area than did *S. japonicus*, although there were also some spatial overlaps. These larval distribution patterns suggest that spawning areas for the 2 species are separate within the main location of the spawning ground.

**Larval distribution in response to oceanographic conditions**

The difference in distribution patterns between species corresponded closely with the optimal habitat temperature of each species. Although there was overlap between 20 and 22°C, the habitat temperature of *Scomber japonicus* larvae showed a lower and broader range than that of *S. australasicus* (15 to 22°C versus 20 to 23°C). Mature *S. australasicus* tends to occur in higher water temperature areas than does *S. japonicus* in the ECS (17 to 25°C versus 15 to 22°C SST) (Yukami et al. 2009), which corresponds with the difference of larval habitat temperature. Water temperature where *S. japonicus* larvae occur are at 15 to 22°C in the Pacific coast off central Japan (Watanabe 1970), 14 to 21.9°C in the eastern North Pacific (Kramer 1960) and 16 to 22°C in the Gulf of California (Esqueda-Escarcega 1995), which corresponds closely with our observations. No information is available on the habitat temperature of *S. australasicus* larvae in the North Pacific and its adjacent waters. In southeastern Australia, they occurred at 17.5 to 20.5°C (Neira & Keane 2008), which is significantly lower than the habitat temperature in our study area, i.e. the optimal temperature for this species appears to be different among populations. Since between-year change of water mass distribution is observed in the southern ECS during winter (Tang et al. 2000, Sassa et al. 2008b), interannual variation would also occur in the area of optimal habitat temperature for *S. japonicus* and *S. australasicus* larvae. This might cause year-to-year variations in the extent of spawning and nursery grounds of each species.

During the early life history, *Scomber* spp. are considered to be passively transported by currents until entering the juvenile stage of 20 mm SL, ~3 to 4 wk after fertilization in laboratory rearing conditions, when they begin to swim actively and form schools (Watanabe 1970, Hunter & Kimbrell 1980, Masuda et al. 2002). There are 2 routes of larval transport from the spawning ground in the southern ECS (our Fig. 2, lower right; Kasai et al. 2008, Sassa et al. 2008b). One route is entrained by the KBCNT; i.e. first, the larvae are transported northward and then northeastward at a slow speed of ~0.2 to 0.5 knots. Another is a rapid northeastward transport by the Kuroshio Current, which sometimes reaches 1.5 to 3 knots. Based on a particle-tracking model, in 17 to 30 d a high percentage of larvae in the southern ECS transported by the former process would recruit into the shelf region of the ECS, and those transported by the Kuroshio Current would reach the Pacific Ocean off the coast of southern Japan (Kasai et al. 2008). In 2004, when an intrusion of the warm KBCNT was evident, *S. australasicus* larvae were transported northeastward as they grew, while they dispersed eastward along the Kuroshio front in 2005 when the intrusion of the KBCNT was weak. Although *S. japonicus* larvae showed a similar pattern, it was much more gradual than that of *S. australasicus*, corresponding with the weaker flow in the northern part of the study area (Tang et al. 2000, Kasai et al. 2008) where *S. japonicus* mainly occurred.

Since the current features in the southern ECS are highly variable and complex on both spatial and temporal scales due to the relative strength of the Kuroshio intrusions, tidal current and mesoscale frontal disturbances (Gong et al. 1997, Tang et al. 2000, Lie & Cho 2002, Sassa et al. 2008b), the larval transport process of the 2 mackerel species would fluctuate over relatively short periods in the southern ECS. Although our study was based on a ‘snapshot’ observation, the results suggest that a small difference in the larval distribution and the oceanographic conditions would lead to a remarkable difference in the transport route and the end point of transportation. Based on a particle-tracking model, Kasai et al. (2008) reported year-to-year variations in transport processes of larval carangids originating from the spawning ground in the southern...
ECS, supporting the above view. Furthermore, as discussed below, the difference in physical oceanographic conditions would also be related to the difference in larval habitat conditions, resulting in between-year variations in the larval growth.

**Larval growth in response to habitat conditions**

This study is the first to report the larval growth of *Scomber japonicus* and *S. australasicus* in the ECS. The exponential model gave the best fits to the laboratory rearing experiments (Hunter & Kimbrell 1980, Mendiola et al. 2009). The daily specific growth rate of *S. japonicus* in our study ranged from 6.2% BL d⁻¹ at the habitat temperature of 17.7 to 18.7°C to 8.2% BL d⁻¹ at 20.3 to 20.4°C (Tables 2 & 4). These values are within the range of the growth rates reported in the rearing conditions, i.e. from 2.9% BL d⁻¹ at 16°C to 8.8% BL d⁻¹ at 22°C (Mendiola et al. 2009). There is no information on the growth of *S. australasicus* larvae under laboratory rearing conditions to allow comparison with our results.

The growth rates of larvae of both species were significantly higher in 2004 than in 2005, and the between-year difference was larger in *S. australasicus* (Table 2). Interannual differences are also found in the growth of the congener Atlantic mackerel larvae in the northwest Atlantic Ocean (Robert et al. 2007). Our results would relate to the between-year difference in habitat temperature and food availability for larvae, since larval growth is a function of these 2 factors (Yamashita et al. 2001, Fuiman & Werner 2002, Takasuka & Aoki 2006). The WMT of *S. japonicus* habitat was 2 to 3°C higher in 2004 than in 2005, corresponding with the higher growth rate in 2004. For *S. australasicus*, on the other hand, there was only a slight difference (0.1 to 0.5°C higher in 2004) in the habitat temperature between the 2 yr, which would be related to the stenothermal nature of this species (Fig. 8).

*Scomber* spp. larvae < 6 mm BL feed mainly on copepod nauplii, and with growth, calanoid copepodites, especially *Paracalanus* spp., and appendicularians become more important as prey (Sassa et al. 2008a). *S. japonicus* and *S. australasicus* larvae were distributed in an area with higher nauplii density in 2004 than the density in the area in which they were found in 2005 (Table 4). There were areas of high nauplii density in 2005, but neither species was abundant in these. In addition, in 2004 larvae of both species occurred in the area of higher chl *a* concentration compared with that in 2005 (Table 4). Okazaki et al. (2006) indicate that the egg production rate of *Paracalanus* spp. was significantly higher in 2004 than in 2005 (25.2 versus 19.4 eggs female⁻¹ d⁻¹) in our study area due to higher temperature and chl *a* concentration. In addition, appendicularian abundance might also have been higher in 2004, since it shows positive relations with temperature and chl *a* concentration (López-Urrutia et al. 2005, Xu & Zhang 2010). These observations suggest that the food availability for both *S. japonicus* and *S. australasicus* in 2004 was better for larval growth compared with that in 2005, although this assumption is based on the limited spatial and temporal data examined.

The difference in food availability between the 2 yr could relate to the difference in the physical oceanographic conditions, i.e. an intrusion of the warm KBCNT in 2004 was much more evident compared with 2005 (Fig. 2). A permanent upwelling is found at the shelf break northeast of Taiwan, centered at ~25° 15' N to ~25° 45' N and ~121° 45' E to ~122° 45' E (Gong et al. 1997, Hsu et al. 2000, Wong et al. 2000). Although the intensity of upwelling is weak during winter when the northeastern monsoon is dominant (Ichikawa & Beardsley 2002), the upwelled waters would be a major source of nutrients that support primary production in southern ECS (Gong et al. 1997, Wong et al. 2000). The intrusion of the KBCNT is considered to bring the nutrient-enriched upwelled subsurface waters northeastward from the upwelling area to our study area (Gong et al. 1997). Additionally, in the KBCNT area where a frontal structure between the Kuroshio and coastal waters is formed, the mixed layer depth becomes relatively shallow and water column stability is kept high even in winter due to the development of a pycnocline (Nakata et al. 2007). The shallow mixed layer depth and high water temperature is also thought to enhance phytoplankton production (Nakata et al. 2007). Thus, the higher chl *a* concentration in 2004 would be related to the stronger KBCNT intrusion, resulting in better conditions for prey production; although, further study is needed for a better understanding of the prey production mechanism.

Since fast-growing larvae are considered to have a higher probability of surviving than slow-growing larvae (Takasuka et al. 2003, Takahashi & Watanabe 2004, Robert et al. 2007), the rate of growth would affect year-to-year variations in survival and recruitment. In the ECS and the western Japan Sea, recruitment success, expressed as recruit per spawner (RPS), of age-0 *Scomber japonicus* and *S. australasicus* in 2004 is calculated to be 2.6 and 2.3 times higher, respectively, than in 2005, based on analysis of population dynamics (Fisheries Agency and Fisheries Research Agency of Japan 2010). The higher recruitment success in 2004 corresponded with the higher larval growth rate in 2004. In *S. scombrus*, during years of higher temperature and food availability, higher larval
growth rates are hypothesized to allow more larvae to survive to the juvenile stage, resulting in higher year-class strength (Ringuette et al. 2002, Castonguay et al. 2008). A similar relationship might explain variations in recruitment of the 2 mackerel species in the ECS. In future, the relationships among the larval habitat conditions, larval growth and the year-to-year variations in their recruitment need to be clarified for *S. japonicus* and *S. australasicus* in the ECS to allow predictive models for their stocks to be developed.

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LITERATURE CITED


Ku JF, Tzeng WN (1985a) Age and growth of common mackerel (*Scomber japonicus* in the waters of northeastern Taiwan, with particular reference to the subpopulation discrimination. J Fish Soc Taiwan 12:1–11

Ku JF, Tzeng WN (1985b) Age and growth of spotted mackerel, *Scomber australasicus* (Cuvier), in the shelf waters of northeastern and southwestern Taiwan. J Fish Soc Taiwan 12:12–26

► Lie HJ, Cho CH (2002) Recent advances in understanding the circulation and hydrography of the East China Sea. Fish Oceanogr 11:318–328


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