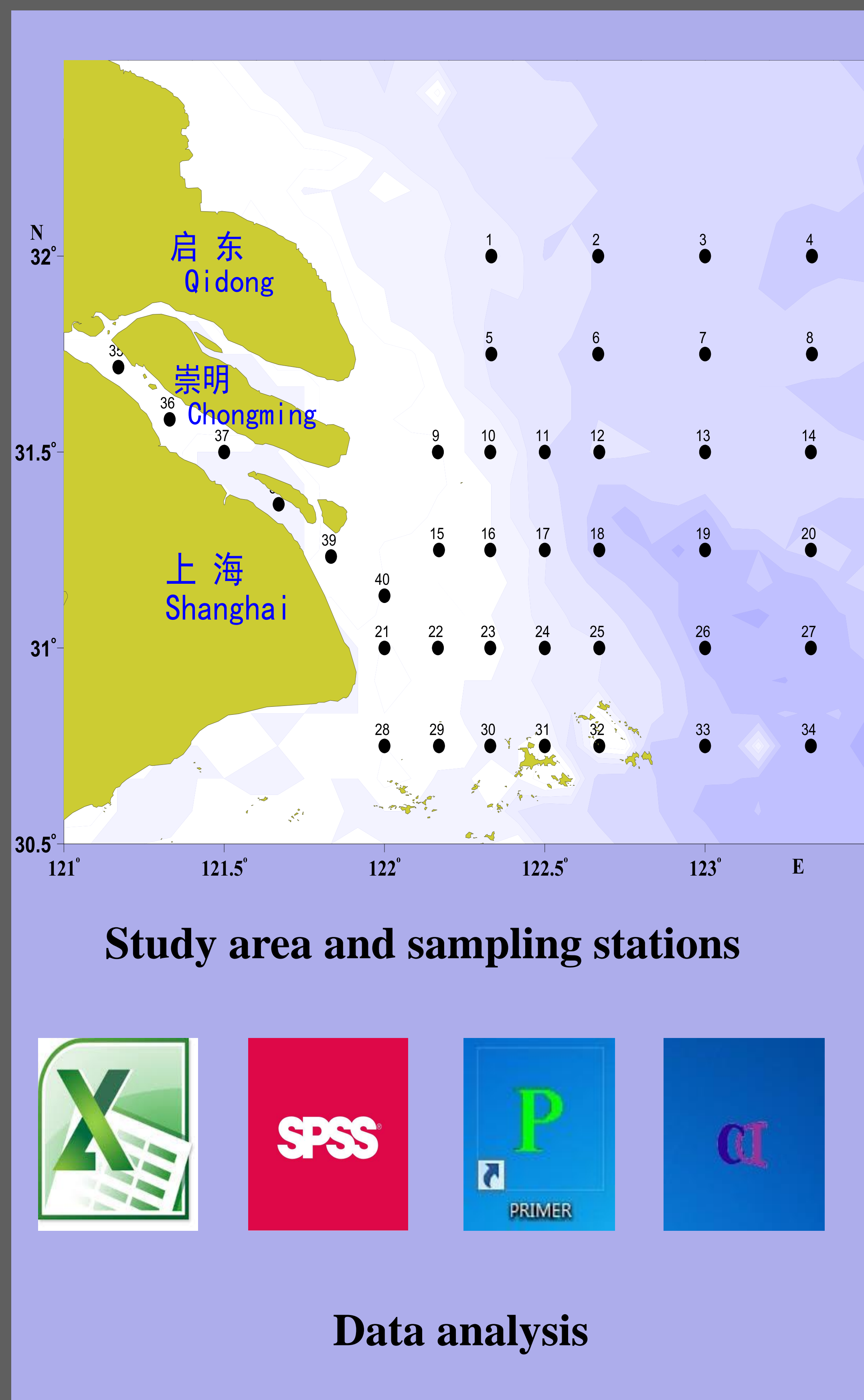


Ichthyoplankton assemblage structure of springs in the Yangtze Estuary revealed by biological and environmental visions

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Introduction

The Yangtze Estuary has been extensively modified and threatened during the last decades. The projects have considerably reduced rivers flowing to the Estuary, brought heavy contamination to estuarine aquaculture and habitation, and deteriorated water quality. This degradation has in turn led to a need for multi-year comprehensive surveys. The present study was based on surveys at four springs in 1999, 2001, 2004 and 2007. Our aims are to provide detailed characterizations of the ichthyoplankton assemblage in springs, examine the long-term dynamics of spring ichthyoplankton assemblages, and evaluate the influence of environmental factors on the spatial distribution and inter-annual variations of ichthyoplankton assemblages associated with the Yangtze Estuary.

Materials and Methods

Biological and oceanographic data were collected during four fishery evaluation cruises in four springs (05/1999, 05/2001, 05/2004, 05/2007) at 40 stations. This study encompasses 160 ichthyoplankton collections from four springs. Ichthyoplankton samples were collected by surface tows of a larva net (0.8m mouth diameter, 2.8 m long, 0.505mm mesh at the body, and 0.505mm mesh at the cod end) equipped with a flow meter. At each station, the net was towed at a depth of about 0.5m from the surface for 10 min against tidal flow. Towing speed was 2-3 knots. The samples were preserved in 5% buffered formaldehyde-seawater solution.

A conductivity, temperature, and depth device was used to measure environmental variables including depth (D), salinity (S), temperature (T), transparency (Trans), dissolved oxygen (DO), pH, total phosphorus (TP), total nitrogen (TN), suspended particulate matter (SPM), chemical oxygen demand (COD), chlorophyll a (Chla) and primary productivity (PP).

Results

Table 1 Ichthyoplankton species information.

Family	Species	EG	Percentage	1999			2001			2004			2007		
				F%	A%	P%	F%	A%	P%	F%	A%	P%	F%	A%	P%
Engraulidae	<i>Engraulis japonicus</i>	MED	20.15	34.38	51.10	32.35	10.60	16.67	11.66	26.32	6.70				
	<i>Coilia mystus</i>	AN	46.30	31.25	21.02	23.53	58.47	19.44	30.06	21.05	32.46				
	<i>Thrasops kammalensis</i>	MED	1.06				2.94			2.63	0.15				
	<i>Setipinna saty</i>	MED	0.01				2.94	1.63							
	<i>Anchovia commersoni</i>	ER	0.33				8.82	0.27		13.89	3.07	5.26	0.58		

Table 2 Dominant species determined by the IRI.

Species	1999		2001		2004		2007	
	IRI	Species	IRI	Species	IRI	Species	IRI	Species
<i>E. japonicus</i>	5,915	<i>C. mystus</i>	5,624	<i>A. bleekeri</i>	4,131	<i>A. bleekeri</i>	4,800	
<i>C. mystus</i>	2,212	<i>E. japonicus</i>	1,783	<i>C. mystus</i>	3,180	<i>C. mystus</i>	3,200	
<i>A. bleekeri</i>	1,085	<i>T. fasciatus</i>	1,405	<i>T. fasciatus</i>	1,354	<i>T. fasciatus</i>	1,100	
<i>C. hexanema</i>	462	<i>A. bleekeri</i>	753	<i>E. japonicus</i>	1,057	<i>E. japonicus</i>	800	

Table 3 Inter-annual comparison of the assemblage structure according to one-way ANOSIM (R value and significance level).

Years	ANOSIM		SIMPER
	R	P	
Global	0.002	0.443	
1999 vs. 2001	0.014	0.235	78.19
1999 vs. 2004	0.007	0.325	75.05
1999 vs. 2007	0.023	0.868	75.12
2001 vs. 2004	0.024	0.193	79.28
2001 vs. 2007	0.006	0.296	79.69
2004 vs. 2007	0.026	0.818	76.05

Table 4 Environmental factors (means and range) in the Yangtze Estuary.

Variables	1999	2001	2004	2007
D (m)	19.52 (5.00-48.00)	20.48 (5.00-50.00)	19.57 (6.00-60.00)	21.45 (5.00-52.00)
S (‰)	15.79 (2.86-27.56) ^A	17.93 (3.09-31.58) ^{A,B}	20.97 (3.09-30.19) ^B	21.46 (3.08-31.25) ^B
T (°C)	19.19 (14.98-20.86) ^A	18.14 (16.37-21.17) ^B	20.79 (19.15-22.22) ^C	18.57 (17.65-20.53) ^D

Table 8 Environmental data (means and range) of the areas resultant of CCA.

Variable	Inner	Central	Outer
	Mean (range)	Mean (range)	Mean (range)
D (m)	8.37 (5.00-10.00) ^A	12.20 (5.00-32.00) ^B	34.17 (12.00-60.00) ^C
S (‰)	4.03 (2.86-18.03) ^A	19.88 (3.30-28.61) ^B	24.50 (6.62-31.58) ^C
T (°C)	20.10 (18.68-22.22) ^A	18.91 (14.98-21.75) ^B	18.84 (15.89-21.64) ^B

Table 5 Results of CCA in the present study.

Eigenvalues	CCA axes				Total inertia
	1	2	3	4	
Eigenvalues	0.490	0.338	0.270	0.190	8.394
Species-environment correlations	0.811	0.772	0.846	0.623	

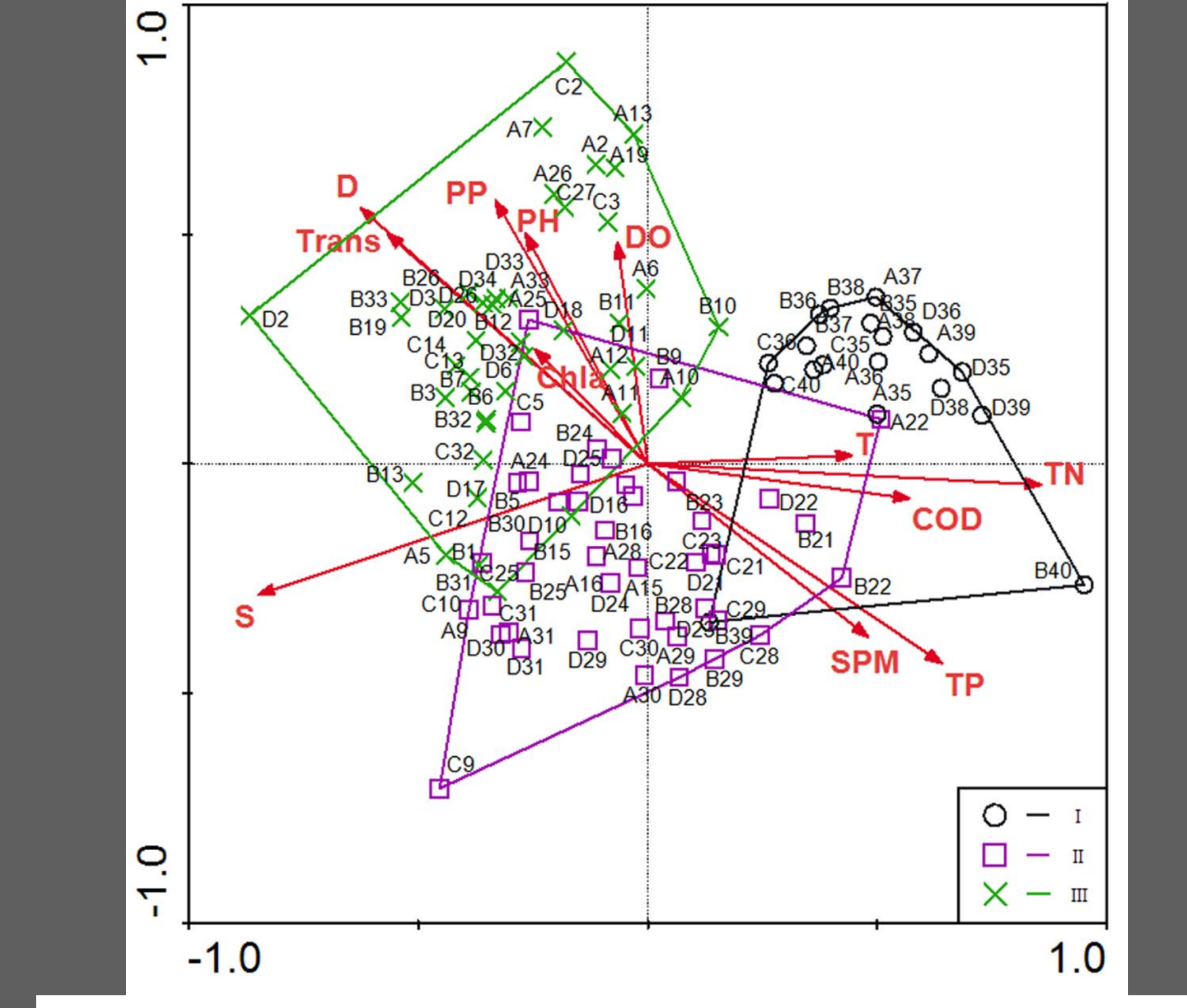
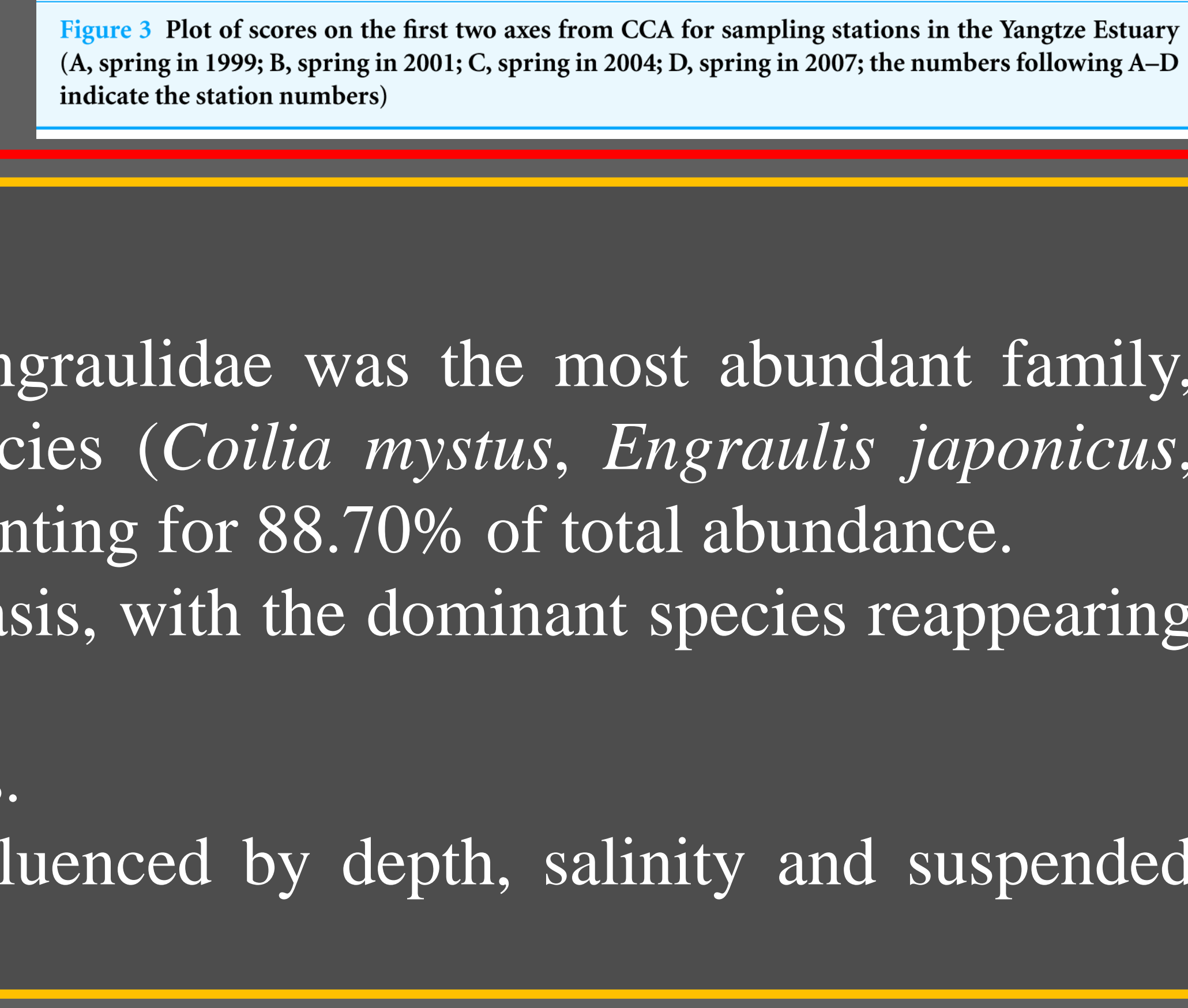


Table 6 Conditional effect of environmental variables, Canonical coefficients and intra-set correlation of environmental variables with the first two axes of CCA.

Variable	Lambda A	p	Coefficient		Correlation	
			Axis 1	Axis 2	Axis 1	Axis 2
D	0.31	0.002	0.20	0.68	-0.51	0.43
S	0.41	0.002	-0.52	-0.91	-0.69	-0.22



Discussion

- Forty-two ichthyoplankton species belonging to 23 families were collected. Engraulidae was the most abundant family, including six species and comprising 67.91% of the total catch. Only four species (*Coilia mystus*, *Engraulis japonicus*, *Trachidermus fasciatus* and *Allanetta bleekeri*) could be considered dominant, accounting for 88.70% of total abundance.
- The structure of the ichthyoplankton spring assemblage persisted on an annual basis, with the dominant species reappearing consistently even though their abundance fluctuated from year to year.
- CCA indicated a spatial structure of the ichthyoplankton assemblage in three areas.
- The observed ichthyoplankton assemblage structure appears to be strongly influenced by depth, salinity and suspended particulate matter gradients.

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