Temporal and Spatial Variations of Picoplankton and Nanoplankton and Short-Term Variability Related to Stormy Weather in the Danshui River Estuary in Northern Taiwan

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ABSTRACT

This study examined the seasonal dynamics of the bacteria, *Synechococcus* spp. and their nanoflagellate grazers in the Danshui River estuary in northern Taiwan and compared the variations in microbes (bacteria, *Synechococcus* spp. and nanoflagellate) dynamics under different (clear vs. stormy) weather conditions. Temporal variations in chlorophyll *a* concentrations, bacterial production, bacteria and *Synechococcus* spp. abundance were related to temperature, with higher values during the warmer seasons. Spatial variations in *Synechococcus* spp. were related to chlorophyll *a* concentrations and in bacterial production and abundance related to chlorophyll *a* concentrations when temperature is above 20°C (April, August, and October). Spatial variations in nanoflagellates were related to oscillations in bacterial abundance or production during non-rainy periods. We conclude that heavy freshwater input to the estuary carries a large quantity of riverine bacteria which might disrupt the relationship between bacterial and nanoflagellate abundance during rainfall events.

Key words: Danshui River estuary, Synechococcus spp., Bacterial abundance, Nanoflagellate

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1. INTRODUCTION

In aquatic environments, phytoplankton derived dissolved organic carbon (DOC) is an important source of energy and material supporting the production of heterotrophic bacteria. Because bacteria can consume 30% to 60% of the primary production via DOC, the presence of phytoplankton, therefore, strongly influences bacterial dynamics (Fuhrman and Azam 1982; Cole et al. 1988; Kirchman et al. 1991; White et al. 1991; Gasol et al. 1998; Nagata 2000).

Generally, bacterial abundance and productivity in aquatic environments may be linearly dependent on phytoplankton biomass and productivity (Gasol et al. 1998; Nagata 2000). In many oligotrophic waters, there is a close association between bacterial and phytoplankton production (White et al. 1991; Coffin et al. 1993; Gasol and Duarte 2000; Duarte et al. 2005). In estuarine environments, however, the relationship between bacteria and phytoplankton is not so closely related (Ducklow and Kichman 1983; Gasol et al. 1998; Caroppo 2002; Schultz et al. 2003). Therefore, bacteria may also process carbon derived from the allochthonous organic matter released by estuaries and sediments or introduced by rivers rather than process only carbon produced by phytoplankton in estuarine systems (Malone et al. 1991; Hoch and Kirchman 1993; Hullar et al. 1996; Kelley et al. 1998; Schultz et al. 2003). It is possible that limits in allochthonous organic matter caused by a low discharge of freshwater runoff may affect the relationship between bacterioplankton and phytoplankton. One important question might be whether phytoplankton is a major determinant of abundance and spatial distribution of estuarine bacteria during non-rainy periods, when there is less inflow of inorganic and organic materials from freshwater runoff.

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Picophytoplankton is an essential component of microbial food webs and carbon flow, particularly in warm oligotrophic waters (Kuosa 1991; Agawin et al. 2000; Worden et al. 2004; Uysal 2006). *Synechococcus* spp. have received much attention in many recent coastal phytoplankton studies (DuRand et al. 2001; Worden et al. 2004; Tsai et al. 2005, 2008; Uysal 2006). Seasonal variation in this neritic picophytoplankton is reported to be greatly influenced by irradiance and water temperature as well as by the increased turbulence, river input, and suspended matter caused by storms for spatial variations (Kuosa 1991; Agawin and Agustí 1997; Agawin et al. 1998; Tsai et al. 2008).

The abundance of nanoflagellates is highly variable. Previous studies have largely been concerned with seasonal variations in this abundance, which has often shown a peak during summer (Sander et al. 1992; Safi and Hall 1997; Tsai et al. 2008). However, some studies have shown that low nanoflagellate abundance in summer coincident with strong stratification in the upper water layer reduces the availability of nutrients (Calbet et al. 2001; Huang et al. 2008). The spatial variation of nanoflagellate abundance is clearly higher in coastal eutrophic environments than in offshore oligotrophic environments (Hwang and Heath 1997; Huang et al. 2008). It has been suggested that temperature, chlorophyll a, nutrient concentrations, and prey biomass (bacteria and picophytoplankton) may be the major controlling factors for the spatial distribution of nanoflagellates (Safi and Hall 1997; Huang et al. 2008). Hwang and Heath (1997) found that copepod abundance has an influence on the spatial distribution of HNF. Top-down controls such as grazing are thought to set a limit on nanoflagellates abundance, while nanoflagellates growth rates are thought to be set by bottom-up factors such as nutrients, prey supply, and temperature.

The Danshui River estuary is the largest estuarine system in Taiwan and is formed by the confluence of the Tahan Stream, Hsintien Stream, and Keelung River. This estuary receives a relatively small mean annual river discharge (210 m³ s⁻¹) compared to other larger estuaries, for example the Columbia River estuary in northwestern United States (6700 m³ s⁻¹) (Simenstad et al. 1990) and Elbe Estuary in Germany (700 m³ s⁻¹) (Ploug et al. 2002). Thus, in waters receiving little allochthonous organic matter as in Danshui River estuary, we hypothesized that most of the dissolved organic matter to support bacteria growth is from phytoplankton. By examining the relationships among the members of microbial loop- Chl a, bacteria, and nanoflagellate, we may better understand the transfer of energy in this marine ecosystem. Therefore, in this study, we investigated the distribution of both picoplankton (bacteria and Synechococcus spp.) and nanoflagellates at different water depths in the Danshui River estuary and studied the possible relationships between these distributional patterns and the inshoreoffshore physical, chemical, and biological gradients. Furthermore, we speculate that storms could lead to dramatic reduction in phytoplankton production, which would in turn influence bacterial growth and subsequent changes at higher consumer levels (i.e., nanoflagellate) and trophic transfer. In order to test this hypothesis, we also collected a sample during a rainy period and a week after the rain ended.

2. MATERIAL AND METHODS

2.1 Sampling

This study had eleven sampling stations (Fig. 1). Nine (stations 1 - 6 and 8 - 10) were located at three transects of the estuary over progressively greater depths of water (10, 30 and 50 m). Two additional stations (stations 7 and 11) were located in deep water (70 m). Complete plankton samples were collected during the four cruises on board the R/V Ocean Researcher II (April, August, October 2007 and January 2008). Two additional samplings were performed during and after a short period of stormy weather, 7 and 15 June 2007, respectively. Seawater was collected using a SeaBird CTD-General Oceanic Rosette assembly with 20 L Go-Flo bottles at 2 m depths. Temperature, salinity, and transmission (TM%) profiles were recorded through the water column (SeaBird CTD-General Oceanic Rosette). A custom-made flow injection analyzer with a detection limit of 0.15 µM was used for nitrate analysis (Gong et al. 1995). Chl a was collected by the filtration (25 mm GF/F) method and then measured with an in vitro fluorometer (Turner Design 10-AU-005) in triplicate using 90% acetone extracts as described in Parsons et al. (1984). Rainfall data were obtained from a local meteorological station.

Bacterial production was estimated by ³H-thymidine incorporation (Fuhrman and Azam 1982) with a conversion factor of 1.18×10^{18} cells mol thymidine⁻¹ (Cho and Azam 1988). Triplicate aliquots of water samples (30 to 40 mL) were incubated with ³H-[methyl]-thymidine (S.A., 6.7 Ci m mol⁻¹; final conc., 20 nM) in clean polycarbonate test tubes at in situ temperature in the dark for 0.5 h. Reactions were stopped by adding formaldehyde (final conc., 1%). The fixed samples, including time zero controls, were filtered through 0.2 µm cellulose nitrate filters. These filters were then rinsed sequentially 3 times each with ice cold 5% trichloroacetic acid and ice cold 80% ethyl alcohol. A scintillation cocktail (6 ml; Ultima Gold, Packard) was added after dried filters were dissolved completely in ethyl acetate (0.5 mL). Radioactivity was determined by liquid scintillation (Packard 1600).

2.2 Plankton Abundance

2.2.1 Flow Cytometric Analysis of Picoplankton

The samples (2 mL) for flow cytometry analysis were fixed with 40 μ L paraformaldehyde (0.2% final conc.), quickly frozen in liquid nitrogen, and stored in a freezer at



Fig. 1. Map of the sampling stations.

-75°C for later analysis. Abundances of picoplankton (bacteria and *Synechococcus* spp.) were measured using a Becton Dickinson LSR 6 Flow Cytometer. The samples were acquired on a low rate setting for 2 min. *Synechococcus* spp. were distinguished according to their position in plots of orange (FL2) and red (FL3) fluorescence. SYBR Green I (Molecular Probes) was used as the nucleic acid stain (Marie et al. 1997) to identify bacteria in plots of FL3 vs. green (FL1) fluorescence. Internal calibration beads (1 μm yellow-green fluorescence beads) were added as a standard.

2.2.2 Epifluorescence Microscopic Analysis of Nanoplankton

Samples for the measurement of nanoplankton quantities were fixed immediately by adding cacodylate-buffered glutaraldehyde. A Millipore filter (0.45 μ m) was used to obtain uniform distribution of cells with low pressure (< 100 mm Hg). Pigmented- and non-pigmented nanoflagellates were determined from samples (20 mL) filtered onto black Nuclepore filters (0.8 μ m). Cells left on the filter membranes were stained with DAPI (4'6-diamidino-2phenylindole) at a final concentration of 1 μ g mL⁻¹ (Porter and Feig 1980), and examined by epifluorescence microscope (1000 ×, Nikon Optiphot-2). Non-pigmented nanoflagellates were identified by their blue fluorescence under UV illumination, and pigmented nanoflagellates were identified by their orange and red autofluorescence under blue light. To obtain reliable estimates of abundance, at least 100 nanoflagellates were counted per sample.

2.3 Statistical Analysis

A correlation analysis with a significance level of 5% was undertaken between physical (temperature, light transmission), chemical (salinity, NO₃) and biological variables (Chl *a*, bacteria, *Synechococcus* spp., bacterial production and nanoflagellate). All statistical operations were performed using Statistica (version 7.0).

3. RESULTS

3.1 Temporal and Spatial Variations of Temperature and Salinity

Precipitation accumulation values were recorded from the Central Weather Bureau of Taiwan during the three-day period prior to each sampling day (Fig. 2). The weather was stormy on 7 June 2007 (Fig. 2). During clear weather, temperature varied less than 0.5°C among stations (Fig. 3), while average surface water temperatures showed marked seasonality, ranging from an average of 18.4°C in January (Fig. 3d) to 27.9°C in August (Fig. 3b). The salinity conditions at Station 4 were clearly influenced by river flow. During clear weather, surface salinity at this station was between 29.9 and 33.9 psu (Fig. 3), dipping to < 32 psu in August and October 2007 (Figs. 3b, c). In times of storms, heavy rains caused marked variation in temperature and salinity at different stations. For example, on 7 June 2007 temperature ranged from 25.3 to 28° C and salinity from 7.5 to 33.8 psu (Fig. 3e). However, one week after the storm (15 June 2007), surface salinity gradients did not vary significantly, except for one low value of < 30 psu at St. 2 (Fig. 3f).

3.2 Temporal and Spatial Variations of Hydrography

Seasonally, average NO₃ concentrations ranged from a low of 2.0 μ M in October to a high of 5.7 μ M in January (Fig. 4a). The highest mean was near the estuarine mouth at St. 4 (10 m deep) (3.1 - 6.9 μ M). Values decreased with increasing water depth, with lowest concentration (0.7 -4.8 μ M) occurring at 70 m (Fig. 4a). Mean NO₃ concentrations varied spatially during the storm on 7 June, consistently with low concentrations ranging from 5.7 to 7.3 μ M at depths from 50 to 70 m at various stations, peaking at 25.7 μ M at St. 4 (Fig. 4d). One week after the storm, however, we found no clear spatial variations in NO₃ concentrations.

Average Chl *a* concentrations ranged from 0.4 mg m⁻³ in January to 1.6 mg m⁻³ in August (Fig. 4b). Spatially, there were no significant difference in Chl *a* concentration among the stations sampled at different times of the year except for August (*ANOVA*, p > 0.05) (Fig. 4b). During the storm, average Chl *a* concentrations ranged from 0.3 to 1.1 mg m⁻³ among stations, but no difference with water depths (*ANO-VA*, p > 0.05) (Fig. 4e).

We averaged measurements taken at each station to derive a general pattern of surface water transmission (TM%) along transects with different water depth. TM (%) was found to be significantly greater in August (*t*-test, p < 0.05) for all sampling water depths except at 10-m depth during non-rainy stable weather, with a mean of 40% in January 2008 and 62% in August 2007 (Fig. 4c). Furthermore, there were significant differences in TM (%) between depths of 10 and 70 m (*t*-test, p < 0.05) (Fig. 4c).

3.3 Temporal and Spatial Patterns in Picoplankton (Bacteria and *Synechococcus* spp.) and Nanoplankton

Bacterial abundance was significantly higher in April and August (6 - 8×10^5 and 7 - 7.5 $\times 10^5$ cells mL⁻¹, re-



Fig. 2. Precipitation accumulation values were recorded in the threeday period prior to the sampling day.



Fig. 3. Non-rainy weather, storm weather and after a week of storm distribution of surface temperature (°C) and salinity (dashed lines). Samples were collected from four cruises during non-rainy weather: (a) April 2007, (b) August 2007, (c) October 2007, and (d) January 2008; Storm weather: (e) 7 June 2007 and after a week of storm: (f) 15 June 2007.



spectively) than in January (4 - 4.7×10^5 cells mL⁻¹) (*t*-test, p < 0.05) (Fig. 5a) during the non-rainy weather. However, bacterial abundance was significantly higher (10 - 16 × 10^5 cells mL⁻¹) in storm on 7 June 2007 than other samples (*t*-test, p < 0.05) (Fig. 5d).

Synechococcus spp. values were higher in August than in other months (*t*-test, p < 0.05) (Fig. 5b). August values ranged between 1.2 and 3.5×10^4 cells mL⁻¹ (mean of 2.8 × 10^4 cells mL⁻¹). We found significant differences between offshore (water 70 m depth) and inshore (water 10 - 30 m depth) distribution of *Synechococcus* spp. in our August and October samples (*t*-test, p < 0.05) (Fig. 5b). During the 7 June storm, abundances (0.7 to 1.5×10^5 cells mL⁻¹) were slightly lower than in August, but were not different from the stations (*ANOVA*, p > 0.05) (Fig. 5e).

Figure 5c shows the spatial and temporal variations in concentrations of total nanoflagellates in clear weather at the study stations. At depths of 10 m, maximum abundance $(> 2 \times 10^3 \text{ cells mL}^{-1})$ was observed in April 2007, while at

other stations they were about 1×10^3 cells mL⁻¹ (Fig. 5c). However, at depths of 50 and 70 m, concentrations of nanoflagellates were higher in August and October than in other months (Fig. 5c) (*t*-test, p < 0.05). However, in stormy weather, there was a general trend of increase in total nanoflagellate abundance with increase of water depth from 10 to 70 m (Fig. 5f). Regardless of the weather conditions, pigmented nanoflagellates dominated the nanoflagellate community (57% - 88%) (data not shown) in all samples.

3.4 Bacterial Production

Figure 6 gives the bacterial production rate of all stations. Bacterial production rates were significantly higher (*ANOVA*, p < 0.05) in August (2.5 - 16.5 mgC m⁻³ d⁻¹) than other months (1 - 11.5 mgC m⁻³ d⁻¹) (Fig. 6). For further analysis of the relation between bacterial production rates and bacterial abundance, we found that in non-rainy weather, except in August, the greater the bacterial abundance,



Fig. 5. Comparison of distributions of bacteria (a) (d), *Synechococcus* spp. (b) (e), and total nanoflagellates (%) (c) (f) along water depths from 10 to 70 m during non-rainy and the rainfall periods. Marks as in Fig. 4.



Fig. 6. Relationship between bacterial production and bacterial abundance. Marks as in Fig. 4.

the greater the bacterial production rate (< 5 mgC m⁻³ d⁻¹). We did not have data of the bacterial production rate for 15 June; however, we found no significant relationship between bacterial production rate and bacterial abundance during the 7 June storm event and bacterial production rates were low (< 5 mgC m⁻³ d⁻¹) (Fig. 6).

4. DISCUSSION AND CONCLUSION

4.1 Temporal and Spatial Variations of Chl a

Fluctuations in Chl *a* concentrations were related to temperature (Table 1). Moreover, as for spatial changes in Chl *a* concentrations, the result showed a negative correlation with salinity in April (Table 2). We found no correlations between Chl *a* concentrations and any environmental parameter [i.e., turbidity (TM%), salinity, nutrient] in other months. Controls on spatial dynamics of phytoplankton in the estuary are more complex than in oligotrophic waters. Generally, growth rates of phytoplankton are controlled by light intensity, temperature, and nutrient concentrations in the euphotic zone. However, photosynthetic studies of the St. Lawrence River transition zone have indicated that the phytoplankton are well adapted to the intermittent exposure to bright light which might occur within the turbid, wellmixed waters (Vincent et al. 1994). Although our samples were all collected during the daytime, we cannot rule out the possibility that Chl *a* dynamics are partly related to the photoadaptation, as has been shown elsewhere (Vincent et al.

Table 1. Pearson correlation coefficients for the temporal relationship between microbes (Chl *a*, bacterial, *Synechococcus* spp., nanoflagellate abundance and bacterial production) and surface water temperature and Chl *a* concentrations.

	temperature	Chl a
Chl a	0.58*	
bacterial abundance	0.63*	0.60*
Synechococcus spp. abundance	0.61*	0.87 **
nanoflagellate abundance		
bacterial production	0.45*	0.82**
* Correlation is significant at the 0.0)5 level (2-tailed).	

* Correlation is significant at the 0.05 level (2-tailed)

** Correlation is significant at the 0.01 level (2-tailed).

Table 2. Pearson correlation coefficients for the spatial relationship between microbes (Chl *a*, bacterial, *Synechococcus* spp., nanoflagellate abundance and bacterial production) and surface water salinity, Chl *a* concentrations, bacterial abundance and production.

	salinity	Chl a	bacterial abundance	bacterial production
Chl a	-0.85** (Apr.)			
bacterial abundance	-0.81** (Apr.)	0.76* (Apr.)		
	-0.95** (<u>storm</u>)	0.60* (Aug.)		
		0.85** (Oct.)		
Synechococcus spp.abundance	0.78* (Apr.)	0.92** (Apr.)		
	0.55* (Oct.)	0.93** (Aug.)		
	0.70* (<u>storm</u>)	0.65* (Oct.)		
	0.90** (after storm)	0.52* (Jan.)		
		0.66* (<u>storm</u>)		
		0.56* (after storm)		
nanoflagellate abundance			0.45* (Apr.)	0.70* (Apr.)
			0.64* (Oct.)	0.57* (Oct.)
				0.67* (Jan)
bacterial production	-0.47* (Apr.)	0.60* (Apr.)	0.55* (Apr.)	
	-0.90** (Aug.)	0.60* (Aug.)	0.62* (Oct.)	
		0.78* (Oct.)		

* Correlation is significant at the 0.05 level (2-tailed).

** Correlation is significant at the 0.01 level (2-tailed).

1994). Other factors, such as loss due to grazing, may also have influenced this spatial pattern of Chl *a*. The density of macrozooplankton and microzooplankton are known to be high in estuaries (Lovejoy et al. 1993; Vincent et al. 1994; Calbet 2001; Winkler et al. 2003), and thus a drop in Chl *a* concentration in these regions (water depth at 10 m) may result from increased grazing. On the other hand, storms can cause a decrease *in situ* light intensity and an increase in suspended matter from terrestrial runoff or turbulence, which can result in lower Chl *a* concentrations as in our study (< 1 mg m⁻³) (Fig. 4e).

4.2 Temporal and Spatial Variations of *Synechococcus* Spp.

Numerous seasonal studies have shown that an abundance of *Synechococcus* spp. is the greatest during warmer months (Ning et al. 2000; Tsai et al. 2005, 2008). The influence of irradiance and water temperature on the seasonal distributions of abundance and biomass of *Synechococcus* spp. has been established (Kuosa 1991; Chang et al. 1996; Agawin et al. 1998; Tsai et al. 2008). Although the relationship between temperature and abundance may be confounded by an autocorrelation between light and temperature, we suggest that temperature, including its relative variability, is a major physical factor affecting the numerical response of *Synechococcus* spp. (Table 1).

We also observed peaks of Synechococcus spp. abundance in summer (August) (Fig. 5b), which may be explained by transient imbalances between growth rates and grazing rates of Synechococcus spp. outpacing nanoflagellate grazing rate and the increased food concentration for nanoflagellates (i.e., increased bacterial production in summer) (Fig. 6). This could result in a release in grazing pressure on Synechococcus spp. because, as Landry et al. (1984) have suggested, the threshold of Synechococcus spp. abundance corresponds significantly to grazing by nanoflagellates at 6×10^4 cells mL⁻¹. It is also evident from Fig. 6 that bacterial abundance did not increase along with increasing bacterial production rate in August, suggesting that the upward transfer of bacterial organic matter in the food web occurred mainly during the warm season (August) in this study.

Synechococcus spp. have not been measured in the Danshui River estuary. While the importance of *Synechococcus* spp. and other picophytoplankton is well documented in the open oceans (Fogg 1995), their importance in estuaries is less well studied (Murrell and Lores 2004). Murrell and Lores (2004) reported that *Synechococcus* spp. in Pensacola Bay in general represented 43% of the total chlorophyll, but up to over 90% in summer. In other estuaries, their contribution to the total chlorophyll appears to be much smaller. For example, in the York River estuary, picophytoplankton comprises 7% of chlorophyll over an annual cycle, peak-

ing at 14% during summer (Ray et al. 1989). In the present study, we found spatial variations in *Synechococcus* spp. to be related to chlorophyll *a* concentrations (Table 2), which suggests that *Synechococcus* spp. dominate the chlorophyll biomass in this estuary. According to literature, contribution of picophytoplankton to bulk chlorophyll is only dominant in oligotrophic environments where chlorophyll levels range from 0.5 to 1 μ g L⁻¹, and their contribution diminishes with increases in trophic state. While this pattern may hold for temperate estuaries, Danshui River estuary and similar subtropical systems such as Pensacola Bay (Murrell and Lores 2004) do not fit this pattern.

4.3 Bacterial Variations

Previous studies have reported temperature to strongly affect estuarine bacterial abundance and growth (Shiah and Ducklow 1994; Schultz et al. 2003; Tsai et al. 2008); nevertheless, Ochs et al. (1995) found and later confirmed by Shiah and Ducklow (1994) that bacterial growth rates were unrelated to temperature above 14°C. Their findings suggest that temperature might control growth activity during the colder months. Growth in the warmer seasons seems to be controlled by factors such as substrate supply or availability of nutrients (Keil and Kirchman 1991). Our results showed that spatial variations in bacterial production and abundance were related to chlorophyll *a* concentrations (Table 2) when temperature is above 20°C (April, August, and October). Our result agrees with those of previous studies reporting temperature not to be an important determinant of bacterial production rates during the warmer seasons (Shiah and Ducklow 1994; Ochs et al. 1995). In addition, during the warmer August season, we found the minor changes in bacterial abundance to be accompanied by high bacterial production (5 - 15 mgC m⁻³ d⁻¹) (Fig. 6), suggesting that predation on bacteria exerted more influence on bacteria dynamics than resource supply during the warmer August season (Boissonneault-Cellineri et al. 2001; Hahn et al. 2001; Schultz et al. 2003).

There appeared to be different factors controlling bacterial production and abundance under stable, clear weather and stormy weather. Primary production appears to have provided most of the carbon needed for bacterial production in stable, clear weather, bacterial production and abundance were all related to Chl *a* concentrations (Table 2). We also found that bacterial abundance was significantly higher (10 - 16 × 10⁵ cells mL⁻¹) during the storm on 7 June than during the non-rainy warm periods in August (Fig. 5d) and that changes in bacterial abundance were mostly closely related to changes in salinity (*p* < 0.05) (Table 2). Previous studies have suggested that the concentration of nutrients and dissolved substrates along the salinity gradient may be important in the control of bacterial growth (Kelley et al. 1998; Schultz et al. 2003); however, in our study there was no evidence to indicate that bacterial production was related to changes in salinity of the storm on 7 June (Table 2). Furthermore, in the Danshui River estuary during periods of heavy rainfall, the average value of bacterial production was too low (about 3 - 5 mgC m⁻³ d⁻¹) (Fig. 6) to significantly raise bacterial abundance $(25 \times 10^5 \text{ cells mL}^{-1})$. On the other hand, we estimated bacterial cell carbon in our study to be 25 fg C cell⁻¹, a value characteristic of bacteria in nutrient-rich waters (Bell 1993). Based on this value, we estimated that during the stormy event on 7 June bacterial growth rate was about 0.12 d⁻¹ (BP/BB). This growth rate was much lower than that reported by Tsai et al. (2008) in their study of the western subtropical coast of the Pacific. Based on our observations, it is possible that the increase in bacterial abundance during the stormy event on 7 June was not as much the result of increased growth as that of heavy freshwater input carrying a large quantity of riverine bacteria into the estuary.

Nanoflagellates provide an important link with the transfer of carbon within the microbial food-web; therefore, it is important to understand the factors which affect their abundance in order to develop an understanding of the flux of carbon in this estuary. Fifty-seven to eighty-eight percent of the nanoflagellates in our samples were pigmented, a percentage similar to that (> 50%) reported by Tsai et al. (2007) for the western subtropical coast of the Pacific. Previous studies of pigmented nanoflagellates have highlighted the potential ecological significance of these organisms in the aquatic environment and have found evidence that many pigmented nanoflagellates are capable of phagotrophy (Bird and Kalff 1986; Caron et al. 1990). In the present study, because we assumed that pigmented nanoflagellates were capable of phagotrophy, we correlated data over all surface data from separate sampling stations, between bacteria abundance/production and total nanoflagellate abundance in our study site. As shown in Table 2, we founded a correlation between bacteria production and nanoflagellate abundance in non-rainy weather (April, October, and January) (Table 2). We suggest that spatial variations of nanoflagellates are related to oscillations of bacterial abundance or production in the non-rainy periods. Moreover, the correlations between nanoflagellates and bacterial abundance may be lost during storms (Table 2), which might explain the disruption in the relationship between bacterial and nanoflagellate abundance to be caused by the sudden influx of riverine bacteria.

In conclusion, our study shows that spatial variations in *Synechococcus* spp. are related to chlorophyll *a* concentrations in the Danshui River estuary. Moreover, spatial variations in bacterial production and abundance are related to chlorophyll *a* concentrations when temperature is above 20°C (April, August, and October). Spatial variations in nanoflagellates are related to oscillations of bacterial abundance or production in the non-rainy periods. Most importantly, we suggest that heavy freshwater input can carry a large quantity of riverine bacteria into the estuary, possibly explaining how riverine bacteria might have disrupted the relationship between bacterial and nanoflagellate abundance during rainfall events.

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