

Temporal and spatial scaling of planktonic responses to nutrient inputs into a subtropical embayment

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ABSTRACT: We carried out a study of the spatial and temporal effects of land-derived material on water column nutrients and plankton dynamics in a subtropical estuary. The study had 2 parts: (1) a 3 yr synoptic monitoring program, and (2) a shorter 1.5 yr study during the second half of the program, which focused on individual pulses driven by discrete rainfall events. Although we found spatial differences in some water column parameters within Kāne'ohe Bay and an adjacent oceanic site, inorganic nutrient levels were generally comparable in the Bay and offshore. One difference was that *Prochlorococcus* spp. numerically dominated the plankton at the oceanic site whereas *Synechococcus* spp. dominated at all Bay sites. The switch in dominance appears to be due to light characteristics and dissolved organic nitrogen (DON), but not dissolved inorganic nutrient availability. There were no annual cycles in water column parameters within the Bay; however, a comparison of dry and wet seasons did show some differences. Planktonic cell abundance was in general lower during the wet season, with the exception of opportunistic diatoms that were more abundant during the wet season. A drought during the study period may have influenced our results. Pulses were characterized by an elevation in inorganic nutrient concentrations in the Bay close to the stream mouth. The general response was an increase in abundance of microphytoplankton and chl *a* after a 3 to 6 d lag following the nutrient increase. Picophytoplankton showed an increase in fluorescence per cell after a 12 to 24 h lag, probably related to a decrease in irradiance associated with turbidity in runoff. The Bay can act as source of dissolved inorganic nutrients and plankton for oceanic waters; however, planktonic populations in the Bay are primarily autochthonous and do not represent an oceanic source of nutrients for plankton consumers within Kāne'ohe Bay.

KEY WORDS: Picoplankton · Phytoplankton · Zooplankton · Nutrients · Runoff

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INTRODUCTION

Coral reef ecosystems are characterized by low net community productivity (similar to the surrounding oligotrophic oceanic water) such that the system accumulates biomass slowly and exports little (Hatcher 1990). Typically, tropical oceanic water has low nutrient concentrations (Crossland & Barnes 1983, Furnas 1992) and is dominated, in terms of biomass and production, by pico- (0.2 to 2 μm) and nanoplankton (2 to 20 μm) (Ducklow 1990, Olson et al. 1990 a,b, Tremblay & Legendre 1994, but see Scharek et al. 1999). Advec-

tion from the surrounding ocean (including upwelling) and terrestrial run-off are considered to be the main sources of new nutrients supporting net production in reef ecosystems (Furnas et al. 2005).

The oligotrophic conditions of oceanic waters surrounding coral reefs can be quite different from relatively nutrient-rich inshore conditions on high islands or atolls. For example, high productivity has been described within French Polynesian atoll lagoons in comparison to oceanic waters (Charpy 1996, Ferrier-Pagès & Furla 2001), and these lagoons can also represent a source of nitrogen for the adjacent oceanic waters

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(Charpy-Roubaud et al. 1990). Short episodes of high stream discharge can be responsible for most of the annual delivery of dissolved and particulate materials into bays or lagoons on high islands (Taguchi & Laws 1989, Dubinsky & Stambler 1996, Hoover 2002, Ringuet & MacKenzie 2005), increasing dissolved nutrient concentrations (N, P and dissolved organic C [DOC]) in the receiving water. However, the scale of spatial and temporal responses of plankton communities to stream discharge and subsequent nutrient inputs are still poorly understood, especially within atolls or bays with high water exchange with adjacent oceanic water.

Kāne'ohe Bay is located on the windward northeast coast of O'ahu, Hawai'i, and is one of the most intensively studied coral reef systems in the world (e.g. Smith et al. 1981, Taguchi & Laws 1987, 1989, Hunter & Evans 1995, Laws & Allen 1996, see also www.hawaii.edu/cisnet). Water circulation in the Bay is primarily influenced by tradewinds that drive oceanic water across the foreereef and out through a northern deep channel (Fig. 1). At our north bay (NB) study site, most of this water flows quickly offshore (Bathen 1968). At the more protected central bay (CB) and, especially, south bay (SB and MP) study sites, water remains in the system for longer periods of time (Bathen 1968, Smith et al. 1981). Stream discharge in Hawai'i responds rapidly to brief, intense rainstorms that occur at unpredictable intervals throughout the year (Kinzie et al. 2006). As stream channels are short and steep and rain events can be locally intense, pulsed discharge into Kāne'ohe Bay can result in marked responses by the Bay system (Banner 1974, Taguchi & Laws 1989, Jokiel et al. 1993, Hoover 2002, Ringuet & Mackenzie 2005). Pulsed discharge can exceed median streamflow by 2 to 3 orders of magnitude (Oki 2004). In addition, channelization may increase the peak discharge and decrease the duration of pulses (Laws & Roth 2004).

The aim of this study was to reveal the pattern of distribution of planktonic communities over a range of stream discharge conditions in Kāne'ohe Bay and over a spatial gradient of exchange with oceanic water. We monitored the temporal dynamics of annual cycles as well as short-term stream discharge pulses. As reported in previous studies (Chisholm 1992, Agawin et al. 2000), we expected that increased nutrients would shift phytoplankton to larger size classes, followed by a similar change in zooplankton size. During periods of low nutrient input, such as during a drought, phytoplankton should be dominated by picoplankton (Raven

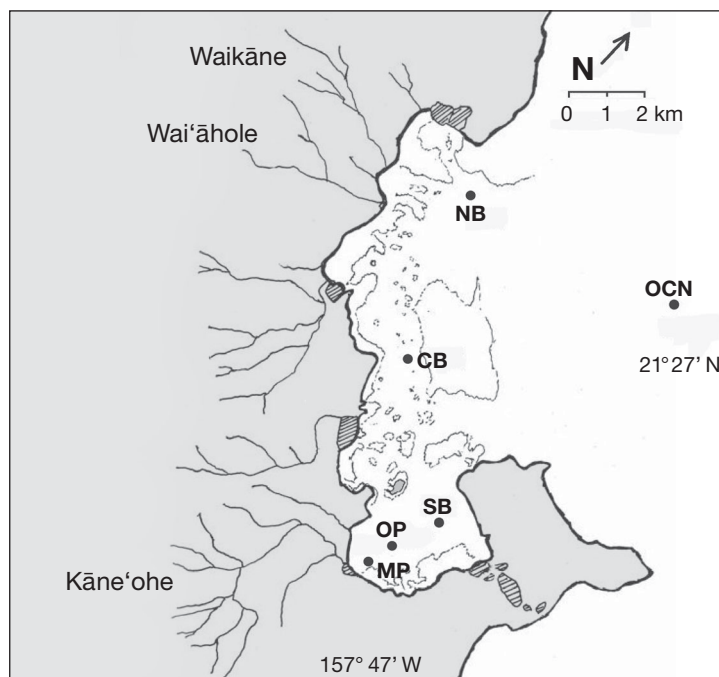


Fig. 1. Kāne'ohe Bay, O'ahu, Hawai'i. Major stream systems, fish ponds (hatched), and reefs (dotted) shown. (●) Sampling site (OCN: oceanic site; NB: north bay; CB: central bay; SB: south bay; OP: outside plume; MP: mid plume)

1998). Our main goal was to reveal the type and temporal and spatial scales of responses of planktonic components to nutrient increase associated with stream discharge, and the effects of oceanic water and Bay circulation patterns on the dissipation of planktonic changes.

MATERIALS AND METHODS

Baseline sampling. From 1998 to 2001, a monitoring program (CISNet Kāne'ohe Bay: www.hawaii.edu/cisnet) collected data for comparison with earlier work during the 'sewage' and 'post-sewage' periods in Kāne'ohe Bay (Smith et al. 1981). Parameters monitored in this program included water column chl *a* and nutrients, as well as physical factors including temperature, salinity, light attenuation and suspended sediments. Sampling was conducted at 5 stations (Fig. 1) and took place every 2 wk for north bay (NB), central bay (CB), south bay (SB) and mid plume (MP) sites and roughly every 3 mo for oceanic (OCN) sites. Depth (lead sounding) and Secchi depth (30 cm diameter white disk) were recorded at each site. Profiles for temperature and salinity were obtained with a SeaBird CTD. Light penetration was determined using paired deck and submerged readings with photosynthetically

active radiation (PAR) meters (LiCor) equipped with upward-facing cosine sensors. Water was collected using a submersible pump, mounted on a float to maintain the intake at a constant depth of 0.5 m. All water column parameters were taken from this 0.5 m depth unless otherwise noted. Water samples were all held in 4 l opaque acid washed plastic bottles, after initial rinsing with water from the pump, and kept on ice until processed. Four water samples were collected at each site.

In the last 1.5 yr of this program, we initiated a sampling project addressing the planktonic communities in greater detail. For this more intense program, pico-, nano- and microplankton (collectively called the microbial fraction) were collected from the water column samples described above. At 3 of the stations (NB, CB, and SB), zooplankton samples were also taken every 2nd wk.

In addition, during this period we intensively monitored plumes arising from rainfall episodes. These plumes developed at the mouth of Kāne'ōhe Stream (Fig. 1) and could be visually followed for hours to days (Jokiel et al. 1993).

Nutrient analysis. Aliquots of approximately 100 ml were taken from the 0.5 m water samples and filtered through a 25 mm GF/F filter in a syringe holder. Samples were collected in acid-rinsed 125 ml plastic bottles, first rinsed with filtered sample water. Samples were frozen and analyzed for inorganic and total nutrients using the methods of Smith et al. (1981) by SOEST Analytical Laboratory (until April 2001) and University of Washington Marine Chemistry Laboratory (from April to July 2001). Parameters measured were: NH_4^+ (referred to as ammonium in this paper), NO_3^- (+ NO_2^-) (as nitrate), PO_4^{3-} (as phosphate), SiO_2 (as silicate), total dissolved nitrogen (TDN), and total dissolved phosphorus (TDP).

Particulate carbon and nitrogen. Total particulate water column carbon and nitrogen were measured by filtering 1 l of water on pairs of pre-combusted GF/F glass fiber filters, with 3 replicates from different water sample bottles. Acidification (10% HCl) was performed on 1 filter of each pair to dissolve carbonates. Filters were dried overnight at 60°C and stored in desiccators until analyzed with a Perkin-Elmer CHN Analyzer.

Chlorophyll analysis. Three size fractions were collected for the CISNet program (<2 μm , >2 μm , and GF/F). Water samples were prefiltered through a 183 μm mesh to remove large particles. Next, approximately 250 ml was filtered onto a 47 mm Whatman GF/F filter and another 250 ml filtered onto a 47 mm 2.0 μm membrane filter (Millipore Isopore). Approximately 100 ml of the filtrate that passed through the 2.0 μm membrane filter was then filtered onto a 47 mm

0.2 μm membrane filter (Millipore Isopore). Volumes filtered were recorded for each sample. All 3 size fractions were obtained in triplicate for each site. Filters were placed into 15 ml plastic centrifuge tubes, and 10 ml of 100% acetone was added to each sample tube. Each sample was sonicated on ice in the centrifuge tubes for 20 s with a HeatSystems W-370 sonicator set at maximum power for the micro tip. Samples were held at -5°C in the dark until reading, usually within 24 h.

Before reading, samples were cleared in a refrigerated centrifuge at 20000 rpm for 20 min and then allowed to come to room temperature. Chlorophyll was determined from fluorescence measured on a Turner Designs 10-AU fluorometer. Fluorescence before and after acidification was used to determine chl *a* using the equations given in Trees et al. (2000). The Turner Designs solid chlorophyll standard was run at the beginning and end of each set of samples. Additionally, a dilution series of pure chl *a* (Sigma) was measured at the end of each sample run.

Pico- and nanoplankton. We used flow cytometry to quantify heterotrophic bacteria, *Prochlorococcus* spp. (hereafter *Prochlorococcus*), *Synechococcus* spp. (hereafter *Synechococcus*), and picoeukaryotes. Three water samples (2 ml from 4 l water sample bottles) were fixed with 1% paraformaldehyde + 0.05% glutaraldehyde (final concentration) and frozen in liquid nitrogen. Samples were stored at -80°C or in dry ice until processing. For determination of bacterial and picoeukaryote abundance we used a B&D FACScalibur bench machine as described in Gasol & Morán (1999). Nanoeukaryote abundance was determined on 20 ml subsamples stained with DAPI and filtered through a 0.2 μm filter (Nucleopore). Stained cells were directly enumerated using epifluorescence microscopy. Cell sizes of heterotrophic bacteria, *Synechococcus*, picoeukaryotes, and nanoeukaryotes were measured on the same filters using framegrabber and software for image analysis as described in Massana et al. (1997). Fluorescence (red indicating chlorophyll and orange indicating phycoerythrin) per cell and relative to beads was calculated from flowcytometry data as reported in Gasol & del Giorgio (2000).

Microplankton (phytoplankton and ciliates). To quantify phytoplankton and ciliate cell numbers, 350 ml water samples (from 4 l water sample bottles) were preserved with Lugol's solution (10% final concentration). Subsamples of 100 ml were transferred to settling chambers, and major groups of microphytoplankton were quantified using an inverted microscope.

Cell sizes (length and width) were measured using an ocular micrometer. Cell biovolumes were estimated from the length and width measurements, assuming

Table 1. C and N conversion factors for selected cell types

Cell type	Carbon	Nitrogen	Reference
Heterotrophic bacteria	9.3 fg cell ⁻¹	1.8 fg cell ⁻¹	Gundersen et al. (2002)
<i>Prochlorococcus</i> spp.	37 fg cell ⁻¹	4 fg cell ⁻¹	Heldal et al. (2003)
<i>Synechococcus</i> spp.	192 fg cell ⁻¹	21 fg cell ⁻¹	Heldal et al. (2003)
Picoeukaryotes & nanoeukaryotes	183 fg μm ⁻³	26.1 fg μm ⁻³	Caron et al. (1995)
Diatoms & dinoflagellates	pg cell ⁻¹ = 0.109 × (μm ³) ^{0.991}	pg cell ⁻¹ = 0.0172 × (μm ³) ^{1.023}	Montagnes et al. (1994)

the nearest geometrical shape (Edler 1979). Carbon (C) and nitrogen content (N) were estimated using a set of conversion factors selected from the literature (Table 1).

Zooplankton samples. Zooplankton was divided into large, medium and small size fractions. Triplicate samples were collected with 10 m vertical tows (approximately 1 m s⁻¹) using a 35 μm mesh net (30 cm diameter). Only NB, CB and SB sites were sampled, as MP was too shallow and time constraints prevented sampling at OCN. The entire sample from each 0.71 m³ tow was first poured onto a 183 μm mesh and rinsed. This provided the >183 μm fraction (large fraction). This size fraction was not retained on the first 7 sampling dates. The fraction that passed through the 183 μm mesh was split into 3 fractions. One was collected on a precombusted 25 mm GF/F filter for CHN analysis. The remaining sample was passed through a 63 μm screen. Material retained was the <183>63 μm sample (medium fraction), while material passing through the screen was the <63>35 μm sample (small fraction). Each biomass sample was dried at 60°C overnight and reweighed. The biomass samples were combusted at 500°C for 4 h and reweighed to provide ash free dry mass (AFDM). Zooplankton biomass data are reported per m³ of Bay water sampled. The CHN samples were processed using the same methods as the water column CHN filters.

The large fraction included material that would not have been sampled in the water column sampling (as those samples were all prefiltered through a 183 μm mesh). This sample usually consisted of chaetognaths, larger copepods and larval crustaceans, as well as occasional larval or juvenile fishes. Because the net diameter and mesh size were small, this fraction is probably not a quantitative sample of the fast-swimming zooplankton. The small fraction

consisted of smaller copepodites and nauplii, while the medium fraction contained mostly adult copepods and some decapod larvae.

Pulse events. During high rainfall events, environmental parameters were measured daily at sites close to the mouth of Kāne'ohe Stream (MP) as well as further offshore and out of the plume (OP). Rainfall data were obtained from the National Weather Service hydronet (www.prh.noaa.gov/hnl) and discharge data from the US Geological Service Kāne'ohe Stream gauge (<http://waterdata.usgs.gov/nwis>). The dispersion of the freshwater plume was estimated by measuring salinity of a shallow scoop of water from the sur-

face layer (<10 cm depth) with a handheld YSI model 55 salinometer. Water for nutrient and plankton sampling was hand collected daily during events from subsurface waters (~50 cm depth) at the MP site. Measurements of pico-, nano- and microplankton, dissolved inorganic nutrients, chl *a*, and particulate carbon and nitrogen were made from these water samples using the methods outlined above. During pulsed stream discharge events, chl *a* samples consisted of 3 replicate samples on 47 mm Whatman GF/F filters. To analyze total suspended solids (TSS), subsamples of approximately 250 ml were filtered through pre-weighed and ashed GF/F filters for subsequent drying at 60°C for measurement of TSS. Pico-, nano- and microplankton were sampled during the April 2000, October 2000 and June 2001 pulse events. Data from pulse events were compared to baseline CISNet data (3 yr means from CISNet sampling period at the MP site).

Statistical analyses. Statistical analyses were performed using SAS. Data were transformed if necessary. Post-hoc tests following ANOVA used Duncan's test with α set at 0.05.

RESULTS

Spatial differences

The most striking spatial difference in water column nutrients during the 3 yr study was the elevated nitrate concentration at the NB site compared to OCN, CB, SB and MP (Table 2). TDN was significantly lower at the OCN site, while MP had the highest concentration (Table 2). Dissolved organic nitrogen (DON, the difference between TDN and nitrate + ammonia) showed a gradual increase from 5.7 μM at the OCN site to 8.3 μM at the MP site. Silicate was significantly higher

Table 2. Comparison of water column and plankton parameters (mean \pm SE) at sampling sites (OCN: oceanic site; NB: north bay; CB: central bay; SB: south bay; MP: mid plume). Letters in parentheses denote significant differences determined by Duncan's test. nd: no data; *Chaetoceros* was not found in the OCN samples

	OCN	NB	CB	SB	MP
Water column					
Salinity (PSU)	35.0 \pm 0.07 (a)	34.7 \pm 0.04 (ab)	35.0 \pm 0.04 (a)	35.1 \pm 0.04 (a)	34.3 \pm 0.22 (b)
N	5	71	71	71	59
TSS (mg l ⁻¹)	0.39 \pm 0.06 (d)	2.7 \pm 0.22 (b)	1.9 \pm 0.12 (c)	2.0 \pm 0.10 (bc)	3.5 \pm 0.22 (a)
N	8	71	71	71	59
Secchi depth (m)	28 \pm 3.1 (a)	6.2 \pm 0.24 (b)	6.6 \pm 0.25 (b)	5.4 \pm 0.20 (b)	3.3 \pm 0.12 (c)
N	6	70	70	70	57
Extinction					
coefficient (m ⁻¹)	0.11 \pm 0.02 (c)	0.29 \pm 0.01 (b)	0.24 \pm 0.01 (b)	0.29 \pm 0.01 (b)	0.47 \pm 0.01 (a)
N	4	62	62	62	54
Phosphate (μ M)	0.09 \pm 0.01 (a)	0.07 \pm 0.005 (b)	0.06 \pm 0.005 (b)	0.08 \pm 0.004 (ab)	0.71 \pm 0.004 (b)
Nitrate (μ M)	0.05 \pm 0.02 (b)	0.28 \pm 0.04 (a)	0.09 \pm 0.03 (ab)	0.05 \pm 0.01 (b)	0.20 \pm 0.08 (ab)
Ammonia (μ M)	0.12 \pm 0.02 (a)	0.14 \pm 0.01 (a)	0.14 \pm 0.02 (a)	0.12 \pm 0.01 (a)	0.10 \pm 0.01 (a)
Silicate (μ M)	2.3 \pm 0.22 (c)	7.1 \pm 0.20 (b)	5.0 \pm 0.24 (bc)	6.9 \pm 0.21 (b)	17 \pm 1.6 (a)
N	8	72	72	72	51
TDP	0.32 \pm 0.02 (a)	0.30 \pm 0.01 (a)	0.30 \pm 0.01 (a)	0.32 \pm 0.01 (a)	0.32 \pm 0.01 (a)
TDN	5.9 \pm 0.53 (c)	7.0 \pm 0.24 (b)	7.2 \pm 0.24 (b)	7.4 \pm 0.23 (b)	8.6 \pm 0.35 (a)
DON	5.7 \pm 0.53 (bc)	6.5 \pm 0.22 (bc)	6.9 \pm 0.25 (b)	7.2 \pm 0.23 (b)	8.1 \pm 0.29 (a)
N	8	39	39	39	31
Particulate C & N (mg m ⁻³)					
PC total	49 \pm 8.4 (c)	190 \pm 15 (b)	180 \pm 12 (b)	200 \pm 84 (b)	290 \pm 16 (a)
PN	5.1 \pm 0.72 (d)	23 \pm 1.2 (b,c)	22 \pm 1.2 (c)	28 \pm 1.5 (b)	46 \pm 2.3 (a)
POC (acidified sample)	54 \pm 14 (c)	140 \pm 11 (b)	140 \pm 11 (b)	170 \pm 8.0 (b)	290 \pm 18 (a)
PN (acidified sample)	5.9 \pm 1.3 (d)	24 \pm 1.1 (c)	24 \pm 1.3 (c)	31 \pm 1.3 (b)	49 \pm 2.7 (a)
N	8	36	36	36	30
Chl a (mg m ⁻³)					
0.2 μ M	0.07 \pm 0.01 (c)	0.38 \pm 0.03 (b)	0.43 \pm 0.04 (b)	0.56 \pm 0.03 (ab)	0.65 \pm 0.04 (a)
2.0 μ M	0.019 \pm 0.004 (c)	0.38 \pm 0.02 (b)	0.26 \pm 0.02 (b)	0.40 \pm 0.03 (b)	0.90 \pm 0.09 (a)
Ratio 0.2:2.0	4.43 \pm 0.47 (a)	1.07 \pm 0.07 (b)	1.83 \pm 0.13 (b)	1.60 \pm 0.08 (b)	0.94 \pm 0.07 (c)
GF/F	0.094 \pm 0.012 (c)	0.65 \pm 0.03 (b)	0.60 \pm 0.04 (b)	0.82 \pm 0.04 (b)	1.4 \pm 0.01 (a)
N	7	71	71	71	59
Plankton					
Heterotrophic bacteria					
ml ⁻¹ ($\times 10^6$)	1.30 \pm 0.20(c)	1.58 \pm 0.61(c)	2.45 \pm 0.74 (b)	3.13 \pm 0.97 (a)	3.35 \pm 1.18 (a)
<i>Prochlorococcus</i> ml ⁻¹ ($\times 10^3$)	254 \pm 115 (a)	3.80 \pm 2.60 (b)	0	0	0
<i>Synechococcus</i> ml ⁻¹ ($\times 10^5$)	0.08 \pm 0.05 (d)	1.53 \pm 1.30 (c)	2.98 \pm 1.03 (b)	3.49 \pm 1.13 (a,b)	3.64 \pm 1.41 (a)
Picoeukaryotes ml ⁻¹ ($\times 10^4$)	0.22 \pm 0.10 (d)	1.17 \pm 0.50 (c)	1.81 \pm 0.85 (c)	2.86 \pm 1.31(b)	3.20 \pm 1.60 (a)
Nanoeukaryotes ml ⁻¹	24 \pm 5 (c)	265 \pm 158 (b)	264 \pm 199 (b)	325 \pm 294 (b)	615 \pm 368 (a)
N	15	96	96	96	96
Pennate diatoms l ⁻¹					
size (μ m)	142 \pm 103 (b)	3840 \pm 2550 (a)	1040 \pm 943 (b)	2040 \pm 1580 (a,b)	3650 \pm 3560 (a)
<i>Chaetoceros</i> l ⁻¹	34 \pm 23 (b)	54 \pm 32 (a,b)	49 \pm 10 (a,b)	77 \pm 47 (a)	75 \pm 37 (a)
size (μ m)	0	602 \pm 2580 (a)	105 \pm 219 (a)	363 \pm 419(a)	879 \pm 2440 (a)
Dinoflagellates l ⁻¹	nd	10 \pm 6 (a)	12 \pm 7(a)	11 \pm 9 (a)	15 \pm 12 (a)
size (μ m)	2410 \pm 1070 (a)	2620 \pm 2020 (a)	2330 \pm 1850 (a)	2930 \pm 2360 (a)	2180 \pm 1190 (a)
Ciliates l ⁻¹	16 \pm 7 (a)	21 \pm 9 (a)	23 \pm 10 (a)	22 \pm 8 (a)	22 \pm 9 (a)
size (μ m)	818 \pm 445 (a)	636 \pm 586 (a)	418 \pm 252 (a)	609 \pm 691(a)	619 \pm 667 (a)
N	26 \pm 7(a)	21 \pm 8 (a)	28 \pm 11(a)	25 \pm 13 (a)	26 \pm 15 (a)
N	10	30	30	30	30
Zooplankton (mg m ⁻³)					
Large >183 μ m		11.0 \pm 5.83 (c)	14.7 \pm 6.00 (b)	19.8 \pm 5.41(a)	
		32	32	32	
Medium <183>63 μ m		16.4 \pm 7.77 (c)	21.2 \pm 10.0 (b)	32.6 \pm 8.20 (a)	
Small <63 μ m		16.9 \pm 5.32 (a)	10.6 \pm 3.00 (b)	12.9 \pm 3.39 (b)	
N		39	39	39	

at MP > NB = SB > CB = OCN. The other inorganic nutrients measured (phosphate, TDP, ammonia) were not significantly different among sites (Table 2). Chl *a* also showed some among-site differences: for the 0.2 μm fraction, MP = SB > CB = NB > OCN; for the 2.0 μm fraction, MP > SB = NB = CB > OCN; and for the GF/F fraction, MP > SB = NB = CB > OCN (Table 2). The ratio between the small and large chl *a* fractions also showed significant differences with OCN > CB = SB = NB > MP (Table 2).

Significant differences in picoplankton abundances were found among sites (Table 2). Mean (\pm SE) cell size was $0.5 \pm 0.3 \mu\text{m}$ for heterotrophic bacteria, $1.2 \pm 0.6 \mu\text{m}$ for *Synechococcus*, $1.4 \pm 0.3 \mu\text{m}$ for picoeukaryotes, and $3.3 \pm 0.7 \mu\text{m}$ for nanoeukaryotes. A general trend was observed with an increase in cell abundance from OCN < NB < CB < SB < MP (Table 2b). The oceanic site had the lowest concentrations of all cell types except *Prochlorococcus*. *Prochlorococcus* was the dominant picoplankton at the oceanic site (2.54×10^5 cells ml^{-1}) but was only detectable within Kāne'ohē Bay at the NB site (3.8×10^3 cells ml^{-1}). Oceanic values for *Synechococcus* were significantly different from the Bay sites, even though these sites are connected by the channels and water flowing over the barrier reef. Within Kāne'ohē Bay, NB had the lowest abundance of most cell types and MP the highest (Table 2). Microplankton also showed significant spatial differences in abundances in pennate diatoms but not in other groups (Table 2). There were no significant spatial differences in cell size for microplankton, although there was a general trend of smallest size at OCN site and largest size at SB and MP (Table 2).

ANOVA indicated significant differences among the 3 sites sampled for zooplankton biomass for the large and medium fractions, with SB > CB > NB (Table 2). The small fraction was greatest at NB, while CB and SB were not significantly different from each other (Table 2).

Carbon and nitrogen content at different sites

Particulate carbon and nitrogen content of the pico-, nano-, and microplankton at the different sites was calculated from the number and size of cells using conversion factors (Table 1). Within Kāne'ohē Bay and oceanic sites, there was an increase in microbial carbon and nitrogen from OCN < NB < CB < SB < MP. At the 3 sites where all plankton fractions (from pico- to zooplankton) were measured (NB, CB, SB), picoplankton was the dominant fraction in terms of carbon (79 to 88%) and nitrogen (76 to 83%, Fig. 2). At all sites, picoplankton populations represented more than 90% of total microbial carbon and nitrogen. Picoplankton

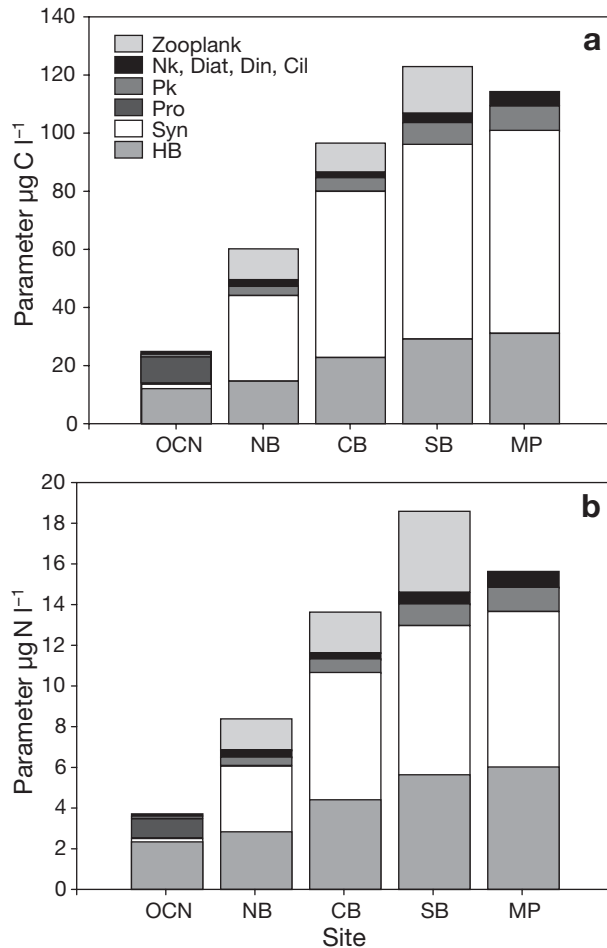


Fig. 2. (a) Carbon and (b) nitrogen content of plankton. Values for microbial plankton derived by cell conversion factors (see 'Materials and methods'), for zooplankton from CHN analysis of dried samples (NB: zooplankton not sampled at OCN or MP sites). Zooplank: zooplankton; Nk: nanoeukaryotes; Diat: diatoms; Din: dinoflagellates; Cil: ciliates; Pk: picoeukaryotes; Pro: *Prochlorococcus* spp.; Syn: *Synechococcus* spp.; HB: heterotrophic bacteria. Site abbreviations as in Fig. 1

carbon and nitrogen at the oceanic site was mainly in heterotrophic bacteria (more than 50% of carbon and nitrogen), with *Prochlorococcus* cells contributing 40% of the picoplankton carbon and 28% of the nitrogen. The dominant component within Kāne'ohē Bay was *Synechococcus*, accounting on average for more than 50% of the picoplankton carbon and nitrogen (Fig. 2).

The particulate carbon and nitrogen content of the water column as estimated from cell count-size-conversion factors was compared to the carbon and nitrogen measured from water column samples (see 'Materials and methods'), where total carbon included both inorganic and organic particulate carbon. An estimation of organic particulate carbon was obtained by acidification treatment to remove carbonates. Carbon values from the water column samples (total and

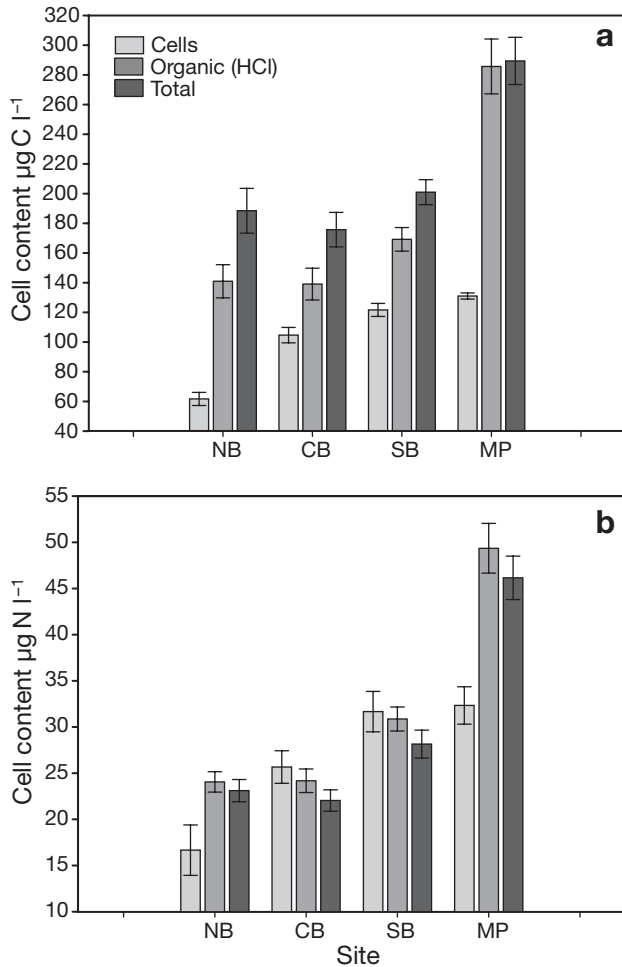


Fig. 3. Comparison of (a) particulate carbon and (b) particulate nitrogen as estimated by the cell conversion method (light gray bars) and water column samples (black bars: samples rinsed with deionized water only; dark gray bars: samples rinsed with hydrochloric acid). Site abbreviations as in Fig. 1

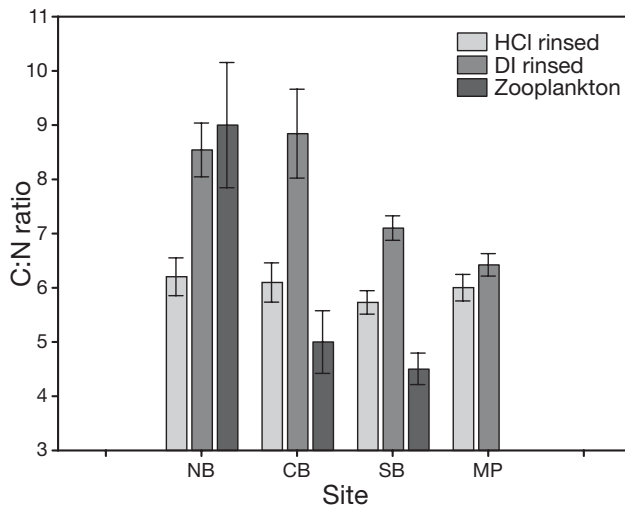


Fig. 4. C:N ratios for water column and zooplankton samples. Site abbreviations as in Fig. 1

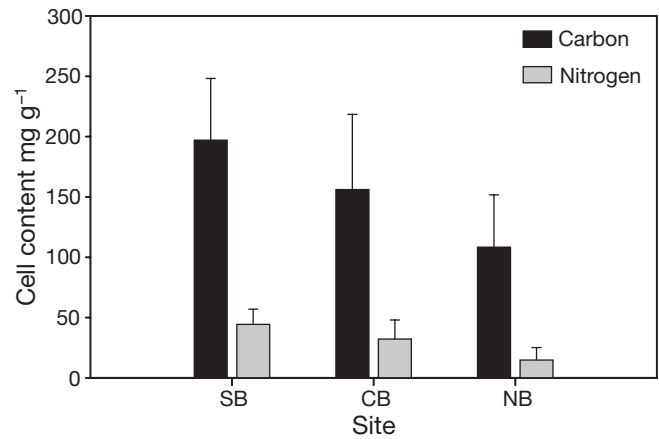


Fig. 5. Zooplankton carbon and nitrogen normalized to zooplankton dry mass. Site abbreviations as in Fig. 1

organic) were significantly higher than the cell count-size-conversion factor method (Fig. 3a). There were no significant differences between total nitrogen and nitrogen after acidification at any site. Nitrogen values calculated by analysis of particulates in filtered water and count-size-conversion factors were the same at the CB and SB sites, but the NB and MP sites showed higher values in water column samples (Fig. 3b), which suggests that detritus represented about 30% of particulate nitrogen in these samples, reflecting the proximity of streams as was also seen in the lower salinity and higher TSS at these sites (Table 2). C:N ratios were higher for non-decalcified samples with values ranging from 6.4 (MP) to 8.8 (CB). When samples were rinsed with HCl, the C:N ratios were close to 6 at all sites (Fig. 4), similar to marine plankton ratios (Atkinson & Smith 1983); this suggests that detritus was probably not from benthic macroalgae, in which case C:N ratios would have been much higher.

Zooplankton samples showed that carbon and nitrogen values ranged from 10 to 15 $\mu\text{C l}^{-1}$ and from 2 to 4 $\mu\text{N l}^{-1}$ (Fig. 2), which represented about 10 to 17% of the carbon and 15 to 22% of the nitrogen at NB, CB and SB where all fractions were analyzed. When zooplankton carbon and nitrogen content were normalized to zooplankton dry mass, there were significant differences in both carbon and nitrogen content, with $\text{SB} > \text{CB} > \text{NB}$ (Fig. 5).

Temporal differences

There were no obvious seasonal patterns in nutrient levels throughout the 3 yr CISNet sampling program at any of the 4 main study sites (www.hawaii.edu/cisnet). This project took place during a drought on O'ahu, with stream discharge substantially below long-term

mean daily discharge (for example, US Geological Survey data for Kāne'ohe Stream: 1976–2001, mean daily discharge $0.286 \text{ m}^3 \text{ s}^{-1}$; CISNet study: 1998–2001, mean daily discharge $0.142 \text{ m}^3 \text{ s}^{-1}$).

However, if the CISNet data were simply grouped into dry and wet seasons (dry: June to September; wet: October to May), we did find some significant differences in nutrient concentrations within Kāne'ohe Bay. Phosphate and TDP were higher during the wet season

(Table 3). Significantly higher nitrate concentrations during the wet season were only found at the NB site. In contrast, higher silicate concentrations during the dry season were found at the CB and SB sites (Table 3). The $0.2 \mu\text{m}$ chl *a* fraction was significantly higher in the dry season at the NB, CB and SB sites, but all other chl *a* fractions and the ratio of small to large chl *a* showed no significant dry versus wet season differences (Table 3). The microbial groups also showed significant differ-

Table 3. Comparison of water column and plankton parameters (mean \pm SE) during dry and wet seasons in Kāne'ohe Bay. Site abbreviations as in Fig. 1. Values in **bold** are significantly higher than those of the alternate season for that site. Site abbreviations as in Fig. 1

	NB		CB		SB		MP	
	dry	wet	dry	wet	dry	wet	dry	wet
Water column								
Salinity (PSU)	34.9 \pm 0.05	34.7 \pm 0.06	35.2 \pm 0.07	35.0 \pm 0.05	35.4 \pm 0.03	35.0 \pm 0.06	34.4 \pm 0.31	34.2 \pm 0.26
N	23	48	23	48	23	48	23	36
Secchi (m)	6.4 \pm 0.3	5.4 \pm 0.4	7.0 \pm 0.3	5.7 \pm 0.3	5.7 \pm 0.3	4.8 \pm 0.2	3.5 \pm 0.6	3.0 \pm 0.2
N	21	47	21	47	21	47	21	35
TSS (mg l^{-1})	2.7 \pm 0.3	2.7 \pm 0.4	1.9 \pm 0.2	1.9 \pm 0.2	2.0 \pm 0.1	2.1 \pm 0.2	3.1 \pm 0.2	4.0 \pm 0.4
N	21	48	21	48	21	48	21	36
Phosphate (μM)	0.045 \pm 0.007	0.082 \pm 0.006	0.039 \pm 0.005	0.076 \pm 0.006	0.059 \pm 0.005	0.083 \pm 0.004	0.053 \pm 0.005	0.083 \pm 0.005
Nitrate (μM)	0.14 \pm 0.03	0.34 \pm 0.05	0.07 \pm 0.03	0.10 \pm 0.04	0.04 \pm 0.01	0.06 \pm 0.02	0.13 \pm 0.09	0.23 \pm 0.12
Ammonia (μM)	0.12 \pm 0.02	0.15 \pm 0.02	0.15 \pm 0.05	0.13 \pm 0.02	0.15 \pm 0.03	0.11 \pm 0.01	0.12 \pm 0.02	0.09 \pm 0.01
Silicate (μM)	7.6 \pm 0.27	6.8 \pm 0.25	6.0 \pm 0.55	4.5 \pm 0.20	7.9 \pm 0.24	6.5 \pm 0.27	20 \pm 2.7	15 \pm 1.9
N	23	49	23	49	23	49	23	36
TDP (μM)	0.25 \pm 0.02	0.32 \pm 0.01	0.27 \pm 0.02	0.32 \pm 0.01	0.29 \pm 0.01	0.34 \pm 0.01	0.29 \pm 0.01	0.34 \pm 0.02
TDN (μM)	7.1 \pm 0.40	6.9 \pm 0.03	7.5 \pm 0.05	6.9 \pm 0.24	7.7 \pm 0.47	7.2 \pm 0.425	8.6 \pm 0.50	8.4 \pm 0.50
N	14	25	14	25	14	25	14	17
POC (mg m^{-3})	170 \pm 26	130 \pm 9	170 \pm 22	120 \pm 1	200 \pm 17	150 \pm 5	330 \pm 33	250 \pm 16
PON (mg m^{-3})	27 \pm 2	22 \pm 1	29 \pm 2	22 \pm 1	37 \pm 2	27 \pm 1	56 \pm 4	45 \pm 3
N	13	23	13	23	13	23	13	23
Chl <i>a</i> (mg m^{-3})								
0.2 μM	0.52 \pm 0.05	0.31 \pm 0.06	0.49 \pm 0.07	0.40 \pm 0.05	0.72 \pm 0.14	0.48 \pm 0.04	0.70 \pm 0.08	0.62 \pm 0.04
2.0 μM	0.44 \pm 0.04	0.36 \pm 0.03	0.26 \pm 0.03	0.26 \pm 0.02	0.44 \pm 0.06	0.38 \pm 0.04	0.94 \pm 0.12	0.87 \pm 0.12
0.2:2.0	1.3 \pm 0.09	0.98 \pm 0.09	2.0 \pm 0.13	1.8 \pm 0.18	1.6 \pm 0.11	1.6 \pm 0.11	0.81 \pm 0.07	1.0 \pm 0.10
GF/F	0.82 \pm 0.04	0.57 \pm 0.04	0.60 \pm 0.04	0.59 \pm 0.05	0.91 \pm 0.06	0.78 \pm 0.05	1.5 \pm 0.17	1.3 \pm 0.13
N	23	48	23	48	23	48	23	36
Plankton								
Heterotrophic bacteria ml^{-1} ($\times 10^6$)	1.83 \pm 0.10	1.44 \pm 0.80	2.97 \pm 0.19	2.20 \pm 0.40	3.70 \pm 0.20	2.80 \pm 0.10	3.80 \pm 0.30	3.10 \pm 0.11
<i>Synechococcus</i> ml^{-1} ($\times 10^5$)	2.49 \pm 0.33	1.03 \pm 0.64	3.47 \pm 0.21	2.72 \pm 0.12	3.94 \pm 0.27	3.26 \pm 0.13	4.53 \pm 0.30	3.17 \pm 0.16
Picoeukaryotes ml^{-1} ($\times 10^4$)	1.44 \pm 0.80	1.02 \pm 0.60	2.22 \pm 0.19	1.59 \pm 0.90	3.51 \pm 0.30	2.53 \pm 0.15	4.11 \pm 0.40	2.73 \pm 0.18
Nanoeukaryotes ml^{-1}	366 \pm 30	218 \pm 20	216 \pm 27	288 \pm 32	339 \pm 73	318 \pm 37	640 \pm 81	603 \pm 51
N	36	60	36	60	36	60	36	60
Pennate diatoms l^{-1}	4920 \pm 1290	3360 \pm 353	1590 \pm 642	476 \pm 133	1690 \pm 279	2330 \pm 367	4320 \pm 2010	3260 \pm 772
<i>Chaetoceros</i> l^{-1}	137 \pm 46	812 \pm 695	0	210 \pm 170	317 \pm 56	398 \pm 106	760 \pm 614	1370 \pm 1190
Dinoflagellates l^{-1}	2950 \pm 888	2480 \pm 384	3290 \pm 1210	1360 \pm 673	4190 \pm 655	1990 \pm 292	3250 \pm 450	1630 \pm 318
Ciliates l^{-1}	528 \pm 206	685 \pm 109	397 \pm 144	440 \pm 178	456 \pm 80	723 \pm 175	554 \pm 242	562 \pm 188
N	10	20	10	20	10	20	10	20
Zooplankton (mg m^{-3})								
Large >183 μm	11.5 \pm 1.63	10.7 \pm 1.36	15.0 \pm 1.60	14.4 \pm 1.45	16.8 \pm 1.51	22.2 \pm 0.93		
N	14	18	14	18	14	18		
Medium								
<183 >63 μm	19.6 \pm 2.17	14.6 \pm 1.42	27.0 \pm 2.86	18.1 \pm 1.66	34.8 \pm 2.41	31.4 \pm 1.52		
Small <63 μm	17.2 \pm 1.36	16.7 \pm 1.11	11.0 \pm 0.56	10.4 \pm 0.69	13.2 \pm 0.74	12.7 \pm 0.75		
N	14	25	14	25	14	25		

ences, with higher concentrations of heterotrophic bacteria, *Synechococcus*, and picoeukaryotes during the dry season (Table 3). For larger cells, the only significant dry versus wet season differences occurred in dinoflagellate abundances at the SB and MP sites (Table 3b). Zooplankton showed higher biomass (AFDM) during the dry season in the large fraction at SB, and in the medium fraction at CB (Table 3).

General pulse features

All pulses of rainfall and increased discharge measured during this period resulted in a plume of brown, turbid water in southern Kāneʻohe Bay. The MP site was within this brown plume during all pulses, as evidenced by dramatically-decreased surface salinity and increased TSS at the beginning of each pulse (Table 4). The duration and extent of the brown plume depended on the amount and timing of increased discharge as well as additional factors, such as wind speed/direction and strength of tidal flushing. All pulses resulted in increased nitrate and TDN, correlated with lowered salinity (Fig. 6: nitrate $r^2 = 0.79$, data from all pulses combined; TDN $r^2 = 0.83$). Likewise, phosphate ($r^2 = 0.50$), TDP ($r^2 = 0.49$), and silicates ($r^2 = 0.83$) also showed an increase following pulses, correlated with decreased salinity (Fig. 6). Chl *a* typically peaked in concentration 3 to 7 d following the initial nutrient inputs. The same pattern was also observed in microphytoplankton groups, specifically in the *Chaetoceros* genus, but not in picoplankton. Picoplanktonic cell abundance did not show a large increase, except following the dry season pulse (June 2001). However, fluorescence for *Synechococcus* (dominant picophytoplankton group) showed an increase 12 to 24 h after rain discharge (Fig. 7), suggesting an increase in pigments per cell.

Individual pulse histories

January 2000 pulse

Heavy rains occurred during the evening of January 19, resulting in over 80 mm of rainfall in 24 h as recorded at the Luluku rain gauge. Discharge from Kāneʻohe Stream increased to $1.84 \text{ m}^3 \text{ s}^{-1}$ (Fig. 8). Surface salinity at the MP site on January 20 dropped to 12 PSU, with salinity at 0.5 m at 27 PSU. Nitrate con-

Table 4. Maximum (or minimum) values for water column parameters during stream pulses. Pulses ranked by maximum discharge. MP: mid plume; OP: outside plume; nd: no data

	Jan 00	Oct 00	Jun 01	Apr 00	Feb 01
MP					
Max. discharge ($\text{m}^3 \text{ s}^{-1}$)	1.841	1.303	0.651	0.566	0.566
Min. salinity (PSU)	12	27	25	19	23
Max. TSS (mg l^{-1})	17	6.4	5.9	12	12
Max. nitrate (μM)	16.5	5.22	2.95	9.85	15.1
Max. TN (μM)	29.0	16.4	16.2	22.6	25.0
Max. phosphate (μM)	0.57	0.44	0.39	0.39	0.61
Max. TP (μM)	0.80	0.71	0.65	0.69	0.85
Max. silicate (μM)	121	54	70	83	86
Max. chl <i>a</i> ($\mu\text{g l}^{-1}$)	4.86	1.87	2.09	5.94	2.52
OP					
Min. salinity (PSU)	28.9	nd	35.3	32.4	34.3
Max. TSS (mg l^{-1})	3.0	nd	2.4	3.8	1.7
Max. nitrate (μM)	6.6	nd	0.22	0.92	0.37
Max. TN (μM)	15.4	nd	9.99	9.21	10.8
Max. phosphate (μM)	0.15	nd	0.09	0.13	0.15
Max. TP (μM)	0.43	nd	0.27	0.45	0.43
Max. silicate (μM)	30	nd	13	75	11
Max. chl <i>a</i> ($\mu\text{g l}^{-1}$)	4.18	nd	1.33	1.57	1.02

centrations on the morning of January 20 were $16 \mu\text{M}$ (87 times baseline), phosphate increased to $0.57 \mu\text{M}$ (8 times baseline), and silicate increased to $121 \mu\text{M}$ and remained elevated for 5 days (Fig. 8). Chl *a* peaked at $4.9 \mu\text{g l}^{-1}$, 3 days following the pulse (Fig. 8). Nitrate and TDN concentrations increased at the OP site, but only following the second rain and discharge event (January 27, Fig. 9). Chl *a* at the OP site had increased 3-fold by 5 days after the first discharge pulse. No microbial samples were collected during this pulse.

April 2000 pulse

Rainfall on March 31 and April 1 (59 mm in 24 h) increased stream discharge to $0.57 \text{ m}^3 \text{ s}^{-1}$ (Fig. 8). Surface salinity at MP on March 31 dropped to 19 PSU, while at 0.5 m depth it was 34 PSU. By April 1, salinity was 21 PSU at 0.5 m depth; by April 3, surface salinity had returned to approximately 31 PSU. Following the initial pulse, nitrate concentration at MP increased to $9.8 \mu\text{M}$ (52 times baseline) and phosphate to $0.39 \mu\text{M}$ (6 times baseline) (Fig. 8). Silicate remained elevated for 6 days following the initial discharge pulse. Chl *a* lagged the nutrient increase, peaking at $5.9 \mu\text{g l}^{-1}$ 6 days after the pulse (Fig. 8). There were small increases in nitrate (to $0.92 \mu\text{M}$) and phosphate (to $0.13 \mu\text{M}$) at the OP site (Fig. 9 & Table 4).

A second period of rainfall on April 9 (26 mm) elevated stream discharge to $0.24 \text{ m}^3 \text{ s}^{-1}$. This was accompanied by a small decrease in surface salinity to

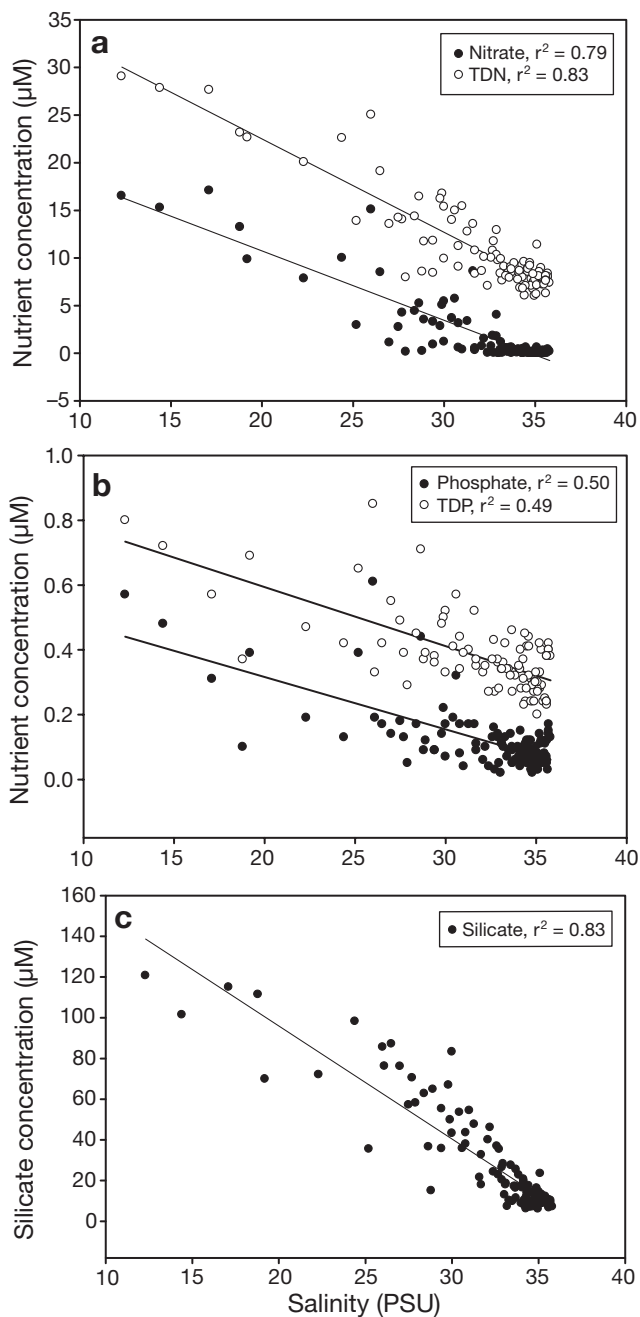


Fig. 6. Relationship between salinity and water column parameters (a) nitrogen, (b) phosphorus and (c) silicate during 1.5 yr time-period of pulse sampling. TDN: total dissolved nitrogen; TDP: total dissolved phosphorus

28 PSU that persisted for several days. Increases in nitrate (to $5.4 \mu\text{M}$) and phosphate (to $0.18 \mu\text{M}$) lagged the second discharge pulse by 3 d. At the OP site there was a small increase in nitrate 6 days after the secondary discharge pulse (Fig. 9).

Microbial sampling began on April 3, 2 days after the initial stream pulse. The initial low abundance of heterotrophic bacteria and *Synechococcus* may have been due

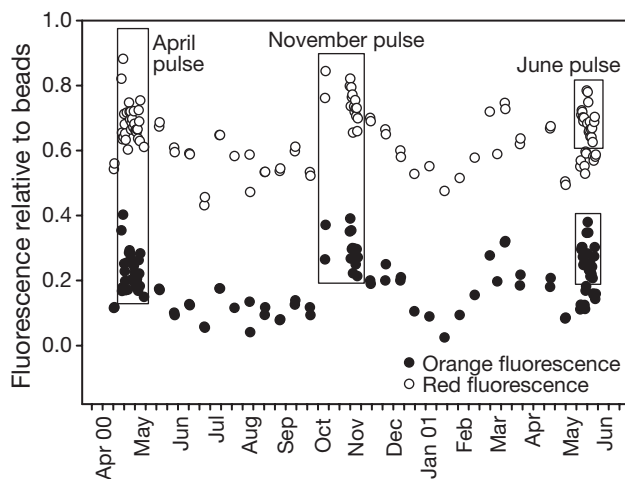


Fig. 7. *Synechococcus* spp. Average autofluorescence signals relative to beads fluorescence by samples run in flowcytometer. Orange fluorescence: chlorophyll; red fluorescence: phycoerythrin. Pulse data is boxed

to flushing and dilution. Abundance also dropped following the second stream discharge on April 9 (Fig. 10). In contrast, nanoeukaryotes at the MP site were elevated above baseline during this pulse (Fig. 10). Counts for microplankton showed a lagged increase in abundance following the stream discharge pulse. The genus *Chaetoceros* responded the most dramatically, with a very large increase in cell number 5 days following the initial pulse (Fig. 11). Pennate diatoms, dinoflagellates, and ciliates also showed a lagged response (5 to 10 d) following the initial stream pulse and showed their highest abundances following the second discharge pulse (Fig. 11). Microbial cell counts at OP generally overlapped with cell counts at the MP site (Fig. 10).

October 2000 pulse

Following 2 days of rain on October 29 and 30 (total 71 mm), discharge increased to $0.62 \text{ m}^3 \text{ s}^{-1}$ (Fig. 8). Sub-surface salinity decreased to 29 PSU, and nitrate increased to $5.2 \mu\text{M}$ (28 times baseline) and phosphate to $0.44 \mu\text{M}$ (6 times baseline) (Fig. 8). Silicate rose to $37 \mu\text{M}$, and remained elevated for most of the sampling period. Chl *a* peaked at $1.9 \mu\text{g l}^{-1}$, 3 days following the initial pulse.

An additional 56 mm of rain fell on November 3, increasing stream discharge to $1.3 \text{ m}^3 \text{ s}^{-1}$. Although sampling commenced on November 3, the drop in salinity and increase in nitrate (19 times baseline) and phosphate (3 times baseline) were not recorded at the MP site until November 7 (Fig. 8). The second discharge was 79% of the rain discharged during the initial 2 d pulse, and although the nitrate increase was 70% of the initial pulse of nitrate, the phosphate was only 43%

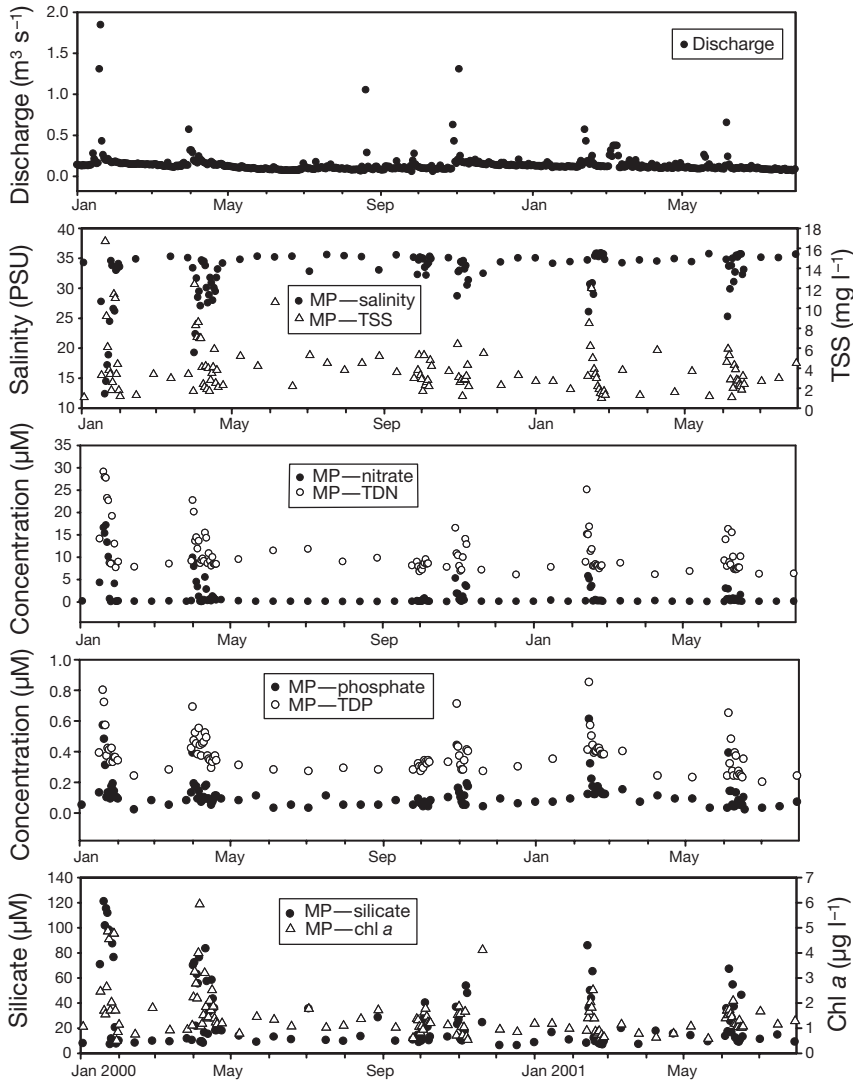


Fig. 8. Water column parameters at mid plume (MP) site during pulse-sampling period. Discharge at Kāne'ohē Stream gauge (US Geological Service data) indicates the pulses followed (NB: August 2000 pulse was not sampled)

of the initial phosphate discharged during the earlier pulse. Only pico- and nanoplankton were sampled from October 31 to November 8. These plankton groups did not show strong responses to the pulse (Fig. 10), which were within the range of baseline values.

February 2001 pulse

February 12 was a regular CISNet sampling day. On February 13, 63 mm of rainfall were recorded at the Luluku gauge, associated with an increase in stream discharge to $0.57 \text{ m}^3 \text{ s}^{-1}$ (Fig. 8). Subsurface salinity dropped from 34 PSU on February 12 to 26 PSU on February 13, and gradually returned to 34 PSU by

February 18. Nitrate jumped from $0.14 \text{ } \mu\text{M}$ during CISNet sampling (February 12) at MP to $15 \text{ } \mu\text{M}$ on the following day. Likewise, phosphate increased from $0.12 \text{ } \mu\text{M}$ during the CISNet sampling to $0.61 \text{ } \mu\text{M}$. Silicate increased from $8 \text{ } \mu\text{M}$ to $86 \text{ } \mu\text{M}$. Chl *a* peaked at $2.5 \text{ } \mu\text{g l}^{-1}$ approximately 5 days following the nutrient peak, but returned quickly to baseline levels (Fig. 8). At the OP site, there was a slight increase in nitrate following the discharge but no significant changes in phosphate, silicate or chl *a* (Fig. 9). No microbial data were collected during this pulse.

June 2001 pulse

June 4, a regular CISNet sampling date, was followed by 2 days of rain, producing a 2 d sum of 51 mm at the Luluku rain gauge. Stream discharge peaked briefly at $0.61 \text{ m}^3 \text{ s}^{-1}$ but rapidly returned to baseline levels (Fig. 8). Subsurface salinity dropped to 25 PSU, but returned to 34 PSU by June 8. Nitrate peaked at $2.95 \text{ } \mu\text{M}$ (16 times baseline) and phosphate peaked at $0.39 \text{ } \mu\text{M}$ (6 times baseline) (Fig. 8). Silicate peaked at $67 \text{ } \mu\text{M}$ on June 7. Chl *a* peaked at $2.1 \text{ } \mu\text{g l}^{-1}$, 6 days following the start of the event. There were no major changes in nitrate, phosphate, silicate or chl *a* at the OP site (Fig. 9).

There were several smaller rain events following the initial pulse, leading to a small increase in stream discharge on June 17, accompanied by a decrease in subsurface salinity to 32 PSU and an increase in nitrate ($1.52 \text{ } \mu\text{M}$), phosphate ($0.10 \text{ } \mu\text{M}$) and silicate ($46 \text{ } \mu\text{M}$).

Picoplankton concentrations were above baseline values following the discharge pulse (Fig. 10). Diatoms and dinoflagellates showed only small increases above baseline values; however, ciliate concentrations reached 12 times the overall mean 5 days after the peak discharge (Fig. 11).

DISCUSSION

Spatial differences

Kāne'ohē Bay receives 2 main sources of allochthonous water, freshwater inputs and oceanic water, that flush the Bay. The effect of runoff from the main streams of Kāne'ohē represents an input of new nutri-

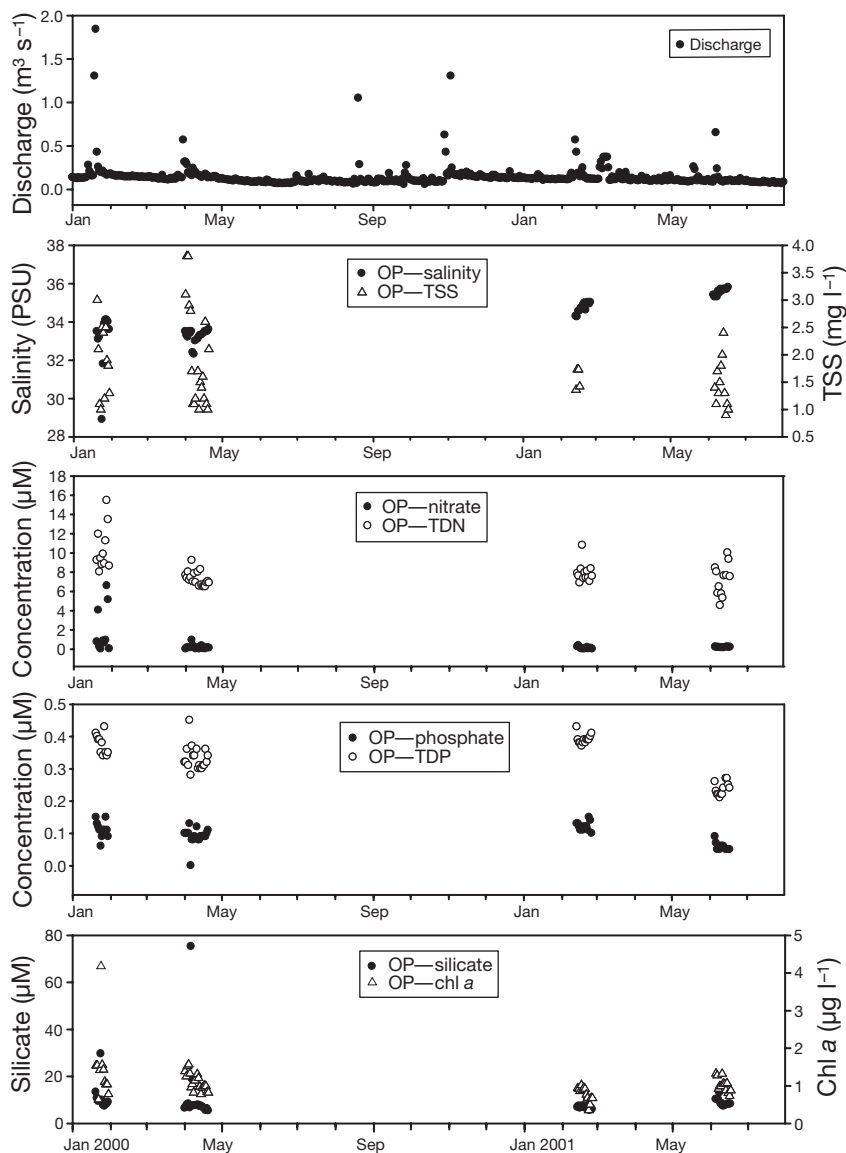


Fig. 9. Water column parameters at outside plume (OP) site during pulse-sampling period. Discharge at Kāne'ohe Stream gauge (US Geological Service data) indicates the pulses followed (NB: August, October and November 2000 pulses were not sampled)

ents to the Bay and, in contrast, the flushing with oceanic water dilutes the Bay water. However, average nutrient concentrations within Kāne'ohe Bay were (with some exceptions) not different from the oceanic values measured in this study. In general, nutrient concentrations within the Bay were low, possibly due to the drought that coincided with our sampling period. The exceptions to the low water column nutrient levels within the Bay were nitrate at the NB site, silicate at the MP, SB, and NB sites, and TDN at the MP, SB and CB sites. NB receives the greatest flushing with oceanic water; however, this site still had the highest nitrate levels, demonstrating the importance of input from

Waikāne and Wai'hole streams emptying into the Bay near the NB site (Fig. 1). Higher silicate concentrations also reflected freshwater inputs. Increased nitrogen may also be associated with the efficient trapping, consumption and remineralization of particulate organic matter by reef biota (Ribes et al. 2005), decomposition in the pore space of the sand and reef framework (Rasheed et al. 2002), or nitrogen fixation by organisms living in the reef environment (Capone 1983, D'Elia & Wiebe 1990, Williams & Carpenter 1998).

All plankton cell types and total plankton biomass (in carbon and nitrogen units) gradually increased in the order OCN < NB < CB < SB < MP. There was a clear dominance of picoplankton in terms of cell concentrations and biomass, typical of oligotrophic systems such as coral reefs. Picoplankton have been previously reported to account for 40 to 45 % of the biomass of phytoplankton in Kāne'ohe Bay (Taguchi & Laws 1989, Laws & Allen 1996). Our sampling period occurred during a drought, and picoplankton constituted more than 90 % of the phytoplankton community, reflecting the dominance of this size class during low nutrient conditions. The zooplankton fraction was about 10 to 20 % of the total plankton biomass, reflecting a planktonic community composition dominated by components < 20 µm, typical of oligotrophic systems such as coral reefs. Dominance of small-sized prokaryotic phytoplankton in terms of biomass and primary production has been also reported for atoll lagoons (Charpy 1996, Charpy & Blanchot 1998) and tropical coastal ecosystems adjacent to high islands (Tada et al. 2003, Agawin et al. 2003).

The switch in dominance of the cyanobacteria from *Prochlorococcus* at the oceanic site to *Synechococcus* in Kāne'ohe Bay was the most characteristic feature of plankton distribution in our study. A similar dominance of *Synechococcus* in atoll lagoons and *Prochlorococcus* in the surrounding ocean waters has been reported on French Polynesian reefs (Charpy & Blanchot 1998). Nitrogen availability and source and light intensity and spectral quality have been discussed as the main factors controlling marine cyanobacteria abundance (Kana & Glibert 1987, Moore et al. 1995, 2002, Ting et al. 2002). *Prochlorococcus* may be insensitive to changes in nutrient concentration, whereas *Synecho-*

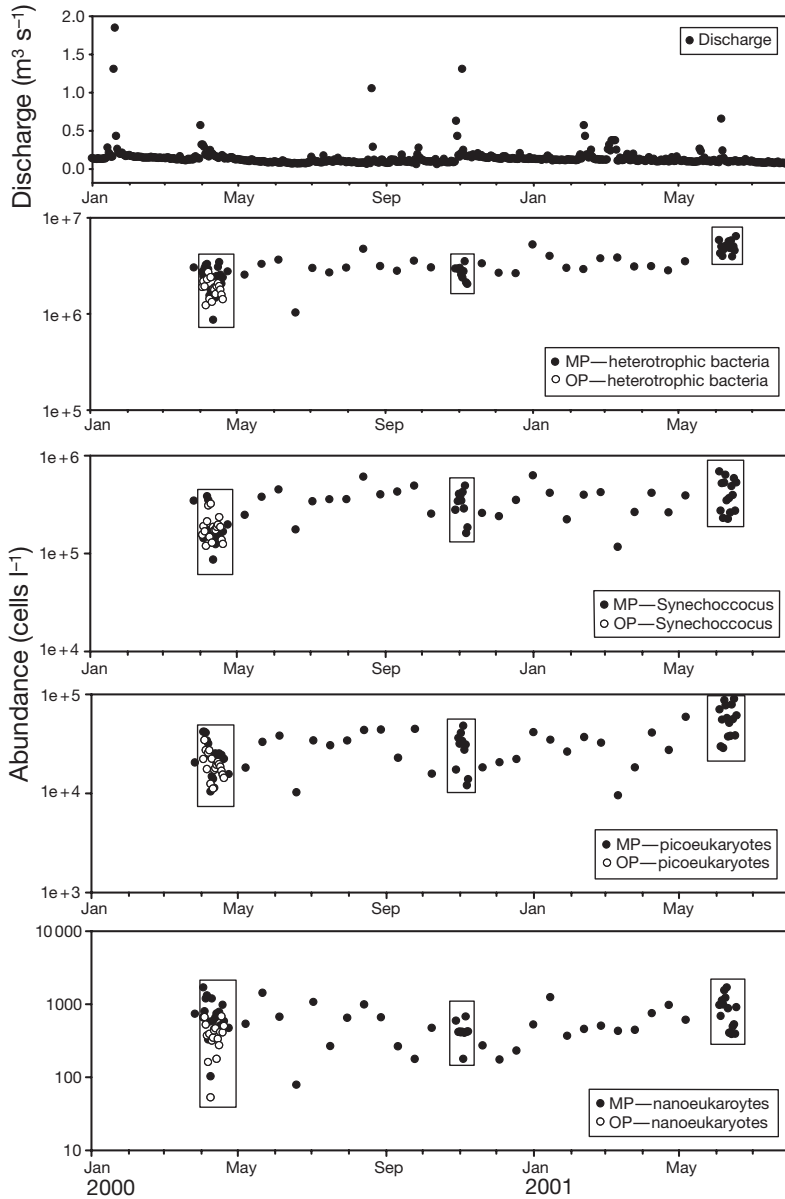


Fig. 10. Microbial fractions at mid plume (MP; ●) and outside plume (OP; ○) site during pulse-sampling period (NB: baseline data at MP were not collected until April 2000, and August 2000 pulse was not sampled)

coccus grows faster in higher nitrate concentrations (Partensky 1999a,b and references therein). Therefore, *Prochlorococcus* is expected to be the dominant cell type in oligotrophic conditions when nitrate $< 0.1 \mu\text{M}$ (Shalapyonok et al. 2001). Transitions to less oligotrophic conditions are typically associated with a decrease in *Prochlorococcus* abundance and an increase of *Synechococcus* and other phytoplankton groups. The oceanic site in our study appears to be permanently oligotrophic (mean nitrate concentration $\sim 0.05 \mu\text{M}$); hence, a dominance of *Prochlorococcus* would be expected. The *Prochlorococcus* abundances

we measured (mean of $2.54 \times 10^5 \text{ cells ml}^{-1}$) are within the range reported for surface oligotrophic waters (Campbell & Vaultot 1993, Campbell et al. 1997, Partensky et al. 1999 a,b, Berninger & Wickham 2005).

Spectral quality is another factor that could determine the differential distribution between *Synechococcus* and *Prochlorococcus*. *Prochlorococcus* is particularly efficient at absorbing blue wavelengths of light that dominate oceanic oligotrophic waters. Although *Synechococcus* does not absorb blue light as well as *Prochlorococcus*, it is capable of absorbing green wavelengths (500 to 550 nm) that predominate in coastal waters (Ting et al. 2002). Current research (E. Hochberg unpubl.) on spectral quality in Kāneʻohe Bay shows that within the Bay there is more attenuation in blue wavelengths (by phytoplankton as well as dissolved organic matter). Secchi depth and attenuation coefficients (Table 2) also demonstrate differences in the light regimes of the oceanic and Bay sites. Therefore, light availability and—in particular—wavelength composition could be factors contributing to the switch from *Prochlorococcus* to *Synechococcus* dominance from oceanic to Bay sites.

The fact that Kāneʻohe Bay waters were devoid of *Prochlorococcus* was previously pointed out in a short-term study by Liu et al. (1995). In our study, *Prochlorococcus* was only found at the NB site, even though it dominated the oceanic site. The presence of *Prochlorococcus* at the NB site may reflect advection into the Bay from oceanic waters, even if populations of *Prochlorococcus* were unable to grow under the environmental conditions found in the northern sector of the Bay. A similar phenomenon has been described for the advection of *Prochlorococcus* into Uchinumi Bay and subsequent failure of population growth (Katano et al. 2005).

Synechococcus at the oceanic site in our study showed higher cell concentrations ($7.8 \times 10^3 \text{ cells ml}^{-1}$) than mean surface values ($1.4 \times 10^3 \text{ cells ml}^{-1}$) reported for oligotrophic Station ALOHA, 100 km north of Oʻahu, suggesting the export of these cells from Kāneʻohe Bay. Within the Bay, *Synechococcus* showed a north to south increase in concentration reaching a maximum at the MP site. A positive relationship between cell abundance and nitrate concentrations has been reported for *Synechococcus* (Agawin et al. 2000,

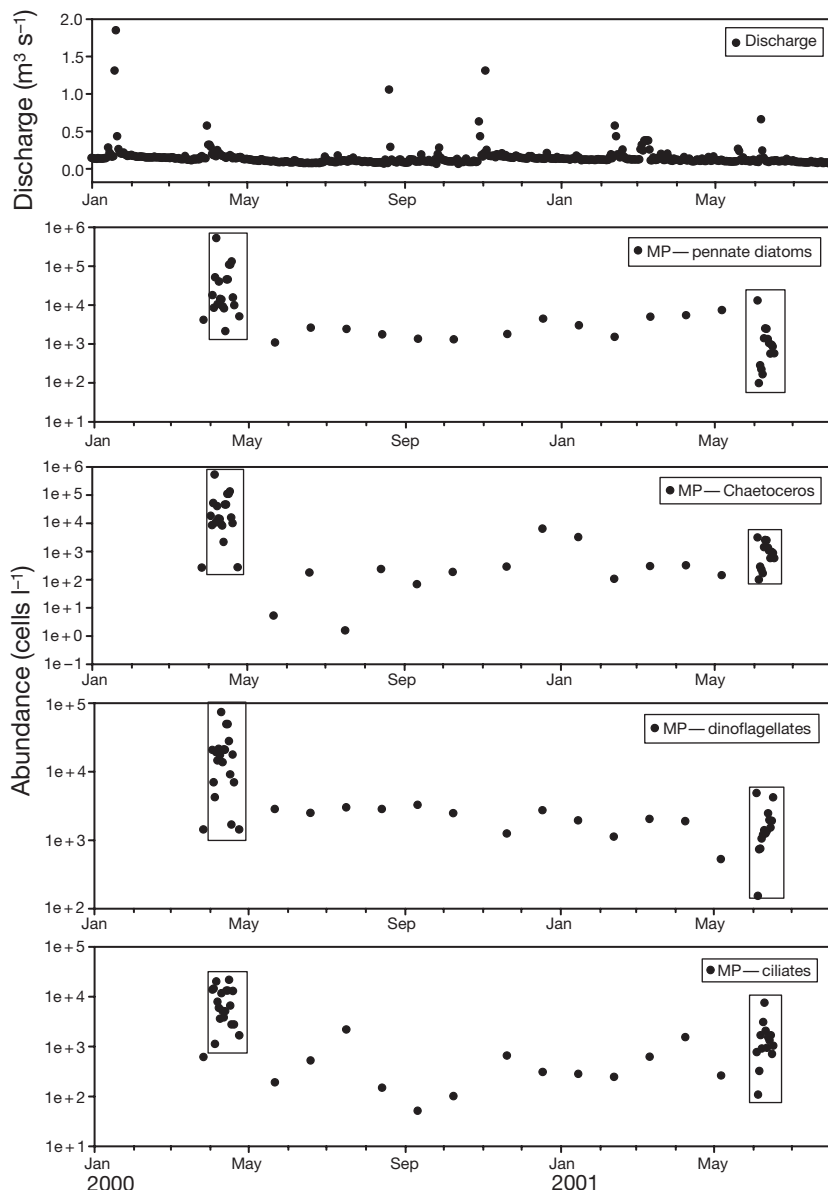


Fig. 11. Microplankton fractions at mid plume (MP) site. Pulse data is boxed (NB: baseline data at MP were not collected until April 2000, and August, October and November 2000 pulses were not sampled)

Shalapyonok et al. 2001). However, in Kāneʻohe Bay, nitrate concentration does not seem to explain the distribution and abundance of *Synechococcus*. Glibert et al. (2004) have shown that *Synechococcus* is capable of utilizing organic sources of nitrogen, and the increase in dissolved organic nitrogen in the order NB < CB < SB < MP suggests that organic sources of nitrogen may play a role in *Synechococcus* abundance in the Bay.

Overall, our data indicate that plankton biomass and composition at the study sites within Kāneʻohe Bay could also be influenced by water circulation patterns and residence time. Longer water residence time in more enclosed areas may permit acclimatization of microbial

communities to local conditions (in particular, light conditions), as well as retaining plankton cells. Longest water residence time, approximately 13 d, is found at the SB and MP sites (Sunn, Low, Tom, & Hara Inc. 1976, Smith et al. 1981), where plankton biomass was highest.

Other biotic factors can also affect plankton biomass and community structure. In coral reef systems many macroinvertebrates feed on picoplankton (Pile 1997, Ribes et al. 2003, 2005). The number and distribution of benthic picoplankton consumers such as sponges and ascidians determine the ability of reef systems to capture plankton in this size range (Ribes et al. 2005). Picoplankton consumers in the water column (i.e. appendicularians, Scheinberg 2004, Scheinberg et al. 2005) could also remove significant prokaryotic biomass from the water column. Zooplankton biomass (>63 μm) in our study mirrored the spatial distribution pattern of microbial communities, as would be expected for consumers of smaller plankton.

Temporal differences — annual

Bi-weekly data did not show clear annual cycles of nutrient concentration and microbial or zooplankton community abundances in Kāneʻohe Bay, as were reported in previous studies (Smith et al. 1981, Taguchi & Laws 1987, 1989). This could be due to the drought during our sampling period, but also demonstrates that response times of the parameters we measured were shorter than the 2 wk sampling interval of this study. This prompted the shorter time scale measurements we recorded during pulses in Kāneʻohe Bay.

When sampling dates were grouped into wet and dry seasons, phosphate and TDP were higher at all sites during the wet season. Nitrate was higher during the wet seasons at the NB and MP sites, as would be expected with discharge from adjacent streams. In contrast, Taguchi & Laws (1987) found a significant seasonal pattern in nitrate, but not in phosphate, in the southern sector of Kāneʻohe Bay. We found that heterotrophic and photosynthetic microbial concentrations were generally higher during the dry season, suggesting that factors other than elevated dissolved nutrients may determine

the abundance of most of microbial groups. An exception was *Chaetoceros*, which showed higher abundances at NB and MP (directly affected by streams) during the wet season. *Chaetoceros* is an opportunistic species that shows high growth rates in response to increased nutrients (Margalef 1958, Brand & Guillard 1981). In our study, zooplankton also showed higher concentrations during the dry season. This fact could be explained by the higher concentration of plankton classes on which zooplankton would be feeding.

Temporal differences—daily, during pulses

Sampling daily during the discharge pulses showed a large increase in nitrate immediately following the rain event. The increases in nitrate concentration varied widely and were not entirely related to the magnitude of the stream discharge, as 4 of the 5 pulses tracked during this study showed a similar peak discharge but produced different increases in nitrate. There was also no consistent relationship between the increase in nitrate and increase in phosphate. Previous history of rain and stream discharge must affect the input of nutrients independently during each pulse. The response of the microbial plankton community was also different among pulses and among plankton groups. Heterotrophic bacteria, *Synechococcus*, and picoeukaryotes did not show strong responses in cell concentration during the pulses, except following the dry season pulse; however, there was an increase in fluorescence of *Synechococcus* hours after the pulse was reported. Ringuet & MacKenzie (2005) also found no change in picoplankton chlorophyll during a bloom in February 2003 in southern Kāneʻohe Bay. Örnólfsson et al. (2004 a,b) found a shift in dominance in biomass from cyanobacteria to diatoms following rivine discharge in Galveston Bay (Texas), and reported that the response was greatest when ambient nitrogen concentrations were lowest. Katano et al. (2005) found that *Synechococcus* did not respond to experimental addition of nutrients, and showed lower cell concentrations following intrusions of nutrient rich bottom waters into Uchiyumi Bay. During our study, large diatoms and dinoflagellates showed a lagged increase in abundance; however, mean silicate concentration at the MP site (17 μM) was never limiting for diatom growth (Dortch & Whitedge 1992). Ringuet & MacKenzie (2005) also found an increase in diatoms and dinoflagellates following the February 2003 discharge pulse. The increase in abundance in our study was especially evident in the opportunistic genus *Chaetoceros*. *Chaetoceros* showed not only a short term response to nutrient enhancement during daily pulse sampling but also seasonal and spatial differences in abundance, associated with nitrate concentrations.

Clearly, responses of picoplankton and other phytoplankton are also influenced by the population dynamics of small grazers. Scheinberg (2005) described increases in copepods, appendicularians, cladocerans, larval bivalves and polychaetes following a storm event in Kāneʻohe Bay in November 2003. The zooplankton community shifted from small copepods to large, gelatinous zooplankton and larval meroplankton, demonstrating that not only bottom-up (increases in nutrients) but also top-down (grazing by zooplankton) factors influence the composition and abundance of plankton. Berninger & Wickham (2005) found that grazing control was stronger for the smaller pico- and nano-plankton groups, whereas microplankton were nutrient limited.

Our data provide new insights into the type and timing of responses of different planktonic groups to environmental changes. Picophytoplankton responded to pulsed discharge with an increase in fluorescence per cell in a time scale of hours, apparently related to reduced light availability following increased TSS during pulse events. Increases in nutrients during pulse events favored microphytoplankton growth with a time response of several days. These shifts may only be evident when pulses are large in magnitude, or water residence time is longer than the lag in response of phytoplankton. Shifts of phytoplankton to larger sizes following nutrient enrichments have been documented in marine waters (Chisholm 1992, Webber et al. 1992, Nakamura et al. 1993, Chang et al. 1996, Agawin et al. 2000, 2003, Örnólfsson et al. 2004 a,b); however, this work provides new insights into the immediate shift in fluorescence, changes in abundance and species composition within the microbial community, and the temporal and spatial extent of these changes. Differences in plankton community composition and species abundance between oceanic and Bay waters seem to be primarily related to physical factors (water circulation patterns and light availability). It seems evident that plankton populations in Kāneʻohe Bay are autochthonous and do not rely upon an external (oceanic) source of nutrients, whereas oceanic sites (at least those within a few kilometres of Kāneʻohe Bay) may receive some nutrients or plankton from the inshore waters.

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