

# Picoplankton carbon flux in Chesapeake Bay

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**ABSTRACT:** Although it is increasingly clear that picoplankton play a major role in the oceanic carbon cycle, relatively little is known concerning the significance of picoplankton in coastal systems subject to significant environmental variance on tidal to interannual scales. Here we report on seasonal and interannual patterns of variability in the productivity and biomass of phototrophic and heterotrophic picoplankton (P- and H-PICO, respectively) and on the flow of carbon from phytoplankton to H-PICO. Annual cycles in the biomass and productivity of both picoplankton trophic levels exhibit winter-spring minima and summer maxima but do not appear to be directly coupled in terms of carbon flow from P- to H-PICO. H-PICO exceeds P-PICO during spring when picoplankton productivity is low, and P-PICO exceeds H-PICO during summer when productivity is high. P-PICO productivity and biomass increase rapidly each year to an early summer peak (20% of total phytoplankton productivity on average) immediately following the collapse of the spring diatom bloom. In contrast, H-PICO productivity and the abundance of bacterioplankton (the predominant group of H-PICO) increase slowly to a late summer peak (equivalent to 16% of phytoplankton productivity on average). Recently released phytoplankton exudates are a major source of dissolved organic carbon with H-PICO taking up an average of 54% during spring and 83% during summer. Variations in H-PICO are closely coupled to the release of DOC, most of which is produced by phytoplankton  $> 2 \mu\text{m}$ . New nutrient input to the Bay appears to be coupled to H-PICO productivity via increases in the productivity and DOC release of phytoplankton  $> 2 \mu\text{m}$  in size.

## INTRODUCTION

Photosynthetic picoplankton (P-PICO,  $< 1$  to  $3 \mu\text{m}$ ), predominantly cyanobacteria (Waterbury et al. 1979, Johnson & Sieburth 1979) and prochlorophytes (Chisholm et al. 1988, Olson et al. 1990), play an important role in the carbon cycles of oceans and lakes (cf. Joint 1986, Stockner & Antia 1986). The proportions of phytoplankton biomass and productivity accounted for by P-PICO typically increase from high to low latitudes (Murphy & Haugen 1985, Joint 1986) and from eutrophic coastal to oligotrophic oceanic systems (Bienfang & Takahashi 1983, Li et al. 1983, Platt et al. 1983, Stockner & Antia 1986, Chavez 1989). In coastal ecosystems, P-PICO has been reported to achieve its annual maximum during summer after the spring bloom in the northern Baltic (Larsson & Hagström 1982) and Celtic Seas (Joint et al. 1986). Little is known concerning the seasonal variability of picoplankton in temperate estuarine systems subject to seasonal variations in incident radiation and nutrient supply.

On average, P-PICO and photosynthetic nanoplank-

ton ( $2$  to  $20 \mu\text{m}$ ) account for most photosynthetic biomass in the oceans although episodic and seasonal peaks in biomass are frequently caused by larger, net-plankton species (Malone 1980a, b, Joint 1986). This is thought to reflect the predominance of small phytoplankton in stable environments where phytoplankton biomass is kept in check by grazing and regenerated production is high relative to new production. Episodic inputs of new nutrients associated with vertical mixing and upwelling events often support increases in the biomass of larger phytoplankton (diatoms in particular) and an uncoupling of phytoplankton production and grazing (Walsh 1976, Malone 1980a, b, Sprules & Munawar 1986, Legendre & LeFevre 1989). In coastal ecosystems, large accumulations of biomass can result in a shift from metazoan food webs that support high fish yields to microbial food webs that lead to greater decomposition and oxygen depletion (e.g. Swanson & Sindermann 1979, Bird & Kalff 1984, Tuttle et al. 1987, Verity 1987).

Free-living heterotrophic bacteria (H-PICO  $< 1 \mu\text{m}$  in size) appear to exhibit patterns of variability that are

similar to P-PICO (cf. Ducklow 1983, 1984). Bacterioplankton account for a significant fraction of particulate organic production in most pelagic ecosystems and can dominate the picoplankton in terms of biomass and the flow of carbon (Hagström et al. 1979, Fuhrman & Azam 1980, 1982, Joiris et al. 1982, Smith et al. 1984, Williams 1984, Ducklow 1986, Fuhrman et al. 1989, Malone & Ducklow 1990). Both P- and H-PICO are important sources of food for heterotrophic nanoplankton (Haas & Webb 1979, Fenchel 1982, Sherr & Sherr 1983, Caron et al. 1985, Goldman et al. 1985, McManus & Fuhrman 1988) and it is generally thought that the microbial food web can be a major source of regenerated nutrients depending on nutrient conditions (Glibert 1982, Azam et al. 1983, Pomeroy 1984, Goldman et al. 1985, Caron et al. 1988, Jumars et al. 1989).

Terrestrially derived dissolved organic carbon (DOC), consisting primarily of humic acids, often accounts for most DOC in estuaries. As indicated by their macromolecular character and by conservative DOC-salinity distributions, humic acids are relatively refractory to microbial decomposition (Mantoura 1981). The principle sources of dissolved organic matter required to support observed levels of bacterioplankton production are believed to be direct release from intact phytoplankton and indirect release through the grazing and excretion activities of zooplankton (Lancelot 1979, Fuhrman et al. 1980, Eppley et al. 1981, Joiris et al. 1982, Larsson & Hagström 1982, Peterson 1984, Caron et al. 1985, Roman et al. 1988). The relative magnitudes of these pathways has important ecological implications in terms of energy flow and the trophic status of aquatic ecosystems (Bird & Kalff 1984, Strayer 1988, Fuhrman et al. 1989, Jumars et al. 1989).

The significance of phytoplankton as the source of dissolved organic substrates for H-PICO metabolism has been deduced from large-scale correlations between phytoplankton productivity, release rate of DOC, and H-PICO productivity (Fuhrman et al. 1980, Bird & Kalff 1984, Ducklow 1984). Measurements of uptake of recently released phytoplankton exudates by H-PICO indicate that this pathway may support from virtually all to a negligible fraction of H-PICO demand (e.g. Lancelot 1979, Nalewajko et al. 1980, Cole et al. 1982, Joiris et al. 1982, Larsson & Hagström 1982, Søndergaard et al. 1985). However, DOC release by phytoplankton is less than 10 % of phytoplankton production under most conditions, and estimates of H-PICO demand often exceed this by a factor of 2 to 5 (Sharp 1977, Fuhrman & Azam 1980, 1982, Sellner 1981, Williams 1981, 1984, Azam et al. 1983, Hagström 1984, Lancelot & Billen 1985, Malone et al. 1986, Bjørnson 1988). Recent studies suggest that much if not most of the DOC required to meet H-PICO demand is generated by the feeding and excretory activities of pelagic

consumers (Roman et al. 1988, Scavia 1988, Strayer 1988, Jumars et al. 1989).

This contribution is the third in a series that addresses the dynamics of phytoplankton and bacterioplankton in the mainstem of Chesapeake Bay (Malone et al. 1986, 1988). The Bay is a eutrophic, partially stratified estuary subject to large seasonal variability in incident radiation, temperature, and freshwater flow (e.g. Kemp & Boynton 1984, Schubel & Pritchard 1986). As shown by Malone et al. (1986, 1988), the annual cycles of fresh water flow, new nutrient input and phytoplankton biomass characteristically exhibit spring maxima while the annual cycles of nutrient regeneration and phytoplankton productivity exhibit summer maxima. The productivity and abundance of bacterioplankton are also high during summer, achieving mean levels of about 20 % and 30 % of phytoplankton biomass and productivity, respectively (Malone et al. 1986, Ducklow et al. 1987, Jonas & Tuttle 1990). Here we report on seasonal and interannual variations in photosynthetic and heterotrophic picoplankton in the context of the annual cycles of phytoplankton biomass and productivity that characterize the mesohaline Bay. The contributions of autotrophic and heterotrophic picoplankton growing in the productive surface waters of the Bay are evaluated in terms of (1) particulate organic carbon production and (2) the flow of organic carbon from phytoplankton to bacterioplankton via recently released phytoplankton exudates.

## METHODS

The experimental design we used is based on a protocol described by Ward (1984) which employs radiotracer and fractionation techniques in an attempt to trace the flow of carbon from the dissolved inorganic pool into phytoplankton size classes and from phytoplankton to DOC and bacterioplankton. Similar experimental approaches have been used by Larsson & Hagström (1982), Jørgensen et al. (1983), and Søndergaard et al. (1985). As discussed below, the results of such experiments are equivocal and must be interpreted with caution given the complex nature of microbial plankton communities (Li 1986). This approach is attractive largely because it is an extension of methodologies that have a long history of use; it has been an important source of information (and debate) concerning the role of microbial food webs in pelagic systems; and it is amenable to field applications requiring frequent measurements. As we hope to show, the approach provides reasonable first order estimates of carbon flows which are useful in elucidating the role of picoplankton in the carbon cycle of Chesapeake Bay.

Experiments were performed at weekly to monthly

intervals on 29 cruises during 1987 to 1989 as part of an interdisciplinary study of nutrient dynamics in Chesapeake Bay. Water samples were collected from 1 to 2 m below the surface at Stn 3 of the 'Chop-Pax' transect (Fig. 1). This station was selected because of its central location within the mesohaline reach of the Bay where most new nutrient input is assimilated and

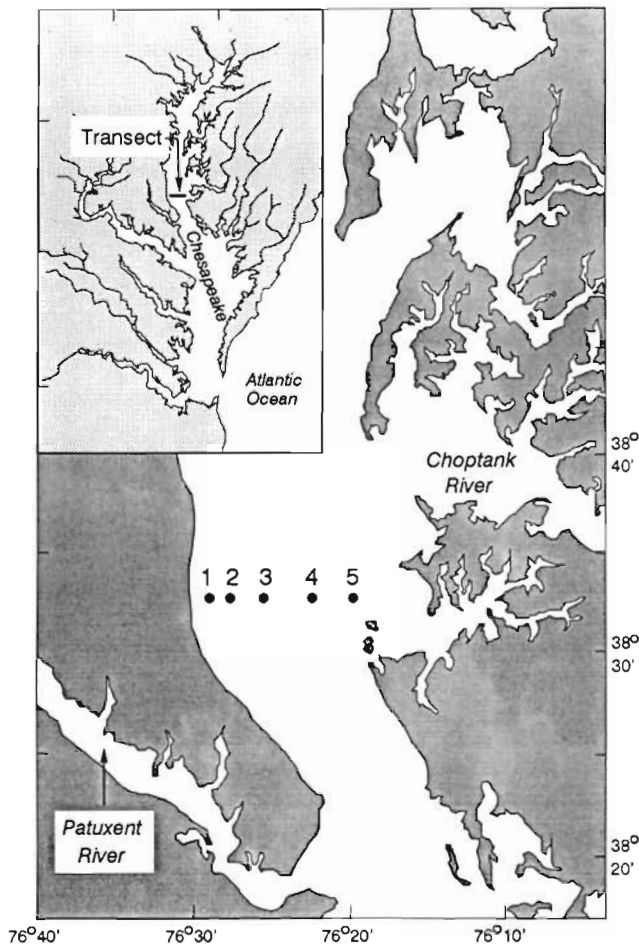


Fig. 1. Chesapeake Bay, USA. Samples were collected at Stn 3 of the Chop-Pax transect

annual production is highest (Boynton et al. 1982, Malone et al. 1988), and it is the site of a time series of measurements of physical and ecological variables initiated in 1984 (Malone et al. 1986).

Each experiment consisted of two 6 h time courses of  $^{14}\text{C}$ -uptake into particulate and dissolved organic pools run in parallel at surface water temperature under 70 % sunlight. Whole (unfractionated) water was used for one time course, and water prefractionated through a 1  $\mu\text{m}$  Nuclepore filter for the second. Prefractionation was done under low vacuum pressure (< 50 mm Hg) using 47 mm, 1  $\mu\text{m}$  Nuclepore filters. Duplicate whole and prefractionated samples were then dispensed into

500 ml acid-cleaned, polycarbonate bottles and inoculated with 220  $\mu\text{Ci}$  of  $\text{NaH}^{14}\text{CO}_3$  (Amersham). Sub-samples of 50 ml were withdrawn from each bottle at Time 0, 1, 2, 3, 4, 5, and 6 h and sequentially filtered through a 25 mm, 1  $\mu\text{m}$  Nuclepore filter (whole water only) and a 25 mm, 0.45  $\mu\text{m}$  Millipore filter (both whole and prefractionated). All filtrations were under low vacuum (< 50 mm Hg) and filtrates were collected for  $\text{DO}^{14}\text{C}$  measurements. Samples collected on 1 and 0.45  $\mu\text{m}$  filters were acidified with 30  $\mu\text{l}$  of glacial acetic acid, exposed to air for 30 min, and placed in 20 ml scintillation vials with 10 ml of Ready-Solv. Filtrates were acidified with 1N HCl to pH 1 and frozen at  $-20^\circ\text{C}$ . Upon returning to the laboratory, the filtrates were thawed, bubbled with filtered air to remove the remaining  $\text{H}^{14}\text{CO}_3^-$ , and dispensed (1 ml aliquots) into 20 ml scintillation vials with 10 ml of Ready-Solv.  $^{14}\text{C}$  activity was measured with an LKB 1212 RackBeta liquid scintillation counter.

Five quantities are measured by this procedure (Fig. 2). Postfractionation time courses yield estimates of (1) photosynthetic carbon incorporation by phytoplankton retained by the 1  $\mu\text{m}$  filter, (2) carbon incorporation by phytoplankton and bacterioplankton passed by the 1  $\mu\text{m}$  filter, and (3) net DOC accumulation by direct phytoplankton release and by indirect release via the microbial food web. Prefractionation time courses yield rates of (4) carbon incorporation by phy-

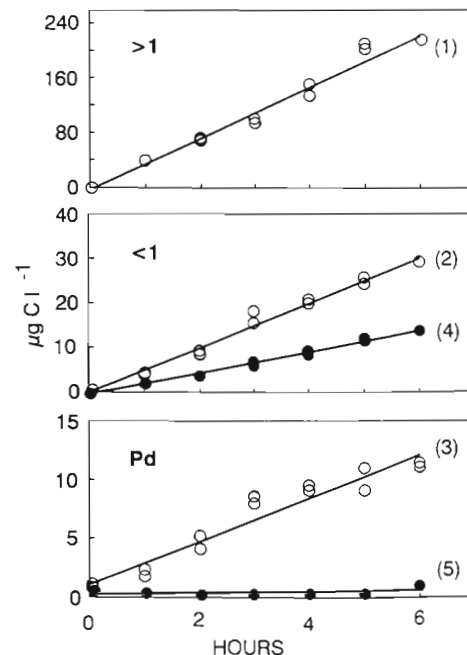


Fig. 2. Representative time courses of  $^{14}\text{C}$ -incorporation into dissolved ( $\text{P}_d$ ) and particulate (> 1  $\mu\text{m}$  and < 1  $\mu\text{m}$ ) pools: time courses 1, 2, and 3 are for post-incubation fractionations; time courses 4 and 5 are for pre-incubation fractionation from an experiment run on 9 June 1988

toplankton and chemosynthetic bacteria passed by the 1  $\mu\text{m}$  filter (assuming a negligible flow of carbon from P-PICO to H-PICO) and (5) net DOC release by picoplankton. As recommended by Li (1986), dark bottle 'controls' were not run. Rather, Time 0 h blanks were subtracted from uptake at each subsequent time point. Although nonlinearities in similar time courses have been reported, particularly for Quantities 3 and 5 (Lancelot 1979), we found uptake into both particulate and dissolved pools to be approximately linear over 6 h (Table 1). Consequently, rates of carbon flow ( $\mu\text{g C}$

Table 1. Summary of coefficients of determination ( $r^2$ ) for 29 time courses of  $^{14}\text{C}$  incorporation into post- and pre-fractionated organic pools as shown in Fig. 2

Fraction	Range	Median	Mean
(1) post- > 1 $\mu\text{m}$	0.90–1.00	0.99	0.97
(2) post- < 1 $\mu\text{m}$	0.72–0.99	0.97	100t95
(3) post-DOC	0.70–0.99	0.93	0.92
(4) pre- < 1 $\mu\text{m}$	0.85–1.00	0.97	0.96

$\text{l}^{-1} \text{h}^{-1}$ ) were calculated from the slope of least square regression of uptake on time (after subtracting Time 0 h blanks) as follows:

$$\begin{aligned} P_p &= (1) + (2) + (3), \\ P_p\text{-PICO} &= (4) + (5), \\ H_p\text{-PICO} &= (2) - [(4) + (5)], \\ P_d &= (3) + [(2) - (4)], \end{aligned}$$

where  $P_p$  = total phytoplankton productivity;  $P_p\text{-PICO}$  = productivity of P-PICO;  $H_p\text{-PICO}$  = uptake of  $P_d$  by H-PICO; and  $P_d$  = total rate of DOC release.

$P_p$ , which was light saturated at the incubation light level (70 %), includes DOC release ( $P_d$ ) estimated as the sum of net DOC release and uptake by H-PICO.  $P_p\text{-PICO}$  is estimated by the sum of particulate uptake and net DOC release in the pre-fractionated time course. Particulate uptake in the unfractionated time course that is in excess of  $P_p\text{-PICO}$  is taken as an estimate of  $H_p\text{-PICO}$ . Note that  $P_p$  and  $P_p\text{-PICO}$  will be overestimated to the extent that anaplerotic fixation occurred. Anaplerotic fixation by bacteria, the most likely source of chemosynthetic assimilation (Li 1986), cancels out in the calculations of  $P_d$  and  $H_p\text{-PICO}$  (2 – 4). As discussed by Li (1986) and others (e.g. Smith et al. 1984), these experiments cannot distinguish between flows of carbon mediated by phytoplankton and microzooplankton when they occur in the same size fraction. Thus, all rates will be underestimated by amounts equivalent to the respective respiratory  $^{14}\text{C}$  loss.

Subsamples of whole water were also used to measure chlorophyll *a* (whole and < 1  $\mu\text{m}$ ) and bacterial abundance and productivity. Chlorophyll *a* (chl) was

measured with a Turner Designs fluorometer following extraction with 90 % acetone. The abundance and productivity of bacterioplankton were estimated from direct counts (Hobbie et al. 1977) and rates of incorporation of methyl- $^3\text{H}$ -thymidine (specific activity > 80 Ci  $\text{mmol}^{-1}$ , New England Nuclear) into cold TCA-insoluble cell (Fuhrman & Azam 1980). Stock solutions of  $^3\text{H}$ -thymidine (TdR) were prepared in 0.2  $\mu\text{m}$  filtered Milli-Q deionized water and stored at 5°C in Teflon bottles. Samples were inoculated with 5 nM TdR and incubated in the dark at ambient temperature for 15 to 45 min. Zero-time uptake was measured to correct for background levels of label. Samples were filtered at low vacuum pressure (< 200 mm Hg) onto 0.45  $\mu\text{m}$  Millipore filters; rinsed 3 times each with cold trichloroacetic acid (TCA), filtered baywater, and 80 % ethanol; and stored dry in scintillation vials. Upon return to the laboratory, filters were prepared for radioassay and counted on an LKB Rack-Beta scintillation counter.

## RESULTS AND DISCUSSION

### Method evaluation

The choice of 1  $\mu\text{m}$  Nuclepore filters to separate picoplankton was based on the retention characteristics of membrane filters and on the size scale proposed by Sieburth et al. (1978), who recommended the term 'picoplankton' for organisms smaller than 2  $\mu\text{m}$ . In contrast to screens for which the stated and effective pore sizes are similar, the effective pore size of membrane filters has been shown to differ from the stated pore size (Sheldon & Sutcliffe 1969). Screens with mesh diameters of 2  $\mu\text{m}$  are available, but we have found that 2  $\mu\text{m}$  Nitex screens pass phytoplankton cells as large as 20  $\mu\text{m}$ . Size fractionation experiments with Nuclepore filters show that 2  $\mu\text{m}$  Nuclepore filters retain only 50 % of 2  $\mu\text{m}$  particles (mean spherical diameter) while 1  $\mu\text{m}$  filters retain 70 % (Sheldon 1972). Consequently, organisms that pass a 1  $\mu\text{m}$  Nuclepore filter have been functionally defined as picoplankton (Joint 1986, Li 1986), a practice that we have adopted here.

One concern was that 1  $\mu\text{m}$  filters would retain a measurable fraction of the bacteria present in the water column. Using surface samples from the mesohaline Bay, we compared the abundance of bacterioplankton in whole, unfractionated samples with the filtrates of 1  $\mu\text{m}$  Nuclepore filters using the AODC technique (Hobbie et al. 1977). The least square regression (model II) of abundance in filtrates on abundance in whole samples gives a slope of 0.96 (SE  $\pm$  0.02) and an intercept of -0.2 (SE  $\pm$  0.3) ( $r^2$  = 0.99,  $n$  = 15), i.e. 96 % of bacteria were passed by 1  $\mu\text{m}$  filters on average. This

result is consistent with the retention efficiencies reported by Sheldon (1972) for bacterial sized particles.

Size fractionation by filtration is subject to errors related to cell breakage and changes in physiology. We attempted to minimize these potential sources of error by using low vacuum pressure ( $< 50$  mm Hg) and large diameter filters (47 mm rather than 25 mm). Waterbury et al. (1986) found that photosynthesis by *Synechococcus* spp. was unaffected by gentle prefiltration through  $1\ \mu\text{m}$  Nuclepore filters (although 20 to 50 % of the cells were retained). An indication that prefiltration had little affect on Chesapeake Bay P-PICO is the strong correlation of  $P_p$ -PICO and the chl content of PICO in samples collected over 3 yr during all seasons ( $r = 0.77$ ,  $n = 27$ ,  $p < 0.001$ ). The intercept of the least square regression (model II) of  $P_p$ -PICO on chl-PICO was not significantly different from 0 ( $-1.0$ ,  $SE = \pm 2.1$ ) and the slope of 6.3 ( $SE = \pm 1.0$ ) approximates the light-saturated, chlorophyll *a* specific  $P_p$ -PICO reported by Joint & Pomroy (1986) for the Celtic Sea in summer. We also have no direct way to evaluate the effects of cell breakage. However, if this was a significant source of error in our estimates of  $^{14}\text{C}$  flow into picoplankton ( $< 1\ \mu\text{m}$ ,  $> 0.45\ \mu\text{m}$ ),  $\text{DO}^{14}\text{C}$  would have increased with time in the prefractionation time courses. This was not observed (e.g. Fig. 2, Quantity 5). Although physiological effects of prefractionation and of cell breakage in postfractionation time courses cannot be ruled out, we believe they were minimal.

The flows of carbon into DOC and H-PICO pools will be underestimated to the extent that isotopic equilibrium is not attained (Wiebe & Smith 1977, Lancelot 1979, Smith 1982). Although an unequivocal demonstration that isotopic equilibrium was achieved in these experiments is not possible, circumstantial evidence indicates that equilibrium was at least approached. First, the time courses were linear, with a resolution of 1 h over 6 h (Fig. 2, Table 1). Second, the accumulation of  $^{14}\text{C}$  in the DOC and H-PICO pools were a nearly constant fraction of total  $^{14}\text{C}$  fixation after the first hour of incubation (Fig. 3). Attainment of equilibrium on these time scales has been observed by a number of workers, particularly in coastal environments where phytoplankton and bacterial productivity are high (Wiebe & Smith 1977, Lancelot 1979, Mague et al. 1980, Smith et al. 1986).

$P_d$  and  $H_p$ -PICO will also be underestimated if there is a substantial amount of DOC released by PICO.  $\text{DO}^{14}\text{C}$  levels in the prefractionated time courses did not increase with time (Fig. 2) and were consistently a small fraction ( $< 1$  to 19 %, median = 6 %, mean = 7 %) of  $\text{DO}^{14}\text{C}$  recovered in the postfractionated experiments. Since carbon flow to phytoplankton to dissolved organic carbon to bacterioplankton initially proceeds as a simple catenary sequence, the appearance of  $^{14}\text{C}$

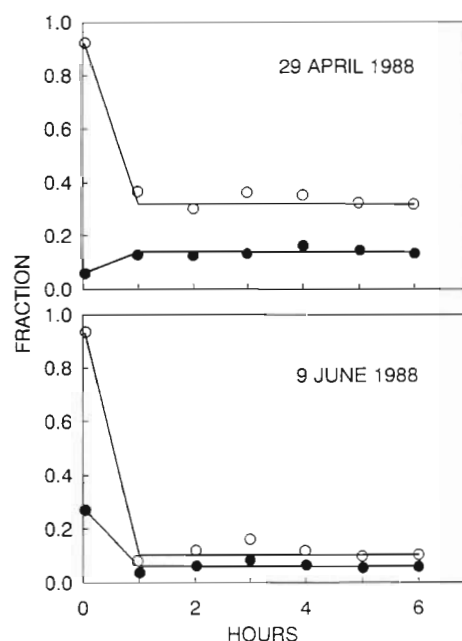


Fig. 3. Representative time courses of the appearance of  $^{14}\text{C}$  in the DOC pool (○) and in bacterioplankton (●) normalized to total  $^{14}\text{C}$  fixation by phytoplankton; calculated from data shown in Fig. 2

in each pool should exhibit a time lag that is related to the number of transfers required to reach that pool (e.g. Li et al. 1983). Unless all  $P_d$  was assimilated by H-PICO with little time-lag ( $< 1$  h) between release and uptake (and the correlation between  $P_p$ -PICO and Chl-PICO was fortuitous), these results indicate that P-PICO was a negligible source of DOC and that  $^{14}\text{C}$  uptake in the prefractionated time courses (Quantity 4 in Fig. 2) provided a reasonable estimate of  $P_p$ -PICO. Most DOC release was apparently by larger plankton, a conclusion reached by Li et al. (1983) working in the Eastern Tropical Pacific.

Comparisons of  $H_p$ -PICO with direct measures of bacterial abundance and  $^3\text{H}$ -thymidine uptake (TdR) suggest that the experimental procedure yields reasonable estimates of carbon flows from phytoplankton to DOC and bacterioplankton.  $H_p$ -PICO was significantly correlated with AODC estimates of bacterial abundance ( $r^2 = 0.57$ ,  $p < 0.01$ ) and with bacterioplankton productivity ( $r^2 = 0.78$ ,  $p < 0.001$ ) as indicated by TdR (Fig. 4). The slope of  $0.04\ \mu\text{g C pmol}^{-1}$  is equivalent to commonly used conversion factors used to calculate bacterioplankton productivity from TdR, e.g.  $4 \times 10^{18}$  cells  $\text{mol}^{-1}$  and a conservative estimate of  $10\ \text{fg C cell}^{-1}$  (Lee & Fuhrman 1987, Fuhrman et al. 1989). The distribution of data in Fig. 4 suggest that conversion factors may have varied from about 0.02 (low growth efficiency) to 0.08 (high growth efficiency) with high



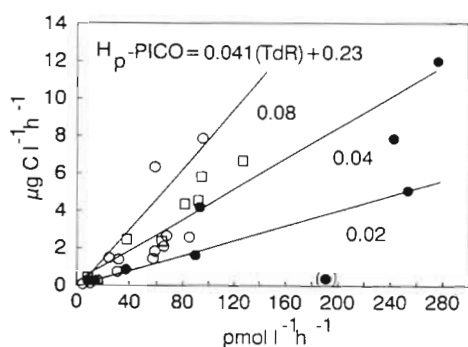


Fig. 4. Comparison of carbon productivity by bacterioplankton calculated from  $^{14}\text{C}$  time courses ( $H_p\text{-PICO}$ ) and uptake rates of  $^3\text{H}$ -thymidine uptake (TdR): the least-square regression is shown by the middle line; lines with slopes of 0.02 and 0.08 are discussed in the text; the point in brackets was not used in the regression analysis. ( $\square$ ) 1987; ( $\circ$ ) 1988; ( $\bullet$ ) 1989

and low growth efficiencies occurring most frequently during 1987–88 and 1989, respectively.

#### Seasonal phytoplankton responses to new nutrient input

In 1987 and 1988, surface phytoplankton biomass and productivity exhibited annual cycles with peaks during spring and summer, respectively (Fig. 5). Chl was highest in May and early June with concentrations of 15 to 20  $\mu\text{g l}^{-1}$ , and  $P_p$  was highest in August with

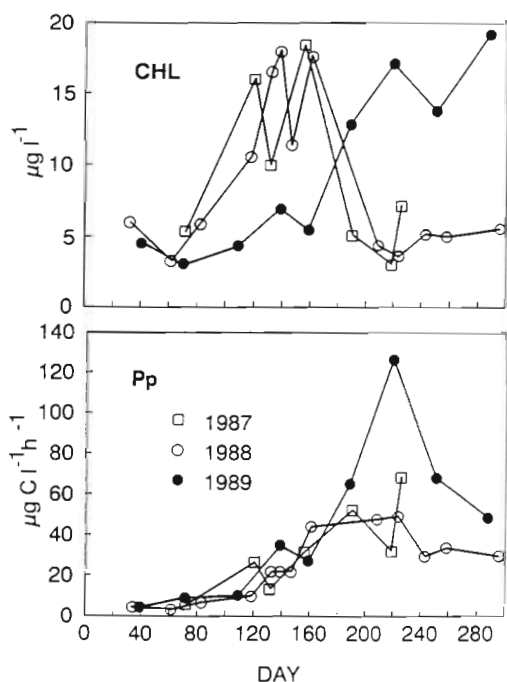


Fig. 5. Surface chlorophyll *a* concentration (CHL) and phytoplankton productivity ( $P_p$ )

rates of 50 to 70  $\mu\text{g C l}^{-1} \text{h}^{-1}$ . This seasonal phasing between chl and  $P_p$  was not observed in 1989 when surface chl exhibited a small spring peak and increased through the summer to an October high of nearly 20  $\mu\text{g l}^{-1}$ . Depth-integrated chl during the 1989 spring bloom was also unusually low (Malone unpubl., Sellner pers. comm.). As in previous years,  $P_p$  peaked in August but at a rate 2 to 3 times the summer peaks of 1987 and 1988.

Annual cycles of water column chl and  $P_p$  are typically out of phase with chl peaking during March to May in response to the seasonal increase in riverborne nutrient input and  $P_p$  during June to August in response to the seasonal increase in incident radiation and temperature (Malone et al. 1988, Malone 1991). The transition from the spring period of high biomass to the summer period of high productivity occurs as surface temperature exceeds 20°C during a 2 wk period in late May–early June and coincides with a change in the distribution of phytoplankton biomass (Malone 1991). During spring, isopleths of chl tend to have a vertical orientation with high chl throughout the water column (in and below the euphotic zone). In contrast, isopleths of chl tend to be horizontally oriented during summer with high chl restricted to the surface layer (within the euphotic zone). The accumulation of biomass during spring is correlated with the volume flow of the Susquehanna River ( $Q_f$ ) which usually reaches its annual maximum in March–April (Schubel & Pritchard 1986, Malone et al. 1988). In 1989,  $Q_f$  did not peak until May (U.S. Geological Survey 1990). Given a 1 mo lag between  $Q_f$  of the Susquehanna river and downstream volume flow in the surface layer of the mesohaline Bay (Malone et al. 1988, Boicourt 1991), the delivery of new nutrients to the mesohaline Bay would not have occurred until after the transition from spring to summer chl distributions. Thus, the small spring bloom and unusually high summer  $P_p$  observed in 1989 probably reflect the timing of the input of new (riverborne) nutrients with respect to the spring–summer transition and the concurrent seasonal increase of incident radiation and temperature.

These contrasts in annual cycles of chl and  $P_p$  between the 1989 and the previous 2 years were not observed in the annual cycles of  $P_p\text{-PICO}$  and  $\text{Chl-PICO}$  (Fig. 6).  $P_p\text{-PICO}$  increased rapidly each year as surface temperature exceeded 20°C in late May–early June. Little or no lag occurred between  $\text{Chl-PICO}$  and  $P_p\text{-PICO}$  which peaked in June–July and July, respectively. As a fraction of  $P_p$ ,  $P_p\text{-PICO}$  was generally less than 2% from February to May, began to increase rapidly in late May–early June, and reached peaks of about 20% in June–July.  $P_p\text{-PICO}$  has been reported to account for similar fractions of summer phytoplankton productivity in temperate continental shelf waters

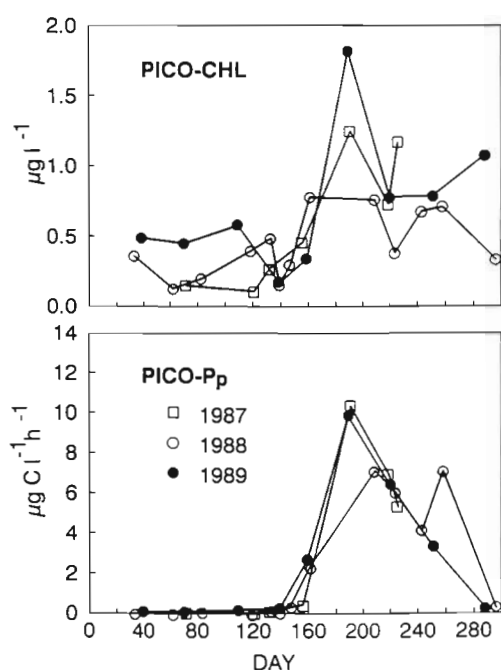


Fig. 6. Surface levels of picoplankton chlorophyll *a* (PICO-CHL) and primary productivity (PICO-P<sub>p</sub>)

(Joint & Pomroy 1983, Douglas 1984, Joint et al. 1986). In Chesapeake Bay, the annual cycle of P<sub>p</sub>-PICO showed remarkably little interannual variability. Apparently, the timing of the increase in P<sub>p</sub>-PICO and the magnitude of the summer maximum was independent of the timing and magnitude of the spring freshwater flow maximum and associated delivery of new nutrients.

The rapid increase in P<sub>p</sub>-PICO reflected an increase in chl-specific P<sub>p</sub>-PICO (P<sub>p</sub><sup>chl</sup>-PICO) from less than 1 µg C (µg chl)<sup>-1</sup> h<sup>-1</sup> at temperatures less than 20°C to rates in excess of 10 µg C (µg chl)<sup>-1</sup> h<sup>-1</sup> at higher temperatures. P<sub>p</sub><sup>chl</sup>-PICO was significantly correlated with temperature ( $r^2 = 0.80$ ,  $p < 0.001$ ) as described by the following least-squares regression (model II, with  $\pm$  SE):

$$\ln(P_p^{\text{chl-PICO}}) = 0.19 \pm 0.02(T) - 3.10 \pm 0.84$$

The exponential constant for this relationship is high compared to values typical of light-saturated photosynthesis and growth of phytoplankton (cf. Malone 1982), perhaps as a consequence of changes in floristic composition. The shift from spring to summer chl distributions coincides with an increase in phytoplankton growth rates from ca 0.1 d<sup>-1</sup> to ca 1.0 d<sup>-1</sup> (Harding et al. 1986, Malone et al. 1988) and with a rapid change in dominant floristic groups from chain-forming diatoms and large dinoflagellates to small, solitary cyanobacteria and  $\mu$ -flagellates (Tyler & Seliger 1978, Van Valkenburg et al. 1978, Malone et al. 1986, Sellner 1987).

Thus, the increase in P<sub>p</sub>-PICO not only signals the rapid transition from spring to summer chl distributions, it marks a major change in floristic composition and a shift from the spring period of slow biomass turnover to the summer period of rapid biomass turnover.

Phytoplankton DOC release (P<sub>d</sub>) roughly paralleled variations in P<sub>p</sub> and was unusually high in late summer 1989 (Fig. 7). Summer rates of 7 to 14 µg C l<sup>-1</sup> h<sup>-1</sup> are

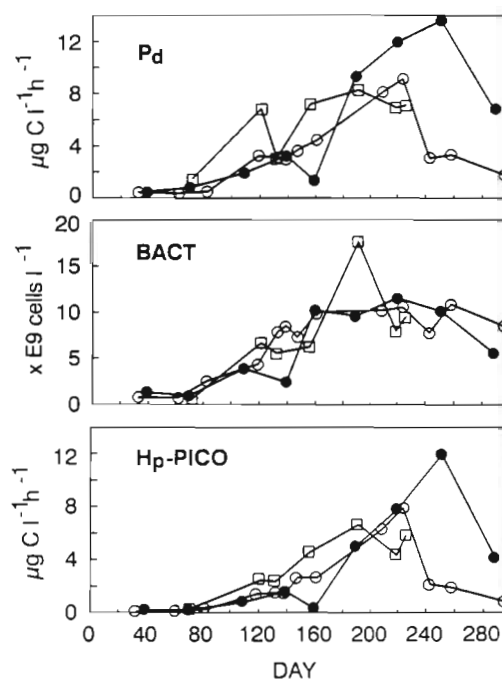


Fig. 7. Release rates of dissolved organic carbon (P<sub>d</sub>), bacterioplankton cell densities (BACT), and uptake rates of dissolved organic carbon by bacterioplankton (H<sub>p</sub>-PICO); symbols as in Fig. 6

also high relative to those reported for other coastal systems (cf. Joint & Morris 1982). P<sub>d</sub> varied between 5 and 35 % of P<sub>p</sub> with the highest percent release occurring in April and July–August. Contrary to many reports (e.g. Thomas 1971, Berman & Holm-Hansen 1974, Søndergaard et al. 1985), percent release was not inversely related to productivity. P<sub>d</sub> was significantly correlated with P<sub>p</sub> ( $r^2 = 0.75$ ,  $p < 0.001$ ) by the least squares regression (model II,  $\pm$  SE)

$$P_d = 0.15 \pm 0.01(P_p) + 0.0 \pm 1.7$$

The slope of the regression gives an estimate of 15 % for the mean fraction of P<sub>p</sub> released as DOC. This is similar to percent release reported for the northern Baltic Sea (Larsson & Hagström 1982) and in the high end of the range reported for phytoplankton in general (cf. Sharp 1977, Lancelot & Billen 1985). Similar experiments in the coastal plume of the Chesapeake Bay indicate release rates of 4 % in April and August (Ducklow & Malone unpubl.).

### Carbon flow to bacterioplankton

Bacterioplankton abundance and uptake of recently released DOC ( $H_p$ -PICO) also increased from winter–spring minima to summer maxima but at a more gradual rate and with a later peak than  $P_p$ -PICO (Fig. 7). The annual cycle of bacterial abundance and the seasonal range of  $10$  to  $20 \times 10^9$  cells  $l^{-1}$  are similar to previous reports for other locations and years in the mesohaline reach of the Bay (Malone et al. 1986, Ducklow et al. 1987, Tuttle et al. 1987, Jonas & Tuttle 1990) and are high compared to other coastal environments (Williams 1984). Like  $P_p$  and  $P_d$ ,  $H_p$ -PICO increased to unusually high rates in late summer 1989 and was significantly correlated with  $P_p$  ( $r^2 = 0.64$ ,  $p < 0.001$ ) by the regression

$$H_p\text{-PICO} = 0.09 \pm 0.01(P_p) - 0.16 \pm 1.82$$

As indicated by the slope, an average of 9% of  $P_p$  cycled through bacterioplankton via recently released DOC (range of 1 to 18%). This is within the range reported for the fraction of phytoplankton production that appears as bacterial production in pelagic systems, and somewhat less than half of that estimated for the mesohaline Bay based on TdR incorporation (Malone et al. 1986, Ducklow et al. 1987, Tuttle et al. 1987, Jonas & Tuttle 1990).  $H_p$ -PICO was better correlated with phytoplankton productivity  $> 2 \mu m$  ( $r^2 = 0.61$ ) than with  $P_p$ -PICO ( $r^2 = 0.39$ ) reflecting the lag between seasonal peaks in  $P_p$ - and  $H_p$ -PICO. At its late summer peak,  $H_p$ -PICO averaged 16% of phytoplankton productivity.

As a fraction of  $P_d$ ,  $H_p$ -PICO was generally between 20 and 60% at temperatures less than  $20^\circ C$  (February to May) and between 50 and 90% at temperatures greater than  $20^\circ C$  (June to September). Similar seasonal ranges have been reported for phytoplankton growing in coastal environments and lakes (e.g. Larsson & Hagström 1982, Søndergaard et al. 1985).  $H_p$ -PICO was significantly correlated with  $P_d$  during spring ( $r^2 = 0.85$ ,  $p < 0.001$ ) and summer ( $r^2 = 0.95$ ,  $p < 0.001$ ) by the least squares regressions (model II,  $\pm$  SE)

$$H_p\text{-PICO} = 0.54 \pm 0.06(P_d) - 0.06 \pm 0.57 \text{ (spring)}$$

$$H_p\text{-PICO} = 0.83 \pm 0.07(P_d) - 0.83 \pm 0.95 \text{ (summer)}$$

The release of DOC by phytoplankton and uptake by bacterioplankton appear to have been closely coupled with  $U_b$  averaging 54% of  $P_d$  at temperatures less than  $20^\circ C$  (spring) and 83% at temperatures greater than  $20^\circ C$  (summer). Lower percent uptake in spring compared to summer may reflect differences in the molecular composition of released compounds (Lancelot 1984), lower bacterial demand, or differences in the  $^{14}C$  labeling kinetics of the DOC and  $H$ -PICO pools (e.g. isotopic equilibrium was more closely achieved during summer than spring). In any case, phytoplankton

exudates were a significant source of organic substrates for bacterioplankton production, particularly during summer.

Taken together, the proportion of total particulate production accounted for by  $P_p$ - and  $H_p$ -PICO increased from less than 10% in late winter to nearly 30% in summer (Fig. 8). Within the picoplankton,  $H$ -PICO accounted for most production during spring when rates were low (Fig. 8). However,  $P_p$ -PICO

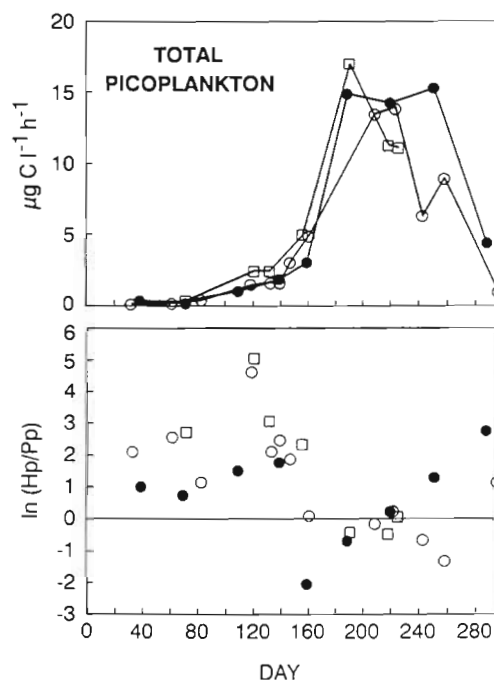


Fig. 8. Picoplankton productivity (phytoplankton + bacterioplankton) and the natural log of the ratio of bacterioplankton to phytoplankton productivity within the picoplankton ( $H_p/P_p$ ); symbols as in Fig. 6

increased relative to  $H_p$ -PICO during the transition from spring to summer chl distributions and  $H_p$ -PICO fluctuated around 50% over much of the summer when turnover rates were high. Summer 1989 was again the exception when  $H_p$ -PICO increased to unusually high rates and accounted for over 70% of the carbon flow through the picoplankton.

### DOC turnover

A rough indication of DOC turnover in the surface layer can be gained by comparing phytoplankton release rates with measured concentrations of DOC. DOC in the mesohaline reach of the Bay varies from 1.5 to  $6.5 \text{ mg } l^{-1}$ , tends to decrease with increasing salinity, and averages about  $3.5 \text{ mg } l^{-1}$  with no obvious seasonal trend (Fisher pers. comm.).  $H_p$ -PICO averaged



1.3 and 5.0  $\mu\text{g C l}^{-1} \text{h}^{-1}$  during spring and summer, respectively. Using the mean DOC concentration of 3.5  $\text{mg l}^{-1}$ , these rates give turnover times of 112 d during spring and 29 d during summer.

Dissolved organic compounds range from complex macromolecules (e.g. humic acids) which turn over slowly to simple monomers (e.g. dissolved free amino acids, glycolate) which turn over rapidly (Mantoura 1981, Lancelot 1984, Fuhrman 1990). Although we do not have direct measurements of either of these general pools of organic compounds, measurements of biological oxygen demand (BOD) from 5 d BOD measurements can be used to roughly estimate the DOC content of the high turnover pool. Jonas & Tuttle (1990) measured the 5 d BOD for surface water of the mesohaline reach of Chesapeake Bay after filtering through Gelman A/E glass fiber filters. Mean spring and summer values were 0.27  $\text{mg C l}^{-1}$  and 0.42  $\text{mg C l}^{-1}$ , respectively. Assuming these to be estimates of the concentration of DOC in the high turnover pool and using 3.5  $\text{mg C l}^{-1}$  as an estimate of total DOC, these results indicate that about 10% of the DOC pool is turning over on a time scale of days or less. Based on mean  $H_p\text{-PICO}$ , turnover times of this pool are on the order of 9 d during spring and 3 d during summer. Assuming that  $P_d = H_p\text{-PICO}$  on a time scale of days, turnover times would be on the order of 4 d during spring and 2 d during summer (mean  $P_d = 2.7$  and 7.0  $\mu\text{g C l}^{-1} \text{h}^{-1}$ ). Although 5 d BOD measurements are at best a rough indication of biologically labile DOC, these calculations indicate that recently released phytoplankton exudates are important in the carbon dynamics of the Bay.

## CONCLUSIONS

Annual cycles of phytoplankton and bacterioplankton productivity are similar, with bacterioplankton productivity averaging 9% of phytoplankton productivity. The transition from a high biomass, slow turnover system during spring to a lower biomass, high turnover system during summer is marked by a rapid increase  $P_p\text{-PICO}$  as surface temperature increases above 20°C.  $H_p\text{-PICO}$  increases more slowly so that  $H_p\text{-PICO}$  decreases from greater than 90% of total picoplankton productivity during spring to about 50% during summer. Thus, carbon flow through the picoplankton, and presumably through the microbial food web as a whole, increases to a summer maximum due to increases in  $P_p\text{-PICO}$  and, to a lesser extent, in  $H_p\text{-PICO}$ .

The extent to which  $P_p\text{-PICO}$  dominates carbon flow through the picoplankton can be modified by the timing of the spring freshet. The high summer  $P_p$  in 1989 appears to be a response to a late freshet which deli-

vered nutrients to the mesohaline Bay following the transition from spring to summer plankton communities. The annual cycle of  $P_p\text{-PICO}$  was unaffected, but  $P_p$  was higher due to an increase in the productivity of phytoplankton  $> 2 \mu\text{m}$ . DOC uptake by bacterioplankton was also higher during the summer of 1989, perhaps in response to higher DOC release rates by phytoplankton. Since picoplankton appear to be an insignificant source of DOC, these results suggest that nutrient input associated with the late 1989 freshet stimulated carbon flow through the microbial loop via an increase in DOC release by phytoplankton  $> 2 \mu\text{m}$  and uptake by bacterioplankton. Bacterioplankton abundance was not higher than in previous years, suggesting that high  $H_p\text{-PICO}$  reflected higher growth rates or lower growth efficiencies of bacterioplankton. If growth rates were higher, bacterioplankton abundance must have been kept in check by a corresponding increase in grazing losses. The possibility of lower growth efficiency is suggested by the observation that in late summer 1989  $H_p\text{-PICO}$  was low for a given rate of TdR incorporation when compared to the previous 2 years, i.e. the rate of TdR incorporation would be related to carbon productivity by a low conversion factor (Fig. 4).

Although the high productivity characteristic of many estuaries is apparently a consequence of the rates of supply and recycling of nutrients (Nixon 1981, Boynton et al. 1982), the role of bacterioplankton is largely unknown. Inputs of new (allochthonous) nutrients have been shown to stimulate the production of particulate organic matter by both phytoplankton and bacterioplankton on seasonal and annual scales (Boynton et al. 1982, Bird & Kalff 1984, Hobbie & Cole 1984). In Chesapeake Bay, seasonal variations in bacterioplankton productivity are not directly related to the input of new nutrients nor to phytoplankton biomass (which peak during spring) but to phytoplankton productivity and DOC release (which peak during summer).

Bacterioplankton take up an average of 54% and 83% of  $P_d$  during spring and summer, respectively. Assuming a bacterial growth efficiency of 50%, recently released phytoplankton exudates could account for roughly 25% (spring) to 40% (summer) of the carbon metabolized by bacterioplankton. The release of DOC by phytoplankton is a major source of the dissolved organic carbon required to support high bacterioplankton productivity in the Bay, particularly during the summer when phytoplankton productivity is high and river flow is low.

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