

Complex interactions in microbial food webs: Stoichiometric and functional approaches

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ABSTRACT

The food web structure in some high mountain lakes deviates from the established tendency of high heterotrophic bacteria: phytoplankton biomass ratios in oligotrophic ecosystems. Thus, the microbial food web in La Caldera Lake is weakly developed, and bacteria constitute a minor component of the plankton community in terms of abundance, biomass and production. Autotrophic picoplankton is absent, and heterotrophic microbial food web is weakly developed compared to a grazing chain dominated by calanoid copepods and a phytoplankton community mainly composed of mixotrophic flagellates. In order to explain the singular food web structure of this lake, functional, stoichiometric and taxonomical approaches are followed to assess, on various temporal and spatial scales, the relevance of stressful abiotic factors (ultraviolet solar radiation and P-limitation) on the structure and functioning of this ecosystem. P-availability was the main factor controlling the algal biomass whereas bacterial P-limitation was a transient phenomenon. The algae-bacteria relationship was predominately commensalistic. In contrast to algae, full-sunlight radiation had no negative effect on bacterial growth but rather enhanced bacterial dependence on the carbon released by algae. The prevalence of the commensalistic-mutualistic relationship and the development of a more complex microbial food web were related to the stoichiometry of algae and bacteria (N:P ratios). The microbial food web only developed at balanced algal and bacterial N:P ratios, with the appearance of ciliates after a nutrient pulse. However, mixotrophic algae dominated the planktonic community under P-deficit conditions, and they were the main factor controlling bacterioplankton. Their regulatory effect has a dual nature: (i) a resource-based control, where bacteria depend on the photosynthetic carbon released by algae, i.e., a commensalistic interaction (“without you I cannot live”); and (ii) a predatory control, where bacteria is a prey for mixotrophs (“with you I die”). Hence, the niche of microheterotrophs (nanoflagellates and ciliates) is occupied by mixotrophs, and there is a resulting simplification of the planktonic structure. With respect to the carbon cycle, mixotrophic bacterivory constitutes a “by-pass” for the flux of C towards the grazing chain, precluding the development of a complex heterotrophic microbial food web. Mixotrophs thereby improve the energetic transfer efficiency in high mountain lakes through a reduction in the number of trophic levels. Antagonistic effects of UVR x P interactions on the algae-bacteria relationship were caused by an enhancement of dual (resource and predation) control. Based on these results, an alternative model for the flux of C in autotrophic high mountain lakes has been proposed.

Key words: Carbon flux, high mountain lakes, microbial loop, nutrient limitation, stoichiometry, ultraviolet radiation.

RESUMEN

La estructura de la red trófica en algunos lagos de alta montaña, se aleja de los patrones establecidos para ecosistemas oligotróficos que proponen el predominio de la red trófica microbiana sobre la cadena de pastoreo. Así, en la laguna de La Caldera las bacterias son el componente minoritario de la comunidad planctónica en términos de abundancia, biomasa y producción. El picoplancton autótrofo está ausente y la red microbiana heterotrófica se encuentra escasamente desarrollada frente a una cadena de pastoreo dominada por copépodos calanoides y algas mixotróficas. Para comprender los mecanismos que determinan esta estructura trófica hemos seguido diferentes aproximaciones de análisis: funcional, estequiométrica y taxonómica sobre distintas escalas espaciales y temporales, en relación con los principales factores de estrés abiótico (radiación ultravioleta y limitación por fósforo) que controlan el funcionamiento de los ecosistemas de alta montaña. Nuestros resultados indican que la disponibilidad de fósforo, de forma generalizada, controla la biomasa algal y de manera transitoria la bacteriana, estableciéndose entre ambas comunidades una relación comensalista. La radiación solar completa no afecta negativamente el desarrollo de las bacterias y si el de las algas y potencia la relación de dependencia por el carbono orgánico (comensalismo) entre algas y bacterias. El predominio de la relación comensalista-mutualista y el desarrollo del bucle microbiano esta relacionado con la estequiometría (razón N:P) de algas y bacterias. Así, sólo cuando la razón N:P de algas y bacterias es equilibrada para crecer, un pulso de

nutrientes permite el desarrollo del bucle microbiano. En condiciones naturales de déficit de P, sin embargo, existe un predominio de "algas" con metabolismo mixotrófico. Las algas mixotróficas ejercen un efecto regulador dual sobre las bacterias que denominamos Ni contigo ni sin ti, (i) control por depredación, donde las bacterias son consumidas por algas mixotróficas ("contigo me muero"), (ii) control basado en los recursos estableciéndose una relación de dependencia de las bacterias sobre del carbono liberado por las algas ("sin ti no puedo vivir"). La mixotrofia supone una simplificación en la cadena trófica microbiana, donde los mixótrofos ocupan el nicho potencial de nanoflagelados y ciliados. Desde un punto de vista energético implica un cortocircuito en el flujo de energía y un incremento en la eficiencia de transferencia energética en ecosistemas ultraoligotróficos y con alta dosis de radiación ultravioleta (RUV). Los efectos de la interacción entre RUV y pulsos de P tienen un efecto antagónico sobre la interacción alga-bacteria, intensificando la interacción comensal-depredadora. A partir de los resultados obtenidos proponemos un modelo alternativo de flujo de energía para ecosistemas autotróficos de alta montaña.

Palabras clave: Bucle microbiano, Estequiometría, Flujo del carbono, Limitación por Nutrientes, Lagos de Alta Montaña, Radiación Ultravioleta

INTRODUCTION

The food web constitutes one of the most complex conceptual phenomena in modern Biology (Pimm *et al.*, 1991). This complexity is increased by the "nodes" of the food web, through which energy and materials flow, are formed by individual organisms of diverse species, each individual is a complex biochemical system, and every species is the product of ongoing evolutionary change (Holt, 1995). Nevertheless, according to Margalef, knowledge of the relations among elements is more important to the understanding of a system than knowledge of the precise nature of its constituents (Margalef, 1992). It is therefore crucial to study the interactions that take place in the food web –an important issue in the emerging science of "Biocomplexity" (Michener *et al.*, 2001)–, and to develop appropriate concepts or key variables for this purpose. Trophic level (tropho-dynamic view, Lindeman, 1942) and body size, which control many of the physiological properties of organisms and influence the trophic structure of the communities (Rodríguez, 1999), are among two of the traditionally considered variables. Recently, Sterner & Elser (2002) proposed stoichiometry as a tool for ecological analysis in order to relate the elemental composition of organisms (carbon [C], nitrogen [N] and phosphorus [P]) to their growth rates and resource availability (Biological stoichiometry; Elser *et al.*, 1996). Furthermore, stoichiometry permits the examination of the balance of energy (usually characterized in carbon terms) with multiple chemical

elements (e.g., nitrogen and phosphorus) in the organisms, and of the relationship between ecological interactions and biogeochemical cycles (Ecological stoichiometry; Sterner & Elser, 2002). More recent studies of food webs suggest that the supposed complexity of the food web is more apparent than real (Elser & Hessen, 2005). On the grounds of biosimplicity, they proposed the consideration of the Darwinian paradigm in food web models alongside elemental composition and the laws of thermodynamics. This multiple approach avoids the bias implied by the use of a single tool to study the complexity of food webs. As these authors put it, "If your only tool is a hammer, then every problem looks like a nail". This multiple approach has been adopted in the food web analysis presented in this review.

WHAT WE KNOW AND WHAT WE DO NOT UNDERSTAND

Food webs in aquatic ecosystems function via the channeling of energy and the flux of materials among diverse organism assemblages organized in two chains: the classical grazing chain (phytoplankton-zooplankton-fish), traditionally considered the main pathway for the flow of energy within these ecosystems; and the so-called "microbial loop", (bacteria-heterotrophic nanoflagellates [HNF] - ciliates). The latter is parallel to and converges with the grazing chain via zooplankton consumption of bacteria, HNF, and ciliates (Pomeroy, 1974; Azam *et al.*, 1983). The clearly established

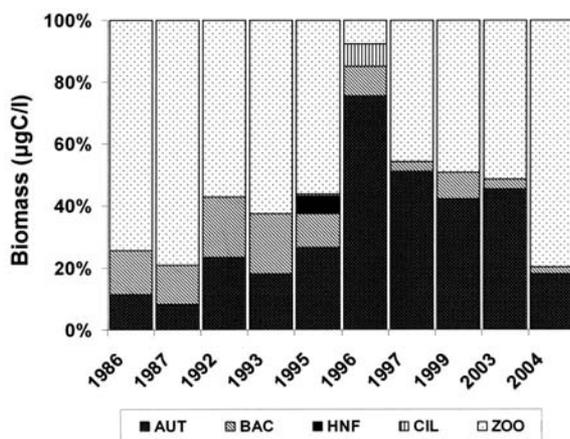


Figure 1. Inter-annual variability of the food web structure of the plankton community in La Caldera Lake. AUT, phytoplankton; BAC, bacteria; HNF, heterotrophic nano-flagellates; CIL, ciliates; ZOO, zooplankton. *Variabilidad interanual de la estructura de la red trófica de la comunidad planctónica de la laguna de La Caldera. AUT, fitoplancton; BAC, bacteria; HNF, nanoflagelados heterotróficos; CIL, ciliados; ZOO, zooplankton.*

dominance of the microbial communities in oligotrophic ecosystems (del Giorgio & Gasol, 1995; Gasol *et al.*, 1997; Vadstein, 2000) derives from their ability to consume organic matter and mineral nutrients at higher rates compared to algae (Biddanda *et al.*, 2001). The dominance of the microbial community in the metabolic balance of oligotrophic ecosystems invites an in-depth study of: the functioning of this compartment, the effects of variations in abiotic factors (i.e. ultraviolet radiation [UVR], nutrient availability) on interactions among its components, and the propagation or attenuation of stress-factor effects throughout the food web. The biological simplicity of the high mountain lakes in Sierra Nevada National Park, which is mediated by strong physical regulation and extreme oligotrophic conditions, makes them an ideal setting for observing and modeling the structure and functioning of the microbial food web. Thus, on an inter-annual scale, the food web structure of one of these ecosystems, La Caldera Lake (Fig. 1), deviates from the trends established for oligotrophic ecosystems, with scant microbial loop development, frequently represented only by bacteria. The first question that arises is then,

¿Why does the microbial loop fail to develop in La Caldera Lake?

Algae-bacteria interactions have been shown to be the key link between the classical grazing chain and the microbial food web, and the coexistence of both trophic levels is a “sine qua non” condition for the persistence of the ecosystem (Daufresne & Loreau, 2001). Different analyses in freshwater and marine ecosystems of the relationship between the biomass/production of bacteria and algae (Cole *et al.*, 1988; Teira *et al.*, 2001; Carrillo *et al.*, 2002) have shown a direct relationship between these organisms. The commensalistic algae-bacteria relationship based on bacterial dependence on organic C released by algae has provided the key explanation of some empirical trends. Accordingly, it is assumed that bacterioplankton is not limited by mineral nutrients (Chrzanowski & Grover, 2001) and establishes an indirect mutualistic interaction with algae, whereby algae provide carbon to sustain bacterial growth and bacteria recycle mineral nutrients to support algal growth (Aota & Nahajima, 2001). This mutualistic interaction is often established in lakes with negligible allochthonous C input, absence of anoxic (or hypoxic) hypolimnion (see discussion in Pace & Cole, 1994), or a high N:P inorganic ratio (Lee *et al.*, 1994), all characteristics of high mountain lakes. Nevertheless, preliminary studies in La Caldera Lake showed a decoupling between the two communities (Reche *et al.*, 1996). Since the algae-bacteria relationship is critical to an understanding of microbial loop development, deeper analysis of the nature of this interaction is required.

Daufresne & Loreau (2001) were the first to propose that the algae-bacteria relationship varies according to stoichiometry of C and nutrient content in the matter transferred from primary producers to decomposers. Because the composition of the released organic matter and the nutritional status of the algae are intrinsically linked (Obernosterer & Herndl, 1997; Caron *et al.*, 2000; Sterner & Elser, 2002), organism stoichiometry may be a useful

tool to understand algae-bacteria interactions, since it yields data on nutrient demands and nutrient limitations to growth. Thus, according to the elemental C:N:P cell content and the availability of substrates, bacteria or algae will be limited by one element or another. For example, if algae are P-deficient (high C:P), the organic matter released will be characterized by a high C:P ratio, leading towards a P-limitation for bacteria. The higher nutrient requirements of bacteria in terms of biomass, C:P and N:P ratios are met by the immobilization of mineral nutrients (Daufresne & Loreau, 2001). Therefore, competition results from a high C:nutrient ratio of the matter transferred between algae and bacteria. Thus, the algae-bacteria relationship can be changed from mutualism to competition for mineral nutrients, especially in oligotrophic environments (Aota & Nahajima, 2001; Cotner & Biddanda, 2002; Joint *et al.*, 2002). Because of the dynamic nature of the algae-bacteria interaction, the relative importance of mutualism versus competition must be studied, as well as the individual and interactive effects of stress-factors (UVR and P-limitation) on this interaction. These fundamental ecological questions will be addressed in this review, allowing us to propose an alternative model for the microbial food web in high mountain lakes.

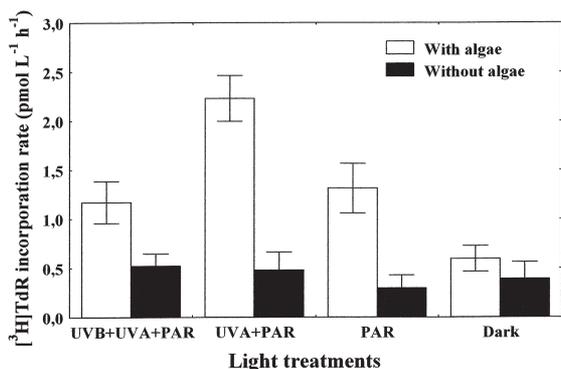


Figure 2. Bacterial [³H] TdR incorporation rates in the presence versus the absence of algae under different light treatments. Error bars indicate means \pm SD, $n = 3$. *Valores de la tasa de incorporación de [³H] TdR por bacterias en presencia vs. ausencia de algas bajo diferentes tratamientos de luz (UVB+UVA+PAR, UVA+PAR, PAR, Oscuridad). Las barras indican medias \pm SD, $n=3$.*

THE MICROBIAL LOOP DOES NOT DEVELOP. DAZZLED BY THE LIGHT?

The high mountain lakes at Sierra Nevada receive a high ultraviolet radiation (UVR) flow due to their altitude (3000 m) and proximity to the subtropics. On the other hand, the low DOC concentration in the water of these lakes (Reche *et al.*, 2001), a consequence of the scarce development of terrestrial vegetation in their catchment areas, allows for a high level of UVR penetration into the water column. Therefore, the low development of the microbial loop might be related to the direct negative effects of ultraviolet-B radiation (<320 nm) (UVB) on these communities. Experimental evaluation of the bacterial activity under different light qualities confirmed our initial hypothesis. Thus, UVB reduced bacterial [³H] TdR incorporation rates by 39-87 % at the surface, although it had no significant effect at intermediate layers (Fig. 2) (Carrillo *et al.*, 2002; Medina-Sánchez *et al.*, 2002). In contrast, either UVA and photosynthetic active radiation (UVA + PAR) or PAR alone exerted a stimulatory effect on [³H] TdR incorporation rates (Fig. 2). This bacterial response was interpreted as the result of photorepair mechanisms (Kim & Sancar, 1993; Vincent & Roy, 1993), which were especially efficient under high light intensities (Carrillo *et al.*, 2002). This photorepair effect is translated into a net stimulatory full-sunlight effect (UVB + UVA + PAR) on bacterial growth (Fig. 2). Nevertheless, an adequate supply of photosynthetic carbon from algae is required to support a net increase in bacterial growth after the recovery of UVB-damaged bacteria. In fact, in experiments made in the absence of algae, this potential recovery did not manifest as enhanced bacterial growth (Fig. 2) (Medina-Sánchez *et al.*, 2002).

Can photosynthetic C supply be inhibited by ultraviolet radiation? Our experimental results showed that UVR inhibits primary production (PP) but increases excretion of organic carbon by algae (EOC) (Carrillo *et al.*, 2002). A higher proportion of this released C is assimilated by bacteria (% PEA) under UVA + PAR or PAR rather than under UVB radiation. These data are compatible with the mechanism proposed

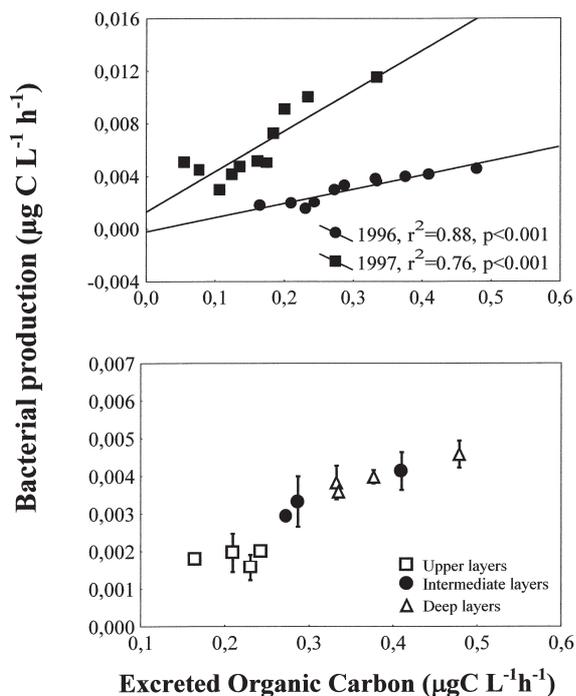


Figure 3. Annual relationship between Bacterial Production and Excretion of Organic Carbon (EOC) for the entire water column (A), and discriminating between depths in 1997 (B). Error bars indicate means \pm SD of bacterial production, $n = 3$. *Relación anual entre Producción Bacteriana y Excreción de Carbono Orgánico (EOC) en la columna de agua completa (A) y discriminando entre profundidades (B). Las barras indican medias \pm SD de producción bacteriana, $n=3$.*

by Berman-Frank & Dubinsky (1999) for the regulation of algae-bacteria interaction in poor-mineral nutrient ecosystems with high UVR flow, and also with the stimulatory effect on bacterial $[^3\text{H}]$ TdR incorporation rates under photorepair light (UVA and/or PAR). The consistency of the results obtained (increased net bacterial activity) using two different methodological approaches suggests a relationship between the consumption of bacterial C and the ability to channel it to new bacterial biomass. For this reason, our group proposed the PB:PEA ratio (as percentage; % CUEb) to quantify the use of photosynthetic carbon by bacteria. Interestingly, UVA + PAR or PAR were shown to increase carbon use efficiency (Carrillo *et al.*, 2002). Therefore, a net increase of bacterial production (PB) under high full-sunlight inten-

sity is possible because the negative effect of UVB radiation on bacteria is counteracted by the positive effect of UVA and PAR (Kim & Sancar, 1993; Kaiser *et al.*, 1997; Davidson & Van der Heijden, 2000), and the presence of higher photosynthetic C availability. A net positive bacterial response to full-sunlight would lead to a coupling between PB and PP (and EOC) (Fig. 3). In fact, when PB measurements were obtained under full-sunlight instead of the usual darkness incubations (Fig. 4), a coupled algae-bacteria relationship was verified. From all of the data above, we conclude that full-sunlight has no negative effect on bacterial development (basis of microbial loop) and increases bacterial dependence on organic carbon released by algae (commensalism). Moreover, the absence of the algal fraction produced a highly significant decrease in PB under different regions of the spectral solar radiation, confirming the dependence of bacteria on carbon released by algae (“without you I cannot live”) (Fig. 2). Further questions arose from these results: (i) What other factors, besides light quality, restrict bacterioplankton and microbial loop development in La Caldera Lake?; (ii) How prevalent is commensalism versus competition between algae and bacteria during the ice-free period? A more detailed analysis of the results obtained in this ecosystem revealed that (i) the stimulation of bacterial growth under full-sunlight only occurred when bacterial elemental composition was balanced and therefore suitable for growth ($\text{N:P} \leq 20\text{-}24$ *sensu* Chrzanowski *et al.*, 1996) (Medina-Sánchez *et al.*, 2002) and (ii) algae can regulate C-release depending on their elemental composition and the nutrient-availability in lake water (Berman-Frank & Dubinsky, 1999; Villar-Argaiz *et al.*, 2002a). This ability permits algae to modify the strength of their commensalistic relationship with bacteria and can even lead to a competitive interaction (Reche *et al.*, 1997; Villar-Argaiz *et al.*, 2002a). These results indicate that mineral nutrient availability and algal and bacterial elemental composition can generate important constraints in the development of the microbial loop.

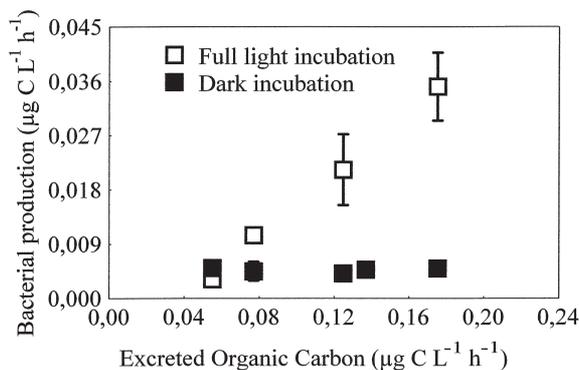


Figure 4. Relationship between Bacterial Production and Excretion of Organic Carbon (EOC) measured at upper depths in 1997, with the bacterial production data obtained under full sunlight or dark incubations. Bars indicate means \pm SD of EOC, $n = 3$. *Relación entre Producción Bacteriana y Carbono Orgánico Excretado (EOC) en la capas superficiales del lago en 1997, con medidas de producción bacteriana obtenidas en incubaciones con luz completa y oscuridad. Las barras indican medias \pm SD de EOC, $n=3$.*

THE MICROBIAL LOOP DOES NOT DEVELOP. IS IT NUTRIENT LIMITED?

Most high mountain lakes in Sierra Nevada are chronically P-deficient ecosystems (DIN:TP > 40; Carrillo *et al.*, 1996; Morales-Baquero *et al.*, 1999). This strong limitation may be the cause of the low bacterial and microbial loop development. Nevertheless, our studies have shown that variances in bacterial activity or biomass in La Caldera Lake are not explained by the concentrations of Total Phosphorus (PT) or Total Dissolved Phosphorus (TDP) (Reche *et al.*, 1997; Medina-Sánchez *et al.*, 1999; Carrillo *et al.*, 2002). Moreover, P-rich atmospheric inputs (Saharan dust) (Carrillo *et al.*, 1990a; Villar-Argaiz *et al.*, 2001, 2002a; Pulido-Villena, 2004) are largely (60-80 %) incorporated into the algal fraction (Fig. 5) (Villar-Argaiz *et al.*, 2001; Villar-Argaiz *et al.*, 2002b). This preferential algal P-consumption has been demonstrated in natural observations (Fig. 5) and after experimental nutrient amendments in the short-term (hours) (Villar-Argaiz *et al.*, 2002a; Medina-Sánchez *et al.*, 2002), medium-term (days), and long-term (weeks-months). These results are consistent with the greater abi-

lity of algae versus bacteria to incorporate limiting nutrients at high P-substrate concentrations (Tarapchak & Moll, 1990). These mechanisms undoubtedly offer an ecological advantage for algae since they can grow by consuming intermittent P-pulses associated with allochthonous inputs (Cotner & Wetzel, 1992; Duarte *et al.*, 2000) and P recycled by zooplankton (Carrillo *et al.*, 1995, 1996; Reche *et al.*, 1997). This generalized response to enrichment suggests that P-availability controls algal biomass in La Caldera Lake but is not a leading factor in the restriction of bacterial activity or biomass on a seasonal or experimental scale.

The stimulatory effect of P on primary producers, in some conditions, is propagated throughout the microbial loop, allowing ciliate development. C-flux quantification allowed us to propose that the complex microbial loop is established by atmospheric nutrient inputs (Medina-Sánchez *et al.*, 1999; Villar-Argaiz *et al.*, 2002a) or P-addition, which stimulate algal growth, and algae-released organic carbon (Medina-Sánchez *et al.*, 2006). The increase in C-availability enhances bacterial production (without changes in bacterial abundance or biomass), enabling ciliate development (Medina-Sánchez *et al.*,

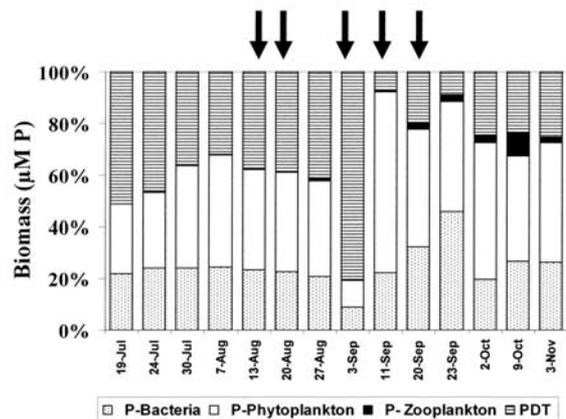


Figure 5. Temporal distribution of phosphorus in particulate (bacteria, phytoplankton, and zooplankton) and dissolved (TDP) fractions during 1996 in La Caldera Lake. Arrows indicate humid P inputs in the Lake area. *Distribución temporal de fósforo en las fracciones particuladas (bacteria, fitoplancton y zooplancton) y disuelta (TDP) durante el año 1996 en la laguna de La Caldera. Las flechas indican entradas atmosféricas húmedas de fósforo en el área de la laguna.*

1999; Villar-Argaiz *et al.*, 2002a). These results confirm that nutrient enrichment can lead to a lengthening of the microbial food chain (Samuelson *et al.*, 2002; Vaqué *et al.*, 2003). Therefore, the scarce microbial loop development can at least be partly ascribed to the severe oligotrophic conditions of this ecosystem. Nevertheless, propagation of the P-pulse effect to ciliate development does not always take place in the system, raising the question as to what constrains the development of a complex microbial loop. Microcosm studies showed that balanced algal (N:P = 16) and bacterial (N:P \leq 20-24) elemental compositions are required, providing further evidence that bacterial elemental composition plays a key role in determining the nature of the algae-bacteria relationship and the subsequent development of ciliates.

Thus, when bacteria (N:P \leq 20-24) and algae (N:P \approx 12) were P-sufficient (i.e., at thaw), algal P-incorporation rates were higher than bacterial rates after P addition (Fig. 6). This P-incorporation resulted in a substantial enhancement of algal abundance (Carrillo *et al.*, submitted). P-enriched algae pursuing a net growth strategy may produce and release enough organic material to meet the P-requirements of bacteria, obviating the need to compete for inorganic nutrients (Caron *et al.*, 2000). In this scenario, bacteria would function as a carbon-link to other trophic levels (ciliate development during thaw experiment) (Fig. 6). In contrast, growth of P-deficient algae (mid-summer) after nutrient pulse generates algal “bloom-growth” that limits the release of organic carbon (Villar-Argaiz *et al.*, 2002a), thereby intensifying the mutualism-commensalism relationship and producing a tendency to bacterial P impoverishment (to see also Caron *et al.*, 2000) that ultimately restricts the development of the microbial loop (Fig. 6).

When bacteria were P-deficient (N:P \geq 20-24, late ice-free period), bacterial P-incorporation rates were significantly higher than algal rates (Fig. 6). Bacterial and primary production, were stimulated after the nutrient pulse, and competition for P was the prevalent relationship between algae and bacteria (Villar-

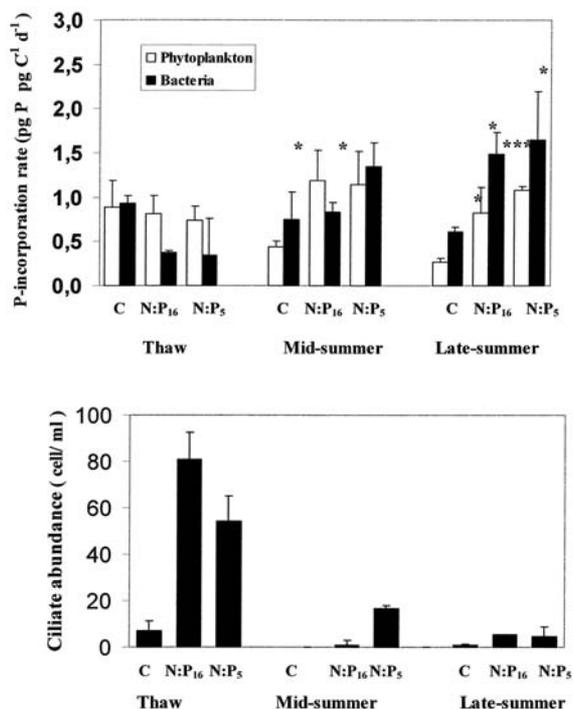


Figure 6. Phytoplankton and bacterial P-incorporation rates in un-enriched (control) and enriched (N:P₁₆ and N:P₅) treatments after short-term incubations [24 h in thaw experiment and 48 h in mid and late ice-free period experiments] (upper panel). Ciliate abundance in un-enriched (control) and enriched (N:P₁₆ and N:P₅) treatments after 15 days incubation period in the thaw, mid and late ice-free period experiments (lower panel). Error bars indicate means \pm SD. Asterisks indicate significantly higher values relative to the controls (one-tailed t-test). * p <0.05; ** p <0.01; *** p <0.001. *Tasas de incorporación algal y bacteriana de fósforo en tratamientos no enriquecidos (control) y enriquecidos en fósforo (N:P₁₆ and N:P₅) [después de 24 horas de incubación en el deshielo y 48 horas en mitad y al final del periodo libre de hielo] (gráfica superior). Abundancia de Ciliados en tratamientos no enriquecido (control) y enriquecido en fósforo (N:P₁₆ and N:P₅) al final de 15 días de incubación en el deshielo, mitad y final del periodo libre de hielo (gráfica inferior). Las barras de error son medias \pm SD. Los asteriscos indican valores significativamente mayores en relación con el control (t-test). * p <0.05; ** p <0.01; *** p <0.001.*

Argaiz *et al.*, 2002a). In spite of the addition of P, bacterial elemental composition did not reach the threshold for balanced growth (Fig. 6). To summarize, the algae-bacteria interaction shifted from mutualism during most of the ice-free period to competition for available P towards the end of the ice-free period. Therefore, limitation by mineral nutrients can be a transient phe-

nomenon that lasts for a shorter time than the total turnover of the system (Daufresne & Loreau, 2001). This interpretation is further sustained by seasonal and inter-annual observations. First, the algal elemental composition is a good predictor of algal growth rate, whereas the bacterial N:P ratio is not related to bacterial growth rate (Fig. 7). This lack of relationship indicates that bacteria are limited by an element other than P, despite the strong P-limitation in La Caldera Lake (DIN:TP>40). Second, there is a coupled relation between bacterial and primary production based on a direct dependence on the organic carbon released by algae (see above) (Fig. 3). The prevalence of commensalism over competition contrasts with the clear bacterial P-limitation established for other oligotrophic ecosystems (Rivkin & Anderson, 1997; Vrede, 1999; Caron *et al.*, 2000; Carlsson & Caron 2001) and with the Light Nutrient Hypothesis (LNH) (Sterner *et al.*, 1997) on the structure and functioning of aquatic ecosystems. We consider that the main difference with our approach is that the above authors did not include irradiance quality effects within the predictive conceptual framework of the LNH. This is especially important because we have shown that UVR favours algae-bacteria commensalism. Therefore, we can affirm that mutualism-commensalism is more prevalent than competition between algae and bacteria.

However, given that bacteria can tolerate UVR-stress and respond positively to a shift in EOC-availability (basis of the predominantly mutualistic relationship), the question arises as to why they do not reach higher development if C exceeds bacterial requirements (Carrillo *et al.*, 2002; Medina-Sánchez *et al.*, 2006) and mineral nutrients do not strongly limit their growth (Villar-Argaiz *et al.*, 2002a). Another relevant question is why bacterial production, C-assimilation and use efficiencies show such low or even decreased values (Carrillo *et al.*, 2002; Medina-Sánchez *et al.*, 2004) after the addition of P (Medina-Sánchez *et al.*, 2002; 2006). It appears that other controls are responsible for the scarce abundance and activity of bacteria in La Caldera Lake.

This paradoxical scenario calls for a shift from a simple trophic to a metabolic approach, whereby the “algal” community is divided into strict-autotrophs or mixotrophs.

THE MICROBIAL LOOP DOES NOT DEVELOP. ARE ALGAE ANIMALS OR PLANTS?

The traditional approach to the energy and matter that flows through an ecosystem involves the classification of organisms into two main trophic groups: osmotrophs/phototrophs (primary producers) and phagotrophs/heterotrophs (consumers). This basic division does not always match the reality of the microbial world because some organisms, such as protista, combine both

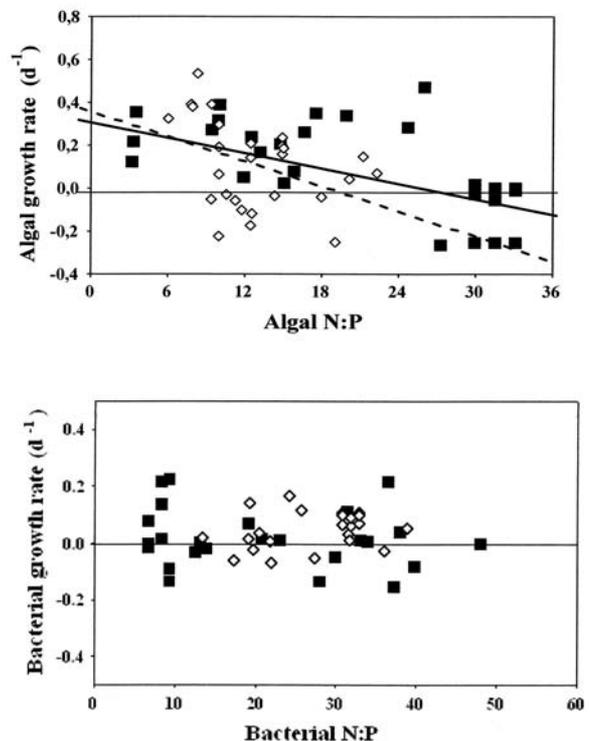


Figure 7. Relationship between algal (upper panel) and bacterial (lower panel) growth rate and elemental composition (N:P ratio) of algae and bacteria, respectively, in nutrient-pulse experiments for the mid-and late ice free period experiments. *Relación entre la tasa de crecimiento y la composición elemental (razón N:P) de algas (gráfica superior) y bacterias (gráfica inferior), en experimentos de nutrientes pulsados llevados a cabo en la mitad y final del periodo libre de hielo.*

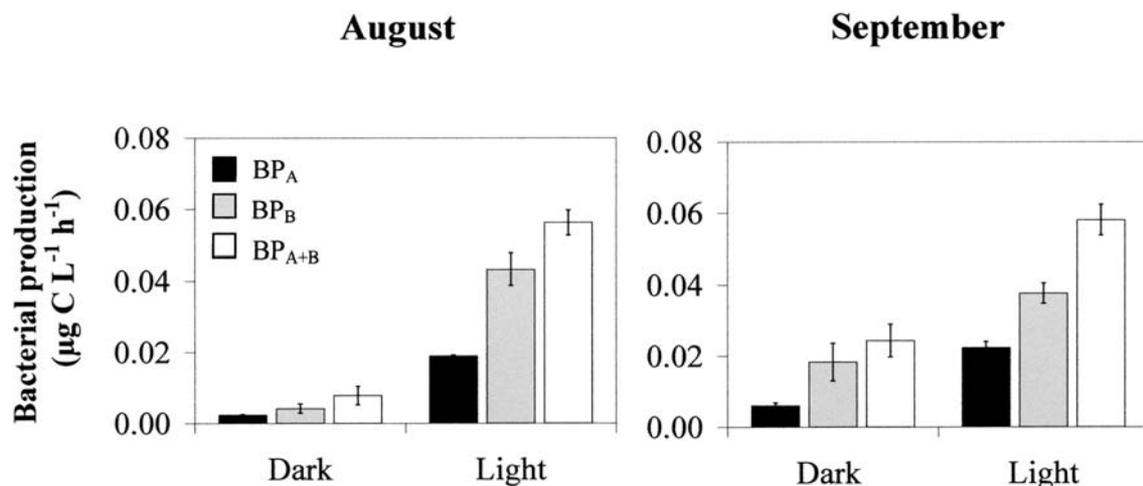


Figure 8. Bacterial production measured as BPA, BPB, and BPA+B in light treatments ('dark' and 'full light') in different moments during the ice-free period. August and September. BPA, Bacterial production incorporated in algal fraction; BPB, bacterial production not incorporated by algae; BPA+B, bacterial production in both the algal and bacterial fractions. Error bars indicate means \pm SD. *Producción bacteriana medida como BPA, BPB y BPA+B en tratamientos de luz (oscuridad y luz completa) en distintos momentos durante el periodo libre de hielo. BPA, Producción bacteriana incorporada en la fracción algal, BPB, Producción bacteriana no incorporada por las algas; BPA+B, Producción bacteriana en las fracciones algales y bacterianas juntas. Las barras de error son medias \pm SD.*

trophic abilities and are designated mixotrophs (Sanders, 1991; Jones, 1997). By definition, a mixotroph is an organism in which both photosynthesis and phagotrophy are possible (Sanders, 1991). Although mixotrophic metabolism is an energetically more demanding strategy (maintaining both photosynthetic and phagotrophic systems in the same cell) compared with strictly autotrophic or heterotrophic metabolism, mixotrophy has evolved as an adaptive strategy for growth in adverse conditions.

Mixotrophs constitute a functional type usually present in oligotrophic and dystrophic lakes (Salonen & Jokinen, 1988; Jansson *et al.*, 1996) and at high latitudes (Duthie & Hart, 1987; Eloranta, 1989, 1995; Lepistö & Rosenström, 1998), and they are frequently dominant in the phytoplanktonic community of high mountain lakes (Sánchez-Castillo *et al.*, 1989; Carrillo *et al.*, 1990 b, 1991 a, b, 1995; De Hoyos *et al.*, 1998; Straskrabová *et al.*, 1999) and in marine and freshwater ecosystems (Raven, 1997; Sanders *et al.*, 2000; Sherr & Sherr, 2002).

The dual nature of mixotrophic (autotrophic and heterotrophic) metabolism offers an advantage in ecosystems where light or mineral nutrients

are scarce, since the phagotrophic ability enables supplementation of autotrophic growth with the extra C and nutrient contents of prey, e.g., picoplankton (Rothhaupt, 1996 a, b; Raven, 1997). In this way, mixotrophs can constitute a key functional type for the structure of the community and the flow of energy through the food web. They not only contribute to the input of autochthonous C into the ecosystem but also, by their interaction with picoplankton, constitute a mechanism for the transfer of nutrients and energy towards upper trophic levels, such as zooplankton (Thingstad *et al.*, 1996; Sherr & Sherr, 2002). On the other hand, it was recently shown that mixotrophy is an adaptive strategy both to deficient environmental conditions and also to excess-induced stress conditions. Thus, Medina-Sánchez *et al.*, (2004) found higher bacterial consumption rates by mixotrophs exposed to stressing-sunlight (UVB+UVA+PAR) (Fig. 8), suggesting that bacterivory by mixotrophs allows them to acquire organic carbon and mineral nutrient contents from their prey under inhibitory light conditions (+UVR) for C fixation (Carrillo *et al.*, 2002) and acquisition of dissolved nutrients (Hessen *et al.*, 1995; Döhler, 1997).

With this background, we propose that mixotrophy is an adaptive strategy of “algae” to stress by ultraviolet radiation (Medina-Sánchez *et al.*, 2004). Mixotrophy implies a strong predatory control on bacteria, especially on the most active bacteria (Sherr & Sherr, 2002).

Since algae act both as predators of, and C suppliers bacteria (dual control), we estimated the relative importance of Predation vs. Resource control by defining a new variable. This variable, designated “algal control”, is the ratio between the potential mixotrophic consumption of bacterial production and the photosynthetic C supplied (i.e. $PMC_{BP}:EOC$, Medina-Sánchez *et al.*, 2004). Figure 9 shows the strong inverse relationship between bacterial production and “algal control” when the algal community is dominated by mixotrophs. Therefore, predation by mixotrophs accounts for the scarcity of bacteria (10^5 cell/ml, lowest end of the values recorded for oligotrophic ecosystems; Cotner & Biddanda, 2002) and the low carbon use efficiency by bacteria.

Hence, a complex regulation is established between algae and bacteria, namely “neither

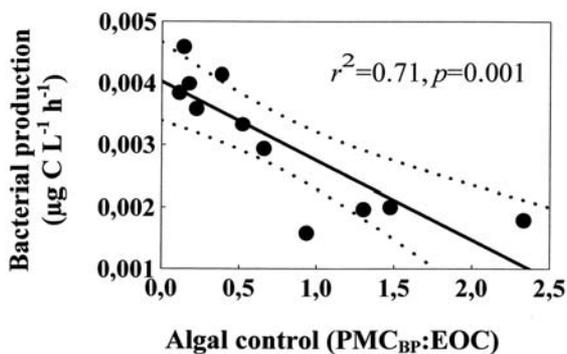


Figure 9. Intraannual relationship between bacterial production and ‘algal control’ measured as the ratio between potential mixotrophic consumption of bacterial C and photosynthetic C supplied by algae (i.e. $PMC_{BP}:EOC$). Dashed lines indicate 95 % confidence intervals around the fitted regression line (solid line). *Variación estacional en la relación entre la producción bacteriana y el ‘control algal’ medido como la razón entre consumo de carbono bacteriano por potenciales mixótrofos y el carbono fotosintético aportados por las algas (i.e. $PMC_{BP}:EOC$). Líneas discontinuas indican el intervalo de confianza del 95 % en relación con la línea de ajuste por regresión (línea continua).*

with nor without you” (Medina-Sánchez *et al.*, 2004). Bacteria are preyed upon by algae (“with you I die”) and simultaneously depend on the organic carbon released by the algae (“without you I cannot live”) (Carrillo *et al.*, 2002; Medina-Sánchez *et al.*, 2002, 2004). According to Thingstad *et al.* (1996), mixotrophs can take advantage in a situation where phytoplankton contribute the organic substrate that bacteria demand, while bacteria can favourably compete with algae for mineral nutrients (Currie & Kalff, 1984; Cotner & Wetzel, 1992) and finally, mixotrophs consume these enriched bacteria. Thus, when sunlight is intense and mineral nutrients are scarce, mixotrophic algae are able to grow because they can feed bacteria with “cheap” organic carbon and consume the “expensive” mineral nutrients packaged in the bacteria (*sensu* Thingstad *et al.*, 1996).

This scenario is consistent with (i) the ability of algae to regulate photosynthetic C release depending on their elemental composition (Villar-Argaiz *et al.*, 2002a); (ii) the ability of bacteria to grow under high full-sunlight intensities (Carrillo *et al.*, 2002, Medina-Sánchez *et al.*, 2002); and (iii) the decrease in bacterial production in the presence of algae after a nutrient pulse (Medina-Sánchez, 2002, 2006). Furthermore, mixotrophs can retain or release the P contained in bacteria depending on their predominant nutrition mode (autotrophic vs. phagotrophic) and nutrient requirements (Rothhaupt, 1997).

These abilities can constitute mechanisms that modulate the competition with bacteria for mineral nutrients, suggesting a higher complexity of the algae-bacteria interaction where algae exert the main control. Knowledge of this complex algae-bacteria interaction further advances the understanding of the planktonic structure and energy flow in this kind of aquatic ecosystem:

- 1) Thus, mixotrophs can overcome the harmful effects of UV radiation and displace: (i) strict autotrophs, less competitive at low concentrations of dissolved mineral nutrients (Rothhaupt, 1996a; Christaki *et al.*, 1999), (ii) picoplankton autotrophs, better adapted to the uptake of dissolved nutrient at low con-

centrations but more sensitive to the inhibitory effect of UVR (Callieri *et al.*, 2001), and (iii) heterotrophic nanoflagellates (HNF), which depend on a higher minimum threshold of bacterial abundance compared with mixotrophs (Havskum & Riemann, 1996).

- 2) With respect to the energy flow, mixotrophic bacterivory acts as a “by-pass” for the flow of energy and nutrients towards the grazing chain (Medina-Sánchez *et al.*, 2004). Hence, mixotrophs would occupy the niche of microheterotrophs and produce a weakening of the traditional microbial loop (Medina-Sánchez *et al.*, 2004). Besides, mixotrophic metabolism may improve the efficiency for the transfer of energy towards upper trophic levels, because the number of trophic steps has been reduced due to the by-pass from the heterotrophic microbial chain to the grazing chain (Fig. 10) (Rivkin & Anderson, 1997; Medina-Sánchez *et al.*, 2004). Considering all of the above, mixotrophy may at least in part explain the scarce development of the microbial loop in La Caldera Lake.

Nevertheless, several important questions are raised by the fact that mixotrophy is an advantageous metabolism strategy for a system stressed by UVR and oligotrophy and that P-atmospheric input of Saharan origin are relatively frequent: How do allochthonous nutrient inputs affect mixotrophs? Do P pulses interact with UVR in modulating this metabolism? The addressing of these questions responds to the current demand for research into the interactive effects of multiple factors on the functioning and structure of ecosystems.

THE MICROBIAL LOOP DOES NOT DEVELOP. INTERACTION OF LIGHT AND NUTRIENTS?

The interaction between UVR and atmospheric nutrient inputs can affect species performance and their relationships in ways that cannot be predicted from single-factor analysis, since the effects of two or more factors are non-additive. The interaction among distinct factors can

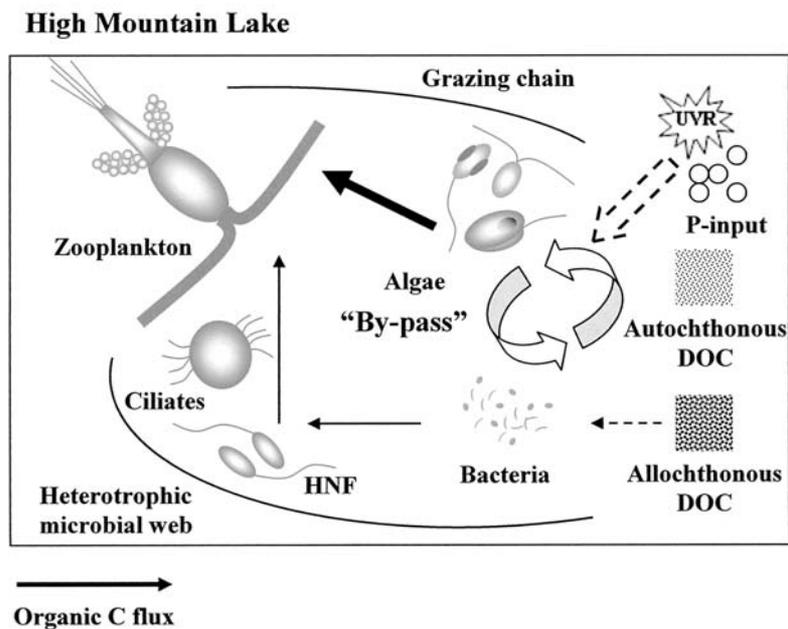


Figure 10. Proposed model of carbon flux through microbial food web in oligotrophic clear-water high mountain lakes. The curved arrows indicate reinforcement of the by-pass from bacteria to mixotrophs as a response to the interactive effect between UV radiation and the nutrient-pulses added. Thickness of arrows indicates the relative importance of C flux. *Modelo propuesto de flujo de carbono en la red trófica microbiana de lagos oligotróficos claros de alta montaña. Las flechas curvas indican refuerzo del corto circuito desde bacterias a mixótrofos como respuesta al efecto interactivo entre la radiación ultravioleta y los aportes pulsados de nutrientes. El grosor de las flechas indica la importancia relativa del flujo del carbono.*

either shift the sign of the effect of a single-factor (antagonistic effect) or, on the contrary, accentuate this effect (synergetic effect). Therefore, the analysis of multiple factors at different rates and scales is crucial for building a realistic model of the functioning of food webs (Breitburg *et al.*, 1999; Folt *et al.*, 1999; Xenopoulos *et al.*, 2003; Medina-Sánchez *et al.*, 2006). The increasing intensity and frequency of atmospheric material inputs from the Sahara desert to Sierra Nevada is associated with periodic climatic variations (North Atlantic Oscillation, NAO) and movements of the Intertropical Zone Convergence (ITZC). Hence, planktonic communities may be controlled not only by the “harmful” effect of UVR but also by the “stimulatory” effect of allocthonous inorganic nutrients. According to our initial hypothesis, P-pulses would reduce the negative effects that UVR exerts on algae, thereby decreasing the bacterivory rate (associated with mixotrophy). As a consequence, the nutrient pulses would modify the algae-bacteria relationship by shifting it from dual control (with bacterivory) to commensalism. Our investigations on short temporal scales showed that:

- 1) The elemental composition of algae and bacteria plays a key role in the ecological and biological response to light quality variation and nutrient availability. UVR x P interactions have an antagonistic effect on primary production, C-release by autotrophs, and bacterial production on both communities in un-enriched conditions. This antagonistic effect implies the attenuation or elimination of UVR effects.
- 2) The interactive effect between solar radiation and atmospheric nutrient inputs on the algae-bacteria link reinforces the dual control that algae exert on bacteria, i.e., it increases both the carbon released by algae and the mixotrophic predation on bacteria (Medina-Sánchez *et al.*, 2006). The by-pass is therefore enhanced, with consequences for the development of the microbial loop.

Finally, our results have implications at the ecosystem level. Thus, the attenuation of the

harmful effects of UVR by P-inputs may smooth UVR stress, which is relevant in a scenario of climatic change with increased UVB fluxes. This buffering effect would be more accentuated in clear-water ecosystems, where the main nutrient inputs have an atmospheric origin, such as in the tropical Atlantic Ocean (Prospero & Lamb, 2003, see also NOAA <http://toms.gsfc.nasa.gov/aerosol>). In the context of carbon flux, this buffering effect of harmful UVR at the base of both aquatic trophic webs (grazing chain and microbial loop) would favor the diversion of a higher proportion of released carbon flux through biotic webs, trapping carbon that would otherwise be susceptible to photolysis and be lost to the atmosphere.

In spite of the light thrown by this study on the processes that regulate the pelagic community of clear-water high mountain lakes, many areas remain in the shadows and new questions have arisen, briefly summarized as follows:

1. Do the results obtained over short time scales apply to long-term scales (days, weeks or months)?
2. How do UVR x nutrient interactive effects propagate along the grazing chain?
3. How universal are the patterns established for high mountain lakes with high UVR doses? Can structural tendencies established across trophic gradients be extrapolated to high-UVR-flux oligotrophic ecosystems?

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