

**From surface Chl a to phytoplankton  
size classes:  
a first step towards size-specific primary production**

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# Context of the Study

- **Utilization of ocean color**

- ⇒ Estimation of **phytoplankton biomass quantity**
- ⇒ Estimation of **primary production**

- **Requirements**

- ⇒ Understanding and prediction of **phytoplankton community composition:**

*Ciotti et al.* (JGR, 1999)

*Gregg et al.* (DSR, 2002)

*Iglesias-Rodriguez et al.* (GBC, 2002)

*Bouman et al.* (MEPS, 2003)

# Objective of the Study

**Surface Chla concentration**  
( $\text{Chla}_{\text{surf}}$ )

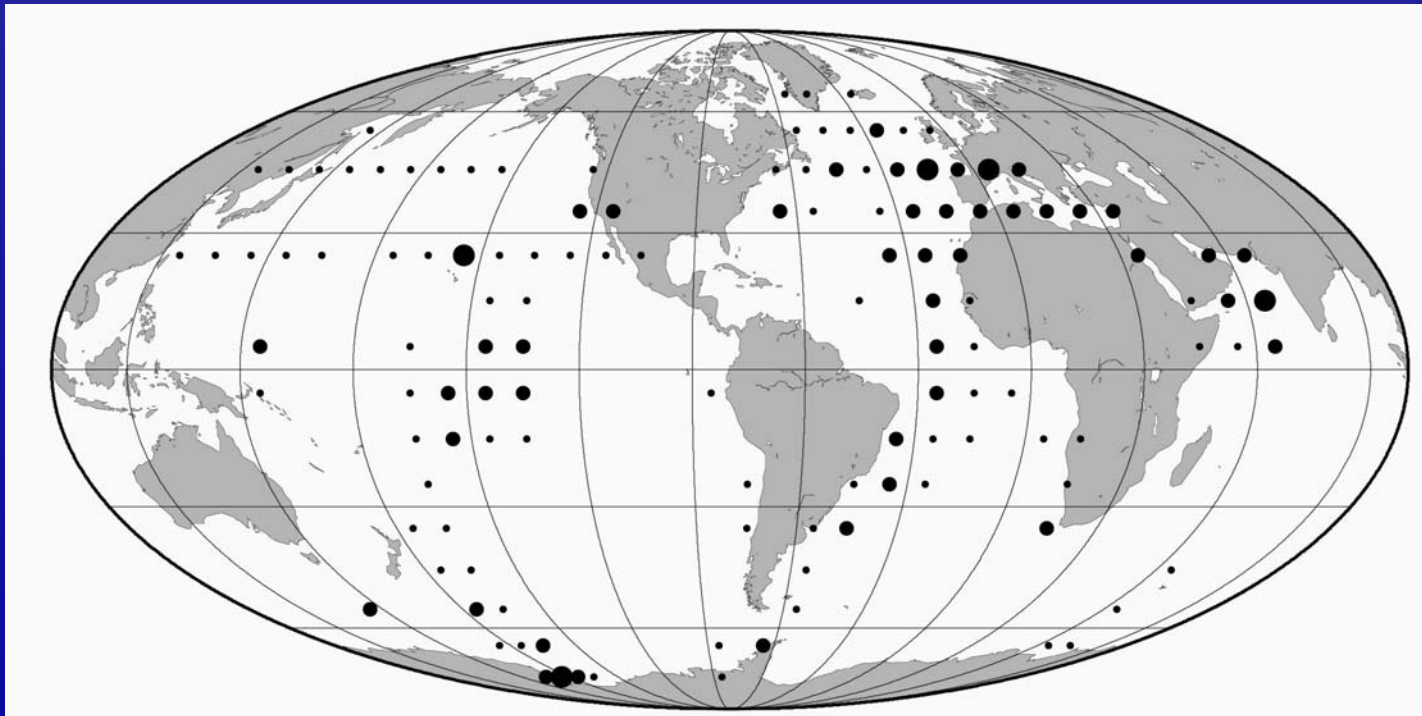


**Phytoplankton biomass quality:**

- Composition
- Vertical distribution

# Data

- **HPLC pigment database**: 2419 stations (1990-2002)




Number of stations per square of  $10^\circ \times 10^\circ$ ; small spots:  $< 10$  stations, medium spots: 10-100 stations, large spots:  $> 100$  stations.

# Method: from diagnostic pigment to phytoplankton classes

## ▪ Sum of the diagnostic pigments :

$$\Sigma = 1.4 \text{ Fuco} + 1.4 \text{ Peri} + 1.3 \text{ Hex} + 0.4 \text{ But} + 0.6 \text{ Allo} + 0.9 \text{ Zea} + 1.0 \text{ TChlb}$$

  
~ Micro > 20 μm                      ~Nano 2-20 μm                      ~ Pico < 2 μm

## ▪ Size classes percentage:

$$\% \text{micro} = 100 * (1.4 \text{ Fuco} + 1.4 \text{ Peri}) / \Sigma$$

$$\% \text{nano} = 100 * (1.3 \text{ Hex} + 0.4 \text{ But} + 0.6 \text{ Allo}) / \Sigma$$

$$\% \text{pico} = 100 * (0.9 \text{ Zea} + 1.0 \text{ TChlb}) / \Sigma$$

## ▪ Chla associated to each size class (sc-Chla):

$$\text{micro-Chla} = \% \text{micro} * \text{Chla}$$

$$\text{nano-Chla} = \% \text{nano} * \text{Chla}$$

$$\text{pico-Chla} = \% \text{pico} * \text{Chla}$$

*After : Claustre (L&O, 1994); Vidussi et al. (JGR, 2001)*

# Method: Standardization and Sorting of the profiles

- **Standardization of the sc-Chla profiles:**

$$\text{zeta} = z / Z_{\text{eu}}$$

$$\text{sc-Chla}(\text{zeta}) / \overline{C}_{Z_{\text{eu}}}$$

- **Interpolation of the dimensionless sc-Chla profiles:**

$$\text{zeta} = 0 \text{ (surface)} - 2 (2 * Z_{\text{eu}}) \longrightarrow 20 \text{ points / profile}$$

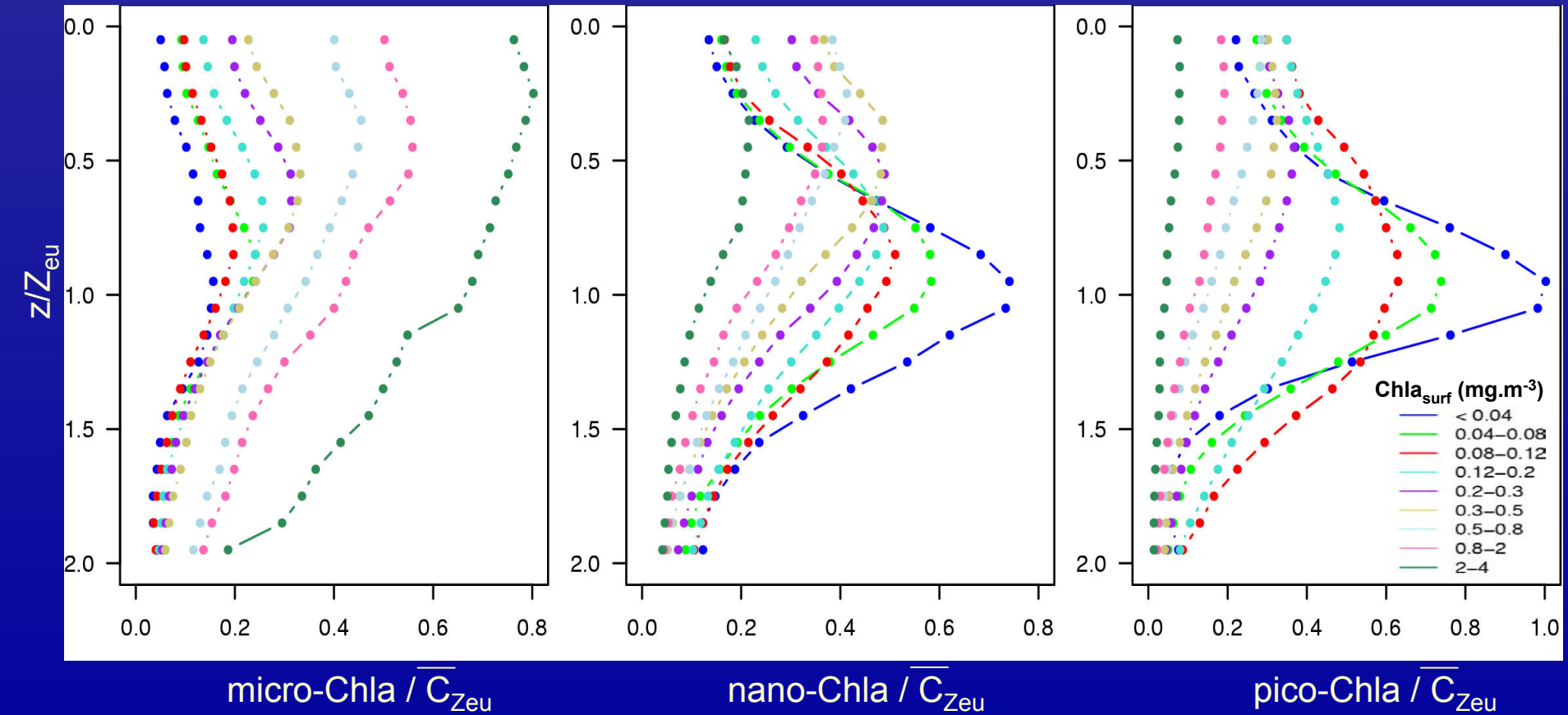
- **Sorting of the interpolated sc-Chla profiles:**

- ➔ according to the hydrological regime: stratified / mixed

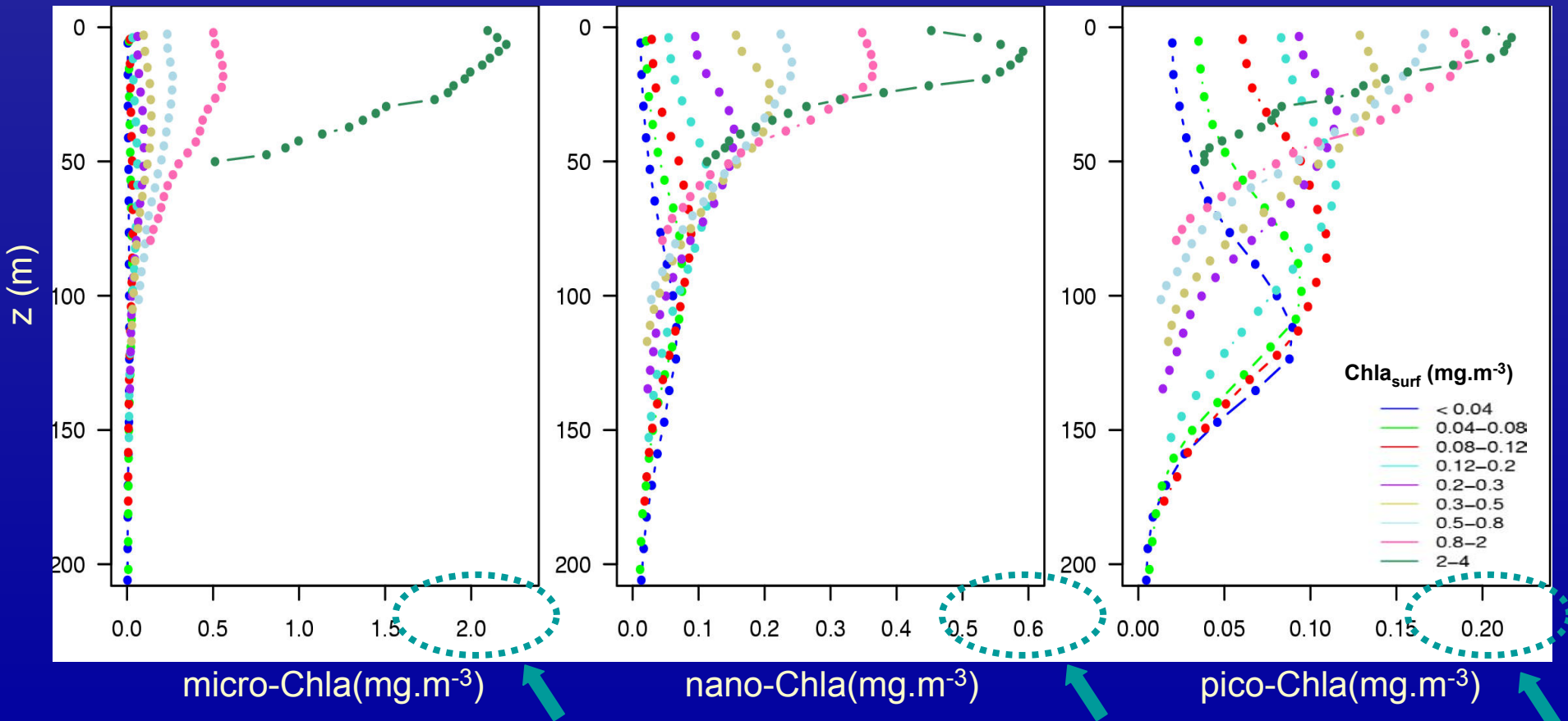
- ➔ according to  $\text{Chla}_{\text{surf}}$ : trophic classes

- **For each trophic class: computation of average sc-Chla profiles**

# Results: Dimensionless mean profiles



# Results: Mean « rescaled » profiles

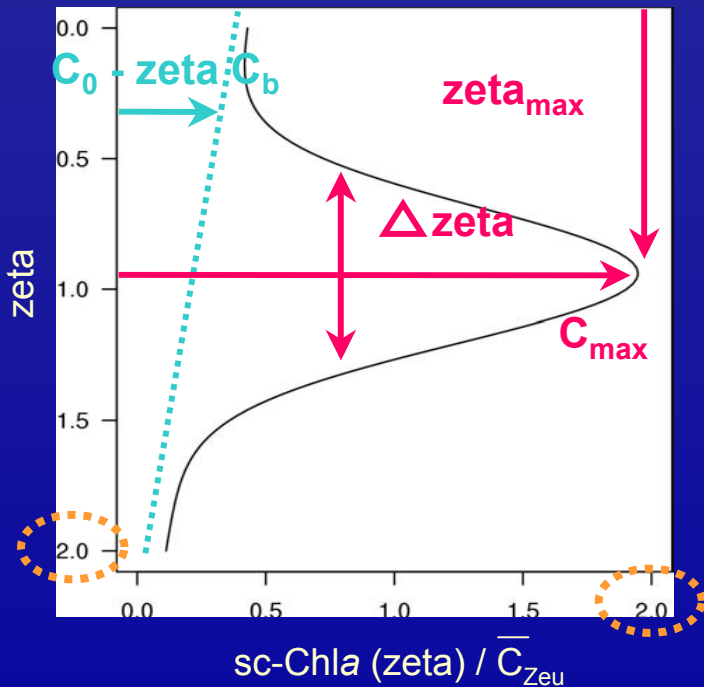




# Parameterization of the profiles

- Equation fitted to each mean sc-Chla profile:

$$\text{sc-Chla}(\text{zeta}) = \underbrace{C_0 - C_b \text{zeta}}_{\text{Background concentration}} + \underbrace{C_{\text{max}} \exp \left\{ - \left[ \frac{\text{zeta} - \text{zeta}_{\text{max}}}{\Delta \text{zeta}} \right]^2 \right\}}_{\text{Gaussian profile}}$$



Background concentration

Gaussian profile

- Computation of the 5 parameters:

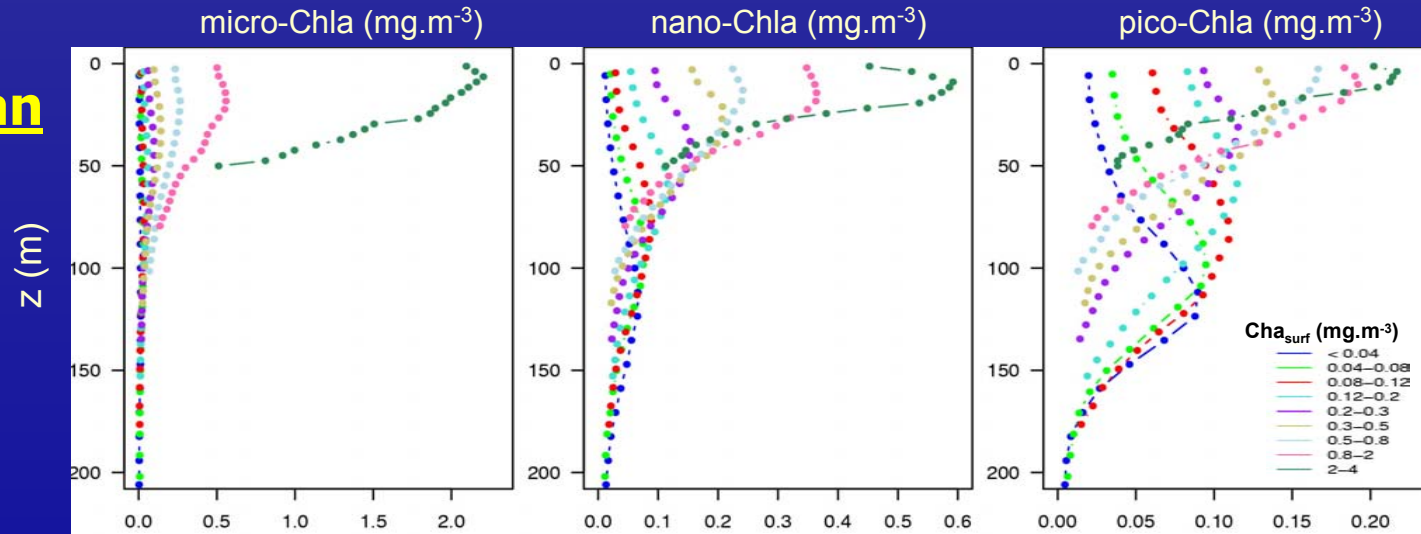
The adjustment procedure allows  $C_0$ ,  $C_b$ ,  $C_{\text{max}}$ ,  $\text{zeta}_{\text{max}}$ ,  $\Delta \text{zeta}$  to be calculated for each trophic class

- Parameterization used in a continuous manner:

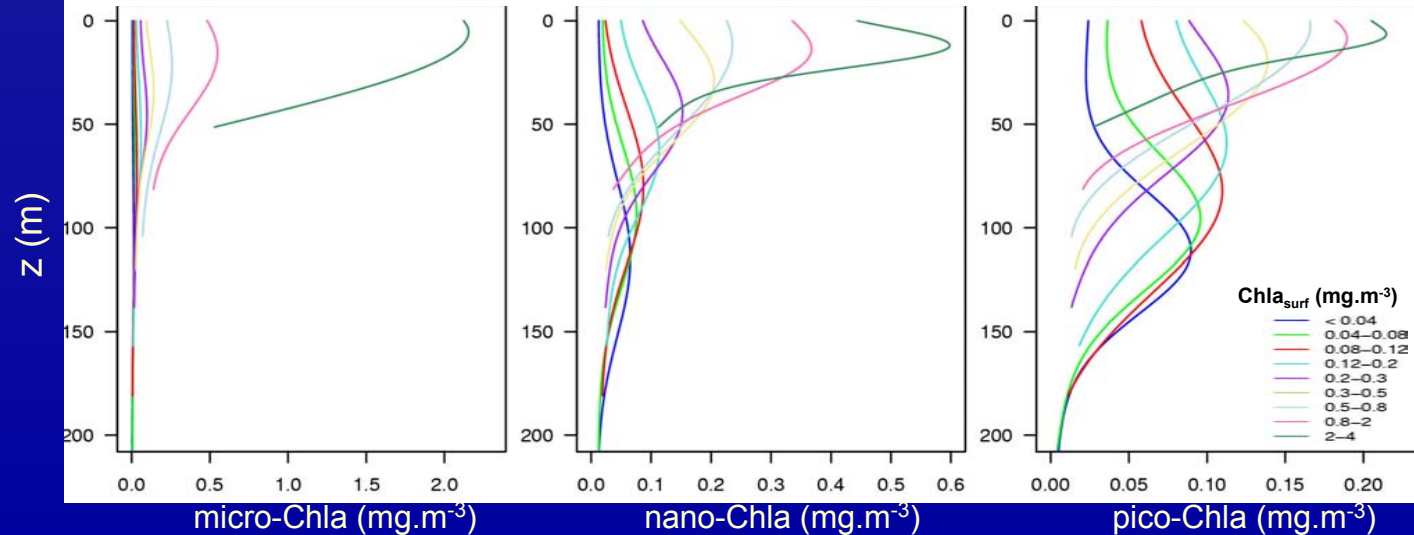
For any  $C_{\text{surf}}$  value, the 5 parameters can be retrieved using an interpolation between the discrete values specific to each trophic class

# Modeled / Measured « rescaled » profiles

**Measured mean profiles**



**Modeled profiles**

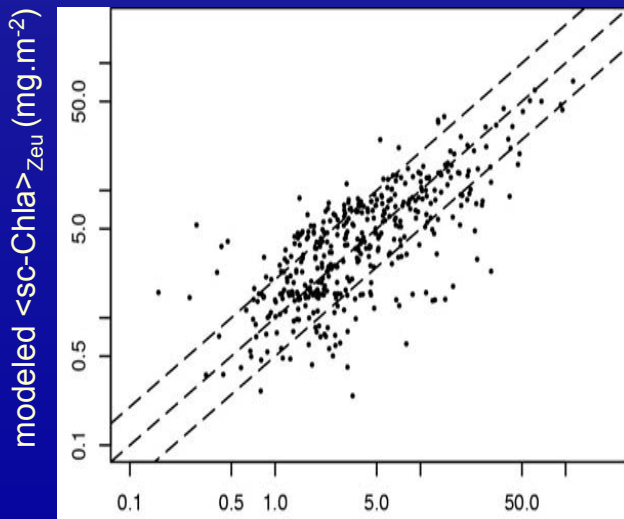


# Validation: Modeled vs measured integrated contents

- **HPLC database:**

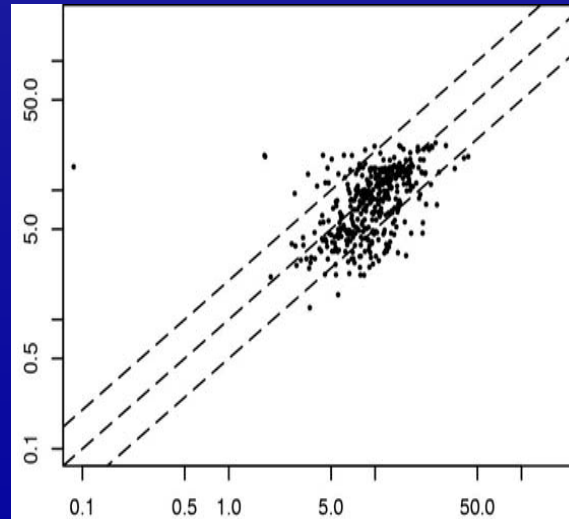
80% for the parameterization / 20% for the validation

Micro



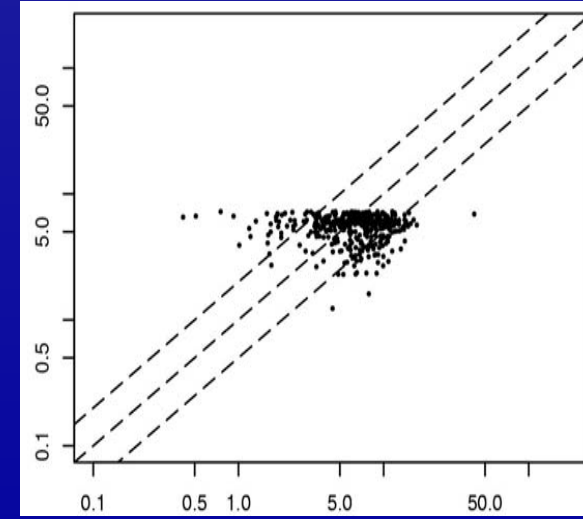
measured <sc-Chla><sub>Zeu</sub>  
(mg.m<sup>-2</sup>)

Nano



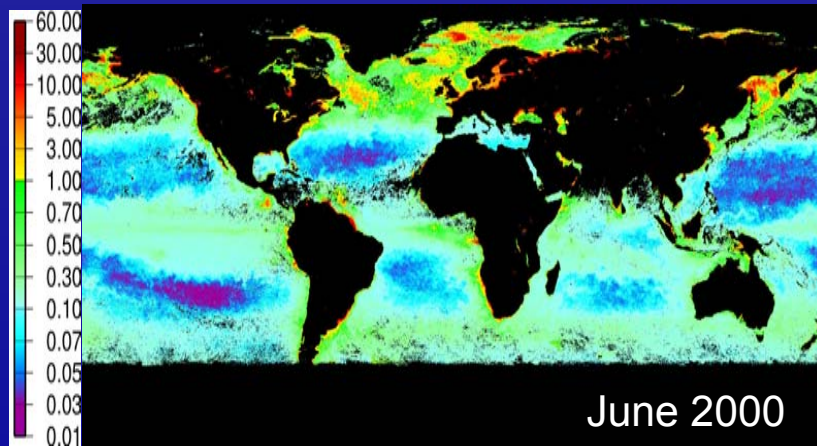
measured <sc-Chla><sub>Zeu</sub>  
(mg.m<sup>-2</sup>)

Pico



measured <sc-Chla><sub>Zeu</sub>  
(mg.m<sup>-2</sup>)

# Application: phytoplankton size classes climatology



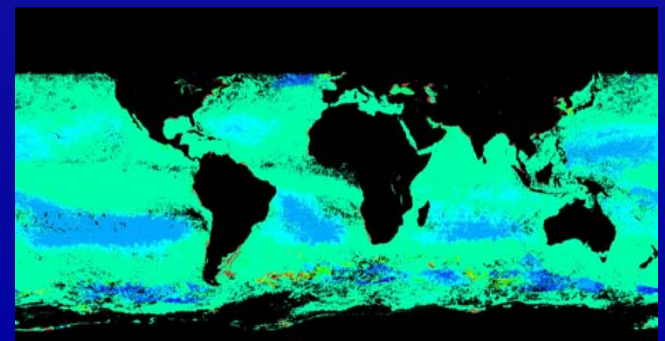
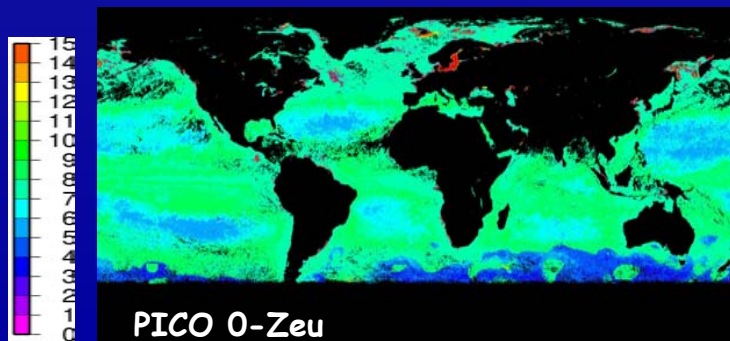
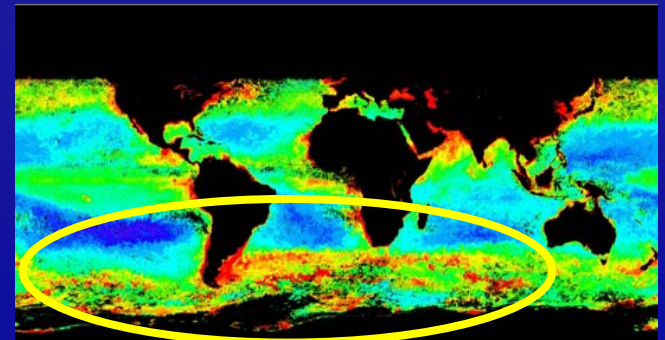
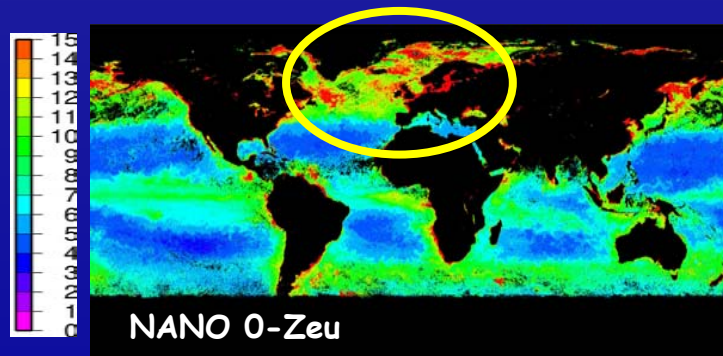
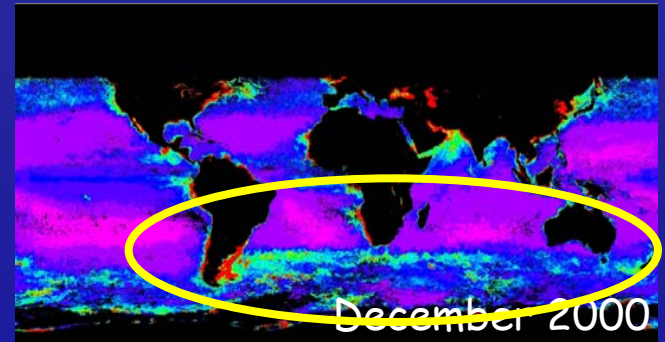
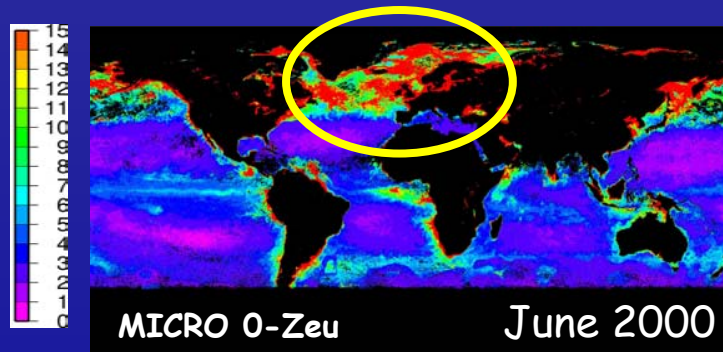
Map of monthly average SeaWiFS Chl<sub>a</sub><sub>surf</sub> (mg.m<sup>-3</sup>)

Statistical model



micro-, nano-, pico-Chla  
vertical profiles

# Application: phytoplankton size classes climatology



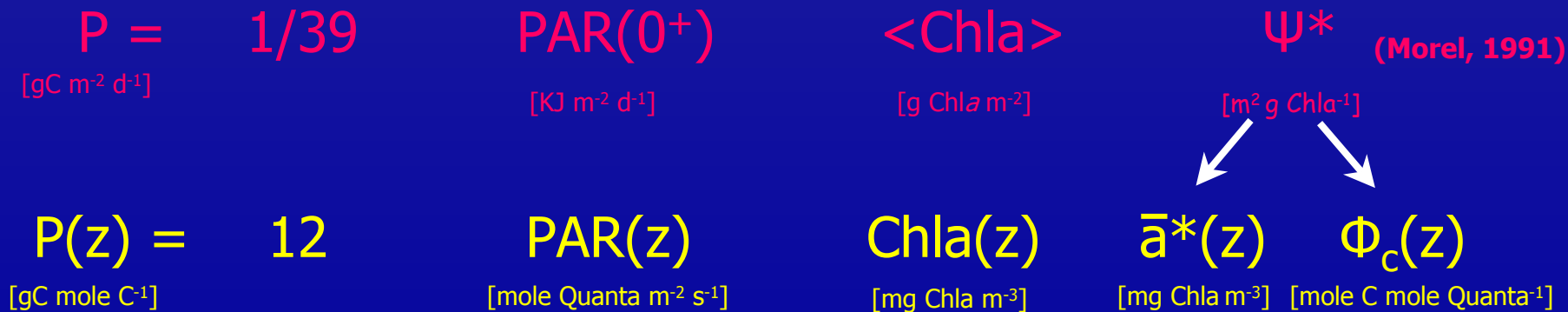
# Perspectives

- **Objective:**

From specific biomass fields (*e.g.* the one inferred from remote sensing) to size-specific primary production

- **How?**

Bio-optical models are a possible link, but currently they work with total algal biomass:



Derived from P vs E curves parameters



# Perspectives

$$P(z) = 12$$

$$PAR(z)$$

$$Chla(z)$$

$$\bar{a}^*(z)$$

$$\Phi_c(z)$$



$$P(z) = \text{micro-}P(z) + \text{nano-}P(z) + \text{pico-}P(z)$$

## Present study:

$$Chla(z) = \text{micro-}Chla(z) + \text{nano-}Chla(z) + \text{pico-}Chla(z)$$

## Question:

Are photo-physiological parameters phytoplankton **community composition dependant** ?

-> Is it possible to **index** some of the bio-optical models **parameters** on some **community composition** indexes?

# Perspectives

## How to check this?

Database comprising **HPLC** data + **P-I** curves parameters + **absorption** data

$$\bar{a} \stackrel{?}{=} \bar{a}_{\text{micro}}^* \text{ micro-Chla} + \bar{a}_{\text{nano}}^* \text{ nano-Chla} + \bar{a}_{\text{pico}}^* \text{ pico-Chla}$$

$$P_m \stackrel{?}{=} P_m^b \text{ micro-Chla} + P_m^b \text{ nano-Chla} + P_m^b \text{ pico-Chla}$$

$$\alpha \stackrel{?}{=} \alpha^b \text{ micro-Chla} + \alpha^b \text{ nano-Chla} + \alpha^b \text{ pico-Chla}$$

$$\rightarrow \Phi_{\text{Cmax}} = \alpha^b / \bar{a}^* \rightarrow \text{micro } \Phi_{\text{Cmax}}, \text{ nano } \Phi_{\text{Cmax}}, \text{ pico } \Phi_{\text{Cmax}}$$

## If it works ...

$$P(z) = \text{micro-P}(z) + \text{nano-P}(z) + \text{pico-P}(z)$$

$$\frac{39 P}{\text{PAR}(0^+)} = \langle \text{Chla} \rangle \Psi^* \quad \curvearrowright$$

$$= \langle \text{micro-Chla} \rangle_{\text{micro}} \Psi^* + \langle \text{nano-Chla} \rangle_{\text{nano}} \Psi^* + \langle \text{pico-Chla} \rangle_{\text{pico}} \Psi^* \quad \curvearrowright$$



# Preliminary results

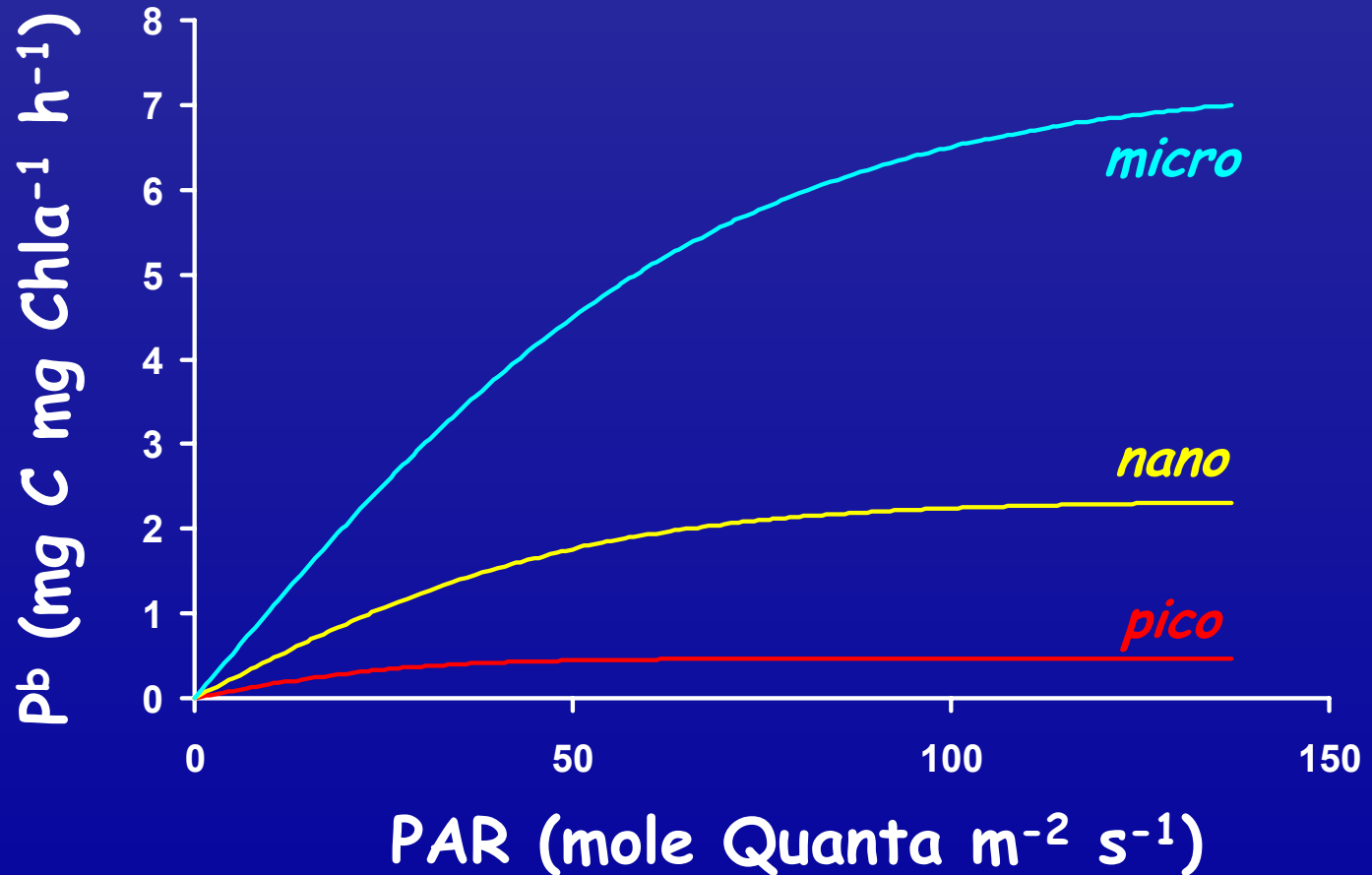
## Results from the POMME dataset (H. Claustre, unpublished data):

	$\bar{a}^*$ m <sup>2</sup> mg Chla <sup>-1</sup>	$P_{\max}^b$ mg C mg Chla <sup>-1</sup> h <sup>-1</sup>	$\alpha^b$ $\frac{\text{mg C mg Chla}^{-1} \text{ h}^{-1}}{\text{mole Quanta m}^{-2} \text{ s}^{-1}}$	$\Phi_{C\max}$ $\frac{\text{mole C}}{\text{mole quanta}}$
micro	0.020 ± 0.002	7.27 ± 0.66	0.105 ± 0.660	0.125
nano	0.021 ± 0.001	2.33 ± 0.24	0.046 ± 0.004	0.051
pico	0.037 ± 0.001	0.47 ± 0.29	0.017 ± 0.005	0.011
	R <sup>2</sup> = 0.91 n = 344	R <sup>2</sup> = 0.73 n = 344	R <sup>2</sup> = 0.72 n = 344	

! 0.125 is a theoretical limit !....

# Preliminary results

*The derived « specific » P vs E curves*



*Low for picophytoplankton....*

# Acknowledgements

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