Retrieval of phytoplankton pigments from underway spectrophotometry in the Fram Strait, Arctic Ocean

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- Introduction

Phytoplankton pigment databases have been extensively used in developing, validating or refining bio-optical algorithms for estimating phytoplankton biomass and functional types.
Here, we investigate the performances of two approaches, i.e. Gaussian decomposition and singuar value decomposition combined with non-negative least squares (SVD-NNLS) in determining the concentrations of either individual pigments or pigment groups from *a*_p(λ) obtained from underway AC-S flow-

Data Collection & Processing (Liu et al., 2018, and references therein) PS93.2 PS99.2 PS107 **Underway AC-S flow-through system** AC-S valve seawater__ debubble controller c tube AC-S data processing to derive $a_{ph}(\lambda)$ $a_{p}(\Lambda)\&c_{p}(\Lambda)$ calculation Spikes TS correction 1-min bin removal ר<u></u> 10⁻ ב

- through system in the Fram Strait.
- ✓ The effect of package effect on the retrieval accuracy was assessed by including a normalization term in $a_{ph}(\lambda)$ (see below).

Gaussian Decomposition

(Chase et al., 2013)

- ✓ $a_p(\lambda)$ was decomposed into 12 Gaussian functions + 1 NAP power law function.
- ✓ Gaussian amplitudes were related to the concentrations of TChla, TChl-b, Chl-c1/2, PSC and PPC.

✓ $a_{ph}(\lambda)$ normalized by package effect: $\stackrel{\land}{a_{ph}}(\lambda) = a_{ph}(\lambda) \frac{0.033 \times TChl - a}{a_{ph}(675)}$







Statistics (leave-one-out cross validation)

✓ relative percentage different (RPD). $RPD = \frac{1}{n} \sum_{i=1}^{n} \frac{C_i^{esti} - C_i^{meas}}{C_i^{meas}} \times 100\%$

✓ **bold**: pigments with better retrieval accuracy after applying the package effect

normalization to $a_{\rm ph}(\lambda)$.

Pigments	Gaussian Decomposition		SVD-NNLS-5		SVD-NNLS-18	
	non-normalized $a_{ph}(\lambda)$	normalized $a_{ph}(\lambda)$	non-normalized $a_{ph}(\lambda)$	normalized $a_{ph}(\lambda)$	non-normalized $a_{ph}(\lambda)$	normalized a _{ph} (λ)
TChl-a	11.9%	2.3%	6%	-0.4%	7.2%	-0.2%
TChl-b	15.3%	12.0%	53.3%	39.1%	93.8%	88.8%
Chlc_1/2	39.8%	33.6%	59.1%	63.1%	163.5%	170.5%
PSC	49.6%	27.6%	34.8%	44.5%	-	-
PPC	33.8%	15.2%	42.2%	37.5%	-	-
Chl_c3	-	-	-	-	280.2%	202.6%
Allo	-	-	-	-	28.2%	37.2%
a_Caro	-	-	-	-	102.0%	71.2%
β_Caro	-	-	-	-	51.0%	48.5%
Diadino	-	-	-	-	41.5%	47.3%
Diato	-	-	-	-	37.1%	60.2%
Fuco	-	-	-	-	56.0%	49.8%
Hex	-	-	-	-	67.1%	51.5%
But	-	-	-	-	249.1%	97.2%
Neo	-	-	-	-	14.8%	12.3%
Lut	-	-	-	-	29.4%	36.2%
Peri	-	-	-	-	64.1%	97.6%
Prasino	-	-	-	-	6.0%	4.4%
Viola	-	-	-	-	56.9%	71.0%
Zea	-	-	-	-	39.2%	39.7%
Others	-	-	49.5%	26.9%	-	-

Conclusion

 Gaussian decomposition was capable of estimating TChl-a, TChl-b, Chl-c1/2, PPC and PSC with a prediction error of less than 50% and outperformed SVD-NNLS in retrieving TChl-b, Chl-c1/2 and PPC.

 ✓ SVD-NNLS enabled the retrieval of a series of phytoplankton pigments with defined

uncertainty (RPD ranges 6-280%).

✓ Lower uncertainties for the retrieval of all the five pigments using Gaussian decomposition and of 9 types of pigments using SVD-NNLS were obtained with the combined use of observed $a_{ph}(\lambda)$ and TChl-a concentration that partially accounts for the package effect across the whole absorption spectra.

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