

- Phytoplankton pigment assemblages in the open ocean identified using hyperspectral data
- Potential usefulness of derivative spectroscopy of absorption and remote-sensing reflectance
- Cluster-based approach to identify the most suitable spectral ranges and derivative parameters

Cluster analysis of hyperspectral optical data for discriminating phytoplankton pigment assemblages in the open ocean

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Abstract

1 Optical measurements including remote sensing provide a potential tool for the identification of
2 dominant phytoplankton groups and for monitoring spatial and temporal changes in biodiversity
3 in the upper ocean. We examine the application of an unsupervised hierarchical cluster analysis
4 to phytoplankton pigment data and spectra of the absorption coefficient and remote-sensing
5 reflectance with the aim of discriminating different phytoplankton assemblages in open ocean
6 environments under non-bloom conditions. This technique is applied to an optical and
7 phytoplankton pigment data set collected at several stations within the eastern Atlantic Ocean,
8 where the surface total chlorophyll-*a* concentration (TChl*a*) ranged from 0.11 to 0.62 mg m⁻³.
9 Stations were selected on the basis of significant differences in the ratios of the two most
10 dominant accessory pigments relative to TChl*a*, as derived from High Performance Liquid
11 Chromatography (HPLC) analysis. The performance of cluster analysis applied to absorption and
12 remote-sensing spectra is evaluated by comparisons with the cluster partitioning of the
13 corresponding HPLC pigment data, in which the pigment-based clusters serve as a reference for
14 identifying different phytoplankton assemblages. Two indices, cophenetic and Rand, are utilized
15 in these comparisons to quantify the degree of similarity between pigment-based and optical-
16 based clusters. The use of spectral derivative analysis for the optical data was also evaluated, and
17 sensitivity tests were conducted to determine the influence of parameters used in these
18 calculations (spectral range, smoothing filter size, band separation). The results of our analyses
19 indicate that the second derivative calculated from hyperspectral (1 nm resolution) data of the
20 phytoplankton absorption coefficient, $a_{ph}(\lambda)$, and remote-sensing reflectance, $R_{rs}(\lambda)$, provide
21 better discrimination of phytoplankton pigment assemblages than traditional multispectral band-
22 ratios or ordinary (non-differentiated) hyperspectral data of absorption and remote-sensing
23 reflectance. The most useful spectral region for this discrimination extends generally from

24 wavelengths of about 425 - 435 nm to wavelengths within the 495 - 540 nm range, although in
25 the case of phytoplankton absorption data a broader spectral region can also provide satisfactory
26 results.

27

28 **1. Introduction**

29 In situ and remotely-sensed optical observations of ocean waters provide information
30 regarding the concentrations of optically significant constituents in seawater, and offer the ability
31 to observe important biological and biogeochemical variables (e.g., Chang et al., 2006).
32 Numerous studies over the past three decades have focused on the development of bio-optical
33 algorithms linking measurable optical properties to the primary pigment in phytoplankton,
34 chlorophyll-*a*, a proxy for the phytoplankton biomass (e.g., Morel, 1988; Bricaud et al., 1998;
35 O'Reilly et al., 2000; Reynolds et al., 2001). In recent years, efforts to expand the use of optical
36 measurements for estimating other biogeochemically important ocean variables and phenomena
37 have increased considerably. For example, optical measurements including satellite remote
38 sensing have been used to detect harmful algal blooms (Cullen et al., 1997; Stumpf et al., 2003),
39 surface concentrations of particulate inorganic and organic carbon (Balch et al., 2005; Stramski
40 et al., 2008), particle size distribution (Kostadinov et al., 2009), phytoplankton community
41 composition and size structure (Alvain et al., 2005; Uitz et al., 2006; Nair et al., 2008; Ciotti and
42 Bricaud, 2006; Aiken et al., 2007), and phytoplankton class-specific primary production (Uitz et
43 al., 2010).

44 Recent advances in measuring ocean optical properties and light fields within and leaving the
45 ocean have included a progressive shift from using multispectral to high spectral resolution
46 (hyperspectral) acquisition systems (Chang et al., 2004). New technologies and the
47 miniaturization of electro-optical components have permitted the development of accurate, low-
48 cost, and energy-efficient hyperspectral sensors suitable for deployments from a variety of
49 platforms such as in-water vertical profiling systems, moorings, drifters, autonomous vehicles,
50 air-borne and space-borne platforms (Perry & Rudnick, 2003; Dickey et al., 2006). The
51 capability to obtain measurements at hundreds of narrow and closely spaced wavelength bands
52 from the ultraviolet to near-infrared, with a resolution better than 10 nm, has become one of the

53 most powerful and fastest growing areas of technology in the field of ocean optics.

54 Hyperspectral optical data provide the opportunity for improvements in spectral shape
55 analysis and subsequent extraction of environmental information compared with low spectral
56 resolution optical data. Derivative spectroscopy is one powerful technique of spectral shape
57 analysis which enhances subtle features in hyperspectral data, and has been used successfully to
58 obtain information about optically significant water constituents. For example, Craig et al.
59 (2006) assessed the feasibility of detection of a toxic bloom of the dinoflagellate *Karenia brevis*
60 from the analysis of the fourth derivative of phytoplankton absorption spectra, estimated from in
61 situ hyperspectral measurements of remote-sensing reflectance $R_{rs}(\lambda)$ (λ is light wavelength in
62 vacuo). The advantages offered by hyperspectral measurements of $R_{rs}(\lambda)$ in combination with
63 derivative spectroscopy for identifying algal blooms were also demonstrated by Lubac et al.
64 (2008), who based their analysis on the position of the maxima and minima of the second
65 derivative of the spectral $R_{rs}(\lambda)$. Louchard et al. (2002) assessed major sediment pigments of
66 benthic substrates from derivative spectra of hyperspectral $R_{rs}(\lambda)$ measured in shallow marine
67 environments. In general, the optical detection of specific algal blooms appears feasible because
68 certain accessory pigments with specific absorption features are unique to individual
69 phytoplankton taxa and can be better differentiated in hyperspectral absorption data than in
70 multispectral data with a limited number of wavelengths.

71 The advantages and increasing availability of high spectral resolution measurements suggest
72 that the effectiveness of hyperspectral optical information for assessing phytoplankton diversity
73 should be further explored. In particular, there is a need to test whether the hyperspectral
74 approach, which has proven useful in inland and coastal waters (e.g., Lee & Carder, 2004;
75 Hunter et al., 2008; Lubac et al., 2008), can be also effective for the identification of different
76 phytoplankton assemblages at large spatial scales in open ocean waters. These tests are also
77 especially important for the common situation in which various phytoplankton groups co-exist at
78 significant concentrations, and no single species dominates the assemblage (i.e., a non-bloom

79 condition).

80 In this study, we analyze phytoplankton pigment data in conjunction with optical data of
81 absorption coefficients and remote-sensing reflectance, which were determined along a north-to-
82 south transect in the eastern Atlantic Ocean. Our primary goal is to examine the feasibility of
83 classifying different open ocean environments under non-bloom conditions in terms of
84 phytoplankton pigment assemblages from analysis of hyperspectral absorption and remote-
85 sensing reflectance measurements. In order to address this question, an unsupervised
86 hierarchical cluster analysis is applied to the pigment data set obtained from High Performance
87 Liquid Chromatography (HPLC) analysis of seawater samples and to the optical data sets
88 including the spectra of absorption coefficients and remote-sensing reflectance and their second
89 derivative spectra. For this analysis, the pigment data and the corresponding optical data were
90 selected to represent distinct differences in major accessory pigments present in the samples. We
91 view our analysis basically as a proof-of-concept study in which our approach is to use a
92 relatively small but carefully selected set of data which exhibits significant contrasts in the
93 composition of pigments, rather than to indiscriminately use large data sets. The pigment-based
94 clusters provide a reference for partitioning the selected data set into distinct subsets, each
95 characterized by different phytoplankton pigment composition. Two indices, cophenetic and
96 Rand, are examined to quantify the degree of similarity between the pigment-based clusters and
97 optical-based clusters, and are ultimately used to illustrate the effectiveness of optical
98 classification. The degree of similarity between clusters was evaluated for calculations involving
99 different spectral ranges of optical data. Because the quality of derivative analysis also depends
100 on parameters involved in data processing and computations, especially smoothing filter size and
101 derivative band separation (Tsai & Philpot, 1998; Lee & Carder, 2002; Vaiphasa, 2006), a
102 sensitivity of cluster analysis to the choice of these parameters was performed.

103

104 **2. Measurements and Data Analysis**

105 The approach in this study consists of three main components: (i) collection of field data of
106 phytoplankton pigments and ocean optical properties and selection of a subset of data
107 characterized by distinct differences in major accessory pigments for the cluster analysis, (ii)
108 radiative transfer modeling to compute hyperspectral remote sensing-reflectance, and (iii) cluster
109 analysis of pigment and optical data. The methodology of each component is described below.

110

111 *2.1. Field measurements*

112 Measurements of phytoplankton pigment composition and seawater optical properties were
113 obtained during the ANT-XXIII/1 expedition of the R/V Polarstern along a north-to-south
114 transect in the eastern Atlantic Ocean during October and November, 2005 (Fig. 1). The
115 investigated area spanned a wide range of different oceanic environments between the English
116 Channel and the waters off the African coast of Namibia. Typically, one full station was
117 conducted daily near local noon throughout the cruise. These full stations consisted of in situ
118 measurements of seawater inherent and apparent optical properties along with laboratory
119 analyses of water samples collected from discrete depths with the ship's CTD/rosette system.
120 For the present study, a subset of nine stations (see Fig. 1 for station locations) was selected for
121 cluster analysis based on the observation of distinct differences in the ratios of dominant
122 accessory pigments to total chlorophyll-*a* (further details in sec. 3.1). The selected data from the
123 nine stations are representative of surface waters within the top 5-10 m of the ocean, as our main
124 interest lies in the methodology for estimating variability in phytoplankton communities from
125 remote-sensing reflectance. A brief description of the measurements is provided in the following
126 three subsections. More methodological details, especially for the radiometric and
127 backscattering measurements, can be found in Stramski et al. (2008).

128

129 2.1.1. HPLC pigment analysis

130 Concentrations of chlorophyll-*a* and accessory pigments in phytoplankton were measured on
131 surface water samples from each station using HPLC techniques. Two sets of replicate samples
132 were collected and analyzed at two laboratories, the Center for Hydro-Optics and Remote
133 Sensing (CHORS) laboratory at San Diego State University (California, USA) and the GKSS
134 Research Centre in Geesthacht (Germany). The CHORS analysis was based on a method
135 described in Van Heukelem and Thomas (2001), and the GKSS samples were analyzed following
136 the method of Zapata et al. (2000). The CHORS analysis included identification and
137 quantification of more pigments (27) than the GKSS method (23) including alternative forms of
138 chlorophyll-*a*, and was chosen as the primary pigment data set for identifying phytoplankton
139 assemblages using cluster analysis. Throughout the rest of this paper, as a measure of
140 chlorophyll-*a* we use the CHORS values of the total chlorophyll-*a*, TChl*a*, which is defined as
141 the summed contributions of concentrations of monovinyl chlorophyll-*a* (MVChl*a*), divinyl
142 chlorophyll-*a* (DVChl*a*), chlorophyllide-*a* (Chlide), and the allomeric and epimeric forms of
143 chlorophyll-*a*.

144 Following completion of our analyses, the potential for errors in the CHORS results was
145 identified. The NASA team tasked with investigating HPLC data quality problems at CHORS
146 recommended that overall “These data are not validated and should not be used as sole basis for
147 a scientific result, conclusion, or hypothesis – independent corroborating evidence is required”
148 (Hooker and Van Heukelem, 2009). Based on field data obtained in the SeaHARRE-3
149 intercalibration experiment, corrections for specific individual pigments (MVChl*a*, DVChl*a*)
150 were developed (C. Trees, personal communication), which are used in our data set in the
151 determination of TChl*a*. These corrections are described in Stramski et al. (2008). The corrected
152 TChl*a* data exhibit reasonable agreement with fluorometrically-derived chlorophyll
153 measurements, and provide realistic estimates of chlorophyll-specific phytoplankton absorption

154 coefficients within the red peak of chlorophyll-*a*. We also applied similar corrections to three
155 other pigments in the data set; monovinyl chlorophyll-*b*, β -carotene, and alloxanthin. We caution
156 that our corrections were developed from a limited set of pigment data and intercalibration
157 results, and should not be used indiscriminately with other data sets affected by the CHORS data
158 quality problems.

159 We also compared the corrected CHORS results with independent pigment determinations
160 done by GKSS. Some differences in the concentrations of individual pigments between the two
161 data sets were observed. For example, the sum of monovinyl chlorophyll-*a* (MVChl*a*) and
162 divinyl chlorophyll-*a* (DVChl*a*) was generally higher (on average by 20% with a standard
163 deviation of 23%, number of samples 25) for CHORS compared with GKSS. However, the
164 CHORS data yielded more reasonable estimates of chlorophyll-specific phytoplankton
165 absorption.

166 Despite such differences in the estimates of some individual pigment concentrations, with
167 regard to the present application it is important to note that both laboratories provided similar
168 characterization of samples in terms of the relative pigment composition as described by ratios of
169 various individual pigments to TChl*a*. The same dominant accessory pigments for any given
170 station and the same trends in the pigment ratios among the stations were obtained from both sets
171 of HPLC analyses. This is an essential result for our study because in the cluster analysis we
172 utilize only pigment ratios, and not individual pigment concentrations. For the nine stations
173 selected in our analysis, the cluster techniques applied independently to the CHORS and GKSS
174 sets of HPLC pigment ratios yielded a very similar partitioning of stations into clusters. We
175 therefore chose the one set of corrected pigment results from CHORS for all subsequent analyses
176 in this paper (sec. 3.1 presents CHORS-based pigment clusters). This choice is further supported
177 by comparisons of cluster analysis of phytoplankton absorption data with the CHORS and
178 GKSS-based pigment clusters. The measures of cluster similarity (see sec. 2.3) between the
179 absorption-based clusters and pigment-based clusters were found to be similar when either the

180 CHORS and GKSS pigment data were used.

181 Because of a large range of pigment compositions across different phytoplankton classes,
182 determinations of phytoplankton composition from HPLC pigment data is not straightforward
183 (e.g., Jeffrey et al., 1997). Whereas certain diagnostic pigments can serve as unambiguous
184 markers for some phytoplankton classes (e.g., peridinin in dinoflagellates, alloxanthin in
185 cryptophytes), many important pigments are shared by more than one algal taxa (e.g.,
186 fucoxanthin in diatoms, haptophytes, chrysophytes, and raphidophytes). Nevertheless, because
187 many of the classes have distinctive suites of marker pigments, HPLC data can be useful for
188 indicating their presence and abundance in a mixed phytoplankton population. Specifically, a
189 useful indication of contributing phytoplankton classes can be obtained from the ratios of the
190 concentrations of specific pigments to chlorophyll-*a* or the ratios of specific diagnostic pigments
191 to the sum of these diagnostic pigments, because these ratios can differ between taxonomic
192 groups (Mackey et al., 1996; Wright et al., 1996; Vidussi et al., 2001).

193 For each of the nine stations selected in this study, we calculated two sets of pigment ratios
194 for subsequent use in the cluster analysis. The first set of pigment ratios consisted of ratios of
195 the concentration of each individual pigment to the TChl*a* concentration, as obtained from HPLC
196 measurements at the CHORS laboratory. The following 24 pigments were included in these
197 calculations: monovinyl chlorophyll-*a*, divinyl chlorophyll-*a*, chlorophyllide-*a*, chlorophyll-*a*
198 allomer, chlorophyll-*a* epimer, monovinyl chlorophyll-*b*, divinyl chlorophyll-*b*, chlorophyll-*c*2,
199 chlorophyll-*c*3, α -carotene, β -carotene, alloxanthin, diadinoxanthin, diatoxanthin, fucoxanthin,
200 19'-hexanoyloxyfucoxanthin, 19'-butanoyloxyfucoxanthin, neoxanthin, prasinoxanthin,
201 violaxanthin, zeaxanthin, peridinin, pheophorbide-*a*, and lutein. The CHORS pigment data set
202 also included a few additional pigments (chlorophyll-*c*1, gyroxanthin-diester, and pheophytin-*a*)
203 which were below the level of detection for the nine stations. These pigments were not included
204 in our cluster analysis as they would have no effect on the results.

205 The second set of pigment ratios was based on 8 diagnostic pigments: divinyl chlorophyll-*a*
206 (DVChl*a*), divinyl chlorophyll-*b* (DVChl*b*), alloxanthin (Allo), fucoxanthin (Fuco), 19'-
207 hexanoyloxyfucoxanthin (Hex), 19'-butanoyloxyfucoxanthin (But), peridinin (Peri), and
208 zeaxanthin (Zea). This list of pigments is consistent with that proposed by Vidussi et al. (2001)
209 with the exception that we added DVChl*a* which is a diagnostic pigment for prochlorophytes.
210 For each station the 8 ratios were calculated by dividing a concentration of a given diagnostic
211 pigment to the sum of the 8 diagnostic pigment concentrations (see Uitz et al., 2006).

212

213 *2.1.2. Inherent optical properties*

214 The inherent optical properties (IOPs) of seawater have a two-fold application in our study.
215 First, the absorption, scattering, and backscattering coefficients are used to define IOP inputs to
216 radiative transfer simulations that generate hyperspectral data of remote-sensing reflectance (sec.
217 2.2). Second, the total absorption coefficient, $a(\lambda)$, and the phytoplankton absorption coefficient,
218 $a_{ph}(\lambda)$, are utilized directly in the cluster analysis (sec. 2.3 and 3.2).

219 The spectral absorption coefficients of particles, $a_p(\lambda)$, and colored dissolved organic matter
220 (CDOM), $a_{cdom}(\lambda)$ in m^{-1} , were determined at 1-nm intervals from high spectral resolution
221 measurements on freshly-collected discrete water samples with a point-source integrating cavity
222 absorption meter (PSICAM) over the range 350-750 nm (Röttgers et al., 2005, Röttgers &
223 Doerffer, 2007). As the PSICAM did not provide data below 350 nm, the $a_p(\lambda)$ values within the
224 300 to 350 nm spectral range were obtained from filter pad measurements on discrete water
225 samples collected on GF/F filters and frozen in liquid nitrogen until analysis with a dual-beam
226 spectrophotometer (Lambda 18, Perkin Elmer). The filter pad measurements were made with the
227 transmittance-reflectance (T-R) technique of Tassan and Ferrari (1995; 2002) using a correction
228 for the pathlength-amplification factor from Stramska et al. (2006). We have chosen to use the
229 PSICAM data of $a_p(\lambda)$ over the majority of the spectrum because the PSICAM technique
230 involves a direct measurement of absorption on particle suspension with minimal scattering

231 artifacts, which is expected to be generally superior to the filter pad measurements. The data of
232 $a_{cdom}(\lambda)$ below 350 nm were obtained from an exponential fit to the PSICAM-measured $a_{cdom}(\lambda)$.
233 A null point correction based on wavelengths in the far red or near-infrared was applied to all
234 $a_p(\lambda)$ and $a_{cdom}(\lambda)$ spectra. The total spectral absorption coefficient, $a(\lambda)$, was determined as the
235 sum of $a_p(\lambda)$, $a_{cdom}(\lambda)$, and the pure water component, $a_w(\lambda)$. The latter was obtained from Pope
236 & Fry (1997) for the spectral range 380-727 nm and from Fry et al. (2006) for the range 300-379
237 nm. We note that our primary interest is in the spectral information contained at wavelengths
238 longer than about 350 nm extending throughout the visible part of the spectrum up to 725 nm
239 where most phytoplankton pigments exhibit significant absorption features. However, data at
240 wavelengths shorter than 350 nm are useful for our analysis, especially in the context of
241 derivative spectra whose discrete values at specific wavelengths were calculated in our study
242 using data covering a bandwidth on the order of 10 - 30 nm.

243 The spectra of the phytoplankton absorption coefficient, $a_{ph}(\lambda)$, were determined as a
244 difference between the absorption coefficient of particles, $a_p(\lambda)$, and the non-phytoplankton
245 component of particulate absorption, $a_d(\lambda)$, which is commonly referred to as detrital absorption.
246 These determinations were based on the T-R filter pad measurements, in which the $a_d(\lambda)$ spectra
247 were measured on GF/F sample filters following treatment with sodium hypochlorite NaClO
248 (Ferrari and Tassan, 1999). In this treatment, the particles on the sample filter were exposed to a
249 small amount of a 2% NaClO solution for several minutes to bleach phytoplankton pigments.

250 The spectral beam attenuation coefficient of particles and CDOM, $c_{p,cdom}(\lambda)$ in m^{-1} , was
251 determined at each station from in situ measurements with two single-wavelength C-Star
252 transmissometers (488 and 660 nm; WET Labs, Inc.). Note that the C-star derived values
253 represent the total beam attenuation, $c(\lambda)$, with pure seawater contribution, $c_w(\lambda)$, subtracted, i.e.
254 $c_{p,cdom}(\lambda) = c(\lambda) - c_w(\lambda) = a_p(\lambda) + a_{cdom}(\lambda) + b_p(\lambda)$, where $b_p(\lambda)$ is the spectral scattering
255 coefficient of particles. The values of $b_p(\lambda)$ at 488 and 660 nm were thus calculated from C-Star

256 attenuation and PSICAM absorption measurements as $b_p(\lambda) = c_{p,cdom}(\lambda) - a_p(\lambda) - a_{cdom}(\lambda)$. A
257 power function fit was then applied to these values to produce the spectral data of $b_p(\lambda)$ over the
258 range 300-750 nm with a 1 nm resolution. The pure seawater scattering coefficient, $b_w(\lambda)$, was
259 calculated using the Buiteveld et al. (1994) equations with measured water temperature and
260 salinity (see Twardowski et al., 2007 and Stramski et al., 2008 for details). The total scattering
261 coefficient, $b(\lambda)$, was obtained as a sum $b_w(\lambda) + b_p(\lambda)$.

262 The spectral backscattering coefficient, $b_b(\lambda)$ in m^{-1} , was determined by combining in situ
263 measurements with three instruments, a Hydroscat-6 and two a-beta sensors (HOBI Labs, Inc.),
264 to yield a total of eight spectral bands: 420, 442, 470, 510, 550, 589, 620, and 671 nm (Stramski
265 et al. 2008). Because $b_b(\lambda)$ is generally expected to be a smooth monotonic function of
266 wavelength, especially in the open ocean under non-bloom conditions, the experimental
267 measurements were fitted to a power function to obtain hyperspectral resolution over the 300-
268 750 spectral range. The spectral backscattering coefficient of particles, $b_{bp}(\lambda)$ in m^{-1} , was
269 determined as a difference between the total and pure seawater backscattering coefficients, $b_b(\lambda)$
270 - $b_{bw}(\lambda)$, in which the pure seawater component, $b_{bw}(\lambda)$, was calculated as $0.5 b_w(\lambda)$.

271 From the values of $b_p(\lambda)$ and $b_{bp}(\lambda)$, we calculated the particle backscatter fraction $B_p(\lambda) =$
272 $b_{bp}(\lambda) / b_p(\lambda)$. These data were then fitted to a power function, $B_p(\lambda) = B_p(\lambda_0) (\lambda_0 / \lambda)^m$, where λ_0
273 is the reference wavelength 550 nm. The backscattering fraction $B_p(\lambda_0)$ at the reference
274 wavelength and the exponent m represent the best fit parameters of the linear regression analysis
275 performed for the log-transformed data of $B_p(\lambda)$ vs. λ for each station. The parameters of the
276 power function fit of $B_p(\lambda)$ were used as input to radiative transfer simulations (see sec. 2.2).

277

278 2.1.3. Remote-sensing reflectance

279 Values of multispectral remote-sensing reflectance, $R_{rs}(\lambda)$ in sr^{-1} , were estimated at each
280 station at 13 wavelengths from in situ measurements of underwater vertical profiles of spectral
281 nadir upwelling radiance, $L_u(\lambda, z)$ in $W m^{-2} sr^{-1} nm^{-1}$, and spectral downwelling plane irradiance,

282 $E_d(\lambda, z)$ in $\text{W m}^{-2} \text{nm}^{-1}$, where z is depth. These measurements were made with a freefall
283 spectroradiometer, the SeaWiFS Profiling Multichannel Radiometer (SPMR, Satlantic, Inc.).
284 The wavelengths for these measurements are 339, 380, 412, 443, 470, 490, 510, 532, 554, 589,
285 619, 666, and 683 nm. The radiometric measurements and data processing were consistent with
286 methods recommended in NASA protocols (Mueller et al., 2003).

287

288 2.2. Modeled hyperspectral reflectance

289 Because hyperspectral radiometric measurements were not conducted during the ANT-
290 XXIII/1 cruise and our primary interest is in the analysis of hyperspectral optical data, we
291 performed numerical simulations of radiative transfer (RT) to estimate the hyperspectral remote-
292 sensing reflectance $R_{rs}(\lambda)$ for each of the nine selected stations. The radiative transfer model
293 Hydrolight/Ecolight version 5.0 (Sequoia Scientific, Inc.) was used (Mobley 1994; 2008). An
294 important prerequisite for undertaking these RT simulations was the availability of a
295 comprehensive suite of IOPs for each station for use as input to the simulations, and also the
296 availability of the multispectral $R_{rs}(\lambda)$ derived from in situ measurements for use in validating
297 the simulated hyperspectral $R_{rs}(\lambda)$.

298 The RT calculations were carried out within the spectral region 300 nm to 725 nm with high
299 spectral resolution (1 nm). Similarly to the absorption, our main interest is in the reflectance data
300 at wavelengths longer than about 350 nm. However, in addition to the requirements associated
301 with derivative calculations, the radiative transfer simulations below 350 nm are needed to
302 account for Raman scattering contributions observed at $\lambda > 350$ nm. The ocean was assumed to
303 be infinitely deep and optically homogeneous, and the simulations included the Raman scattering
304 and fluorescence by colored dissolved organic matter within the ocean. The sea surface
305 boundary conditions were estimated from observations of wind speed and sky conditions
306 (cloudiness) at each station site, and the solar zenith angle was calculated for the corresponding

307 date and geographic coordinates. The inherent optical properties of the water column required as
308 input to the simulations were derived from the IOP measurements in the surface waters and
309 additional relevant determinations of $a(\lambda)$ and $b(\lambda)$ as described in sec. 2.1.2. The selection of
310 the particulate scattering phase function, which is also part of IOP inputs to the RT simulations,
311 was based on the particle backscatter fraction $B_p(\lambda)$. We used the Fournier-Forand phase
312 functions which are parameterized in terms of $B_p(\lambda)$ and are built into the Hydrolight/Ecolight
313 model.

314 Fig. 2 compares the model-simulated hyperspectral $R_{rs}(\lambda)$ with the measured multispectral
315 $R_{rs}(\lambda)$ for two selected stations. The model results compare reasonably well with measurements,
316 which lends confidence to the use of hyperspectral $R_{rs}(\lambda)$ in our cluster analysis. This level of
317 consistency between the model and measurements suggests that the suite of parameters used as
318 input to the RT simulations realistically represent the actual field conditions. The ability to
319 define realistic inputs derives, in turn, from a comprehensive suite of IOP measurements that
320 were carried out during the cruise.

321

322 *2.3. Hierarchical cluster analysis and similarity indices between dendrograms*

323 A hierarchical cluster analysis (HCA) was used to classify the 9 selected stations into distinct
324 groups on the basis of several types of input data vectors (or objects), which included the HPLC
325 pigments and optical data derived from spectral absorption coefficients and remote-sensing
326 reflectance. For a given type of data, the input to the cluster analysis consisted of 9 numerical
327 data vectors, each representing one of the 9 stations. For the input data representing the ratio of
328 individual pigment concentrations to TChla, an object for a given station is a data vector $\{p_1, p_2,$
329 $p_3, \dots, p_{24}\}$ where the consecutive elements p_i represent the ratio of each of the 24 individual
330 pigment concentrations to TChla concentration. Another type of pigment data vector used in the
331 cluster analysis is of the form $\{d_1, d_2, d_3, \dots, d_8\}$, where the consecutive elements represent a ratio

332 of one of the 8 diagnostic pigment concentrations to the sum of diagnostic pigment
333 concentrations.

334 Several types of optical data vectors were used as input to the HCA analysis, including
335 objects consisting of hyperspectral data of the remote-sensing reflectance, $R_{rs}(\lambda)$, the
336 phytoplankton absorption coefficient, $a_{ph}(\lambda)$, the sum of pure water and phytoplankton
337 absorption coefficients, $a_w(\lambda) + a_{ph}(\lambda)$, and the total absorption coefficient, $a(\lambda)$. The input
338 characterizing the hyperspectral remote-sensing reflectance for any given station was used in the
339 form of the following data vector $\{R_{rs}(\lambda_1)/R_{rs}(555), R_{rs}(\lambda_2)/R_{rs}(555), R_{rs}(\lambda_3)/R_{rs}(555), \dots,$
340 $R_{rs}(\lambda_n)/R_{rs}(555)\}$, where the consecutive elements represent the values of R_{rs} at successive light
341 wavelengths normalized to R_{rs} at 555 nm over the spectral range from λ_1 to λ_n . Similar input
342 vectors were created for the different components of spectral absorption. Because our analysis is
343 focused on the spectral shapes, all the optical spectra used in the cluster analysis were
344 normalized by the value of the optical variable at 555 nm at which variations in R_{rs} within the
345 open ocean are generally small. The spectra involving the absorption coefficients were
346 additionally normalized by TChl a concentration to minimize variability in absorption associated
347 with changes in phytoplankton biomass. The rationale for selecting the data of $a(\lambda)$, $a_w(\lambda) +$
348 $a_{ph}(\lambda)$, and $a_{ph}(\lambda)$ to create input data vectors for the cluster analysis stems from the fact that the
349 variation in the spectral shape of $a(\lambda)$ is typically a major determinant of the variation in the
350 spectral shape of $R_{rs}(\lambda)$. In turn, the variations in the spectral shape of $a_w(\lambda) + a_{ph}(\lambda)$ or $a_{ph}(\lambda)$
351 can be viewed as an important or dominant source of variation in the spectral shape of $a(\lambda)$ in
352 open ocean situations.

353 We also created vectors from the second derivative spectra of the hyperspectral reflectance
354 and absorption objects for input into the cluster analysis. The estimation of the second derivative
355 spectra from these data was made with a finite divided difference algorithm, the so-called “finite
356 approximation”, which computes the changes in curvature of a given spectrum over a sampling

357 interval ($\Delta\lambda$) or band separation (BS) defined as $\Delta\lambda = \lambda_j - \lambda_i$, where $j > i$. Because the
358 identification of spectral details in the derivative spectra depends on the selection of the band
359 separation, we tested the sensitivity of cluster results to the choice of BS . The derivative
360 technique is also sensitive to signal noise, thus smoothing was applied to the hyperspectral
361 optical data prior to computation of derivative spectra. Specifically, a mean-filter smoothing
362 method was used in which the extent of spectral smoothing depends on the size of the filter
363 window (WS) used for averaging. We tested the sensitivity of cluster results to different values
364 of WS . The sensitivity analysis over a range of BS and WS values allowed us to achieve the best
365 compromise between the ability to resolve fine spectral details and the reduction of noise effects
366 in the second derivative spectra. As discussed below (sections 3.2 and 3.3), the optimal values of
367 BS and WS chosen in this study for the derivative analysis of absorption data are 9 nm. For the
368 derivative analysis of reflectance data, these values are 27 nm. Therefore, although the
369 derivative calculations were made using data from the spectral range 300 – 725 nm, our results
370 from the derivative analysis for absorption will be reported between the wavelengths of $\lambda_{\min} =$
371 309 nm ($\equiv 300 + 9$) and $\lambda_{\max} = 716$ nm ($\equiv 725 - 9$). For the reflectance derivative, the results
372 will be reported between $\lambda_{\min} = 327$ nm ($\equiv 300 + 27$) and $\lambda_{\max} = 698$ nm ($\equiv 725 - 27$).

373 With regard to the analysis of remote-sensing reflectance, the cluster analysis was also
374 applied to the multispectral reflectance data obtained from in situ SPMR measurements at
375 several discrete wavelengths. We examined the 3-element objects $\{R_{rs}(443)/R_{rs}(554),$
376 $R_{rs}(490)/R_{rs}(554), R_{rs}(510)/R_{rs}(554)\}$, which consist of band ratios that are similar to those used
377 in current research based on satellite ocean color observations such as the Sea-viewing Wide
378 Field-of-View Sensor (SeaWiFS). We also examined the vectors consisting of 13 band ratios of
379 remote-sensing reflectance with $R_{rs}(554)$ in the denominator, as determined from SPMR
380 measurements at 13 wavebands.

381 The HCA method, schematically presented in Fig. 3, was applied using the above defined
382 pigment and optical data vectors as input objects. This method utilizes an unsupervised

383 classification algorithm which creates a hierarchical cluster tree (dendrogram) that partitions a
 384 given set of input data into clusters or groups of objects (Jain et al., 1999; Berkhin, 2006). Each
 385 group includes objects that are similar to each other, but different from objects in other groups.

386 The cluster tree is obtained using a linkage algorithm based on initial calculations of the
 387 pairwise distance between all objects included in the input data set. In this study the similarity
 388 between each pair of objects was the cosine distance, d , calculated as one minus the cosine of the
 389 angle θ between each pair of objects:

$$390 \quad d(x_1, x_2) = 1 - \cos \theta = 1 - \left(\frac{x_1 \cdot x_2}{\|x_1\| \times \|x_2\|} \right) \quad (1)$$

391 where the objects x_1 and x_2 include the two considered input data vectors and the cosine of the
 392 angle between the vectors is obtained as the ratio of the dot product of the vectors to the product
 393 of norms of the vectors. Note that as the angle between the objects decreases the cosine
 394 approaches 1, resulting in a smaller distance between the input data vectors and therefore higher
 395 similarity. We also tested other measures of similarity between the input objects, e.g., Euclidean
 396 distance using a similar approach to that proposed in Robila (2005). The cosine distance was
 397 selected as the most appropriate measure for our study because it reflects mainly the differences
 398 in the spectral shape of optical data rather than magnitude. The cosine distance is also
 399 advantageous because it is scale invariant, i.e., insensitive to normalization of optical spectra at a
 400 specific wavelength.

401 As a linkage algorithm, the shortest distance D , also referred to as the nearest neighbor, was
 402 computed to measure the distance between two clusters of objects in the tree:

$$403 \quad D(a, b) = \min [dist(x_{ai}, x_{bj})] \quad i \in (1, \dots, n_a) \text{ and } j \in (1, \dots, n_b) \quad (2)$$

404 where x_{ai} is the i th object in cluster a and x_{bj} is the j th object in cluster b . In the traditional
 405 graphical representation of a dendrogram, the individual objects appear at one end and a single
 406 cluster containing all objects at the other end (e.g., Jain et al., 1999). In our presentation of

407 dendrograms, pairs of objects showing a small cosine distance between them (i.e., with similar
 408 pigment composition or spectral properties) provide small linkage distance and therefore appear
 409 closer to each other in the cluster tree.

410 To evaluate the utility of optical data for discriminating phytoplankton pigment assemblages,
 411 we compared the dendrograms obtained for the different spectral optical data with a reference
 412 dendrogram obtained using the pigment composition data (see final step in Fig. 3). For this
 413 analysis, we utilize two objective criteria of cluster similarity, the cophenetic index (Sokal &
 414 Rolf, 1962) and the Rand index (Rand, 1971).

415 The cophenetic index (r_c) is a measure of how precisely two dendrograms preserve the
 416 pairwise distances between data objects. This index is computed from the cophenetic matrix (C)
 417 associated with each dendrogram. The elements of a cophenetic matrix ($c_{i,j}$) encode the distance
 418 between two objects (i, j), representing in the dendrogram the height of the link at which those
 419 two objects are first joined. This height is the distance between the two clusters that are merged
 420 by this link. The cophenetic index r_c represents the correlation between two cophenetic
 421 matrices (C_1 and C_2):

$$422 \quad r_c = \frac{\sum_i \sum_j (c_{1,i,j} - \bar{c}_1)(c_{2,i,j} - \bar{c}_2)}{\sqrt{\left(\sum_i \sum_j (c_{1,i,j} - \bar{c}_1)^2\right) \left(\sum_i \sum_j (c_{2,i,j} - \bar{c}_2)^2\right)}} \quad (3)$$

423 where \bar{c}_1 and \bar{c}_2 are the mean values of the elements of the matrices C_1 and C_2 , respectively.

424 The Rand index (r_r) provides a measure of the similarity between two hierarchical
 425 dendrograms in terms of the proportion of pairs of objects whose relationship is the same in both
 426 dendrograms. The r_r value of 1 means that all pairs of objects are clustered in the same way in
 427 both dendrograms. Note that this index has to be computed using all dendrograms cut
 428 horizontally at a level (i.e., at a specific linkage distance) which yields the optimal number of
 429 clusters (k). Otherwise, r_r would always provide a proportion of 100% because a complete
 430 dendrogram always decomposes the input data all the way through the lowest level (i.e., until the

431 branches consist only of single objects). Detecting natural groupings in the dendrogram and
432 selecting the optimal number of clusters is performed by analyzing a diagram of the increasing
433 linkage distances along the dendrogram. Based on the points at which the linkage distances
434 between the objects change abruptly (which is associated with a steep increase of the within
435 cluster variance), the optimal number of clusters k is determined and all objects located below
436 the point where the hierarchical tree is cut off are assigned to a single cluster (Salvador & Chan,
437 2004).

438

439 **3. Results and discussion**

440

441 *3.1. Classification of stations based on pigment composition*

442 For the 9 stations selected in the study, the estimate of the TChl a concentration ranges from
443 about 0.11 mg m⁻³ at the southernmost station 59 in the open ocean off the coast of Namibia to
444 0.62 mg m⁻³ at the northernmost station 1 in the English Channel (Fig. 1). The variability in
445 pigment composition for the 9 stations is summarized in Table 1, which provides the ratios of the
446 concentration of several dominant pigments to TChl a . Apart from MVChl a , which is a principal
447 pigment common to all phytoplankton, the second most important pigment at different stations
448 was either DVChl a , zeaxanthin (Zea), 19'-hexanoyloxyfucoxanthin (Hex), or fucoxanthin
449 (Fuco). Table 1 also identifies the two dominant pigments (excluding MVChl a) which yield the
450 highest ratio to TChl a at each station. The values for the ratios of the two dominant pigments to
451 TChl a were used as a basis for selecting the 9 stations. As these stations represent different
452 pigment compositions, we assigned a class label A, B, C, D, E, or F to each station.

453 Most stations visited during the cruise within the tropical and subtropical regions of the
454 Atlantic were dominated by Zea and DVChl a , but the relative predominance of these two
455 pigments varied between the stations. These pigments are diagnostic of picophytoplankton that

456 include DVChla- and Zea-containing prochlorophytes and Zea-containing cyanobacteria (mainly
457 *Synechococcus* in the open ocean waters). We selected 5 stations (6, 12, 37, 44, and 46) to
458 represent this type of pigment assemblage. Note that 4 stations dominated by Zea and DVChla
459 with fairly similar ratios DVChla/TChla and Zea/Chla are grouped within the same class C with
460 a label C1, C2, C3, and C4. The station 6, where DVChla/TChla is significantly higher than
461 Zea/TChla, is considered as a separate class B. This station (or class B) is dominated by
462 prochlorophytes as DVChla is an unambiguous marker of this group. The class C stations also
463 show significant role of prochlorophytes. However, this class exhibits a relatively higher
464 contribution of Zea than class B, which is likely indicative of an increased role of cyanobacteria.

465 Fuco and MVChlb are the predominant accessory pigments at station 1 (class A). As these
466 two pigments are not confined to one phytoplankton class, this station could have been
467 dominated by Fuco-rich diatoms, haptophytes, and/or dinoflagellates, as well as MVChlb-rich
468 prasinophytes and/or chlorophytes. The predominant accessory pigment at stations 48 (class D)
469 and 51 (class E) is Hex, which suggests that haptophytes and/or chrysophytes are major
470 phytoplankton groups at these locations. These stations are designated as different classes
471 because they clearly differ in accessory pigments that follow Hex in ranking. Zea and Fuco are
472 the second most important diagnostic pigments at stations 48 (class D) and 51 (class E),
473 respectively. Finally, station 59 (class F) also shows a significant role of Hex-rich phytoplankton
474 although Zea is the most important diagnostic pigment at this location, indicating potential
475 significance of cyanobacteria and/or prochlorophytes.

476 Fig. 4a shows the hierarchical cluster tree obtained for the input data consisting of the ratios
477 of concentrations of 24 individual pigments to TChla at each station. The optimal number of
478 clusters (k) is derived from a diagram of linkage distances along the dendrogram (Fig. 4b). The
479 first steep increase in the linkage distance observed in this diagram, which is associated with an
480 increase of the within cluster variance, suggests an optimal partitioning of the pigment data into
481 5 clusters. The linkage distance of 0.023 can be selected to characterize this steep increase in

482 variance (see dashed lines in Fig. 4a and 4b). For the dendrogram cut at a level of linkage
483 distance of 0.023, all clusters are single object (i.e., single station) clusters, except for a multi-
484 object cluster that includes stations C1, C2, C3, C4, and B. The results of this cluster analysis
485 are quite consistent with the preliminary classification obtained by considering just two dominant
486 diagnostic pigments (see Table 1). Note that stations C1, C2, C3, C4, and B are all characterized
487 by relatively high ratios of DVChl*a* and Zea to TChl*a*. Some differences between these stations
488 in terms of the relative roles of DVChl*a* and Zea do not, however, produce significant distances
489 between the corresponding pigment data vectors and hence these 5 stations are grouped into a
490 single cluster. The dendrogram also indicates that the stations classified as A, D, E, and F
491 display significant dissimilarities between each other and when compared to the stations
492 classified as B and C. Note that the stations 48 (class D) and 59 (class F) have the Hex and Zea
493 as dominant diagnostic pigments, albeit in reverse ranking (see Table 1), so these stations appear
494 closer to one another in the dendrogram (Fig. 4a).

495 Fig. 4c also depicts results from the cluster analysis of pigment data but for the input data
496 vectors consisting of the ratios of concentrations of the 8 diagnostic pigments to the sum of the 8
497 diagnostic pigments. The partitioning of the stations obtained on the basis of these pigment
498 ratios is qualitatively identical to the partitioning based on the ratios of 24 pigments to TChl*a*.
499 The stations C1, C2, C3, C4, and B are again grouped within a single cluster and each of the
500 remaining stations represent a separate cluster. The results in Figs. 4a and 4c along with the
501 preliminary classification of stations based on two dominant diagnostic pigments (Table 1)
502 suggest that there is a certain degree of flexibility in the selection of pigment ratios as a basis for
503 discriminating different pigment assemblages in a consistent fashion. In the following analysis
504 of optical data as a means for assessing differences in pigment assemblages, the pigment-based
505 cluster partitioning obtained with the 24 pigment ratios (as shown in Fig. 4a) is used as a
506 reference.

507

508 3.2. Classification of stations based on absorption spectra

509 Fig. 5 shows hyperspectral data of absorption coefficients first normalized at 555 nm and
510 then divided by the total chlorophyll-*a* concentration for the nine stations. These spectra are
511 referred to as the spectral chlorophyll (Chl)-specific normalized absorption coefficients.
512 Specifically, we examine the Chl-specific normalized coefficients for the total absorption, $a_n^*(\lambda)$,
513 the absorption of pure seawater plus phytoplankton, $a_{n,w+ph}^*(\lambda)$, and the absorption of
514 phytoplankton alone, $a_{n,ph}^*(\lambda)$. The differences in the shape of phytoplankton absorption in the
515 UV and blue spectral regions are generally quite large between most stations (Fig. 5a). With the
516 addition of the pure water contribution, differences in the spectral shape of $a_{n,w+ph}^*(\lambda)$ continue
517 to be seen but are considerably smaller (Fig. 5b). Finally, upon further addition of the
518 contributions associated with non-phytoplankton particles and CDOM, the spectral shape of total
519 absorption again shows larger differences between the stations at wavelengths shorter than the
520 normalization point at 555 nm (Fig. 5c). From the visual inspection of these plots it is, however,
521 difficult to deduce to what extent the observed differences might be consistent with the
522 classification of stations based on pigment composition.

523 Fig. 6 illustrates the results from cluster analysis applied to the absorption spectra presented
524 in Fig. 5 and the corresponding second derivative spectra over the entire spectral range from 300
525 nm to 725 nm. In nearly all cases (Fig. 6a-e), the absorption-based cluster trees differ
526 significantly from the pigment-based cluster tree shown in Fig. 4a. Thus, the full hyperspectral
527 data of $a_n^*(\lambda)$, $a_{n,w+ph}^*(\lambda)$, and $a_{n,ph}^*(\lambda)$ as well the second derivative spectra of $a_n^*(\lambda)$ and
528 $a_{n,w+ph}^*(\lambda)$ do not provide useful information for discriminating the differences in pigment
529 assemblages at the examined stations. The only case in which stations are classified within the
530 dendrogram in a similar way to the pigment-based cluster tree is when the second derivative of
531 phytoplankton absorption spectra is considered (Fig. 6f). When this absorption-based
532 dendrogram is cut horizontally at a level of linkage distance of 0.023 that yields 5 clusters, the

533 same stations are grouped in separate clusters as in the pigment-based cluster tree. This result
534 supports the potential usefulness of the second derivative of phytoplankton absorption spectra for
535 discriminating different pigment assemblages.

536 In the analysis above we considered a spectral range from 300 nm to 725 nm, which is much
537 broader than the spectral region where specific absorption imprints caused by accessory
538 pigments occur. It is therefore useful to examine whether the cluster analysis of absorption data
539 yields similarity with pigment-based clusters if different, narrower spectral ranges are
540 considered. Fig. 7 illustrates the degree of similarity between the absorption-based and pigment-
541 based cluster trees for different spectral ranges of absorption data. The degree of similarity is
542 shown in terms of cophenetic and Rand indices. The three absorption spectra, $a_n^*(\lambda)$,
543 $a_{n,w+ph}^*(\lambda)$, and $a_{n,ph}^*(\lambda)$, are considered in this analysis. In each graph, the distribution of
544 values for the cophenetic or Rand index is shown as a function of the spectral range considered,
545 with the lower limit of the spectral range, λ_{\min} , displayed along the y-axis (ordinate) and the
546 upper limit, λ_{\max} , along the x-axis (abscissa). The similarity indices are thus shown for many
547 spectral ranges represented by many combinations of λ_{\min} and λ_{\max} . The higher values of indices,
548 depicted by darker areas in the graphs, correspond to better similarity between a given
549 absorption-based cluster tree and pigment-based tree. The best degree of similarity is obtained
550 when the indices are close to 1, indicated by the nearly black areas in the graphs. According to
551 the distributions of cophenetic index, this is the case when the phytoplankton absorption
552 spectrum $a_{n,ph}^*(\lambda)$ is analyzed over the spectral range approximately from $\lambda_{\min} = 425$ nm to λ_{\max}
553 $= 540$ nm (Fig. 7e). The distribution of the Rand index indicates that the best similarity between
554 the $a_{n,ph}^*(\lambda)$ -based cluster tree and the pigment-based tree occurs within a broader spectral
555 region, approximately between $\lambda_{\min} = 390$ nm and $\lambda_{\max} = 610$ nm (Fig. 7f). These optimal
556 spectral regions generally overlap with the wavelength range where absorption characteristics of
557 main accessory pigments appear (e.g., Bricaud et al., 2004). The remaining results in Fig. 7

558 (panels a, b, c, d) show generally poor similarity between the absorption data of $a_n^*(\lambda)$ or
559 $a_{n,w+ph}^*(\lambda)$ and pigment composition, regardless of the spectral range considered.

560 Fig. 8 depicts similar results, but for the similarity between pigment composition and the
561 second derivative spectra of $a_n^*(\lambda)$, $a_{n,w+ph}^*(\lambda)$, and $a_{n,ph}^*(\lambda)$. The use of derivative spectra
562 generally improves the similarity as indicated by the presence of darker areas or the larger extent
563 of dark areas in the distributions of the cophenetic and Rand indices. For example, compared to
564 the results for the ordinary spectra of $a_{n,w+ph}^*(\lambda)$ in Fig. 7d, a significant increase in the Rand
565 index is observed for the second derivative spectra of $a_{n,w+ph}^*(\lambda)$ within the spectral range from
566 $\lambda_{\min} = 440$ nm to $\lambda_{\max} = 650$ nm (Fig. 8d). The improvement is even more striking for the results
567 involving the second derivative spectra of phytoplankton absorption (Fig. 8e, f). The spectral
568 regions where the cophenetic and Rand indices assume high values near or equal to 1 are much
569 larger compared with the analysis of ordinary (non-differentiated) spectra of $a_{n,ph}^*(\lambda)$. The
570 darkest areas in the distributions of the indices in Fig. 8e, f cover a broad spectral range,
571 approximately from $\lambda_{\min} = 370$ nm to $\lambda_{\max} = 716$ nm. This result indicates that a high degree of
572 similarity between the second derivative spectra of $a_{n,ph}^*(\lambda)$ and pigment composition can be
573 obtained for many different combinations of spectral ranges of absorption data (i.e., different
574 combinations of λ_{\min} and λ_{\max}).

575 The improvement in the similarity between the pigment-based and absorption-based cluster
576 trees achieved as a result of utilization of second derivative spectra compared to the ordinary
577 spectra of phytoplankton absorption, $a_{n,ph}^*(\lambda)$, is clearly illustrated by histograms of cophenetic
578 index obtained on the basis of all possible combinations of spectral ranges examined (Fig. 9).
579 We note that the cophenetic index does not require a priori selection of the optimal number of
580 clusters which is somewhat subjective and, therefore, this index facilitates the comparison of
581 results for ordinary and derivative spectra shown in Fig. 7e and Fig. 8e. Compared with the
582 histogram obtained using the ordinary spectra (Fig. 9a), the histogram for the derivative spectra
583 (Fig. 9b) shows a substantial shift to higher values of the cophenetic index. Specifically, there is

584 a significant increase in the frequency of occurrence of high values (> 0.9) of the cophenetic
585 index. This increased frequency is associated with a broader overall spectral region, or
586 equivalently a larger number of spectral ranges, for which the index is higher than 0.9.

587 Whereas the above results illustrate the advantages of derivative analysis, such analysis can
588 be highly sensitive to parameters chosen for the calculation of derivative spectra, specifically the
589 size of the filter window (WS) used in the spectral smoothing of the ordinary spectra and the
590 band separation (BS) used in the calculation of derivatives. To examine the sensitivity of cluster
591 analysis to the selection of these parameters, we computed the distribution of cophenetic index
592 between the pigment-based and absorption-based cluster trees using the second derivative
593 spectra of $a_{n,ph}^*(\lambda)$ within the spectral range of 420 - 515 nm as input (Fig. 10). This spectral
594 range is adequate for this sensitivity analysis because it showed very high values of cophenetic
595 and Rand indices in Fig. 7e and Fig. 8e. The distribution of cophenetic index in Fig. 10 is shown
596 with the smoothing parameter WS varying from 1 to 29 consecutive samples (with a step of 2
597 samples) along the y-axis and the band separation parameter BS varying also from 1 to 29
598 samples with a step of 2 samples along the x-axis. Note that the number of consecutive samples
599 is equivalent to the wavelength interval in nanometers because our spectral data (samples) have
600 the resolution of 1 nm. The highest values of the cophenetic index are obtained for intermediate
601 values of WS and BS around 9 nm – 10 nm. These are the optimal values for our derivative
602 analysis of absorption spectra. This result is consistent with the general expectation that if the
603 values of WS and BS are too small, the derivative spectra are sensitive to noise and exhibit false
604 spectral features, and on the other hand if the WS and BS are too large, the real significant
605 spectral features get smoothed out and essentially removed from the analysis. As the best
606 compromise, the cluster analyses presented in Fig. 8 and 9b were obtained with WS and BS of 9
607 nm.

608

609 *3.3. Classification of stations based on remote-sensing reflectance*

610 The relationship between the spectral remote-sensing reflectance, $R_{rs}(\lambda)$, of the ocean and
611 phytoplankton pigment composition is less direct and far more complicated than that for the
612 spectral phytoplankton absorption coefficient, mainly due to the presence of many optically
613 significant non-phytoplankton constituents in seawater. The investigation of $R_{rs}(\lambda)$ is of
614 particular interest, however, because information contained in this measurement provides a
615 potential means for remote-sensing applications. Fig. 11 shows the Hydrolight-simulated $R_{rs}(\lambda)$
616 spectra normalized at 555 nm for the nine stations identified as classes from A (station 1) through
617 F (station 59). In general, there are significant differences in the UV and blue spectral regions
618 between these normalized spectra, with the largest contrast between the classes A (station 1) and
619 C1 (station 12).

620 The dendrograms obtained from cluster analysis as applied to four different sets of input data
621 vectors containing information about remote-sensing reflectance (as described in sec 2.3) are
622 displayed in Fig. 12. The limited spectral information, i.e., the three reflectance band ratios used
623 commonly in satellite ocean color applications (Fig. 12a) and the 13 band ratios corresponding to
624 multispectral measurements with the SPMR instrument (Fig. 12b), provide a very dissimilar
625 classification of stations compared with the pigment-based cluster analysis (see Fig. 4a). All
626 stations, with only the exception of class A, show very little separation in the cluster-tree based
627 on multispectral reflectance data. The high spectral resolution (1 nm) normalized reflectance
628 spectra over the entire spectral range 300 – 725 nm also produce a dendrogram (Fig. 12c) that is
629 very different from the pigment-based cluster tree. Although the stations belonging to class C
630 are closer to one another compared with the multispectral-based cluster tree, they are grouped
631 together with two other stations (class E and D). In addition, station B forms a separate single-
632 object cluster in Fig. 12c, whereas it is grouped in a single multi-object cluster together with the
633 stations from class C in the pigment analysis. The only case when the cluster analysis of
634 reflectance data provides a high degree of similarity with pigment analysis (i.e., Rand index of

635 0.78) is for the second derivative of hyperspectral normalized reflectance over the entire spectral
636 range 327 - 698 nm (Fig. 12d). We note that the derivative reflectance spectra were calculated
637 with the parameters *WS* and *BS* of 27 nm (as supported by the sensitivity analysis discussed
638 below). The stations A, E, and F in Fig. 12d form single-object clusters at a significant distance
639 from the remaining stations. Similarly to pigment analysis, the stations C1, C2, C3, C4, and B
640 are grouped relatively close to one another. However, station D also belongs to that group,
641 which is not the case in the pigment-based cluster tree. This may be attributable to the fact that
642 *Zea* and *DVChla*, which are the two most dominant diagnostic pigments at stations C1, C2, C3,
643 C4, and B, also play a significant role at station D where they are ranked as the second and third
644 most important diagnostic pigments (see Table 1).

645 The progression of linkage distances corresponding to the four dendrograms from Fig. 12
646 clearly illustrates the advantage of the second derivative spectra over the multispectral data or
647 non-differentiated spectra of reflectance (Fig. 13). The improved separation seen in terms of the
648 larger linkage distances between the clusters of stations obtained with derivative spectra
649 indicates that this approach enables better identification of the differences in the magnitude and
650 shape between the high resolution spectra. In contrast, in the analysis of multispectral data and
651 ordinary spectra, stations are linked at a very small distance, which indicates that these types of
652 reflectance data will be essentially useless for obtaining information about pigment assemblages
653 from cluster analysis.

654 The improvement in the similarity between the pigment-based and reflectance-based
655 classification achieved with the use of second derivative spectra as opposed to multispectral data
656 or ordinary spectra of reflectance is presented in Table 2. The results for the derivative analysis
657 of hyperspectral normalized $R_{rs}(\lambda)$ over the entire spectral range 300 - 725 nm show a significant
658 increase in both the cophenetic and Rand index when compared with multispectral and ordinary
659 spectral data. However, the best performance is obtained when the derivative analysis is

660 restricted to the spectral range from 435 to 495 nm (as supported by the sensitivity analysis
661 discussed below). In this case, the similarity indices are highest.

662 Fig. 14 shows distributions of the cophenetic and Rand index which identify the optimal
663 spectral ranges for the cluster analysis of the second derivative of remote-sensing reflectance.
664 The cophenetic index is generally close or slightly higher than 0.5 for most spectral ranges
665 examined, that is for most combinations of λ_{\min} and λ_{\max} (Fig. 14a). In the spectral region from
666 435 nm to 510 nm, this index is about 0.65. This can be considered as an optimal spectral range
667 for the application of derivative approach with potential for good similarity between the
668 pigment-based and reflectance-based cluster trees. This result is also supported by very high
669 value of the Rand index of 0.86 in that spectral range (Fig. 14b). The Rand index attains even
670 higher value of about 1 within somewhat narrower wavelength range from $\lambda_{\min} = 435$ nm to λ_{\max}
671 $= 495$ nm, which defines an alternative optimal spectral range. In addition, the Rand index
672 suggests good performance of the derivative-based analysis over a broader spectral region
673 including shorter wavelengths of λ_{\min} from the near-UV. In general, this index is quite high for
674 λ_{\min} varying between 350 nm and 450 nm, for example as high as 1 when the spectral range is
675 from $\lambda_{\min} = 365$ nm to $\lambda_{\max} = 480$ nm. However, because the optical roles of different diagnostic
676 pigments in the near-UV are insignificant or certainly less important than in the blue region, the
677 use of the spectral range 435 – 510 nm or 435 – 495 nm, where both the cophenetic and Rand
678 indices are relatively high, appears to be most reasonable.

679 The effects of the size of the filter window (*WS*) used in the smoothing of the ordinary
680 spectra and the band separation (*BS*) used in the calculation of derivatives on the similarity
681 between the cluster trees from the analysis of pigments and the second derivative reflectance
682 spectra is illustrated in Fig. 15. These results are shown for the derivative spectra calculated over
683 one of the optimal spectral ranges, specifically 435 - 495 nm, which showed high values for both
684 cophenetic and Rand indices. The best similarity with cophenetic index of about 0.65 and the
685 Rand index of 1 is obtained when the calculations of second derivative spectra are made with

686 relatively large values of *WS* and *BS*. For example, a very good result is obtained if both *WS* and
687 *BS* assume a value of 27 consecutive spectral samples (i.e., 27 nm as the resolution of our
688 hyperspectral reflectance data is 1 nm). We recall that this value was used to compute the results
689 pertinent to the derivative reflectance spectra presented in Figs. 12 - 14. Similarly good results
690 are obtained with a smaller *WS* (~14 nm) and a larger *BS* (~37 nm), or vice versa. In contrast, if
691 both *WS* and *BS* are small (less than about 10 nm) or large (above ~40 nm) the cophenetic and
692 Rand indices are reduced significantly. This sensitivity analysis, in agreement with similar
693 analysis for absorption spectra, supports strong dependence of the derivative-based cluster trees
694 on the selection of parameters *WS* and *BS* for the derivative calculations.

695

696 **4. Conclusions**

697 By applying the unsupervised hierarchical cluster analysis to pigment and optical data from
698 the eastern Atlantic Ocean we demonstrated the potential usefulness of hyperspectral data of
699 absorption coefficient and remote-sensing reflectance for discriminating different phytoplankton
700 pigment assemblages in the open ocean under non-bloom conditions. The most promising
701 results were obtained with the second derivative spectra of phytoplankton absorption coefficient
702 covering the spectral range as wide as 370 nm - 725 nm (or narrower spectral regions from
703 within that range), and the second derivative spectra of remote-sensing reflectance over the
704 spectral range from about 435 nm to 510 nm. Our ability to discriminate different phytoplankton
705 pigment assemblages from the derivative-based cluster trees was optimized by selecting the most
706 suitable parameters used in the spectral derivative calculations (see also Torrecilla et al., 2009).
707 In particular, for these optical data with a 1 nm resolution, we determined that the optimal values
708 for the smoothing filter window and band separation used in the calculations of second derivative
709 spectra are 9 nm for absorption and 27 nm for remote-sensing reflectance. These derivative
710 spectra are presented in Fig. 16 for the nine Atlantic stations selected for this study. The

711 assessment of similarity between these derivative spectra and phytoplankton pigment
712 composition was made using the similarity-based cluster algorithm and two indices, cophenetic
713 and Rand. In this cluster algorithm, the partitioning of data into clusters is based on the
714 determinations of the angular distance between each pair of examined input data vectors. This
715 type of similarity-based cluster technique accounts for complete spectral behavior of optical data
716 with no need to identify specific spectral features. This approach is particularly useful in cases
717 when it is difficult or impossible to define explicitly a set of unambiguous diagnostic spectral
718 features in the original optical data (Duin et al., 1997; Pekalska and Duin, 2000).

719 In addition to the second derivative spectra of phytoplankton absorption and remote-sensing
720 reflectance, we examined other absorption and reflectance data but they generally showed either
721 more limited value or no usefulness at all for discriminating phytoplankton pigment
722 assemblages. For example, the cluster analysis of the ordinary (non-differentiated) reflectance
723 spectra at 1 nm resolution or multispectral (13 wavebands) reflectance data showed very poor
724 similarity with pigment-based clusters. Similar results were obtained for the ordinary spectra of
725 the total absorption coefficient. However, the ordinary spectra of phytoplankton absorption are
726 useful, especially within the spectral range 425 nm - 540 nm.

727 We demonstrated that the quantification of similarity between the optical-based clusters and
728 pigment-based clusters with the cophenetic and Rand indices provides a valuable methodology
729 for identifying optical variables and their spectral ranges most suitable for characterizing the
730 phytoplankton pigment assemblages, and for selecting the optimal values of the parameters used
731 in the calculation of derivative spectra. Whereas the present study uses a limited data set, albeit
732 carefully selected to represent distinct differences in phytoplankton assemblages in terms of
733 dominant accessory pigments, further work is needed to evaluate or refine the proposed
734 methodology with larger data sets from various oceanic environments. Given significant interest
735 in the development of the capabilities for large-scale characterization of phytoplankton
736 biodiversity from optical measurements including remote-sensing observations, one may expect

737 further expansion of comprehensive databases consisting of simultaneously collected pigment
738 and hyperspectral optical data in the near future. We expect that this will support further work on
739 the cluster-based approach and other techniques, such as neural networks (Raitsos et al., 2008;
740 Aymerich et al., 2009), which exploit optical measurements as a source of information on
741 phytoplankton community composition.

742

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752

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921

922 Table 1. Summary of phytoplankton pigment data for the nine stations selected in this study.
 923 The stations are sorted into different classes characterized by differing pigment assemblages
 924 based upon the ratios of the concentrations of two dominant accessory pigments to total
 925 chlorophyll-*a*, TChl*a* (see the 2nd and 3rd columns from the left). The ratios of the
 926 concentrations of six dominant pigments to TChl*a* are also displayed, with the two most
 927 dominant accessory pigments indicated within the shaded areas. Pigment abbreviations are:
 928 MVChl*a* = monovinyl chlorophyll-*a*, DVChl*a* = divinyl chlorophyll-*a*, Fuco = fucoxanthin, Hex
 929 = 19' hexanoyloxyfucoxanthin, But = 19' butanoyloxyfucoxanthin, MVChl*b* = monovinyl
 930 chlorophyll-*b*, Chl*c*2 = chlorophyll-*c*2, Zea = zeaxanthin, Pra = prasinoxanthin, Dia =
 931 diadinoxanthin, and α -caro = α -carotene.

932

Station ID	2 dominant pigments	Class	TChl <i>a</i> [mg/m ³]	Ratio of concentrations of dominant pigments to TChl <i>a</i>					
				MVChl <i>a</i>	Fuco	MVChl <i>b</i>	Chl <i>c</i> 2	Pra	Hex
1	Fuco \approx MVChl <i>b</i>	A	0.62	0.95	0.22	0.21	0.11	0.07	0.07
6	DVChl <i>a</i> > Zea	B	0.28	0.50	0.47	0.31	0.18	α -caro	Chl <i>c</i> 2
12	DVChl <i>a</i> \approx Zea	C1	0.14	0.54	0.44	0.40	0.24	0.09	0.09
37	DVChl <i>a</i> \approx Zea	C2	0.15	0.63	0.58	0.40	0.17	0.10	α -caro
44	DVChl <i>a</i> \approx Zea	C3	0.22	0.56	0.50	0.42	0.17	0.08	0.08
46	DVChl <i>a</i> \approx Zea	C4	0.14	0.52	0.50	0.49	0.20	0.09	0.08
48	Hex > Zea	D	0.21	0.76	0.48	0.25	0.21	0.18	0.16
51	Hex > Fuco	E	0.26	0.86	0.35	0.24	0.23	0.18	0.13
59	Zea \approx Hex	F	0.11	0.80	0.39	0.34	0.21	0.18	0.12

933
 934
 935

936 Table 2. A comparison of similarity indices between pigment-based clusters and reflectance-
 937 based clusters for the different sources of reflectance data that are depicted in Fig. 12. For the
 938 case of the second derivative of hyperspectral reflectance, the result of computations for two
 939 different spectral regions is given.

Reflectance data	Rand index	Cophenetic index
3 band ratios based on 4 SeaWiFS bands	0.69	0.39
Multispectral (13 bands)	0.69	0.39
Hyperspectral (325 bands, range 300-725 nm)	0.69	0.39
Hyperspectral 2nd derivative (300-725 nm)	0.78	0.53
Hyperspectral 2nd derivative (435-495 nm)	1	0.65

940

941

942 **Figure captions**

943 Figure 1. Map depicting the location of full stations sampled along the north-to-south ANT-
944 XXIII/1 cruise track in the eastern Atlantic during October and November, 2005. Each full
945 station consisted of in situ optical measurements accompanied by discrete water sample analyses.
946 Stations chosen for use in the cluster analysis are identified by filled circles and labeled with the
947 station ID.

948

949 Figure 2. Hyperspectral (1 nm) determinations of the remote-sensing reflectance $R_{rs}(\lambda)$ obtained
950 from radiative transfer simulations (solid line) compared with in situ multispectral measurements
951 at 13 discrete bands (solid circles). Each panel illustrates a different station location.

952

953 Figure 3. A schematic diagram illustrating the general approach to hierarchical cluster analysis
954 and similarity determination. The dendrogram obtained with pigment composition as the input
955 (upper pathway) is used as the reference for comparison with results obtained utilizing various
956 optical data as input.

957

958 Figure 4. (a) Dendrogram obtained for the nine stations using 24 pigment to total chlorophyll-*a*
959 (TChl*a*) ratios determined from the CHORS HPLC analysis. (b) Linkage distances obtained
960 from the cluster analysis shown in (a) as a function of distance along the dendrogram. (c)
961 Similar to (a), but using input ratios calculated from 8 diagnostic pigments (see text for details).

962

963 Figure 5. Chlorophyll-specific normalized absorption coefficients for the nine stations

964 corresponding to: (a) absorption of phytoplankton, $a_{n,ph}^*(\lambda)$, (b) absorption of pure seawater plus
965 phytoplankton, $a_{n,w+ph}^*(\lambda)$, and (c) total absorption, $a_n^*(\lambda)$.

966

967 Figure 6. Results of cluster analysis applied to absorption data from the nine stations. The left
968 panels represent dendrograms obtained using the different absorption components of (a) $a_n^*(\lambda)$,
969 (c) $a_{n,w+ph}^*(\lambda)$, and (e) $a_{n,ph}^*(\lambda)$, and the right panels (b, d, f) illustrate results obtained using
970 each respective component's second derivative spectrum.

971

972 Figure 7. Similarity indices between absorption-based and pigment-based cluster trees obtained
973 for the nine stations using different combinations of spectral range for (a) $a_n^*(\lambda)$, (b) $a_{n,w+ph}^*(\lambda)$
974 and (c) $a_{n,ph}^*(\lambda)$. The y-axis indicates the lower limit of the spectral range (λ_{\min}) and the x-axis
975 the upper limit of the spectral range (λ_{\max}) utilized in the cluster analysis. Left and right panels
976 depict the cophenetic and Rand indices, respectively.

977

978 Figure 8. Similar to Fig. 7, but based on cluster trees obtained using different spectral range
979 combinations for the second derivative spectra of (a) $a_n^*(\lambda)$, (b) $a_{n,w+ph}^*(\lambda)$ and (c) $a_{n,ph}^*(\lambda)$.
980 Optimal values for band separation and window size determined from prior analyses ($BS = WS =$
981 9 nm, see text for details) were used in the calculation of derivative spectra.

982

983 Figure 9. Histograms of cophenetic indices obtained for all combinations of spectral ranges
984 shown in Fig. 8 based on (a) the absorption of phytoplankton, $a_{n,ph}^*(\lambda)$, and (b) its second
985 derivative spectra.

986

987 Figure 10. Cophenetic indices obtained from the cluster and similarity analysis of the second
988 derivative spectra of $a_{n,ph}^*$, in which different parameter sets for the derivative analysis are
989 considered. The analysis was conducted using the optimal spectral region from 420 to 515 nm.
990 The y-axis indicates the size of the filter window used in smoothing of the absorption spectra
991 (WS), and the x-axis represents the band separation used in the calculation of the derivative
992 calculation (BS).

993

994 Figure 11. Hydrolight-simulated $R_{rs}(\lambda)$ spectra, normalized at 555 nm, computed for the nine
995 stations using measured IOPs as input.

996

997 Figure 12. Dendrograms resulting from cluster analysis of the nine stations calculated using four
998 different sets of input data vectors: (a) three reflectance band ratios of $R_{rs}(\lambda)$ based on 4 SeaWiFS
999 wavebands obtained from measurements with SPMR instrument, (b) 13 band ratios
1000 corresponding to multispectral measurements of $R_{rs}(\lambda)$ with SPMR instrument, (c) hyperspectral
1001 (1 nm) ordinary (non-differentiated) normalized $R_{rs}(\lambda)$ spectra computed from the Hydrolight
1002 simulations, and (d) second derivative of hyperspectral normalized $R_{rs}(\lambda)$ spectra obtained using
1003 optimal values for band separation and smoothing filter window (i.e., $BS = WS = 27$ nm).

1004

1005 Figure 13. Linkage distances as a function of distance along the dendrogram for each cluster tree
1006 depicted in Fig.12.

1007

1008 Figure 14. Similarity indices between reflectance-based and pigment-based cluster trees
1009 obtained for the nine stations using different combinations of spectral ranges for the second
1010 derivative of the hyperspectral normalized $R_{rs}(\lambda)$. Optimal values of $BS = WS = 27$ nm,
1011 determined from prior analyses, were used for the calculation of derivative spectra. The y-axis
1012 indicates the lower limit of the spectral range (λ_{\min}) and the x-axis the upper limit of the spectral
1013 range (λ_{\max}) utilized in the cluster analysis. Panels (a) and (b) depict the cophenetic and Rand
1014 indices, respectively.

1015

1016 Figure 15. (a) Cophenetic and (b) Rand indices obtained from the comparison of pigment-based
1017 cluster trees with trees computed from the second derivative of hyperspectral normalized $R_{rs}(\lambda)$,
1018 in which different choices for parameters of the derivative analysis are considered. The analysis
1019 has been carried out for the optimal spectral region from 435 and 495 nm. The y-axis indicates
1020 the size of the filter window used in the smoothing of spectra (WS), and the x-axis the band
1021 separation used in the derivative calculation (BS).

1022

1023 Figure 16. The second derivative spectra at each station of the (a) chlorophyll-specific
1024 normalized phytoplankton absorption coefficient, $a_{n,ph}^*(\lambda)$, and (b) normalized hyperspectral
1025 remote-sensing reflectance, $R_{rs}(\lambda)/R_{rs}(555)$. The derivative spectra are depicted for the optimal
1026 spectral ranges of 370 - 716 nm for absorption, and 435 - 510 nm for reflectance. Optimal values
1027 for band separation and smoothing filter window were used for the derivative calculations, i.e.,
1028 $BS = WS = 9$ nm for the absorption data and $BS = WS = 27$ nm for the reflectance data.

Figure 1
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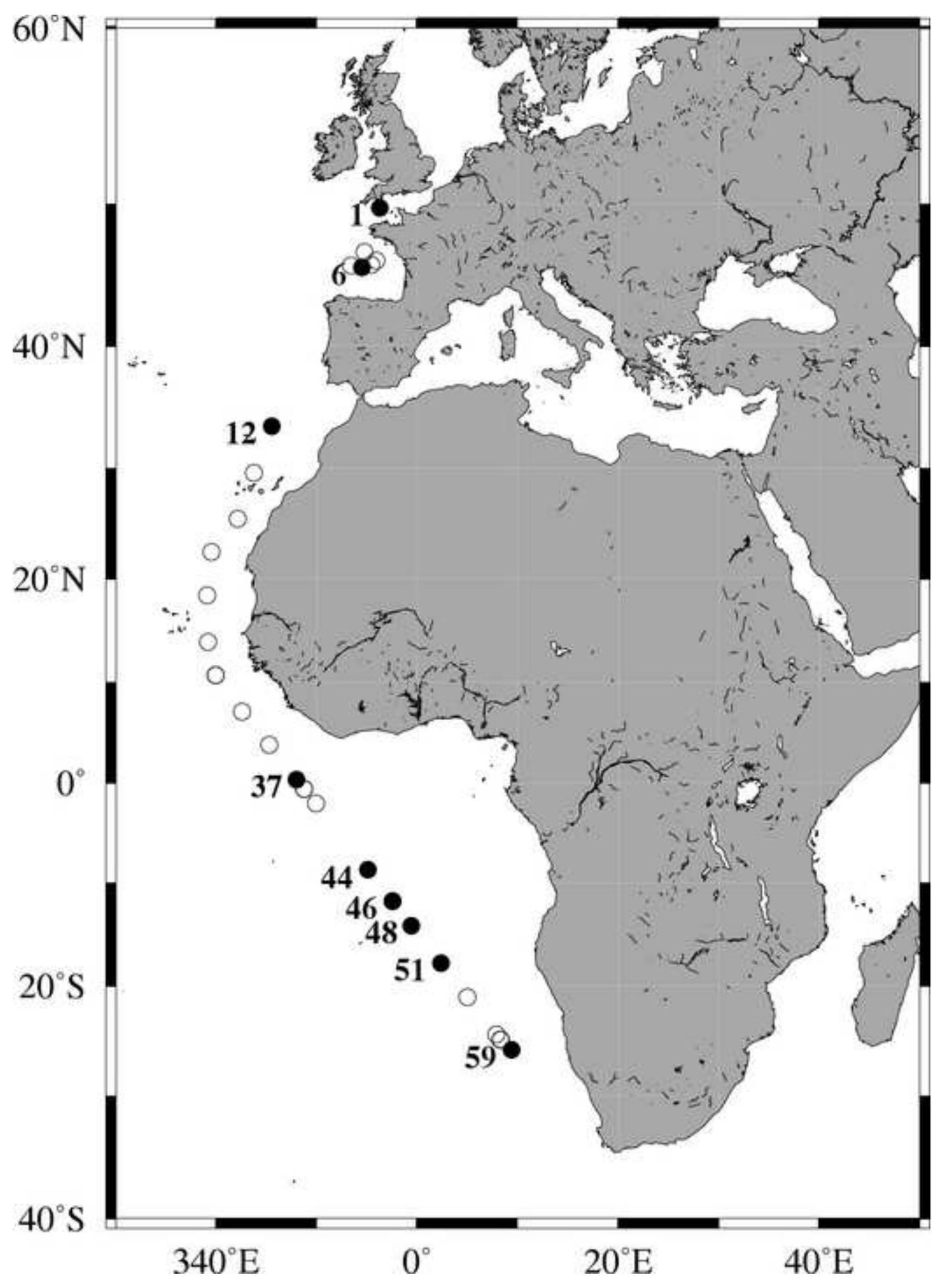


Figure 2

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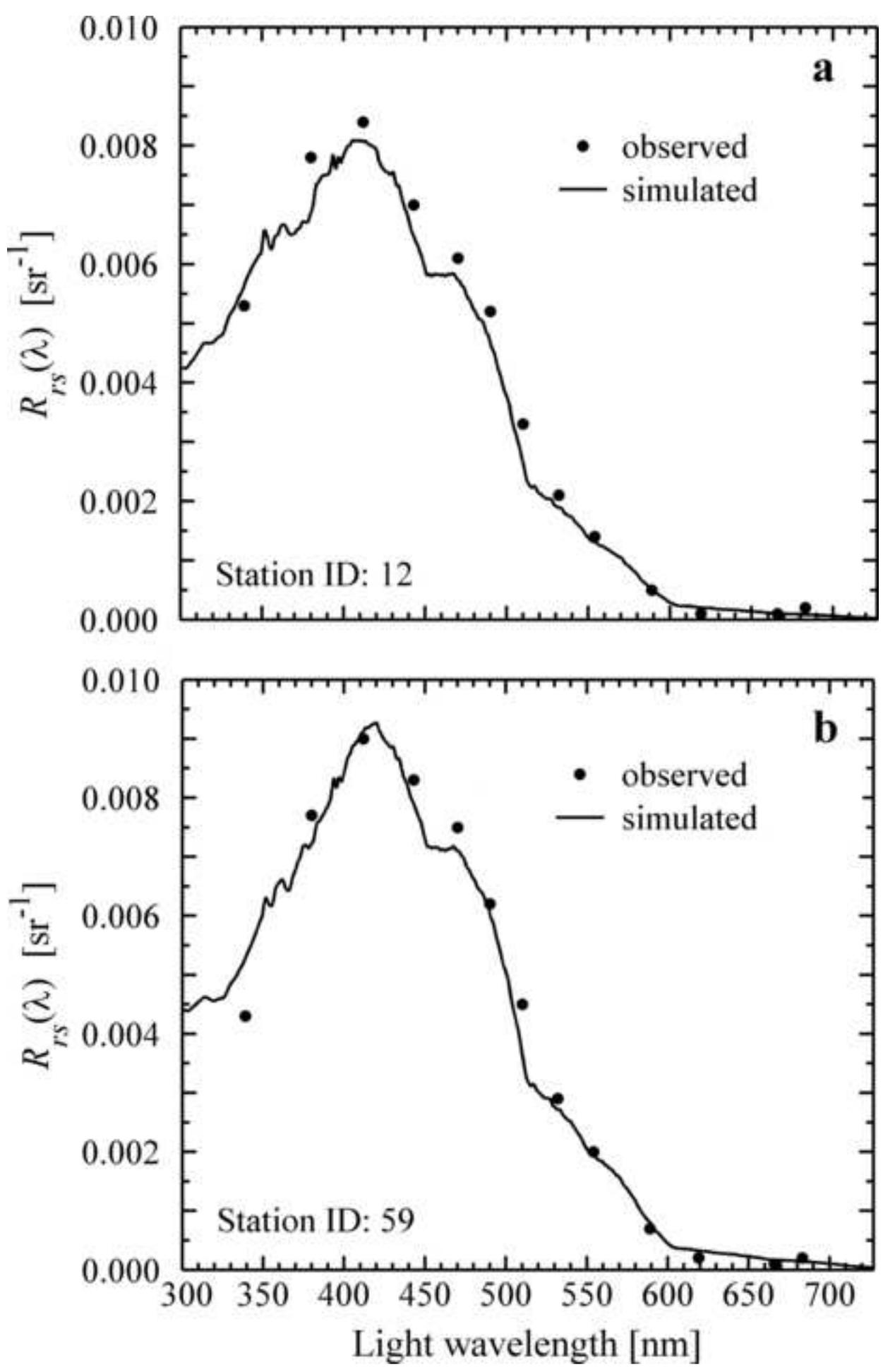


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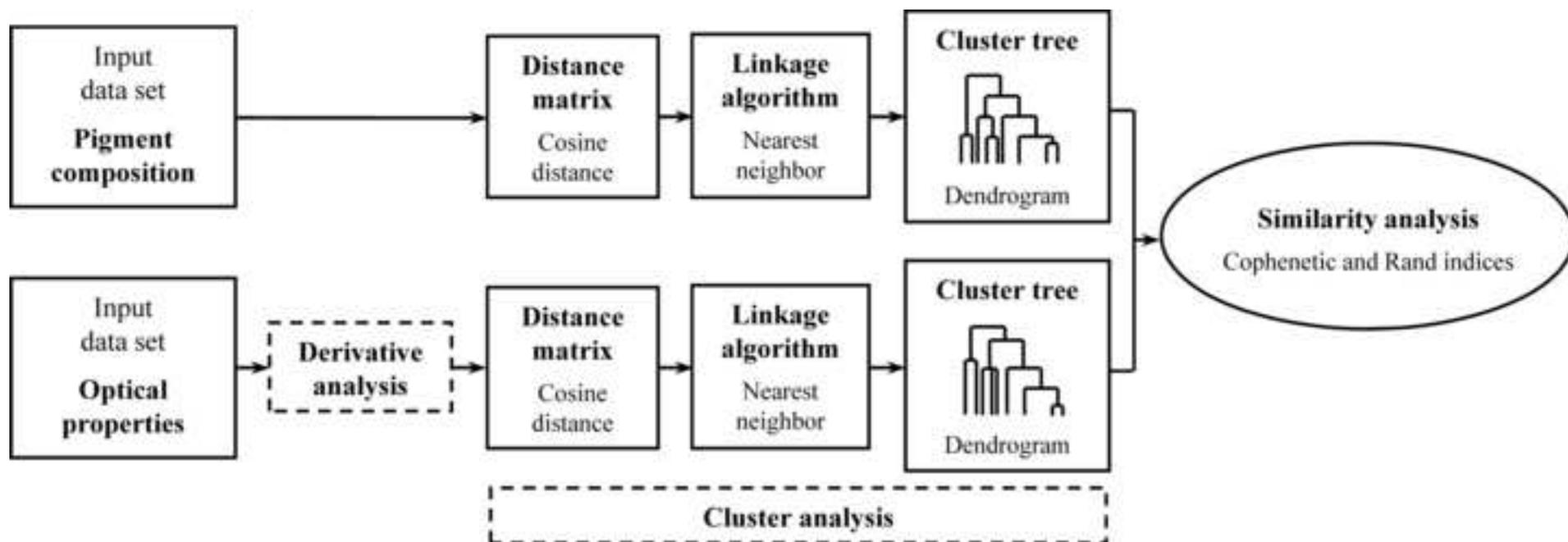


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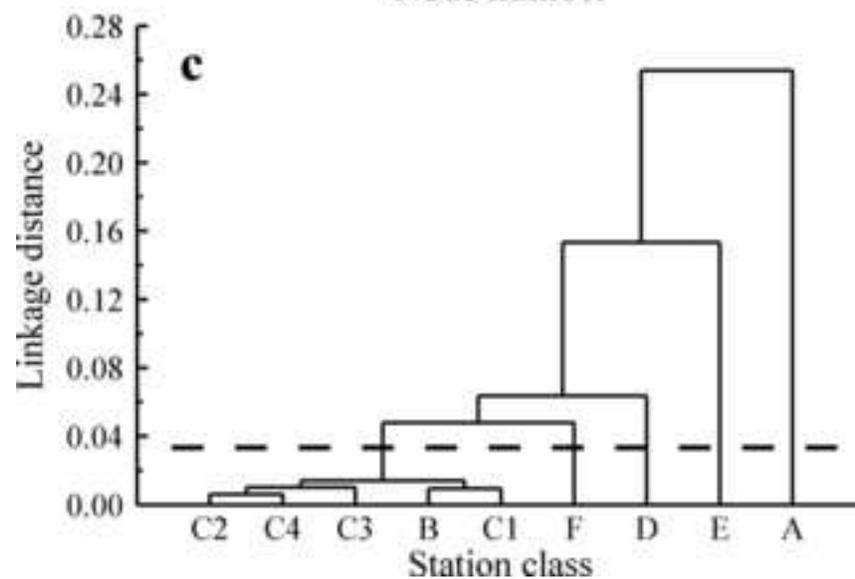
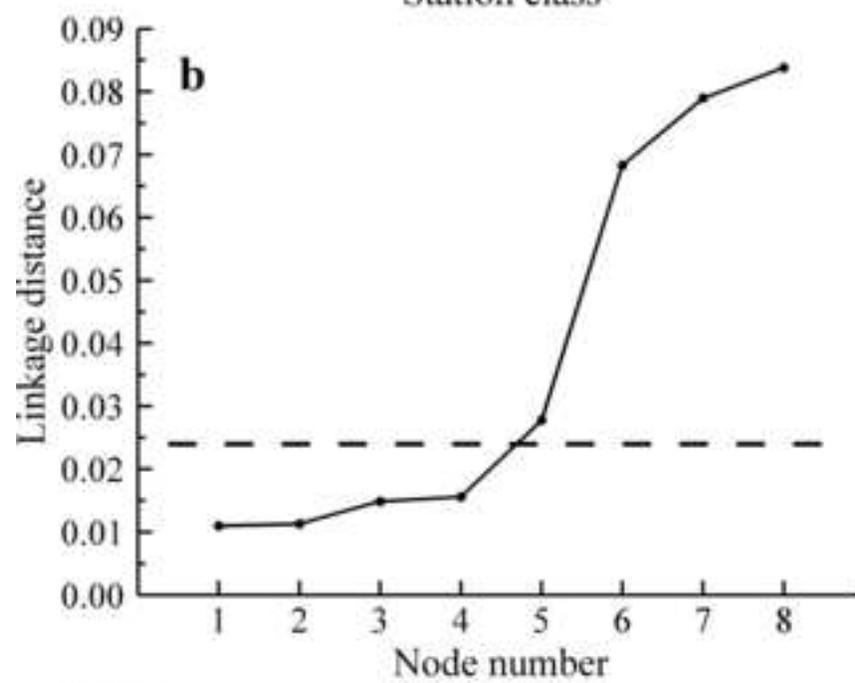
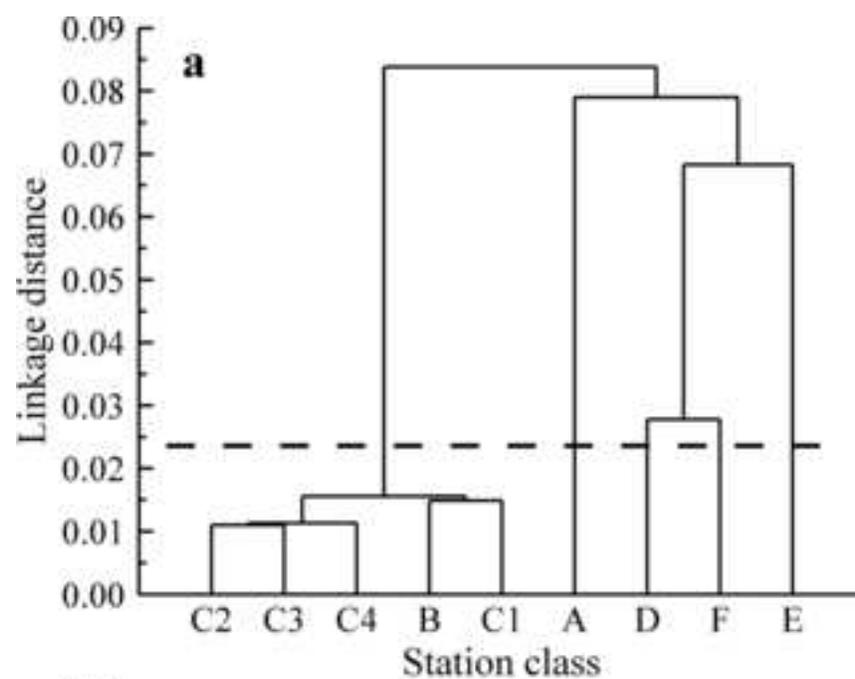


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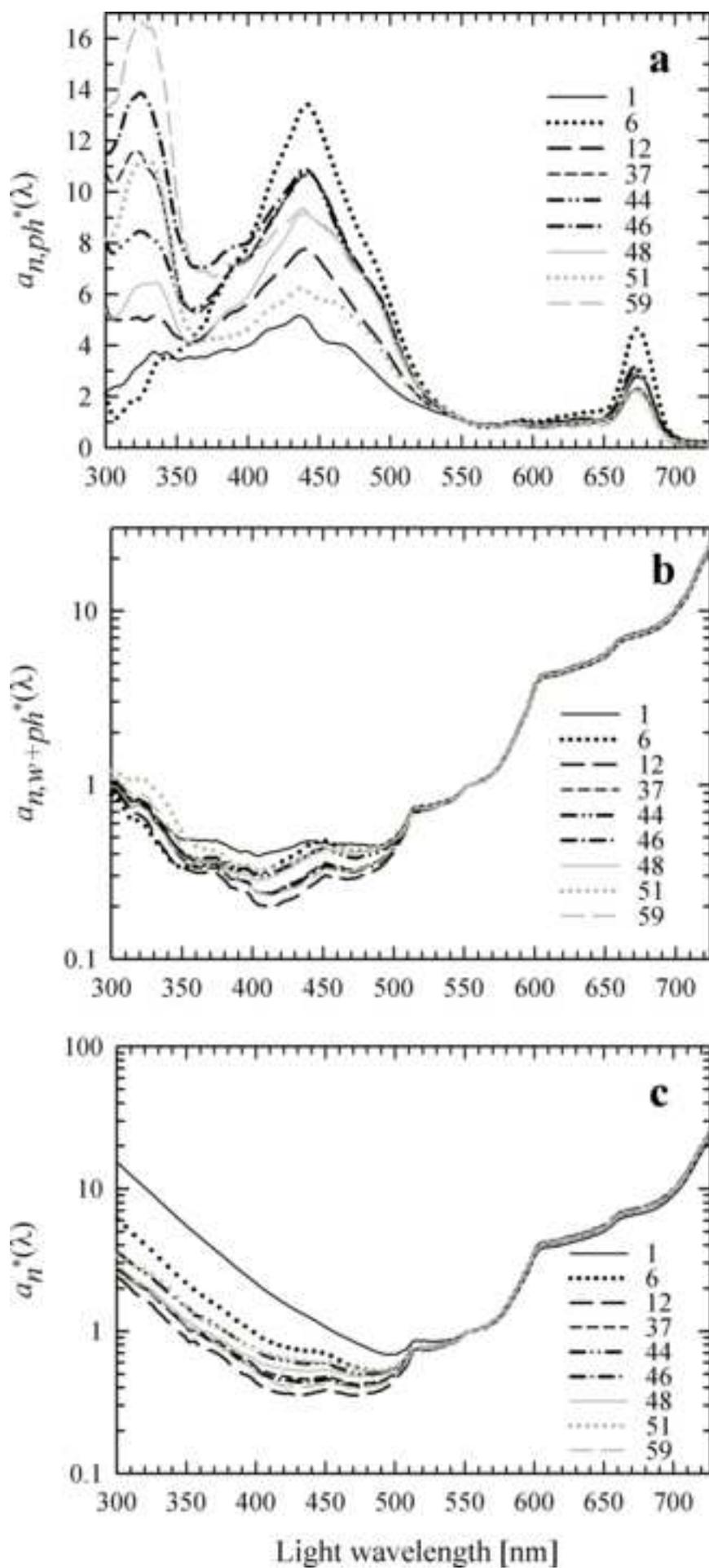


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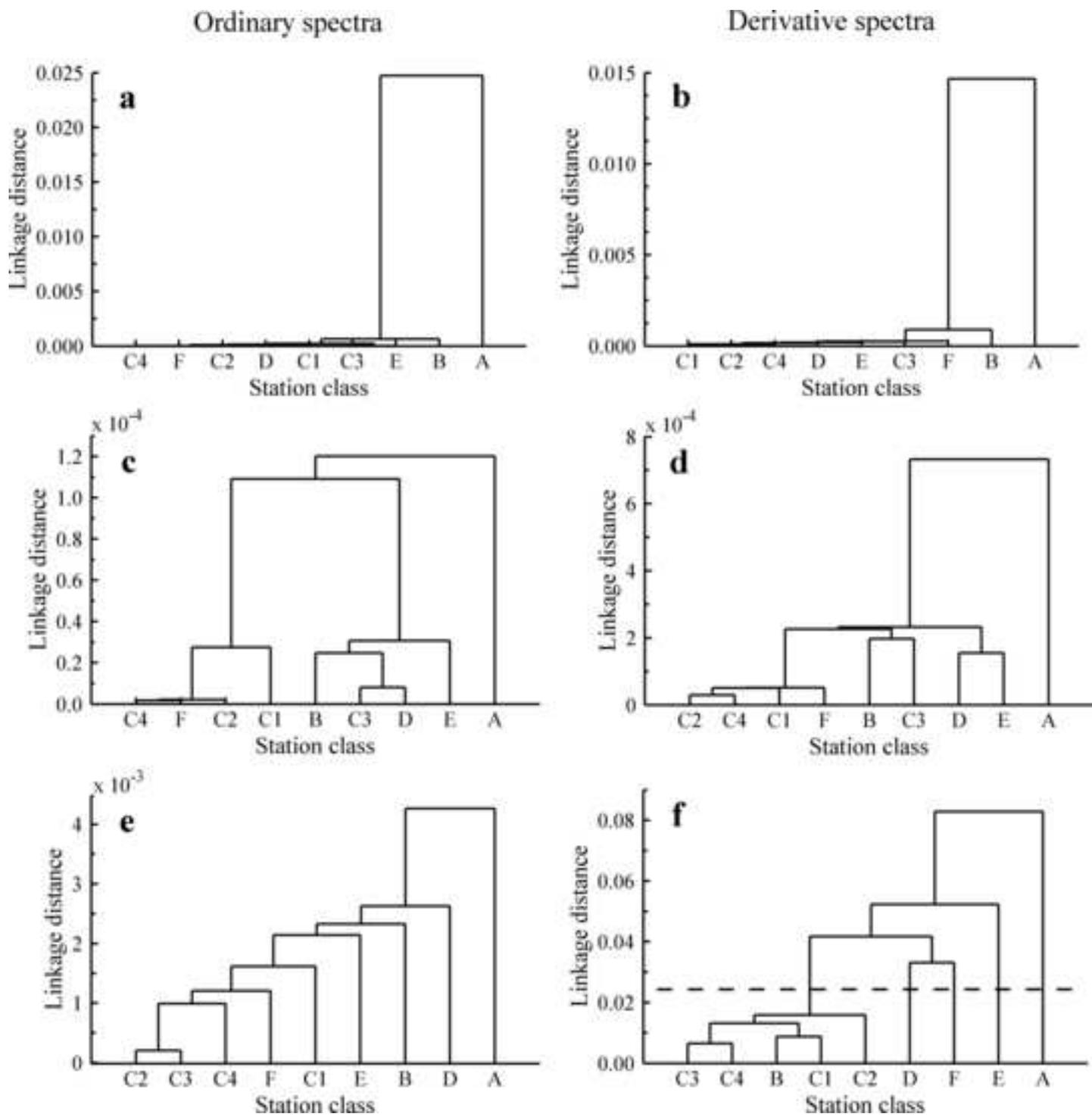
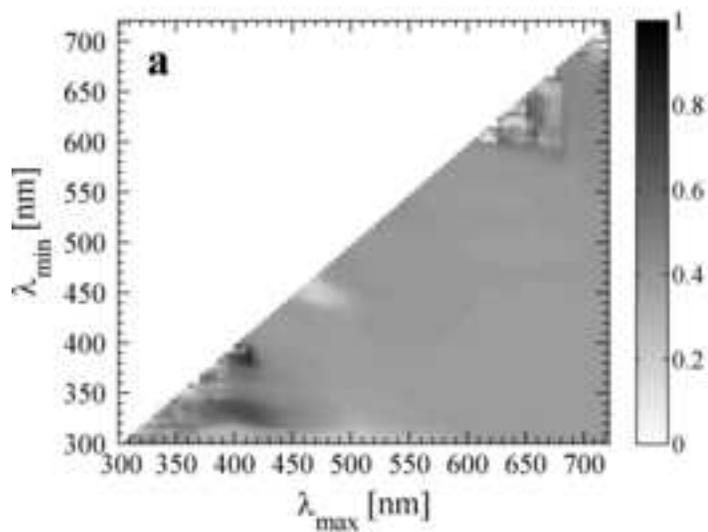


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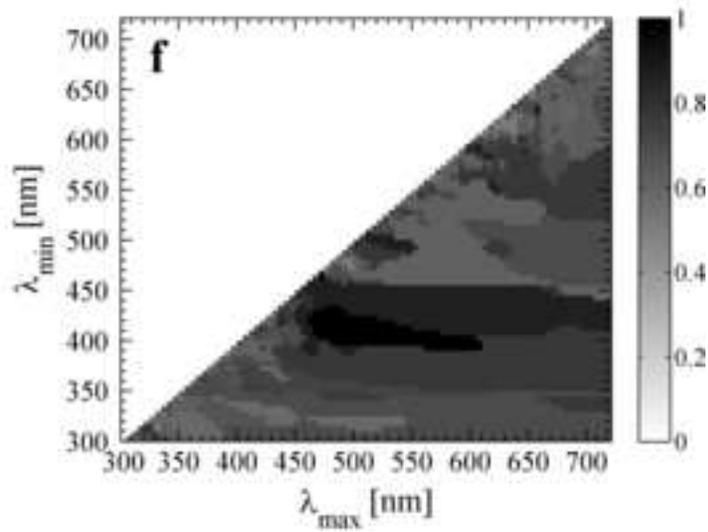
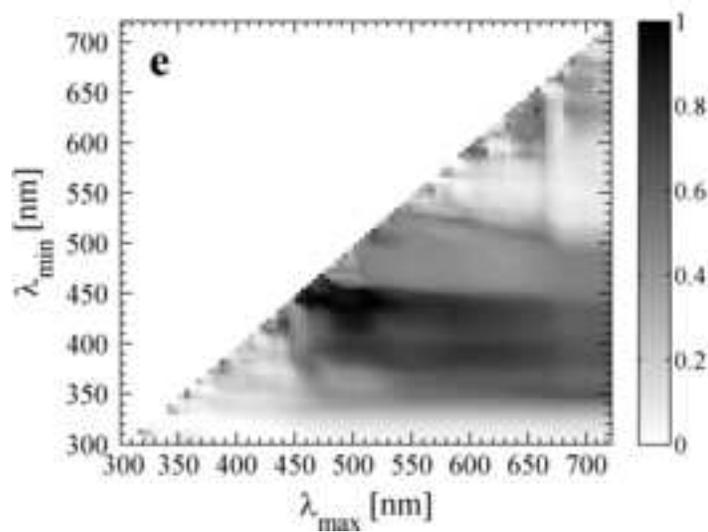
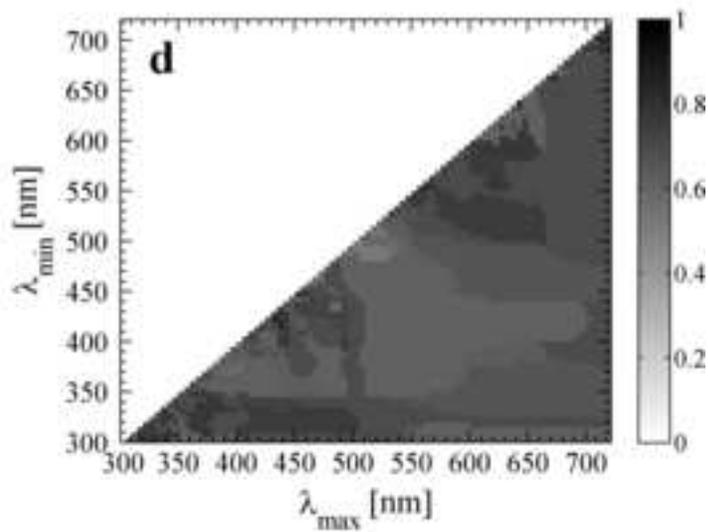
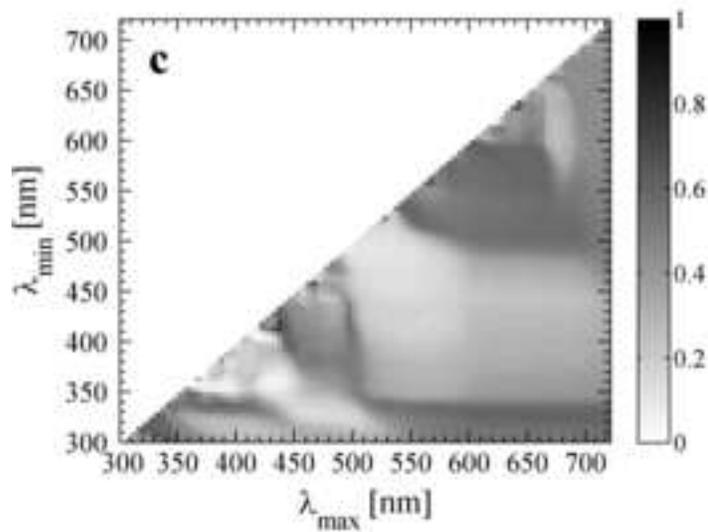
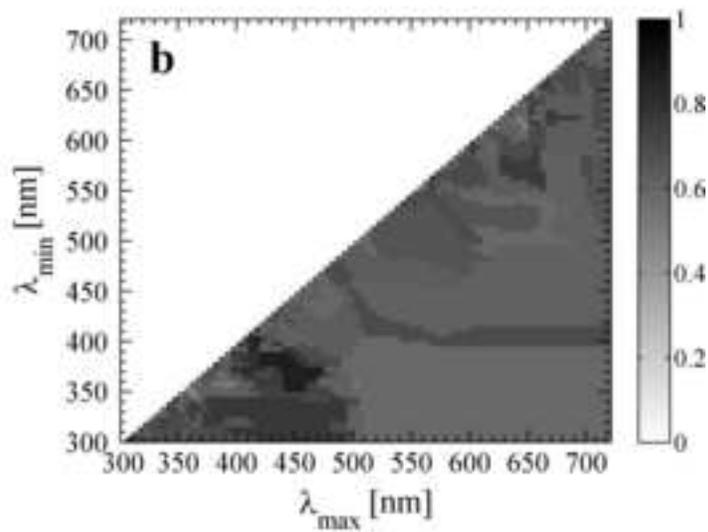


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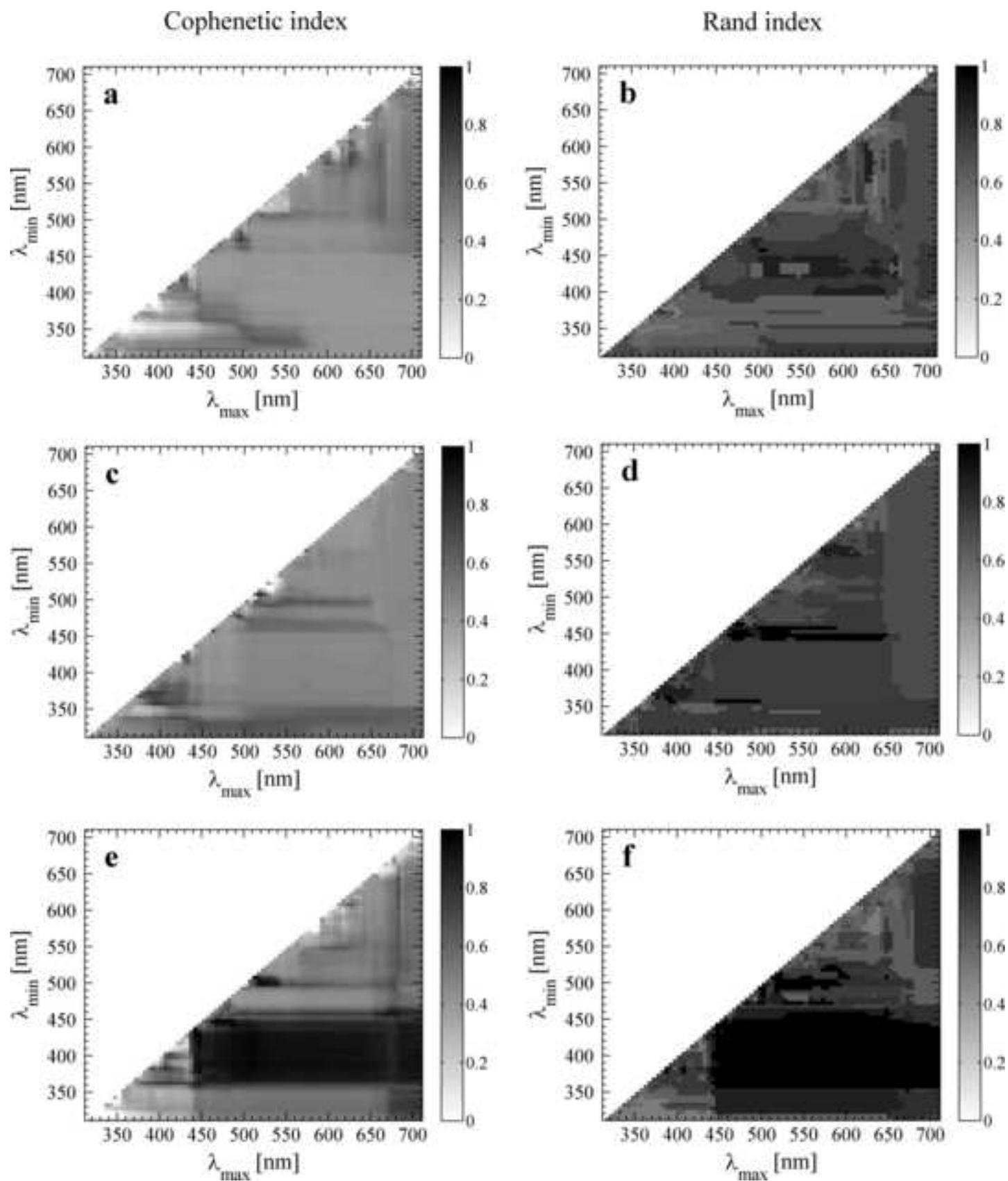


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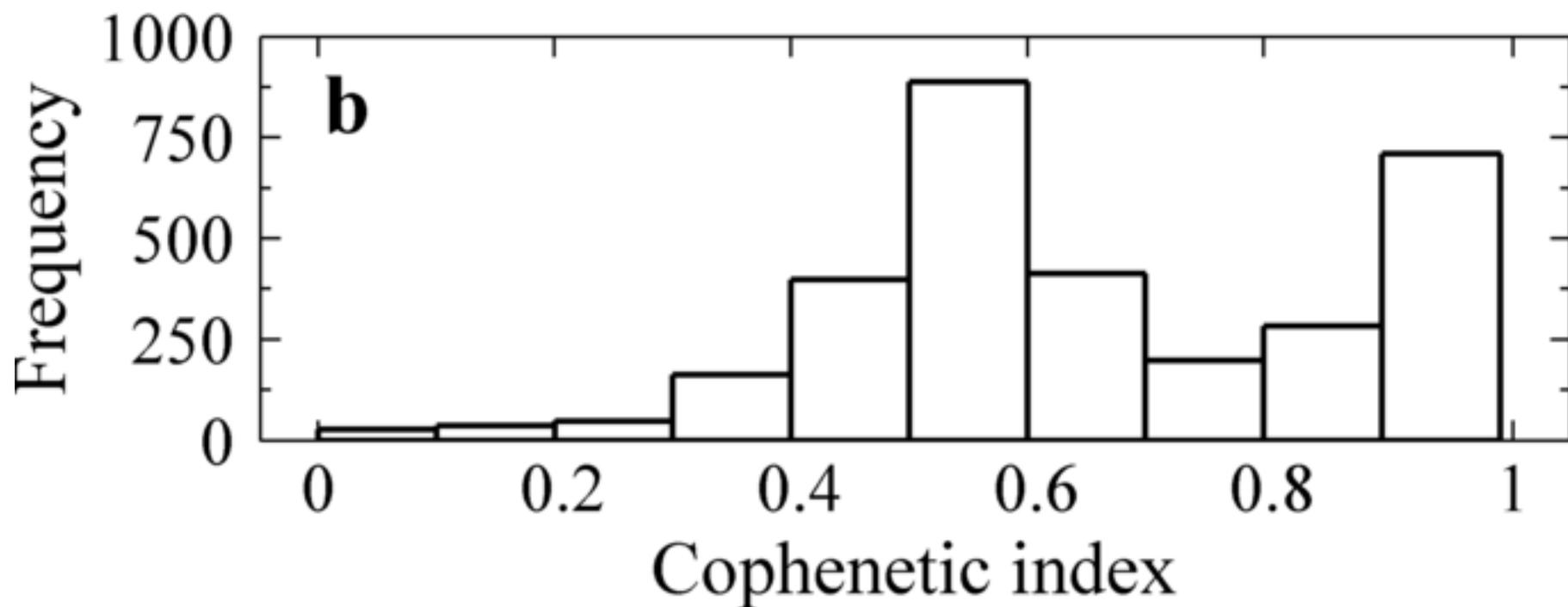
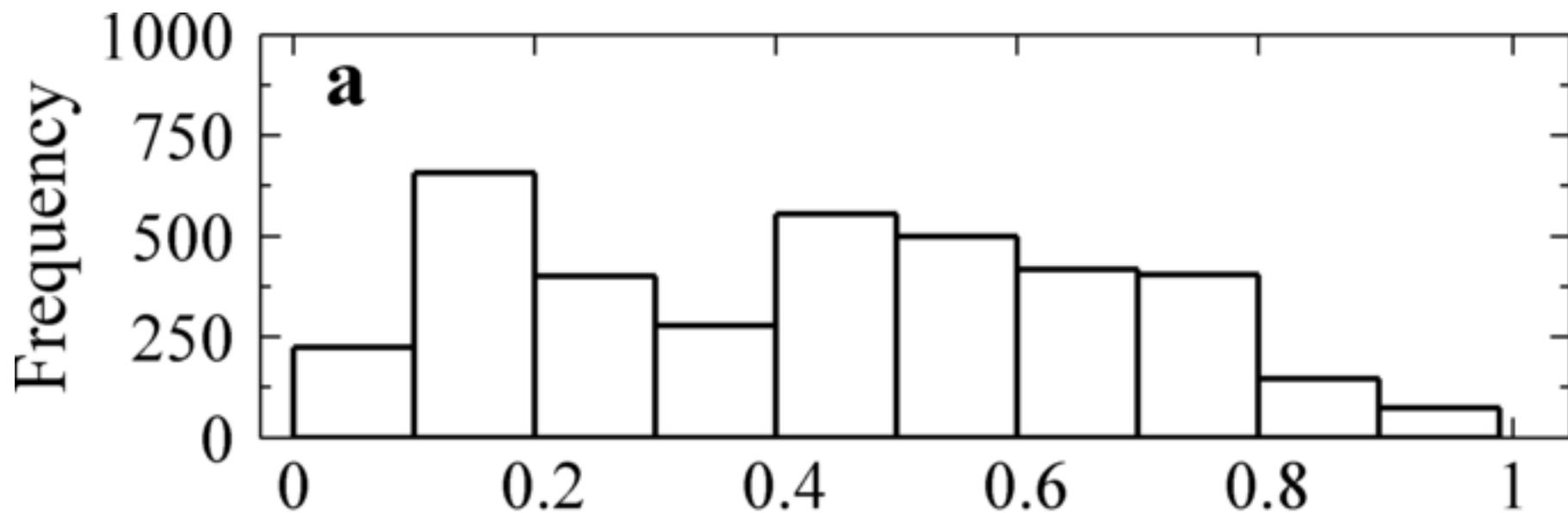


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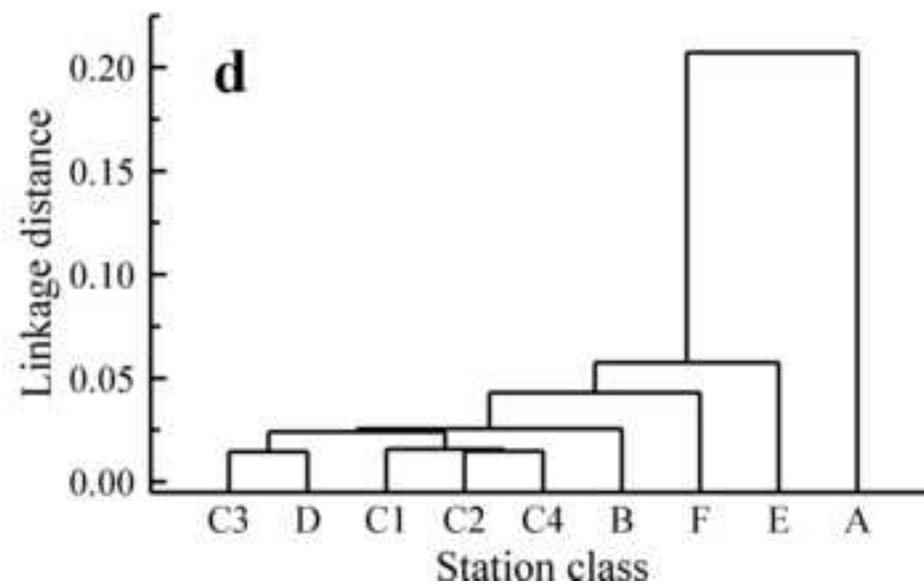
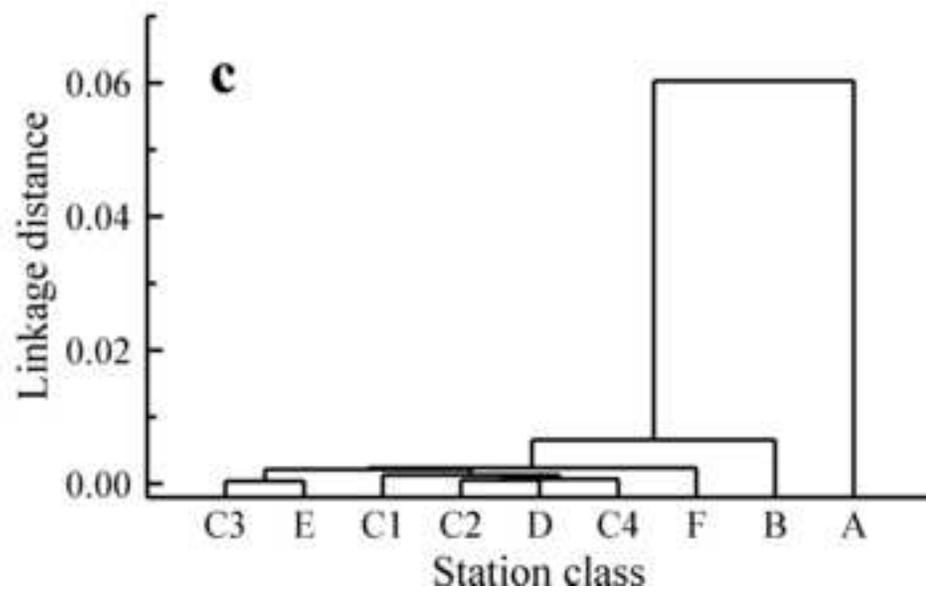
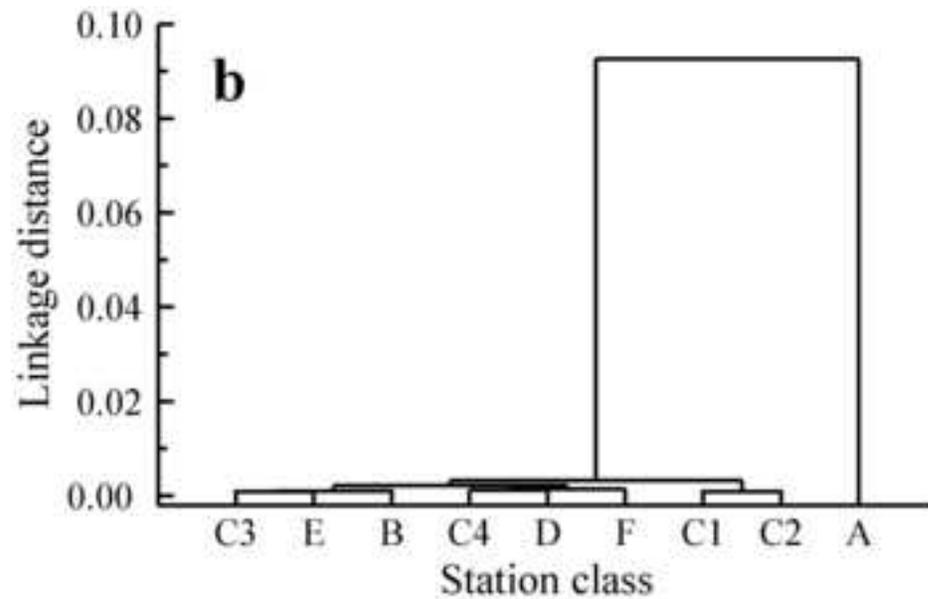
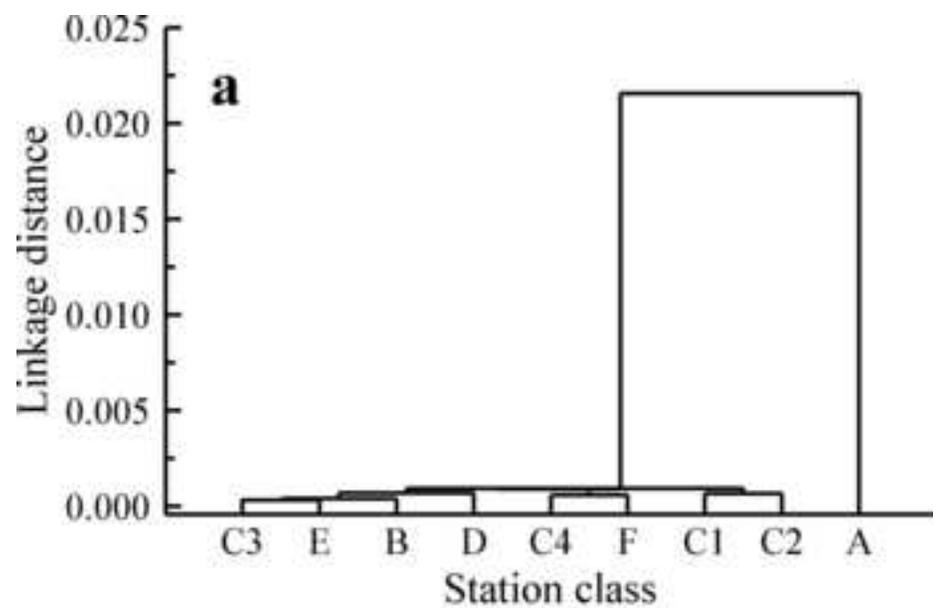


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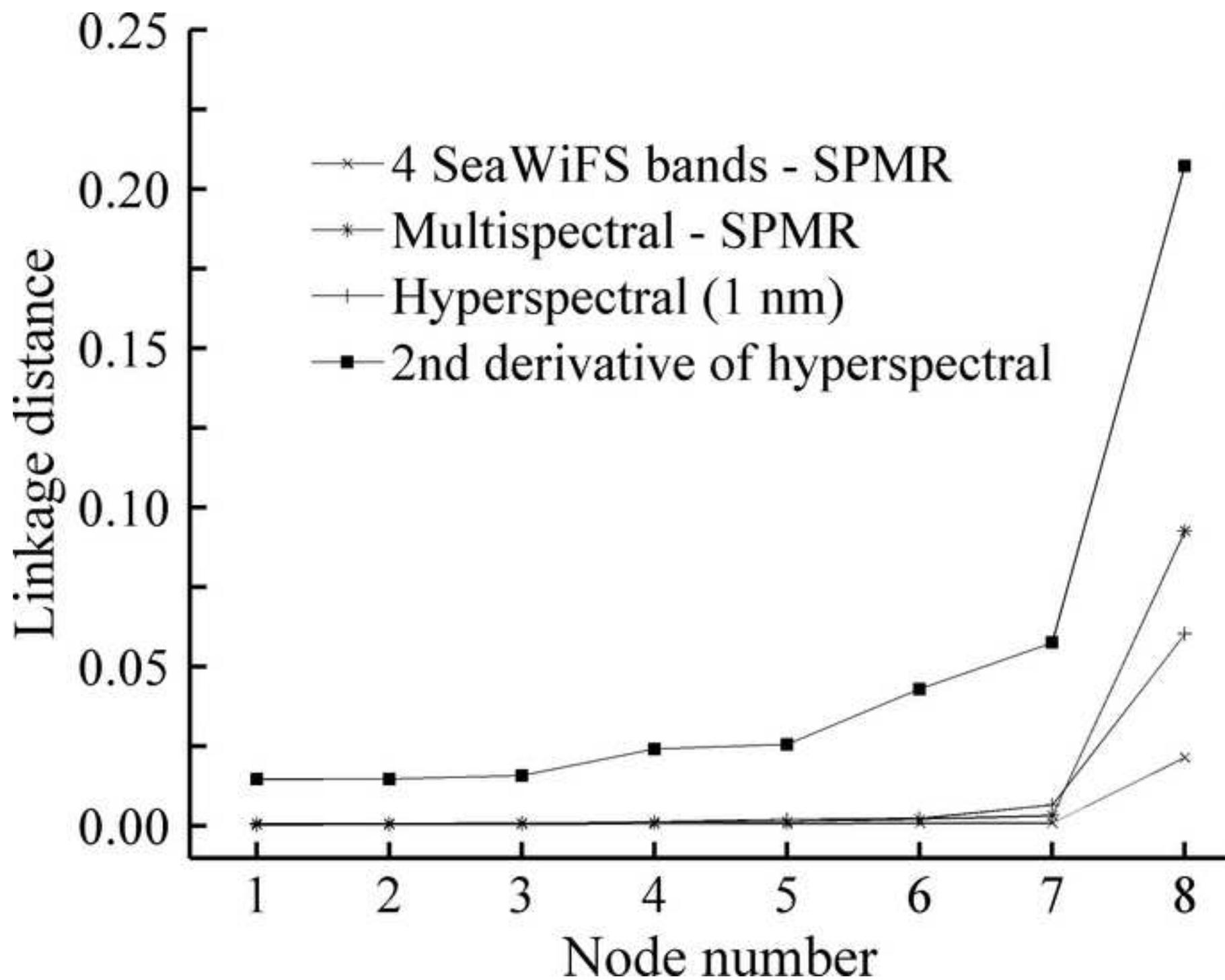


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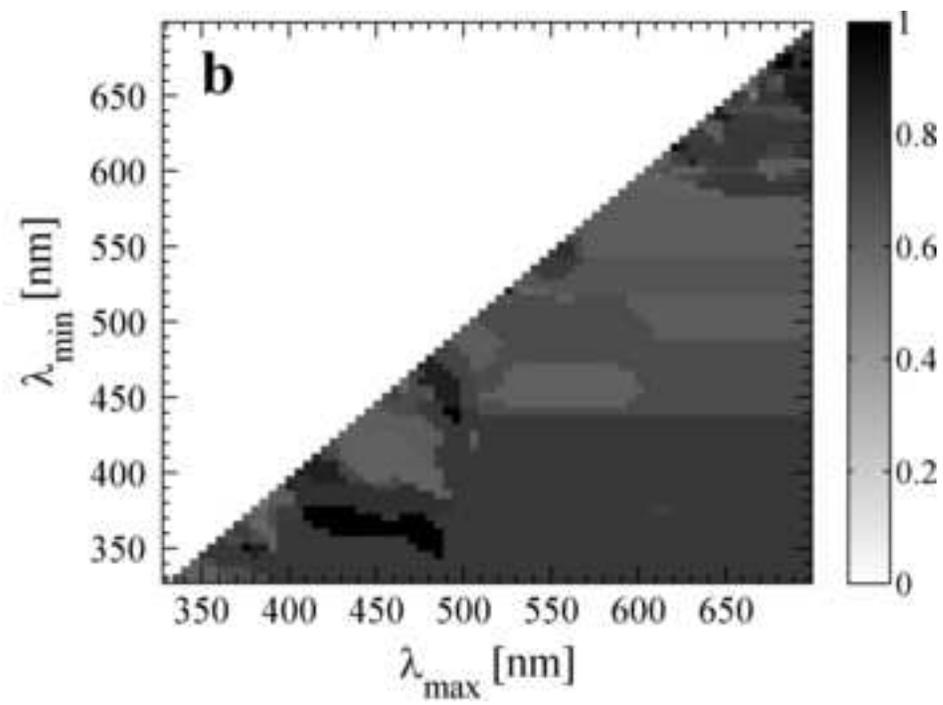
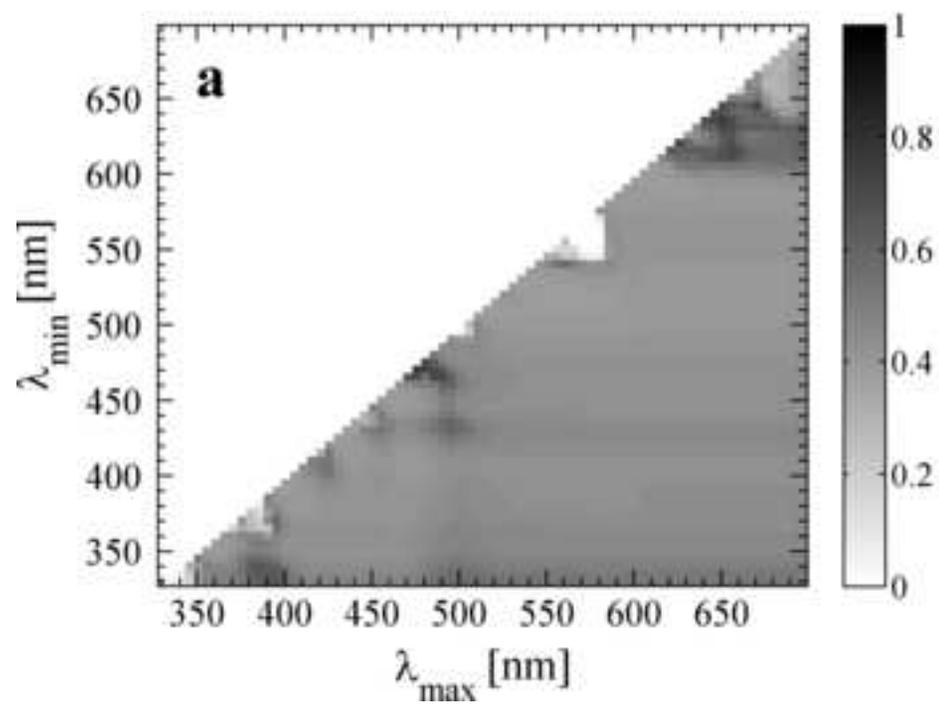


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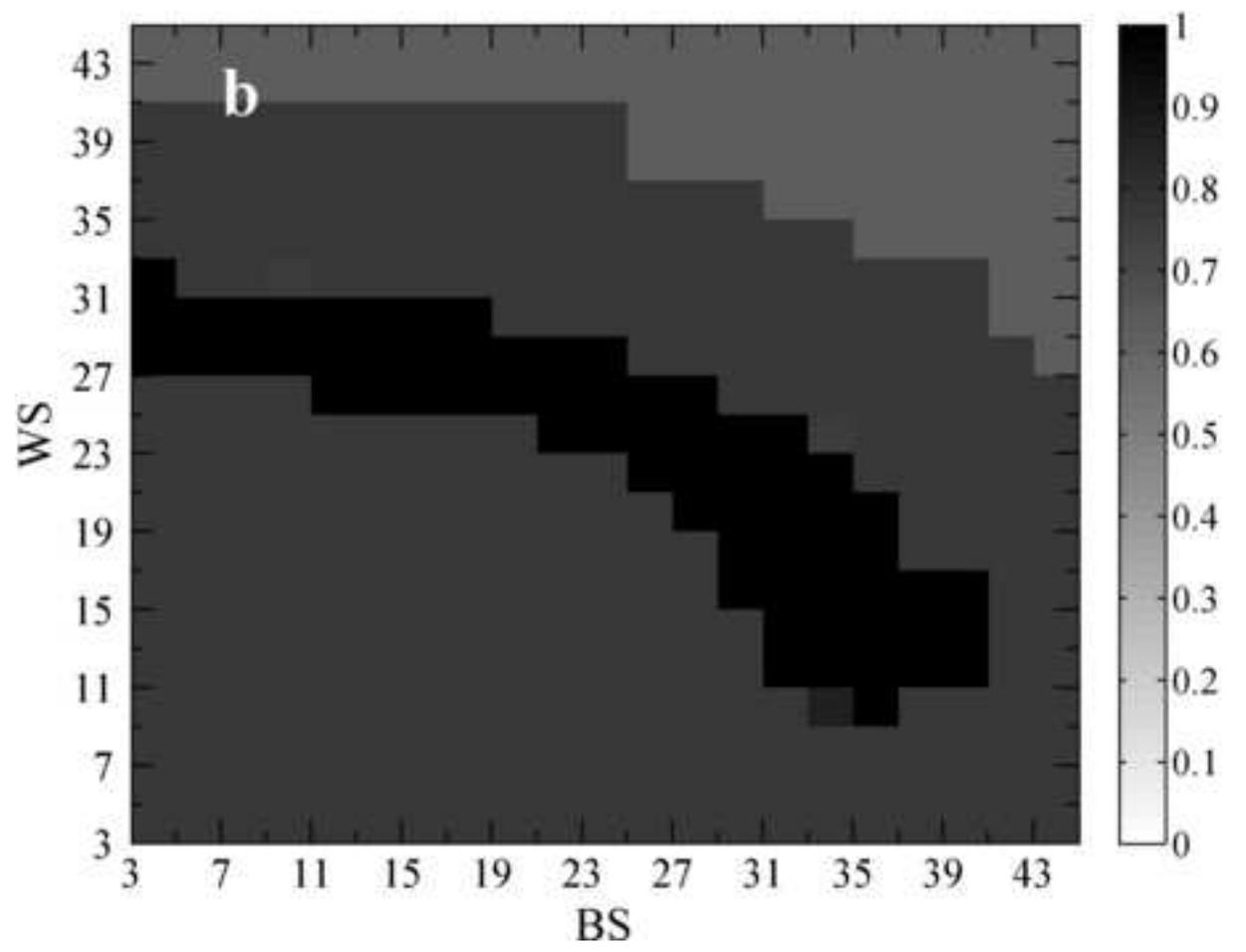
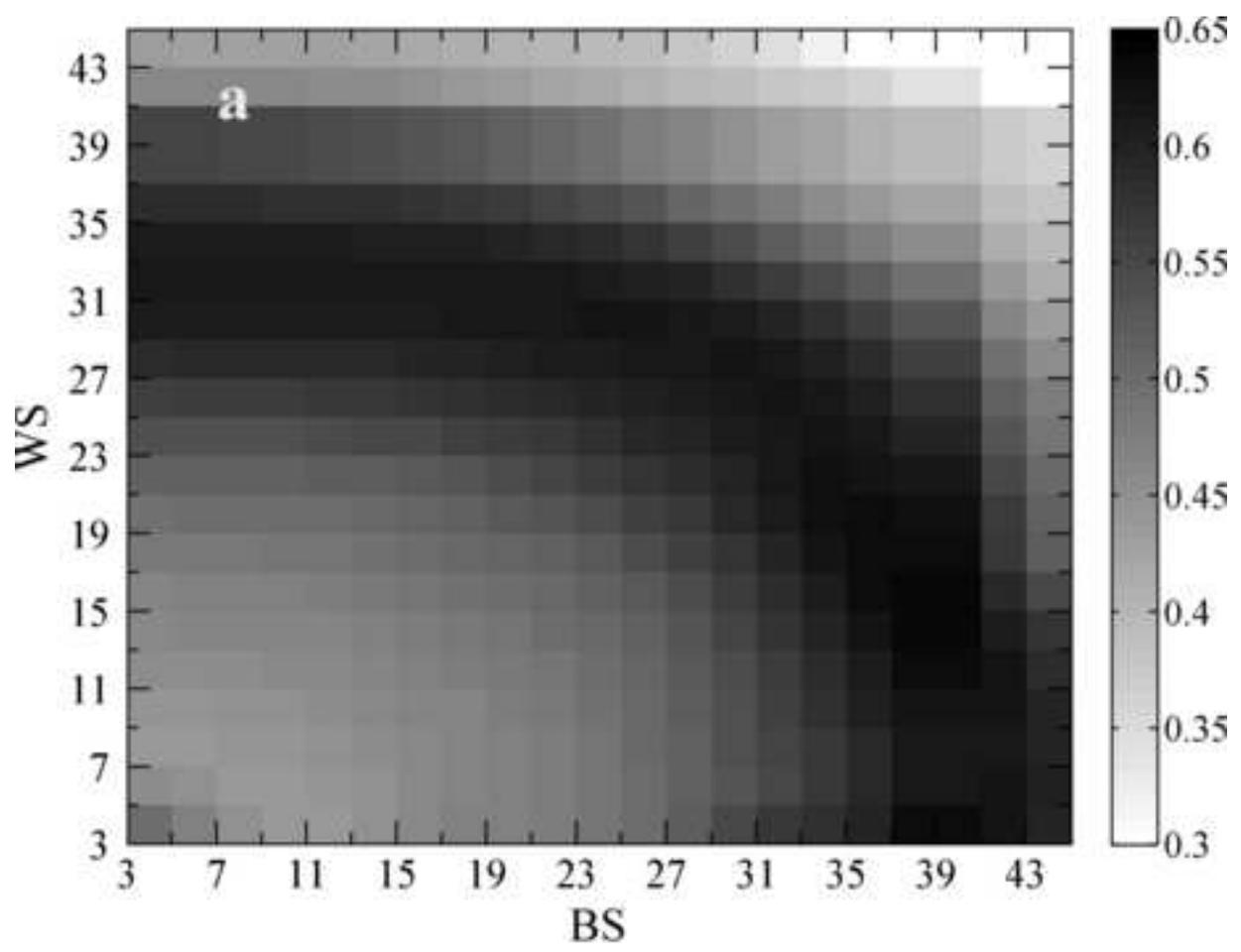


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