

1 **Rapid formation of large aggregates during the spring bloom of**
2 **Kerguelen Island: observations and model comparisons.**

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34 **Abstract**

35 While production of aggregates and their subsequent sinking is known to be one pathway for
36 the downward movement of organic matter from the euphotic zone, the rapid transition from
37 non-aggregated to aggregated particles has not been reported previously. We made one
38 vertical profile of particle size distributions (PSD, sizes ranging from 0.052 to several mm in
39 equivalent spherical diameter) at pre-bloom stage and seven vertical profiles three weeks later
40 over a 48 h period at early bloom stage using the Underwater Vision Profiler during the
41 Kerguelen Ocean and Plateau Compared Study cruise 2 (KEOPS 2, October–November
42 2011). In these natural iron-fertilized waters southeast of Kerguelen Island (Southern Ocean),
43 the total particle numerical abundance increased by more than 4 fold within this time period.
44 A massive total volume increase associated with particle size distribution changes was
45 observed over the 48 h survey, showing the rapid formation of large particles and their
46 accumulation at the base of the mixed layer. The results of a one dimensional particle
47 dynamics model support coagulation as the mechanism responsible for the rapid aggregate
48 formation and the development of the V_T subsurface maxima. The comparison of V_T profiles
49 between early bloom stage and pre-bloom stage points out an increase of particulate export
50 below 200 m when bloom has developed. These results highlight the role of coagulation to
51 form large particles and to trigger carbon export at early stage of a natural iron fertilized
52 bloom, while zooplankton grazing may dominate later in the season. The rapid changes
53 observed points out the critical need to measure carbon export flux with high sampling
54 temporal resolution. Our results are the first published in situ observations of the rapid
55 accumulation of marine aggregates and their export and the general agreement of this rapid
56 event with a model of phytoplankton growth and coagulation.

57 **1 Introduction**

58 Biological particle production and sedimentation out of the euphotic layer to underlying
59 waters is the major mechanisms for atmospheric CO₂ removal and the redistribution of carbon
60 and associated nutrients in the ocean. The fate of this exported particulate carbon is a function
61 of the plankton community producing them in the upper layer and particle transformation by
62 microbes and zooplankton during their descent to the deep sea. Physical aggregation of
63 particles is one key process in this transformation and transport and can explain the rapid
64 formation and export of large particles during bloom conditions.

65 The Southern Ocean is the largest High Nutrients Low Chlorophyll (HNLC) region of the
66 global ocean. However, several areas in this biological desert display strong seasonal
67 phytoplankton blooms. Since the HNLC regions result from low supplies of the crucial
68 nutrient iron, the hypothesis is that these blooms are supported by natural sources of iron,
69 most likely supplied from local islands and shallow sediment (Moore and Abbott, 2002;
70 Tyrrell et al., 2005; Blain et al., 2007; Pollard et al., 2007).

71 The impact of iron on the biological carbon pump has been investigated in these natural
72 bloom regions (Blain et al., 2007; Pollard et al., 2007) and in patches formed by adding iron
73 to localized HNLC regions (Boyd et al., 2000, 2004; Gervais et al., 2002; Buesseler et al.,
74 2004, 2005; de Baar et al., 2005; Hoffmann et al., 2006; Smetacek et al., 2012; Martin et al.,
75 2013). The observations made during the natural iron fertilization programs KEOPS1 and
76 CROZEX (CROZet natural iron bloom and EXport experiment) documented a two-fold
77 greater carbon export flux downward from the mixed layer (ML) in the natural iron fertilized
78 bloom relative to that in unfertilized surrounding waters (Jouandet et al., 2008, 2011; Savoye
79 et al., 2008; Pollard et al., 2009). An increase in POC flux after artificial fertilization
80 experiments was detected only during SOFex (Southern Ocean Fe Experiment, Buesseler et
81 al., 2005) and EIFEX (European Iron Fertilization Experiment, Smetacek et al., 2012).

82 Optical examination of particles trapped in polyacrylamide gels during KEOPS1 found that
83 export at 100-430 m was dominated by fecal pellets and fecal aggregates (Ebersbach and
84 Trull, 2008) which can be considered as a form of indirect export. (Note that we consider
85 direct export to be the flux of phytoplankton cells, either alone or in aggregates). By contrast,
86 the CROZEX experiment observed direct export of surface production by a diverse range of
87 diatoms (Salter et al., 2007), consistent with phytoplankton aggregation enhancing particulate

88 flux. The lack of phytoplankton aggregation due to insufficient biomass has been invoked as
89 the reason for which carbon export flux in SOIREE (Southern Ocean Iron Release
90 Experiment) were not enhanced (Waite and Nodder, 2001; Jackson et al., 2005). The different
91 results for these systems reflect differences in physical forcing factors, experimental duration,
92 and seasonal evolution of the biological community.

93 Because of the complexity of the export system, there are still extensive unknowns about the
94 effect of iron fertilization on carbon export from the surface to the bottom layer. The aim of
95 our study is to investigate processes responsible for the formation of large particles ($> 52 \mu\text{m}$)
96 at a short time scale during bloom development in the surface ML.

97 We combine multiple vertical profiles of large particles size spectra collected over a relatively
98 short period during KEOPS2 with a one dimensional particle dynamics model that
99 incorporates phytoplankton growth as a function of light and nitrate concentration and
100 coagulation as function of aggregate size. We measured particle distributions using the
101 Underwater Vision Profiler (UVP) deployed at a bloom station above the Kerguelen plateau
102 under pre-bloom conditions and at an early bloom stage during a period of rapid change. The
103 coagulation model used here is an extension of a zero-dimensional model that simulates
104 abundances of phytoplankton cells in the surface mixed layer as well as the size distributions
105 of settling particles (e.g., Jackson et al., 2005; Jackson and Kiørboe, 2008). Here it has been
106 extended into a one-dimensional model to describe the vertical distribution of phytoplankton
107 in the mixed layer and the formation, distribution, and flux of aggregates. The comparison
108 between observed and modelled particle size distribution provides a unique opportunity to test
109 the usefulness of the coagulation theory to explain rapid formation of large aggregates during
110 the early stage of a phytoplankton bloom.

111

112 **2 Material and methods**

113 **2.1 Field measurements**

114 The station A3 ($50^{\circ}380 \text{ S}$, $72^{\circ}050 \text{ E}$), located above the Kerguelen plateau, is characterized
115 by a weak current (speed $< 3 \text{ cm s}^{-1}$, Park et al., 2008b), which results in a water mass
116 residence time of several months. This long residence time allows the bloom to develop and
117 persist over an entire season in response to natural iron fertilization (Blain et al., 2007).

118 During KEOPS2, Station A3 was sampled first during pre-bloom conditions on 21 October
119 2011 (A3-1) and was revisited during the early bloom from 15 to 17 November (A3-2), 2
120 weeks after the bloom had started. High sampling frequency started during the second visit at
121 midnight on 15 November (Table 1).

122 The Underwater Vision Profiler 5 (UVP 5 Sn002) used in the present study was a component
123 of the rosette profiler system. The UVP5 detects and measures particles larger than 52 μm on
124 images acquired at high frequency (Picheral et al., 2010). Images were taken and data
125 recorded at a frequency of 6Hz, corresponding to a distance of 20 cm between images at the 1
126 m s^{-1} lowering speed of the CTD. The observed volume per image is 0.48 dm^3 ; the total
127 volume sampled for the 500 m depth profiles at Station A3 was 1.2 m^3 . The instrument takes
128 a digital picture of a calibrated volume lit from the side. The image is scanned for particles
129 and particle dimensions are measured. The pixel area (S_p) for each object is converted to
130 cross-sectional area (S_m) using the calibration equation $S_m = 0.00018 S_p^{1.452}$. An equivalent
131 spherical diameter d is calculated for that cross-sectional area. Hydrographic and
132 biogeochemical properties, including density, fluorescence, turbidity (as determined by a
133 transmissometer using a wavelength of 660nm and a 25 cm path length), were measured
134 simultaneously with a conductivity-temperature-depth system (Seabird SBE-911+CTD)
135 linked to a Seapoint Chelsea Aquatracka III (6000 m) chlorophyll fluorometer and a WET
136 Labs C-Star (6000 m) Transmissiometer.

137 We also present selected results of chlorophyll a (Chl a) and nitrate concentrations, as well as
138 relative biomass of different phytoplankton size classes. Chl a and pigment concentrations
139 were measured using high performance liquid chromatography (HPLC) following the method
140 described in Lasbleiz et al. (2014); the fraction of a phytoplankton group relative to the total
141 biomass was calculated using the model of Uitz et al. (2006). Nitrate was analysed with a
142 Technicon autoanalyzer as described in Tréguer and Le Corre (1975).

143 **2.2 Data processing**

144 The particles in each 5 m depth interval, with depth determined from the associated CTD
145 measurements, were sorted into 27 diameter intervals (from 0.052 to 27 mm, spaced
146 geometrically), and concentrations calculated for each diameter and depth interval. We further
147 analysed size spectra having a minimum of 5 particles per size bin and depth interval; this
148 criterion eliminated bins with $d > 1.6$ mm. The depth distributions of particles are

149 summarized in terms of their total number N_T ($\# L^{-1}$) and volume V_T ($mm^3 L^{-1} = ppm$)
150 concentrations.

151 Particle number distributions (n) were calculated by dividing the number of particles (ΔN)
152 within a given bin by the width of the ESD bin (Δd) and the sample volume. The resulting
153 units are $\# cm^{-4}$. The distributions are usually plotted in a loglog plot because of the large
154 ranges in n and d . To compensate for these ranges, the results are often displayed as nVd
155 spectra, where n is multiplied by the median diameter (d) and the spherical volume $V = \pi/6 d^3$
156 for the particle size range. This form of the particle size distribution has the advantage that the
157 area under the curve is proportional to the total particle volume concentration when nVd is
158 plotted against $\log(d)$. The carbon export flux F_{POC} ($mg C m^{-2} d^{-1}$) can be estimated from the
159 size spectra using the empirical relationship:

$$F_{POC} = Ad^b \quad (1)$$

160 where d is the diameter in mm, $A = 12.5$ and $b = 3.81$ (Guidi et al., 2008). Guidi et al. (2008)
161 developed the relationship by comparing particle size spectra to sediment trap collection rates
162 at locations around the world. The value of b is less than the value of 5 expected for spherical
163 particles of constant density (for which mass increases as d^3 , and sinking speed as d^2). It is
164 consistent with marine aggregates having increasing porosity with increasing size (e.g.,
165 Alldredge and Gotschalk, 1988).

166

167 **2.3 Model equations and parameterization**

168 The biological model describes the growth rate of phytoplankton in the water column as a
169 function of light and nutrient (nitrate) concentration. The model uses a maximum
170 phytoplankton specific growth rate $G_{max} = 0.45 d^{-1}$ (Timmermans et al., 2004; Assmy et al.,
171 2007). Phytoplankton cells are transformed into aggregates by differential settling and shear
172 using the standard coagulation model of Jackson (1995). Aggregates are also fragmented in
173 two similar parts using size-dependent disaggregation rates (Jackson, 1995). The primary
174 phytoplankton cells are chosen to match the size of *Fragiliaropsis kerguelensis* which was the
175 dominant species under pre-bloom conditions (Armand, Pers. Com.). A single phytoplankton
176 cell has $d_l = 20 \mu m$, a density $1.0637 g cm^{-3}$, and a resulting settling speed $v_l = 1.05 m d^{-1}$. The
177 probability that two particles colliding stick together, α , is assumed to be 1. The average

178 turbulent shear rate is $\gamma = 1 \text{ s}^{-1}$ (Jackson et al., 2005). The initial abundance of phytoplankton
179 is 10 cells cm^{-3} . These and other parameter values are shown in Table 2. The one-dimensional
180 model simulates the distribution of particles of different sizes, including solitary
181 phytoplankton cells, and nitrate concentrations at 2 m depth intervals within the 0–150 m
182 layer. This depth range corresponds to the average surface ML thickness during the survey
183 (Table 1). Neither zooplankton grazing nor particle transformation by bacterial processes is
184 included in these simulations. The model is described in greater detail in Appendix A.

185 While the concept of spherical diameter is simple for a solid sphere, it is not for irregular
186 marine aggregates, with different shapes, assembled from multiple sources, having water in
187 the interstices between their components and yielding different sizes for different
188 measurement techniques (e.g., Jackson, 1995). The simplest diameter is the conserved
189 diameter d_c , the diameter if all the solid matter was compressed into a solid sphere. It has the
190 advantage that when two particles collide and form a new particle, the conserved volume of
191 the new particle is the sum of the conserved volumes of the two colliding particles. The
192 particle diameter d determined by the UVP is larger than d_c because aggregate size is
193 determined from the outer shape of the aggregate and thus contains pore water between
194 source particles. The relationship between the two measures of particle diameter is described
195 using the fractal dimension (see Appendix A). The model calculations use d_c . However, all
196 model results shown here use the apparent diameter d_a , which is used to approximate the
197 diameter reported by the UVP. The value of d_a is calculated from d_c using the fractal
198 relationship and a fractal dimension of 2 (Appendix A). Note that reported values of the
199 fractal dimension vary widely, from 1.3–2.3 (Burd and Jackson, 2009). The value of 2 used
200 here is in this range and yields peaks in the nVd distributions similar to those determined from
201 UVP measurements, unlike values of 2.1 and 1.9 (not shown).

202

203 **3 Results**

204 **3.1 Observations**

205 **3.1.1 Biogeochemical and physical context**

206 The water column was characterized by a deep mixed layer (~150 m) during the pre-bloom
207 and early bloom surveys, with a range of 120 to 171 m (Figs. 1 and 2). Isopycnal

208 displacements of up to 50 m can be seen in the density profiles. Such vertical movements are
209 known to result from semi-diurnal internal tides in this region (Park et al., 2008a). The
210 fluorescence and Chl *a* concentrations show a 4-fold increase from A3-1 (21 October) to A3-2
211 (15–17 November), with Chl *a* concentrations at the surface increasing from 0.5 to ~2 $\mu\text{g L}^{-1}$
212 (Figs. 1 and 2). The Chl *a* profile determined using bottle samples for station A3-2 was
213 characterized by a subsurface maximum at 170 m, at the bottom of the mixed layer (Fig. 3).
214 The chlorophyll profiles determined using the in situ fluorometer were either relatively
215 constant or had maxima at 50 m or shallower (Figs. 1 and 2). Variations in the maximum
216 depth of fluorescence from the in situ profiles were associated with temporary deepening of
217 the mixed layer at 7.50 AM and 7.15 PM on 16 November and at 5.30 AM on 17 November.
218 In the surface mixed layer, the beam attenuation coefficient (turbidity) had a similar
219 distribution as fluorescence (Figs. 1 and 2). The two were, in fact, highly correlated in the
220 surface mixed layer ($r = 0.95$), which was not always the case in deeper layers. Nitrate
221 concentrations at A3-1 were 28 to 30 μM in the mixed layer, and then decreased by 4 μM at
222 A3-2 (Fig. 3a). Pigment analysis (Fig. 3) and cell counts of phytoplankton captured in nets
223 (Armand., Pers. Com.) showed that the phytoplankton community was dominated by diatoms,
224 *Fragilariopsis* at A3-1 and an assemblage of *Fragilariopsis*, *Chaetoceros* and *Pseudonitschia*
225 at A3-2. The zooplankton community was dominated by copepods with a mixture of adult
226 (50.5 %) and copepodites stage (49.5 %) at A3-2 (Carlotti et al., 2014). Zooplankton biomass
227 increased from 1.4 gC m^{-2} at A3-1 to 4.1 gC m^{-2} at A3-2 over the 0–250 m layer, and was thus
228 more than 2 fold lower than the mean biomass of 10.6 gC m^{-2} measured at A3 in summer
229 during KEOPS1 (Carlotti et al., 2008).

230 **3.1.2 Evolution of the total abundance and volume distributions in the mixed layer**

231 In the pre-bloom profile, total particle abundance (N_T) and volume (V_T) distributions at station
232 A3 were characterized by a two layer structure (Fig. 1B). The shallower layer had relatively
233 uniform N_T (V_T) values of 90 ± 5 particles L^{-1} ($0.3 \pm 0.1 \text{ mm}^3 \text{ L}^{-1}$) between 0 and 100 m; the
234 second layer, from 100 m to the base of the ML (166 m), had subsurface N_T and V_T maxima
235 of 142 particles L^{-1} and $0.45 \text{ mm}^3 \text{ L}^{-1}$. There was a two-fold increase in N_T at the first cast of
236 the early bloom (A3-2/1), with values reaching $200 \pm 7 \text{ # L}^{-1}$ in the first hundred meters and a
237 subsurface maximum of 300 # L^{-1} (Fig. 4). V_T also increased by one order of magnitude
238 reaching a value of $3 \text{ mm}^3 \text{ L}^{-1}$ at the depth of the subsurface maxima (Fig. 4). In subsequent
239 casts, there was a 40 m thick surface layer with constant N_T and V_T and a subsurface

240 maximum present at variable depths. Particularly striking was the rapid and continuous
241 increase of both N_T and V_T from A3-2/1 to A3-2/5 over a roughly 24 h time period. This was
242 more than a redistribution of aggregates, as N_T and V_T integrated over the ML increased from
243 282 to 743 # m⁻² and from 101 x 10³ to 1500 x 10³ mm³ m⁻². There was a further increase by
244 the end of the survey in the maximum V_T to mm³ L⁻¹, almost two orders of magnitude greater
245 than for the pre-bloom situation.

246

247 **3.1.3 Evolution of size distributions with depth and time during the early bloom phase**

248 The particle size distributions (PSD) calculated from the UVP observations provide additional
249 insight to the change in particle abundance during the 2 d spring observation period. In order
250 to display the vertical structure of PSD, we compare the average over the nominal euphotic
251 zone (0 to 40 m) to the average over the 40 m centred on the subsurface particle maximum.
252 Particles larger than 129 µm were more abundant in the subsurface layer (Fig. 5A). Consistent
253 with the analysis in the previous section, the smallest difference between the 2 layers occurred
254 during the pre-bloom sampling (A3-1). The maximum increases were in the 0.128 - 0.162 mm
255 and 0.204 - 0.257 mm size classes, with abundance increases of 66 # L⁻¹ and 62 # L⁻¹ for A3-
256 2/3. The increase was also substantial in the 0.4-0.5 mm size range. The cumulative volume
257 distribution in the 0-40 m euphotic zone shows that increased particle volumes resulted from
258 formation of larger particles (Fig. 5b).

259 Within the vertical particle maxima, half of V_T was in particles with $d > 0.5$ mm at the start of
260 the survey, while these larger particles provided more than 80% at the end. The largest change
261 in size spectra was in the approximately 17.5 h period between morning (A3-2/2) and middle
262 of the night (A3-2/5) of 16 November.

263 The nVd size distribution for profile A3-2/5 is shown in detail in Fig. 6. The area under the
264 curve at a constant depth is proportional to the particle volume V_T at that depth. Between the
265 surface and 60 m most particle volume is in the smallest size class with particles d ranging
266 between 200 and 500 µm. Massive changes occurred with depth with an increase of the
267 volume and d . The volumes from 60 m to 150 m are supported by larger particles ranging
268 between 0.65 mm to 1.1 mm, with a peak of 30 ppm for a d of 1 mm.

269

270 **3.1.4 Particle distributions below the ML**

271 In the first 50 m below the ML, V_T values mirrored those in the overlying waters, increasing
272 to 20 ppm by the end of the survey period (A3-2/7) (Fig. 7). V_T decreased from the base of the
273 ML to 200 m by about a factor of 20 for A3-2/6 and A3-2/7. Below 200 m, the depth limit for
274 winter mixing, there was no change in V_T during the two days survey. The average V_T was
275 0.40 ± 0.10 and $0.38 \pm 0.10 \text{ mm}^3 \text{ L}^{-1}$ at 250 and 350 m. There was an increase in V_T at about
276 475 m caused by resuspension from the bottom, as documented during KEOPS1 (Chever et
277 al., 2010; Jouandet et al., 2011). The particle number distribution (n) decreased from the base
278 of the mixed layer to 350 m in all size classes, particularly for particles larger than 500 μm ,
279 which were no longer detectable (Fig. 7B).

280

281 **3.1.5 Relationship between particle volume and fluorescence**

282 As mentioned, there was no simple correlation between V_T and fluorescence. However,
283 separating the observations by depth layers (the mixed layer, the base of the ML to 200 m and
284 deeper than 200 m) reveals a pattern (Fig. 8). In the shallowest layer, there was an increase
285 from the pre-bloom values of low fluorescence and particle volume for A3-1 (21 October) to
286 high fluorescence and low particle volume for A3-2/1 (15 November, 11.20 PM). This is
287 consistent with an increase in phytoplankton biomass but no aggregate production. For A3-
288 2/2, there are hints of an increase of V_T , which became pronounced in subsequent casts. The
289 increased particle concentrations were accompanied by a slight decrease in fluorescence. For
290 the seven casts performed during the early bloom stage, the correlations between fluorescence
291 and V_T were negative (-0.53), with a slope of $-0.015 \mu\text{g Chl mm}^{-3}$. In the second layer,
292 immediately below the surface mixed layer, fluorescence and V_T increased together, with a
293 positive correlation coefficient (0.68) and a slope of $0.036 \mu\text{g Chl mm}^{-3}$ (Fig. 8). This is
294 consistent with no phytoplankton growth in this depth layer, but with phytoplankton and
295 aggregates arriving together from above, presumably in aggregates. There was no correlation
296 between fluorescence and V_T below 200 m during this period.

297

298 **3.1.6 POC flux**

299 The flux at 200 m computed from the UVP particle size distributions increased from 1.8 mg
300 $\text{m}^{-2} \text{d}^{-1}$ during pre-bloom conditions to 23 $\text{mg C m}^{-2} \text{d}^{-1}$ during the early bloom (last cast of the
301 survey). This increase over time as estimated from UVP measurements was also evident at
302 400 m but with a smaller change, with F_{POC} increasing from 1.04 to 3.5 $\text{mg C m}^{-2} \text{d}^{-1}$ at 400 m
303 (Table 3).

304 Our POC flux estimates at 200 m for the spring bloom are in the range of the POC flux
305 estimated from the sediment trap PPS3/3 ($27 \pm 8 \text{ mg C m}^{-2} \text{d}^{-1}$) and below the estimates made
306 from the gel trap ($F_{\text{POC}} = 66 \text{ mg C m}^{-2} \text{d}^{-1}$) and from the thorium deficit ($F_{\text{POC-Th}} = 32 \text{ mg C m}^{-2} \text{d}^{-1}$) (Laurenceau et al., 2014; Planchon et al., 2014). $F_{\text{POC-Th}}$ at 100 m increased from pre-
307 bloom to early bloom but stayed unchanged at 200 m. The $F_{\text{POC-Th}}$ at 200 m was estimated at
308 A3-2/1, consistent with UVP observations that did not record any V_T increase.

310

311 **3.2 Simulations**

312 **3.2.1 Development of the phytoplankton bloom**

313 The phytoplankton in the model grew exponentially in the upper part of the water column for
314 the first eight days of the simulation, slowing down as light limitation became important (Fig.
315 9A). The specific rate of integrated population growth (0 to 150 m) was $\sim 0.06 \text{ d}^{-1}$ for this
316 initial period. The peak phytoplankton biomass was $2 \mu\text{g Chl L}^{-1}$ at about 10 m depth on day
317 13. The phytoplankton biomass decreased slightly when coagulation became an important
318 removal mechanism by day 20, with surface phytoplankton biomass of $1.7 \mu\text{g Chl L}^{-1}$, a
319 maximum concentration of $1.9 \mu\text{g Chl L}^{-1}$ at 15 m, and a minimum concentration of $0.2 \mu\text{g}$
320 Chl L^{-1} at 150 m. Surface nitrate concentrations decreased from the initial 30 to 25 μM by day
321 20 (Fig. 9B).

322

323 **3.2.2 Development of the aggregate volume**

324 Aggregates with $d_a > 100 \mu\text{m}$ appeared by day 14, when the total volume peaked at 1.3 ppm
325 at 40 m (Fig. 9C). As the phytoplankton biomass increased, the maximum total volume also
326 increased. The depth of the aggregate maximum deepened as the aggregates sank, becoming
327 17 ppm below 140 m on day 18. By day 20, the initial rapid coagulation phase ended, with the

328 maximum phytoplankton biomass decreasing slightly in the upper 50 m and the aggregates at
329 the base of the mixed layer slowly decreasing. The vertical size distribution at day 20
330 provides further details on the system (Fig. 10). The nVd_a size distribution shows the
331 distribution of particle volume, with the area under the curve being proportional to the particle
332 volume when displayed with a logarithmic d_a axis, as here (Fig. 10). Most particle volume at
333 the surface is in the smallest particles, the single phytoplankton cells. At 10 m depth,
334 aggregates appear with a maximum nVd_a value at $d_a = 200 \mu\text{m}$. With increasing depth, the
335 total volume and the diameter of the maximum nVd_a both increase. The d_a at the maximum
336 became 0.9 mm at about 70 m depth, remaining constant with increasing depth, even though
337 the total volume continued to increase with depth.

338

339 **4 Discussion**

340 **4.1 Role of coagulation in the rapid changes observed**

341 There are several striking correspondences between the observations at A3 during KEOPS2
342 and the one-dimensional coagulation model used here. First, the formation of large aggregates
343 observed over the short timescale (<2 d) was mimicked by the model. The simulation results
344 highlight the ability of coagulation to change the system state on short times that require a
345 frequent sampling regime to observe. The shapes of the nVd spectra at the base of the mixed
346 layer, centred at 0.9 mm for the model and 1 mm for the observations, with half widths of 1
347 mm for the model and 0.6 mm for A3-2/5 (Fig. 6, 10), were very similar. The transition to
348 rapid coagulation took place when relatively little of the initial nitrate had been consumed in
349 the model (4 μM), consistent with the 3.6 μM decrease observed from A3-1 to A3-2 (Fig. 3 &
350 9).

351 Coagulation theory has been used to predict the maximum phytoplankton biomass in the
352 ocean (e.g., Jackson and Kiørboe, 2008). Coagulation of phytoplankton cells is a non-linear
353 process. Rates increase dramatically as phytoplankton biomass increases, eventually
354 removing cells as fast as they divide. The volume concentration at which this occurs is the
355 critical volume concentration (Jackson, 2005):

$$V_{\text{cr}} = \pi\mu(8\alpha\gamma)^{-1} \quad (2)$$

356

357 For this calculation, we assumed an average specific growth rate for the population increase
358 rate $\mu = 0.1 \text{ d}^{-1}$, in agreement with measurements made by Closset et al. (this volume), $\alpha=1$,
359 and $\gamma=1 \text{ s}^{-1}$. Note that the average increase rate is not the same as the peak rate G_{max} . For a
360 POC: volume ratio of 0.17 g C cm^{-3} (Jackson and Kiørboe, 2008) and a carbon to chlorophyll
361 ratio of 50 g C: g Chl , this is equivalent to $1.5 \mu\text{g Chl a L}^{-1}$. This value for V_{cr} is remarkably
362 close to the maximum concentrations of $2\text{-}2.2 \mu\text{g Chl a L}^{-1}$ observed during the particle
363 formation at A3-2.

364 The rapid production of aggregates at station A3 observed in this study provides an
365 impressive example of the importance of coagulation in controlling PSD and vertical export
366 of primary production.

367 The nature of the exported material collected in a free drifting sediment gel trap at 210 m
368 supports also the importance of algal coagulation in forming the exported material
369 (Laurenceau et al., 2014). Their analysis shows that the particle flux number and volume were
370 dominated by phytoaggregates over the $0.071\text{-}0.6 \text{ mm}$ size range.

371 **4.2 Limitations of the model**

372 There are, not unexpectedly, differences between model results and observations. To start,
373 fluorescence profiles are relatively constant through the surface mixed layer in the
374 observations, but have a pronounced shallow subsurface chlorophyll maximum in the model
375 because of the higher light levels near the surface. Increased mixing in the model could
376 smooth the chlorophyll profiles, as well as the distribution of particle volume. Simulations
377 made using a much larger mixing coefficient ($1000 \text{ m}^2 \text{ d}^{-1}$) yield a smaller difference in
378 chlorophyll between the surface and 150 m, but there is still a difference of $0.8 \mu\text{g Chl L}^{-1}$
379 over the depth range (results not shown). The vertical mixing rate estimated for the iron
380 fertilization experiment EIFEX, $29 \text{ m}^2 \text{ d}^{-1}$, was actually smaller than that used in these
381 simulations, $100 \text{ m}^2 \text{ d}^{-1}$ (Smetacek et al., 2012). A previous model of phytoplankton growth in
382 the Kerguelen region discussed large scale horizontal patterns but unfortunately did not display
383 vertical distribution (Mongin et al., 2008). Whatever the reason for the relatively uniform
384 fluorescence profile, it is not simply a faster diffusive mixing rate. Those differences illustrate
385 the difficulty of building a realistic phytoplankton growth model in the region to drive the
386 coagulation model. The shallower phytoplankton distribution does affect the distribution of
387 aggregates as well.

388 In a model such as the one used in the present study, there are many parameters and modelled
389 processes that influence the final results. These include parameters such as the fractal
390 dimension, the size of the phytoplankton cells, or processes to describe diatom chains growth,
391 disaggregation rates, and grazing. While the parameters could be tuned systematically to give
392 an improved fit, what is striking is the similarity between observations and the model without
393 such a systematic fitting procedure. One important parameter that was varied during model
394 development to adjust the results was the fractal dimension. Decreasing it decreased the
395 diameter of the peak value of nVd . The value that was chosen, $D_{fr}=2$, was similar to some of
396 the estimates of fractal dimension noted above and did provide the correct nVd distribution
397 when coagulation occurred.

398 Other processes are known to affect particle concentrations and fluxes, most notably physical
399 process such as advection and biological processes such as zooplankton grazing and fecal
400 pellet production (e.g., Lampitt et al., 1993; Stemmann et al., 2000; Turner et al., 2002). The
401 importance of advection could be inferred from time series measurements of LADCP. The
402 results indicated a current below 0.1 m s^{-1} , with negligible changes over the survey in the 0-
403 200 m depth layer (Park, pers.com.). The abundance and volume of zooplankton larger than
404 0.7 mm, as well as fecal sticks/pellets and aggregates, were estimated from the identification
405 of organism in the vignettes recorded by the UVP using the Zooprocess imaging software (see
406 Picheral et al., 2010). The volume of copepods did not increase through the early bloom
407 survey, suggesting that they were not responsible for the observed rapid increase in particles.
408 Ingestion rates were also estimated from zooplankton biomass using the relationship detailed
409 in Carlotti et al. (2008) using the biomass results integrated over the 0–250 m layer. The
410 ingestion rate was 1.36 mg C d^{-1} during the early bloom cast and lower than during the
411 KEOPS1 summer cruise. In addition, fecal pellet production should have a diurnal signal
412 (Carlotti et al., 2014), which was not observed in the V_T profiles. Lastly, fast sinking fecal
413 pellets are much smaller than the aggregates observed here. For example, fecal pellets falling
414 at 100 m d^{-1} are typically $2\text{-}5 \times 10^6 \text{ } \mu\text{m}^3$, equivalent to $d = 200 \text{ } \mu\text{m}$ (Small et al., 1979),
415 compared to the mm sized aggregates dominating at A3. Thus, changes in zooplankton
416 populations can be ruled out to explain the observed V_T increase at this time, although not
417 through the entire season. Modelling the dynamics of the entire season would require
418 integrating zooplankton activity.

419

420 4.3 Comparison with other studies

421 4.3.1 KEOPS 1

422 The comparison of our results with the size spectra obtained from UVP measurements at
423 Station A3 during the early bloom (KEOPS2) and the late stage of the bloom (KEOPS 1)
424 allows us to investigate the seasonal variability of particle production in the 0-200 m layer
425 and the POC export flux (Fig. 11, Table 3). During summer (KEOPS1), the phytoplankton
426 community was also dominated by *Chaetoceros* but shifted to *Eucampia antarctica* by the end
427 of the bloom (Armand et al., 2008). Zooplankton abundance was 10-fold higher than during
428 the early bloom and the community was dominated by copepods at copepodite stage (Carlotti
429 et al., 2008). The mixed layer decreased from 150 m during early bloom to 70 m during
430 summer. During KEOPS 2, V_T increased more than 20-fold from pre-bloom conditions,
431 probably as a result of the higher diatom biomass (Armand., Pers. Com.), and coagulation as
432 described in section 4.1. The value of V_T achieved by the time of the bloom decline in
433 February (KEOPS1) was quite similar to that measured during early bloom for KEOPS2 but
434 the vertical structure was different, with two subsurface maxima during KEOPS1, the first
435 one present at the base of the ML (70 m). The larger V_T measured in January was associated
436 with an increase in the fraction of large particles (Fig. 11c).

437 Below 200 m depth, V_T was still 10 times higher during the peak bloom as compared to early
438 bloom. This resulted in 40- (at 200 m) and 10-fold (at 400 m) higher carbon export fluxes
439 during the peak bloom than the early bloom (Table 3). During the decline of the bloom, V_T
440 and POC flux were still higher than during early bloom. This is consistent with the general
441 scheme of low production - high export at the end of the bloom put forward by Wassmann
442 (1998). Our results provide insights on particle production and size distributions at different
443 stages of the seasonal bloom. The early bloom occurs before zooplankton grazing dominates.
444 This leads to a large increase in diatom abundance resulting in rapid aggregate formation and
445 export from the surface ML. Later in the season, export becomes controlled by zooplankton
446 grazing and fecal pellet production, as found from the gel trap analysis (Ebersbach and Trull,
447 2008). Despite the importance of zooplankton grazing in the late season, the presence of V_T
448 maxima at the base of the ML indicates that coagulation still occurred during summer. An
449 increase of aggregate formation through coagulation as result of high cell numbers in the ML
450 and their disappearance due to grazing between the base of the mixed layer and 200 m traps
451 could also explain the dominance of fecal aggregates in the gel traps during the summer

452 deployments. Combining KEOPS cruises to describe temporal scales of particle production
453 and export (transient versus seasonal) is useful as a first step, but our limited observations
454 highlight the need for high frequency data collection over long periods.

455 **4.3.2 Potential impact of coagulation after iron fertilization**

456 Our results can be compared to those from other iron fertilization experiments to understand
457 the relative roles of coagulation and zooplankton grazing on particle export during different
458 parts of the bloom cycle. However, it must be remembered that fertilization experiments
459 differ in important aspects, including location, physical and chemical regimes, and
460 observational techniques applied to determine stocks and fluxes. In addition, conclusions
461 about carbon export from the surface often depend on sediment traps that are usually located
462 well below the euphotic zone or surface ML, sampling events that have been filtered by
463 intervening processes and offset by transit times. With this preamble, we compare our results
464 to those from other iron fertilization studies by classifying them into those with phytodetritus
465 export driven by diatoms and the rest, including those with a zooplankton-mediated export.

466 The artificial iron fertilization experiment SOIREE (February 1999) found an increase in
467 phytoplankton biomass ($\text{Chl } a = 2 \text{ mg m}^{-3}$) as a result of the iron addition, but no rapid
468 removal of phytoplankton production. The export flux was low and driven by phyto-detrital
469 aggregates (Waite and Nodder, 2001). Jackson et al. (2005) argued that the final abundance of
470 phytoplankton cells was too low for rapid coagulation and sinking. There was a change in
471 diatom settling rate associated with a change in iron status. The persistence of the bloom after
472 iron was depleted implies that zooplankton grazing was not removing the particulate material.

473 The EIFEX (February–March 2004) environment was remarkably similar to that of KEOPS2
474 (Smetacek et al., 2012). The mixed layer was slightly shallower during EIFEX (100 m) than
475 during KEOPS2 (150 m), but still relatively deep; the phytoplankton accumulation rates were
476 also similar (0.03 to 0.11 d^{-1}). Iron fertilization stimulated a large diatom bloom that reached
477 concentrations of about $2 \text{ mg Chl } a \text{ m}^{-3}$ four weeks after the fertilization started. There was
478 little effect on vertical export during the first four weeks, but export then increased rapidly to
479 $110\text{--}140 \text{ mmol C m}^{-2} \text{ d}^{-1}$. This change was associated with mass mortality of several diatom
480 species that formed rapidly sinking, mucilaginous aggregates of entangled cells and chains
481 (Smetacek et al., 2012). This pattern of rapid formation of algal cells late in the bloom is
482 similar to what we observed.

483 CROZEX investigated the impact of high biomass ($\text{Chl } a = 2\text{mg m}^{-3}$) associated with the
484 bloom decline on carbon export during 2 legs (November 2004 and January 2005) (Venables
485 et al., 2007). Carbon export fluxes estimated from a sediment trap in the highly productive
486 naturally iron fertilized region of the sub-Antarctic waters were two to three times larger than
487 the carbon fluxes from adjacent HNLC waters (Pollard et al., 2009). Vertical flux was
488 dominated by a diverse range of diatoms, which suggests an important role for direct export,
489 such as by coagulation.

490 In contrast, the particulate flux in the SAZ-Sense in a region of elevated biomass ($\text{Chl } a = 1.9$
491 mg m^{-3}) in the Sub Antarctic Zone east of Tasmania fuelled by enhanced iron was dominated
492 by fecal aggregates (Ebersbach et al., 2011).

493 The LOHAFEX iron fertilization experiment was one of the few to use a particle measuring
494 system for the water column, also the UVP (Martin et al., 2013). A cyclonic eddy low in silica
495 on the Antarctic Polar Frontal Zone was fertilized with iron. In response, phytoplankton
496 biomass almost doubled to $1\text{-}1.5\text{mg Chl } a \text{ m}^{-3}$, but 90% of it was in flagellates less than 10
497 μm instead of diatoms. There was no observable change in concentrations of particles larger
498 than $100 \mu\text{m}$ or in vertical particle flux. There were several reasons proposed for the low
499 export, including the lack of diatoms in the low silicate water and intense particle
500 consumption, particularly at the base of the mixed layer (66 m).

501

502 **5 Conclusions**

503 It is clear that particle flux in the ocean is the result of many interacting processes, and none
504 of these has been identified dominant across systems. In the present study, we were able to
505 observe rapid aggregate formation and sedimentation of high concentrations of diatoms from
506 the euphotic zone. Our observations are consistent with results from a one-dimensional model
507 that includes only phytoplankton growth and coagulation. Our results demonstrate the utility
508 of coagulation theory in understanding vertical flux and its importance to initiate the
509 formation of large particles in the mixed layer and their subsequent transfer to depth during a
510 bloom. Nevertheless, efforts are still required to measure large aggregates distribution at a
511 high frequency to fill the temporal window between these short time events taking place
512 during the early bloom and the possibly slower dynamics of summer. In addition, more effort
513 is required to understand better vertical variations at a fine scale for all times and particularly

514 to estimate the transformative roles of microbes and zooplankton in decreasing the total
515 particle volume exported from the euphotic zone.

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683 **Tables**

684 Table 1: Date and time of the casts performed at Station A3.

685

Station	Date (dd-mm-yyyy)	Time (hh:mm)	Mixed layer depth (m)
A3-1	21-10-2011	2:20 AM	165
A3-2/1	15-11-2011	11:20 PM	143
A3-2/2	16-11-2011	7:50 AM	171
A3-2/3	"	11:30 AM	138
A3-2/4	"	7:15 PM	147
A3-2/5	17-11-2011	1:10 AM	123
A3-2/6	"	5:30 AM	163
A3-2/7	"	2:30 PM	124

686

687 Table 2: Symbols and parameter values used for the model. Conversion constants include:

688 Carbon to chlorophyll = 50 g C: g Chl *a*; carbon to nitrogen = 106 mol C:16 mol N.

689

Symbol	Quantity	Value	Units	Reference
d_c	Conserved diameter		cm	
d_a	Apparent diameter		cm	
d_l	Median algal diameter	20	μm	
D_{fr}	Fractal dimension	2	-	
G	Specific growth rate		d^{-1}	
G_{max}	Maximum specific growth rate	0.45	d^{-1}	Timmermans et al.
I	Irradiance		ly d^{-1}	
I_o	Surface irradiance		ly d^{-1}	Evans & Parslow
k	Total light attenuation= k_w+k_rP		m^{-1}	
k_r	Coefficient for light attenuation	0.03	$\text{m}^2 (\text{mmol N m}^{-3})^{-1}$	Fasham et al. 1990
k_w	Light attenuation of water	0.04	m^{-1}	Fasham et al. 1990
K_d	Half saturation constant	1	mmol N m^{-3}	Fasham et al. 2006
K_z	Eddy diffusivity	100	$\text{m}^2 \text{d}^{-1}$	Park et al. 2008a
m	Particle mass		g	
$n(d)$	Number spectrum for diameter		cm^{-4}	
$n(m)$	Number spectrum for mass m		$\text{cm}^{-3} \text{g}^{-1}$	
N	Nitrate concentration		mmol N m^{-3}	
Q_i	Particle mass in i th section		g	
r	Phytoplankton mortality rate	0.04	d^{-1}	Assmy et al. 2007
r_p	Relative light limitation		-	
r_n	Relative nitrate limitation		-	
v_i	Settling velocity for particle in i th section		m d^{-1}	
V	Particle volume			
α_l	Slope of photosyn. curve		0.04 ly^{-1}	Evans & Parslow
α	Stickiness	1	-	Jackson et al. 2005
β	Coagulation kernels			
${}^1\beta_{i,j,i}, {}^2\beta_{i,i,i}, {}^3\beta_{i,i,i}, {}^1\beta_{i,i,i}$	Sectional coefficients			
ϕ	Phytoplankton concentration		mmol N m^{-3}	
λ_i	Disaggregation coef. for i th section		d^{-1}	Jackson 1995
γ	Fluid shear	1	s^{-1}	Jackson et al. 2005
μ	Average algal growth rate		d^{-1}	

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691 Table 3: Comparison of the POC fluxes (F_{POC} in $\text{mg m}^{-2} \text{d}^{-1}$) derived from particle size
 692 distributions from the UVP, particle distributions from gel-filled sediment traps and sediment
 693 trap PPS3/3 Technicap Inc, France (Laurenceau et al., this volume) during KEOPS2 and
 694 KEOPS1 (Jouandet et al., 2011, Ebersbach et al., 2008).

		Winter KEOPS2	Spring KEOPS2	Mid summer KEOPS 1	End summer KEOPS1
F_{POC} at 200 m ($\text{mg m}^{-2} \text{d}^{-1}$)	F=Ad ^b	1.75	23.11	869	58
	Gel trap		66		
	Trap PPS3		27 ± 8		
F_{POC} at 350 m ($\text{mg m}^{-2} \text{d}^{-1}$)	F=Ad ^b	1.04	3.50	326	67

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697 **Figure captions**

698 **Figure 1:** Vertical distribution of sigma (black line), fluorescence (green line) and turbidity
699 (blue line) (A) and vertical profiles of total abundance (N_T) and total volume (V_T) at the first
700 cast of A3, during winter (A3-1, 21 of October).

701 **Figure 2:** Temporal evolution of density (A), fluorescence (B) and turbidity (C) during the
702 spring survey. The red line shows the mixed layer depth.

703 **Figure 3:** Vertical distribution of the concentration of NO_3 (A); Total Chl a ($T_{\text{chl}a}$), and $T_{\text{chl}a}$
704 associated with micro- ($T_{\text{chl}a_{\text{micro}}}$), nano- ($T_{\text{chl}a_{\text{nano}}}$), and picophytoplankton ($T_{\text{chl}a_{\text{pico}}}$) (B).
705 The filled symbols indicate pre-bloom stage; the hollow symbols indicate early bloom stage.

706 **Figure 4:** Vertical distribution of N_T and V_T for the different casts during early bloom stage.

707 **Figure 5:** Difference of the size spectra abundance between the depth of the volume maxima
708 (Z_{max}) and the euphotic layer (Z_e) (A) and cumulative volume distribution (B) in the euphotic
709 layer (dashed line) and at the depth of the V_T sub surface maxima (solid line).

710 **Figure 6:** Volume distribution size spectra along vertical axis on the 17 of November at 1:10
711 AM (A3-2/5). The white line indicates values at 150 m, the bottom of the model regime.

712 **Figure 7:** Distribution of V_T below the surface mixed layer (A). Normalized particles size
713 spectra abundance average over the 320-350 layer (dotted line) and 100-200 m layer (solid
714 line) (B).

715 **Figure 8:** Scatter plots of fluorescence and V_T for the 3 layers: surface to base of ML (A),
716 base of ML to 200 (B) and >200 m (C). Large symbols indicate the means for a profile in the
717 panel depth range.

718 **Figure 9:** Model results for vertical distribution through time of phytoplankton ($\mu\text{g Chl L}^{-1}$)
719 (A; phytoplankton concentration does not include any algae present in aggregates), nitrate
720 (μM) (B), and V_{Ta} (ppm) (C). Contour interval is $0.1 \mu\text{g Chl L}^{-1}$ (A), $0.5 \mu\text{M}$ (B), 1 ppm (C).
721 The calculation assumes that the UVP only measures aggregates larger than $100 \mu\text{m}$.

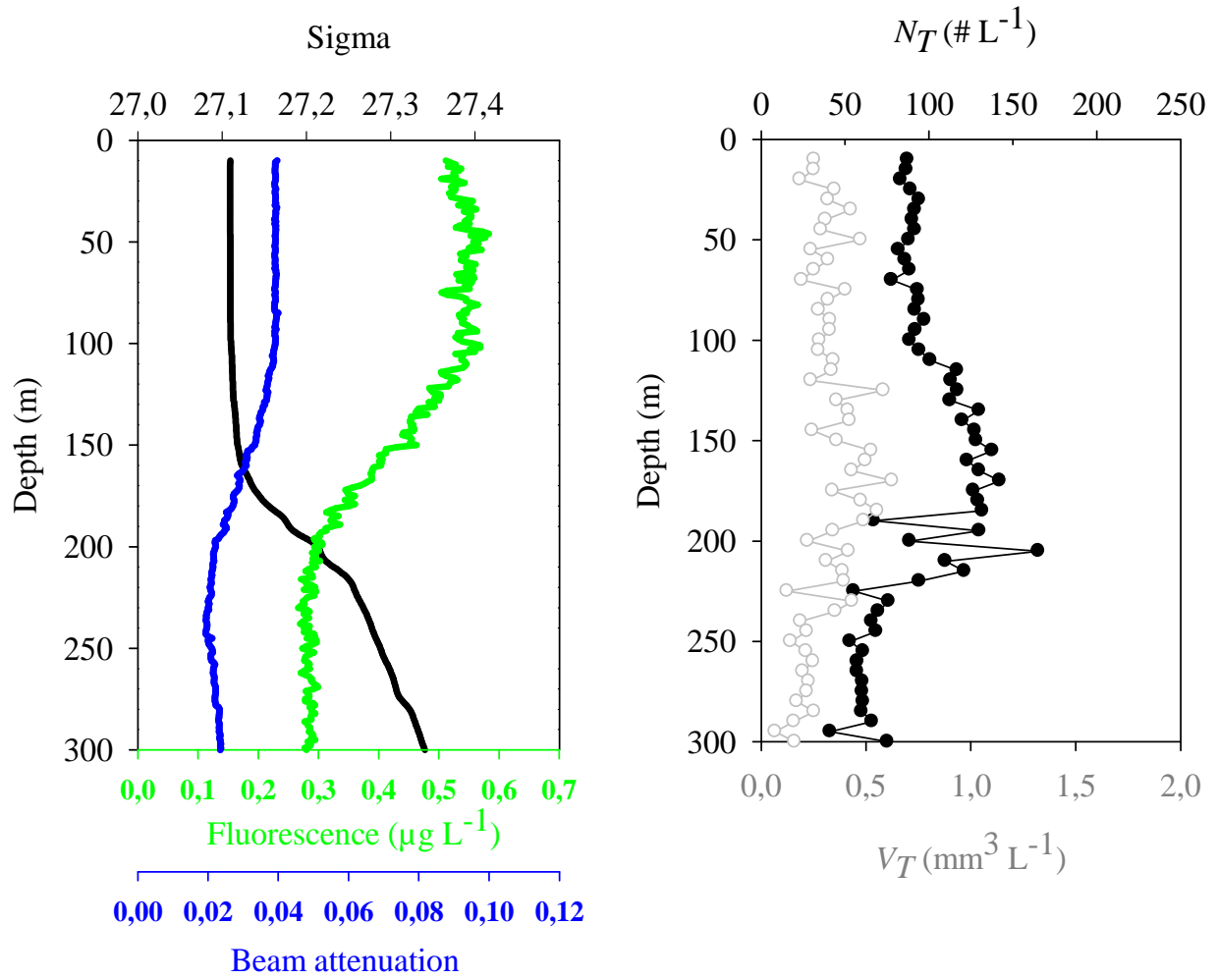
722 **Figure 10:** Distribution of apparent particle volume, nVd_a , as a function of depth and d_a as
723 calculated by the model at 20 d. Because the value of d_a is plotted on a logarithmic scale, the
724 area under the curve for each depth is proportional to total particle volume V_{Ta} .

725 **Figure 11:** A, B: comparison of the total volume profiles measured during KEOPS2 in
726 October (A3-1, blue), November (A3-2/7, green), and during KEOPS1 in January (red) and
727 February (brown). The depth scale for B is expanded to cover only 200-500m. C, D:
728 comparison of the normalised size spectra in the 0-200 m (C) and 200-400 m layer (D). The
729 colours indicate profiles as in A, B.

730 Figure 1

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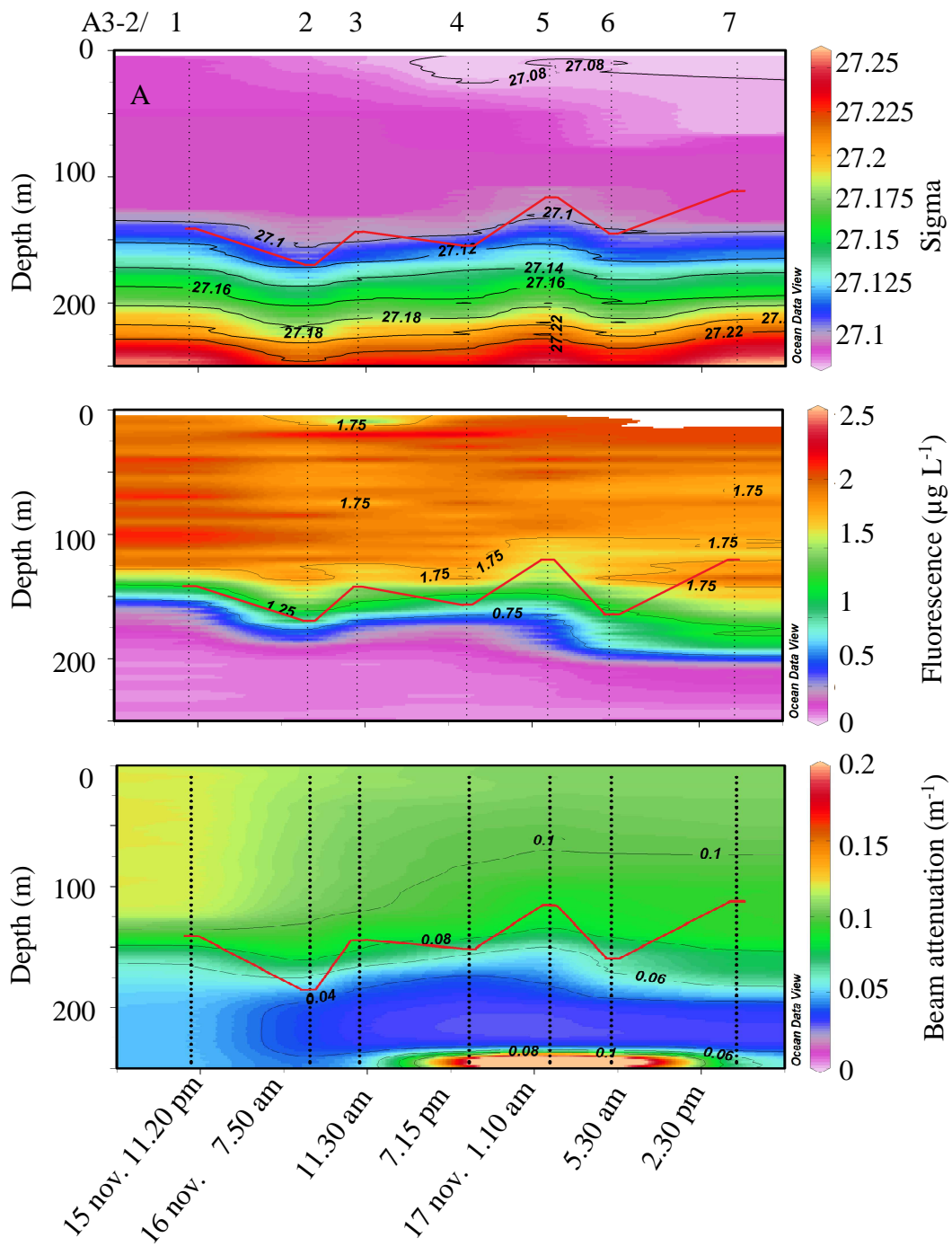
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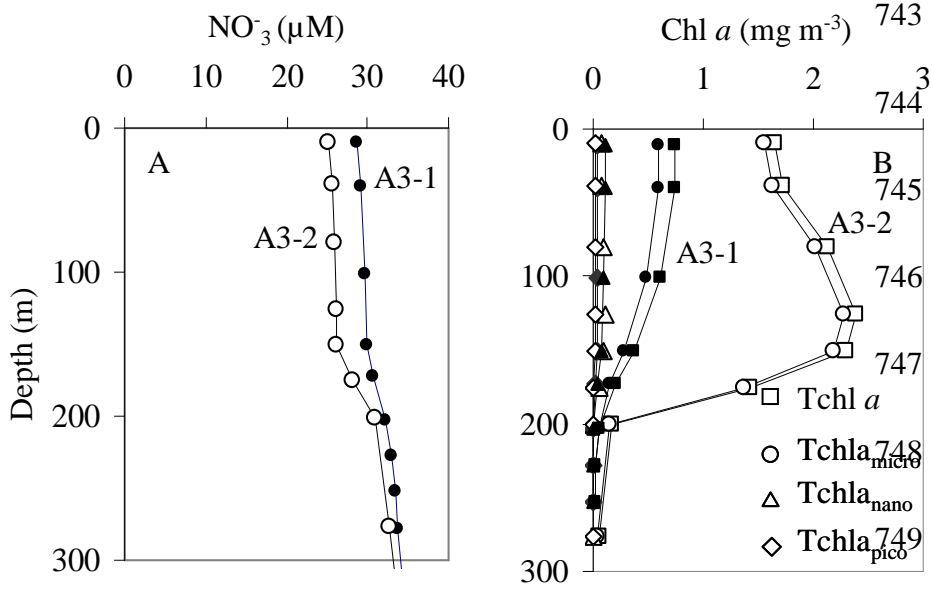
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740 Figure 2



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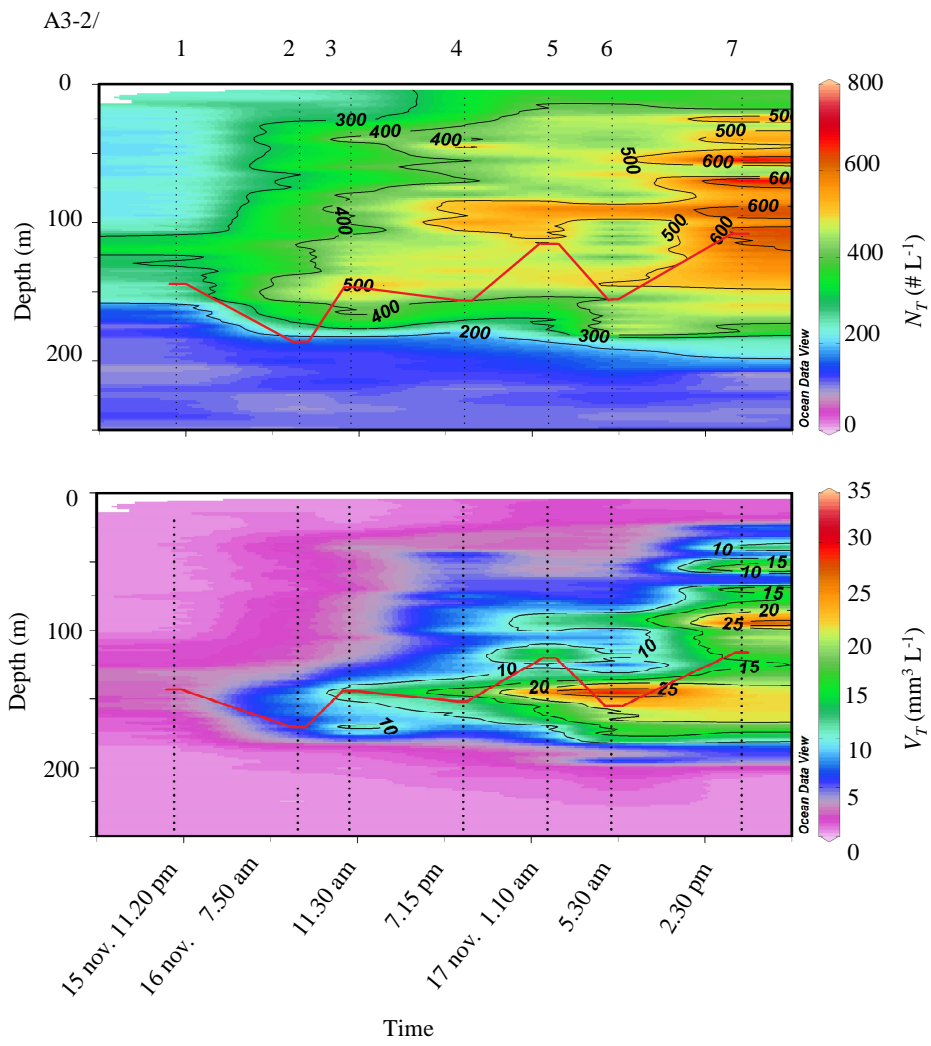


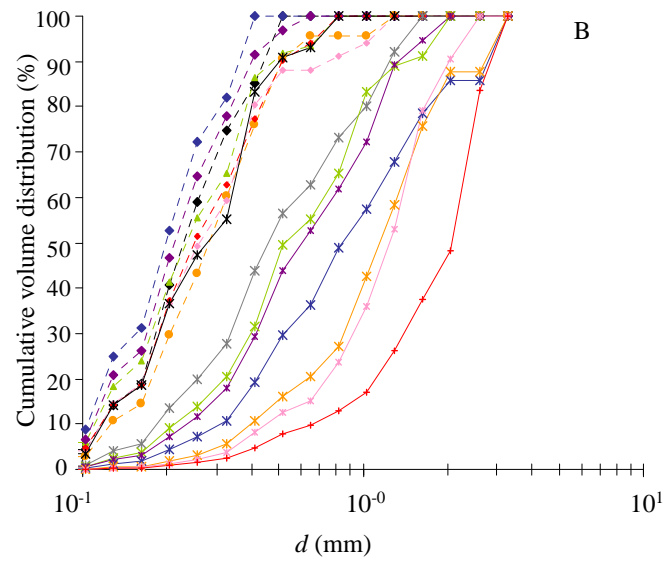
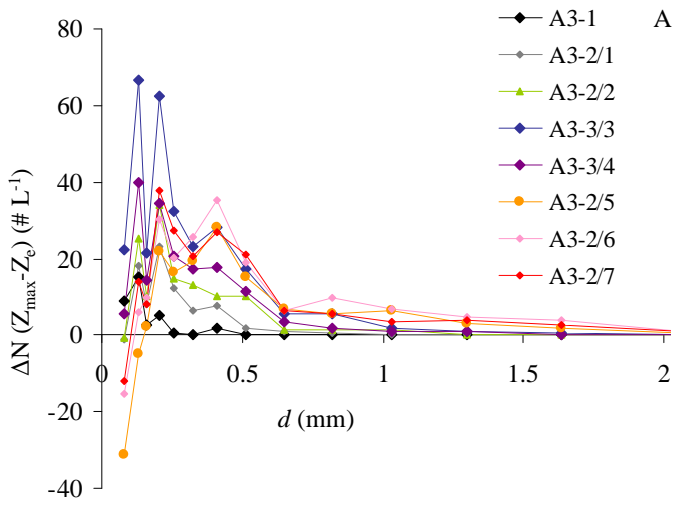
752 Figure 4

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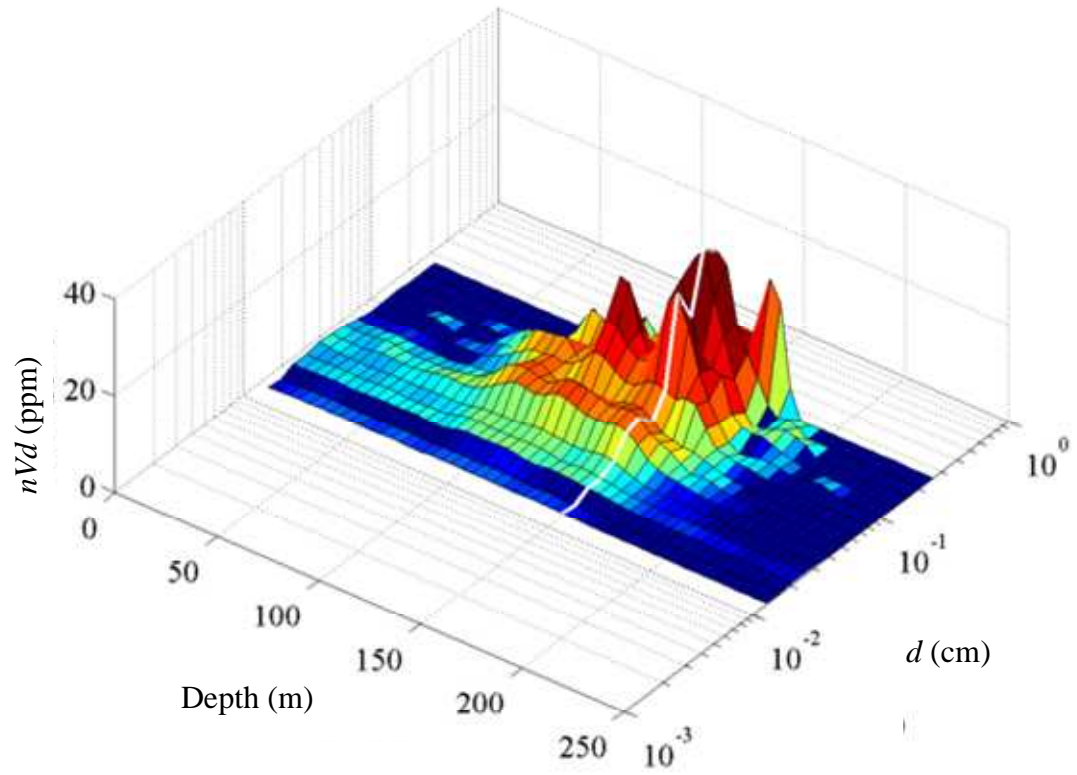
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774 | Figure 7

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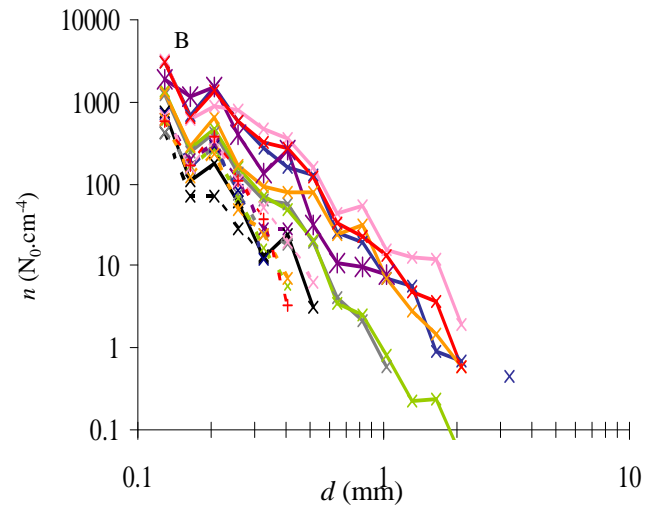
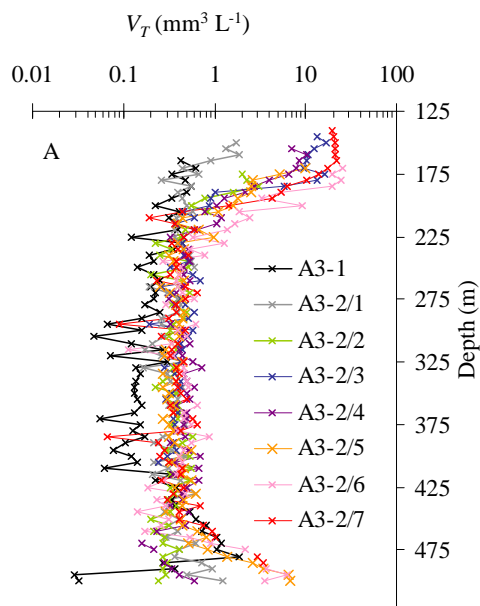
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785 Figure 8

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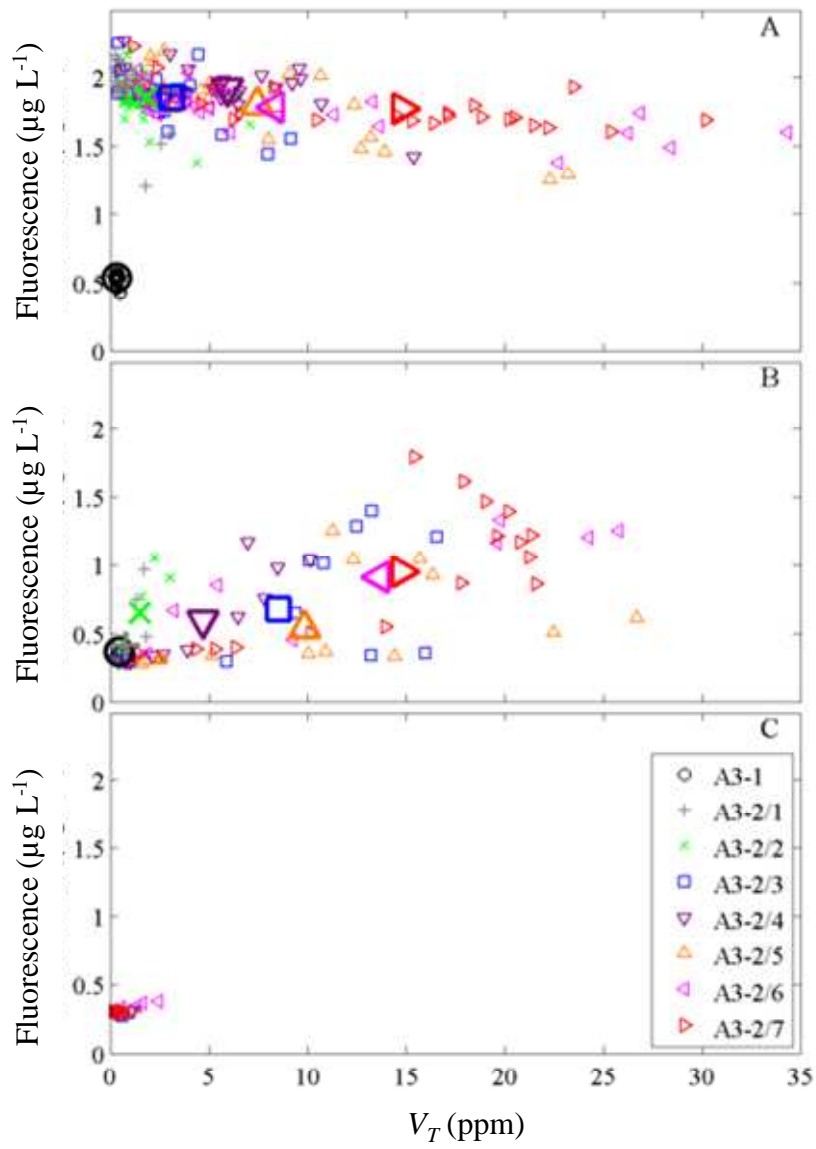
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803 Figure 9

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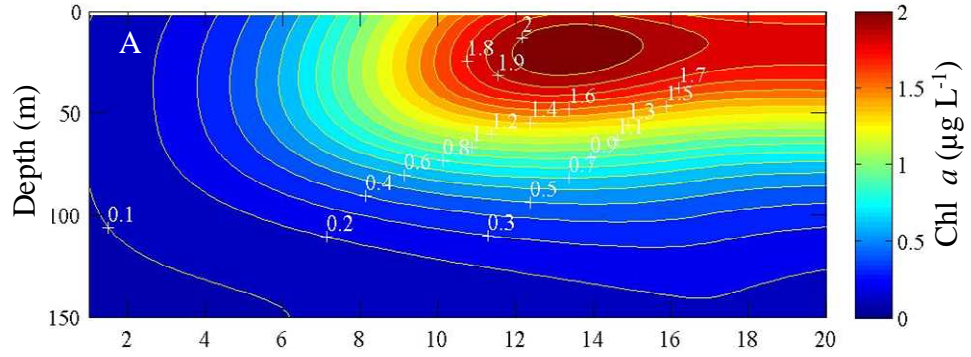
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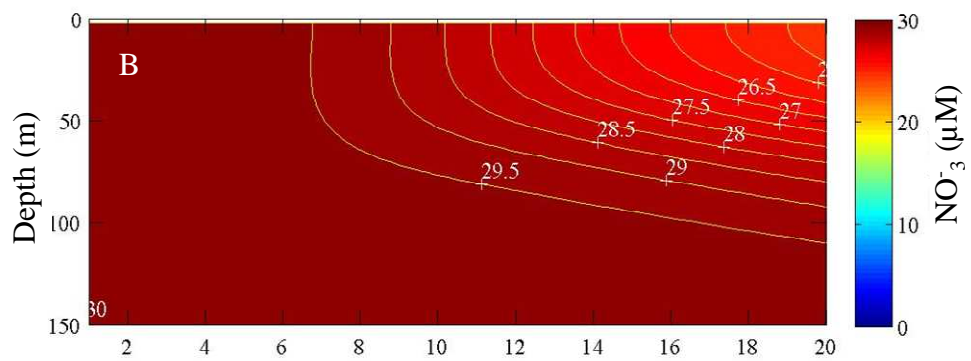
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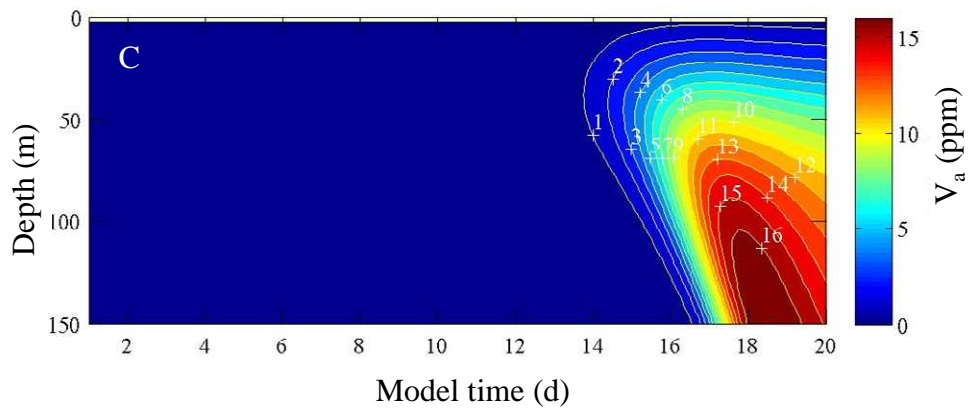
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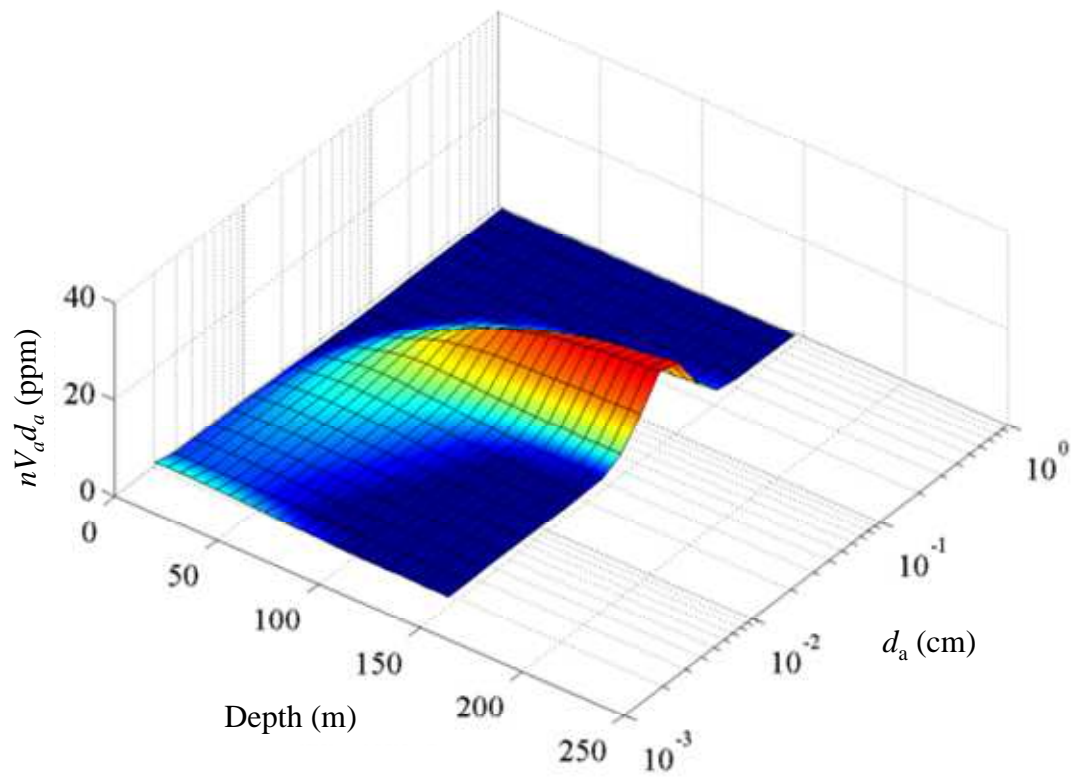
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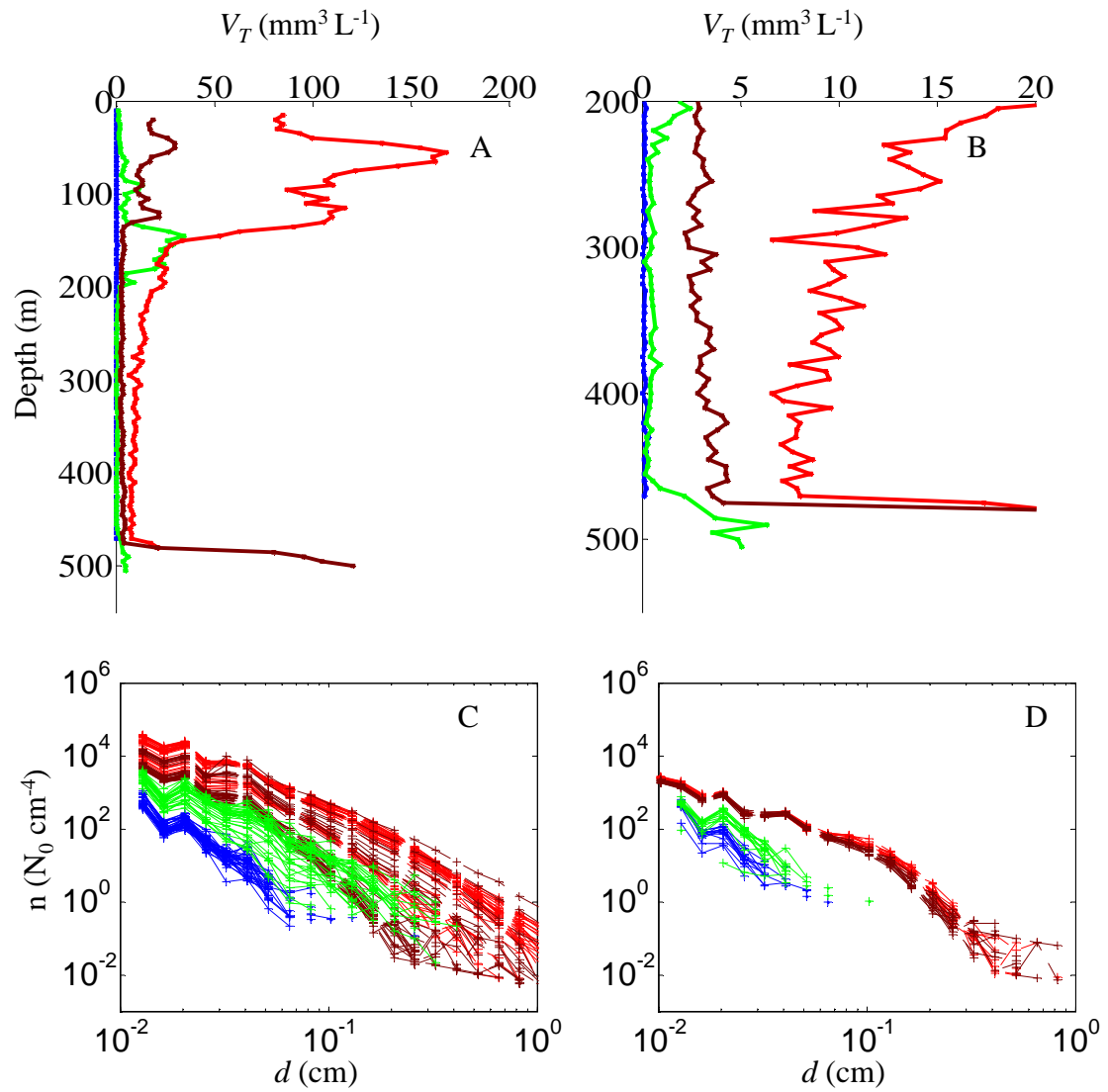


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823 Figure 11

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