

Fluorescent dissolved organic matter as a multivariate biogeochemical tracer of submarine groundwater discharge in coral reef ecosystems.

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1 **Abstract**

2 In Hawai'i and other Pacific high islands submarine groundwater discharge (SGD)
3 can be a significant and continuous source of solutes to nearshore reefs and may
4 play a key role in the structure and function of benthic coral and algal communities.
5 Identifying SGD sources and linking them to reef biogeochemistry is technically
6 challenging. Here we analyzed spectra of fluorescent dissolved organic matter
7 (fDOM) in coral reefs in the context of a suite of biogeochemical parameters along
8 gradients of SGD to characterize fDOM composition and evaluate the utility of fDOM
9 signatures in tracking groundwater dispersal and transformation. We spatially
10 mapped water column chemistry in Maunalua Bay, O'ahu, Hawai'i by collecting 24
11 water samples in grids at each of two ~0.15 km² regions during both high and low
12 tides over a two-day period. We observed clear horizontal gradients in the majority
13 of 15 measured parameters, including inorganic and organic solutes and organic
14 particles that tracked concentrations of conservative SGD tracers (radon, salinity
15 and silicate). Multivariate scanning excitation-emission fluorometry successfully
16 differentiated two distinct groundwater sources and delineated regions of SGD
17 dispersion in each reef from the surrounding water column samples without
18 detectable groundwater. Groundwater was consistently depleted in DOC and
19 enriched in nutrients; although the two SGD sources varied widely in fDOM quantity
20 and fluorophore proportions, indices of humification were consistently elevated in
21 SGD at both sites. Our results provide a robust spectral characterization of fDOM in
22 SGD-influenced coral reefs and indicate the potential for this rapid and cost-effective
23 measurement technique to be useful in tracking SGD dispersal in nearshore
24 ecosystems.

1 **Introduction**

2 Coastal ecosystems experience dynamic inputs of material from benthic, fluvial,
3 groundwater and offshore habitats. Groundwater can be a significant and
4 continuous source of solutes to nearshore reefs and may play a key role in the
5 structure and function of benthic coral and macroalgal communities, as well as
6 influencing local coastal oceanography and planktonic communities. Groundwater
7 nutrient and organic matter pollution, whether through agricultural fertilization, on-
8 site sewage disposal or runoff from industrial/urban land uses, is a major
9 eutrophication concern for coral reefs because of their adaptation to relatively low
10 nutrient conditions (Fabricius, 2005; Lapointe, 1997). However, identifying
11 groundwater sources and linking them to reef biogeochemistry is technically
12 challenging.

13
14 Coral reefs are highly productive ecosystems adapted to oligotrophic oceans, and it
15 remains an open question how they acquire sufficient macro- and micro-nutrients
16 to maintain high productivity in low-nutrient waters (e.g., Alldredge et al., 2013).
17 Submarine groundwater discharge (SGD) is a phenomenon common to the Hawaiian
18 Islands (Dollar and Atkinson, 1992; Johnson et al., 2008; Street et al., 2008;
19 Swarzenski et al., 2013) and other Pacific high islands (Kim et al., 2011) wherein
20 groundwater is continuously and directly discharged into shallow coastal reef
21 ecosystems. SGD is assumed to be a fundamental feature of reef ecosystems where
22 fluxes are significant (Cyronak et al., 2014; Paytan et al., 2006), and tracking the rate
23 and extent of groundwater dispersion in coastal regions has been an area of
24 significant active research in Hawai'i and elsewhere (Johnson et al., 2008; Knee et
25 al., 2010; Moore, 2010; Street et al., 2008). Current techniques to understand where
26 and when SGD is diffused throughout the nearshore habitat include thermal imaging
27 (Johnson et al., 2008), dye tracer studies (Burnett et al., 2006), geophysical
28 exploration (Dimova et al., 2012), radioisotopic tracers (Charette et al., 2008), and
29 mapping of conservative solute concentrations (Street et al., 2008).

30

31 The influence of SGD on the structure and function of coral reefs is poorly
32 understood. The elevated levels of nitrate and phosphate found in SGD in many
33 regions of Hawai'i (Johnson et al., 2008; Knee et al., 2010; Street et al., 2008) have
34 been implicated as a key factor in coastal eutrophication (Dailer et al., 2010),
35 changes in benthic algal composition (Smith et al., 2010; Stimson and Larned, 2000)
36 and alteration of nearshore plankton biomass and community structure (Fabricius,
37 2005; McCook, 1999; Parsons et al., 2008). Despite our conceptualization of SGD as
38 driving eutrophication, we have few studies mapping the distribution of organic
39 matter in the water column of reefs experiencing significant SGD inputs (Tedetti et
40 al., 2011). A key question for the role of SGD in coral ecosystems is how SGD may
41 influence the organic composition of coral reefs, both through allochthonous
42 subsidies and through stimulation of autochthonous productivity.

43
44 Dissolved organic matter (DOM) in aquatic ecosystems is a significant component of
45 the total organic content of marine ecosystems. The pool of DOM in the oceans is
46 vast, containing carbon equivalent to the CO₂ in the Earth's atmosphere, and
47 compositionally complex, with degradation time scales that vary greatly from hours
48 to many years (Hansell and Carlson, 2014). A portion of DOM fuels food webs
49 through metabolism by single-celled osmotrophs such as heterotrophic Bacteria and
50 Archaea that are subsequently grazed by microbial eukaryotes, a process known as
51 the "microbial loop" (Azam et al., 1983). The organic matter content of groundwater
52 can vary widely depending on the geological and hydrological factors defining
53 groundwater catchments and biogeochemical processes altering solutes within the
54 subterranean estuary (STE) (Kim et al., 2012). SGD in island systems can be sourced
55 from a variety of different ages and levels of human impact (Knee et al., 2010;
56 Wolanski et al., 2009), and little is known about the characteristics of groundwater
57 organic matter in Pacific islands (Tedetti et al., 2011). SGD passes through the STE,
58 a biogeochemical reactor that is analogous in metabolic complexity to surface
59 estuaries where terrestrial freshwater and recirculated seawater mix, differing
60 markedly with regards to sunlight exposure, residence time and redox conditions.
61 The sources of DOM in the STE can be diverse and include terrestrial inputs (Tedetti

62 et al., 2011), locally produced DOM within the STE (Santos et al., 2009), and marine
63 DOM, which enters the STE via seawater recirculating through the coastal aquifer
64 (Beck et al., 2007; Goñi and Gardner, 2003; Kim et al., 2012). Direct allochthonous
65 DOM subsidies from SGD may have varying degrees of lability relative to ambient
66 DOM depending on age and composition (Burdige et al., 2004; Kim et al., 2012); if
67 allochthonous DOM in SGD is labile, it could stimulate the microbial loop in reefs
68 thereby supporting a portion of the reef food web. Autochthonous production
69 stimulated by SGD nutrient subsidies may also produce labile DOM that supports
70 higher trophic levels (Johnson and Wiegner, 2013; Lee et al., 2010). Both subsidies
71 of DOM may have significant impacts on reef ecosystem function, and understanding
72 the relationship between SGD and DOM in reefs is an important step toward
73 understanding how SGD influences reef ecosystems and how groundwater
74 contamination may alter ecosystems processes.

75

76 The composition of DOM in aquatic environments is known to be highly complex,
77 comprising a diverse suite of thousands of molecules ranging in molecular weight
78 across many orders of magnitude (Hansell and Carlson, 2014). One method of
79 characterizing DOM is through spectral analysis of a subset of DOM that exhibits
80 autofluorescence, typically stimulated by ultraviolet and blue light (Coble, 1996).
81 This fluorescent DOM (fDOM) can exhibit variable fluorescence across a range of
82 excitation and emission wavelengths, and scanning fluorescence spectroscopy can
83 produce a three dimensional map of the fDOM in a sample that varies through space
84 and time according to subtle shifts in chemical composition of the complex
85 molecular assemblage (Nelson and Coble, 2009). Analysis of the multivariate
86 spectral characteristics of fDOM by generating an excitation-emission matrix (EEM)
87 from a sample is a cost-effective analysis that requires minimal laboratory training
88 and equipment and can produce a suite of informative data about the organic matter
89 chemistry of the water. In marine ecosystems, the variation in fDOM characteristics
90 has been used to differentiate between a variety of DOM sources including
91 terrestrial (Coble, 1996), algal (Determann et al., 1998), microbial (Stedmon and
92 Markager, 2005) and anthropogenic (Dabestani and Ivanov, 1999; Ferretto et al.,

93 2014). Additionally, fDOM characterization has proven useful in differentiating
94 contributions from rivers, groundwater, coastal margins and reefs, and the open
95 ocean (Chen et al., 2003; Helms et al., 2013; Osburn et al., 2013; Tedetti et al., 2011).
96 If fDOM exhibits clearly defined characteristics across gradients of SGD influence it
97 has the potential to serve as a promising tool for understanding the role of
98 groundwater in reef ecosystems.

99

100 The present study sought to examine the relationship between SGD inputs and the
101 field of particulate, dissolved and fluorescent organics in coral reef ecosystems. We
102 identified two regions of a single contiguous reef system with relatively predictable
103 nearshore inputs of SGD (Maunalua Bay, O'ahu, Hawai'i), which range from 12,000
104 to 16,000 m³ d⁻¹ (Holleman, 2011). SGD here is composed of brackish groundwater
105 (salinity 2-5) discharging through channelized groundwater conduits (Dimova et al.,
106 2012) thus bypassing STE processes typical for tidal flats. Since the water contains
107 minimal recirculated seawater and an extensive STE is absent, its terrestrial DOM
108 signature is preserved making it a unique SGD tracer. We first mapped the extent of
109 SGD by collecting water samples in a grid centered on an identified spring discharge
110 site at low and high tide and measuring a suite of inorganic solute concentrations,
111 including relatively conservative groundwater tracers (salinity, radon and silicate)
112 and various chemical species of N and P. From these same water samples we
113 collected a suite of dissolved and particulate organic matter samples, including bulk
114 measurements of DOM, particulate organic C and N, chlorophyll and cytometric
115 counts of picoplankton and bacterioplankton. We tested whether these parameters
116 correlated spatially and by site with the inorganic solutes measured, examining if
117 samples from different regions of the reef clustered together in patterns consistent
118 with SGD influence on the organic matter field. Finally, we conducted spectral
119 analyses on fDOM to understand the characteristics of fDOM in different
120 groundwater sources and evaluate the potential for tracking SGD dispersal and
121 alteration across the reef ecosystem. Our results demonstrate that SGD in coral reef
122 habitats alters not only the composition of inorganic solutes such as salinity, silicate
123 and nutrients, but also bulk concentrations of dissolved and particulate organics and

124 the spectral characteristics of fDOM. We discuss the potential for fDOM
125 measurements to be developed into a cost-effective tool for tracking SGD in similar
126 coral reefs dominated by spring groundwater inputs.

127 **Methods**

128

129 *Water collection*

130 We collected water samples at each of two nearshore fringing coral reef sites within
131 Maunalua Bay, O'ahu, Hawai'i (Figure 1a) over a two day period; 24 samples were
132 collected near Wailupe Beach Park on 28 May 2014 and 24 samples were collected
133 at Black Point on 29 May 2014 (Figure 1b,c). The majority of samples (32 of 48)
134 were collected synoptically during morning low tides (-6 to -9 cm) with an
135 additional subset (16) collected during afternoon high tides (+67 cm; Fig. S1).
136 Depths at these sites are generally < 2 m, and the majority of samples (38) were
137 collected from the surface (0.2m below sea level); a subset of samples were
138 additionally collected from bottom waters (roughly 0.2 m above the benthos). Using
139 kayaks, surface water samples were hand-collected into 5 L high-density
140 polyethylene carboys (73062, US Plastics, Lima, OH, USA); bottom water samples
141 were collected in 5 L horizontal Niskin samplers (101005H, General Oceanics,
142 Miami, FL, USA) and immediately siphoned to carboys for processing. Carboys and
143 Niskin samplers were previously conditioned with seawater, soaked overnight in
144 10% HCl, thoroughly rinsed with low-organic deionized water (DIW; Barnstead
145 Nanopure Diamond, Thermo Fisher Scientific, Asheville, NC, USA) and stored dry
146 before sampling. All sample containers were also triple-rinsed with sample water
147 before filling. All filtration and subsampling of water was done on site within 2 h of
148 collection.

149

150 *Sample collection and storage*

151 All storage vials were acid soaked, thoroughly rinsed with DIW, air dried, and then
152 triple rinsed with sample water before collection. Samples for total alkalinity (TA)
153 were collected first directly from the carboy spigot. Duplicate TA samples (125 mL
154 each) were transferred into polypropylene sample bottles (Huang et al., 2012) and

155 amended with 50 μL of half-saturated HgCl_2 . All subsequent samples were
156 transferred to long-term storage vials via gentle (1 mL s^{-1}) peristaltic pumping
157 directly from the carboy through platinum-cured silicone tubing over a period of
158 roughly 1 h. For dissolved nutrient analyses, the filtrate from a $0.2 \mu\text{m}$
159 polyethersulfone filter (Sterivex, Millipore, Billerica, MA, USA) was collected in acid-
160 washed and sample rinsed 50 mL polypropylene tubes and frozen to $-20 \text{ }^\circ\text{C}$.
161 Samples for fDOM analysis were collected from the $0.2\mu\text{m}$ polyethersulfone filtrate
162 (after a minimum of 250 mL sample flushing to avoid filter DOM leaching) in amber
163 glass vials with teflon-lined septate caps (acid-washed and DIW rinsed) and stored
164 in a dark refrigerator free of volatile organics. For dissolved organic carbon (DOC)
165 analyses the filtrate from glass fiber filters (Whatman GF/F, GE Life Sciences,
166 Pittsburgh, PA, USA) was collected in glass vials with teflon-lined septate caps (acid-
167 washed and DIW rinsed) and frozen to $-20 \text{ }^\circ\text{C}$ in an organic-free freezer. All glass
168 vials and filters were pre-combusted within days of sampling (2 h at $400 \text{ }^\circ\text{C}$) and
169 stored in a laboratory free of volatile organics. For analysis of chlorophyll *a* and
170 particulate organic carbon/nitrogen, 600 to 1000 mL of sample water for each
171 sample was filtered onto a 25 mm GF/F filter, folded in half, wrapped in Al foil and
172 frozen to $-20 \text{ }^\circ\text{C}$. For flow cytometry (FCM), 1.5 mL of unfiltered water was fixed at
173 0.5% paraformaldehyde (amended with 100 μL 8% ampulated paraformaldehyde,
174 Electron Microscopy Sciences, Hatfield, PA, USA) in a 2 mL polypropylene cryovial,
175 mixed briefly and then frozen to $-80 \text{ }^\circ\text{C}$. All samples were immediately refrigerated
176 in the field and frozen or refrigerated within 6 hours of collection for long-term
177 storage.

178

179 *Inorganic nutrient and organic matter concentration measurements*

180 Nutrient samples were thawed to room temperature, mixed thoroughly and
181 analyzed on a Seal Analytical Segmented Flow Injection AutoAnalyzer AA3HR for
182 simultaneous determination of soluble reactive phosphate (PO_4^{3-}), ammonium
183 (NH_4^+), nitrate+nitrite (N+N; $\text{NO}_3^- + \text{NO}_2^-$), silicate (SiO_4) and total dissolved
184 nitrogen and phosphorus (TDN, TDP; via in-line persulfate/UV oxidation). DOC and

185 TDN samples (GF/F filtrate frozen in glass vials) were measured as non-purgeable
186 organic carbon and nitrogen via acidification, sparging and high-temperature
187 catalytic oxidation on a Shimadzu TOC-L with TMN-L attachment, ensuring that
188 Deep Seawater Reference waters from the University of Miami Consensus Reference
189 Materials Project measured within specifications in each run to facilitate
190 comparison of results to those obtained by the international DOM community.
191 Chlorophyll *a* (Chl_a) concentrations were measured by acetone extraction and
192 fluorescence spectroscopy on a modified Turner 10-AU fluorometer following
193 Welschmeyer (1994). Particulate organic carbon (POC) and nitrogen (PON)
194 concentrations were determined via filter combustion on an Exeter Analytical CE
195 440 Elemental Analyzer after acid fumigation to remove particulate inorganic
196 carbon, drying, weighing and packing into tin capsules. TA samples were analyzed
197 using open cell potentiometric titrations on a 166 Mettler T50 autotitrator and
198 calibrated against a Certified Reference Material (Dickson et al., 2007). Salinity was
199 measured as electrical conductivity with a combination platinum ring electrode –
200 thermistor (Metrohm 6.0451.100) on a Metrohm conductivity module using Tiamo
201 software (v2.4). All solute and particulate samples were analyzed in the SOEST
202 Analytical Laboratory (<http://www.soest.hawaii.edu/S-LAB/>).

203

204 To assess field and technical replicability, a separate nutrient sample was collected
205 *in situ* at a subset of 15 locations and filtered immediately through a
206 polyethersulfone 0.45 µm groundwater cartridge filter (AquaPrep 600, Pall Life
207 Sciences, Ann Arbor, MI, USA). Samples were stored refrigerated in 250 mL
208 polyethylene bottles for 1 month and analyzed for salinity and by flow injection
209 autoanalyzer in parallel with the primary samples. The replicated samples were
210 representative of the range of biogeochemical zones, spanning 75% of the
211 lognormal data range for each parameter. Linear models of log₁₀-transformed data
212 collected by the two methods demonstrated strong congruency. The two methods
213 were highly correlated ($r > 0.98$ for TDN, TDP, PO₄³⁻, N+N, SiO₄, Salinity and $r = 0.85$
214 for NH₄⁺). In addition, both least-squares and orthogonal (Model II or reduced major

215 axis) regression model slopes for all 7 parameters ($p < 0.001$) were not significantly
216 different from 1 (95% confidence intervals bracketed 1). Intercepts were nearly all
217 non-significant (intercept $p > 0.25$, except TDP $p = 0.0027$), indicating no offset
218 between the two sample sets; when intercepts were constrained to zero, slopes
219 remained not significantly different from 1. This analysis indicates that
220 measurements of standard solutes were robust to minor variation in sample
221 collection and storage as well as field variation in sampling; the primary samples
222 were used in all subsequent analyses.

223

224 Dissolved organic N and P (DON, DOP) were calculated as the difference between
225 TDN, TDP and inorganic species of N and P: $DOP = TDP - PO_4^{3-}$ and $DON = TDN -$
226 $N+N - NH_4^+$. TDN was measured using two separate methods: via high temperature
227 catalytic oxidation (HTCO) and subsequent ozonation chemiluminescence of 0.7 μm
228 filtrate (GF/F) on a Shimadzu TMN-L and via persulfate alkaline oxidation and
229 subsequent colorimetric cadmium reduction of 0.2 μm filtrate (Sterivex) on a Seal
230 Analytical AA3HR (described above). The two methods yielded highly correlated
231 measures of TDN ($r = 0.97$, $n = 46$) but the HTCO method yielded consistently lower
232 estimates of TDN (Model II lognormal regression slope of 0.85 with 95 % confidence
233 interval of 0.79 to 0.91). Estimates of DON derived from the two TDN measurements
234 did not covary and were not significantly related to N+N or DOC ($p > 0.05$). When
235 N+N concentrations exceeded 40 $\mu mol L^{-1}$ estimates of DON from HTCO were
236 negative; for all subsequent analyses, Seal autoanalyzer TDN measurements were
237 used to maintain methodological consistency with other N and P measures.

238

239 *Radon activity measurements*

240 Coastal radon activities were measured using a RAD AQUA closed air loop
241 continuous equilibrium exchanger accessory for the RAD7 radon detector
242 (DurrIDGE Inc., Billerica, MA). The system was mounted on a small boat hand-pulled
243 along the shoreline and in perpendicular transects. The system was mounted on a
244 small boat and hand-pulled along pre-determined GPS transects. Coastal springs and

245 diffuse seepage was identified by moving along the shoreline and shore-
246 perpendicular transects were used to determine the extent of significant
247 groundwater plumes at the two focus areas. The air-water exchanger of the RAD-
248 AQUA was fed by water using a submersible bilge pump submersed 0.2 m below the
249 water surface. The instrument recorded radon in 5-minute integrated intervals
250 providing a spatial resolution of 50-100 meters. Radon in air values were converted
251 to radon in water activities using temperature and salinity recorded by a YSI (V2-2)
252 multiparameter probe (Schubert et al., 2012). It has been shown previously that the
253 nearshore water residence time at the sampling sites is one tidal cycle so the
254 reported radon values are not corrected for radon decay and evasion to the
255 atmosphere (Holleman, 2011). The radon survey covered the whole bay area but
256 only results relevant to Black Point and Wailupe are included in this analysis.

257

258 *Flow cytometry*

259 Flow cytometry was used to measure both autofluorescent and total nucleic-acid
260 stained cell concentrations in fixed unfiltered water samples (Nelson et al., 2011).
261 Samples were thawed and placed in 250 μL aliquots in 96-well autosampler plates
262 in duplicate; one of the two wells was mixed with SYBR Green I stain (1X final
263 concentration) within 2.5 h of analysis. Samples were analyzed on an Attune
264 Acoustic Focusing Cytometer with Autosampler Attachment (Life Technologies,
265 Eugene, OR, USA). Samples were run at flow speeds of 100 $\mu\text{L min}^{-1}$ on standard
266 sensitivity; 150 μL of sample was aspirated, 75 μL was counted and data was
267 collected only from the last 50 μL (event rates were empirically determined to be
268 steady only after 25 μL of continuous sample injection). For SYBR-stained cells a
269 blue laser (488 nm, threshold 10,000 rfu, voltage 2300mV) was used to excite the
270 dye and cell counts obtained by increasing the voltage to maintain event counts of
271 blank controls (SYBR-stained 0.2 μm filtered DIW) below 100 events s^{-1} and event
272 counts of environmental samples below 1500 events s^{-1} . This allowed for clear
273 gating of plankton cells as populations distinct from instrument noise in bivariate
274 plots of sidescatter and green fluorescence (530/30 nm bandpass fluorescence; BL1

275 channel). For autofluorescent cells a combination threshold on Violet (405 nm) OR
276 Blue (488 nm) laser excitation and red emission (600 nm and 640 nm longpass
277 filters, respectively VL3 and BL3 channels, 1000 rfu, 2500 mV) was used, and size-
278 based sidescatter gating was applied to differentiate autofluorescent photosynthetic
279 bacterioplankton (PBact) from photosynthetic autofluorescent picoeukaryotes
280 (PEuks). Concentrations were corrected for stain and paraformaldehyde dilution
281 factors, and heterotrophic bacterioplankton counts (HBact) were calculated as the
282 difference of SYBR and total autofluorescent counts. These settings were empirically
283 tested for streamwater, coastal and open ocean heterotrophic and autofluorescent
284 bacterioplankton from the North Pacific Subtropical Gyre down to depths of 4000 m
285 with densities ranging from 100 to 2000 cells μL^{-1} for SYBR-stained cells and 1 to
286 500 cells μL^{-1} for autofluorescent cells; counts matched those derived from
287 epifluorescent microscopy within 10% in all cases.

288

289 *Fluorescent dissolved organic matter (fDOM) measurement*

290 Analysis of fDOM was conducted on a Horiba Aqualog scanning fluorometer with
291 150W Xe excitation lamp, Peltier-cooled CCD emission detector and simultaneous
292 absorbance spectrometer. Samples were warmed to room temperature (22 °C) for 2
293 hours while the Xe bulb warmed. Excitation-emission matrices (EEMs) were
294 measured from each of 48 samples in a 1 cm DIW-leached and rinsed quartz cuvette
295 (3-Q-10, Starna Cells, Atascadero, CA, USA) with 4 DIW blanks run at the start and
296 end of the contiguous 3 h analysis period. Water was excited through a 5nm
297 bandpass subtractive double monochromator in declining 5nm sequence intervals
298 from 500 to 240 nm and emission was integrated 4s at each step and binned in 4.65
299 nm intervals (8-pixel bins) from 250 to 800 nm. Scans were processed using custom
300 scripts in Matlab (v2007b) as follows: 1) first inner filter effect correction was
301 applied to account for the quenching of fluorescence by absorbance following the
302 recommendations of Kothawala, et al. (2013) by multiplying by the antilog of the
303 average of absorbances at the wavelengths of excitation and emission for each
304 fluorescence data point, 2) next EEMs were scaled to Raman units (RU) by dividing

305 by the integrated emission range of 381 to 426 nm at an excitation of 350 nm in
306 averaged DIW blanks (Lawaetz and Stedmon, 2009; Murphy et al., 2010) and 3)
307 average DIW blank EEMs were subtracted from each sample.

308

309 *fDOM modelling and indices*

310 We used Parallel Factor Analysis (PARAFAC) to derive four modelled fDOM
311 components with the DOMFluor toolbox (v1.7; Stedmon and Bro, 2008), trimming
312 Rayleigh and Raman scatter, testing for outliers (none were identified), deriving up
313 to 6 PARAFAC components then using split-half validation and random initialization
314 to determine the appropriate number of modeled components (in this case only the
315 first 4 could be validated). We also calculated a suite of derived indices from each
316 EEM that are commonly used to differentiate aspects of fDOM character and help
317 interpret DOM sources. The ratio of marine-derived to terrigenous fDOM (e.g. M:C)
318 was calculated as the ratio of fluorescence at Ex310/Em410 divided by fluorescence
319 at Ex345/Em445 (Burdige et al., 2004). The M:C has had utility in differentiating
320 between marine- and terrestrial-derived fDOM (Burdige et al., 2004; Helms et al.,
321 2013). The Fluorescence index (FI; McKnight et al., 2001), calculated as the ratio of
322 fluorescence at 470 nm to 520 nm under 370 nm excitation (Cory et al., 2010; Maie
323 et al., 2006), expresses the ratio of terrigenous vs. autochthonous-produced humic
324 DOM. Similarly, the fluorescent biological index (BIX), which is associated with
325 microbially-derived and autochthonous DOM, was calculated as the ratio of
326 fluorescence at 380 nm to 430 nm under 308 nm excitation (Huguet et al., 2009).
327 BIX >1 can indicate a strong signal of recent autochthonous DOM production,
328 whereas those <0.7 reflect older autochthonous DOM (Huguet et al., 2009). Lastly,
329 the fluorescent humification index (HIX), often used to estimate the extent of DOM
330 diagenesis or maturation in soils, was calculated as the integrated fluorescence from
331 434 to 480 nm divided by the integrated fluorescence from 300 to 346 nm under
332 254nm excitation (Zsolnay et al., 1999). High HIX values (>10) indicate aromatic
333 DOM (potentially from terrestrial or marine humic acids) whereas low values (<4)
334 reflect more autochthonous origin (Birdwell and Engel, 2010). Lastly, we used the

335 absorbance spectra to calculate specific ultraviolet absorbance (SUVA₂₅₄) by
336 dividing the linear absorbance (m⁻¹) by DOC (mg L⁻¹) (Weishaar et al., 2003).

337

338 *Statistical analyses*

339 All nutrient, organic, carbonate, fDOM indices and flow cytometry parameters were
340 log₁₀-transformed to better approximate a gaussian (normal) distribution before
341 statistical analysis; raw fDOM values were normally distributed and were not
342 transformed for statistical analysis. Hierarchical clustering (Ward's minimum
343 variance method) was used to group samples according to similarity in multiple
344 biogeochemical parameters as a way to define clusters of samples with similar
345 properties. Each parameter was first standardized (by subtracting the column mean
346 and dividing by the column standard deviation) to avoid weighting clusters by
347 absolute measurement values. To conservatively define groups of samples
348 according to relative proportion of SGD influence based on inorganic chemical
349 composition, samples were initially clustered by the full suite of 7 standard
350 inorganic solute measurements made (PO₄³⁻, N+N, NH₄⁺, SiO₄, Salinity, TA and Rn).
351 This clustering approach differentiated samples into spatially-distributed
352 "biogeochemical provinces" interpreted as SGD Springs, Transition SGD mixing
353 Zones, Diffuse SGD Zones and Ambient Reef waters (detailed further in results).
354 Analysis of variance (ANOVA) was then used to test if mean values of organic
355 parameters differed among the inorganic biogeochemical provinces and sites, with
356 Tukey and Dunnet's *post hoc* tests used to assess pairwise differences among groups
357 at $\alpha = 0.05$. Chi-square tests were used to assess similarity in cluster assignment of
358 samples using different suites of organic variables. Pearson correlation and linear
359 regression models (least squares and orthogonal/reduced major axis/Model II
360 approaches) were used to assess covariance among variables.

361

362 **Results:**

363

364 *Distributions of dissolved inorganic solutes and delineation of groundwater influence*

365 At each site SGD sources and dispersal patterns were clearly visualized by contour
366 mapping of conservative inorganic solute tracer concentrations in surface samples
367 at low tide (Figure 2a-f). Concentration gradients were consistent with rapid
368 dilution within 200m of the source springs at each site. Contours of the fDOM
369 humification index (HIX) closely tracked these conservative solute gradients across
370 the reef platform (Figure 2g,h), and HIX was highly correlated with salinity and
371 silicate consistently at both sites ($r > 0.75$; Figure S2), demonstrating that fDOM
372 parameters tracked salinity.

373

374 Hierarchical clustering of samples according to the suite of 7 standard inorganic
375 solute measurements (SiO_4 , Salinity, Rn, PO_4^{3-} , N+N, NH_4^+ and TA) separated
376 samples into six distinct groups, which we refer to subsequently as “biogeochemical
377 provinces” because of their spatial differentiation (Figure 3a). Groundwater springs
378 at Wailupe and Black Point were distinct (BP Spring and WL Spring Provinces),
379 areas of significant SGD mixing at Wailupe and Black Point were distinct (BP
380 Transition Zone and WL Transition Zone Provinces), while Diffuse SGD Zones and
381 Ambient Reef provinces did not differ between sites (Figure 3b,c). Figure 3d,e
382 provides a conceptual spatial schematic of the biogeochemical provinces defined in
383 Figure 3a that are referenced throughout this study (e.g. Spring, Transition Zone,
384 Diffuse Zone and Ambient Reef).

385

386 At the Springs, silicate concentrations were $> 500 \mu\text{mol L}^{-1}$, salinities < 10 and radon
387 activities $> 150 \text{ dpm L}^{-1}$ while Ambient Reef waters had silicate concentrations < 5
388 $\mu\text{mol L}^{-1}$, salinities near 30 and radon activities $< 20 \text{ dpm L}^{-1}$; Transition and Diffuse
389 Zones exhibited characteristic intermediate silicate concentrations and did not
390 differ significantly from Ambient Reef sources in salinity or radon (Table 1, Figure
391 S3). Sites did not differ significantly in any of the inorganic tracer solutes except that
392 Transition Zone waters had more radon at Black Point (mean 148 dpm L^{-1}) than at
393 Wailupe (mean 43 dpm L^{-1} ; Figure S3). Springs at both sites were significantly
394 higher in N+N ($> 50 \mu\text{mol L}^{-1}$) and PO_4^{3-} ($> 1.5 \mu\text{mol L}^{-1}$) than any other samples;

395 nearby Transition Zone samples remained significantly higher (> 5X) than adjacent
396 Diffuse Zone and Ambient Reef waters that did not differ significantly from each
397 other (< 1.5 $\mu\text{mol L}^{-1}$ N+N and < 0.15 $\mu\text{mol L}^{-1}$ PO_4^{3-} ; Figure S4). In contrast, NH_4^+
398 concentrations were depleted in Springs (near limits of detection) relative to the
399 adjacent Transition Zone and Diffuse Zone waters, both of which were strikingly
400 enriched in NH_4^+ (mean 0.9 $\mu\text{mol L}^{-1}$) above Ambient Reef samples (mean 0.3 μmol
401 L^{-1} ; Table 1 and Figure S4). TA in the groundwater Springs differed markedly
402 between the two sites, being significantly elevated at Black Point (mean 2826 μmol
403 kg^{-1}) and significantly depleted at Wailupe (mean 1616 $\mu\text{mol kg}^{-1}$) relative to all
404 other biogeochemical provinces at both sites (which did not differ significantly;
405 mean concentrations 2250 $\mu\text{mol kg}^{-1}$; Table 1 and Figure S4).

406

407 *Distributions of particulate and dissolved organics:*

408 Hierarchical clustering of samples according to a suite of 9 measured dissolved and
409 particulate organic matter concentrations and ratios (Chl *a*, Picoeukaryotic
410 phytoplankton, autotrophic bacterioplankton, heterotrophic bacterioplankton, DOC,
411 DON, DOP, DOC:N and TN:TP) yielded 6 distinct groups (Figure 4a) with spatial
412 distributions of sample types consistent with SGD gradients (Figure 4b,c). Group
413 assignment of samples by clustering on inorganic solutes (Fig. 3a) and organic
414 matter (Fig. 4a) was highly congruent (Contingency $R^2 = 0.63$, Pearson Chi-square
415 0.96 and $p < 0.0001$ for Low Tide samples), with 75% of the samples assigned to
416 identical groups (Figure 4a). Spring samples were all assigned perfectly, but Black
417 Point samples were more homogenous spatially in terms of organic matter than
418 observed with inorganic solutes and did not separate clearly among Transition
419 Zone, Diffuse Zone and Ambient Reef types, potentially indicating a more extensive
420 influence of SGD on the reef organic field (Figure 4b). Wailupe spatial patterning of
421 organic matter was consistent with inorganic solutes (Figure 4c).

422

423 DOC was significantly depleted in both Springs relative to the surrounding waters
424 (mean 85 $\mu\text{mol L}^{-1}$); concentrations in Wailupe Springs (mean 20 $\mu\text{mol L}^{-1}$) were

425 more than twice as low as those in Black Point Springs (mean 47 $\mu\text{mol L}^{-1}$; Table 1
426 and Figure S5). The two Springs had very different DON concentrations, both
427 significantly different from the surrounding waters (mean 6.5 $\mu\text{mol L}^{-1}$), with Black
428 Point highly enriched (mean 34 $\mu\text{mol L}^{-1}$) and Wailupe significantly depleted (mean
429 1.4 $\mu\text{mol L}^{-1}$). Dissolved organic phosphorus was unresolvable in Springs and near
430 detection limits in Transition Zone regions, and did not differ among Diffuse Zones
431 and Ambient Reef waters (mean 0.3 $\mu\text{mol L}^{-1}$). Particulate organic concentrations
432 (POC, PON and chl *a*; Fig. S6) and flow cytometry (Picoeukaryotic phytoplankton,
433 Autotrophic and Heterotrophic bacterioplankton; Fig S7) pairwise differences
434 among biogeochemical provinces were mostly non-significant due to high variance,
435 but overall exhibited trends of particulate depletion in Springs and plankton
436 enrichment in the surrounding Transition Zone waters (Figs. S6, S7).

437

438 *fDOM characteristics and distributions:*

439 The PARAFAC modelling validated 4 fluorescence components that covaried with
440 fluorescence regions widely identified from marine systems (Regions: A - humic-like
441 UV excitation; M - visible, blue-shifted, marine humic-like; C - visible excitation,
442 humic-like; and T - aromatic amino protein-like; Coble 1996) both in terms of
443 spectral characteristics (Fig. S8) and in terms of standardized distributions among
444 variables within this dataset (Figure S8e). In addition, the spectral loadings of our
445 PARAFAC components (Fig. S8a-d) matched PARAFAC components found in marine
446 systems in various recent reviews: Component 1 (Em(2°)/Ex: 260(375) nm/375
447 nm) is consistent with component C1 from Jørgensen et al. (2011) and component
448 C2 from Ishii and Boyer (2012). Our Component 2 (Em(2°)/Ex: <250(325) nm /
449 400-480 nm) corresponds to component C4 from Jørgensen et al. (2011) and
450 component C3 from Ishii and Boyer (2012). Component 3 (Em/Ex(2°): 300-
451 380nm/510(480nm) corresponds to component C4 in Kowalczyk et al. (2009).
452 Component 4 (Ex/Em(2°): 260nm/330(510nm) corresponds to component C2 in
453 Jørgensen et al. (2011).

454

455 Each of the four PARAFAC components differed significantly among the
456 biogeochemical provinces (ANOVA $p < 0.0001$). At Black Point both Spring and
457 Transition Zone provinces were enriched relative to Diffuse Zone and Ambient Reef
458 waters for all components, indicating that total fDOM was elevated in the
459 groundwater. In contrast, Wailupe Transition and Diffuse Zones were enriched
460 relative to both Spring and Ambient Reef waters (Fig. 5), consistent with the idea of
461 production of fDOM in the SGD-influenced reef waters of Wailupe. Notably, at Black
462 Point, although fDOM decreased from Spring to Ambient Reef waters, DOC exhibited
463 some enrichment in the transition and diffuse zones, suggesting that autochthonous
464 production of non-fluorescent DOM may have occurred in the diffuse zones.
465 Within any of the four biogeochemical provinces there were clear site differences in
466 fDOM quantity: Black Point Spring samples were enriched in all four components
467 relative to Wailupe Springs while the reverse was true in Diffuse Zone samples, with
468 Wailupe enriched for components A, M and C (Fig. 5), again consistent with the idea
469 of production of fDOM in the SGD-influenced reef waters of Wailupe.

470

471 The four ratio-based fDOM indices exhibited very different patterns within sites,
472 emphasizing that each index is assessing a different aspect of the character of fDOM
473 (Figure 6). In SGD Springs, the humification index, HIX, was more than double (> 7)
474 Ambient Reef values (< 3) at both sites (Table 1). HIX was highly correlated with
475 salinity and silicate consistently between sites ($r > 0.75$; Figure S2) and declined
476 continuously with distance from the Springs (Figure 2). The M:C index covaried with
477 HIX ($r = 0.83$) and was also significantly enriched in Springs and in Transition Zone
478 waters relative to the Diffuse Zone and Ambient Reef waters. Relative to ambient
479 waters, the Wailupe Spring had a significantly higher fluorescence index (FI) and
480 $SUVA_{254}$ was significantly greater in the Black Point Spring. BIX ranged from 0.76-
481 0.86 across both sites, and was generally elevated in Black Point relative to Wailupe,
482 but did not differ significantly among water types. Indices did not differ between
483 sites within a given biogeochemical province (Figure 6).

484

485 Hierarchical clustering of samples according to a suite of 9 fDOM-derived
486 parameters (4 PARAFAC components, 4 fDOM indices and SUVA₂₅₄) separated
487 samples into 6 groups (Figure 7a) with spatial distributions of sample types
488 consistent with SGD gradients (Figure 7b,c). Group assignment of samples by fDOM
489 characteristics was generally congruent with clustering by inorganic solutes (Fig. 3)
490 and organic matter (Fig. 7; Contingency $R^2 = 0.52$ and 0.50 , respectively, Pearson
491 Chi-square 93 and 78, respectively and $p < 0.0001$ for Low Tide samples), with 70%
492 of the samples assigned to identical groups. As with the organic matter clustering,
493 Spring samples were all assigned perfectly, but at both sites the other
494 biogeochemical provinces were less clearly differentiated spatially in terms of fDOM
495 than observed with inorganic solutes. Notably, both Transition Zone and Diffuse
496 Zone samples appeared to be different in fDOM parameters between Wailupe and
497 Black Point (Figure 7a), potentially indicating a more extensive influence of SGD on
498 the reef fDOM field. The two SGD springs (and the two diffuse zones) may be
499 differentiated according to their fDOM amount (i.e. fluorescence intensities of
500 fluorophores) (Fig. 5) but cannot be clearly discriminated by fluorophore ratio
501 indices (Fig. 6).

502

503 **Discussion**

504

505 *Biogeochemical characteristics of groundwater entering Maunalua Bay*

506 Groundwater discharging from springs in Maunalua Bay showed some consistencies
507 and differences between the Black Point and Wailupe sites. Both sites released
508 groundwater with biogeochemical profiles consistent with previous studies of SGD
509 in coral reefs (Swarzenski et al., 2013), including low salinities and elevated
510 concentrations of radon and silicate, elevated PO_4^{3-} and $\text{N}+\text{N}$, depleted NH_4^+
511 concentrations and depleted DOC, POC, chl *a* and phytoplankton cells (Table 1, Figs.
512 2-4 and S3-S7). However, some measurements were strongly and significantly
513 different between the two Springs. First, the Wailupe Springs were significantly
514 depleted in TA and DON relative to the adjacent waters whereas the Black Point TA
515 and DON were significantly greater than the adjacent waters, suggesting a

516 fundamentally different hydrological origin (Figs. S4, S5). Second, while both
517 Springs were depleted in DOC relative to the adjacent waters, the Wailupe site had
518 nearly half the DOC concentrations of the Black Point site (Fig. S5), yet Black Point
519 Springs were strongly depleted in bacterioplankton, nearly 4 times less than
520 Wailupe Springs or the surrounding ocean (Fig. S7). Aside from the unexpectedly
521 high concentrations of bacterioplankton in the Wailupe Springs, these patterns
522 suggest that groundwater flowpaths at these two sites are very different, potentially
523 capturing differences in land-use and geology in these two parts of the watershed.
524 Black Point has a higher density of on-site sewage disposal (septic and cesspool)
525 systems (Whittier and El-Kadi, 2009), but we do not have evidence that the high
526 nutrient or fDOM levels are due to these potential sources. The difference in radon
527 concentrations between the springs (Table 1) suggest that groundwater at Black
528 Point flows through rocks generating more radon (i.e., more enriched in U and Ra)
529 than the rocks and sediments at Wailupe, indicating differences in the geologic
530 make-up of the aquifer. Land-use, including density of septic systems, presence of
531 historic agricultural sites, etc. is a likely cause for differences in nutrient and organic
532 matter levels.

533

534 *fDOM characteristics of groundwater entering Maunalua Bay*

535 Across sites, SGD was significantly enriched in aromatic/humic components (e.g.,
536 Regions A and M, HIX, M:C) and had higher specific ultraviolet absorbance (Table 1).
537 The Black Point Springs had significantly more of all 4 fDOM components than the
538 Wailupe Springs and were enriched in all components relative to the Ambient Reef
539 waters (Fig. 5); the Wailupe Springs had fDOM quantities identical to Ambient Reef
540 waters for all four components. The lack of significant differences in fDOM indices
541 between these two sites within any given biogeochemical province (Fig. 6) suggests
542 that SGD fDOM molecular composition was similar. However, groundwater DOC
543 concentrations overall were significantly lower than the overlying reef, and when
544 combined with elevated fDOM (at least at Black Point) and significantly higher ratios
545 of humic compounds (HIX Index, Figs 2 and 6) our data indicate that a greater
546 proportion of the DOC in the groundwater is fluorescent. Indeed, ratios between any

547 of the four PARAFAC components and DOC concentrations, a proxy for specific
548 fDOM fluorescence, were significantly enriched in both Springs relative to the
549 adjacent waters (ANOVA with Tukey *post hoc* $p < 0.0001$), indicating that
550 groundwater DOC has a much higher fluorescence than marine DOC. Although both
551 overall fDOM and DOC concentrations were significantly higher in the Black Point
552 site (Figs 5 and S5), ratios of fDOM components to DOC did not differ between sites,
553 only between biogeochemical provinces, again emphasizing that the DOC in SGD is
554 consistently highly fluorescent with a strong humic component (Figs 2, 6, S2).

555

556 *Dispersal and biogeochemical influence of submarine groundwater discharge*

557 In an observational study it is difficult to separate cause from effect, and in the case
558 of the current dataset we cannot definitively determine whether parameters that
559 differ significantly in the Transition and Diffuse Zones from the Ambient Reef waters
560 are driven by dilution of SGD or stimulation of biogeochemical processes that
561 subsequently alter the characteristics of the water. However, the vast majority of the
562 measured parameters exhibited statistically robust gradients spatially concordant
563 with SGD dispersal across the reef. A handful of parameters did not follow this
564 trend, and it is likely that these represent reef-specific production or consumption
565 processes. Ammonium was the only inorganic solute that exhibited enrichment in
566 the Transition Zone and Diffuse Zone provinces above both Spring and Ambient
567 Reef endmembers, suggesting a production process (Figure S4). Moreover, this
568 parameter was only significantly enriched in the Transition Zone province, not in
569 the adjacent Reef waters, possibly indicating rapid consumption and
570 ammonification of DON, dissimilatory reduction of nitrate to ammonia (DNRA), or
571 recycling of nitrate to ammonia through organic assimilation and remineralization.
572 A similar pattern was observed in particulate organic matter, with the Wailupe site
573 exhibiting enriched Chl *a* and eukaryotic phytoplankton counts suggestive of some
574 stimulation of water column productivity by the groundwater nutrient delivery.
575 Finally, there was consistent enrichment of fDOM components A,C and T in the
576 Transition Zone and Diffuse Zone provinces above levels found in the Springs and
577 Ambient Reef zones (Fig. 5). Region C has been associated with microbial

578 production processes in marine systems and Region T is generally associated with
579 proteinaceous material because of the similarities to pure tryptophan fluorescence
580 (Coble et al., 2014). One interpretation of these patterns is that these components
581 represent fDOM being produced by the reef habitat; whether that production is
582 influenced directly or indirectly by SGD inputs or is simply a characteristic of reefs
583 generally is not currently known. Certainly there are potential physicochemical
584 changes in DOM across these gradients due to photodegradation, metal-ligand
585 bonding, pH, and salinity shifts that impart changes to fDOM character (Helms et al.,
586 2013; Osburn et al., 2013).

587

588 *Recommendations for the use of fDOM to track groundwater discharge in reefs*

589 This study shows clearly how fDOM spectral analyses can be used to differentiate
590 water masses according to the degree of influence of SGD in two sites with very
591 different SGD organic matter profiles, consistent with previous work conducted in
592 coral reef environments with allochthonous DOM inputs (Tedetti et al., 2011). Based
593 on our observations, it is clear that groundwater entering Maunalua Bay contains a
594 significant quantity of fDOM with more than double the humification index (HIX) of
595 the receiving Diffuse Zone and Ambient Reef waters (Figure 6, Table 1), consistent
596 with fDOM from sedimentary and volcanic sources in reef ecosystems (Tedetti et al.,
597 2011). This single feature of the fDOM spectra (i.e., elevated HIX) was strongly
598 correlated with both salinity and silicate in both sites with identical slopes, and from
599 this simple index there is the potential to model these inorganic solute
600 concentrations from the HIX value (Figure S2). In addition, the groundwater sources
601 to Black Point stood out clearly from the surrounding reef in humic fluorescence
602 regions A and M (Figure 5), though not at Wailupe. These differences suggest that
603 fDOM characteristics may be able to differentiate groundwater according to land
604 use, hydrology, or other factors, allowing the development of fDOM as a
605 groundwater source-tracking tool in concert with other biogeochemical parameters.
606 Future studies should consider the use of continuous sensor monitoring of coastal
607 fDOM (our PARAFAC component C1 exhibits a secondary emission peak
608 corresponding to the excitation-emission maxima of commercial DOM sensors) and

609 examination of the interacting roles of photobleaching and residence time in
610 defining the extent of fDOM distributions in coastal waters. Because fDOM samples
611 are relatively easy to collect (filtering a few mL of water into glass vials and dark
612 refrigerated storage), are unaffected by gas exchange and quick to analyze (the
613 scans took less than 5 minutes each), fDOM may prove a cost-effective and efficient
614 monitoring tool for mapping groundwater dispersal in reefs. Analyzing a sample
615 EEM spectra also provides a wealth of additional ratio-based indices and values of
616 the literature-derived identified spectral regions and our results hint that with
617 larger datasets from more reefs fDOM may be used as cost-effective monitoring tool
618 to identify new and promising indices to differentiate SGD within coastal waters.

619

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621

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FIGURE LEGENDS

Figure 1. Maps of the sampling locations. Panel (a) shows the location of the two sampling sites in Maunalua Bay, O‘ahu, with the inset showing the location within the Main Hawaiian Islands. Water samples were collected at each site, Black Point to the West (b) and Wailupe to the East (c), with red markers at each location where water was collected.

Figure 2. Fluorescent dissolved organic matter (fDOM) in the spatial context of Submarine Groundwater Discharge (SGD) in Maunalua Bay. Contour plots of conservative solutes and fDOM solutes at Black Point (a,c,e,g) and Wailupe (b,d,f,h) 28-29 May 2014, including salinity (a,b), silicate concentrations (c,d) ($\log_{10} \mu\text{mol L}^{-1}$), radon concentrations (e,f) (dpm L^{-1}) and the fDOM humification index (g,h) (HIX). Contour gridding and interpolation was done with the kriging function (spherical semivariogram model) in ArcGIS 10.3 Spatial Analyst. All contour plots were generated from samples collected at low tide.

Figure 3. Hierarchical clustering of samples into biogeochemical provinces according to inorganic solute concentrations. Panel (a) is a hierarchical clustering dendrogram grouping samples according to similarity in \log_{10} concentrations of 7 inorganic solutes (shown in heat map with legends at right; X indicates no data). The tips of the dendrogram are colored to match the dendrogram clusters and define biogeochemical provinces by sets of samples with similar chemistry. WL refers to Wailupe and BP refers to Black Point. Panels (b) and (c) illustrate the spatial extent of each biogeochemical province by mapping the sample points color-coded by dendrogram clusters - defined with large colored boxes in panel (a) - at Black Point and Wailupe, respectively. Panels (d) and (e) provide a conceptual illustration of the spatial arrangement of biogeochemical provinces.

Figure 4. Hierarchical clustering of samples according to particulate and dissolved organic matter concentrations. Panel (a) is a hierarchical clustering dendrogram grouping samples (tips colored according to biogeochemical provinces delineated in Fig. 3) according to similarity in \log_{10} concentrations of 8 organic measurements (shown in heat map with legends at right; X indicates no data). Panels (b) and (c) define the spatial extent of organic provinces by mapping the sample points color-coded by dendrogram clusters - defined with large colored boxes in panel (a) - at Black Point and Wailupe, respectively.

Figure 5. Comparison of fDOM PARAFAC components among the biogeochemical provinces in each reef site. Boxes depict standard interquartile ranges with medians and are labeled at top with letters for ANOVA Tukey *post hoc* tests; all ANOVA models $p < 0.0001$ and samples with different letters are significantly different at $\alpha = 0.05$.

Figure 6. Comparison of fDOM indices among the biogeochemical provinces in each reef site. Boxes depict standard interquartile ranges with medians and are

labeled at top with letters for ANOVA Tukey *post hoc* tests; all ANOVA models $p < 0.01$ and samples with different letters are significantly different at $\alpha = 0.05$.

Figure 7. Hierarchical clustering of samples according to fDOM spectral characteristics. Panel (a) is a hierarchical clustering dendrogram grouping samples (tips colored according to biogeochemical provinces delineated in Fig. 3) according to similarity in fDOM PARAFAC components and ratio-based indices (shown in heat map with legends at right; X indicates no data). Panels (b) and (c) define the spatial extent of fDOM character by mapping the sample points color-coded by dendrogram clusters – defined with large colored boxes in panel (a) – at Black Point and Wailupe, respectively.

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	Wailupe				Black Point			
	Spring	Transition	Diffuse	Reef*	Reef	Diffuse	Transition	Spring
Practical Salinity	4.2	24.4	30.4	28.8	30.5	32.6	25.5	7.1
SiO ₄ (μmol L ⁻¹)	688.5	225.2	32.2	2.8	3.2	16.6	106.8	626.9
Rn (dpm L ⁻¹)	160.4	30.4	23.7	5.6	1.7	14.2	127.2	269.7
N+N (μmol L ⁻¹)	61.63	8.50	0.24	0.14	0.23	1.16	21.15	114.16
PO ₄ ³⁻ (μmol L ⁻¹)	1.89	0.49	0.05	0.04	0.10	0.15	0.68	3.09
NH ₄ ⁺ (μmol L ⁻¹)	0.05	0.89	0.43	0.26	0.34	0.52	0.94	0.22
TA (μmol kg ⁻¹)	1614	2143	2234	2280	2238	2263	2362	2824
DOC (μmol L ⁻¹)	20.0	70.8	91.4	83.7	77.3	88.3	88.5	45.6
DON (μmol L ⁻¹)	1.4	5.0	6.8	5.7	6.1	6.6	6.6	33.6
Chl a (μg L ⁻¹)	0.06	0.29	0.12	0.08	0.07	0.09	0.14	0.03
HBact (cells mL ⁻¹)	4.5×10 ⁵	2.3×10 ⁵	5.0×10 ⁵	3.2×10 ⁵	3.7×10 ⁵	2.3×10 ⁵	2.2×10 ⁵	1.0×10 ⁵
PBact (cells mL ⁻¹)	1.6×10 ³	7.2×10 ³	1.8×10 ³	4.0×10 ³	1.8×10 ³	1.9×10 ³	1.4×10 ³	7.2×10 ²
PEuks (cells mL ⁻¹)	7.7×10 ³	9.2×10 ⁴	6.6×10 ³	6.0×10 ³	4.9×10 ³	5.2×10 ³	6.8×10 ³	5.4×10 ³
fDOM: A	0.028	0.061	0.056	0.025	0.029	0.037	0.052	0.075
fDOM: M	0.036	0.073	0.065	0.029	0.034	0.043	0.062	0.091
fDOM: C	0.006	0.018	0.018	0.007	0.009	0.011	0.015	0.017
fDOM: T	0.008	0.026	0.026	0.016	0.017	0.021	0.027	0.023
SUVA ₂₅₄	0.95	0.87	0.82	0.65	0.98	0.72	0.88	1.44
M:C	1.25	1.11	1.05	1.02	1.01	1.03	1.07	1.20
FI	1.82	1.72	1.73	1.73	1.79	1.75	1.75	1.76
BIX	0.83	0.79	0.82	0.80	0.84	0.83	0.82	0.85
HIX	7.78	4.28	3.60	2.35	2.74	3.00	3.47	6.98

Table 1. Mean values of each parameter in each of the biogeochemical provinces shaded according to magnitude of significant differences among regions. Rows are parameters, columns are biogeochemical provinces, and values are geometric means. Shaded cells are significantly different from the Ambient Reef (WL) province (denoted at top by *; Dunnet's *post hoc* test), with color and intensity scaled by mean log-ratio relative to Ambient Reef waters (legend at left). POC, PON and DOP are excluded due to a lack of data to properly test each province.

Figure 1



Figure2

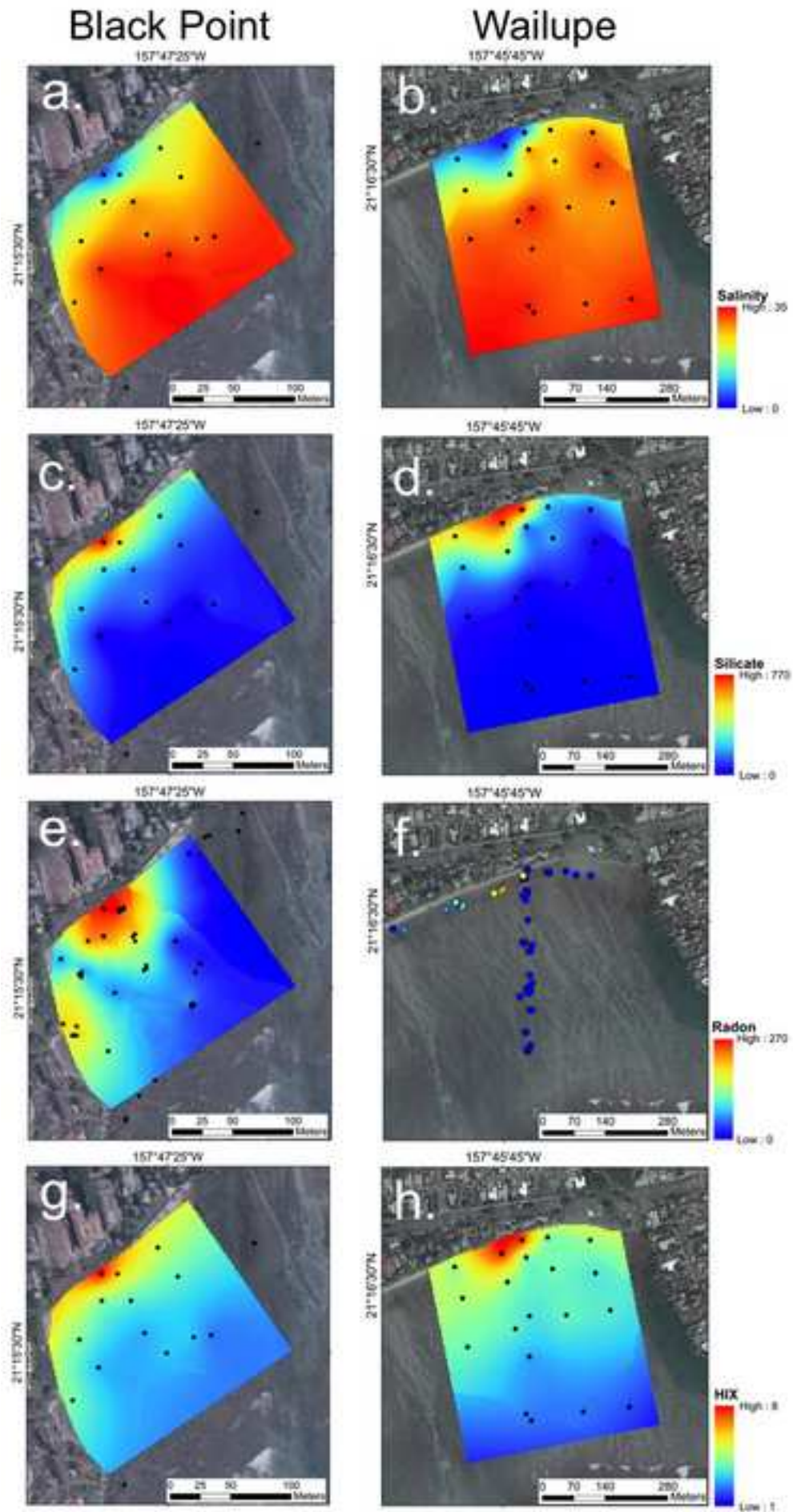


Figure3

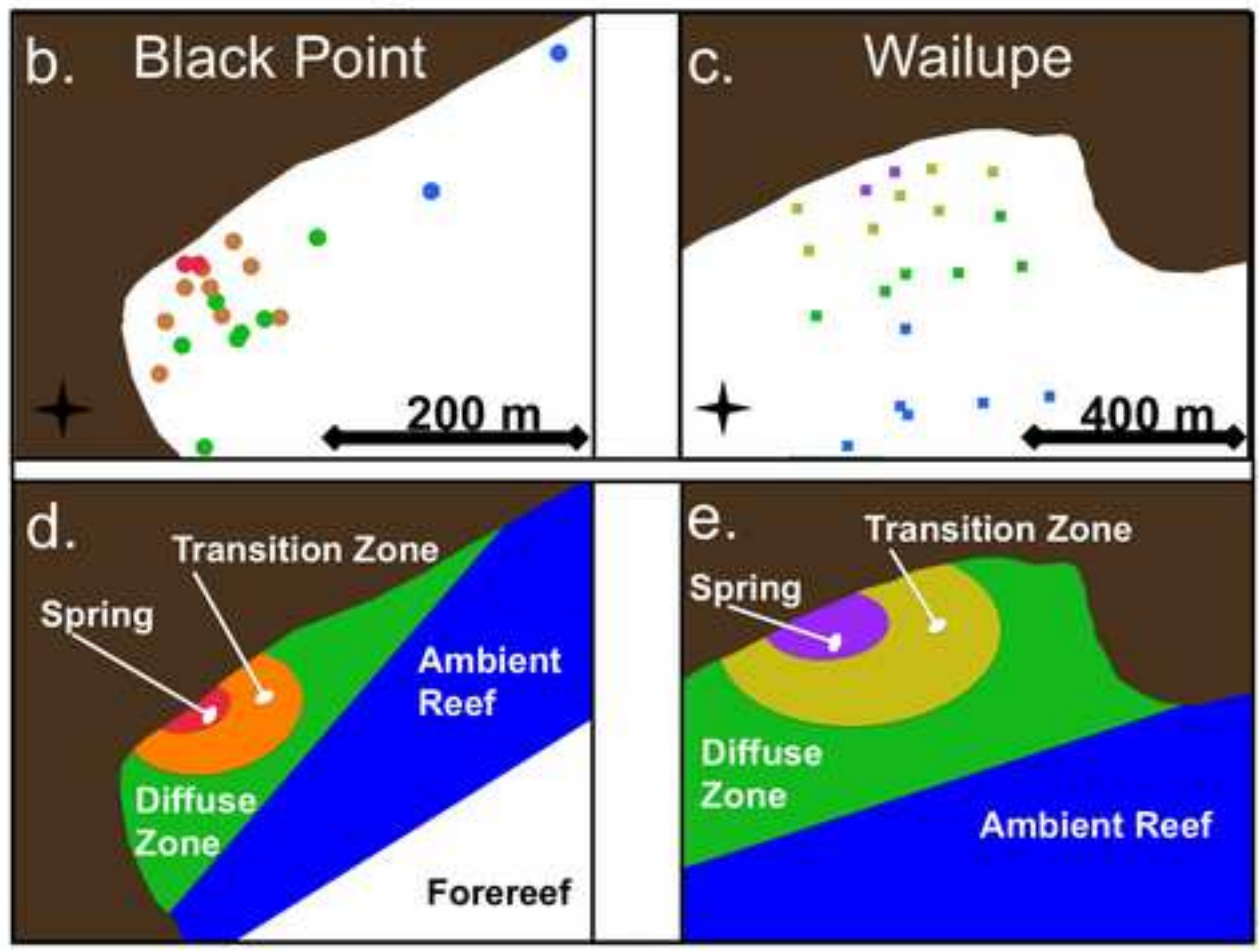
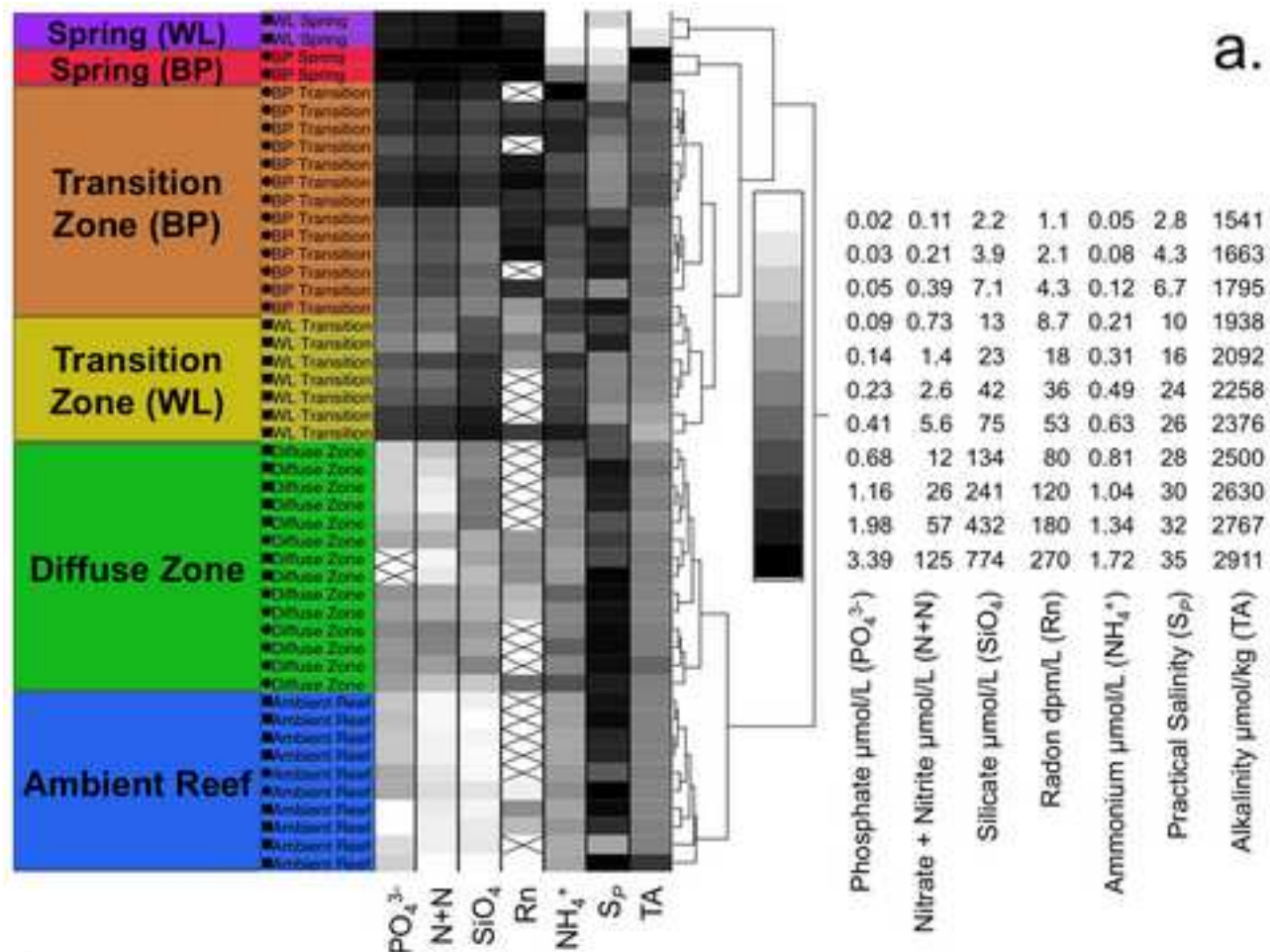


Figure 4

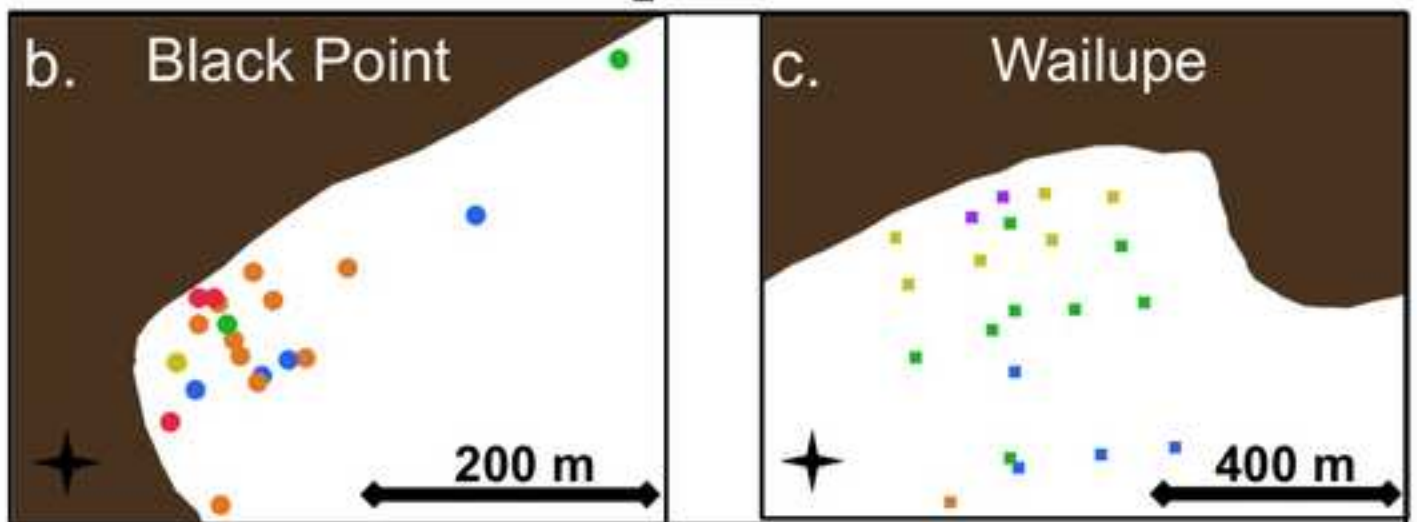
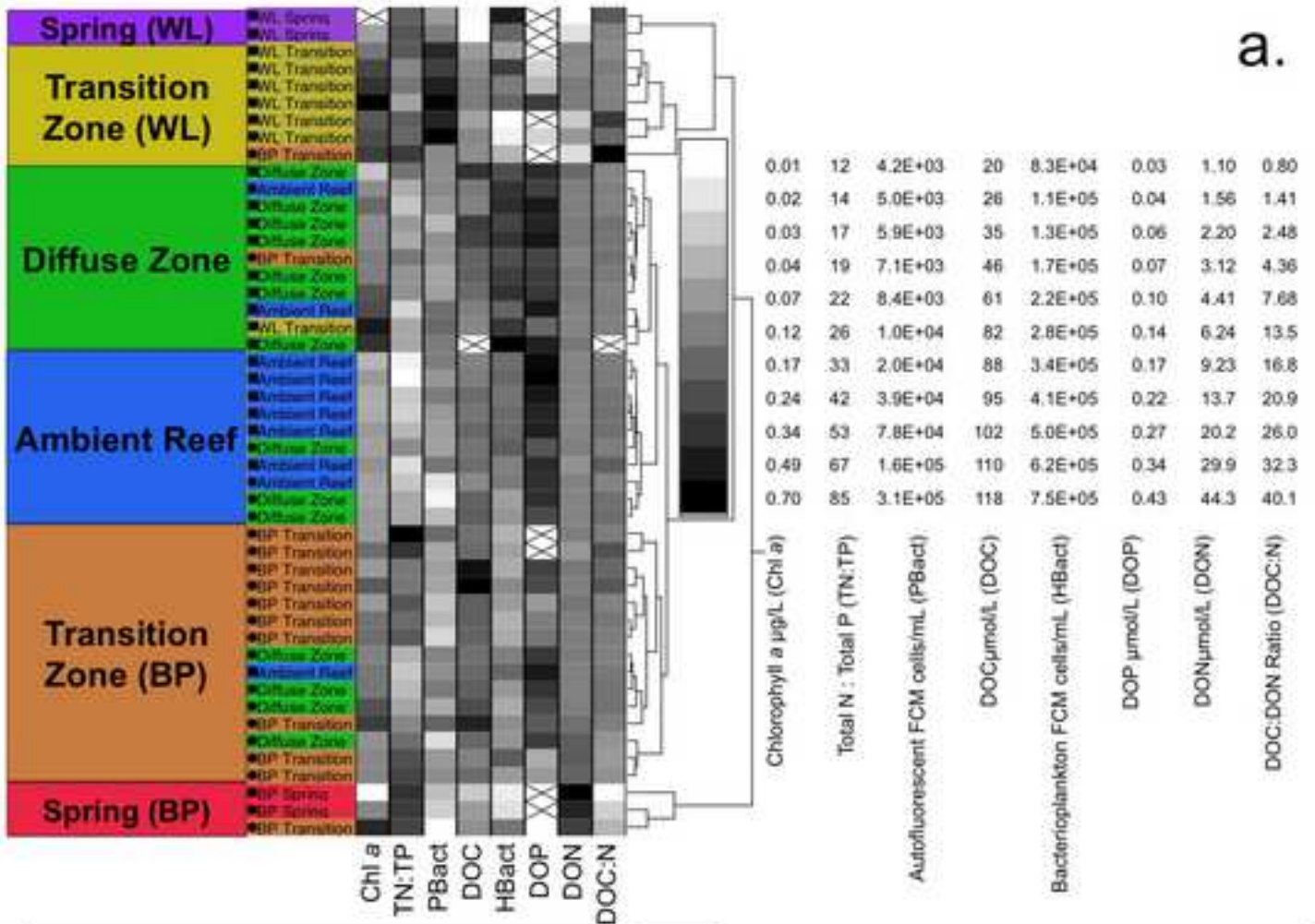


Figure 5

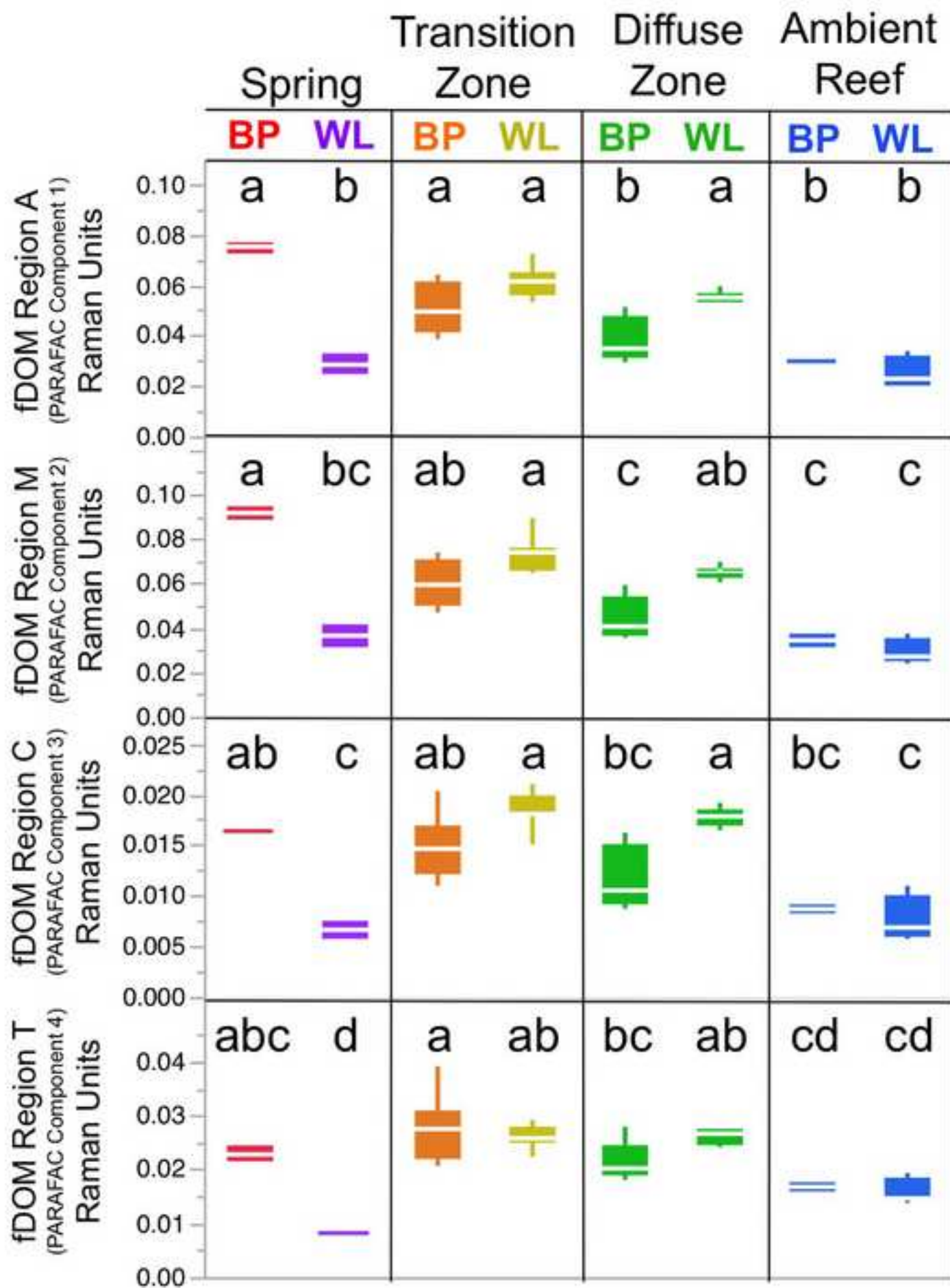


Figure 6

