

1 **Potential bioavailability of representative pyrogenic organic matter**
2 **compounds in comparison to natural dissolved organic matter pools**

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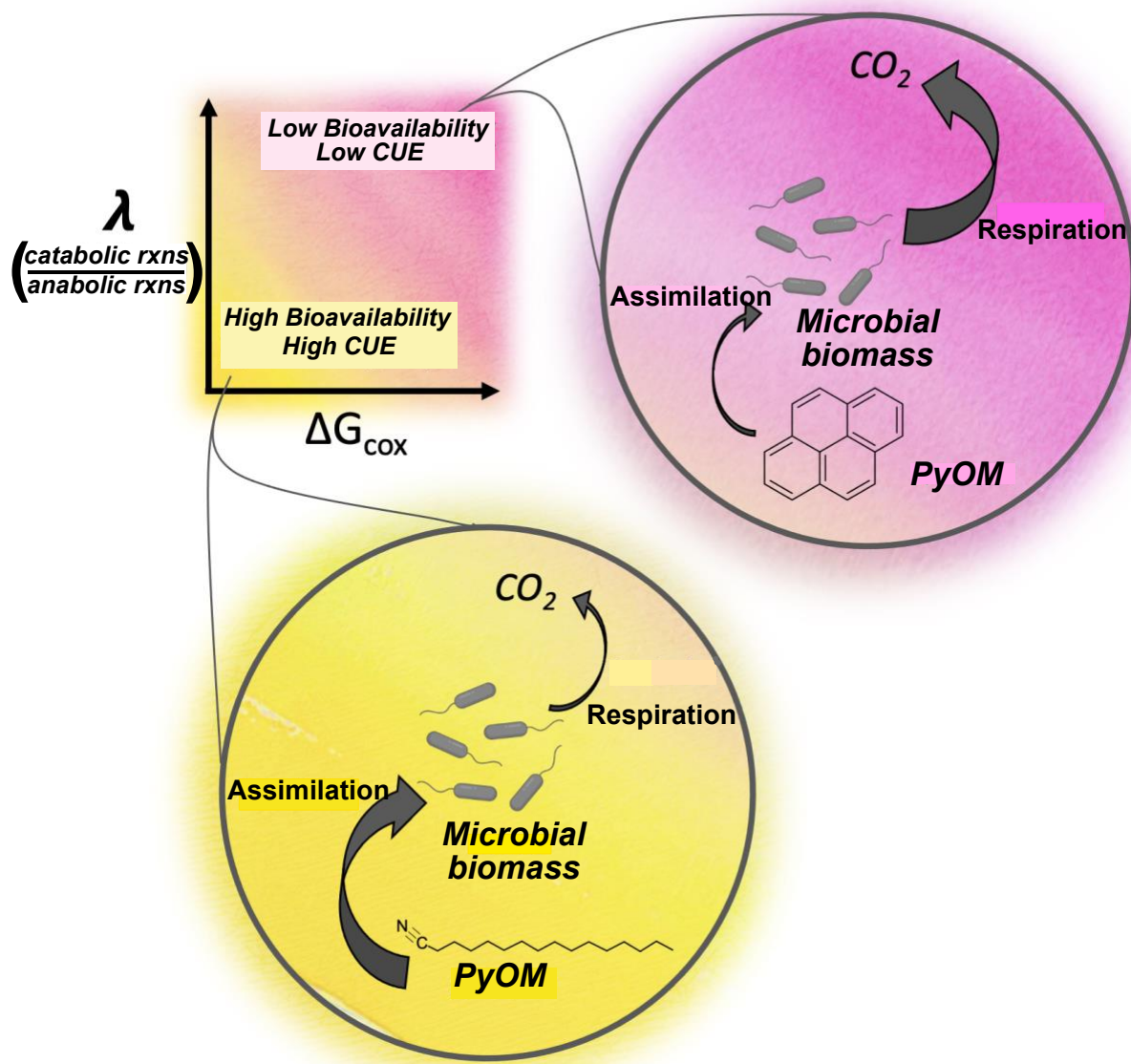
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20 GRAPHICAL ABSTRACT



21

22

23 **Abstract.** Pyrogenic organic matter (PyOM) from wildfires impacts river corridors globally and
24 is widely regarded as resistant to biological degradation. Though recent work suggests PyOM
25 may be more bioavailable than historically perceived, estimating bioavailability across its
26 chemical spectrum remains elusive. To address this knowledge gap, we assessed potential
27 bioavailability of representative PyOM compounds relative to ubiquitous dissolved organic
28 matter (DOM) with a substrate-explicit model. The range of potential bioavailability of PyOM
29 was greater than natural DOM; however, the predicted thermodynamics, metabolic rates, and
30 carbon use efficiencies overlapped significantly between all OM pools. Compound type (e.g.,
31 natural vs. PyOM) had approximately 6-fold less impact on predicted respiration rates than
32 simulated carbon and oxygen limitations. Within PyOM, the metabolism of specific chemistries
33 differed strongly between unlimited and oxygen limited conditions—anhydrosugars, phenols,
34 and polycyclic aromatic hydrocarbons degradations were more favorable under oxygen
35 limitation than other molecules. Notably, amino sugar-, protein-, and lignin-like PyOM had
36 lower carbon use efficiencies relative to natural DOM of the same classes, indicating potential
37 impacts in process-based model representations. Overall, our work illustrates how similar PyOM
38 bioavailability may be to that of natural DOM in the river corridor, furthering our understanding
39 of how PyOM may influence riverine biogeochemical cycling.

40 **1 Introduction**

41 Wildfires have burned an average of 1.8-million ha year⁻¹ in the United States alone over
42 the past 80 years, dramatically impacting terrestrial and aquatic ecosystems (Bladon et al., 2014;
43 Shakesby and Doerr, 2006; Randerson et al., 2006; Verma and Jayakumar, 2012). As wildfire
44 activity continues to increase in response to climate change (Pierce et al., 2004; Bowman et al.,
45 2020; Flannigan et al., 2009), its impact on river corridor biogeochemistry is receiving
46 significant attention (Wagner et al., 2018; Abney et al., 2019).

47 Pyrogenic organic matter (PyOM) generated by wildfires influences river corridor
48 biogeochemistry due to the importance of organic matter as a carbon (C) and energy source.
49 Though there is substantial uncertainty in the quantification of PyOM, in part due to its chemical
50 diversity, estimates suggest that 116–385 Tg of condensed PyOM is generated per year. This
51 amounts to 300 to 500 Tg C stored in sediments, soils, and waters (Jaffé et al., 2013; Dittmar et
52 al., 2012; Hockaday et al., 2007; Santin et al., 2016) and ~10% of dissolved organic C pools in
53 surface waters (Jaffé et al., 2013). Given that organic matter drives aquatic biogeochemical
54 cycles, the loading of PyOM into river corridors has the potential to produce substantial impacts
55 on ecosystem functions and downstream drinking water treatability (Emelko et al., 2011; Hohner
56 et al., 2017).

57 While some estimates place aquatic residence times at thousands of years (Meyer and
58 Wells, 1997; Elliott and Parker, 2001; Bigio et al., 2010; Kuzyakov et al., 2014), recent work has
59 shown that PyOM may be more bioavailable (i.e., biolabile or able to be accessed and degraded
60 by microorganisms) than previously thought (Myers-Pigg et al., 2015; Norwood et al., 2013;
61 Zimmerman and Ouyang, 2019). This inference is also supported by research on biochars,
62 highlighting the diverse reactivities of combustion byproducts (Sohi et al., 2010; Mia et al.,

63 2017). PyOM bioavailability may, therefore, play an unrecognized role in global biogeochemical
64 cycles and climate feedbacks. Yet, we are just beginning to understand its potential
65 bioavailability (Zimmerman and Mitra, 2017; Wozniak et al., 2020; Masiello, 2004).

66 We used a substrate-explicit model to assess the potential bioavailability of PyOM in
67 comparison to dissolved organic matter (DOM) from global surface waters and sediments. We
68 derived 16,971 representative PyOM compounds from the primary literature describing plant
69 charcoals, biochar, smoke, burnt soil, and their leachates (Table S1). Natural DOM pool
70 composition was derived from a recent high-resolution survey of river corridor sediments and
71 surface waters (Garayburu-Caruso et al., 2020a, Figure S1). The substrate-explicit model used
72 molecular formulae to predict energetic content, metabolic efficiency, and rates of aerobic
73 metabolism for organic substrates. It has proven useful in linking DOM composition to aerobic
74 metabolism in natural environments (Song et al., 2020; Graham et al., 2017; Garayburu-Caruso
75 et al., 2020b). It provides a systematic way to formulate reaction kinetics and is agnostic of many
76 factors that have complicated a universal understanding of OM bioavailability; including
77 molecular structure, chemical inhibition, mineral-associations and physical protection, terminal
78 electron acceptors, microbial community composition and accessibility, and abiotic reactions
79 (reviewed in Arndt et al. (2013)). We hypothesize that PyOM compounds have a greater
80 similarity in their potential bioavailability to natural DOM pools than expected based on
81 historical literature.

82

83 **2 Results and discussion**

84 Our model-based approach enabled us to directly compare known combustion products to
85 thousands of ubiquitous DOM compounds, which would have been unfeasible to directly assess

86 in a laboratory setting. Key outputs from the model include lambda (λ , the number of catabolic
87 reactions that must occur to provide the energy required for the synthesis of one mole of biomass
88 carbon), Gibbs free energy of C oxidation (ΔG_{Cox}), under standard conditions with a modification
89 to pH 7, carbon use efficiency (CUE), and the rate of aerobic metabolism (as oxygen consumed
90 per mol-C biomass produced) under three scenarios: (a) no limitations, (b) C-limitation and (c)
91 oxygen (O_2) limitation. Though the relative magnitude of these metrics will vary based on the
92 specific stoichiometry of a molecule, highly bioavailable compounds are indicated by low λ and
93 ΔG_{Cox} coinciding with high CUE and metabolic rates. Low λ values denote efficient energetics
94 of catabolism in producing biomass through anabolism. Low ΔG_{Cox} denotes high thermodynamic
95 favorability in an electron donor half reaction associated with organic matter, and high CUE
96 reflects more C assimilated into biomass per unit C respired.

97

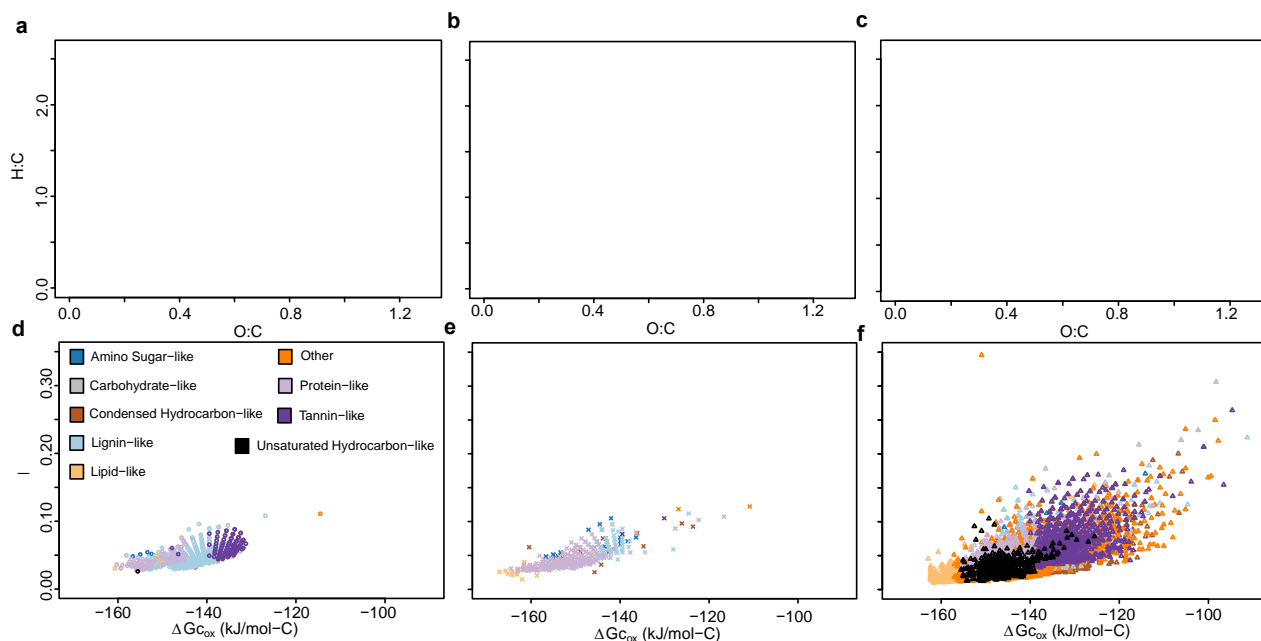
98 *2.1 Potential Bioavailability of Pyrogenic Organic Matter*

99 Though previous work has shown that sediment and surface water DOM is altered by
100 wildfires (Cawley et al., 2018; Jaffé et al., 2013; Wagner et al., 2018), our results suggest that the
101 chemically distinct pools of PyOM transported to aquatic systems may have similar potential
102 bioavailability to DOM. We found that the ranges of ΔG_{Cox} , λ , and CUE overlapped substantially
103 between PyOM and DOM in sediments and surface waters (Figure 1 and 2a-b, Figure S2).
104 Predicted CUE of PyOM classes was also comparable to literature values reported by others
105 (Saifuddin et al., 2019; Domeignoz-Horta et al., 2020; Pold et al., 2020). While λ , ΔG_{Cox} and
106 CUE were significantly different when comparing all three groups (ANOVA, $P < 0.001$, $F =$
107 16.12, 604.9, and 461.2 respectively), λ did not vary significantly between sediment DOM and
108 PyOM (Tukey HSD, $P = 0.21$) and was more similar between sediment DOM and PyOM than

109 between the surface water and sediment DOM pools (mean difference, surface water-sediment
110 DOM = 0.0025, sediment DOM-PyOM = 0.0019) . The range of all PyOM metrics was greater
111 than that of surface water or sediment DOM, reflecting the known heterogenous nature of PyOM
112 chemistries. Though the λ of many PyOM compounds was higher than the maximum observed
113 in either DOM pool, minimum values of λ were lowest for PyOM (surface water DOM = 0.026,
114 sediment DOM = 0.015, PyOM = 0.010), indicating that a portion of wildfire-derived molecules
115 may be more bioavailable than extant DOM pools. It is also interesting that surface water and
116 sediment DOM had greater dissimilarity in ΔG_{Cox} and CUE than any comparison involving
117 PyOM. For example, the mean differences in ΔG_{Cox} and CUE between surface water and
118 sediment DOM were 7.34 kJ/mol-C and 0.058. The differences between PyOM and surface
119 water were 4.28 kJ/mol-C for ΔG_{Cox} and 0.024 for CUE. Overall, statistical differences were not
120 surprising given an extremely large sample size (sediment $n = 398$, surface water $n = 811$, PyOM
121 $n = 16,971$), and the low effect sizes denote that differences in bioavailability between PyOM
122 and DOM were minimal despite statistical separation.

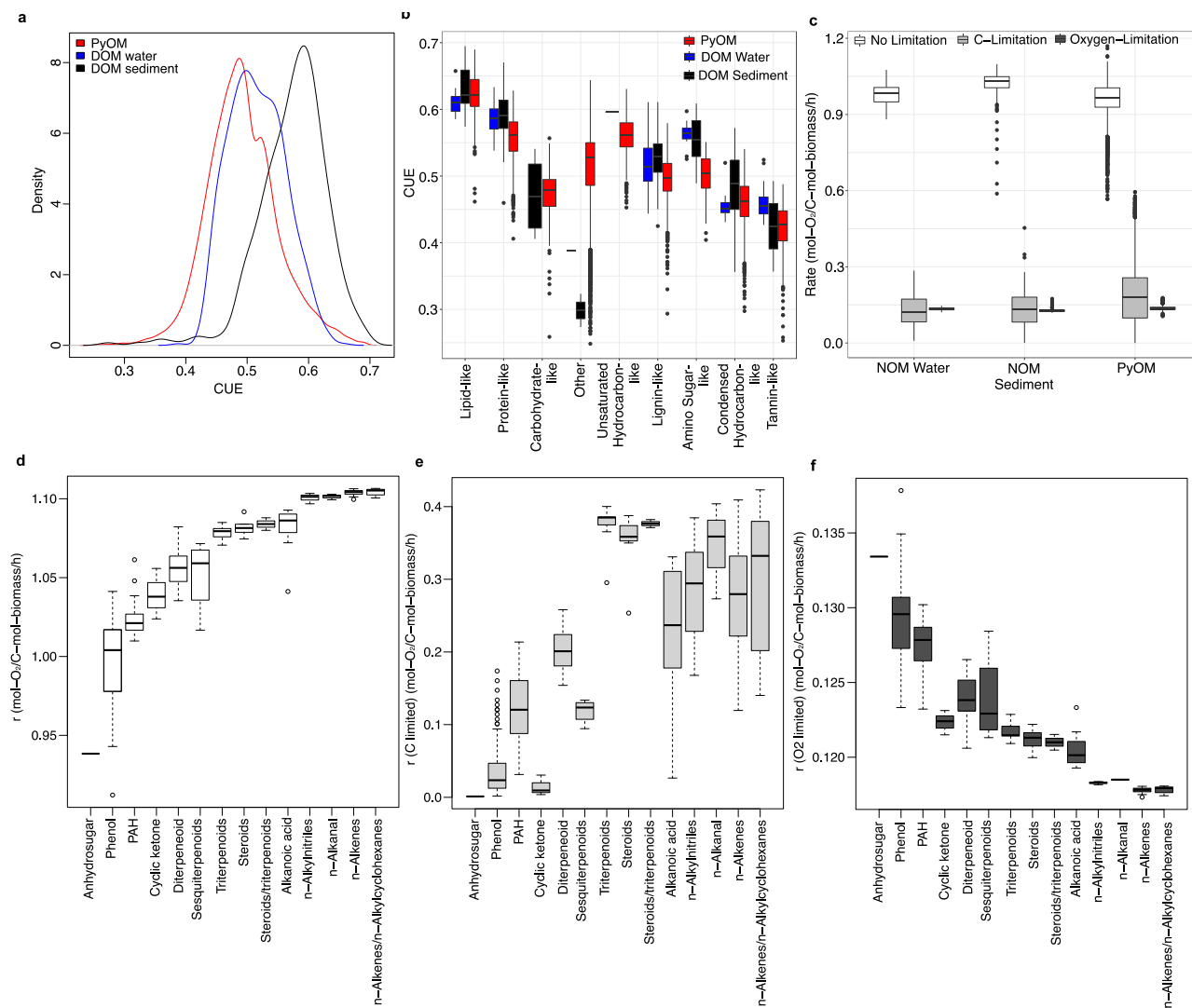
123 These results signal a strong overlap between the potential bioavailabilites of PyOM and
124 DOM pools; however, within PyOM compounds, there was variability in ΔG_{Cox} , λ , and CUE
125 consistent with heterogeneous chemistries (Figure 1 and 2). This is not surprising, given the
126 diversity of PyOM molecules generated by wildfires of different burn severities and source
127 materials (Wagner et al., 2015; Wagner et al., 2018; Neary et al., 2005), some of which are
128 chemically similar to natural DOM.

129



130

131 **Figure 1.** Comparison of natural DOM and PyOM. (a-c) Van Krevelen diagrams of molecules
 132 that were present in at least 95% of (a) surface water or (b) sediment samples versus (c)
 133 representative PyOM. (d-f) Distribution of energy content (ΔG_{CoX}) vs. metabolic efficiency (λ) in
 134 (d) surface water and (e) sediment DOM vs (f) representative PyOM. Higher values on the x-axis
 135 (ΔG_{CoX}) represent less energetically favorable molecules, while higher values on the y-axis (λ)
 136 represent higher rates of catabolism relative to anabolism. Colors for all panels indicate inferred
 137 chemical classes.



138

139 **Figure 2.** Predicted CUE and metabolism of DOM and PyOM. (a) shows the probability density
 140 function (PDF) of CUE in surface water DOM (blue), sediment (black) DOM, and PyOM (red).
 141 The PDF reflects the relative likelihood that value of a random sample drawn from a particular
 142 group would equal the value on the x-axis. (b) depicts CUE of surface water DOM, sediment
 143 DOM, and PyOM by inferred chemical class. (c) depicts the predicted metabolism of surface
 144 water DOM, sediment DOM, and PyOM under no elemental limitations (no fill), C-limitation
 145 (light grey fill), and oxygen-limitation (dark grey fill). (d-f) represent metabolic rates for PyOM
 146 molecules with known chemical classes under (d) no elemental limitations, (e) C-limitation, and
 147 (f) oxygen-limitation. For boxplots, median values are denoted by a bar, hinges correspond to the
 148 first and third quartiles, and whiskers extend to the largest/smallest value no further than 1.5
 149 times the inter-quartile range from the hinge. Data beyond the end of the whiskers are plotted
 150 individually.

151

152

153 Interestingly, the large overlaps in predicted CUE across PyOM and DOM pools suggests
154 that PyOM decomposition in rivers could emit proportionally similar amounts of CO₂ to natural
155 DOM (Figure 2a-b). CUE is used in many microbially-explicit decomposition models to
156 constrain organic matter bioavailability (reviewed in Graham and Hofmockel, 2022). Therefore,
157 predicted CUE offers a path for assimilating PyOM in microbially-explicit models. Such an
158 approach could be used to directly evaluate the impact of PyOM on global C cycles, and help
159 lead to better incorporation of PyOM impacts in models (Santin et al., 2020). We specifically
160 highlight lower CUE of amino sugar-like, protein-like, and lignin-like PyOM in comparison to
161 corresponding classes of natural DOM, pointing to potential shifts needed in the representation
162 of these classes in process-based models in ecosystems impacted by wildfires.

163

164 *2.2 Inferred Metabolism of Pyrogenic Organic Matter.*

165 Predicted rates of PyOM metabolism were also similar to DOM pools (Figure 2c),
166 reinforcing comparable bioavailability. Elemental limitations had a ~6-fold larger impact on
167 predicted metabolic rates than changes in compound type (e.g., surface water DOM, sediment
168 DOM, and PyOM). Metabolic rates averaged across all compound types were 0.965, 0.179, and
169 0.136 mol O/mol C biomass produced for unlimited, C-limited, and oxygen-limited conditions
170 respectively. In contrast when metabolic rates were averaged across elemental limitations within
171 each compound group, we found that surface water DOM, sediment DOM, and PyOM had
172 nearly identical mean predicted rates of metabolism (ANOVA $P = 0.14$ $F = 1.922$, all pairwise
173 comparisons $P > 0.10$). While metabolic rates within a given limitation scenario were statistically
174 different across compound types (ANOVA, all $P < 0.001$, $F = 197.4$, 145.9 , and 265.9), effect
175 sizes were low in comparison to changes across elemental limitations. Still, PyOM displayed

176 statistically elevated rates of metabolism under C- and oxygen-limitation versus both surface
177 water and sediment DOM (Tukey HSD, $P < 0.001$). Given that riverbed sediments can reach
178 anoxia within millimeters of the sediment-water interface, elevated rates of PyOM metabolism
179 under oxygen limitation may be ecologically relevant at ecosystem scales.

180 When considering the impact of elemental limitations on PyOM metabolism, predicted
181 rates were strongly inhibited under low C and oxygen conditions. Predicted PyOM metabolism
182 was over 5 times lower when C or oxygen was scarce. Low decomposition rates under C and
183 oxygen limitation could be one reason for the observed persistence of PyOM in depositional
184 features that tend to be anoxic. Still, it is worthwhile to note that metabolism of all PyOM classes
185 under low C or oxygen was predicted to be substantially slower than without elemental
186 limitations, indicating PyOM compounds may both actively cycle in well-oxygenated surface
187 waters with fresh C inputs and persist over long periods of time in O₂-limited sediments.

188 When examining PyOM with known chemistries (e.g., instead of inferred chemical
189 classes from FTICR-MS), anhydrosugars, phenols, and polycyclic aromatic hydrocarbons
190 (PAHs) were less negatively impacted by oxygen limitation than any other group (Figure 2 d-f).
191 Previous work has demonstrated that microorganisms are capable of decomposing chemically
192 complex organic molecules, such as long-chained and/or aromatic hydrocarbons under low
193 oxygen availability (Bushnell and Haas, 1941; Pozdnyakova, 2012; Rabus et al., 2016; Coates et
194 al., 1997). Similar microbial metabolic pathways may also be capable of degrading pyrogenically
195 modified phenols, and PAHs in natural settings and could be investigated with future laboratory
196 work.

197 Notably, our work also supports the notion that black nitrogen could be more bioavailable
198 than other PyOM pools. While we only examined one class of PyOM molecules containing

199 nitrogen in detail (n-alkylnitriles), it had among the highest predicted CUE and metabolic rates.
200 We posit this may be due, in part, to its chemical structure that includes pyrrole-type moieties,
201 which are relatively biodegradable (Knicker, 2010; De La Rosa and Knicker, 2011).

202

203 *2.3 Correspondence to Empirical Investigations.*

204 While the substrate-explicit modelling approach used here has been validated in natural
205 settings, we acknowledge that it does not account for all chemical and physical attributes
206 influencing PyOM bioavailability. This includes DOM structure and size, abiotic reactions, and
207 chemical complexation with minerals and particulates. Below we note aspects of the model
208 predictions that are inconsistent with experimental evidence and highlight the role of laboratory
209 studies in evaluating PyOM bioavailability.

210 For instance, n-alkenes and related compounds tended to have high modelled
211 bioavailability despite being relatively stable in the environment (Wiesenberg et al., 2004;
212 Smittenberg et al., 2004). These compounds are characterized by carbon-carbon double bonds,
213 which are not considered by the model and may decrease bioavailability. However, n-alkanes
214 generated through combustion tend to have reduced chain length in comparison to their unburned
215 counterparts (Knicker et al., 2013), and thus may be relatively bioavailable compared to
216 unburned n-alkanes. Additionally, we note that previous work has shown fast degradation of
217 combustion-derived lipids in soils (Knicker et al., 2013); as well as high n-alkene metabolism in
218 anaerobic sediments and high lipid reactivities at the sediment-water interface (Grossi et al.,
219 2008; Wilkes et al., 2016; Yongdong et al., 2015; Mbadanga et al., 2011; Canuel and Martens,
220 1996). While work on aerobic n-alkene metabolism is limited, the comparative bioavailability of

221 n-alkenes and known degradation pathways suggests that sediment microbiomes may metabolize
222 them as part of natural biogeochemical cycles.

223 Another notable discrepancy is the low potential bioavailability of anhydrosugars when
224 compared to other PyOM compounds. Experimentally, anhydrosugars are highly bioavailable in
225 oxic conditions, with a half-life of less than seven days (Norwood et al., 2013). The model may
226 therefore not adequately account for some enzyme-catalyzed reactions such as levoglucosan
227 kinase or levoglucosan dehydrogenase that may be common enzymes in aquatic microorganisms
228 (Bacik and Jarboe, 2016; Suciu et al., 2019).

229 Because of these nuances, the analysis presented here is best used as bounding estimates
230 for experimental validation and as a holistic comparison to DOM bioavailability. Still, the span
231 of compounds investigated here, and their comparison to DOM pools, provides a breadth of
232 investigation that is unfeasible without model-based approaches.

233

234 **3 Conclusions**

235 Our work supports the hypothesis that PyOM may have similar overall bioavailability as
236 compared to natural sources of DOM and provides a foundation for targeted experiments
237 investigating specific components of the PyOM continuum. Globally intensifying wildfires are
238 increasing the production of PyOM with potential implications for water supplies, which are
239 critical for domestic, industrial, agricultural, and ecological needs. Yet, many fundamental
240 questions such as “how much” PyOM exists in ecosystems, “how fast” it cycles, and “how old”
241 it is remain largely unknown (Abiven and Santin, 2019). Our work provides the first
242 comprehensive computational assessment of the potential bioavailabilities of various PyOM
243 chemistries in comparison to natural DOM pools. The comparable potential bioavailability to

244 DOM revealed that PyOM may be actively transformed in aquatic ecosystems and may be an
245 increasing source of C emissions to the atmosphere as the prevalence of wildfires increases.

246

247 **Code and Data Availability**

248 Code is available at: <https://github.com/hyunseobsong/lambda>. Data describing DOM pool
249 chemistry are published as a data package (Goldman et al., 2020) (available at:
250 doi:10.15485/1729719) and are discussed in more detail by Garayburu-Caruso et al. (2020a).

251

252 **Author Contributions**

253 EBG conceived of the manuscript and was responsible for writing the manuscript and generating
254 all figures. HSS performed all modelling. SG determined PyOM compounds for modelling based
255 on extensive literature review, with guidance from AMP. VGC and JCS contributed data and
256 insight on DOM pool chemistry. All authors contributed to revisions.

257

258 **Competing Interests**

259 The authors declare that they have no conflict of interest.

260

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269 Environmental Molecular Science Laboratory User Facility.

270

271 **APPENDIX. Materials and methods**

272 An extended version of our methods is available in the Supporting Information.

273 To assess the potential bioavailability of PyOM, we searched primary literature for
274 representative compounds of the PyOM continuum. Specifically, we targeted characteristic
275 organic compounds from controlled burns of various fuel types representing a range of moisture,
276 temperature, and oxygen conditions (Table S1). The chosen compounds focused on biomass
277 burning alteration products derived from plant charcoals, biochars, smoke, burnt soil and their
278 leachates, which are often used to characterize PyOM in the environment. This included
279 compounds such as theoretical black carbon-like compounds (defined here as condensed
280 aromatic core structures polysubstituted with O-containing functionalities (Wagner et al., 2017),
281 anhydrosugars, and polycyclic aromatic hydrocarbons (PAHs). The list also included compounds
282 created and/or transformed from biomass burning, such as those derived from biopolymers like
283 lignin (e.g., methoxyphenols), waxes (e.g., n-alkenes from thermal dehydration of n-alkanols),
284 and resins (e.g., thermally oxidized diterpenoids) (Oros and Simoneit, 2001b, a). We also include
285 molecular formula assigned via FTICR-MS spectra collected on biochar leachates to increase
286 data type comparability with common methods used in natural organic matter studies. The
287 selected set of compounds spans the chemical continuum of PyOM but is not intended to be
288 exhaustive. In total, our literature search for PyOM chemistries yielded 16,971 unique molecular

289 formulae. When a formula was found in more than one study or material type, this was noted in
290 Table S1, however only unique formulae were considered for the model.

291 After generating a set of representative compounds, we used a substrate-explicit
292 modelling framework developed by Song et al. (2020) to characterize the potential
293 bioavailability of each compound and predict its rate of decomposition. The model uses
294 molecular formulae to predict energetic content, metabolic efficiency, and rates of aerobic
295 metabolism, while it does not account for structural components of organic molecules (e.g.,
296 double bonds, folding patterns, cross-linkages). This enables flexibility in application to high-
297 throughput mass spectrometry techniques that yield chemical formulae but not structural
298 information (e.g., FTICR-MS) for comparison to environmental DOM. Despite its limitations,
299 the substrate-explicit model used here has proven useful in linking DOM composition to aerobic
300 metabolism in natural environments (Song et al., 2020; Graham et al., 2017; Garayburu-Caruso
301 et al., 2020b), and its structure is consistent with Harvey et al. (2016) who argued for the
302 importance of thermodynamic estimates of PyOM bioavailability that underlie this model. It was
303 chosen to allow for comparison of PyOM to the most comprehensive assessment of global
304 aquatic DOM pools to date (Garayburu-Caruso et al., 2020a).

305 Briefly, the substrate-explicit model uses the elemental stoichiometry of organic
306 molecules, based on molecular formulae, to predict the number of catabolic reactions that must
307 occur to provide the energy required for the synthesis of one mole of biomass carbon. This
308 quantity is described by the parameter lambda (λ) in which lower λ values denote more efficient
309 energetics of catabolism in producing biomass through anabolism. The model also predicts the
310 Gibbs free energy of C oxidation (ΔG_{Cox}), under standard conditions with a modification to pH 7
311 adjusted from LaRowe and Van Cappellen (2011) by Song et al. (2020), as well as C use

312 efficiency (CUE) as defined by Saifuddin et al. (2019). Lower ΔG_{Cox} denotes higher
313 thermodynamic favorability in an electron donor half reaction associated with organic matter,
314 and higher CUE reflects more C assimilated into biomass per unit C respired. We also predicted
315 the rate of aerobic metabolism (as oxygen consumed per mol-C biomass produced) under three
316 scenarios commonly observed in aquatic ecosystems: (a) C-limitation, (b) oxygen (O_2)
317 limitation, and (c) both C and O_2 -limitation. For more details of the substrate-explicit modelling
318 approach used, please see Song et al. (2020). Each metric (λ , ΔG_{Cox} , CUE, metabolic rates)
319 denotes a different aspect of potential bioavailability. Though the relative magnitude of the
320 metrics in comparison to each other will vary based on the specific stoichiometry of a molecule,
321 highly bioavailable compounds are indicated by low λ and ΔG_{Cox} coinciding with high CUE and
322 metabolic rates.

323 Three sets of organic molecules were used as model inputs: measured global dissolved
324 (1) surface water and (2) sediment DOM pools, extracted in H_2O and analyzed by FTICR-MS as
325 per Garayburu-Caruso et al. (2020a); and (3) literature-derived PyOM compounds as described
326 above. Inputs to the model from the PyOM compounds were unique molecular formulae,
327 grouped in subsequent analysis by their corresponding compound classes (Table S1). If one
328 molecular formula was represented by several PyOM compounds (e.g., $C_{10}H_{16}O_2$, which
329 corresponds to the sesquiterpenoid cis-Thujan-10-oic acid and 3-, 4- substituted methylcatechol
330 phenols), we assigned multiple compound classes to that molecular formula. Surface water and
331 sediment DOM pools were filtered to compounds occurring in 95% of samples to yield a dataset
332 of globally ubiquitous DOM (surface water $n = 811$ molecules, sediment $n = 398$ molecules).
333 The purpose of this filtering was to focus on widespread natural organic matter chemistries by
334 removing compounds that were specific to subsets of rivers. Formulae assignment and inferred

335 chemical classes via van Krevelen diagrams in DOM pools are described by Garayburu-Caruso
336 et al. (2020a). As a whole, 16,332 compounds were found only in PyOM, 197 were found only in
337 surface water DOM, and 167 were found only in sediment DOM. We compared modelling
338 outputs from representative PyOM to outputs of ubiquitous DOM pools to infer relative
339 bioavailability using ANOVA and Tukey HSD statistical tests with R software v 4.1.0 (R Core
340 Team, 2021). Visualizations were produced using base R and ‘ggplot2’ (Wickham, 2011). All
341 model outputs are available in Tables S2–S4.

342

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