

1 **Molecular fingerprinting of particulate organic matter as a new tool**
2 **for its source apportionment: changes along a headwater drainage in**
3 **coarse, medium and fine particles as a function of rainfalls**

4 Laurent Jeanneau^{1*}; Richard Rowland²; Shreeram Inamdar²

5 ¹ Univ rennes, CNRS, Geosciences Rennes - UMR 6118, F-35000 Rennes, France

6 ² Water Science & Policy Program, University of Delaware, Newark, USA

7 *Correspondence to:* Laurent Jeanneau (laurent.jeanneau@univ-rennes1.fr)

8 **Abstract.** Tracking the sources of particulate organic matter (POM) exported from catchments is important to understand the
9 transfer of energy from soils to oceans. The suitability of investigating the molecular composition of POM by thermally
10 assisted hydrolysis and methylation using tetramethylammonium hydroxide directly coupled to gas chromatography and
11 mass spectrometry is presented. The results of this molecular fingerprint approach were compared with previously published
12 elemental (%C, %N) and isotopic data ($\delta^{13}\text{C}$, $\delta^{15}\text{N}$) acquired in a nested headwater catchment in Piedmont region, Eastern
13 United States of America (12 and 79 ha). The concordance between these results highlights this molecular tool as a valuable
14 method for source fingerprinting of POM. It emphasizes litter as the main source of exported POM at the upstream location
15 ($80 \pm 14\%$) with an increasing proportion of stream bed (SBed) sediments remobilization downstream ($42 \pm 29\%$),
16 specifically during events characterized by high rainfall amounts. At the upstream location, the source of POM seems to be
17 controlled by the maximum and median hourly rainfall intensity. An added-value of this method is to directly investigate
18 chemical biomarkers and to mine their distributions in term of biogeochemical functioning of an ecosystem. In this
19 catchment, the distribution of plant-derived biomarkers characterizing lignin, cutin, and suberin inputs were similar in SBed
20 and litter, while the proportion of microbial markers was 4 times higher in SBed than in litter. These results indicate that
21 SBed OM was largely from plant litter that has been processed by the aquatic microbial community.

22 1 Introduction

23 Particulate organic matter (POM) plays key-roles in aquatic ecosystems, controlling the transfer and the bioavailability of
24 energy, nutrients and micropollutants. The flux of POM from soils to oceans has been estimated at 0.2 GtC per year (Ludwig
25 et al., 1996) with 80 % coming from biospheric inputs and the complement from petrogenic inputs (Galy et al., 2015).
26 Assuming that the energy provided by natural organic matter is equivalent of the energy provided by the combustion of
27 wood, this flux of POM corresponds to an energy of 2.8 EJ, that is to say less than 2 days of the global energy consumption
28 by the humanity in 2015 (yearbook.enerdata.net). This export mainly occurs during storm events, those hot moments being
29 responsible for up to 80% of annual particulate organic carbon (POC) export depending on the investigated catchment
30 (Dhillon and Inamdar, 2013; Jeong et al., 2012; Jung et al., 2012; Oeurng et al., 2011).

31 Among these hot moments, extreme events, defined as storm flow exceeded less than 10 % of the time (IPCC, 2001), seem
32 to play a dominant role. In two contrasted catchments, a mountainous one in South-Korea and a lowland one in the Eastern
33 United States of America (USA), the specific POC flux (flux per unit area of the catchment) has been shown to be non
34 linearly related to total rainfall with a threshold value beyond which the slope increased sharply (Dhillon and Inamdar, 2013;
35 Jung et al., 2014). The threshold value (approx. 70 mm in the American catchment and approx. 120 mm in the South-Korean
36 catchment) and the magnitude of this increase differed between both catchments and are probably watershed-dependant. Is
37 the non linearity of the relationship between rainfall amount and POC export observed previously linked to a modification of
38 the source of POM? POM in a river system is a combination of allochthonous and autochthonous OM. The former is derived
39 mainly from the soils and banks erosion, while the latter can be composed of fresh aquatic living organisms and bed
40 sediments. The balance between these different sources is controlled (i) by the catchment' size and morphology and (ii) by
41 the rainfall event characteristics (Tank et al., 2010).

42 Tracking the sources of POM can be done indirectly by investigating the sources of suspended matter. This can be done
43 through the analysis of fallout radionuclides such as Beryllium-7, Lead-210 and Cesium-137 (Ritchie et al., 1974; Wallbrink
44 and Murray, 1996; Walling, 1998) or by geochemical fingerprinting of rare elements (Collins and Walling, 2002). It can also
45 be done directly by investigating the composition of POM using bulk-scale descriptors such as OC and Nitrogen
46 concentrations, C/N ratio and stable isotopes $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ (Fox and Papanicolaou, 2008). Molecular biomarkers analyses
47 have also been used. They are based on specific molecular classes such as lipid or lignin biomarkers (Goñi et al., 2013; Jung
48 et al., 2015). Thermochemiolysis using tetramethylammonium hydroxide coupled to gas chromatography and mass
49 spectrometry has already been applied to the investigation of the fate of river DOM (Jeanneau et al., 2015) and POM
50 (Mannino and Harvey, 2000). This analytical technique is widely used to investigate the biogeochemistry of soil organic
51 matter (Derenne and Quénéa, 2015) and, coupled to a principal component analysis (PCA), it has been shown to be valuable
52 for forensic soils applications (Lee et al., 2012). An advantage of such an analysis is to generate a distribution of more than
53 hundred identified target compounds with small amount of particulate matter (from 5 to 10 mg) (Jeanneau et al., 2014),

54 giving a dataset rich enough to differentiate between sources (Walling, 2013). Here this analytical approach is combined with
55 a principal component analysis (PCA) to determine the main sources of POM as a function of the sediment size, the
56 catchment size and the rainfall characteristics.

57 The first objective of this paper is to test the suitability of molecular biomarkers derived from THM-GC-MS as a tool to
58 determine the sources of river POM. The second objective is to investigate how the sources of POM changed as a function of
59 the catchment size, particle size of the sediment, and the hydrological characteristics of the rainfall events. This study is
60 based on a subset of samples used to investigate the sources of POM exported during storm events using ^{13}C and ^{15}N as
61 tracers (Rowland et al., 2017). We hypothesized that molecular biomarkers provide important insights into sources of POM
62 and can be used as complimentary tracers for POM alongside or in addition to stable isotopes.

63 **2 Material and methods**

66 **1 Site description**

67 This study was conducted in a 79 ha watershed (second order stream) located in the Piedmont physiographic region of
68 Maryland, USA (Figure 1). The watershed drains into the Big Elk Creek which discharges into the Chesapeake Bay. For a
69 detailed description of the study site, refer to Rowland et al. (2017). Briefly, the watershed is predominantly forested with
70 pasture along the outer periphery. Dominant canopy species include *Fagus grandifolia* (American beech), *Liriodendron*
71 *tulipifera* (yellow poplar), and *Acer rubrum* (red maple). Bedrock formations consist of metamorphic gneiss and schist and
72 soils are coarse loamy, mixed, mesic lithic inceptisols on slopes and oxyaquic inceptisols in saturated valley bottoms.
73 Elevations in the watershed range from 77 to 108 m with slope gradients ranging from 0.16 to 24.5° (mean 6.3°). Mean
74 annual precipitation from 1981 to 2010 in this region was 1173.5 mm, with late spring and late summer as the wettest and
75 driest periods, respectively, and mean annual temperature is 13°C (Delaware State Climatologist Office Data Page, 2016).

79 **2.2 Watershed monitoring and sampling strategy**

80 Detailed information on monitoring and sampling is provided in Rowland et al. (2017). Climatological data was obtained
81 from a local station maintained by the Delaware Environmental Observing System approximately 450 m from the 79 ha
82 catchment outlet. This consists of temperature and GEONOR gage hourly rainfall measurements. Stream discharge estimates
83 were obtained at 20-minute intervals using a Parshall flume at 12 ha stream location (nested within the 79 ha watershed,
84 Figure 1) and a discharge rating curve calculated from paired pressure transducer and acoustic Doppler velocity meter
85 measurements at a rectangular concrete culvert at the 79 ha location.

86 Suspended sediments were collected using in-situ samplers made of 10 cm diameter capped PVC pipes placed vertically in
87 the middle of the stream. The upstream face of the pipes was perforated with 1.5 cm diameter holes beginning ~10 cm above

88 the stream bed. During periods of elevated discharge, stream stage rose above the perforations, trapping suspended sediment
89 within the sampler. The trapped sediment thus represented a time-integrated composite sediment sample (CSS). Such a
90 method induces modification of the velocity profile around the sampler, which could result in grain size fractionation. All
91 CSS were retrieved within 24 hours of the end of an event and frozen prior to processing and analysis. In this study POM
92 was defined by this sampler as the organic matter in the objects (natural debris, soil particles, colloids) that were trapped .
93 The slots on the samplers were approximately 1.5 cm which represents the higher threshold. The samples were dried before
94 further analysis and then included the smallest fractions defined as colloidal OM and dissolved OM.
95 Seven potential sediment sources were identified within the catchment and have been sampled at three locations to integrate
96 their spatial heterogeneity (Rowland et al., 2017). These included the stream bed (SBed), exposed stream bank A (BaA) and
97 B (BaB) horizons, valley-bottom wetland surficial soils (W), forest floor litter (Li) and humus (FH) and the upland A
98 horizons (Up). Sampling was conducted during the summer of 2015. 500-750 g of each end-member were sampled using an
99 ethanol-cleaned trowel or auger from both of the main tributary branches of the watershed. Stream beds were sampled from
100 areas without major backwatering or pooling, as POM may undergo diagenesis here, and were composited along a three by
101 three-point grid within the channel. Bank sediments were collected from exposed incised banks with three points composited
102 from the A and B horizons. Forest floor litter and humus, valley-bottom wetland soils and upland A horizons samples were
103 composited from five points along 20 m transects in low gradient locations in order to integrate their spatial heterogeneity.
104 End-member soil and sediment samples and CSS were dried in acid-cleaned Pyrex dishes in an oven at 45°C until visibly
105 dry. Oven-dry CSS samples were partitioned into coarse (CPOM) > 1000 µm, medium (MPOM) 1000-250 µm and fine
106 (FPOM) < 250 µm size classes via dry sieving. Dry masses were recorded for particle size class from which the fractional
107 mass percent was calculated for each class in each CSS sample. End-member samples were pre-sieved at 2 mm to remove
108 large organic debris such as roots. Aliquots were lyophilized overnight and preserved in a desiccator cabinet until elemental,
109 isotopic and molecular analyses. CSS and end-member samples were pulverized and homogenized using a ceramic mortar
110 and pestle that was cleaned with ethanol between samples.

111 **2.3 Analytical methodology**

112 For elemental and isotopic analyses, please refer to Rowland et al., (2017). The thermochemiolysis using
113 tetramethylammonium hydroxide (TMAH) coupled to gas chromatography and mass spectrometry (THM-GC-MS) was
114 performed according to Jeanneau et al. (2014). Briefly we introduced approximately 5 mg of freeze-dried solid residue into
115 an 80 µL aluminum reactor with an excess of solid TMAH (ca. 10 mg) and 10 µl of a solution of dihydrocinnamic acid d9
116 (CDN Isotopes, ref. D5666) diluted at 25 µg/ml in methanol as an internal standard. The THM reaction was performed on-
117 line using a vertical micro-furnace pyrolyser PZ-2020D (Frontier Laboratories, Japan) operating at 400°C. The products of
118 this reaction were injected into a gas chromatograph (GC) GC-2010 (Shimadzu, Japan) equipped with a SLB 5MS capillary
119 column in the split mode (60 m × 0.25 mm ID, 0.25 µm film thickness). The temperature of the transfer line was 321°C and
120 the temperature of the injection port was 310°C. The oven was programmed to maintain an initial temperature of 50°C for 2

121 minutes, then rise to 150°C at 15°C min⁻¹, and then rise to 310°C at 3 °C min⁻¹ where it stayed for 14 minutes. Helium was
122 used as the carrier gas, with a flow rate of 1.0 ml/min. Compounds were detected using a QP2010+ mass spectrometer (MS)
123 (Shimadzu, Japan) operating in the full scan mode. The temperature of the transfer line was set at 280°C, the ionization
124 source at 200°C, and molecules were ionized by electron impact using an energy of 70 eV. The list of analyzed compounds
125 and m/z ratios used for their integration are given in the supplementary materials (Table S1). Compounds were identified on
126 the basis of their full-scan mass spectra by comparison with the NIST library and with published data (Nierop et al., 2005;
127 Nierop and Verstraten, 2004). They were quantified assuming similar ionization and detection efficiencies between all
128 compounds. This assumption means that the concentrations must be handled as rough estimations.

129 Target compounds were classified into four categories: low molecular weight organic acids, phenolic compounds including
130 lignin and tannin markers, carbohydrates and fatty acids. The peak area of the selected m/z (mass/charge) for each compound
131 was integrated and corrected by a mass spectra factor calculated as the reciprocal of the proportion of the fragment used for
132 the integration and the entire fragmentogram provided by the NIST library (Table S1). The proportion of each compound
133 class was calculated by dividing the sum of the areas of the compounds in this class by the sum of the peak areas of all
134 analyzed compounds expressed as a percentage. The analytical uncertainty for this analytical method, expressed as a relative
135 standard deviation ranged from 10 to 20% depending on the samples and the target compounds. The use of THM-GC-MS to
136 investigate the sources of POM meant that it was necessary to assume that matrix effects are equivalent for all analyzed
137 compounds in all samples.

138 **2.4 Statistical analyses and calculation of the proportions of the main sources of POM in CSS**

139 Statistical analyses were performed using XLSTAT (version 19.01, Addinsoft). First a principal component analysis (PCA)
140 was performed using the end-members as individuals and CSS as additional individuals. The relative proportions of the 112
141 target compounds and the sum of their concentrations in ng/mg of freeze-dried matrix were used as variables. The relative
142 distribution of target compounds allows the direct comparison of the different samples without concentration effect, while
143 using the sum of their concentrations takes into consideration the fact that the concentration of target compounds differed
144 from a sample to another.

145 The first PCA allows identifying the correlated variables on the basis of a modulus of the Pearson coefficient > 0.9. When
146 two variables were correlated, the least abundant was removed. Then a second PCA was performed. The variables with a
147 correlation lower than 0.4 with the two first factors (F1: 29.8%; F2: 17.2% of variance) were removed, resulting in a new set
148 of 71 variables. A third PCA was calculated and a hierarchical ascendant classification (HAC) was calculated using the
149 coordinates of the individuals (end-members and CSS) on the 9 first factors that explained 90.5% of the variance of the
150 dataset. This HAC identified Upland soils and Stream bank sediments as minor contributors. Consequently a fourth PCA was
151 calculated removing Upland soils and Stream bank sediments from the potential end-members. Similarly to the three
152 previous PCA, CSS were considered as additional individuals. The coordinates of CSS on the two first factors (on 10) of this
153 PCA (F1: 40.1%; F2: 24.0% of variance) were used to calculate the proportion of the three main sources of POM in CSS

154 identified as 1. stream bed sediments, 2. litter and 3. forest floor humus + wetland soil, resolving a system of equations with
155 three unknowns. To solve this system, the coordinates of end-members must be specified. The heterogeneity of the
156 distribution of target compounds resulted in an area for each end-member. To calculate the proportions and uncertainties, the
157 coordinates of end-members were randomly selected ten times in the areas defined by the 95% IC. When the calculation
158 gave a negative contribution for an end-member, it was set at 0 and the two others contributions were recalculated to sum at
159 100. Finally the contributions of those three sources were approximated for the bulk POM by using the proportion and the
160 OC content of each fraction. From the third PCA to the end of the procedure, this treatment was also performed adding TOC,
161 $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ from Rowland et al. (2017) as variables.

162 In order to test the efficiency of the source apportionment calculated with the molecular data, the proportions of end
163 members and their isotopic values (Rowland et al., 2017) were used in an end-member mixing approach to model the $\delta^{13}\text{C}$ of
164 CSS. Modeled values were compared to measured values reported by Rowland et al. (2017) by calculating the relative
165 standard deviation (RSD) and against a linear regression model.

166 **3 Results**

168 **3.1 Rainfall and hydrology**

169 The molecular composition of POM in coarse, medium and fine size classes was investigated for four events. The rainfall
170 and discharge characteristics recorded for those events are indicated in Table 1. The total rainfall ranged from 40.1 (E4) to
171 148.9 (E1) mm, the maximum hourly rainfall (I_{max}) ranged from 19.9 (E1) to 31.3 (E3) mm h^{-1} and the median hourly
172 rainfall (I_{med}) ranged from 0.4 (E3) to 2.2 (E2) mm h^{-1} . The maximum discharge for those events ranged from 15.6 (E4) to
173 150.1 (E1) l s^{-1} . Then the four events can be distinguished as follows. E1 was characterized by high rainfall, a low maximum
174 intensity (I_{max}), an intermediate median intensity (I_{med}) and an intermediate antecedent precipitation index (API7). E2 was
175 characterized by mean total rainfall, a mean I_{max} , a high I_{med} and a mean API7. E3 was characterized by high rainfall and
176 I_{max} , low I_{med} and high API7. Finally E4 was characterized by low rainfall and I_{max} , a high I_{med} and a dry antecedent
177 conditions (API7 = 0 mm). E2 and E4 were comparable in terms of precipitation regime but can be differentiated by the
178 API7, E4 occurring after 7 days without precipitation.

183 **3.2 Size distribution**

184 CSS were separated into coarse (>1 mm), medium (>250 μm) and fine (<250 μm) fractions, with the exception of CSS at the
185 downstream (79 ha) location for the fourth event (Table 1). In the 12 ha sub-catchment, the coarse, medium and fine
186 fractions represented 22 ± 20 , 22 ± 4 and 55 ± 21 % of particulate matter, respectively, while in the 79 ha catchment, they
187 represented 61 ± 19 , 22 ± 10 and 18 ± 10 % of particulate matter, respectively. In the 12 ha sub-catchment, the relative

188 standard deviation (RSD) of those proportions was 90, 17 and 37 % for the coarse, medium and fine fractions, respectively,
189 while in the 79 ha catchment it was 31, 45 and 55 %, respectively.

190 3.3 Molecular composition of end-members

191 The number of detected target compounds ranged from 49 (SBed#1) to 112 (FH). A Dixon test for extreme value identified
192 the lowest value (SBed#1) as an outlier (p -value = 0.011). Once this value removed, the number of detected target
193 compounds ranged from 75 (BaB) to 112 (FH). The low value recorded for one of the SBed could be due to a combination of
194 a low OC content with a low analytical efficiency. This sample was removed from the dataset.

195 The distribution of target compounds into chemical families gives a first overview of the molecular composition of OM in
196 the different end-members (Figure 2). In W, Li and FH, the main compounds are phenolic compounds and high molecular
197 weight fatty acids ($> C_{20}$, HMW) that represent more than 30% of target compounds. In BaA and BaB, the proportion of
198 phenolic compounds was lower (22 ± 4 and 19 ± 1 %, respectively; mean \pm SD) than in W, Li and FH and the proportion of
199 low molecular weight ($< C_{20}$, LMW) fatty acids was higher (27 ± 17 and 35 ± 9 %, respectively). In Up, compared to W, Li
200 and FH, the proportion of HMW fatty acids increased (57 ± 19 %), while the proportion of phenolic compounds decreased
201 (13 ± 8 %). In SBed, the main identified target compounds were LMW fatty acids (72 ± 8 %), while phenolic compounds
202 and HMW fatty acids represented 15 ± 2 % and 9 ± 4 %, respectively.

203 HMW fatty acids was composed of linear n -alkanoic acids from n -C_{20:0} to n -C_{32:0} with an even-over-odd predominance
204 characteristic of plant-derived inputs (Eglinton and Hamilton, 1967), linear ω -hydroxyacids and α,ω -diacids from n -C₁₆ to n -
205 C₂₈, 10,16-dihydroxyC_{16:0} and 9,10,18-trihydroxyC_{18:0} characteristic of plant-derived aliphatic biopolymers cutin and suberin
206 (Armas-Herrera et al., 2016; Kolattukudy, 2001). These two latter hydroxyacids were the main compounds among HMW
207 fatty acids. The proportion of ω -hydroxyacids and α,ω -diacids among HMW fatty acids is higher in roots than in leaves and
208 can be used to differentiate between suberin from roots and cutin from shoots (Mueller et al., 2012). This proportion
209 decreased from soils (Up, FH and W) and bank sediments to litter and was minimal for SBed (17 ± 8 %), highlighting that
210 the proportion of cutin decreased from SBed, Li to bank sediments and soils.

211 Phenolic compounds included of methoxy-benzene, -acetophenone, -benzaldehyde and -benzoic acids. These compounds
212 derived from lignin and tannins and are characteristic of plant-derived OM. The main compounds were guaiacyl-like
213 structures: 3,4-dimethoxybenzaldehyde, 3,4-dimethoxybenzoic acid methyl ester, *erythro* and *threo*-1,2-dimethoxy-4-(1,2,3-
214 trimethoxypropyl)benzene and syringil-like structures: 3,4,5-trimethoxybenzaldehyde and 3,4,5-trimethoxybenzoic acid
215 methyl ester, which is typical of the THM-GC-MS of OM deriving from woody plants (Challinor, 1995). Benzoic acid was
216 not classified in this chemical family since it was negatively (slope of the linear regression model: -0.20; -0.18; -0.17) and
217 poorly correlated (Pearson coefficient, p -value: 0.14, 0.002; 0.14, 0.002; 0.21, <0.001) with 3,4-dimethoxybenzoic acid
218 methyl ester, 3,4,5-trimethoxybenzoic acid methyl ester and 3-(3,4-dimethoxyphenyl)prop-2-enoic acid methyl ester,

219 respectively, that are the main representatives of the three types of lignin units analyzed by THM-GC-MS (Challinor, 1995).
220 As a consequence, it was not considered to calculate the proportion of molecules coming from lignins and tannins.
221 LMW acids included *n*-alkanoic acids from *n*-C_{6:0} to *n*-C_{19:0}, *iso* and *anteiso* C_{13:0}, C_{15:0} and C_{17:0}, *iso* C_{14:0} and C_{16:0} and *n*-
222 alkenoic acids *n*-C_{16:1} and *n*-C_{18:1}. The LMW fatty acids with less than 13 C atoms can derive from microbial or plant-derived
223 inputs, while the LMW fatty acids with more than 13 C atoms are known as phospholipid fatty acids and are microbial
224 biomarkers (Frostegård et al., 1993) with the exception of *n*-C_{16:0} and *n*-C_{18:0} that can derive from plant-derived inputs. The
225 proportion of microbial markers among target compounds was calculated according to Jeanneau et al. (2014). It increased
226 from litter and soils (<15%) to bank sediments (18 ± 12 % and 25 ± 7 % in BaA and BaB, respectively) to SBed (48 ± 15 %).

227 3.4 Molecular composition of stream suspended sediments

228 The distribution of target compounds into the five chemical families previously described changed with the catchment size as
229 illustrated on Figure 3. At the 12 ha location, this distribution was fairly homogenous across the particle classes. When
230 averaged across size fractions and events, the THM-GC-MS of the POM of CSS sampled at the 12 ha location mainly
231 produced phenolic compounds (48 ± 6 %, mean ± SD) and HMW fatty acids (22 ± 10 %). The relative standard deviation
232 weighted by the proportion (RSDp) was 13, 14 and 22 % for C, M and F fractions, respectively, which highlights a low inter-
233 event variability of this distribution. At the 79 ha location, the distribution of target compounds was dominated by LMW
234 fatty acids (41 ± 20 %) and phenolic compounds (37 ± 9 %). It was almost stable between the three size fractions with a
235 higher proportion of LMW fatty acids in the M fraction. However, the RSDp was 50, 55 and 23 % for C, M and F fractions,
236 respectively, which means a higher inter-event variability than at the 12 ha location.

244 3.5 End-members contributions

245 A hierarchical ascendant classification (HAC) was performed using the coordinates of end-members and stream sediments
246 (CSS) on the nine first factors (90.5 % of variance) of the PCA, which were calculated with the relative proportions of target
247 compounds and the sum of their concentrations as variables. Three classes were isolated. The first one included the three Li,
248 one FH and one W as end-members, the size fractions of CSS from the 12 ha location and 3 size fractions of CSS from the
249 79 ha location. The second group included two W, two FH and the three BaA, BaB and U end-members. Finally the third
250 group included the SBed end-members and the size fractions of CSS from the 79 ha location. Based on this HAC, Up, BaA
251 and BaB were considered as minor contributors to the POM exported from the 12 ha and 79 ha locations.

252 An additional PCA was then calculated using SBed, Li, FH and W as individuals, CSS as additional individuals, and the
253 previously defined list of 71 variables. The two first factors of this PCA explained 64.1 % of the variance of this final
254 dataset. The projection of end-members and CSS on the plan obtained with these two factors is illustrated on Figure 4. This
255 projection allows differentiating: (i) the three groups of end-members, Li, SBed and a combination of FH and W, denoted
256 FH-W and (ii) POM from the two sampling locations. Moreover the size classes were also separated. From this 2D
257 projection, an area was defined for each end-member corresponding to the 95% confidence interval. The results of the source

258 apportionment calculated using this 2D projection are listed in Table 2. Some CSS plotted outside the triangle formed by
259 end-members most probably because (1) the litter end-member did not capture the full compositional diversity of the
260 catchment and (2) end-member composition was investigated on bulk samples.

261 At the 12ha location, as an average of the four sampled events, from FPOM to CPOM, the proportion of OM coming from
262 SBed decreased from 17 ± 16 % (mean \pm SD) to 1 ± 1 %, the proportion of OM coming from FH-W decreased from 16 ± 16
263 % to 8 ± 12 % and the proportion of OM coming from Li increased from 67 ± 7 % to 90 ± 11 %. The large uncertainties
264 quantified by the mean RSD (78 ± 53 %, mean \pm SD, $n = 9$) reflected the inter-storm variability of this source
265 apportionment. Bulk POM was mainly inherited from Li with contributions ranging from 65 to 92 %.

266 At the 79ha location, as an average of the four sampled events, CPOM was mainly inherited from Li (63 ± 28 %) and SBed
267 (36 ± 30 %). MPOM was mainly due to SBed inputs (49 ± 39 %) and received a substantial contribution of FH-W (17 ± 31
268 %). Similarly to CPOM, FPOM was mainly inherited from Li (55 ± 15 %) and SBed (38 ± 24 %). Similarly to the source
269 apportionment at the 12ha location, the large uncertainties (RSD = 97 ± 57 %, $n = 9$) were due to inter-storm variability.
270 Bulk POM was mainly inherited from Li with contributions ranging from 42 to 89 % and SBed with contributions ranging
271 from 8 to 57 %.

272 4 Discussions

273 4.1 What are the main sources of POM for the watershed?

274 The HAC identified four main end-members for the stream water POM: litter (Li), the surface horizon of forest soils (FH)
275 and wetland soils (W) and stream bed sediments (SBed). Li was the main source of POM identified along the catchment
276 representing 80 ± 14 % and 49 ± 24 % of the POM exported from the 12 ha and 79 ha catchments, respectively. These high
277 proportions of Li-derived POM is in accordance with the results of Jung et al. (2015) where isotopic and *n*-alkanes
278 fingerprints of POM exported from a mountainous forested headwater catchment highlighted similarities with litter and
279 surface soils. Moreover the decrease in the proportion of Li-derived OM along the catchment fits well with the observation
280 of Koiter et al. (2013) where the contribution of topsoil sources of suspended sediments decreased from 75 to 30 % when
281 moving downstream.

282 Stream bank A and B horizons and the surface horizons of upland soils did not group with any CSS, which would mean that
283 they were minor contributors for the investigated samples. This seems to be in contradiction with the documented impact of
284 bank erosion on the mobilization of particulate organic matter (Adams et al., 2015; Nosrati et al., 2011; Tamooh et al., 2012).
285 This apparent contradiction could be due to the catchment's size. Contrary to the previously cited investigations (Adams et
286 al., 2015; Nosrati et al., 2011; Tamooh et al., 2012), this present study focused on a headwater catchment (0.79 km²). In these
287 small catchments, POM mainly comes from the erosion of surrounding soils as observed for monsoon floods in Laos
288 (Gourdin et al., 2015; Huon et al., 2017) or from a combination of bedrock and surface erosion in an Alpine catchment with

289 relative proportions controlled by the precipitations (Smith et al., 2013). However, in this catchment, the mobilization of
290 stream banks has been shown to be effective in winter due to freeze-thaw process (Inamdar et al., 2017). This present study
291 analyzed four events sampled in spring and summer. The lower contribution of stream bank erosion could then be due to
292 seasonal variability.

293 The relative proportion of phenolic compounds compared to HMW fatty acids plotted against the proportion of α,ω -diacids
294 and ω -hydroxyacids with more than 20 C atoms among HMW fatty acids resulted in a visual differentiation of Li and SBed
295 from wetland (W), forest humus (FH), River bank horizons A (BaA) and B (BaB) and from Upland soil (Up) (Figure 5). This
296 observation highlights Li as the main origin of SBed plant-derived OM, which fits well with the high proportion of Li-
297 derived POM in CSS from both catchments. Moreover from Li to SBed, (i) the ratio of coumaric and ferulic acids to
298 vanillaldehyde, acetovanillone and vanillic acid, commonly noted C/V, decreased from 0.79 ± 0.26 to 0.20 ± 0.07 , denoted
299 that lignins were more biodegraded in SBed than in Li and (ii) the proportion of microbial markers among the target
300 compounds increased from 12 ± 5 to 48 ± 15 %. Both of these observations highlight the recycling of terrestrial plant-
301 derived OM in river sediments from a headwater catchment, and are in accordance with the higher mineralization rate of soil
302 organic carbon in river sediments (Wang et al., 2014).

308 **4.2 Are molecular data in accordance with isotopic and elemental data?**

309 A four-step analysis was performed to determine if the molecular data produced by THM-GC-MS were in accordance with
310 the isotopic results (Rowland et al., 2017) previously acquired on those samples.

311 The first one consists in a point-by-point comparison of the source apportionments resulting from the two approaches. Four
312 main observations were reported by Rowland et al. (2017) using the isotopic approach. First, “the litter layer was a dominant
313 contributor to CPOM, especially for the upstream locations”. This is in agreement with our data: the proportion of Li-derived
314 CPOM was 90 ± 11 % and 63 ± 28 % for the 12ha and the 79ha catchments, respectively. Secondly, “the proportional
315 contributions of SBed and banks to MPOM and FPOM increased downstream”. This is also in agreement with molecular
316 data, however stream banks were not considered as a main contributor through the present statistical treatment. The
317 proportion of SBed-derived POM increased from 8 ± 8 % to 49 ± 39 % and from 17 ± 16 % to 38 ± 24 % between the 12 ha
318 and the 79 ha catchments in MPOM and FPOM, respectively. Thirdly, “no appreciable shift was observed in CPOM source”.
319 This is partly in agreement with the molecular data. The main contributor to CPOM was Li in the two locations but the
320 proportion of SBed-derived CPOM increased downstream. Finally, the highest contribution of forest floor humus was
321 observed in MPOM and FPOM for E4. This is in agreement with the source apportionment in this study since the proportion
322 of FH-W-derived POM was the highest for this event in CPOM, MPOM and FPOM from the 12 ha catchment and in MPOM
323 and FPOM from the 79 ha catchment.

324 In a second step, the quality of the source apportionment calculated from the end member mixing approach was investigated
325 by modeling the $\delta^{13}\text{C}$ of the samples using the isotopic fingerprint of end members. These modeled values were compared to

326 the measured values used in the isotopic fingerprinting approach (Rowland et al., 2017). The relative standard deviation was
327 1.1 ± 0.2 % (mean \pm 95% CI; $n = 20$) and the linear regression resulted in a slope of 1.01 ($R^2 = 0.58$; p -value < 0.0001 ;
328 Figure S1) highlighting a fairly good agreement between the model and the data, that is to say between the source
329 apportionment using molecular data and measured $\delta^{13}\text{C}$.

330 In a third step, TOC, $\delta^{13}\text{C}$, $\delta^{15}\text{N}$ and C/N were added as variables in the PCA treatment. In a first PCA, W, FH, Li, SBed,
331 BaA, BaB and Up were considered as potential end members. A HCA using the nine first PCA factors (90.4 % of the
332 variance) highlighted BaA, BaB and Up as minor contributors, similarly to this step performed on molecular data alone.
333 Then a second PCA was calculated with FH, W, Li and SBed as potential end members and the CSS as additional
334 individuals. The two first factors represented 64.4 % of the variance and resulted in a clear differentiation between Li, SBed
335 and FH-W. The same approach was then applied using the molecular data alone, resulting in the calculation of the
336 proportions of those three end members in the CSS for ten different combinations of the position of end members in the 2D
337 plan created by the two first factors of the PCA. For each CSS sample a set of ten values was created for Li-, SBed- and FH-
338 W-derived POM (Table S2). Student T-test was used to compare these distributions between the modality “molecular data”
339 and the modality “molecular + isotopic, elemental data”. A p -value was calculated for each sample. They ranged from 0.08 to
340 0.49 (0.25 ± 0.03 ; mean \pm 95% CI), highlighting that there were no significant differences between the two approaches
341 (Table S3).

342 The final step aimed at investigating to what extent the molecular data are representative of bulk POM. The linear regression
343 between the sum of the concentrations of target compounds (expressed in $\mu\text{g/g}$ of dry solid) and the total organic content
344 (expressed in % of dry solid) resulted in a correlation coefficient of 0.94 (p -value < 0.0001 ; Figure S2). This correlation
345 between bulk scale and molecular analyses has already been highlighted for sedimentary and dissolved OM (Jeanneau and
346 Faure, 2010; Jeanneau et al., 2014) and emphasizes the suitability of molecular investigations to determine the sources of
347 OM.

348 Once validated by this four-step comparison, what are the insights provided by the molecular approach on the source
349 apportionment of CPOM, MPOM and FPOM along this Piedmont headwater catchment?

350 **4.3 Modification of the source apportionment as a function of rainfall parameters**

351 These present results may be valuable to investigate the relationships between the sources of exported POM and rainfall
352 characteristics. However they have been acquired on only four events and this part of the discussion should be enriched by
353 future investigations.

354 Rainfall is the primary driver for C export since it controls soil erosion and stream discharge (Raymond and Oh, 2007).
355 Rainfall amount and API7 have been shown to control the export of POC from headwater catchments (Dhillon and Inamdar,
356 2013, 2014; Jung et al., 2014). Moreover Imax and Imed have also been identified as important drivers for soil erosion since

357 they control the rainfall erosivity (Wischmeier, 1959). The four investigated events represented a range of rainfall amounts,
358 maximal hourly intensity (I_{max}), median hourly intensity (I_{med}) and antecedent precipitation index (API7).
359 Linear regression were performed between the proportions of Li-, SBed- and FH-W-derived POM in CPOM, MPOM and
360 FPOM from both catchments against rainfall amount, I_{max} , I_{med} and API7 (Table 1). With only four investigated events,
361 only relationships characterized by Pearson coefficient higher than 0.8 were considered. p -Values were not calculated for
362 those regressions since they would not have had any statistical value. With only four events the highlighted relationships
363 must be handled with care and may be seen as guidelines for future works.

364 In the 12 ha catchment, SBed-derived OM was positively related to I_{max} and API7 and negatively related to I_{med} . The
365 positive relationship with API7 was recorded in C and F fractions, while the positive relationship with I_{max} and the negative
366 relationship with I_{med} were recorded only in the F fraction. In the M fraction, SBed-derived OM was related to the total
367 rainfall. However since this fraction represented 22 ± 4 % (mean \pm SD) of the exported particles, this relationship was not
368 considered as representative. In the 12 ha catchment the export of SBed-derived OM would be favored by rainfall
369 characterized by high I_{max} occurring after a period of dryness (Figure 6a). Moreover the proportion of FH-W-derived OM
370 was positively related to I_{med} in F fraction. This fraction represented 55 ± 21 % (mean \pm SD) of the exported particles,
371 giving some representativity to this observation. A deeper analysis of the relationship between I_{med} and the proportion of
372 FH-W-derived OM in the different fractions from the 12 ha catchment highlights a concomitant control of API7 (Figure 6b).
373 For similar I_{med} (E2 versus E4), the proportion of FH-W-derived OM increased in the three fraction with dry antecedent
374 conditions. The activation of the soil reservoir seems to be controlled by both I_{med} and API7, which could be interpreted as
375 the necessity of a dry period to replenish a stock of soil OM available for soil erosion and that intensive and regular rainfalls
376 could result in higher soil erosion.

377 In the 79 ha catchment, the proportions of Li and FH-W were negatively related to the rainfall amount and the proportion of
378 SBed was positively related to this variable. These relationships were recorded in the C and M fractions, with the exception
379 of FH-W (only in the C fraction). A deeper analysis of the link between the POM source apportionment and the rainfall
380 amount highlights different threshold values for C, M and F fractions (Figure 6c). In M and F fractions, there was a sharp
381 modification of the source of POM between E4 (40.1 mm) and E2 (43.9 mm). The proportion of FH-W-derived POM
382 decreased from 64 ± 20 % to 0 ± 1 % and from 21 ± 22 % to 1 ± 2 %, in the M and F fractions, respectively. These decreases
383 were concomitant with increases in the proportion of SBed-derived POM from 0 ± 0 % to 43 ± 8 % and from 2 ± 2 % to $48 \pm$
384 9 %, in the M and F fractions, respectively. The source apportionment of FPOM remained unchanged by further increases of
385 the rainfall amount, while for MPOM the source apportionment was clearly modified during E1, which was characterized by
386 the highest rainfall amount (148.9 mm). The proportion of Li-derived POM decreased to 0 ± 1 % and the proportion of
387 SBed-derived POM increased from 58 ± 9 % to 95 ± 7 %. The source apportionment of CPOM drastically changed between
388 E2 and E3 (97.4 mm). The proportion of Li-derived POM decreased from 95 ± 8 % to 47 ± 9 % and the proportion of SBed-
389 derived POM increased from 2 ± 4 % to 53 ± 9 %. This source apportionment remained unchanged between E3 and E1.
390 Since the C fraction was the most important during events 1, 2 and 3, its source apportionment was an important driver of the

391 source of total POM. It was mainly modified between events 2 and 3 with a decrease in the proportion of Li-derived POM
392 and an increase in the proportion of SBed-derived POM. From these observations, the threshold value of 75 mm previously
393 found in this catchment with an increase in the slope of the POC exported in kg/ha as a function of the rainfall amount
394 (Dhillon and Inamdar, 2013) falls in the range from 43.9 mm (E2) to 97.4 mm (E3), where the main modifications of the
395 source of POM exported from the 79 ha catchment were observed. The increase in the proportion of SBed-derived POM
396 accompanied with the increase in the proportion of the C fraction could be the result of the exceeding of a threshold value of
397 the hydrodynamism for sediment remobilization.

398 **4.4 Benefits and limitations of this molecular fingerprinting approach**

399 The present molecular fingerprinting method has benefits and limitations. Among the benefits, when the analysis is
400 performed on-line, that is to say, when the products of the THM are directly sent to the GC, then the analysis needs low
401 sample mass, in the order of 5 to 10 mg. Then this method is based on the molecular composition of OM, which is perfectly
402 suitable to investigate the fate of POM. Moreover it takes advantage of the differences of chemical composition between
403 living organisms (microorganisms versus plants) and in their different parts (leaves versus roots). As a consequence the
404 recorded modifications can be discussed in term of biogeochemistry of POM.

405 However limitations must be considered. Seasonal variability of the molecular fingerprint could exist especially for quickly
406 reactive reservoir such as litter (Williams et al., 2016). In soils, the turnover of OM takes time (> 50 years; Frank et al.,
407 2012). Consequently their molecular fingerprints may be less sensitive to seasonal variations, with the exception of
408 agricultural soils subject to changes in vegetation cover. This limitation can be easily avoided by sampling the most reactive
409 end-members at different seasons. The second and third limitations come from the method itself. First this is a time-
410 consuming method because each compound must be determined with care in each sample. For an analysis, approximately
411 two hours are necessary. Finally, because it is not only a value given by an analytical tool, using it asks having an expertise
412 in organic geochemistry.

413 When benefits and limitations are well considered, this molecular fingerprinting approach may be particularly suitable to
414 investigate the sources of POM in combination with other fingerprinting approaches.

415 **5 Conclusion**

416 This study emphasizes the suitability of molecular analysis of POM using THM-GC-MS to investigate the sources of POM
417 in headwater catchments. This analytical technique needs less than 5 mg of freeze-dried matter, which makes it realistic in
418 regard of the amount of suspended sediment exported and simple with only freeze-drying as a preparing step. With
419 approximately hundred of target compounds, the provided chemical fingerprint allows for the differentiation of the main
420 sources of exported POM, specifically between litter, surface soils, and in-channel sediments. The fairly good relationships
421 obtained by comparison with the conclusions gained by the isotopic-elemental investigation provide additional evidence in

422 favor of this organic fingerprinting approach. The present data highlight plant litter as the main source of exported POM with
423 an increasing contribution of stream bed sediments downstream. This latter contribution seems to be controlled by the
424 rainfall amount with a threshold phenomenon already observed for quantitative data. The contribution of soil erosion could
425 be controlled by both the median intensity of rainfall and the amount of rain in the previous 7 days. The investigation of
426 additional events in different catchments will be necessary to determine if those results are generic.

427 **Data availability**

428 Data are available on request from the corresponding author.

429 **Acknowledgements**

430 This study was funded by NSF ESPCoR Grant # IIA 1330238 (NEWRnet) and USDA NIFA Grant # 2015-67020-23585. We
431 would like to thank the Fair Hill Natural Resources Management Area for allowing us to conduct this study in the Fair Hill
432 Nature Preserve. Many thanks to students who assisted with sampling including Erin Johnson, Catherine Winters, Chelsea
433 Krieg, Shawn Del Percio, Margaret Orr and Daniel Warner. The authors also thank the three anonymous reviewers who
434 participate in the improvement of the quality of this paper.

435 **References**

- 436 Adams, J. L., Tipping, E., Bryant, C. L., Helliwell, R. C., Toberman, H., and Quinton, J.: Aged riverine particulate organic
437 carbon in four UK catchments, *Sci. Total Environ.*, 536, 648–654, doi:10.1016/j.scitotenv.2015.06.141, 2015.
- 438 Armas-Herrera, C. M., Dignac, M.-F., Rumpel, C., Arbelo, C. D., and Chabbi, A.: Management effects on composition and
439 dynamics of cutin and suberin in topsoil under agricultural use, *Eur. J. Soil Sci.*, 67(4), 360–373, doi:10.1111/ejss.12328,
440 2016.
- 441 Challinor, J. M.: Characterisation of wood by pyrolysis derivatisation—gas chromatography/mass spectrometry, *J. Anal.*
442 *Appl. Pyrolysis*, 35(1), 93–107, doi:10.1016/0165-2370(95)00903-R, 1995.
- 443 Collins, A. L. and Walling, D. E.: Selecting fingerprint properties for discriminating potential suspended sediment sources in
444 river basins, *J. Hydrol.*, 261(1), 218–244, doi:10.1016/S0022-1694(02)00011-2, 2002.
- 445 Derenne, S. and Quéneá, K.: Analytical pyrolysis as a tool to probe soil organic matter, *J. Anal. Appl. Pyrolysis*, 111, 108–
446 120, doi:10.1016/j.jaap.2014.12.001, 2015.
- 447 Dhillon, G. S. and Inamdar, S.: Extreme storms and changes in particulate and dissolved organic carbon in runoff: Entering
448 uncharted waters?, *Geophys. Res. Lett.*, 40(7), 1322–1327, doi:10.1002/grl.50306, 2013.

449 Dhillon, G. S. and Inamdar, S.: Storm event patterns of particulate organic carbon (POC) for large storms and differences
450 with dissolved organic carbon (DOC), *Biogeochemistry*, 118(1), 61–81, doi:10.1007/s10533-013-9905-6, 2014.

451 Fox, J. F. and Papanicolaou, A. N.: Application of the spatial distribution of nitrogen stable isotopes for sediment tracing at
452 the watershed scale, *J. Hydrol.*, 358(1), 46–55, doi:10.1016/j.jhydrol.2008.05.032, 2008.

453 Frank, D. A., Pontes, A. W., and McFarlane, K. J.: Controls on Soil Organic Carbon Stocks and Turnover Among North
454 American Ecosystems, *Ecosystems*, 15(4), 604–615, doi:10.1007/s10021-012-9534-2, 2012.

455 Frostegård, Å., Tunlid, A., and Bååth, E.: Phospholipid Fatty Acid Composition, Biomass, and Activity of Microbial
456 Communities from Two Soil Types Experimentally Exposed to Different Heavy Metals, *Appl. Environ. Microbiol.*, 59(11),
457 3605–3617, 1993.

458 Galy, V., Peucker-Ehrenbrink, B., and Eglinton, T.: Global carbon export from the terrestrial biosphere controlled by erosion,
459 *Nature*, 521(7551), 204–207, 2015.

460 Goñi, M. A., Hatten, J. A., Wheatcroft, R. A., and Borgeld, J. C.: Particulate organic matter export by two contrasting small
461 mountainous rivers from the Pacific Northwest, U.S.A., *J. Geophys. Res. Biogeosciences*, 118(1), 112–134,
462 doi:10.1002/jgrg.20024, 2013.

463 Gourdin, E., Huon, S., Evrard, O., Ribolzi, O., Bariac, T., Sengtaheuanghoung, O., and Ayrault, S.: Sources and export of
464 particle-borne organic matter during a monsoon flood in a catchment of northern Laos, *Biogeosciences*, 12(4), 1073–1089,
465 doi:10.5194/bg-12-1073-2015, 2015.

466 Huon, S., Evrard, O., Gourdin, E., Lefèvre, I., Bariac, T., Reyss, J.-L., Henry des Tureaux, T., Sengtaheuanghoung, O.,
467 Ayrault, S., and Ribolzi, O.: Suspended sediment source and propagation during monsoon events across nested sub-
468 catchments with contrasted land uses in Laos, *J. Hydrol. Reg. Stud.*, 9, 69–84, doi:10.1016/j.ejrh.2016.11.018, 2017.

469 Inamdar, S., Johnson, E., Rowland, R., Warner, D., Walter, R., and Merritts, D.: Freeze–thaw processes and intense rainfall:
470 the one-two punch for high sediment and nutrient loads from mid-Atlantic watersheds, *Biogeochemistry*, doi:
471 10.1007/s10533-017-0417-7, 2017.

472 IPCC, 2001. *Climate Change. The IPCC Third Assessment Report. Volumes I (Science), II (Impacts and Adaptation) and III*
473 *(Mitigation Strategies)*. Cambridge Univ Press, Cambridge.

474 Jeanneau, L., Jaffrezic, A., Pierson-Wickmann, A.-C., Gruau, G., Lambert, T., and Petitjean, P.: Constraints on the Sources
475 and Production Mechanisms of Dissolved Organic Matter in Soils from Molecular Biomarkers, *Vadose Zone J.*, 13(7),
476 doi:10.2136/vzj2014.02.0015, 2014.

477 Jeanneau, L., Denis, M., Pierson-Wickmann, A.-C., Gruau, G., Lambert, T., and Petitjean, P.: Sources of dissolved organic
478 matter during storm and inter-storm conditions in a lowland headwater catchment: constraints from high-frequency
479 molecular data, *Biogeosciences*, 12(14), 4333–4343, doi:10.5194/bg-12-4333-2015, 2015.

480 Jeong, J.-J., Bartsch, S., Fleckenstein, J. H., Matzner, E., Tenhunen, J. D., Lee, S. D., Park, S. K., and Park, J.-H.:
481 Differential storm responses of dissolved and particulate organic carbon in a mountainous headwater stream, investigated by

482 high-frequency, in situ optical measurements, *J. Geophys. Res. Biogeosciences*, 117(G3), n/a-n/a,
483 doi:10.1029/2012JG001999, 2012.

484 Jung, B.-J., Lee, H.-J., Jeong, J.-J., Owen, J., Kim, B., Meusburger, K., Alewell, C., Gebauer, G., Shope, C., and Park, J.-H.:
485 Storm pulses and varying sources of hydrologic carbon export from a mountainous watershed, *J. Hydrol.*, 440, 90–101,
486 doi:10.1016/j.jhydrol.2012.03.030, 2012.

487 Jung, B.-J., Lee, J.-K., Kim, H., and Park, J.-H.: Export, biodegradation, and disinfection byproduct formation of dissolved
488 and particulate organic carbon in a forested headwater stream during extreme rainfall events, *Biogeosciences*, 11(21), 6119–
489 6129, doi:10.5194/bg-11-6119-2014, 2014.

490 Jung, B.-J., Jeanneau, L., Alewell, C., Kim, B., and Park, J.-H.: Downstream alteration of the composition and
491 biodegradability of particulate organic carbon in a mountainous, mixed land-use watershed, *Biogeochemistry*, 122(1), 79–99,
492 doi:10.1007/s10533-014-0032-9, 2015.

493 Koiter, A. J., Lobb, D. A., Owens, P. N., Petticrew, E. L., Tiessen, K. H. D., and Li, S.: Investigating the role of connectivity
494 and scale in assessing the sources of sediment in an agricultural watershed in the Canadian prairies using sediment source
495 fingerprinting, *J. Soils Sediments*, 13(10), 1676–1691, doi:10.1007/s11368-013-0762-7, 2013.

496 Kolattukudy, P.: Polyesters in Higher Plants, in: *Biopolyesters*, vol. 71, edited by W. Babel and A. Steinbüchel, Springer,
497 Berlin Heidelberg, 1-49, 2001. [online] Available from: http://dx.doi.org/10.1007/3-540-40021-4_1

498 Lee, C. S., Sung, T. M., Kim, H. S., and Jeon, C. H.: Classification of forensic soil evidences by application of THM-
499 PyGC/MS and multivariate analysis, *J. Anal. Appl. Pyrolysis*, 96, 33–42, doi:10.1016/j.jaap.2012.02.017, 2012.

500 Ludwig, W., Probst, J.-L., and Kempe, S.: Predicting the oceanic input of organic carbon by continental erosion, *Glob.*
501 *Biogeochem. Cycles*, 10(1), 23–41, doi:10.1029/95GB02925, 1996.

502 Mannino, A. and Harvey, H. R.: Terrigenous dissolved organic matter along an estuarine gradient and its flux to the coastal
503 ocean, *Org. Geochem.*, 31(12), 1611–1625, doi:10.1016/S0146-6380(00)00099-1, 2000.

504 Mueller, K. E., Polissar, P. J., Oleksyn, J., and Freeman, K. H.: Differentiating temperate tree species and their organs using
505 lipid biomarkers in leaves, roots and soil, *Org. Geochem.*, 52, 130–141, doi:10.1016/j.orggeochem.2012.08.014, 2012.

506 Nierop, K. G. J. and Verstraten, J. M.: Rapid molecular assessment of the bioturbation extent in sandy soil horizons under
507 pine using ester-bound lipids by on-line thermally assisted hydrolysis and methylation-gas chromatography/mass
508 spectrometry, *Rapid Commun. Mass Spectrom.*, 18(10), 1081–1088, doi:10.1002/rcm.1449, 2004.

509 Nierop, K. G. J., Preston, C. M., and Kaal, J.: Thermally Assisted Hydrolysis and Methylation of Purified Tannins from
510 Plants, *Anal. Chem.*, 77(17), 5604–5614, doi:10.1021/ac050564r, 2005.

511 Nosrati, K., Govers, G., Ahmadi, H., Sharifi, F., Amoozegar, M. A., Merckx, R., and VanMaerke, M.: An exploratory study
512 on the use of enzyme activities as sediment tracers: biochemical fingerprints?, *Int. J. Sediment Res.*, 26(2), 136–151,
513 doi:10.1016/S1001-6279(11)60082-6, 2011.

514 Oeurng, C., Sauvage, S., Coynel, A., Maneux, E., Etcheber, H., and Sánchez-Pérez, J.-M.: Fluvial transport of suspended
515 sediment and organic carbon during flood events in a large agricultural catchment in southwest France, *Hydrol. Process.*,
516 25(15), 2365–2378, doi:10.1002/hyp.7999, 2011.

517 Raymond, P. A. and Oh, N.-H.: An empirical study of climatic controls on riverine C export from three major U.S.
518 watersheds, *Glob. Biogeochem. Cycles*, 21(2), n/a-n/a, doi:10.1029/2006GB002783, 2007.

519 Ritchie, J. C., Spraberry, J. A., and McHenry, J. R.: Estimating Soil Erosion from the Redistribution of Fallout ¹³⁷Cs₁, *Soil*
520 *Sci. Soc. Am. J.*, 38(1), 137–139, doi:10.2136/sssaj1974.03615995003800010042x, 1974.

521 Rowland, R., Inamdar, S., and Parr, T.: Evolution of particulate organic matter (POM) along a headwater drainage: role of
522 sources, particle size class, and storm magnitude, *Biogeochemistry*, 133(2), 181–200, doi:10.1007/s10533-017-0325-x, 2017.

523 Smith, J. C., Galy, A., Hovius, N., Tye, A. M., Turowski, J. M., and Schleppe, P.: Runoff-driven export of particulate organic
524 carbon from soil in temperate forested uplands, *Earth Planet. Sci. Lett.*, 365, 198–208, doi:10.1016/j.epsl.2013.01.027, 2013.

525 Tamoo, F., Van den Meersche, K., Meysman, F., Marwick, T. R., Borges, A. V., Merckx, R., Dehairs, F., Schmidt, S.,
526 Nyunja, J., and Bouillon, S.: Distribution and origin of suspended matter and organic carbon pools in the Tana River Basin,
527 Kenya, *Biogeosciences*, 9(8), 2905–2920, doi:10.5194/bg-9-2905-2012, 2012.

528 Tank, J. L., Rosi-Marshall, E. J., Griffiths, N. A., Entekin, S. A., and Stephen, M. L.: A review of allochthonous organic
529 matter dynamics and metabolism in streams, *J. North Am. Benthol. Soc.*, 29(1), 118–146, doi:10.1899/08-170.1, 2010.

530 Wallbrink, P. J. and Murray, A. S.: Distribution and Variability of ⁷Be in Soils Under Different Surface Cover Conditions and
531 its Potential for Describing Soil Redistribution Processes, *Water Resour. Res.*, 32(2), 467–476, doi:10.1029/95WR02973,
532 1996.

533 Walling, D. E.: Use of ¹³⁷Cs and other fallout radionuclides in soil erosion investigations: progress, problems and prospects,
534 Joint FAO/IAEA Division of Nuclear Techniques in Food and Agriculture, International Atomic Energy Agency (IAEA),
535 Vienna (Austria). [online] Available from:
536 http://www.iaea.org/inis/collection/NCLCollectionStore/_Public/29/049/29049354.pdf, 1998.

537 Walling, D. E.: The evolution of sediment source fingerprinting investigations in fluvial systems, *J. Soils Sediments*, 13(10),
538 1658–1675, doi:10.1007/s11368-013-0767-2, 2013.

539 Wang, X., Cammeraat, E. L. H., Romeijn, P., and Kalbitz, K.: Soil Organic Carbon Redistribution by Water Erosion – The
540 Role of CO₂ Emissions for the Carbon Budget, *PLoS One*, 9(5), e96299, doi:10.1371/journal.pone.0096299, 2014.

541 Williams, J. S., Dungait, J. A. J., Bol, R., and Abbott, G. D.: Contrasting temperature responses of dissolved organic carbon
542 and phenols leached from soils, *Plant Soil*, 399, 13–27, doi:10.1007/s11104-015-2678-z, 2016.

543 Wischmeier, W. H.: A Rainfall Erosion Index for a Universal Soil-Loss Equation¹, *Soil Sci. Soc. Am. J.*, 23(3), 246–249,
544 doi:10.2136/sssaj1959.03615995002300030027x, 1959.

545 **Figure captions**

546 Figure 1: Location of the study watershed in the Piedmont region of Maryland. Composite suspended sediments were
547 sampled at the 12 and 79 ha locations (grey circles). The sites of collection of end-members are indicated with triangles:
548 violet for wetland soils (Wet), blue for bed sediments (SBed), green for forest soil humus (FH) and litter (Li), orange for
549 upland soils (Up) and yellow for bank sediments from horizons A and B (BaA and BaB).

550 Figure 2: Relative proportions of low organic acids (LOA), phenolic compounds (PHE), low molecular weight and high
551 molecular weight fatty acids (LMW and HMW FA) and carbohydrates (CAR) among identified target compounds in the end
552 members. Uncertainties correspond to standard deviation of sampling triplicates (duplicates for bed sediments SBed).

553 Figure 3: Relative proportions of low organic acids (LOA), phenolic compounds (PHE), low molecular weight and high
554 molecular weight fatty acids (LMW and HMW FA) and carbohydrates (CAR) among identified target compounds in the
555 coarse, medium and fine fractions of CSS. Uncertainties correspond to the inter-event standard deviation.

556 Figure 4: Plan defined by the two first factors of the PCA calculated using the distribution of target compounds. Squares
557 represent end members Li (green), FH-W (red) and SBed (blue). The area characteristic of each end member is defined by the
558 95% confident interval. Circles represent CSS from the 12 ha (orange) and the 79 ha (purple) locations. The mean positions
559 for each size fraction are represented by large circles and uncertainties correspond to inter-event standard deviation.

560 Figure 5: 2D plot illustrating the variability of the distribution of plant-derived markers using the relative proportion of
561 phenolic compounds (PHE) against HMW fatty acids and the proportion of α,ω diacids and ω OH fatty acids among HMW
562 fatty acids (denoted HMW FA ratio).

563 Figure 6: Illustration of the most significant correlations between the source apportionments performed using the molecular
564 data and rainfall characteristics. At the 12 ha location, positive correlations (a) between the proportion of Sbed-derived POM
565 and I_{max} and (b) between the proportion of FH-W-derived POM and I_{med} . At the 79 ha location, positive correlation
566 between Sbed-derived POM and rainfall amount (c). Coarse, medium and fine fractions are depicted by the dark grey, light
567 grey and white circles, respectively and the composite POM by the black diamond.

Table 1. Rainfall characteristics, discharge and proportion of coarse, medium and fine fractions for the 4 investigated storm events.

	Event 1	Event 2	Event 3	Event 4				
	<i>May 1, 2014</i>	<i>Apr. 21, 2015</i>	<i>July 3, 2015</i>	<i>Sept. 30, 2015</i>				
Rainfall								
total (mm)	148.9	43.9	97.4	40.1				
max (mm h ⁻¹)	19.9	20	31.3	20.2				
median (mm h ⁻¹)	1.3	2.2	0.4	2.1				
API7 (mm)	9.7	10.4	68.2	0				
Discharge (12 ha catchment)								
max (l s ⁻¹)	150.1	68.3	87.4	15.5				
Particle size distribution								
	<i>12 ha</i>	<i>79 ha</i>	<i>12 ha</i>	<i>79 ha</i>	<i>12 ha</i>	<i>79 ha</i>	<i>12 ha</i>	<i>79 ha</i>
Coarse (%)	52	81	20	43	12	58	6	nd
Medium (%)	22	13	22	32	27	20	18	nd
Fine (%)	27	7	59	25	61	21	75	nd

568

569

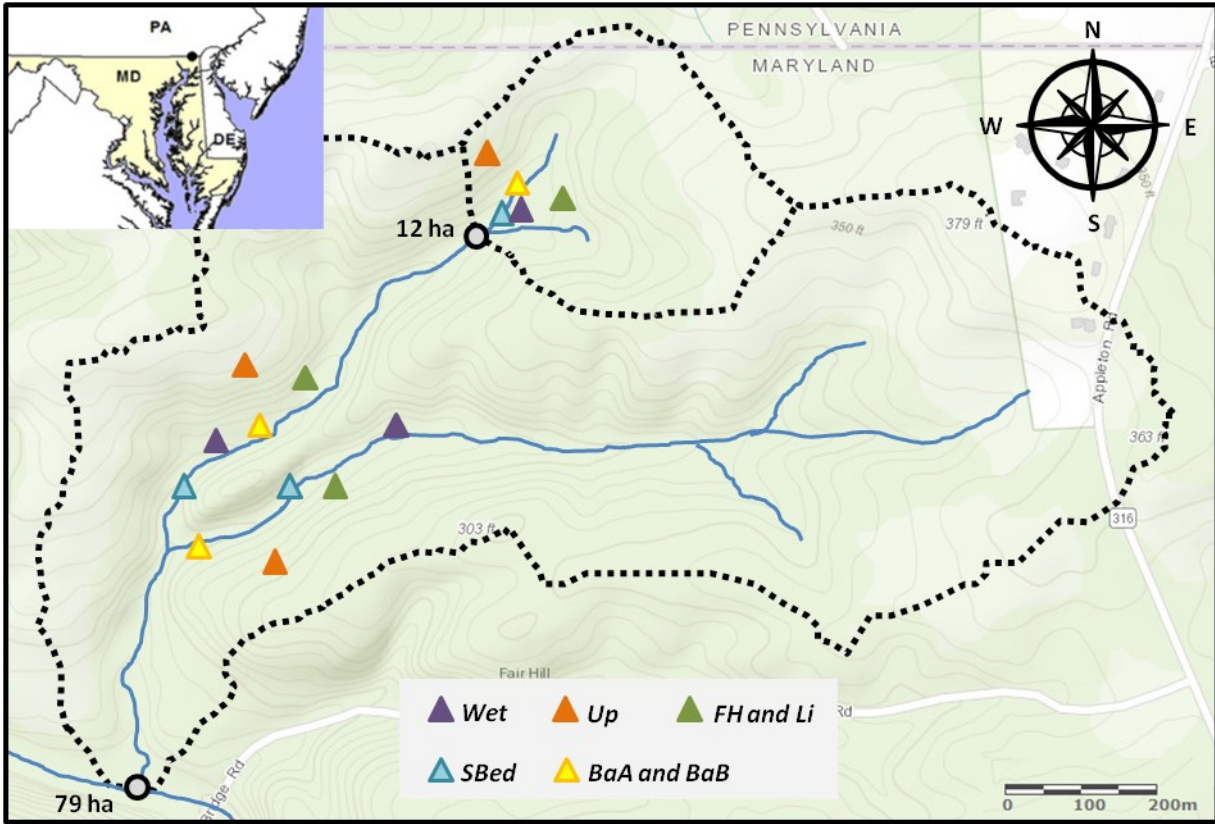
Table 2. Source apportionment calculated using the molecular data.

		<i>12 ha location</i>			<i>79 ha location</i>		
		Li (%)	Sbed (%)	FH-W (%)	Li (%)	Sbed (%)	FH-W (%)
Event 1 <i>May 1, 2014</i>	C	97 ± 7	1 ± 2	3 ± 7	45 ± 9	55 ± 9	0 ± 0
	M	78 ± 7	18 ± 5	4 ± 8	0 ± 1	95 ± 7	4 ± 7
	F	76 ± 12	13 ± 6	11 ± 16	48 ± 9	52 ± 9	0 ± 0
	POM	92 ± 9	4 ± 4	4 ± 11	42 ± 6	57 ± 8	0 ± 2
Event 2 <i>Apr. 21, 2015</i>	C	95 ± 8	2 ± 3	3 ± 8	95 ± 8	2 ± 4	3 ± 8
	M	94 ± 9	2 ± 3	4 ± 9	57 ± 9	43 ± 8	0 ± 1
	F	69 ± 16	15 ± 6	17 ± 20	51 ± 10	48 ± 9	1 ± 2
	POM	86 ± 11	6 ± 4	8 ± 12	89 ± 9	8 ± 7	3 ± 4
Event 3 <i>July 3, 2015</i>	C	96 ± 5	3 ± 4	1 ± 5	47 ± 9	53 ± 9	0 ± 0
	M	87 ± 6	10 ± 5	3 ± 7	42 ± 9	58 ± 9	0 ± 0
	F	61 ± 8	39 ± 7	0 ± 1	45 ± 11	51 ± 9	4 ± 6
	POM	81 ± 6	17 ± 6	2 ± 4	46 ± 10	53 ± 9	2 ± 2
Event 4 <i>Sept. 30, 2015</i>	C	73 ± 22	0 ± 0	27 ± 22	fraction not available		
	M	70 ± 22	0 ± 0	30 ± 22	36 ± 20	0 ± 0	64 ± 22
	F	62 ± 23	0 ± 0	38 ± 23	77 ± 20	2 ± 2	21 ± 22
	POM	65 ± 23	0 ± 0	35 ± 23	-	-	-

570

571

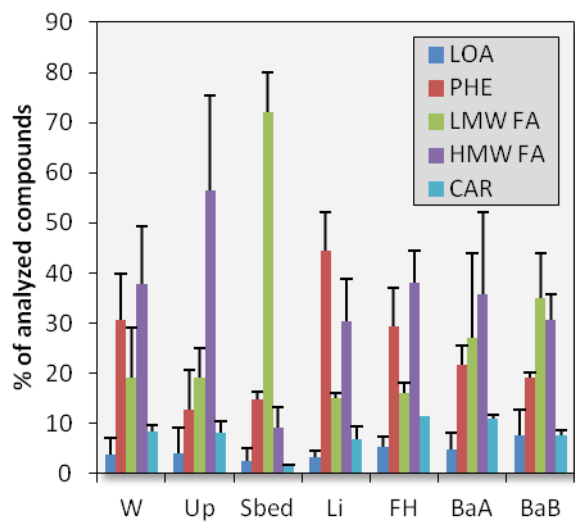
572 Figure 01



573

574

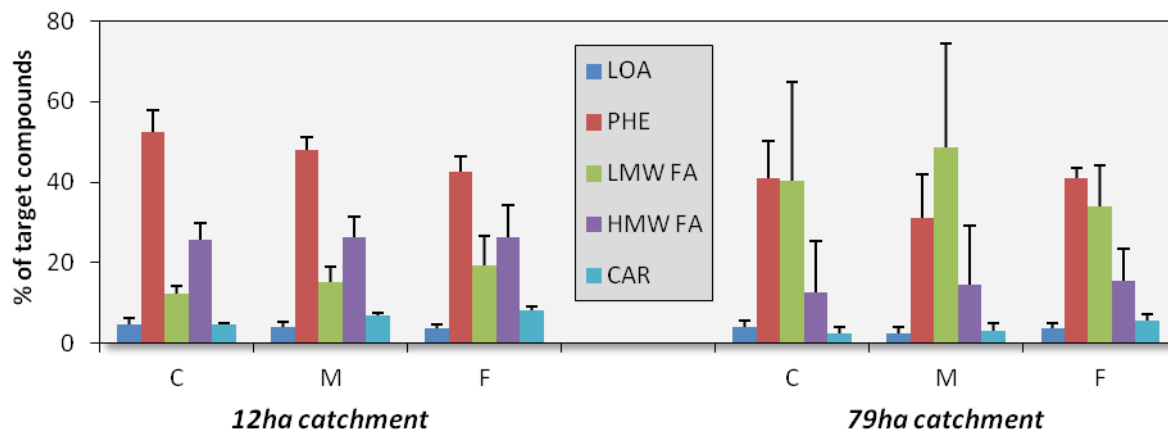
575 **Figure 02**



576

577

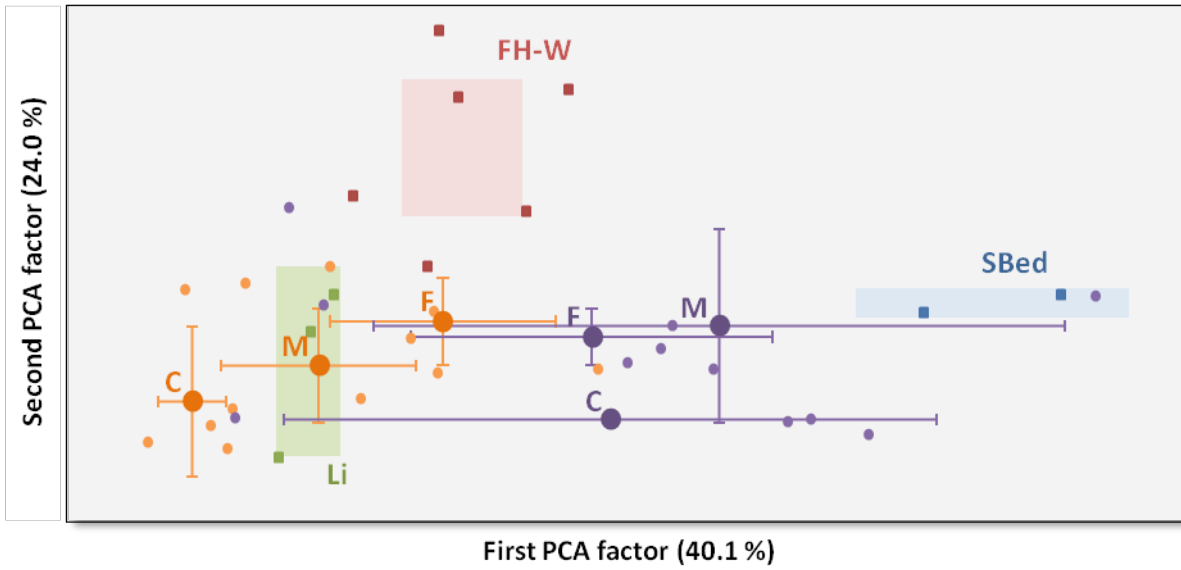
578 **Figure 03**



579

580

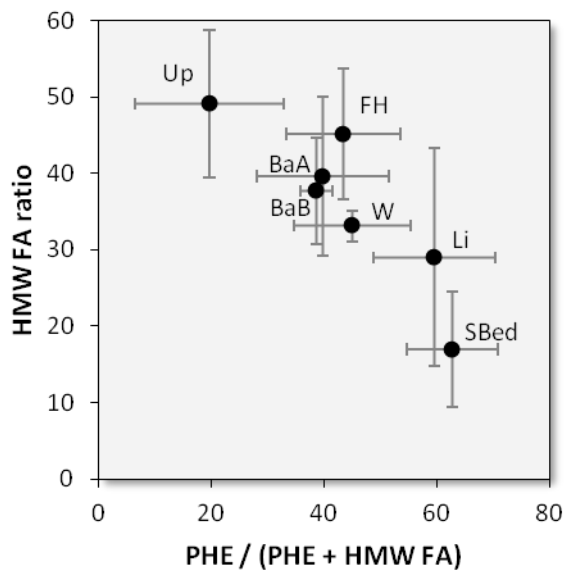
581 Figure 04



582

583

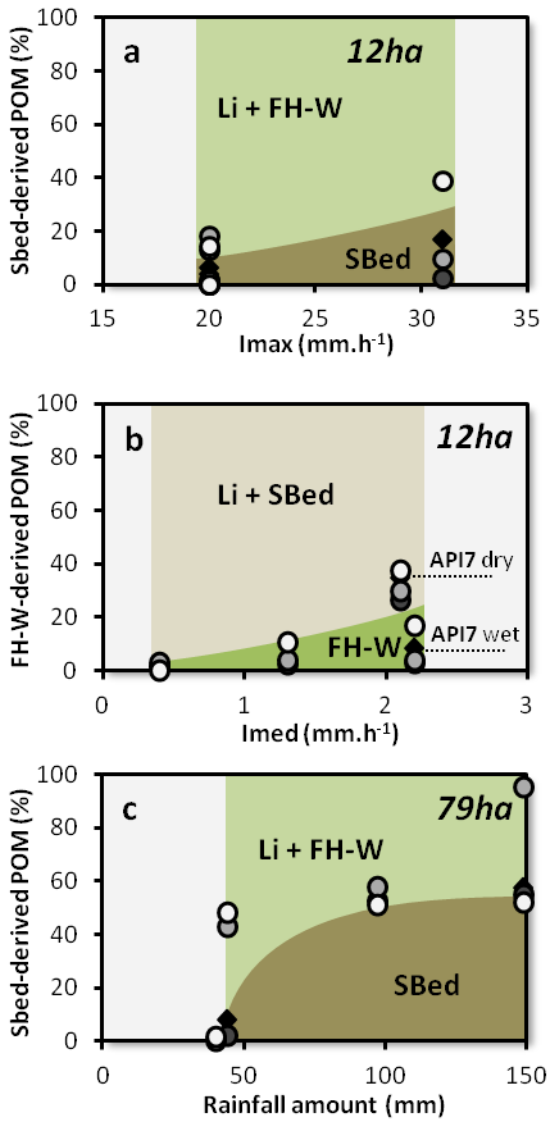
584 **Figure 05**



585

586

587 **Figure 06**



588

589

590