| 1 | The <i>n</i> -alkane and sterol composition of living fen plants as a potential |
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| 2 | tool for palaeoecological studies |
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13 ABSTRACT

In groundwater-fed fen peatlands, the surface biomass decays rapidly and as a result, highly 14 15 humified peat is formed. A high degree of humification constrains palaeoecological studies because reliable identification of plant remains is hampered. Organic geochemistry 16 techniques as a means of identifying historical plant communities have been successfully 17 applied in bog peats. The method has also been applied to fen peat, but without reference to 18 the composition of fen plants. In this study we have applied selected organic geochemistry 19 methods to determine the composition of neutral lipid fractions from 12 living fen plants, to 20 21 investigate the potential for the distributions to characterize and separate different fen plants 22 and plant groups. Our results show correspondence with previous studies, e.g. C23 and C25 nalkanes dominate Sphagnum spp. and C27 to C31 alkanes dominate vascular plants. However, 23 24 we also found similarities in *n*-alkane distributions between *Sphagnum* spp. and the below ground parts of some vascular plants. We tested the efficiency of different *n*-alkane ratios to 25 separate species and plant groups. The ratios used in bog studies (e.g. $n-C_{23}/n-C_{25}$ and $n-C_{23}/n-C_{25}$ 26 $C_{23}/n-C_{29}$) did not work as consistently on fen plants. Some differences in sterol distribution 27 28 were found between vascular plants and mosses; in general vascular plants had a higher concentration of sterols. When distributions of *n*-alkanes, *n*-alkane ratios and sterols were all 29 30 included as variables, redundancy analyses (RDA) separated different plant groups into their own clusters. Our results imply that the pattern in bog biomarkers cannot directly be applied 31 to fen environments. Nevertheless, they encourage further testing to determine whether or not 32 the identification of plant groups, plants or plant parts from highly humified peat is possible 33 by applying fen species-specific biomarker proxies. 34

Keywords: biomarker, geochemistry, palaeoecology, peatland, fen, *Sphagnum*, vascular
plant.

37 **1. Introduction**

Northern peatlands comprise a large store of carbon, 547 (473-621) Pg (Yu et al., 2010); 38 acute and contemporary questions related to human-induced changes in climate have 39 therefore emphasized the importance of thoroughly understanding peatland dynamics, past 40 and present. Peatland carbon balance is highly sensitive to moisture conditions (e.g. Alm et 41 al., 1999; Waddington and Roulet, 2000): the water table depth defines both the prevailing 42 plant assemblages and the thickness of the oxic layer, where most biological production and 43 decomposition take place. Hydrology and the source of nutrients are the main factors 44 controlling the type of peatland and the occurrence of species (Wheeler and Proctor, 2000, 45 46 Økland et al., 2001). The nutrient (trophic) level of a peatland is described as a gradient from 47 nutrient rich to nutrient poor: eutrophic, minerotrophic, mesotrophic, oligotrophic and ombrotrophic. Nutrient poor bogs receive water and nutrients only through precipitation 48 while nutrient rich fens receive water and nutrients from atmospheric input, groundwater and 49 50 underlying and surrounding mineral soils (Rydin et al., 2006). Bogs are characterized by dry and wet microhabitats: hummocks (surface 20-50 cm above the water table), intermediate 51 lawns (5-20 cm above the water table) and wet flarks where the water table is at the surface, 52 each maintaining specific plant assemblages. Fens on the other hand do not have such 53 distinguishable microhabitat formation (Rydin et al., 2006, Laine et al., 2009). Given the vital 54 55 role of vegetation in the peatland carbon budget (e.g. Riutta et al., 2007) and the fact that bryophyte and vascular plant dominated communities differ in their CO₂ and CH₄ dynamics 56 (Laine et al., 2007, Levy et al., 2012) it is important to understand past mechanisms that have 57 58 controlled the vegetation dynamics. Historical variations in climate and hydrology are 59 preserved in peat layers as alterations in the assemblages of different biological organisms. In particular, past vegetation assemblages have been a key proxy for reconstructing past 60 61 moisture conditions in a range of sites (e.g. Barber et al., 1998; Mauquoy et al., 2002; Tuittila et al., 2007; Väliranta et al., 2007). This reflects the slow and incomplete decomposition of 62 peat in bog environments, meaning that bogs usually contain relatively well preserved plant 63 material for palaeoecological examination. In contrast, in fen environments surface decay is 64 rapid and a major part of the peat below the surface layer is highly humified (Moore et al., 65 66 2007). Fen peats thus tend to lack identifiable plant remains. Given that all bogs are underlain by a fen peat phase and a major proportion of the northern peatlands are still in a fen phase, 67 there is considerable spatio-temporal restriction for palaeoecological applications based on 68 69 identifiable plant remains alone.

71 Studies of bog peats have shown that plant biomarkers, i.e. species-specific compounds, can be successfully applied to less-humified peat to identify fossil plant groups (e.g. Xie et al., 72 2000; Avsejs et al., 2002; Pancost et al., 2002, 2003; Nichols et al., 2006; Jia et al., 2008; 73 McClymont et al., 2008; Bingham et al., 2010). Different plant groups can be separated, for 74 75 instance by comparing *n*-alkane distributions and ratios, e.g. the difference in concentration of low molecular weight (LMW) n-C₂₃ and n-C₂₅, and high molecular weight (HMW) n-C₂₉ 76 77 and $n-C_{33}$ can be used to separate contributions from Sphagnum and non-Sphagnum species (Pancost et al., 2002, Nichols et al., 2006, Vonk and Gustafsson, 2009, Lopez-Diaz et al., 78 79 2010, Ortiz et al., 2011, Andersson et al., 2011). Studies have also shown that some moss species can be distinguished down to species level (Jia et al., 2008; Bingham et al., 2010), 80 e.g. $n-C_{23}/n-C_{25}$ alkane ratio in bog peat may track changes in Sphagnum fuscum abundance 81 82 (Bingham et al., 2010).

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A thorough investigation of the lipid distributions in fen plants has not, to our knowledge, been performed. As a result, it is not clear whether or not the application of biomarker ratios from ombrotrophic peat plants would be a robust approach for the characterization of peatlands including fens (Andersson et al., 2011). In this study we have applied selected organic geochemical analyses to living fen plant species, excluding the litter. Specifically, we aimed to define whether or not (i) the analyses could separate bryophytes from vascular plants and (ii) there are specific fen plant proxies.

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- 92 **2.** Material and methods
- 93 *2.1. Sampling*

Samples of living plants were collected from three individual but closely located fens from
the Siikajoki commune (64°45′N, 24°42′E) in the mid-boreal bio-climate zone in Finland

96 (Fig. 1). The water level of fens is on average 10 cm below the soil surface and the pH of the
97 water squeezed from the mosses is between 4 and 4.3. A detailed description of the sites
98 (SJ2-4) is given by Leppälä et al. (2011) and Laine et al. (2011).

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Twelve plant species typical of fens were chosen: five bryophyte species and seven vascular 100 plant species (Table 1). Bryophytes were treated as whole plants. Vascular plants, sedges and 101 forbes were divided into above and below ground parts because previous studies had shown 102 that the *n*-alkane concentration might vary between different plant parts (Dawson et al., 2000, 103 Jansen et al., 2006). In fen environments sedge and forb roots may also form a substantial 104 105 contribution to the organic matter (OM) input to the upper peat (cf. Saarinen, 1996, Moore et al. 2002, Andersson et al., 2011, Huang et al., 2011). To assess methodological 106 reproducibility we repeated the analyses with six randomly selected samples [Sphagnum 107 108 papillosum, Warnstorfia exannulata, Carex rostrata (above and below ground parts), Potentilla palustris and Menyanthes trifoliata (above ground parts). Compound 109 110 concentrations are as mean values, and the standard error of the mean (SE) is reported when 111 the compound was found in both the original and repeated analyses. Moreover, we collected and analyzed a selection of species (W. exannulata, and the below ground parts of C. 112 rostrata, C. livida, C. nigra, C. lasiocarpa, E. angustifolium and M. trifoliata) from a nearby 113 peatland. This procedure was executed in order to test for location-related variation in 114 compositions. Total organic carbon (TOC) was measured to test whether or not the lipid 115 concentration between sampled plants/plant parts differed because of TOC content or 116 concentration calculated from dry weight. 117

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119 *2.2. Solvent extraction*

120 The plant parts were separated and washed with distilled water. Lipids were extracted from ca. 0.2 g of the freeze dried and ground samples using repeated ultrasonication (20 min) with 121 6 ml CH₂Cl₂/MeOH (3:1, v/v). Samples were saponified with 0.5 M methanolic (95%) NaOH 122 123 for 2 h at 70 °C and the neutral lipids extracted using hexane. The neutral lipids were further separated into apolar and polar compounds using activated Al₂O₃ columns, eluting with 124 hexane/CH₂Cl₂ (9:1, v/v) and CH₂Cl₂/MeOH (1:2, v/v), respectively. Prior to analysis using 125 126 gas chromatography (GC) and GC-mass spectrometry (GC-MS) the polar fractions were derivatised using bis(trimethylsilyl)trifluoroacetamide (Sigma Aldrich). 127

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129 2.3. GC-MS

Apolar and polar fractions were analyzed using GC-MS with the gas chromatograph 130 131 equipped with flame ionisation detection (GC-FID) and split/splitless injection (280 C). Separation was achieved with a fused silica column (30 m x 0.25 mm i.d) coated with 132 133 0.25µm 5% phenyl methyl siloxane (HP-5MS), with He as carrier gas, and the following oven temperature programme: 60 - 200 °C at 20 °C/min, then to 320 °C (held 35 min) at 134 135 6°C/min. The mass spectrometer was operated in full scan mode (50-650 amu/s, electron voltage 70eV, source temperature 230 °C). Compounds were assigned using the NIST mass 136 spectral database and comparison with published spectra (e.g. Goad and Akihisa, 1997; 137 138 Killops and Frewin, 1994). Quantification was achieved through comparison of integrated peak areas in the FID chromatograms and those of internal standards of known concentration 139 140 $(5-\alpha$ -cholestane for apolars and 2-nonadecanone for polars). Concentration values are given as concentration per dry weight of extracted material. The concentration from replicate 141 142 samples was averaged in the statistical analysis.

144 *2.4. Statistical analysis*

We applied multivariate analysis to study the variation within the biomarker data. To test 145 whether or not the variation was related to the specific plant groups or their component parts, 146 we applied redundancy analysis (RDA) with three plant groups: mosses, above ground and 147 below ground vascular plant parts. We conducted a series of RDA determinations. First, we 148 analyzed the data and tested the significance separately for different variables: n-alkanes, n-149 alkane ratios, *n*-alcohols and sterols; we then applied RDA for different compound 150 combinations to find the solution best explained by the three plant group variables. To link 151 152 biomarker composition to plant species we included the latter as a passive variable into the analysis. A Monte Carlo permutation test was used to test the significance of the RDA 153 solutions in all of the analyses. To make the analyses robust, compounds detected in fewer 154 155 than four samples were excluded, because they might skew the result in favor of those 156 samples in which they existed. This means that some species-specific markers were not included in the statistical analysis, but they are mentioned when individual plant analyses are 157 discussed. The statistical analyses were conducted using Canoco for Windows 4.52 (ter Braak 158 and Smilauer, 2002). 159

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161 **3. Results**

We found *n*-alkanes within range of n-C₁₇ to n-C₃₅ in the apolar fraction. A few samples contained taraxer-14-ene, taraxast-20-ene, and an unidentified triterpanoid and taraxeroid. In the polar fraction we found sterols and *n*-alcohols. The absolute concentrations of compounds did differ between sample sets (original, repeated and replicate), but the dominance order of compounds was maintained (the full data set can be downloaded from www.pangaea.de, reference PANGAEA PDI-4071). The samples contained no evidence of bacterial activity as no traces of hopanoids or archaeol were found. A few replicate samples contained stanols and ketones. This suggests a low level of degradation and that the samples contained compounds solely from the plants under study (e.g. Nishimura 1977, Lehtonen and Ketola 1993, Jiao et al. 2008). There was a linear correlation between concentration calculated as mass per dry weight (μ g g⁻¹) and as mass per total organic carbon (μ g TOC), indicating no bias due to selective preservation of OM between plant species or plant groups (Fig. 2).

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The total concentration of *n*-alkanes (C_{17} - C_{35}) in moss species varied between 86.4 µg g⁻¹ (*Sphagnum fimbriatum*) and 9.2 µg g⁻¹ (SE 3.8) (*W. exannulata*). The distribution of *n*alkanes of *Sphagnum* species showed an odd/even preference (Fig. 3). The C_{23} *n*-alkane dominated in *S. subsecundum* (23.6 µg g⁻¹), *S. riparium* (12.4 µg g⁻¹), and *S. papillosum* (32.6µg g⁻¹), whereas *n*- C_{25} dominated in *S. fimbriatum* (21.3 µg g⁻¹). In *W. exannulata*, C_{27} and C_{25} *n*-alkanes dominated (2.3 and 1.5 µg g⁻¹, respectively; Fig. 3). *W. exannulata* was the only moss species where taraxast-20-ene was detected (4.7 µg g⁻¹).

In above ground sedge parts an odd predominance was also present. Total *n*-alkane concentration was highest in *Carex rostrata* (332.7 μ g g⁻¹, SE 44.5) and lowest in *C. nigra* (21.7 μ g g⁻¹). The C₂₇ *n*-alkane dominated in above ground parts of *C. livida* (7.1 μ g g⁻¹), *C. nigra* (12.3 μ g g⁻¹) and *Eriophorum angustifolium* (10.2 μ g g⁻¹) and C₂₉ in *C. rostrata* (188.8 μ g g⁻¹, SE 30.8) and *C. lasiocarpa* (52.8 μ g g⁻¹) (Fig. 3).

The *n*-alkane distributions in the below ground sedge parts were more complex. The short chain *n*-alkanes (C₁₉, C₂₁ and C₂₃) were more abundant (Fig. 3) than the long chain *n*-alkanes (C₂₇, C₂₉ and C₃₁). However, for instance, *C. nigra* had the highest concentration of *n*-C₂₇ (5.2 μ g g⁻¹) but the short chain *n*-alkanes were also present. In *C. lasiocarpa*, C₂₃ had the highest concentration (15.7 μ g g⁻¹); in *C. rostrata* the most abundant alkane was also *n*-C₂₃ 193 (24.7 μ g g⁻¹; SE 10.2). *C. livida* was dominated by *n*-C₂₁ (5.7 μ g g⁻¹). The below ground parts 194 *C. lasiocarpa* was the only sample where small amounts of taraxer-14-ene (4.4 μ g g⁻¹), an 195 unidentified triterpenoid (1.5 μ g g⁻¹) and taraxast-20-ene (0.6 μ g g⁻¹) were found. *E.* 196 *angustifolium* was dominated by *n*-C₂₇ (14.7 μ g g⁻¹; Fig. 3).

197 *Menyanthes trifoliata* above ground parts had the lowest total *n*-alkane concentration (5.6 μ g 198 g⁻¹; SE 0.3). In contrast, below ground plant parts had a much higher total concentration of *n*-199 alkanes (89.7 μ g g⁻¹) than the above ground parts. Short chain *n*-C₂₁ and *n*-C₂₃ alkanes 200 dominated below ground plant parts (38.0 and 25.5 μ g g⁻¹, respectively), while long chain *n*-201 alkanes were present in small amount (Fig. 3).

In *Potentilla palustris*, the above ground and below ground parts were dominated by the long- chain *n*-alkanes and *n*-C₃₁ had the highest concentration in both (497.6 μ g g⁻¹, SE 182.6 and 14.7 μ g g⁻¹, respectively). *Potentilla palustris* above ground parts had the highest total concentration of *n*-alkanes (985.6 μ g g⁻¹; SE 350.4; Fig. 3).

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3.2. n-Alkane ratios

Ratios between different *n*-alkanes have been found to be useful markers for distinguishing
species in bog environments (Nott et al., 2000, Ishiwatari et al., 2005, Jansen et al., 2006,
Nichols et al. 2006, Zheng et al., 2007, Vonk and Gustafsson, 2009, Bingham et al., 2010,
Andersson et al., 2011). The ratios calculated here were used in these studies.

Below ground parts of *Carex nigra, C. livida* and *C. lasiocarpa* showed the highest $n-C_{23}/n-C_{23}$ C₂₅ (ca. 5 to 7) ratio, whereas *Sphagnum* spp. had lower values (ca. 0.6 to 3). The ratios $n-C_{23}/n-C_{27}$, $n-C_{23}/n-C_{29}$ and $n-C_{23}/n-C_{31}$ were all low for the above ground plant parts (around 0) and high in *Sphagnum* species (> 10), especially *S. papillosum*. However, $n-C_{23}/n-C_{27}$ for 216 *C. lasiocarpa* below ground parts (ca. 20) and $n-C_{23}/n-C_{29}$ and $n-C_{23}/n-C_{31}$ for *M. trifoliata* 217 (ca. 40 and 70 respectively) were higher than the values in *Sphagnum* spp. The $n-C_{25}/n-C_{29}$ 218 ratio had a similar distribution pattern to the three above ratios, but with smaller values and 219 more complex distribution in *Sphagnum* spp. (ca. 6 to 15). *Potentilla palustris* above ground 220 and below ground parts were clearly separated from other samples via $n-C_{31}/n-C_{27}$ (> 5) and 221 $n-C_{31}/n-C_{29}$ (PANGAEA PDI-4071).

The $n-C_{23}/(n-C_{23} + n-C_{29})$ and $n-C_{25}/(n-C_{25} + n-C_{29})$ ratios distinguished Sphagnum spp. as 222 their own group (ca. 0.9). The pattern was clearest for $n-C_{25}/(n-C_{25} + n-C_{29})$ where Sphagnum 223 spp. ratio values (> 0.8) consistently exceed higher plant values, excluding M. trifoliata 224 below ground parts, which was equals to moss values (Fig.4). For $n-C_{23}/(n-C_{27}+n-C_{31})$, M. 225 trifoliata had the highest value (ca. 11), and S. papillosum stood out owing to a higher 226 value(ca. 9) than the rest of the mosses and vascular plants. $P_{aq} [(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25})/(n-C_{23}+n-C_{25}+n-C_{25})/(n-C_{23}+n-C_{25}+n-C_{25}+n-C_{25})/(n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}+n-C_{25}$ 227 228 $C_{25}+ n-C_{29}+ n-C_{31}$ did not seem to separate plant species. However, S. fimbriatum and S. papillosum had higher P_{aq} values (ca. 1) than the rest of the samples. $P_{wax} [(n-C_{27}n+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_{29}+n-C_$ 229 $C_{31}/(n-C_{23}+n-C_{25}+n-C_{27}+n-C_{29}+n-C_{31})$] showed low values for Sphagnum spp. and M. 230 trifoliata and C. lasiocarpa below ground parts (max. 0.2) and high values for most of the 231 vascular plant above ground parts, and *W. exannulata* (> 0.8; PANGAEA PDI-4071). 232

According to previous studies the average *n*-alkane chain length (ACL) should separate mosses and vascular plant leaves from each other (Zhou et al., 2005). In our samples the ACL of the mosses and below ground plant parts, except *E. angustifolium* and *C. nigra*, was < 26. Vascular plant above ground parts recorded ACL values > 26 (Fig. 4).

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238 *3.3.Polar fraction*

239 The *n*-alcohol concentration had only minor differences between different plant types and the dominant compounds often overlapped. Among the Sphagnum mosses the total concentration 240 of sterols was 2100 to 2600 µg g⁻¹. It seems that Sphagna had no dominant sterol, but 241 generally similar contributions from campesterol [campest-5-en-3β-ol], stigmasterol [(24*E*)-242 stigmasta-5,22-dien-3 β -ol] and β -sitosterol [(3 β)-stigmast-5-en-3-ol] were observed. 243 Brassicasterol [(22*E*)-ergosta-5,22-dien-3β-ol], 24-methylcholest-7-en-3β-ol, obtusifoliol 244 $[4\alpha, 14\alpha$ -dimethyl- 5α -ergosta- $8, 24(24^1)$ -dien- 3β -ol], ergost-8, 24(28)-dien- 3β -ol were typical 245 for Sphagnum spp., but were occasionally also detected in small amounts in some of the 246 vascular plants. W. exannulata samples were also characterized by high concentrations of 247 campesterol, stigmasterol and β-sitosterol but, in contrast to the Sphagnum spp., obtusifoliol 248 was not detected and the concentration of phytol was clearly highest (2035.9 μ g g⁻¹, SE 249 913.3, but 437.9 µg g^{-1}_{rep}) among all the mosses. Gramisterol [4 α -methyl-5 α -ergosta-250 7,24(24¹)-dien-3 β -ol], albeit in low concentration (ca. 10-60 μ g g⁻¹), was detected in all the 251 mosses but not the vascular plants (Table 2). 252

All vascular plant below ground parts, excluding C. nigra, had a higher concentration of 253 sterols than above ground parts. Above parts were dominated by β-sitosterol, with the 254 occasional presence of the associated stanol (3-stigmastanol) and cycloartenol (5α-cycloart-255 24-en-3β-ol). Sedge below ground parts were similar to the above ground parts, only with a 256 smaller amount of phytol [(3,7,11,15-tetramethylhexadec-2-en-1-ol; ca. 10-60 μ g g⁻¹] and 257 higher amount of lupeol [5α-lup-20(29)-en-3β-ol; 20-250 μg g⁻¹]. M. trifoliata above and 258 below ground parts were dominated by stigmasterol (1483.6 μ g g⁻¹ and 3647.4 μ g g⁻¹). β -259 260 sitosterol and obtusifoliol were absent from all M. trifoliata samples, whereas the schottenol $[5\alpha$ -stigmast-7-en-3 β -ol] was present only in *M. trifoliata* samples (above ground 678.0 μ g g⁻ 261 ¹ and below ground 1029.4 μ g g⁻¹; PANGAEA PDI-4071). 262

263 To copherols- δ and - γ [(2R)-2,8-dimethyl-2-((4R,8R)-4,8,12-trimethyltridecyl)-6-chromanol (2R)-2,7,8-trimethyl-2-(4R,8R)-4,8,12-trimethyltridecyl)-6-chromanol, respectively] 264 and were only detected in *M. trifoliata*: in above ground parts tocopherol- δ 44.1 µg g⁻¹ (SE 41.8) 265 and tocopherol- γ 20.5 µg g⁻¹, and from both original and replicate below ground part samples 266 to copherol- δ 288.4 (SE 41.8) and 100.4 $_{rep}$ $\mu g^{\text{-1}},$ and to copherol- γ 40.2 and 225.8 $_{rep}$ μg $g^{\text{-1}}.$ 267 Triterpenoids were present in *M. trifoliata* and the highest concentration was in *M. trifoliata* 268 below ground parts (788.8 μ g g⁻¹). *Potentilla palustris* above and below ground parts were 269 dominated by β -sitosterol, below ground parts having more than double the concentration as 270 271 the above ground parts (4767.8 µg g⁻¹ and 1945.6, SE 196.4, respectively) (PANGAEA PDI-4071). 272

273 Phytol was recorded in every sample. Above ground parts of both sedges and *M. trifoliata* 274 were dominated by phytol, the concentration being highest in *C. nigra* above ground parts 275 (7510.9 μ g g⁻¹), while below ground parts had a lower concentrations (sedges ca. 7 to 35 μ g 276 g⁻¹; *M. trifoliata* 297.4 μ g g⁻¹).

277

278 *3.4. RDA results*

RDA showed that the variation in each compound type (*n*-alkanes, *n*-alkane ratios, *n*-alcoholsand sterols) was related to plant components (Table 2).

We found that the best result was achieved by combining *n*-alkanes, *n*-alkane ratios and sterols in one analysis. Monte Carlo permutation test of the RDA solution showed that all canonical axes were significant (T 0.245, F 2.597, *p* 0.0020) and the three variables explained 25% of the variance. Analysis separated mosses and vascular plants as their own clusters along the first axis. Vascular plant below ground parts and mosses, however, partly overlapped over axis 1. Mosses formed a more compact cluster than vascular plants that was
also distributed along axis 2. The second axis reflected the differences between the below and
above ground parts; they were separated to the opposite ends of the axis (Fig. 5b.).

We present compounds which explained > 20% of variation detected in plants position in 289 ordination, i.e. 30 compounds (Fig. 5a.). Compounds such as lupeol, 3-stigmastanol and β -290 sitosterol seemed to be descriptive for vascular plants in general. Vascular plant above 291 ground parts were characterized by *n*-alkanes in the range $n-C_{26}$ to $n-C_{29}$, $n-C_{23}/n-C_{21}$ and the 292 phytol concentration (Fig.5a.). The bryophyte cluster seemed to be formed on the basis of 293 ergost-8,24(28)-dien-3 β -ol, obtusifoliol, and *n*-C₂₅/*n*-C₂₉ (Fig.4b.). Some compounds, such as 294 295 the C₂₃ *n*-alkane, several *n*-alkane ratios and brassicasterol, commonly detected in vascular 296 plant below ground parts and in mosses, plotted mid-way between these two groups (Fig.5a, b). The $n-C_{23}/n-C_{25}$ ratio, and lupeol and β -sitosterol concentrations were the main patterns 297 298 describing, and consequently separating, vascular plant below ground parts from mosses (Fig. 5a, b). 299

300

301 **4. Discussion**

Our results support the observation that, by using *n*-alkane ratios, different plant group 302 303 contributions to peat can be separated (Nott et al. 2000, Ishiwatari et al., 2005, Jansen et al., 2006, Nichols et al., 2006, Zheng et al., 2007, Vonk and Gustafsson, 2009, Bingham et al., 304 2010, Andersson et al., 2011). However, they also showed that, when a wider combination of 305 306 plants and peat habitats is included, the absolute values which affect ratios and the relationships between plant types can change. For most of the ratios tested, vascular plant 307 308 above ground parts and mosses were separated as different groups. When the contribution of the vascular plant below ground parts were taken into consideration, the published *n*-alkane 309

310 ratios for bog peat plants were less able to separate vascular plants from *Sphagnum* spp., due to overlap in the distribution patterns between Sphagnum, sedge and M. trifoliata below 311 ground parts. Some ratios, such as $n-C_{23}/n-C_{27}$ (Fig. 4) may potentially be used to separate S. 312 papillosum from other Sphagnum spp. but the probable existence of vascular plant below 313 ground parts in peat might lead to false conclusions about the prevailing vegetation 314 assemblage. The $n-C_{23}/n-C_{25}$ ratio, which has been applied in previous studies (e.g. Bingham 315 316 et al., 2010) as a marker for Sphagnum spp., seems to be effective for the fen environment for separating below ground parts (< 3) from other plants sampled (>3) (Fig. 4). Based on the 317 318 Pwax ratio, it might be possible to separate vascular plant above ground parts with lower values (< 0.9) from below ground parts and Sphagnum spp. (Fig. 4); this agrees with Zheng 319 et al. (2007), who connected high (0.7) P_{wax} values with dry conditions in peat. Thus, where 320 321 the P_{wax} ratio can be measured, we would predict a high contribution of vascular plant above ground material, which is consistent with a drier environment (Strak et al. 2006). 322

Our P_{aq} results agree with Ficken et al. (2000): we found similar values for most of the higher plant above ground parts (< 0.1; PANGAEA PDI-4071), while mosses and below ground parts gave values (0.4-1) close to plants in wet habitats (submerged and floating plants in lake ecosystems) as in Ficken et al. (2000). Due to the high concentration of C_{31} *n*-alkane, the *n*- C_{31}/n - C_{29} and *n*- C_{31}/n - C_{27} ratios show potential for distinguishing *P. palustris* from other species. This corresponds to some previous studies describing *n*- C_{31} as a marker for higher plants (Jansen et al., 2006).

Our results agree with previous studies of bog peats which have shown that LMW *n*-alkanes (*n*-C₂₃ and *n*-C₂₅) are important biomarkers for *Sphagnum* spp. and HMW *n*-alkanes (*n*-C₂₇ to *n*-C₃₁) for above ground parts of vascular plants (Ficken et al., 1998; Baas et al., 2000; Pancost et al., 2002; Nichols et al., 2006). Furthermore, they agree with findings that the *n*- 334 alkane distribution and concentration in vascular plant below ground parts differ from those of above ground parts (Huang et al., 2011, Dawson et al., 2000, Pancost et al., 2002, Jansen et 335 al., 2006). ACL could also be a useful proxy for separating Sphagnum spp. from vascular 336 337 plants in fen environments (Zhou et al., 2005, Andersson et al., 2011). The LMW n-alkane distribution in vascular plant roots has been addressed before and, like studies related to 338 Sphagnum.spp., the dominance of LMW n-alkanes, e.g. n-C₂₃, seems to be related to wet 339 340 environments (Huang et al., 2011, Xie et al., 2004). Huang et al. (2012) concluded that plants growing in water saturated conditions are unlikely to synthesize longer chain *n*-alkanes in 341 342 order to prevent water loss. Thus, the presence of LMW n-alkanes is consistent with the presence of wet conditions. An additional complicating issue in terms of palaeoecological 343 application is that C₂₃ has also been found in significant concentration in *Betula* spp. leaves 344 345 (Sachse et al., 2006).

346 Non species-specific or group-specific *n*-alcohol markers were detected, and the dominant homologue within one group varied. Although *n*-alcohols can be distinguished they have not 347 been shown to have great potential when compared with other biomarkers (e.g. Xie et al., 348 2004). Our study revealed that potential plant group-specific markers may be found among 349 350 sterols such as gramisterol, which was found only in mosses, and tocopherols and schottenol, 351 which were found only in *M. trifoliata*, and lupeol which was not detected in any of the 352 mosses. Otherwise, most of the sterols were commonly present in most of the samples, although concentrations differed considerably, e.g. in the case of β -sitosterol. In agreement 353 354 with Huang et al. (2011) we detected a higher concentration of sterols in the vascular plant below ground parts than in above ground parts. For sterol distributions to be used as a proxy 355 for past vegetation inputs to a fen environment, it is important that either the original sterol or 356 357 corresponding degradation product(s) can be identified within core materials. It has been shown that microbial hydrogenation of sterols within peats can lead to the production of 358

stanols from the original Δ^5 -sterols (e.g. Andersson and Meyers, 2012). It might be expected 359 that with greater degradation of organic matter in a fen environment there will be greater 360 transformation of sterols to stanols. However, if both the sterols and their corresponding 361 stanols can be identified and quantified in a fen core, it may be possible to both assess the 362 degree of organic matter degradation and identify the original vegetation contributions to the 363 peatland. This requires further testing, but our data suggest that, if sterols and stanols are 364 present in peat, they may provide additional information about the contributing vegetation 365 366 (Meyers 2003).

367 The differences detected between mosses and vascular plants, as well as the similarities between Sphagnum spp. and below ground vascular plant parts could spring from the 368 differences in the surrounding hydrological conditions. Mosses and below ground plant parts 369 370 are under the influence of stagnant water in fens, where the water table can be close to the mire surface throughout the growing season (Laine et al., 2012). These plants and plant parts 371 in fens might therefore produce wax with a higher abundance of LMW n-alkanes for 372 protection against micro-organisms and degradation than in moderately drier habitats, i.e. 373 bogs. In the future, one way to study the source of water and the hydrological environment of 374 375 different *n*-alkanes in peat is to examine the δD values of different plant *n*-alkanes (e.g. Xie et al., 2004, Nichols et al., 2010, Garcin et al., 2012). 376

The data presented here shows that there are differences in biomarker distributions between fen plants, but also between species which live in fens and bogs. This means that the application of biomarker distributions from plants living in bog environments to cores from fens could give mis-leading information about past vegetation contributions. In order to apply the data presented here to a fen environment may not be straightforward, however. Although some promising individual biomarkers were found, a better way to identify species and plant 383 groups appeared to be to combine the variables and apply constrained multivariate analysis, such as RDA, as applied here. As a result of the similarity in *n*-alkane distributions, and 384 consequently *n*-alkane ratios, similarities between mosses and vascular plant below ground 385 386 parts remained apparent, but the differences in sterol compositions separated these two groups (Fig. 5b). As a result of our investigations, we would recommend that potential target 387 ratios or markers for down-core analyses might include: (i) lupeol and a high concentration of 388 β -sitosterol, together with n-C₂₃/n-C₂₅ > 3, for prevalence of vascular plant below ground 389 parts; (ii) high $n-C_{23}/n-C_{31}$ value, combined with the presence of obtusifoliol and gramisterol 390 391 to indicate the presence of Sphagnum mosses; and (iii) a high concentration of HMW nalkanes, as in previous studies, for a dominance of vascular plant above ground parts. The 392 degradation of the compounds, especially sterols, has to be considered as they might not be 393 394 preserved in fen environments, due to a high rate of humification or possible transport in the 395 system. A detailed study of this matter is in progress. The effect of peat humification on lipid concentration should also be taken into account by calculating concentration relative to TOC 396 content. This procedure normalizes the results so that different layers with different extent of 397 degradation become more comparable (Meyers 2003; Ortiz et al., 2010). Given the freshness 398 of our samples, this did not impact on our results, but should be considered for palaeo-399 studies. 400

401

402 **5.** Conclusions

We found no clear difference in the sterol composition of the living fen plants but, when comparing *n*-alkanes and their ratios, vascular plant above ground parts could be separated from mosses. However, due to the similar *n*-alkane composition between most of the vascular plant below ground parts and mosses and consequently similar *n*-alkane ratios, separating

407 these two groups from highly humified peat can be challenging. When *n*-alkanes, their ratios 408 and sterols of the plants were compared, together with redundancy analysis, three groups 409 were formed: mosses, above ground and below ground. Thus RDA, or a comparable 410 approach, has potential for also differentiating plant groups in fossil peats. Our results also 411 show that the existing biomarker proxies for peatlands are challenged when a wider 412 combination of plants and peat environments is taken into account.

413

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585 **Captions:**

- Fig. 1. Map of study site. Samples were collected from the fens of the Siikajoki commune
 (64°45´N, 24°42´E), Finland, Northern-Europe.
- 588
- **Fig. 2.** Correlation between concentration $\mu g g^{-1}$ dry wt and $\mu g g^{-1}TOC$. R² values of compared variables: *n*-C₂₃0.99, *n*-C₃₁1, β-sitosterol 0.99, campesterol 0.99.

591

Fig.3. Concentration (μ g g⁻¹ dry wt.) for C₁₇ -C₃₅ *n*-alkanes of plants. Mosses (top), above ground and below ground parts of sedges (middle) and forbes (bottom) are shown. Black bars

represent original samples, samples which were re-analyzed have error bars and sampleswhich were replicated are represented with white bars.

596

Fig. 4. Ratios of *n*-alkanes for mosses and (A) above ground and (B) below ground parts of 597 sedges and forbes: *n*-C₂₃/*n*-C₂₅ (Bingham et al., 2010), C₂₃/*n*-C₂₇, *n*-C₂₃/*n*-C₃₁ (Nott et al., 598 599 2000), n- n-C₂₃/(n-C₂₃ + n-C₂₉) (Nichols et al., 2006), n-C₂₅/n-C₂₉ (Jansen et al., 2006), n-C31/n-C27 (Janssen et al., 2006) and Pwax (Zheng et al., 2007) and ACL of the plant 600 components. Values of $n-C_{23}/n-C_{31}$ for Warn. exannulta, Carex nigra and C. lasiocarpa 601 below ground parts are 0. For ACL, Sphagnum papillosum, Carex rostrata above ground and 602 below ground parts and P. palustris and M. trifoliata below ground parts standard error of 603 mean is < 0.5. 604

605

Fig. 5. RDA (F-ratio 2.597, *p*-value 0.002) shows the distribution of *n*-alkanes, *n*-alkane ratios, sterols and sampled plants (A, above-ground; B, below-ground). Groups: moss, above ground and below ground plant parts, were used as environmental variables. Only compounds with a fit of > 20% are shown (altogether 30 compounds).

610

611 **Table 1**

Studied plants and their status along the nutrient gradient from poor to rich: ombro-, oligo-,
meso-, minero-, eutrophic, and their typical location in microhabitats from dry to wet:
hummock, lawn, flark.

615

616 **Table 2**

- 617 Results RDA with Monte Carlo permutation test to test the significance of plant components:
- 618 mosses, leaves and roots for the variation in compounds a .



Fig 1.







