Diverse organic carbon dynamics captured by radiocarbon analysis of distinct compound classes in a grassland soil

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15 Abstract. Soil organic carbon (SOC) is a large, dynamic reservoir composed of a complex mixture of plant and microbe 16 derived compounds with a wide distribution of cycling timescales and mechanisms. The distinct residence times of individual 17 carbon components within this reservoir depend on a combination of factors, including compound reactivity, mineral 18 association, and climate conditions. To better constrain SOC dynamics, bulk radiocarbon measurements are commonly used 19 to trace biosphere inputs into soils and estimate timescales of SOC cycling. However, understanding the mechanisms driving 20 the persistence of organic compounds in bulk soil requires analyses of SOC pools that can be linked to plant sources and 21 microbial transformation processes. Here, we adapt approaches, previously developed for marine sediments, to isolate organic 22 compound classes from soils for radiocarbon $({}^{14}C)$ analysis. We apply these methods to a soil profile from an annual grassland 23 in Hopland, California (USA) to assess changes in SOC persistence with depth to 1 m. We measured the radiocarbon values 24 of water extractable organic carbon (WEOC), total lipid extracts (TLE), total hydrolysable amino acids (AA), and an acid-25 insoluble (AI) fraction from bulk and physically separated size fractions ($<2 \text{ mm}, 2 \text{ mm}-63 \mu\text{m}$, and $<63 \mu\text{m}$). Our results 26 show that Δ^{14} C values of bulk soil, size fractions, and extracted compound classes became more depleted with depth, and 27 individual SOC components have distinct age-depth distributions that suggest distinguishable cycling rates. We found that AA 28 and TLE cycle faster than the bulk soils and the AI fraction. The AI was the most ¹⁴C depleted fraction, indicating it is the 29 most chemically inert in this soil. Our approach enables the isolation and measurement of SOC fractions that separate 30 functionally distinct SOC pools that can cycle relatively quickly (e.g., plant and microbial residues) from more passive or inert 31 SOC pools (associated with minerals or petrogenic) from bulk soils and soil physical fractions. With the effort to move beyond 32 SOC bulk analysis, we find that compound class ¹⁴C analysis can improve our understanding of SOC cycling and disentangle the physical and chemical factors driving OC cycling rates and persistence. 33

34 **1 Introduction**

35 Soil organic carbon (SOC) is a large and complex terrestrial reservoir of Earth's organic carbon (OC) (Jobbágy and 36 Jackson, 2000). It is a highly dynamic and open pool with inputs from decaying plant material, living roots, and soil microbes, 37 and with losses driven by microbial activity that includes the degradation and transformation of compounds (Angst et al., 38 2021). The result of these processes is a heterogenous mixture of organic compounds with different radiocarbon $({}^{14}C)$ ages 39 and reactivities (Lehmann and Kleber, 2015; Shi et al., 2020; Trumbore and Harden, 1997; Gaudinski et al., 2000; McFarlane 40 et al., 2013). This complexity obscures the mechanisms that control overall OC persistence in soils, resulting in a continued 41 debate over the degree to which environmental factors, physical protection, and chemical composition influence SOC reactivity 42 and persistence (Lützow et al., 2006; Lehmann et al., 2020; Schmidt et al., 2011).

43 Bulk analysis methods do not satisfactorily demonstrate how physical protection and chemical composition interact to 44 influence SOC persistence, and so novel organic matter characterization methods can shed light on how different compound 45 classes of OC are preserved in soils and through what mechanisms. For example, we need to understand how the chemical 46 structure of OC influences interactions with mineral surfaces, such as aggregation or sorption, as well as how the environment 47 influences the decomposition and resource availability of certain OC compounds and functional groups (Lehmann and Kleber, 48 2015; Schmidt et al. 2011; Kleber et al., 2021). However, it has been difficult to isolate, identify, and quantify pools of OC 49 that directly link towithout altering OC molecular chemistry intact or undisturbedsitu OCOC compounds within the 50 soilchemical compounds (Von Lutzow et al., 2007). Thus, specific organic compounds isolated from soils, such as amino acids 51 and lipids (Rethemeyer et al., 2004), can provide information on how OC is stabilized in different environments. Therefore, 52 multiple approaches, such as a physical separation followed by a chemical separation, -are needed to fully understand the 53 interplay between chemical compound reactivity and how carbon-mineral interaction functions as part of SOC persistence in 54 soil.

55 One approach used to investigate the controls on SOC persistence is to separate soil into operationally defined carbon 56 pools (e.g., size or density fractions) and characterize the resulting fractions. This approach has demonstrated that association 57 of OC with soil minerals is a critical mechanism for C stabilization (Vogel et al., 2014; Mikutta et al., 2007), as ¹⁴C data 58 indicate that some mineral-associated C can persist for thousands of years (Torn et al., 2009). However, ¹³C labelling 59 experiments show that some mineral-associated C cycles quickly, within months to years (Keiluweit et al., 2015; De Troyer 60 et al., 2011). Some biomolecules form strong associations with mineral surfaces, such as long-chain lipids with iron oxides 61 (Grant et al., 2022), while other compounds only loosely associate with minerals such as through hydrophobic interactions 62 with other OC compounds (Kleber et al., 2007). Therefore, physically isolated mineral-associated OC is still a heterogenous 63 mixture of OC molecules that have a distribution of turnover times, rather than a single homogenous and intrinsically stable 64 SOC pool (Stoner et al., 2023; Van Der Voort et al., 2017).

65 Another approach that can yield finer resolution of OC turnover than traditional techniques is to isolate and measure the

66 isotopic signature of specific compounds (Von Lutzow et al., 2007). In marine, riverine, and lacustrine systems, compound 67 specific radiocarbon analysis (CSRA) has been used monitor the degradation of organic carbon through the marine water 68 column (Loh et al., 2004), characterize marine particulate OC (Hwang and Druffel, 2003), constrain terrestrial OC burial and 69 export from river systems (Galy et al., 2015; Galy et al., 2008; Repasch et al., 2021, Smittenberg et al., 2004), and determine 70 effect of OC export and burial on precipitation patterns and climate (Hein et al., 2020; Eglinton et al., 2021). Different types of compounds including plant or microbial lipid biomarkers (Douglas et al., 2018; Huang et al., 1996), amino acids (Bour et 71 72 al., 2016; Blattmann et al., 2020), lignin (Feng et al., 2017; Feng et al., 2013), certain carbohydrate compounds (Kuzyakov et 73 al., 2014; Gleixner, 2013), and pyrogenic or black carbon (Coppola et al., 2018) can be isolated and analysed for ¹⁴C leading 74 to a more detailed understanding of the cycling of targeted compounds in the environment.

Each of these specific compounds can provide information related to the persistence, source, and potential fate of OC in soils. For instance, lipids are found in plant cell walls and microbial cell membranes and are used for energy storage. Amino acids are necessary for protein formation, are enriched in nitrogen relative to other plant and microbial residues, and likely play an important role in nitrogen mining and recycling. These two compound classes have diverse chemical reactivities which allows for insight into chemical compound persistence. Understanding the abundance and age of these two biomarkers in soils can help differentiate the source of C used by soil microbes for metabolism and growth (e.g., new C inputs vs older, recycled soil C) as well as the transformation pathways that yield persistent SOC.

82 Recently, CSRA approaches developed for these environments have been applied to soil showing promise for identifying 83 distinct ages of plant and microbial biomarkers in SOC (Gies et al., 2021; Grant et al., 2022; Van Der Voort et al., 2017; Jia et al., 2023; Douglas et al., 2018). Most of these CSRA studies applied to SOC have targeted specific, individual biomarkers 84 85 in soils, which generally contribute less than 5% of the entire carbon pool (Lützow et al., 2006; Kögel-Knabner, 2002). This 86 approach can be too specific to elucidate holistic mechanisms for SOC persistence and turnover that pertain to the majority of SOC. While individual biomarker ages, such as single ages of a particular lipid or single amino acid, can be useful in some 87 88 contexts, comprehensive understanding of carbon compound class persistence is vital for understanding and modelling the 89 vulnerability of soil carbon to degradation.

90 To strike a balance between too specific and too broad, some researchers have characterized broader compound classes rather than isolating a single biomarker. For example, this ¹⁴C-compound class approach has been applied to marine dissolved 91 92 and particulate OC with a range of compounds, such as total lipids and total amino acids, to provide a broader understanding 93 of OC persistence in oceans (Wang et al., 2006; Wang et al., 1998; Wang and Druffel, 2001; Loh et al., 2004). Wang et al. 94 (1998) established a sequential extraction procedure to analyse 14 C abundance of total lipids, amino acids, carbohydrates, and 95 a residual acid insoluble fraction from marine POC and sediments. This approach yielded distinct differences in ¹⁴C age and 96 abundance of the amino acids, lipids, and the acid insoluble fraction in POC from the marine water column and sediment, as 97 well as in coastal versus open ocean environments. Loh et al. (2004) found the lipid fraction of dissolved OC and POC to be 98 the oldest fraction measured in both the Atlantic and Pacific oceans, while the acid insoluble fraction was intermediate in age, 99 and the amino acids and carbohydrates contained a significant contribution of modern carbon. Wang and Druffel (2001) also 100 used this approach and found that the lipids were the oldest compound class from sediments in the Southern Ocean, but the 101 acid insoluble residue was very similar in age to the lipid fraction. These studies suggest that compound classes can have 102 independent cycling rates, but these cycling rates can be influenced by the environment.

103 Here, we apply a ¹⁴C compound class approach to soils to more broadly understand SOC turnover mechanisms. We 104 characterize the distribution and ¹⁴C age of multiple SOC pools with depth in a well-studied annual grassland in California, 105 using soil physical fractionation (McFarlane et al., 2013; Poeplau et al., 2018) and modified compound class extraction 106 methods previously detailed for marine sediments (Wang et al., 1998). We measured the radiocarbon values of water 107 extractable organic carbon (WEOC), total lipid extracts (TLE), total hydrolysable amino acids (AA), and an acid-insoluble 108 (AI) fraction from bulk and physically separated size fractions (bulk soil, sand, and silt+clay). We expected the TLE to be 109 older than its source fraction (bulk soil, sand, or silt+clay), to be older with depth as the decline in plant inputs necessitates 110 recycling and use of older SOC, and to be older in the silt+clay fraction as its high surface area should result in mineral-OC 111 associations that protect SOC from soil microbes- (Grant et al., 2022; Van der vort et al., 2017). We expected the AA to evelo 112 fasterbe younger than the TLE fraction and the bulk SOC pool based on the young ¹⁴C ages found for AA extracted from in 113 marine sediments (Wang et al., 1998; Wang and Druffel, 2001), but hypothesized that recycling of amino acids at depth by 114 soil microbes might result in an increase in the age of AA below 50 cm. Finally, we expected AI to have old C, similar to the 115 TLE, as seen found in marine sediments (Wang et al., 1998). Here, we describe the relative abundance and radiocarbon content 116 of WEOC, total lipid, and acid insoluble compound class extracts in bulk soils and compare carbon storage 117 and cycling rates within soil size fractions. - These data provide a foundation for the continued application of compound class ¹⁴C work to the understanding and modelling of soil OC persistence. 118

119 2 Materials and Methods

120 **2.1 Site and Sample Description**

121 Soil samples were collected from the University of California's Hopland Research and Extension Center (HREC) in January 122 2022. The site is an annual grassland with a Mediterranean-type climate, where the mean annual precipitation (MAP) is 940 123 mm per year and mean annual temperature is 15°C (Nuccio et al., 2016). The underlying geology consists of mixed sedimentary 124 rock of the Franciscan formation. The soils are designated Typic Haploxeralfs of the Witherall-Squawrock complex (Soil 125 Survey Staff, 2020). The samples were collected from the "Buck" site (39.001°, -123.069°) where the vegetation is dominated 126 by annual wild oat grass, Avena barbata (Kotanen, 2004; Bartolome et al., 2007). Soils were collected from a freshly dug soil 127 pit at four depths: 0-10 cm, 10-20 cm, 20-50 cm, and 50-100 cm. The site is dominated by annual grasses, shallow rooted 128 herbs, and forbs, and we did not observe roots below 10 cm. Thus, root derived inputs of OC are important near the soil surface.

but do not directly affect deeper soils at this site. Samples were stored in sealed plastic bags at ambient temperature and transported to the laboratory in Livermore, CA. Soil samples were air dried, homogenized, and sieved to 2 mm, with the >2 mm fraction retained for further analysis. Samples were subdivided for soil characterization, physical size separations, chemical compound extractions, and density fractionation.

133 2.2 Physical Fractionation

134 To compare compound classes between mineral-associated OC and mineral-free OC, we used a salt-free and chemical-free 135 method for isolating the mineral-associated organic matter from the free particulate organic matter (Fig. 1a). Under the 136 assumption that mineral-associated carbon is primarily found in the silt+clay (<63 µm) particle size fraction, we used a size 137 fractionation sieving method where air-dried samples were dry-sieved into three size fractions: bulk soil (<2 mm), sand (2 mm 138 - 63 μm), and silt+clay (<63 μm) (Lavallee et al., 2020; Poeplau et al., 2018). Additionally, because the majority of free 139 particulate organic carbon (POC) is contained in the sand faction, we used a "water density" separation to remove the low 140 density POC from the mineral matter in this fraction, by suspending the sand fraction in 18.2 M Ω and removing the floating 141 OC,- resulting in a POC (<1g mL⁻¹) fraction and a POC-free (>1g mL⁻¹) sand fraction.

To further characterize these soils and aid in interpretation of our data, we compared the size fractionated samples to samples separated by density using sodium polytungstate (SPT-0 adjusted to a density of 1.65 g ml⁻¹) (Poeplau et al., 2018) (see SI Section 1.1 for detailed methods). We chose to focus our compound class extraction efforts on size fractionated samples to avoid chemical alteration of SOC during exposure to SPT, since SPT has a high ionic strength and low pH.

- 146 To constrain any contributions of <u>OC from</u> parent materials to SOC, we processed and analyzed the rock fraction (> 2mm) 147 (Agnelli et al., 2002; Trumbore and Zheng, 1996). Rocks were washed with 18.2 M Ω water in an ultrasonic bath to remove 148 surface contamination, rinsed with 1N HCl to remove any additional weathered material loosely adhered to the surface, dried 149 at 60°C, then manually crushed.
- 150 A large, representative aliquot (~10 g) of the bulk and each physical fraction were ball milled and measured for total organic
- 151 carbon (TOC, wt %), C/N ratio, δ^{13} C and Δ^{14} C (Section 2.6). In addition, we analyzed the bulk soils at each depth with nuclear
- 152 magnetic resonance (¹³C NMR) to assess the broad structural complexity of the OC in the bulk soil (SI Section 2).
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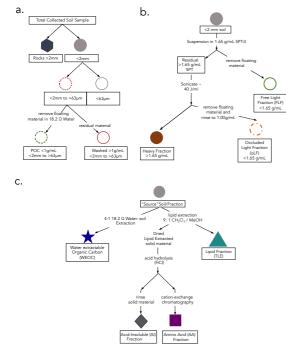




Figure 1: Schematics of protocols used in this study for a) fractionation by size, b.) density separation (details in SI methods), and c) extraction of targeted compound classes. The "<u>sourceparent</u> soil/<u>fraction</u>" refers to the soil from which the different compound classes are extracted. All compound extractions and physical fractionations were applied to the <2 mm bulk soil; total lipid extract (TLE), amino acid (AA), and acid insoluble (AI) compound classes were also extracted from the silt+clay fraction; and only the TLE was extracted from the dense fraction (DF).

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161 **2.3 Water-extractable organic carbon (WEOC)**

162 The water-extractable organic carbon (WEOC) fraction was collected from 80 g of bulk soil with 18.2 M Ω water using a 4:1 163 water to soil ratio (Van Der Voort et al., 2019; Lechleitner et al., 2016; Hagedorn et al., 2004). Saturated soil samples were 164 shaken for 1 hour and then filtered through a pre-rinsed 0.45 µm polyethersulfone (PES) Supor filter under vacuum. An aliquot 165 was taken for dissolved organic carbon (DOC) measurement on a Shimadzu TOC-L combustion catalytic oxidation instrument. 166 Sample concentrations were determined using a nine-point DOC calibration curve ranging from 0–200 mgC L⁻¹. The WEOC 167 fraction was dried using a Labconco CentriVap centrifugal drying system at 40°C and subsequently transferred with 0.1N HCl 168 into pre-combusted quartz tubes to eliminate any inorganic carbon dissolved in the aqueous fraction. The acidified WEOC 169 fractions were then dried down using the CentriVap. Dried samples were flame sealed under vacuum (Section 2.6) for 170 subsequent carbon isotope analyses.

171

172 **2.4 Total Lipid Extraction (TLE)**

173 Total lipids (TLE) were extracted from the soil samples using an Accelerated Solvent Extraction (ASE) system (Dionex 350, 174 Thermo Scientific) in duplicate. The TLE was extracted from the bulk, sand, silt+clay, and the dense fraction (> 1.65 g ml⁻¹; 175 DF). An aliquot of 10-30 g of soil was loaded into a stainless-steel ASE extraction cell depending on TOC content (Rethemeyer 176 et al., 2004). The ASE was set to extract the sample for 5 minutes with a holding temperature of 100°C at 1500 PSI. Lipids 177 were extracted using a 9:1 ratio of dichloromethane (DCM or syn: methylene chloride) to methanol (Wang et al., 1998; Van 178 Der Voort et al., 2017; Grant et al., 2022). The TLE was dried under constant ultra-pure N₂ flow at 40°C using a nitrogen dryer 179 (Organomation Multivap Nitrogen Evaporator). The TLE was resuspended in ~5ml of 9:1 DCM:Methanol then transferred to 180 pre-combusted quartz tubes, dried again, and analyzed for ¹⁴C as described below (Section 2.5). Total CO₂ produced by the combustion of the TLE was measured manometrically on the ¹⁴C vacuum lines during graphitization. Process blank samples 181 182 were analyzed with each batch (SI Section 3.1).

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184 2.5 Amino Acid (AA) Extraction

185 Amino acids (AA) were extracted from the lipid-extracted residual bulk and silt+clay size fraction with an acid hydrolysis 186 procedure, desalted, and isolated with cation exchange chromatography using methods modified from those used in marine 187 systems (Wang et al., 1998; Ishikawa et al., 2018; Blattmann et al., 2020). Briefly, a 500 mg soil aliquot was hydrolyzed with 188 6N HCl (ACS grade) under an N₂ atmosphere for 19-24 hours at 110°C. After hydrolysis, amino acids in solution were 189 separated from the solid acid insoluble (AI) fraction via centrifugation for 5 minutes at 2500 rpm. The AI fraction was 190 subsequently washed at a minimum three additional times with 0.2N HCl to ensure complete AA recovery. The supernatant 191 was collected in a single pre-combusted vial and then filtered through a pre-combusted quartz wool fiber plug to remove 192 extraneous sediment particles. The filtered hydrolysate was dried using a CentriVap at 60°C for 4 hours. The dried supernatant 193 was redissolved in 1 ml 0.1N HCl and loaded onto a preconditioned resin column (BioRad 50WX8 200-400 mesh resin) to 194 isolate the AA from other hydrolyzed organic matter and remove excess chloride. Details of the procedure can be found in 195 Ishikawa et al., 2018. Briefly, once the sample was loaded on the column, it was rinsed with three bed volumes (~6 ml) of 18.2 196 $M\Omega$ H₂O. The free AA were eluted with 10 ml of 2N ammonium hydroxide (NH₄OH), then transferred into pre-baked quartz 197 tubes, dried at 60°C in the CentriVap, and finally sealed and combusted for isotopic analysis. The remaining rinsed solid 198 residual after hydrolysis is the acid-insoluble (AI) fraction. These are processes as a solid sample for isotopic analysis.

199

200 **2.6 Isotopic and elemental analysis**

All samples were analyzed for radiocarbon (¹⁴C) at the Center for Accelerator Mass Spectrometry (CAMS) at Lawrence Livermore National Lab (LLNL) in Livermore, California. Samples were either measured on a 10 MV Van de Graaf FN or 1MV NEC Compact accelerator mass spectrometer (AMS) (Broek et al., 2021), with average errors of $F^{14}C = 0.0035$. For

204 solid soil analysis, 10 to 250 mg of ground material was weighed into a pre-combusted quartz tubes along with 200 mg CuO 205 and Ag, flame sealed under vacuum, then combusted at 900°C for 5 hours. The CO₂ was reduced to graphite on preconditioned 206 iron powder under H₂ at 570°C (Vogel et al., 1984). Measured ¹⁴C values were corrected using δ^{13} C values and are reported as age-corrected Δ^{14} C values using the following the conventions of Stuiver and Polach (1977). Extraneous C was quantified 207 208 for the TLE and AA extractions (SI Table 4 and SI Section 3). For ease of reference, we included conventional radiocarbon 209 ages in our figures and tables. We quantified turnover times using the single pool turnover model described in Sierra et al. 210 (2014) and Van Der Voort et al. (2019) and explained in detail in Trumbore (2000) and Torn et al. (2009). This approach generates two solutions for pools with Δ^{14} C > 0 ‰, one corresponding to each side of the atmospheric ¹⁴C-CO₂ curve over the 211 212 last 70 years (Hua et al., 2022). Unfortunately, we cannot identify the correct solution (McFarlane et al., 2013; Trumbore, 213 2000), especially for TLE and AA fractions from the top 20 cm, as we do not have multiple time points or additional constraints 214 such as pool-specific input or decomposition rates (see Section 2.8). Therefore, our data analysis and interpretations rely on 215 the reported Δ^{14} C values. All individual ¹⁴C measurements used in this study are listed in the Supplementary Information (SI 216 Table 1 and 2).

For each solid sample, a dried homogenized aliquot was analyzed for TOC concentration and δ^{13} C using an elemental analyzer (CHNOS) coupled to an IsoPrime 100 isotope ratio mass spectrometer at the Center for Stable Isotope Biogeochemistry (CSIB) at the University of California, Berkeley. Samples are assumed to have no inorganic carbon based on acid leaching tests and previously published ¹⁴C work at this site (Finstad et al, 2023, Foley et al., 2023). δ^{13} C was measured in duplicate for each solid sample and errors represent the standard deviation of the mean. δ^{13} C values of WEOC, TLE, and AA extracts were measured on a split of the cryogenically purified CO₂ and were analyzed at the Stable Isotope Geosciences Facility at Texas A&M University on a Thermo Scientific MAT 253 Dual Inlet Stable Isotope Ratio Mass Spectrometer (SI Table 1).

224

225 2.7 Data analysis

Data was analyzed using MATLAB version R20223 and R v. 3.614 (R Core Team, 2019). Linear regressions were calculated between the sample depth mid-point and Δ^{14} C values from both the size fractions as well as the extracted compounds (WEOC, TLE, AA, AI) from the different size fractions. This was done to directly compare the difference in Δ^{14} C value between the compound classes. Correlation coefficients, p-values and r² are provided in SI Table 3. Analysis of Variance (ANOVA) was used to assess differences in Δ^{14} C with depth, between TLE and AA, and between soil fractions. ANOVA tests were performed in R v. 3.614 (R Core Team, 2019). In the text, results are reported as means followed by one standard error when n = 2 or 3 or by analytical error when n = 1.

233

234 **2.8 Interpretation of radiocarbon data**

- -In the interpretation of soil ¹⁴C activity, we must consider how ¹⁴C created during atmospheric nuclear weapons may 235 have affected the isotopic signatures of SOC at our study site. Significantly elevated "bomb" derived ¹⁴C was released into the 236 237 environment during atmospheric nuclear weapons testing during the mid-20th century. This atmospheric radiocarbon spike has 238 been continuously incorporated into carbon reservoirs including vegetation, soils, and oceans (Levin and Hessshaimer, 2000). 239 Plants assimilate CO_2 with the ¹⁴C signature of the current year's atmosphere during photosynthesis and thus incorporate the current atmospheric ¹⁴C signature into their tissues and root exudates. This signature then cycles into and through soils as this 240 241 plant-derived organic matter decays, is processed by microbes, and enters stable soil organic matter pools (Torn et al. 2009). 242 Since the termination of atmospheric weapons testing in the 1960s and with continued fossil fuel emissions, the ¹⁴C of 243 atmospheric CO₂ has decreased to approximately pre-1950 values with $0 \pm 1\%$ reported for the 2019 Northern Hemisphere growing season (Hua et al. 2022). Thus, soil carbon pools with ¹⁴C signatures above 0% can be interpreted as decadal-aged or 244 245 decadal cycling C and pools with ¹⁴C signatures below 0‰ cycle on century to millennial timescales.
- 246

247 **3 Results**

248 **3.1 Radiocarbon values and characterization of the physical fractions**

249 We used soil size and density fractionation to separate the bulk soil into fractions with different degrees of mineral protection. Radiocarbon content for the bulk soil, sand, and silt+clay (SI Table S3) became more ¹⁴C depleted (older) with increasing 250 251 depth (Table 1, Fig. 2). SOC in the silt+clay was consistently younger than in the bulk soil, with the average difference in Δ^{14} C 252 values increasing from 4‰ at the surface to 87‰ at depth. In the sand fraction, the Δ^{14} C values of POC were consistently near current atmospheric values ($2 \pm 3\%$) and were not significantly correlated with depth. In contrast, the Δ^{14} C values of the POC-253 254 free sand-sized fraction declined with depth ($25 \pm 3\%$ to $-510 \pm 2\%$, p = 0.006) and were indistinguishable from the POC-255 free sand fraction (Fig. 2). Density fractionation of the bulk soil resulted in most of the sample mass (> 98%) and OC (75-256 83%) recovered in the DF at all depths (SI Fig. S2).

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L	Э	1

	bu	lk (<2mm)	sand-sized (2mm to 63µm)							silt+clay (<63µm)			
						POC-free >1g mL ⁻¹		POC <1 gmL ⁻¹					
Depth	%OC	Δ ¹⁴ C ± err (‰)	mass f	%OC	Δ ¹⁴ C ± err (‰)	%OC	Δ ¹⁴ C ± err (‰)	%OC	Δ ¹⁴ C ± err (‰)	mass f	%OC	Δ ¹⁴ C ± err (‰)	
0-10 cm	3.14	31 ± 3	0.71	2.68	25 ± 3	2.08	25 ± 3	25.69	19 ± 3	0.29	4.25	34 ± 3	
10-20 cm	1.22	-22 ± 3	0.69	0.94	-38 ± 3	0.77	-35 ± 3	25.99	-5 ± 3	0.31	1.84	-13 ± 3	
20-50 cm	0.50	-116 ± 3	0.75	0.39	-142 ± 3	0.38	-149 ± 2	n.m.	4 ± 3	0.25	0.85	-79 ± 3	
50-100 cm	0.25	-468 ± 3	0.79	0.23	-496 ± 3	0.18	-510 ± 2	n.m.	-10 ± 3	0.21	0.35	-380 ± 3	

258 Table 1. Carbon concentrations, mass fractions, and radiocarbon values for the size separations from the Buck Pit

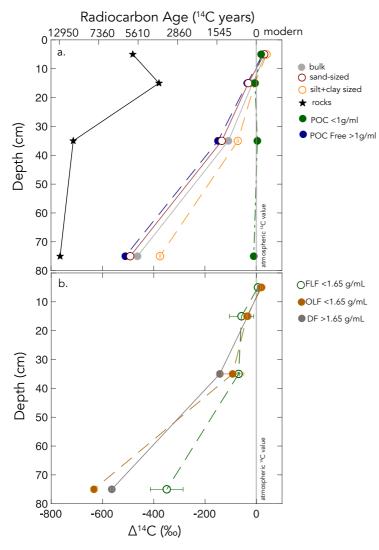
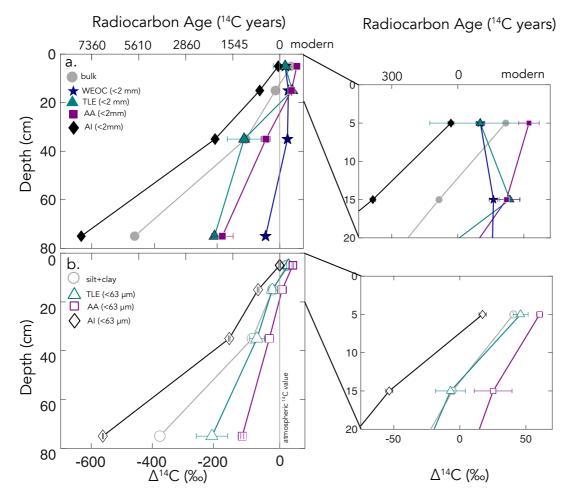




Figure 2: Δ^{14} C values by depth for a) size-fractions. b) density-fractions from the Buck soil pit. Conventional ¹⁴C ages are provided for reference. <u>The following abbreviations appear in the legend: particulate organic carbon (POC), free light fraction (FLF),</u> occluded light fraction (OLF), and dense fraction (DF).

3.2 Compound Class results from bulk soil and silt+clay

- In both the bulk soil and silt+clay fraction, the extracted compound classes became ¹⁴C-depleted with depth except for the WEOC, which had ¹⁴C values that reflected C inputs recently fixed from the atmosphere throughout the soil profile (Fig. 2; SI tables). The Δ^{14} C values of the WEOC ranged from 14 ± 4‰ at the surface to -46 ± 4‰ at depth, and the DOC concentrations ranged from 43.2 to 6.7 mg C g soil⁻¹ at the surface and at depth, respectively.
- The TLE from the bulk soil had Δ^{14} C values that range from 17 ± 27 to $-208 \pm 6\%$ (n = 2; \pm SE) in the surface and deepest sample, respectively. In comparison, the TLE from the silt+clay fraction was modern at the surface and became more ¹⁴C depleted with depth (p < 0.001), from 46 ± 4 to -204 ± 36 ‰. The slopes of the linear regressions of Δ^{14} C with depth were indistinguishable in TLE from the bulk soil and silt+clay. In addition, the TLE from the bulk TLE and silt+clay fraction TLE (SI Tables) had very similar Δ^{14} C values, but the bulk soil had less lipid-C extracted during each experiment (280 µg g C⁻¹ in the 0–10 cm vs. 150 µg g C⁻¹; SI Table 2).
- 279 The Δ^{14} C values of the AA extracted from the bulk soil ranged from 54 ± 5 to -183 ± 24 (n = 2, SE) with depth (Fig. 280 3, SI Table S3). Similarly, the Δ^{14} C value of the AA fraction extracted from silt+clay declined with depth from $60 \pm 3\%$ (n = 281 2, SE) at the surface to -106 ± 4 % (n = 2, SE) at 50–100 cm depth. The slopes of the AA extracted from the bulk and silt+clay-282 size fractions were statistically different, indicating that the AA extracted from the bulk soil became more depleted with depth 283 than that extracted from the silt+clay (SI Table S3). Furthermore, AA fractions were enriched in ¹⁴C relative to the TLE or AI 284 fraction (p < 0.01 for bulk soil and p < 0.05 for silt+clay). The AI fraction was the oldest fraction found in our study at each 285 depth. The Δ^{14} C values of the AI fraction ranged from $-5 \pm 2\%$ to $-633 \pm 2\%$ (analytical error, n=1) and declined with depth 286 (p < 0.01) for bulk soil and silt+clay (Fig. 3; SI Table S3).



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Figure 3: a) Δ^{14} C by depth for bulk soil and four compound class fractions extracted from bulk soil for the entire depth profile with the inset of the top 20 cm. b)) Δ^{14} C by depth for the silt+clay (<63 µm) fraction and three compound classes extracted from the silt+clay for the entire depth profile with the inset of the top 20 cm. For total lipid extract (TLE) and amino acid (AA) fractions (n=2) and error bars represent the standard error from duplicate measurements. For the <2 mm, water extractable organic carbon (WEOC), and acid insoluble (AI) fractions (n=1) and error bars represent analytical error. Error bars are smaller than the marker width where not shown.

294 295

296 4 Discussion

297 4.1 Variability of ¹⁴C in compound classes in soils

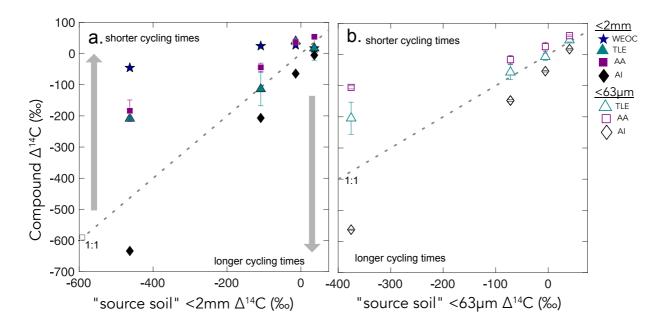
We measured radiocarbon content of four distinct soil chemical extracts: water extractable organic carbon (WEOC), total lipid extract (TLE), free amino acids (AA), and the acid insoluble fraction (AI), each of which had distinct Δ^{14} C values compared to the <u>sourceparent</u> soil it was extracted from (bulk or silt+clay; Fig. 4a and 4b). The central questions of this study 301 are: What are the differences in cycling time/age between various organic compounds in the soil? Do these differences in 302 cycling time change with depth? As expected, Δ^{14} C values of TLE, AA, and AI became more depleted with depth (Fig. 2). 303 More interestingly, the differences between the ¹⁴C content of parent-source soil and the extracted compounds were not consistent with depth (Fig. 3a and 3b). This divergence in Δ^{14} C values reflects differences in turnover times among compound 304 305 classes, which can be influenced by the sources of OC to each of these pools and by differences in the stabilization mechanisms 306 protecting those compounds from decay. In this annual grassland, plant inputs should have a greater influence on SOC pools 307 near the surface, which we confirmed with near modern Δ^{14} C signatures in the 0–10 cm depth for all compound classes and 308 size fractions (Fig. 3b and 3c). Furthermore, at deeper depths, new vegetation inputs should be less readily available, which 309 results in more depleted Δ^{14} C signatures at depth and could necessitate microbial use and recycling of older SOC.

310 We found that, averaged across depths, the Δ^{14} C values of the TLE were more depleted than those of the AA, though both 311 compound classes were more enriched in Δ^{14} C than the bulk soil or silt+clay from which they were extracted. The extracted 312 AAs are the foundational units of hydrolysed proteins and found in both plant and microbial biomass (Blattmann et al., 2020). 313 As in marine studies, we found the AAs to be the youngest compound class fraction (of the TLE and AI) in these soils. The 314 AA pool likely reflects a more actively cycling microbial pool especially at depth, as AA are enriched in nitrogen compounds 315 and likely microbes are both preferentially mining and recycling these compounds (Moe, 2013). The divergence from bulk ¹⁴C 316 values indicate that even at depth in the soil, the AAs are either continuously replenished from transport of AAs from surface 317 horizons or re-synthesized with relatively ¹⁴C enriched sources such as the WEOC.

318 Based on published data for both soils and marine sediments, we expected the TLE to be older than both the AAs and the bulk soil, however we found that all TLE samples, no matter what fraction we measured, were more ¹⁴C enriched than the bulk 319 320 soil. TLE is composed of a continuum of lipids from plant and microbial materials, ranging from leaf waxes to microbial cell 321 structural components (Angst et al., 2021; Angst et al., 2016), that cycle at different rates and likely interact with mineral 322 surfaces. Previous studies where individual lipid biomarker Δ^{14} C values were measured in soils on either short chain or long 323 chain fatty acids found a divergence in Δ^{14} C values between these two pools, with short chain lipids generally having enriched ¹⁴C values and long chain lipids having more depleted ¹⁴C values (Grant et al., 2022; Van Der Voort et al., 2017). For example, 324 325 long-chain lipid biomarkers, primarily thought to be plant derived, had consistently older ¹⁴C ages than bulk soil (Van Der 326 Voort et al., 2017). Short-chain lipids, which can be microbial or root derived (Rethemeyer et al., 2004), were found to be 327 younger than long-chain lipids throughout the soil profiles and younger than bulk soil at depth (Van Der Voort et al., 2017). 328 However, microbial cell wall lipid biomarkers (glycerol dialkyl glycerol tetraethers, GDGTs) had older ¹⁴C ages than bulk 329 soils (Gies et al., 2021). With this consideration, our result of more enriched ¹⁴C of the TLE could be an indication of a 330 predominance of short chain lipids and suggested higher abundance of microbially-derived lipids than plant-derived lipids. 331 However further study of specific lipid abundance (e.g., *n*-alkanes, fatty acids) in these soils are necessary, as it is unclear to 332 what degree lipids are older than bulk soils with depth because of preservation of these compounds through mineral association 333 or because of microbial use of aged OC sources for growth.

334 We found that AI, the residual sample after both the TLE and AA have been extracted (Wang et al., 1998; Wang et al., 335 2006), was the most ¹⁴C depleted OC fraction measured at each soil depth (Fig. 3, 4) The AI fraction was far more depleted 336 relative to the bulk soil (Fig. 3a and 4a) than observed in marine studies with acid-insoluble OC (Wang et al., 2006; Wang and 337 Druffel, 2001). In these marine studies, the ¹⁴C of the AI varied in age depending on sampling depth and location. The 338 significant depletion of the AI in our soils suggests that these chemically stable compounds are not oxidized in soil. 339 Importantly, our AI samples are older than the other chemical and physical soil fractions that we measured in the soil, 340 consistent soil, consistent with the general expectation that aromatic compounds can be difficult to degrade in in soils (Ukalska-341 Jaruga et al., 2019).

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Figure 4. Δ^{14} C values of the three extracted compound classes, the water extractable organic carbon (WEOC), the total lipid extract (TLE), the amino acid (AA) fraction, and the acid insoluble (AI) fraction, (y-axis) compared to the Δ^{14} C values of the parent or source soil/fraction (x-axis) for a) bulk soil and b) silt+clay. The grey dashed lines show the 1:1 line where bulk sample Δ^{14} C equals compound class Δ^{14} C. Gray arrows point to regions where data plot above or below the 1:1 line, suggesting that a given compound class has shorter and longer carbon turnover times than bulk soil, respectively.

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350 4.2 Differential OC cycling between the different "parent" fractions fractionation methods

Our results suggest different OC cycling timescales for the different physical fractions representing the "<u>sourceparent</u>"
 fractions. Here, we focus on the silt+clay fraction as an operationally defined mineral-associated OC pool. Numerous soil

353 physical fractionation schemes have been applied to soils and disparities in methods challenge interpretation and 354 intercomparison of results from different studies using different approaches. We compared the size-based soil fractionation to 355 the density fractionation to aid in interpretation and comparability of our findings to other studies. Our silt+clay fraction had 356 higher Δ^{14} C values than the sand, POC-free sand, and the dense fraction (DF)DF. Our silt+clay fraction could include free 357 organic matter that passed through the 63 µm sieve but that would have floated off the DF during density fractionation. For 358 reference, the free light fraction (FLF) has higher Δ^{14} C values than the mineral-associated pools and bulk soils (Fig. 5), but 359 also has high C:N reflecting the high OC content and dominantly plant origin of this fraction (SI Table S1). We assume that 360 this small-size free OC is a small fraction of the total silt+clay OC as no small fragments of organic matter were visible and 361 because the C:N ratios of the silt+clay fractions are only slightly elevated compared to the bulk soil and sand fractions (SI 362 Table S1). Rather, the silt+clay fractions may have higher Δ^{14} C values relative to the POC-free sand and bulk soil because 363 higher surface area in the silt+clav may facilitate mineral association with surface derived OC-with minerals (e.g., from the 364 WEOC fraction).

365 Additionally, our TLE comparison between different size and density fractions highlights the important influence that 366 method selection has over experimental results. Across studies, the mineral-associated OC is not a uniformly defined pool, 367 and the observed results are a consequence of the methodology used to separate the samples (Fig. 6). The mineral-associated 368 TLE cycled more rapidly than the bulk soil no matter which "mineral-associated" fraction (the silt+clay or the dense fraction DF) was chosen (Fig. 6). The $\Delta^{14}C$ values of TLE from the bulk, sand, and silt+clay fractions were indistinguishable 369 370 from one another, possibly because the size fractionation scheme did not effectively separate distinct lipid pools. However, 371 the Δ^{14} C values of TLE from the DF were significantly more ¹⁴C depleted than TLE from the silt+clay size fraction (Fig. 6), 372 suggesting there were older lipids in the DF relative to the silt+clay. However, more depleted ¹⁴C values found in the TLE 373 from the DF compared to the silt+clay could have resulted from the DF being exposed to SPT and/or ground after drying and 374 before lipid extraction. It is possible that grinding the DF prior to lipid extraction increased the exposed surface area and 375 resulted in a larger fraction of old SOC or rock-derived OC being incorporated into the TLE than if the DF had not been 376 ground. Clearly, the approach used to fractionate soils influences experimental results and must be considered when interpreting differences in persistence across operationally defined OC pools. We hesitate to definitively choose a best method 377 378 for fractionation because each soil environment and experiment require careful methodological consideration and selection. 379 However, given the clear differences in results between MAOM derived from size and density fractionation, it appears grinding 380 the samples prior to extraction had significant effects on the age of the resulting TLEWe hesitate to definitively choose a best 381 method for fractionation because each soil environment and system make require methodological alteration, however given 382 the clear differences in results between the methods. It appears grinding the samples prior to extractions have significant effects 383 on the lipid results. Clearly, the approach used to fractionate soils influences experimental results and must be considered 384 when interpreting differences in persistence across operationally defined OC pools.

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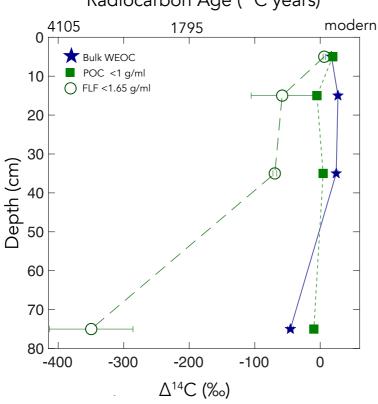
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386 **4.3 Variation in OC cycling throughout the depth profile**

387 The WEOC (extracted from bulk soils) and POC (<1g mL⁻¹ floated off the sand-size fraction) had the highest Δ^{14} C values 388 throughout the soil profile, reflecting a predominance of modern carbon from plant detritus and root exudates to these pools. 389 WEOC fractions can comprise a complex mixture of molecules with different structures (Hagedorn et al., 2004; Bahureksa et 390 al., 2021), which are common only in their ability to be mobilized and dissolved in water. WEOC can mobilize and percolate 391 down the soil profile with sufficient precipitation to allow vertical transport. Both the POC and WEOC fractions supply OC 392 that is readily accessible for microbial degradation and microbial utilization – resulting in the rapid turnover and relatively 393 high Δ^{14} C values of these two pools (Marin-Spiotta et al., 2011). Occurrence of young OC in deep soils may be driven by 394 microbial uptake of this young and bioavailable DOC or POC. Additionally, we found that the free light-density fractions were 395 depleted in ¹⁴C relative to the WEOC and POC (Fig. 5). We suspect this is due to colloidal particles in the FLF, which are not 396 dispersed or dense enough to settle in the SPT.

397 The study site has a Mediterranean climate, and these soils undergo seasonal wetting and drying cycles that may intensify 398 in the future (Swain et al., 2018), potentially shifting the composition or amount of OC that percolates down the soil column, 399 which could shift the age of the OC that the microbial community accesses at depth. When soil is already moist, subsequent 400 rainfall may mobilize both OC and colloidal sized mineral material under reducing conditions, which may interact to form 401 stable mineral-OC colloids that can enhance the transport of OC down the soil profile and out of the system (Buettner et al., 402 2014). With prolonged dry periods, water soluble OC may be more susceptible to microbial decomposition or oxidation because anaerobic preservation is removed (Heckman et al. 2022) This seasonal wetting and drying mechanism likely controls 403 404 what types of organic matter are transported down the soil profile. Deeper in the soil profile, greater reactive mineral surface 405 area and lower microbial activity can enhance carbon stabilization in subsoils (Homyak et al., 2018; Dwivedi et al., 2017; Pries 406 et al., 2023). Further research is needed to understand the effects of seasonal wetting and drying on the behaviour of water-407 soluble OC in the soil profile.

408 In general, the Δ^{14} C values of the TLE, AA, and AI fractions decreased with increasing depth in the profile. While all 409 extracted compounds followed this trend, the degree of ¹⁴C depletion with depth varied somewhat between the different 410 compound classes and between the bulk and silt+clay sourceparent fractions. The TLE extracted from the bulk and from the 411 silt+clay fraction had similar slopes with depth. This suggests that depth has more influence than fraction size on resulting 412 lipid ¹⁴C content, possibly because of limited transport of lipids down the soil profile. The AAs extracted from the bulk and 413 from the silt+clay fraction differed from one another in that the AA extracted from the bulk soil became more depleted with 414 depth than the AA extracted from the silt+clay. This suggests that at depth, AAs from the silt+clay fraction cycle more quickly 415 than AA's extracted from the bulk soil, possibly indicating that the silt+clay fraction is more directly influenced by microbial 416 activity than the sand fraction. At depths greater than 30 cm, the TLE and AA fraction were markedly younger than the bulk 417 soil, possibly resulting from transport of lipids and amino acids from surface horizons down profile, rapid recycling of these 418 compounds at depth, the use of a relatively modern C source for lipid and amino acid synthesis at depth, or most likely, a 419 combination of these. At all depths the AI was significantly older than the sourceparent fraction, indicating that throughout 420 the soil profile the AI contains an old and stable pool of OC.



Radiocarbon Age (¹⁴C years)

422 Figure 5: Particulate organic carbon (POC) (floated from the sand, n = 1), free light fraction (FLF) (from bulk soil, n = 3, and error 423 bars indicate standard error on the mean), and water extractable organic carbon -(WEOC) (from bulk soil, n =1). Δ^{14} C values by 424 depth. For POC and WEOC, error bars indicate analytical error are generally smaller than the symbols.

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4.4 Compound class Δ^{14} C values in mineral-associated SOC 426

427 To investigate the effect of mineral interaction on the Δ^{14} C values or persistence of the TLE, AA, and AI, we measured 428 these extracted compound classes from physical fractions intended to yield approximate mineral-associated carbon pools. We 429 focused primarily on the silt+clay size fraction as the physical fraction that best approximates a mineral-associated OC pool 430 derived from microbially processed plant inputs (Poeplau et al., 2018; Lavallee et al., 2020) and assume that after size 431 fractionation most of the free organic matter in the bulk soil was in the sand size fraction. We compared the silt+clay size fraction Δ^{14} C values to the bulk Δ^{14} C values to determine if the material extracted from the isolated mineral-associated fractions of the soil had greater OC persistence or if these compounds cycled indiscriminate of mineral association (Fig. 2).

While the TLE from the silt+clay and bulk soil had similar Δ^{14} C values, the AA from the silt+clay size fraction was enriched in ¹⁴C compared to the AA from bulk soil (r² = 0.98, p<0.05). This suggests that AAs cycle faster in the silt+clay mineral pool than in the bulk soils. While mineral surfaces usually are thought to promote stability and persistence of OC, in some soil systems, mineral associations may not be the single defining factor of OC persistence (Rocci et al., 2021) and could have a more nuanced role influencing OC cycling in soils.

Our data suggests there is a continuum of compounds that exist with different ¹⁴C values in the mineral-associated pool, because in the silt+clay fraction, the TLE, AA, and AI have significantly different ¹⁴C values (Fig. 4b). For instance, the mineral-associated TLE and AA fractions are enriched in ¹⁴C relative to the silt+clay fraction, suggesting both are cycling faster than the average mineral associated pool. However, the AI from the silt+clay fraction is cycling slower than solid sample it was extracted from, and when we compare the AI from the bulk soil to the AI from the silt+clay, the AI from the silt+clay slightly more ¹⁴C enriched. This suggests that there is slight ¹⁴C enrichment across compounds in the silt+clay fraction relative to sand and bulk soil.

We also compared the TLE extracted from the silt+clay to that extracted from the DF because both fractions are often 446 considered mineral associated. Across studies, the mineral associated OC is not a uniformly defined pool, and the observed 447 results are a consequence of the methodology used to separate the samples (Fig. 6). The DF TLE A¹⁴C is significantly older 448 449 than the silt+elay TLE (Fig. 6b) and the TLE of the bulk soil at depth (Fig. 6). This-Our data suggests that lipids in mineralassociated OC pools vary in cycling rates. This is complementary to findings from other studies where ¹⁴C values from different 450 451 lipid biomarkers are divergent from the bulk soils (Gies et al., 2021) and indicates the necessity of looking at entire compound 452 class pools for understanding soil carbon persistence. Further investigation into the composition and age-distribution of 453 compounds within mineral associated-OC is needed to better quantify the distribution of cycling rates within mineral 454 associated OC pools.

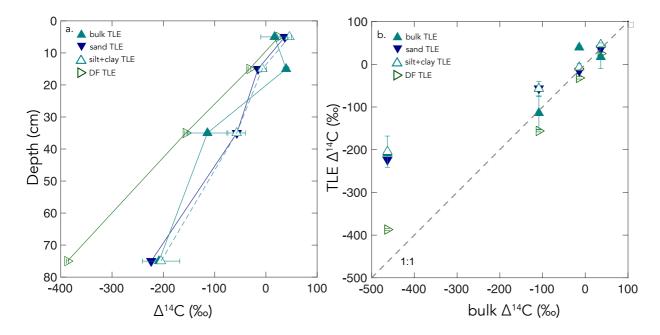


Figure 6: a) Δ^{14} C versus soil depth measured for <u>total lipid extractions (TLE)</u> extractions from four soil size/density fractions, <u>the</u> bulk (<2mm), sand (63 µm to 2mm), and dense fraction (DF). b) A comparison of the bulk soil Δ^{14} C values to the TLE from the four size/density fractions.

460 4.5 Persistent and Petrogenic OC

461 The most persistent, oldest OC was found in the AI fraction. Because carbon in the AI cycles more slowly than other 462 components of this grassland soil, it is important to understand what structural components make up the AI and where these 463 compounds are sourced from. Historically, tThe chemical structure of the AI fraction has been difficult to characterize. Hwang 464 and Druffel (2003) argued that the AI is a lipid-like portion of the ocean OC. However, in soils, the AI can be composed of a mixture of lipid-like compounds and aromatic compounds (Silveira et al., 2008). In our soil, the ¹³C-NMR spectra of the AI 465 466 from 0–10 cm depth show a significant, broad peak in the 100–165 ppm range, indicative of aromatics (SI Fig. 3) (Baldock 467 and Preston, 1995; Baldock et al., 1997). While it is possible that some condensed aromatic compounds form during the 468 hydrolysis procedure used to remove AAs, the AI may also contain naturally occurring aromatic compounds that could include 469 pyrogenic or petrogenic OC.

The parent material of our site is a mixture of sandstone, shale, greywacke, and schist (Foley et al., 2022), so it is possible that some of the OC in our soils is ancient, rock-derived, petrogenic carbon that has been incorporated into the soil profile through pedogenesis progresses (Grant et al., 2023). Comparison of the AI to the rock (>2 mm) fraction shows that the AI is younger than the OC contained in the rock fraction (SI Table 1), with the rock fraction Δ^{14} C values ranging from -481 to - 474 765₀₀. To calculate the contribution of OC_{petro} into the AI fraction, we used a binary mixing model with endmembers of 475 OC_{petro} and aged SOC based on the method in Grant et al. (2023). The $\Delta^{14}C$ value of the OC_{petro} ¹⁴C endmember is -1000 $\%_0$. 476 which is by definition ¹⁴C free, and and the Δ^{14} C value of the biospheric endmember was set as either the measured TLE Δ^{14} C value or the bulk Δ^{14} C value from each depth. This comparison of these two different biospheric endmembers allowed 477 478 us to calculate a possible range of values for the OC_{petro} contribution (Table 1). In the AI extracted from the silt+clay fraction, 479 the OC_{petro} contribution was 4-5% from 0-10 cm depth and 40-53 % in the 50-100cm depth. In AI extracted from the bulk 480 soil, the OC_{petro} contribution was 0-1 % in the 0-10 cm depth, and 17-44 % in the 50-100 cm depth. Therefore, while the AI 481 fraction likely contains OC_{petro}, it is primarily composed of OC compounds derived from more recent plant and microbial 482 inputs that are highly resistant to acid hydrolysis either because of their chemical structure or their strong associations with 483 minerals.

484

485 5 Conclusions and Continued soil radiocarbon compound class characterization

486 In this study, we characterized a soil carbon profile using compound-class ¹⁴C analyses. We found that our extraction 487 methods yielded fractions with ¹⁴C signatures distinctly different from the sourceparent soil from which they were extracted. 488 We found that in this annual grassland soil, the AA and the TLE fractions cycle more rapidly than the bulk soil throughout the 489 soil profile. At each depth, the AI fraction is the oldest fraction and contains a combination of slowly cycling SOC and ancient 490 petrogenic C. These results show that soil compound classes cycle differently than similar components in marine systems. Our 491 results also show that mineral-associated SOC contains a mixture of carbon compounds with distinctly different ages and sources that drive turnover and persistence. Compound-specific ¹⁴C approaches hold promise for improving our understanding 492 493 of the chemical structure of SOC, as well as the connection between carbon degradation and preservation in soils. A molecule-494 resolved understanding of the relationship between compound classes and carbon persistence will also give insight into the 495 fate and turnover time of specific organic biomarkers found in plant residues or the biomass of bacteria, fungi and microfauna. 496 These techniques can also help to determine mechanisms promoting mineral stabilization of soil carbon, especially when 497 combined with soil physical fractionation.

Results from this study highlight that radiocarbon measurements of specific organic compounds and compound classes in soil provide valuable insights into the persistence and decomposition rates of soil organic carbon. To improve our ability to model the future of soil carbon stocks and soil quality in the face of a changing global climate, we need further research that interrogates the composition, radiocarbon content, and cycling rates of soil organic carbon and mechanistically links these rates to physical and chemical drivers.

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513 7 Supplemental Tables/Data Availability

A list of all radiocarbon data, stable carbon, and total OC values with a CAMS tracking number for each of the analyses used in this publication.

8 Author Contributions: KJM, KMF, TABB, JP, and KEG conceptualized the study. KJM, KMF, TABB, JP secured funding for the project. KEG designed the method and carried out the extractions with input from KJM, KMF, and TABB. CJL carried out the density separations. MNR carried out the water extractions. JDK and MM ran the NMR experiments. KEG, KJM, KMF interpreted the data. KEG prepared the paper with contributions of all co-authors.

- 520
- 521 9 Competing interests. The authors declare that they have no conflict of interest.
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