



1 **Root turnover and soil indicators capture belowground**
2 **recovery following saltmarsh restoration**

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15



16 **Abstract**

17 Coastal wetlands, including saltmarsh, are highly productive ecosystems, with carbon- and
18 nutrient-rich soils supporting biodiversity. Beyond carbon stocks and sequestration, the
19 responses to restoration of these nutrient-rich and structurally complex soils remain poorly
20 defined for coastal wetlands, especially in saltmarsh ecosystems restored by exclusion
21 fencing. This study used a space-for-time approach to evaluate belowground responses in
22 *Salicornia quinqueflora*-dominated saltmarsh 25 years after ungulate exclusion in Swan Bay,
23 Victoria, Australia. We monitored surficial soil physicochemical characteristics, root and
24 standardised litter decomposition, and root molecular composition across Grazed, Restored,
25 and Natural Reference sites. Restored and Reference sites had $\geq 20\%$ higher vegetation cover
26 and 2-3-fold higher percent soil carbon and nitrogen content, with 2.5-fold lower shear vane
27 soil strength compared to Grazed sites. However, carbon and nitrogen stocks in the top 10 cm
28 were not significantly different across sites (means ranging 30-36 Mg C ha⁻¹) due to elevated
29 bulk density at Grazed sites caused by compaction from ungulates. *Salicornia quinqueflora*
30 root litter decomposition was slowest in Natural Reference sites, with molecular composition
31 showing preservation of recalcitrant lignin in the Reference and Restored sites, indicating
32 greater soil carbon preservation capacity. In contrast, roots decomposing in Grazed sites
33 showed increased nitrogen and phenolic compounds indicating greater microbial-driven
34 turnover. This study demonstrates that exclusion fencing can restore saltmarsh soil function
35 and promote long-term resilience, particularly through improved preservation of recalcitrant
36 organic matter material decades after intervention. By highlighting shifts in surface soil
37 structure and organic matter preservation, this study shows why soil quality metrics beyond
38 carbon stocks are essential for accurately evaluating restoration outcomes.



39 **1. Introduction**

40 Coastal wetlands comprise supratidal forests, saltmarshes, mangrove forests and seagrass
41 meadows. These ecosystems protect our coastlines and provide habitat for a large range of
42 plants and animals, making these ecosystems biodiversity hotspots (He et al. 2025). However,
43 since the early 1900s, nearly half of coastal wetlands have been lost globally due to climate
44 change and human activity (Davidson 2016). Coastal wetland loss can occur through
45 complete vegetation removal or more gradually through habitat fragmentation (Laegdsgaard
46 2006). Such coastal wetland degradation can compromise key ecosystem functions, including
47 food chain supply, biodiversity conservation, resilience to sea level rise, water filtration, and
48 carbon accumulation and storage (Pralhad 2014). Therefore, understanding the extent and
49 impact of wetland degradation and restoration is critical for the planning and implementation
50 of future management strategies.

51 Restoration of coastal wetlands has the potential to reestablish a range of essential ecosystem
52 functions, including vegetation production and carbon and nutrient cycling, which in turn
53 supports food web complexity, biodiversity and carbon sequestration (Meli et al. 2014). There
54 is an increasing demand for large-scale wetland restoration and protection (Macreadie et al.
55 2021). The exclusion of ungulates by fencing is one such restoration approach that is
56 attractive for the restoration of marginal saltmarsh ecosystems due to its relatively low costs
57 and minimal intervention (Rowland and Lovelock 2024). Over the last few decades, studies
58 have quantified a range of benefits of ungulate exclusion or abandonment to aboveground
59 vegetation (e.g. survival, growth, diversity, successional shifts, and litter production;
60 Laegdsgaard 2006, Rupprecht et al. 2015, Wasson et al. 2021) and soil structure (e.g.
61 increased elevation, moisture, and aeration; Veenklaas et al. 2015, Chang et al. 2016).



62 Recovery of aboveground vegetation biomass and diversity are the most commonly
63 quantified metrics for ecosystem recovery after ungulate exclusion (Rowland and Lovelock
64 2024), and such responses are used to suggest or estimate belowground indicators of soil
65 recovery and function (e.g. carbon accumulation and stocks) (Legesse et al. 2024, Xiao et al.
66 2025). For example, the Australian Blue Carbon Accounting Model estimates the soil carbon
67 benefits of coastal wetland restoration through tidal reinstatement using aboveground
68 vegetation metrics (Lovelock et al. 2022a). However, such correlative approaches do not
69 account for the complex belowground processes influencing wetland recovery and health, and
70 remain poorly understood in the coastal wetland restoration context (Bayraktarov et al. 2020).
71 As such, there is a need to improve the breadth and depth of data on indicators of
72 belowground ecological functions under current and emerging restoration methods to
73 enhance accounting model accuracy (Lovelock et al. 2022a), as well as improving the
74 reliability of restoration assessments that inform model predictions and policy.

75 For saltmarsh ecosystems, static belowground ecological functions like soil carbon stocks are
76 most often reported for restoration monitoring (Bayraktarov et al. 2020, Cadier et al. 2020,
77 Rowland and Lovelock 2024). In contrast, process-based functions like the decomposition of
78 root or soil organic matter (OM) are less commonly quantified despite being essential for
79 nutrient regeneration, soil formation and responses to changes in inundation (i.e. subsidence
80 vs. accumulation) (Meier and Bowman 2008, Kirwan et al. 2013). Understanding the influence
81 of root litter quality on decomposition, and *vice versa*, is required to comprehend the impact of
82 management on belowground OM cycling. Analytical pyrolysis techniques such as pyrolysis-
83 GC-MS (Py-GC-MS) and thermally assisted hydrolysis and methylation (THM-GC-MS) can
84 be used to characterise a wide range of plant biomolecular constituents that indicate the quality
85 of OM, as well as their response to management practices, by showing changes like microbial
86 necromass accumulation, loss of labile constituents, and the selective preservation of relatively



87 recalcitrant plant constituents like root-specific suberin (Kolattukudy 2001, Ferreira et al. 2009,
88 Pineiro-Juncal et al. 2021). In addition, belowground OM availability and cycling can be
89 influenced by localised, site-specific characteristics like soil compaction, as measured by bulk
90 density and soil strength, that can change in response to restoration interventions (Brooks et al.
91 2022). Therefore, standardised OM like rooibos and green tea that have uniform chemical
92 properties can be used in decomposition studies to help distinguish the effects of site-specific
93 environmental factors from indicators of soil processes and land management practices
94 (Keuskamp et al. 2013).

95 While it can take years or decades after restoration to detect belowground functional
96 responses comparable to levels of natural reference ecosystems, space-for-time experimental
97 designs allow us to compare sites that represent different restoration ages to natural reference
98 sites and degraded baseline sites and therefore provide a practical approach to assess
99 restoration success over time (Gulliver et al. 2020, Carnell et al. 2022). Accordingly, we used
100 a space-for-time approach to evaluate the belowground responses across actively grazed
101 saltmarshes, sites excluded from grazing for approximately 25 years, and natural reference
102 saltmarshes. We hypothesise that belowground decomposition and OM quality will be
103 sensitive and detectable indicators of soil responses to exclusion fencing restoration. We
104 focused on the changes in soil properties within surficial soils (top 10 cm), where active root
105 inputs are more likely to affect soil carbon turnover and formation. We measured indicators,
106 including root and standardised OM decomposition and molecular shifts for short-term OM
107 turnover and preservation, soil strength and dry bulk density for soil structure, and soil carbon
108 and nitrogen content for longer-term OM preservation. By taking a multi-indicator approach,
109 this study will advance our understanding and application of robust soil indicators for
110 saltmarsh restoration.

111



112 **2. Materials and methods**

113 **2.1. Site description and experimental design**

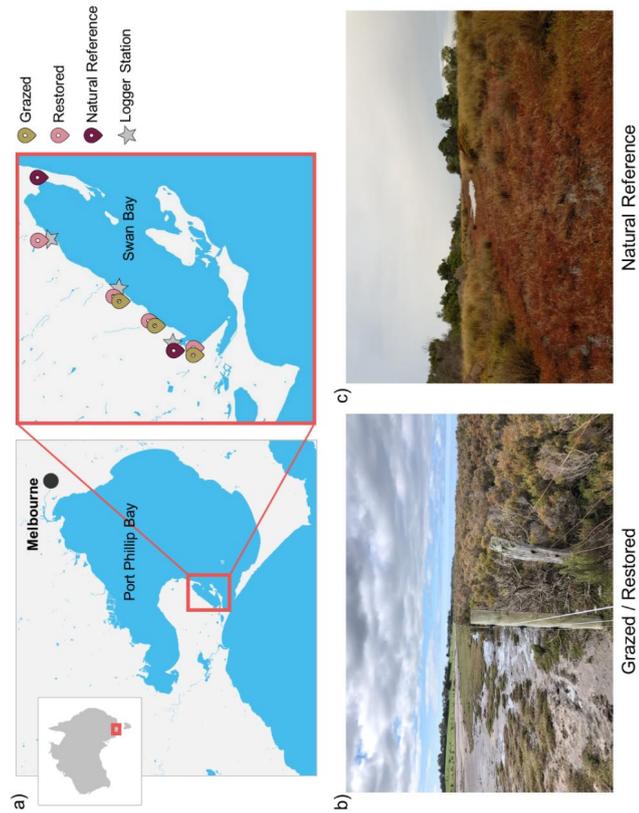
114 All sites are located within Swan Bay, a tidally influenced embayment in Port Phillip Bay,
115 Victoria, Australia (Figure 1a). The coastline, which featured saltmarsh ecosystems, was
116 heavily modified during European settlement for livestock grazing until Swan Bay became a
117 marine reserve in 1979. In 1996, marginal land within Crown Boundaries on the western
118 shore was fenced by Parks Victoria to restrict access and allow saltmarsh to return (i.e. >25
119 year Restored sites; Figure 1b). Private lands adjacent to Crown Land are still being grazed,
120 but are at elevations suitable for saltmarsh to re-establish, indicated by access to tidal flow
121 encroachment of pioneer saltmarsh species such as *Salicornia quinqueflora* (*S. quinqueflora*)
122 (i.e. Grazed sites; Figure 1b). In contrast, some areas of Swan Bay on the northern and
123 southern ends have had little to no modification in the last 50-60 years (landholder and Parks
124 Victoria pers. comm.), and for the purposes of this study will act at a Natural Reference site
125 (i.e. Natural Reference sites; Figure 1c).

126 We chose at least two sites from each rehabilitation condition category to assess belowground
127 responses to grazing and exclusion fencing activities: Grazed (G), Restored (>25y; R), and
128 Natural Reference (NR). All Grazed sites had encroaching saltmarsh growth at least ≥ 0.5 m
129 into the property from nearby marsh. One of the historically Grazed sites (G2) was fenced 3
130 months prior to the start of the experiment, so we also monitored this site as a historically
131 grazed area that may provide information on early responses to exclusion fencing restoration.
132 Restored and Reference sites are classified at Environmental Vegetation Class 9 (Coastal
133 Saltmarsh Community) and comprised of *Salicornia* and *Tecticornia* (Table 1). While we
134 tried to reduce variability in elevation across sites and treatments, the Grazed sites are
135 landward of the Restored and Natural Reference sites, and typically of higher elevation. To



136 reduce variability in belowground responses due to tidal inundation, we chose areas within
137 the site that were seaward and showed evidence of saltmarsh encroachment by pioneer
138 species *S. quinqueflora*.

139 In February 2022, we installed five 40 cm x 40 cm experimental plots at each site,
140 approximately 1 m apart, in a transect running parallel to the shoreline. We sampled soil at
141 each plot to characterise surficial carbon and nitrogen stocks, soil strength and dry bulk
142 density. While site-level vegetation varied across the rehabilitation categories (Table 1), to
143 help standardise the conditions and be relevant to in situ saltmarsh root decomposition, we
144 chose transects where *S. quinqueflora* were the dominant taxa. Within each plot at 3, 6 and 12
145 months after deployment, we monitored aboveground vegetation cover and diversity, as well
146 as decomposition and litter chemistry of root and standardised green and rooibos tea litters
147 over the course of 1 year.



148

149 Figure 1. Site map. (a) Swan Bay is an embayment in Port Phillip Bay, Victoria, Australia. Points represent site rehabilitation category and
150 location of logger stations (b) A site in Swan Bay, showing a currently grazed area on the left, and a >25-year restored area on the right,
151 following the installation of a fence by Parks Victoria in 1996 to restrict access and allow saltmarsh to return. (c) A natural reference site has had
152 little to no modification in the last 50-60 years.



153 Table 1: Site descriptions. Elevation was measured at each plot. Grazed sites have cattle on
 154 the land with some saltmarsh encroachment from the neighbouring restored sites. The
 155 exception is G2 which was <1 year fenced at the time of the study. Grazers had been
 156 excluded from the Restored sites for approximately 25 years, while natural reference sites
 157 were approximately ≥ 60 years old.

Rehabilitation Treatment	Site Name	Year of establishment	Elevation Range (m)	Vegetation
	G1	0	0.581 - 0.594	<i>Salicornia, Suaeda</i>
Grazed	G2	2021	0.792 - 0.81	<i>Salicornia, Distichlis</i>
	G3	0	0.706 - 0.79	<i>Salicornia</i>
	R1	1996	0.544 - 0.587	<i>Salicornia</i>
	R2	1996	0.455 - 0.485	<i>Salicornia, Suaeda</i>
Restored				<i>Salicornia, Tecticornia, Frankenia,</i>
	R3	1996	0.574 - 0.606	<i>Suaeda</i>
	R4	1996	0.668 - 0.729	<i>Salicornia, Suaeda, Frankenia</i>
Natural reference	NR1	n/a	0.499 - 0.551	<i>Salicornia, Tecticornia, Frankenia</i>
	NR2	n/a	0.44 - 0.507	<i>Salicornia, Suaeda, Frankenia</i>

158



159 **2.2. Site characteristics**

160 **2.2.1. Elevation, water level and physicochemical measurements**

161 The elevation at each plot was measured using an EMLID Reach RS+ RTK-GNSS unit.
162 Elevation ranges of the plots were 0.58 – 0.81 m for Grazed, 0.46 – 0.73 m for Restored and
163 0.44 – 0.55 m for Natural Reference sites (Table 1). Soil temperature loggers (HOBO
164 Pendant – Temperature/Light data logger UA-002-64, Onset) were buried at a 5 cm depth at
165 each site. Temperature was recorded every 10 minutes for the duration of the experiment.

166 Logger stations were set up along the length of Swan Bay at NR1, R1 and R2 to understand
167 the water level and soil redox (ORP) along the coastline of Swan Bay (Figure 1a). The data
168 were logged every 10 minutes for at least a full tidal cycle during the initial, 6-month and 12-
169 month samplings. First, REED data loggers coupled with Paleoterra redox and reference
170 probes were deployed at 10 cm. HOBO U20L water level data loggers were placed in
171 perforated PVC casings, and the sensor location was 15 cm below the surface. The elevation
172 of the water present at the deployment site was measured using an EMLID Reach RS+ RTK-
173 GNSS unit. One dedicated water level logger was installed 50 cm above the surface to record
174 barometric pressure variability of Swan Bay. The elevation of the water body and barometric
175 pressure data were used in post-processing to correct and calculate water level measurements.

176 The water level loggers and water elevation data were used to calculate the cross-shore
177 elevation profile at the nine different sites: Mean Higher High Water (MHHW), High Water
178 (HW), Mean Sea Level (MSL), Low Water (LW), Mean Lower Low Water (MLLW) (Figure
179 S1, Table S1). Plots located above MHHW, ranging from + 0.0079 to + 0.0621 m (G1, R1,
180 R2, R3, NR1, and NR2), experience less frequent and shorter periods of tidal flooding
181 compared to those below MHHW ranging from -0.112 to -0.262 m (G2, G3, and R4) (Table
182 S1). The profiles also reveal variations in elevation along the cross-shore transects. G1, G2,



183 and G3 profiles show relatively flat topography, with elevations close to or slightly above
184 MHHW level. R1, R2, and R3 profiles exhibit more pronounced changes in elevation, with
185 the landward portions of the transects rising above the MHHW level and the seaward portions
186 below MSL and MLLW. R4 profile is similar to R1 – R3 profiles but with a more gradual
187 slope. NR1 and NR2 profiles display the most significant changes in elevation along the
188 cross-shore transects. Both profiles start at elevation near the MHHW level on the landward
189 side and then steeply drop below the MLLW level towards the seaward end of the transects.
190 The NR2 profile, in particular, shows a more pronounced drop in elevation compared to NR1.



191 **2.2.2. Soil cores and strength**

192 Soils at each site were analysed for elemental organic carbon (%C) and total nitrogen (%N)
193 as well as soil strength. Soil strength, measured with a shear vane tester (19 mm vane; CMT
194 equipment, Australia), provides information on the mechanical strength in kPA needed to
195 move the vane. Soil strength was measured at 10 and 20 cm depths (8-10 cm and 18-20 cm,
196 respectively, accounting for vane height).

197 Surficial soil cores were taken directly adjacent to each plot using a 10 cm length syringe
198 core, to capture the depth range of root production and decomposition. Cores were sliced at
199 0-2, 2-5 and 5-10 cm intervals and dried at 60 °C to calculate dry bulk density (DBD, g cm⁻³).
200 Samples were ground to a fine powder (Retch RM200) and tested for carbonates using
201 diluted hydrochloric acid. Effervescence after contact with acid indicated inorganic carbon as
202 carbonates were present. Percent organic carbon and total nitrogen analyses were performed
203 on a LECO elemental analyser at the CSIRO Land and Water Facilities (Adelaide). Briefly,
204 total carbon and nitrogen were determined by high temperature combustion in an atmosphere
205 of oxygen using a Leco TruMAC. For samples that did not indicate presence of inorganic
206 carbon, total carbon was assumed to be organic carbon. For carbonate-positive samples,
207 inorganic carbon as carbonates was determined by reacting the sample with acid in a sealed
208 container and measuring the pressure increase (Sherrod et al. 2002, Rayment and Lyons
209 2011). The organic carbon is calculated as (Total carbon – Carbonate*0.12) (as per
210 communication by CSIRO). Organic carbon and nitrogen density were calculated using DBD
211 (g C or N cm⁻³) and converted to mean stock in the top 10 cm at each site (e.g. Mg C ha⁻¹)
212 (Howard et al. 2014).

213

214



215 **2.3. Vegetation monitoring**

216 **2.3.1. Aboveground vegetation**

217 Vegetation diversity and cover at each site were determined visually using a 100 x 100 cm
218 quadrat thrown at random 10 times per site during site assessment. A quadrat (40 x 40 cm),
219 matching the plot size, was used to take plot photos of each plot at the 3, 6, and 12 months
220 sampling times. The photos were then used to determine changes in vegetation diversity and
221 cover within each plot over time.

222 **2.3.2. Root and tea litter decomposition**

223 *Salicornia quinqueflora* roots and standardised green and rooibos tea litters were incubated in
224 the soils to investigate organic matter turnover via decomposition. The standardised tea litter
225 approach has been used extensively to investigate cross-ecosystem and regional/global
226 within-ecosystem drivers of coastal wetland decomposition (Mueller et al. 2018, Trevathan-
227 Tackett et al. 2024). Lipton green and rooibos teas from the original Tea Bag Index product
228 numbers (~0.25 mm mesh, 0.6 mm²; (Keuskamp et al. 2013)) had changed from a nylon
229 mesh to a biodegradable mesh material and was repackaged in nylon mesh tea bags (~0.17 x
230 0.24 mm mesh, 0.4 mm²) to persist in a longer-term incubation. *Salicornia quinqueflora* roots
231 were collected, cleaned of attached soil and dried before adding ~1.7 g dry weight to nylon
232 mesh tea bags.

233 Litter bags were buried in the soil at ~5 cm depth. Each of the five plots per site contained
234 one root litter, green tea and rooibos tea bag per sampling time. Bags were collected at 3, 6
235 and 12 months, cleaned of attached soil and dried at 60 °C. Litter samples were further
236 cleaned of living root contamination before obtaining dry mass. Proportion mass remaining at
237 each time point was calculated and used to calculate single exponential decay rates, using
238 $W_0/W_t = e^{-kt}$, whereby W_0 was the initial mass, W_t was the mass at time t and k is the decay



239 rate in proportion per day (d^{-1}), with the model inclusive of 100% mass at day 0 (Trevathan-
240 Tackett et al. 2021). k was calculated separately for each litter type for each site and
241 rehabilitation category. Root litter carbon and nitrogen were quantified using the elemental
242 analysis described above for soils. Molar C:N ratios were calculated. Percent carbon and
243 nitrogen data were normalised to sample mass, and proportion carbon and nitrogen remaining
244 were used to calculate single exponential decay rates.

245 **2.3.3. Py-GC-MS and THM-GC-MS of *S. quinqueflora* roots**

246 In order to understand the impact of management on the decomposition processes in the
247 incubated root tissues, a subset of *S. quinqueflora* root samples (2 sites from each
248 rehabilitation category and from 3 and 12 month samplings) were analysed for molecular
249 composition via pyrolysis-GC-MS (Py-GC-MS) and thermally-assisted hydrolysis and
250 methylation (THM-GC-MS). Briefly, 1 mg of sample was introduced into quartz wool-
251 containing fire-polished quartz tubes, placed into a Pyroprobe 5000 (CDS Analytical)
252 interface (at 325 °C), and pyrolyzed at 650 °C set-point temperature for 20 s. The pyrolysis
253 products were swept into an Agilent 5977 GC-MS instrument by 1 ml/min helium flow. The
254 pyrolysis and THM products were separated on a HP-5MS GC column and identified using
255 EI at 70 eV, scanning in the m/z 50-500 range. For THM-GC-MS, the analysis was preceded
256 by the addition of an aliquot of aqueous tetramethyl ammonium hydroxide (TMAH, 25%
257 from Sigma-Aldrich).

258 The 109 pyrolysis (Table S2) and 105 THM (Table S3) GC-MS products were semi-
259 quantified based on the peak areas of their dominant m/z fragments. Relative proportions
260 were calculated as the % of total quantified peak area, for Py- and THM-GC-MS separately.
261 These datasets are “closed datasets” that allow the study of organic matter composition and
262 variation therein. The products were grouped into several main classes (Tables S2 and S3).



263 Using (conventional) Py-GC-MS, the macromolecules in a sample are fragmented thermally
264 in absence of reagents (Wampler 2006, Voorhees 2013), whereas in THM-GC-MS, the
265 presence of the strongly alkaline TMAH causes hydrolysis and derivatization (methylation)
266 of hydrolysable functional groups, such as hydroxyl to ether and carboxyl to ester (Challinor
267 2001, He et al. 2020) The set of compounds generated, and their chromatographic behavior,
268 are very different for the two techniques, which therefore have different sensitivities to
269 different OM types (Kaal et al. 2020).

270

271 **2.5. Data analysis**

272 Statistical analyses and PCA were performed in R (v. 4.4.2) using car, lme4, glmmTMB,
273 emmeans and FactoMineR packages. Generalised linear mixed effect models were used to
274 test for the main and interaction effects of rehabilitation category (Grazed, Restored, and
275 Natural Reference) and variable-specific main effects for soil parameters (depth and
276 rehabilitation category), final litter mass remaining and k-rate (litter type and rehabilitation
277 category), and elemental carbon and nitrogen for *S. quinqueflora* roots (sampling time
278 excluding initial data and rehabilitation category). Site was a random factor in all models. For
279 the molecular analyses of *S. quinqueflora* root material that had reduced replication, one-way
280 linear models were used separate for time and rehabilitation category condition test. A Tukey
281 adjustment was made for all pairwise analyses.

282 We performed principal component analysis (PCA) to examine the important variables
283 contributing to the variability in belowground soil characteristics and the relationships
284 between those variables with rehabilitation categories, simultaneously. The eleven variables
285 included ten belowground variables (organic carbon stock (g OC cm^{-3}), nitrogen stock (g N
286 cm^{-3}), percent organic carbon, percent total nitrogen, soil C:N ratio, soil strength (kPA), DBD



287 (g cm⁻³), soil elevation (m), soil temperature (°C), and the proportion of mass remaining of *S.*
288 *quinqueflora* root) and one aboveground variable (vegetation cover percent). The organic
289 carbon stock, nitrogen stock, percent organic carbon, percent total nitrogen, soil C:N ratio,
290 and DBD were averaged across the 10 cm core sections for the PCA. Vegetation cover
291 percent was averaged across the three time points, 3, 6 and 9 months.

292

293 **3. Results**

294 **3.1. Site characteristics**

295 **3.1.1. Soil redox and temperature (abiotic indicators)**

296 There was minimal fluctuation in redox potential across the tidal cycles measured at the three
297 logger stations. Redox potential measurements were positive at all sites and time points. At 6
298 months in winter August 2022, the redox potential measurements were lowest for the Natural
299 Reference (NR1; 103 ± 0.71 mV mean and SEM) and Grazed (G2; 121 ± 1.4 mV) sites,
300 while the restored site (R1) had approximately 2-fold higher redox potentials (234 ± 4.9 mV),
301 suggesting more oxidized soil conditions. In summer February 2023, the redox potential
302 measurements were highest for the Grazed site (267 ± 3.9 mV), in contrast to the relatively
303 more reduced Restored (106 ± 0.5 mV) and Reference sites (54 ± 0.9 mV).

304 Soil temperature across all sites and rehabilitations categories followed a consistent seasonal
305 trend. Summer temperatures maximums were around 25-30 °C, with peak monthly mean
306 temperatures in January 2023 (~21 – 24 °C; Figure S2). Extreme winter temperature lows
307 were approximately 0-5 °C, with minimum monthly mean temperatures in July 2022 (~-10-12
308 °C; Figure S2). The Grazed plots (G1, G2, and G3) exhibited slightly higher monthly mean
309 temperatures, particularly for sites G1 and G2. The Restored plots (R1, R2, R3, and R4) also
310 display similar seasonal patterns, with R2 and R3 having slightly higher summer



311 temperatures than R1 and R4. The similarity in soil temperature patterns among the different
312 rehabilitation categories (Grazed, Restored and Reference) suggests that factors such as
313 climate and seasonal weather variations have a dominant influence on soil temperature,
314 regardless of the plot's specific characteristics (e.g. elevation) or management status.

315

316 **3.1.2 Soil carbon, nitrogen, bulk density and strength (physiochemical indicators)**

317 In the top 10 cm, percent soil OC (%OC) and N (%N), as well as DBD, were significantly
318 different across rehabilitation category and depths (main effects $p \leq 0.002$; Figure 2). %OC
319 and %N significantly decreased with depth, while DBD increased (Figure 2a, b, e). Grazed
320 sites had significantly lower %OC and %N but higher DBD compared to both Restored and
321 Natural Reference sites (Figure 2a, b, e), with Natural Reference soil visibly appearing
322 peatier and rich in root material (Figure S4). There were no significant differences between
323 Restored and Natural Reference sites for these soil parameters. When normalised with DBD,
324 carbon and nitrogen densities (g cm^{-3}) were not significantly different across rehabilitation
325 categories, but significantly decreased with depth (Chisq = 58.7, $p = 1.8 \times 10^{-13}$ and Chisq =
326 71.8, $p = 2.6 \times 10^{-16}$, respectively; Figure 2c,d). Total carbon stocks in the top 10 cm were not
327 statistically different among Grazed ($36.1 \pm 3.5 \text{ Mg C ha}^{-1}$), Restored ($30.4 \pm 1.4 \text{ Mg C ha}^{-1}$)
328 and Natural Reference ($31.1 \pm 1.2 \text{ Mg C ha}^{-1}$) categories (Chisq = 1.28, $p = 0.52$).

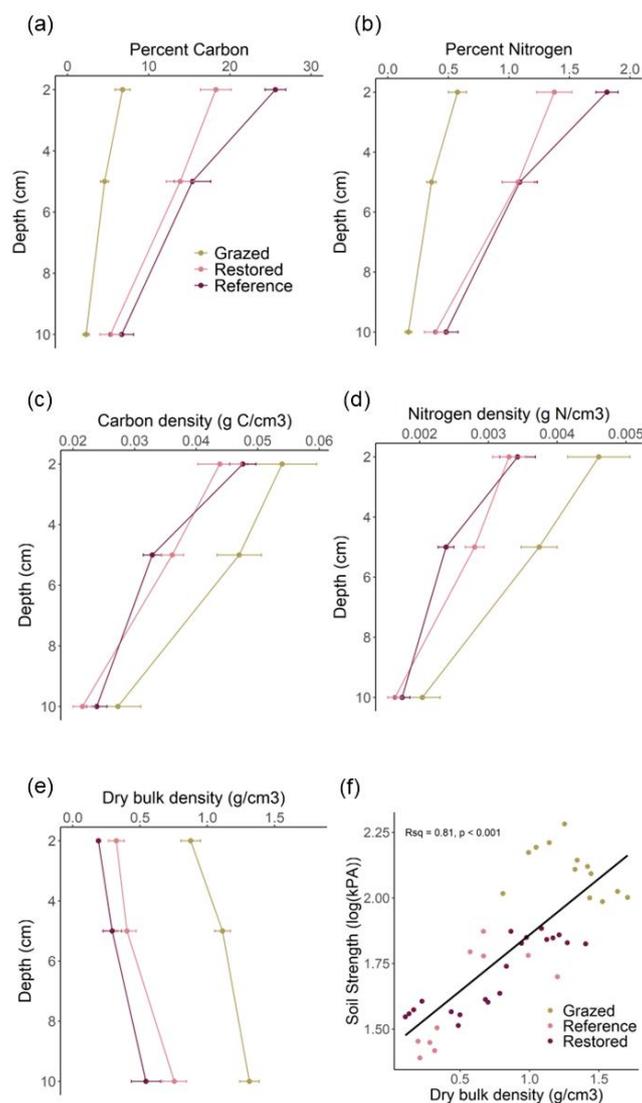
329 Soil strength was significantly different across rehabilitation categories for both 10 and 20 cm
330 depths (Chisq = 18.0, $p = 0.00012$ and Chisq = 17.5, $p = 0.00016$, respectively; Table S4).
331 Grazed sites had the highest soil strength at 10 and 20 cm depths (124 ± 8.0 and 157 ± 8.4
332 kPA, respectively), while soil strength means at Restored and Natural Reference sites were
333 1.7-3.6-fold lower (Table S4). We found a significant and strong positive correlation between
334 DBD and soil strength at 10 cm (Pearson's $t = 8.7$, $p = 7.2 \times 10^{-11}$, $R^2 = 0.806$; Figure 2f).



335

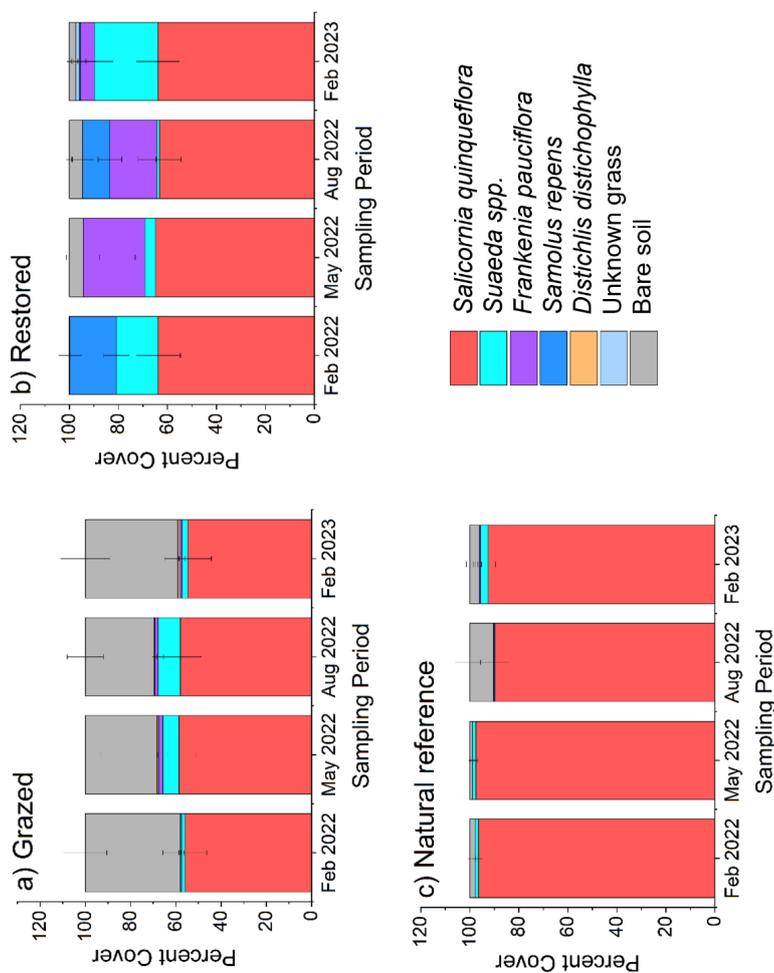
336 **3.2. Aboveground vegetation (biodiversity indicator)**

337 *Salicornia quinqueflora* dominated our experimental plots with 50% or more of cover
338 throughout the year (Figure 3). The Natural Reference sites had the highest *S. quinqueflora*
339 cover ($\geq 90\%$) leading to the lowest alpha diversity of the three rehabilitation categories
340 (Figure 3, Figure S3). Restored sites had the highest alpha diversity that changed slightly in
341 composition over the course of a year (e.g. *Frankenia pauciflora* and *Suaeda* spp). Grazed
342 sites had the highest cover of bare, unvegetated areas of the plots (means ranged from 30-
343 42%) with ephemeral *Suaeda* cover in addition to *Salicornia* patches in our plots (Figure 3).



344

345 Figure 2: Top 10 cm saltmarsh soil characteristics at Grazed, Restored and Natural Reference
346 sites. (a) Elemental percent organic carbon. (b) Elemental percent nitrogen. (c) Organic
347 carbon density. (d) Nitrogen density. (e) Dry bulk density. Data represent means and standard
348 error. (f) Relationship between dry bulk density and soil strength at ~10 cm depth. Note the
349 log scale of soil strength. The line represents Pearson's correlation.



350

351 Figure 3: Saltmarsh vegetation cover over one year at (a) Grazed, (b) Restored and (c) Natural Reference sites. Vegetation surveys were taken of
 352 the experimental plots at initial, 3 month, 6 month and 12 month sampling periods. Data represent mean and standard error.



353 **3.3. Litter decomposition**

354 **3.3.1. *Salicornia quinqueflora* root and tea litter decomposition (biological turnover)**

355 The proportion of mass remaining at the end of the study was significantly different among
356 litter types ($p < 0.001$; Figure 4a), with rooibos tea the highest (0.611 ± 0.012), followed by *S.*
357 *quinqueflora* roots (0.310 ± 0.008) and green tea (0.108 ± 0.012). An interaction effect for
358 litter type and rehabilitation category ($\text{Chisq} = 27.923$, $p < 0.001$) was driven by a
359 significantly higher mass remaining for green tea litter at the Grazed sites compared to
360 Restored sites ($z\text{-ratio} = 4.636$, $p < 0.0001$) and Natural Reference sites ($z\text{-ratio} = 3.791$, $p =$
361 0.0004). There was no significant effect of rehabilitation category on decay rates, but a highly
362 significant effect of litter type ($\text{Chisq} = 303.9692$, $p < 2e^{-16}$), with the decay rates of green tea
363 nearly 8-fold and 2-fold higher than rooibos and *S. quinqueflora* roots, respectively (Figure
364 4b, Table S5).

365 Organic carbon, nitrogen and C:N ratios of *S. quinqueflora* roots had a significant
366 time*rehabilitation category interaction (Figure 5). For rehabilitation category, Grazed sites
367 were significantly higher for nitrogen than Restored sites at the final 12-month sampling ($p =$
368 0.0222), while significantly lower than Restored sites for C:N ratios ($p = 0.0239$; Figure
369 5b,c). Over time, root carbon significantly increased as decomposition progressed for each
370 rehabilitation category (Figure 5a), but nitrogen changed differently according to
371 rehabilitation category (Figure 5b,c). Specifically, the Natural Reference sites did not
372 significantly change over time, while nitrogen increased in Grazed sites but decreased at
373 Restored sites (Figure 5b). These relative changes in root elemental content led to a
374 significant increase in C:N ratios at the Restored sites but not Grazed or Natural Reference
375 sites (Figure 5c). While both carbon and nitrogen decay rates were lowest for the Natural



376 Reference sites, they were not significantly different to the decay rates at the Grazed and
377 Restored sites (Table S5).

378

379 **3.3.2. *Salicornia quinqueflora* root molecular shifts during decomposition (molecular**
380 **preservation indicator)**

381 The molecular analysis of the decaying *S. quinqueflora* roots indicated significant shifts over
382 time, and limited changes related to rehabilitation category (Figure 6, Figure 7 & Figure S5).

383 The multivariate analyses on the Py-GC-MS data showed that 31% of the variance (PC1) was
384 driven by changes in time, while PC3 (11% of variance) was indicative of nitrogen-rich
385 microbial compounds and phenol accumulation at the Grazed sites, in contrast to the enriched
386 syringyl lignin in restored/reference sites (Figure 7, see Figure S5i,j for principal component
387 loadings). PC2 (19 %) did not depend statistically on incubation time or treatment and had
388 high positive loadings for lignin products. Specifically, we found that only nitrogen
389 compounds (i.e. pyrroles, pyridines, acetamide, indoles) and phenols formed upon Py-GC-
390 MS were significantly different among rehabilitation categories, both of which became
391 enriched at the Grazed sites (Figure 7, Figure S5d, e). We noted marginally insignificant
392 effects for N-compounds between Grazed and Restored ($p = 0.051$) and Reference ($p = 0.72$)
393 treatments in pairwise analysis (Figure S5d).

394 Throughout the decomposition incubation of the root litter across all rehabilitation categories,
395 relative concentration of carbohydrate products from Py-GC-MS (including acetic acid,
396 furans, pyrans, anhydrosugars; Figure S5a) and THM-GC-MS (methylated metasaccharinic
397 acids; Figure S5k) significantly decreased, indicating loss of polysaccharides at the 3- and 12-
398 month decay periods. Alkyl-polycyclic aromatic hydrocarbons (PAHs) and unsaturated fatty
399 acids were preferentially eliminated (Figure S5f,o), with the former probably indicating



400 leaching and/or decomposition of resin or resin-like organic matter. Other methylene chain
401 compounds (MCCs; alkanes, alkenes, fatty acids, methylketones) from Py-GC-MS
402 significantly increased throughout the year of decomposition (Figure S5c). These results
403 aligned with relative increases in THM-GC-MS-detected MCCs specific of the relatively
404 recalcitrant root-derived macromolecule suberin (Figure S5m), in particular long-chain (C_{16} ,
405 C_{18} , $C_{18:1}$, C_{20} , C_{22}) fatty diacids (Kolattukudy 2001) (Figure S5n). In contrast, lignin and the
406 Py-GC-MS ratios of lignin oxidation (4-acetylguaiacol/guaiacols and 4-
407 acetylsyringol/syringols significantly increased only in the first 3-months (Figure S5b,g,h), as
408 corroborated by the analogous methoxybenzenes detected by THM-GC-MS (Figure S5l);
409 both indicating an increase in guaiacyl- and syringyl-type lignin products.

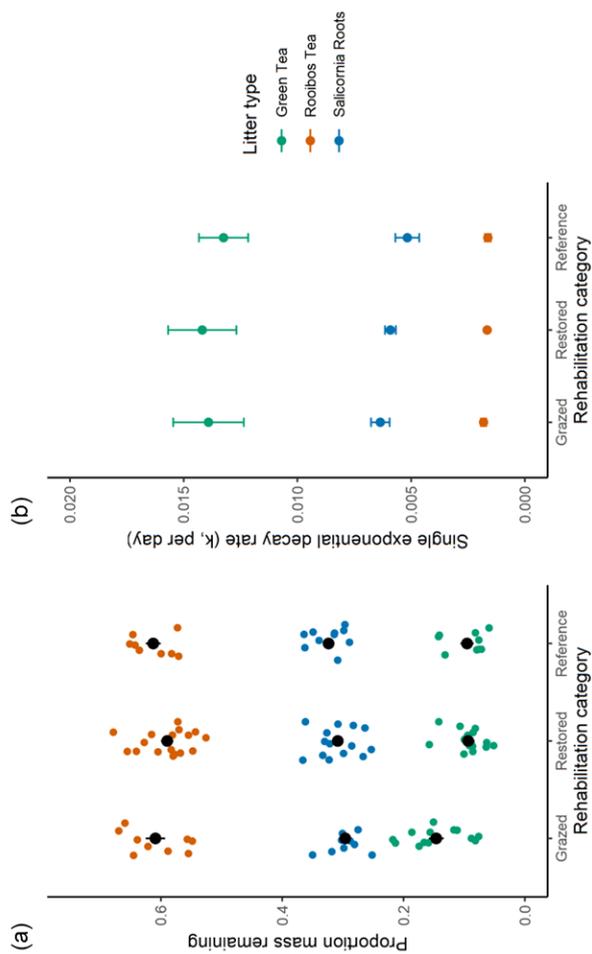
410

411 **3.4. Belowground responses to land-use and restoration (indicator integration)**

412 A total of 70.8% of the belowground variability was explained by the first two principal
413 components. PC1 explained 48.5% of variation and was driven by positive correlations for
414 percent organic carbon and percent nitrogen and by negative correlations with DBD, soil
415 strength and soil elevation (Figure 7). PC2 (22.3%) was dominated by carbon stock and
416 nitrogen stock which were correlated along the positive axis, as well as vegetation cover
417 percent with negative correlation (Figure 7). The squared cosine (Cos^2) shows the importance
418 of a component for a given observation in PCA, and hence, the value of Cos^2 can help to find
419 the components that are important in driving differences in rehabilitation categories. Cos^2
420 analyses suggest carbon stock, followed by DBD and nitrogen stock, are the most significant
421 in contributing to the variability of the soil characteristics (Figure 7). The Restored and
422 Natural Reference sites overlapped with each other, whereas the Grazed sites plotted with the
423 least overlap with the other two. Carbon stock, nitrogen stock, soil strength and DBD are the

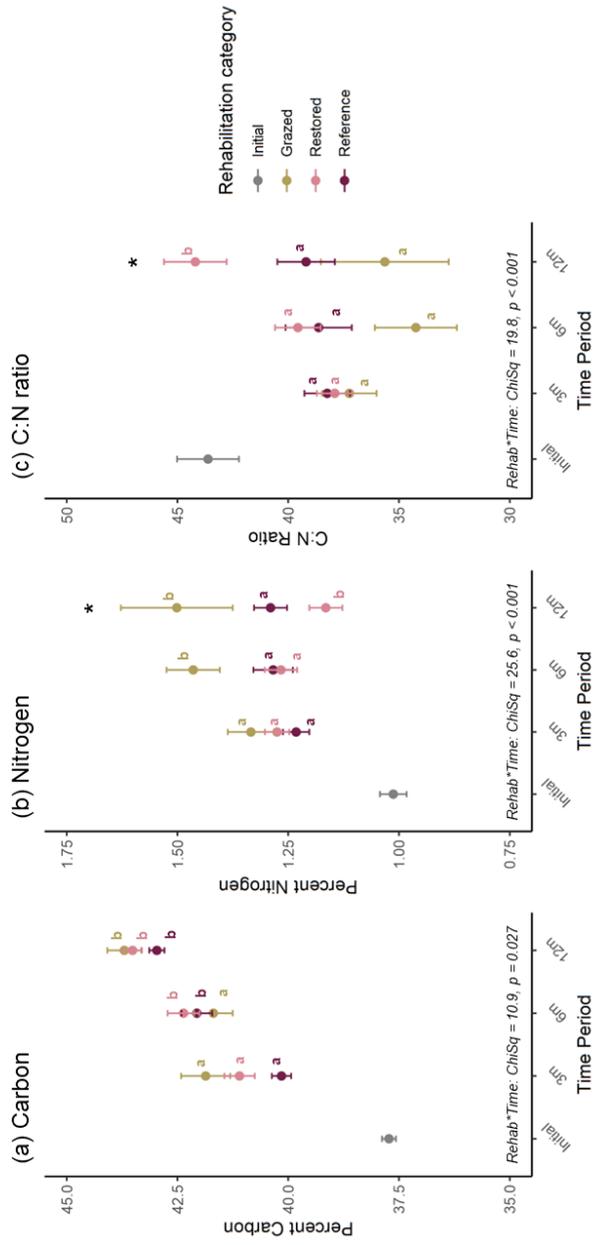


424 significant variables that make the Grazed sites different from the Restored and Natural
425 Reference sites. Conversely, percent organic carbon, percent total nitrogen, and vegetation
426 cover percent are the significant variables mostly influenced by Restored and Natural
427 Reference sites.



428

429 Figure 4: Decomposition parameters over 1 year of green tea, rooibos tea and *S. quinqueflora* root litters. (a) Proportion of mass remaining of
430 litters after 12 months of decay. Coloured circles are raw values with central black circles representing mean and standard error for the litters at
431 each rehabilitation category. (b) Decay rate constant (k , proportion mass lost per day) from single exponential decay fit. Data represent means
432 and standard error.



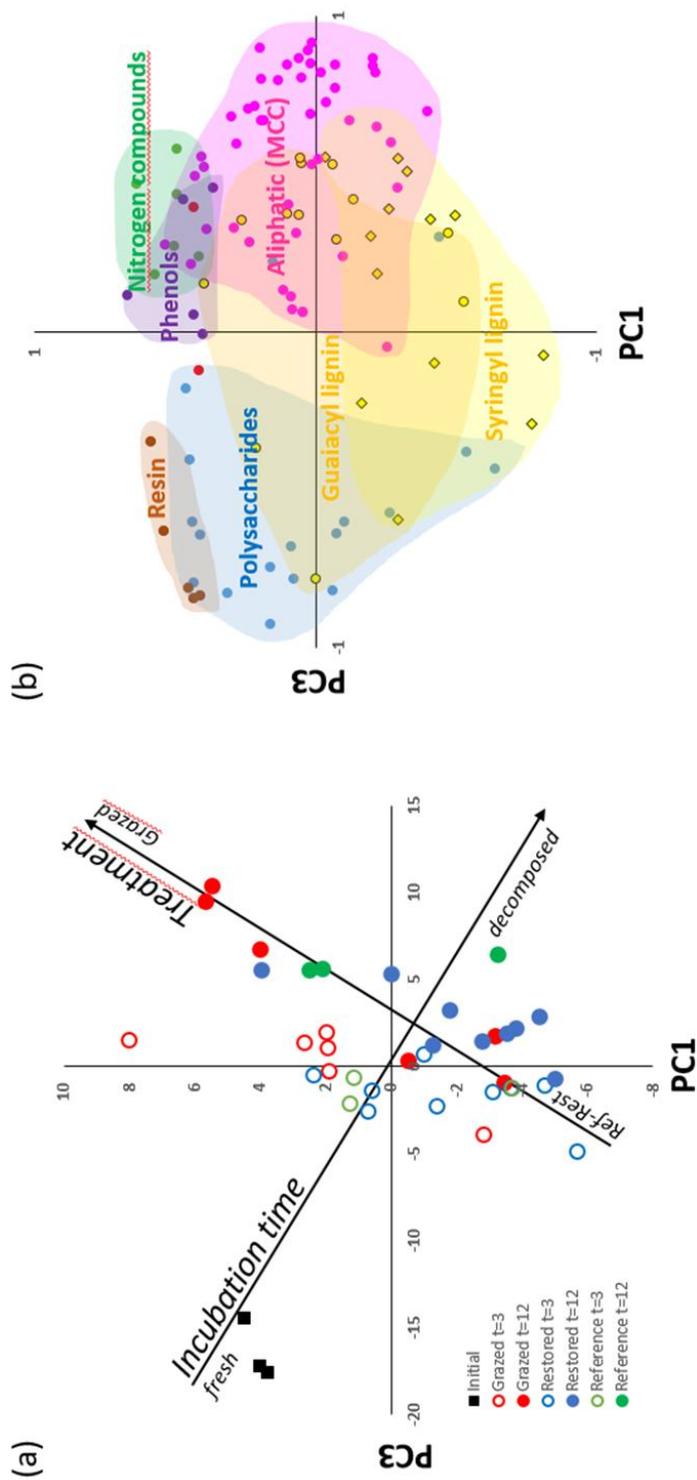
433

434 Figure 5: Elemental content of *S. quinqueflora* roots throughout 1-year of decomposition. (a) Percent organic carbon. (b) Percent nitrogen. (c)

435 Molar Carbon:Nitrogen ratios. Letters indicate post-hoc differences over time for each rehabilitation category individually, not across categories

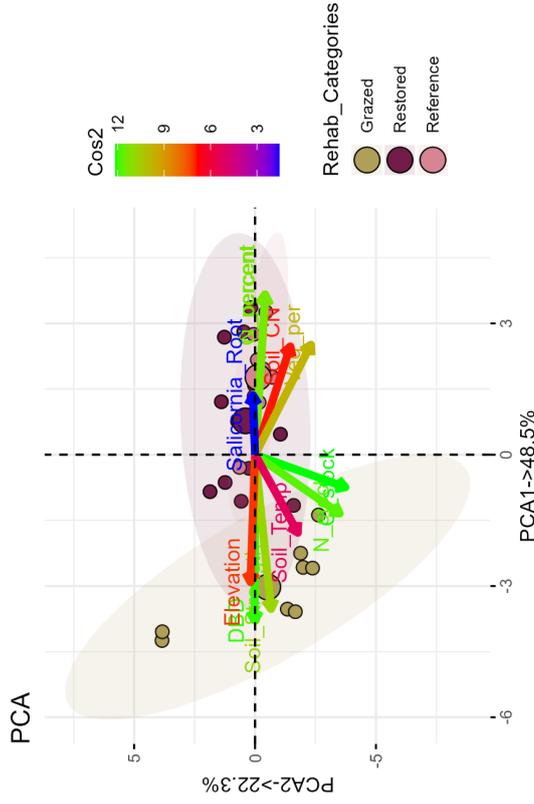
436 for each time point. Asterisks indicate a significant difference at 12 months between Grazed sites and both Restored and Natural Reference sites.

437 Data represent mean and standard error.



438

439 Figure 6: PC1-PC3 plots from principal components analysis of the molecular composition using pyrolysis-GC-MS of *S. quinqueflora* roots
 440 during one year of decomposition. a) PC scores of the samples; b) PC loadings of the pyrolysis products. MCC refers to methylene chain
 441 compounds (aliphatic organic matter).



442

443 Figure 7: Biplots of principal component analysis (PCA) of eleven belowground characteristic variables. Scores of the first two principal
444 components for eleven variables in different saltmarsh rehabilitation categories (Grazed, Restored and Reference) are shown. Cos^2 analysis
445 results of all the eleven variables are shown with the gradient colour bar. Variable include C_stock: carbon stock; N_stock: nitrogen stock;
446 C_percent: percent organic carbon; N_percent: percent total nitrogen; Soil_CN: soil C:N ratio; Soil_Strength: soil strength; DBD: dry bulk
447 density; Elevation: soil elevation; Soil_Temp: soil temperature; Veg_per: vegetation cover percent; Salicornia_Root: proportion of mass
448 remaining of *S. quinqueflora* root).



449 **4. Discussion**

450 **4.1. Site characteristics after exclusion fencing**

451 Restored and Natural Reference saltmarsh had lower soil strength and DBD, likely due to the
452 relief of pugging-related compaction and the development of extensive root biomass and
453 peat-rich soils with increased vegetation cover and diversity (Fig. 3). Grazing restrictions
454 have similarly shown to lower DBD and increase %OC and %N over time in both coastal and
455 terrestrial ecosystems (Sloane et al. 2021, Xiao et al. 2025). Low-laying, herbaceous or
456 succulent saltmarsh plant diversity was higher at Restored compared to Natural Reference
457 sites, likely in part due to our experimental focus on *S. quinqueflora*-dominant areas, which
458 could indicate a high diversity of ground-dwelling species and be linked to successional and
459 trophic shifts at the newly Restored sites (Nordström et al. 2015). Vegetation cover was
460 similar between Restored and Natural Reference sites, and the well-developed root layers in
461 the soil reflects the similar %OC and %N content of the Restored and Natural Reference soils
462 in the top 5 cm depths (Burger et al. 2024). Despite higher %OC and %N, Restored and
463 Natural Reference sites had similar carbon and nitrogen densities due to the high DBD at the
464 Grazed sites (Gifford and Roderick 2003), highlighting how surficial carbon stocks can
465 misrepresent ecological condition without structural and biological metrics. Together,
466 vegetation cover, diversity and surface %OC and %N similarities between the Restored and
467 Natural Reference suggest decadal timescales of vegetation and soil recovery, similar to those
468 found for other saltmarsh (Santini et al. 2019, Ledford et al. 2021) and mangrove (Adame et
469 al. 2018), and seagrass ecosystems (Greiner et al. 2013).

470 A strong positive correlation between DBD and soil strength showed significant reductions
471 after exclusion fencing, likely improving erosion resilience and root development (Daniel et
472 al. 2002, Cahoon et al. 2021). Although moderate compaction has been shown to reduce



473 erosion and indirectly promote vegetation growth (Pages et al. 2019), the soil strength at the
474 Grazed sites (mean 120 kPA) greatly exceeded that of vegetated saltmarsh areas (9-47 kPA)
475 (Turner 2011, Pages et al. 2019, Chirol et al. 2021, Pennings et al. 2021), which, along with
476 the lack of vegetation cover and visibly low presence of roots, indicates the degraded
477 condition of the sites. Our findings support the literature in showing how restoration by
478 exclusion fencing is a simple restoration method, that not only reduces compaction but also
479 helps re-establish soil conditions that support vegetation recovery and ecosystem functions.
480 These findings suggest that soil strength as a metric could be a good, simple indicator for
481 future monitoring of a range of belowground recovery responses.

482

483 **4.2. Belowground decomposition dynamics after exclusion fencing**

484 The significant differences in *S. quinqueflora* root chemistry among the rehabilitation
485 categories after 1 year of decomposition point to different processes influencing belowground
486 OM turnover. In the Restored and Natural Reference sites, the higher root C:N ratios and
487 relatively higher lignin compounds indicate selective the preservation of recalcitrant plant-
488 derived carbon (Arnaud et al. 2024). In contrast, the roots decomposing in the Grazed soils
489 had higher nitrogenous compounds and phenolic content and lower C:N ratios, pointing to
490 increased microbial metabolism and N-rich biomass accumulation (Meier and Bowman 2008,
491 Olsen et al. 2011). Given the Grazed soils also have low %CN content, together these
492 findings point to the mechanisms that underpin enhanced microbial turnover in nutrient-
493 limited, disturbed soils, while in restored and natural saltmarsh soils recalcitrant root carbon
494 is able to contribute to surficial carbon stocks. These differences in the molecular
495 composition of the decomposed root litter indicates some recovery of belowground



496 ecosystem functions at Restored sites through the putative re-establishment of microbial
497 processes and preservation of recalcitrant carbon (Meier and Bowman 2008).

498 In contrast to shifts in *S. quinqueflora* root chemistry during decay, we only found that mass
499 loss of the saltmarsh roots indicated a trend towards slower decomposition at the Natural
500 Reference sites. The root decay rate constant (k) did not differ significantly across
501 rehabilitation categories, and were overall higher (mean range 0.0052-0.0064 d^{-1}) than a
502 congener (*S. fruticosa* $\sim 0.0008 \text{ d}^{-1}$) and the global saltmarsh root average (0.0012 d^{-1}) (Curcó
503 et al. 2002, Scarton et al. 2002, Ouyang et al. 2017). We hypothesise that the differences in
504 our study are due to the relatively higher carbohydrate content in our mixed root materials
505 (~ 0.5 lignin:carbohydrate ratio), compared to more lignin-rich content of specific root types
506 (1.1-1.6 lignin:polysaccharide ratio) (Arnaud et al. 2024).

507 Interestingly, the standardised tea litter decay rates were less sensitive to rehabilitation
508 category than the native roots. Additionally, the mass remaining of the green tea litter was
509 higher in the Grazed soils, the opposite trend to root litter. These results emphasise how the
510 standardised litters, particularly green tea in short-term incubations, are highly influenced by
511 soil moisture and temperature, which enhances leaching (Petraglia et al. 2019, Trevathan-
512 Tackett et al. 2020a). As such, short-term studies (≤ 1 y) using standardised litters are likely
513 more influenced by abiotic drivers than biotic (Trevathan-Tackett et al. 2021), and thus not
514 sensitive to geographically-constrained space-for-time restoration studies (Ibanez-Alvarez et
515 al. 2022).

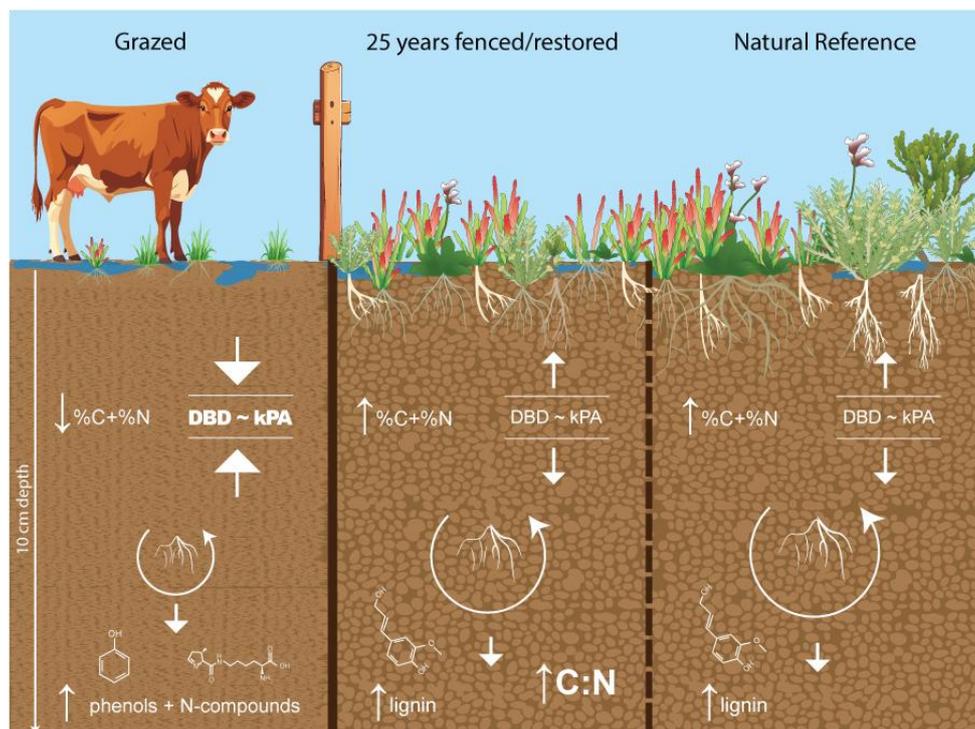
516

517 **4.3. Multi-indicator approach to assess saltmarsh soil recovery after restoration**

518 By combining multi-indicator and space-for-time approaches, we showed that exclusion
519 fencing improves physical, chemical and biological soil properties over 25 years, with the



520 restored temperate saltmarshes closely resembling natural saltmarsh ecosystems (Figure 8).
521 Removal of ungulates enhances surficial soil quality through preservation of recalcitrant
522 carbon, which is likely correlated to both enhanced vegetation cover that promotes root
523 production and allochthonous OM capture (Figure 8) (Mueller et al. 2019). In contrast, the
524 highly compacted, historically grazed soils in this study have higher saltmarsh root turnover,
525 potentially due to a combination of low nutrients, less frequent inundation and higher
526 temperatures (Figure 8). Moreover, disturbance of coastal soils releases reactive organic
527 substances, leading to disturbance in the microbial community composition that stimulates
528 CO₂ emission by using complex organic compounds that remain unreactive in the absence of
529 such disturbance (Ward et al. 2019, Macreadie et al. 2025). These conditions promoted
530 microbial decomposition of the fresh root material and indicate that the elevated carbon and
531 nitrogen stocks at the Grazed sites are reflective of changes in soil physical structure rather
532 than improvements in soil quality or ecosystem function. Any form of disturbance, including
533 changes in temperature, nutrient levels or oxygen availability, can alter microbial community
534 composition, structure and metabolism, potentially triggering the mineralisation of otherwise
535 stable carbon. Thus, the present study attests to the capacity of restoration practices, such as
536 exclusion fencing, for the stabilisation of organic carbon.



537

538 Figure 8: Conceptual design of the physical, chemical and biological surficial belowground
539 responses to saltmarsh exclusion fencing. Ungulate removal leads to reduced soil compaction
540 evidenced through reduced dry bulk density (DBD) and soil strength (kPA), concomitant with
541 enhanced vegetation, root biomass and soil carbon (%C) and nitrogen (%N). Restored and
542 natural saltmarsh conditions were also characterised by improved carbon stability through
543 reduced root decomposition and selective preservation of recalcitrant lignin. Root
544 decomposition in Grazed soils was characterised by increased microbial nitrogen and
545 phenolic compounds.

546

547 Some surficial soil metrics, such as soil strength, soil/root percent carbon and nitrogen
548 content and chemical shifts during root decay, are more sensitive soil indicators of exclusion
549 fencing restoration than others, particularly standardised litter decay and surficial carbon



550 stocks. For the latter, carbon stocks integrate bulk density and carbon content, and each varies
551 in their response to grazing and restoration. In our study, grazed soils had higher bulk density
552 but lower carbon content, whereas restored and reference soils showed the opposite pattern
553 (Figure 8). The combined effect on calculated carbon stocks may then show no difference (as
554 in Harvey et al. 2019), despite clear differences in each component. While it is worth noting
555 that the depths we investigated (0-10 cm) were shallower than standard for blue carbon stock
556 assessments (30 – 100 cm), the surficial soils were expected to be the most responsive to
557 changes to land management, including vegetation establishment and growth resulting in root
558 turnover and soil capture (Burden et al. 2013, Gulliver et al. 2020, Arnaud et al. 2024), as
559 well as compaction impacts of ungulates. By taking a targeted and multi-parameter approach
560 to measuring surficial soil condition, our study reveals novel, quantifiable changes in
561 microbial processes and carbon preservation 25 years after restoration intervention that
562 underpin soil function and carbon sequestration (Figure 8).

563 Importantly, this study provides a snapshot of recovery 25 years after restoration intervention,
564 showing Restored sites approaching Natural Reference conditions. While encouraging, it
565 does not clarify if recovery occurs gradually or follows a nonlinear trajectory. By linking
566 low-cost monitoring methods (e.g. soil strength) with more advanced metrics of recovery
567 (e.g. decay metrics and molecular indicators of OM composition) and understanding the
568 interaction between these, more accessible approaches are realised for continuous monitoring
569 of responses to restoration beyond that of carbon stocks and vegetation assessments. This
570 integrated approach better captures recovery dynamics, informing restoration benefits and the
571 condition of ecosystem functions, focusing on carbon quality over quantity (Strobl et al.
572 2019). As such, our study provides novel molecular indicators, linked to more commonly
573 used soil and decomposition metrics, to produce insight into the mechanisms that contribute
574 to soil formation and function in restored saltmarsh ecosystems.



575

576 **Data availability**

577 Data will be made available upon request to the corresponding author(s).

578

579 **Supplementary material**

580 [Link inserted by Copernicus publications]

581

582 **Author Contribution statement**

583 S.K.B.O., P.I.M., P.E.C. & S.T-T. conceptualised and designed the study. S.K.B.O. and S.T-T.

584 collected the data. S.T-T., A.A., J.K. & S.N. analysed the data. S.K.B.O., A.A. & S.T-T. wrote

585 the first draft of the manuscript. All authors contributed to the drafting and revising of the

586 manuscript.

587

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606

607



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