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1 **Carbon sequestration and biogeochemical cycling in a saltmarsh subject to coastal managed**
2 **realignment**

3 A. Burden ^{a*}, A. Garbutt ^a, C. Evans ^a, D. L. Jones ^b, D. Cooper ^a,

4 ^a *Centre for Ecology and Hydrology, Environment Centre Wales, Deiniol Road, Bangor, Gwynedd, LL57*
5 *2UW, UK*

6 ^b *School of the Environment, Natural Resources & Geography, Bangor University, Bangor, Gwynedd,*
7 *LL57 2UW, UK*

8 * Corresponding author: Email: anrd@ceh.ac.uk; Phone: +44 (0) 1248 374537 ; Fax: +44 (0) 1248
9 362133

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11 realignment

12 Regional index terms: UK, east coast, Tollesbury

13 **Abstract**

14 Globally, wetlands provide the largest terrestrial carbon (C) store, and restoration of degraded
15 wetlands provides a potentially important mechanism for climate change mitigation. We examined
16 the potential for restored saltmarshes to sequester carbon, and found that they can provide a
17 modest, but sustained, sink for atmospheric CO₂. Rates of C and nutrient cycling were measured and
18 compared between a natural saltmarsh (high- and low-shore locations), claimed arable land on
19 former high-shore saltmarsh and a managed realignment restoration site (high- and low-shore) in
20 transition from agricultural land to saltmarsh 15 years after realignment, at Tollesbury, Essex, UK.
21 We measured pools and turnover of C and nitrogen (N) in soil and vegetation at each site using a
22 range of methods, including gas flux measurement and isotopic labelling. The natural high-shore site
23 had the highest soil organic matter concentrations, topsoil C stock and below-ground biomass,
24 whereas the agricultural site had the highest total extractable N concentration and lowest soil C/N
25 ratio. Ecosystem respiration rates were similar across all three high-shore sites, but much higher in
26 both low-shore sites, which receive regular inputs of organic matter and nutrients from the estuary.

27 Total evolution of ^{14}C -isotopically labelled substrate as CO_2 was highest at the agricultural site,
28 suggesting that low observed respiration rates here were due to low substrate supply (following a
29 recent harvest) rather than to inherently low microbial activity. The results suggest that, after 15
30 years, the managed realignment site is not fully equivalent to the natural saltmarsh in terms of
31 biological and chemical function. While above ground biomass, extractable N and substrate
32 mineralisation rates in the high-shore site were all quite similar to the natural site, less dynamic
33 ecosystem properties including soil C stock, C/N ratio and below-ground biomass all remained more
34 similar to the agricultural site. These results suggest that reversion to natural biogeochemical
35 functioning will occur following restoration, but is likely to be slow; we estimate that it will take
36 approximately 100 years for the restored site to accumulate the amount of C currently stored in the
37 natural site, at a rate of $0.92 \text{ t C ha}^{-1} \text{ yr}^{-1}$.

38 **1 Introduction**

39 Globally, wetlands provide the largest terrestrial carbon stores, and restoration of degraded
40 wetlands provides a potentially important mechanism for climate change mitigation. To date, much
41 research has focused on restoring degraded peatlands, for example through re-wetting. However,
42 this research has highlighted uncertainties regarding its overall impact on C and greenhouse gas
43 balances, due to the potential for enhanced release of CH_4 following re-wetting (Strack et al., 2004;
44 Baird et al., 2009). There is an increasing in the potential for restored coastal wetland systems to
45 sequester large amounts of carbon (Craft et al., 2003; Shepherd et al., 2007; Santin et al., 2009;
46 Livesley and Andrusiak, 2012). Additionally, restoring coastal wetlands may avoid the offsetting
47 effects of enhanced methane production associated with peat re-wetting, due to the presence of
48 sulphates which allows sulphate-reducing bacteria to outcompete methanogens for energy sources
49 (Poffenbarger et al., 2011; Bartlett et al., 1987; Andrews et al., 2006). Therefore, per unit area,
50 restoration of coastal wetlands such as saltmarshes may contribute more to C sequestration, and
51 therefore to climate regulation, than peatlands. However, at present, evidence is sparse.

52 As well as carbon sequestration, saltmarshes provide a range of other ecosystem services.
53 These include immobilisation of pollutants (e.g. retention of diffuse nutrient and faecal pollutants
54 into accumulating sediments), flood defence and shore line erosion control and they are a significant
55 reservoir of wild species diversity (Jones et al., 2011). However historically, human activity has
56 focused on the land-claim ('reclamation') of saltmarsh for agriculture, and more recently for port
57 development leading to an estimate by French (1997) that 25% of the world's intertidal estuarine
58 habitat had been lost due to land claim. Accelerated sea-level rise also poses a threat to existing
59 saltmarsh through coastal squeeze, as sea defences restrict their natural landward migration to
60 higher elevation (Blackwell et al., 2004). Globally, efforts are now being made to restore and create
61 saltmarshes to mitigate historic losses and on-going development. Since the early 1990s, the driving
62 force for restoration was the unsustainable increasing cost of maintaining and upgrading existing sea
63 defences (Andrews et al., 2006). However, managed realignment is also undertaken for purposes of
64 habitat or biodiversity enhancement or restoration, for example in Europe, salt-marsh restoration
65 allows government compliance with the European Union Habitats Directive (C.E.G., 1992) which
66 states there should be 'no further net loss of coastal marsh' (UK Biodiversity Group, 1999). UK
67 targets aim to create 2240 ha of saltmarsh between 1999 and 2015, primarily via a process known as
68 'managed realignment'; the landward retreat of coastal defences and subsequent tidal inundation of
69 previously-claimed agricultural land (Garbutt et al., 2006).

70 In general, managed realignment schemes in the UK and elsewhere have shown that, with
71 relatively minimal pre-treatment and/or management of the area, allowing tidal ingress through a
72 breach of the existing seawall onto low-lying agricultural land will quickly produce intertidal mudflats
73 that are colonised by saltmarsh plants (French et al., 2000; Wolters et al., 2005). Managed
74 realignment sites are sinks of sediment and, given time, representative saltmarsh plant, invertebrate
75 and bird communities can become established (Garbutt et al., 2006). Newly created saltmarsh also
76 acts as a natural sea defence by attenuating tidal amplitude (Pethick, 2002). Self-sustaining plant
77 communities are often the primary goal of restoration efforts as they perform some of the desirable

78 functions of wetland ecosystems (Craft et al., 2002; Möller et al., 1999; van Andel, 1998). However,
79 many physical and functional processes such as nutrient cycling in these sites are poorly understood,
80 and it has yet to be shown that restored saltmarshes are functionally equivalent to referenced
81 systems and therefore whether they do effectively compensate for the loss of habitat as intended. In
82 particular, the capacity of managed realignment schemes to accumulate carbon following
83 conversion from agricultural land to saltmarsh has not been fully quantified.

84 This study measures and compares biogeochemical functioning between a 15 year old managed
85 realignment site in a state of transition from agricultural land to saltmarsh, relative to adjacent areas
86 of natural saltmarsh and arable land on former saltmarsh. Our three main objectives were: 1) to
87 compare general soil characteristics between the restored saltmarsh, natural saltmarsh and
88 agricultural sites; 2) to quantify and compare the organic matter, carbon and nitrogen pools at all
89 sites, and estimate how far soil carbon stocks at the restored site have progressed along a trajectory
90 between its former agricultural condition and the natural saltmarsh; 3) to investigate potential
91 differences in the dynamics of organic matter cycling by measuring in situ ecosystem respiration and
92 carbon mineralisation rates.

93 **2 Materials and methods**

94 **2.1 Site description**

95 This study was undertaken at the Tollesbury managed realignment site, adjacent natural marshes
96 and arable land of the Blackwater Estuary, south-east England (51°46'N, 0°51'E, Fig. 1) in July 2010.
97 The 21-ha restoration site had originally been a saltmarsh, but was claimed for agriculture in the late
98 18th century (Boorman et al., 1997). The sea defences were breached in August 1995, leaving a 50-
99 m wide opening and allowing tidal ingress to the site for the first time in over 150 years. The
100 construction of a new sea defence landward of the old embankment prevented tidal flooding of the
101 neighbouring arable fields which were claimed from saltmarsh at the same time as the managed
102 realignment site. The altitude of the site ranges from 0.9 m to 3.0 m above Ordnance Datum (OD),
103 with the major part of the site lying below 2.0 m OD (Garbutt et al., 2006). Mean high water neap

104 (MHWN) and mean high water spring (MHWS) tide levels for the Blackwater estuary are 1.50 m and
105 2.60 m OD, respectively (Pye and French, 1993).

106 There are two dominant plant communities within the managed realignment site. Above
107 1.75m the upper part of the site (referred to as 'restored high') is dominated by a species poor
108 *Puccinellia maritima* dominated community with occasional *Atriplex portulacoides*, *Spergularia*
109 *media* and *Suaeda maritima*. At the same elevation on the adjacent natural saltmarshes (referred to
110 as 'natural high') the plant communities are characterised by a diverse mix of saltmarsh plant species
111 with abundant *Limonium vulgare* and *P. maritima*, frequent *Salicornia europaea* agg. and
112 *Sarcocornia perennis* and occasional *Armeria vulgare*, *A. portulacoides*, *S. maritima* and *Triglochin*
113 *maritima*. Below 1.75m the lower part of the site (referred to as 'restored low') is dominated by
114 *Spartina anglica* and abundant *S. europaea* agg. At the same elevation on the adjacent natural
115 saltmarshes (referred to as 'natural low') the plant communities are dominated by *S. europaea* agg.
116 with occasional *S. maritima*. Landward of the managed realignment site land claimed from saltmarsh
117 in the 1800s is farmed for wheat (*Triticum aestivum*) and other crops.

118 **2.2 Experimental design**

119 Two different elevations (2.5 and 1.75 m above OD) within the managed realignment site were
120 chosen to best represent the dominant plant communities as described above. The equivalent
121 elevations on the adjacent marshes were determined through a topographic survey. Elevation was
122 used as a surrogate for tidal inundation to ensure that the plant communities within the de-
123 embankment sites and reference marshes received equivalent submergence frequencies, and was
124 checked by observing the depth and extent of the incoming tide for each site. No visual differences
125 were observed. At the time of survey, the wheat crop had been harvested from the agricultural land
126 adjacent to the site, however, the soil had not been tilled for subsequent crops and stubble
127 remained at the surface.

128 A split-plot experimental design was used with six locations sampled at each site location.
129 Sampling locations were situated at the same elevation for the two high shore sites, and for the two

130 low shore sites. The six sampling locations at each site were in two clusters of three, with the two
131 clusters separated by 150m and within-cluster spacing of 10m. This arrangement provides an
132 estimate of spatial variability (Fig. 1). With six sampling locations in two high shore, two low shore
133 and one agricultural site, this gave 30 individual sample sites in all. Soil cores (4 cm diameter by 30
134 cm depth) were taken from within the footprint of each of the greenhouse gas monitoring chambers
135 (see below) after the third day of gas sampling. Each core was split into 3 sections (0-10 cm, 10-20
136 cm and 20-30 cm) which were analysed separately. A second soil core was taken for below ground
137 plant biomass measurements only. All field work took place in July 2010.

138 **2.3 Soil characteristics**

139 Electrical conductivity (EC) and pH were measured in 1:1 (w/v) soil:distilled water extracts (Smith
140 and Doran, 1996). Moisture content was determined by measuring the weight loss after drying the
141 soil at 105°C overnight. Organic matter content was determined as the percent weight loss after
142 ignition overnight at 375°C. Available ammonium (NH_4^+) and nitrate (NO_3^-) were determined in 1:5
143 (w/v) soil: 0.5 M K_2SO_4 extracts following the method described in Jones et al. (2005) and the
144 colorimetric analysis procedures of Mulvaney (1996) and Miranda et al. (2001). Water soluble
145 phosphorous (P), sodium (Na), potassium (K) and calcium (Ca) were determined using 1:5 (w/v)
146 soil:distilled water extracts following shaking (1 h) and centrifugation (6000 g, 15 min) to remove
147 particulate material. P was determined colorimetrically using the molybdate blue/ascorbic acid
148 procedure of Murphy and Riley (1962) while Na, K and Ca were determined with a PFP7 Flame
149 Photometer (Jenway flame photometer (Bibby Scientific Ltd, Staffs, UK). Soluble humic substances in
150 the water extracts were estimated by measuring the UV absorbance of the extracts at 254 nm (US-
151 EPA, 2005). Bulk density was measured using bulk density rings with a volume of 45.2 cm³. Samples
152 were collected from the soil surface only. Samples were dried at 105°C for 72 hours and the dry
153 mass divided by the volume of the bulk density ring. Total soil C and N were measured by
154 combustion on a TruSpec CN Analyser (Leco Corp, St Joseph, MI).

155

156 The soil carbon pool was estimated by multiplying the bulk density by the percentage carbon figures.
157 It was therefore only estimated for the surface of the soil where bulk density values were available.
158 The restored high marsh per year increase in carbon was derived by taking the difference in the soil
159 carbon pool between the agricultural and restored high shore sites and dividing by the number of
160 years since managed realignment (i.e. 15 y). The estimate of how long it would take for the soil
161 carbon pool of the restored high shore site to become equivalent to the natural high shore site was
162 calculated by dividing the difference between the agricultural and natural high shore site soil carbon
163 pool by the per year increase of the restored high shore site, assuming that this was equivalent to
164 the agricultural site pre-restoration.

165 **2.4 Gas flux measurements**

166 Ecosystem CO₂, CH₄ and N₂O emissions were measured using dark static chambers within two hours
167 of high water. Gas sampling with the static chambers was carried out on 3 successive days to take
168 account of temporal variability. Placement of the static chambers was marked with canes allowing
169 return to the exact same position on consecutive days. On the first day both clusters of 3 at each
170 land use/elevation combination were sampled giving 30 sample locations in total. On days two and
171 three only one cluster of each land use/elevation combination was sampled, giving 15 sample
172 locations on the second and third days. After placement of the chambers on the soil surface, gas
173 samples were taken with a syringe from each chamber after 0, 15, 30, 45 and 60 minutes, injected
174 into evacuated gas chromatograph vials and analysed within one week. In addition to these samples,
175 duplicate samples were taken from one chamber for each treatment to test reproducibility. All gas
176 samples were analysed using a Perkin Elmer Clarus 500 Gas Chromatograph (GC) equipped with a
177 Poropak QS (80-100 mesh) analytical column and turbomatrix 40 headspace auto analyser. N₂O was
178 detected using an electron capture detector (ECD) at 400°C, sample oven at 40°C; CH₄ was detected
179 using a flame ionisation detector (FID) at 375°C, sample oven at 40°C equipped with a methaniser.
180 Carrier gas pressure was 138 kPa, and injection pressure 160 kPa, all other controls were as Perkin
181 Elmer standard setup. The GC was calibrated using bottled gas with a known concentration of CO₂,

182 CH₄ and N₂O (CryoService Ltd., Worcester, UK) and this gas was used for quality control (QC) at set
183 points throughout each sample run. Gas fluxes were then calculated on an hourly basis using the
184 following calculations:

$$185 \quad C_m = (C_v \times M \times P) / (R \times T)$$

186 where C_m is the mass per volume (expressed as mg m⁻³), C_v is the gas concentration (expressed as
187 mg dm⁻³), M is the molecular weight of the trace species (e.g. 12 for Carbon), P is the barometric
188 pressure, T is the chamber temperature in Kelvin (°C + 273.15) at time of sample, and R is the gas
189 constant.

190 The per hour flux (F , mg m⁻² h⁻¹) was then calculated by:

$$191 \quad F = V \times C_{rate} / A$$

192 where V is the internal volume of the enclosure including collar volume (expressed as m³), A is the
193 area of the collar enclosed surface (m²), C_{rate} is the change in gas concentration (i.e. $C_m t_1 - C_m t_0$)
194 over the enclosure period.

195 **2.5 Carbon substrate mineralisation**

196 A ¹⁴C-isotopically labelled C substrate was used to estimate carbon mineralisation rates in soil as
197 described in Simfukwe et al. (2011). The C substrate consisted of ¹⁴C-labelled shoots of *Lolium*
198 *perenne* (L.) with a specific activity of 12.3 kBq g⁻¹. The ¹⁴C-enrichment of *Lolium perenne* plant
199 material was performed by pulse labelling with ¹⁴CO₂ at a constant specific activity according to Hill
200 et al. (2007). To characterise the ¹⁴C label in the plant material, a sequential chemical fractionation
201 was performed according to Jones and Darrah (1994). Briefly, 50 mg of finely ground plant material
202 was sequentially extracted in 8 ml deionised water for 30 min at 85°C, 8 ml 20% ethanol for 30 min
203 at 80°C, 5 ml 0.3% HCl for 3 h at 95 °C and 5 ml 1 M NaOH for 1 h at 95 °C. After each extraction
204 step, the sample was centrifuged (5000 *g*, 15 min), the supernatant removed and its ¹⁴C content
205 determined using Optiphase 3[®] Scintillation fluid (PerkinElmer, Waltham, MA) and a Wallac 1404
206 Liquid Scintillation Counter (PerkinElmer Corp., Waltham, MA). For each soil, 10 g was placed into a
207 sterile 50 cm³ polypropylene container and 100 mg of the ¹⁴C-labelled complex C substrate was then

208 added to the soil. A vial containing 1 M NaOH was then placed above the soil and the polypropylene
209 containers hermetically sealed. The $^{14}\text{CO}_2$ capture efficiency of the NaOH traps was >95%. The soils
210 were then placed in the dark in a climate-controlled room (10°C) and the NaOH traps exchanged
211 every 3 days for 24 days. The $^{14}\text{CO}_2$ in the NaOH traps was determined by liquid scintillation counting
212 as described above.

213 Of the total ^{14}C contained in the plant material and subsequently added to soil, $32.9 \pm 1.5\%$
214 was extractable by water, $4.2 \pm 0.2\%$ by ethanol, $16.8 \pm 0.6\%$ by HCl, $27.5 \pm 0.4\%$ by NaOH and $18.5 \pm$
215 2.2% was insoluble residue. These components approximately correspond to the readily
216 decomposable or neutral-detergent soluble C (water and ethanol soluble), cellulose and
217 hemicellulose (HCl soluble) and lignin (NaOH soluble and insoluble) fractions of organic matter
218 respectively (Domisch et al., 1998; Ekschmitt et al., 2008; Moorhead and Sinsabaugh, 2006).

219 **2.6 Statistical analysis**

220 A linear mixed effects model (lme) was used to describe the data using R version 2.13.2
221 ($y \sim \text{Site} * \text{Depth}, \text{random} = \sim 1 | \text{Location/sample}$). We also fitted models excluding respectively the
222 depth and site effects to test the need for their inclusion in the model. On all but one occasion both
223 regime and depth were significant ($p < 0.05$), confirming the need for both fixed effects to be
224 included. For each variable we also tested for the significance of differences between each regime
225 and depth pair. Separate analyses by depth and by regime were also carried out. A similar approach
226 was taken for the gas measurements replacing 'depth' with 'day' in the analysis. As there was no
227 significant difference in gas measurements between days within sites, the average flux over the 3
228 days of measurement was analysed. For the carbon substrate mineralisation, the final data points
229 were used in the lme model (i.e. total evolution of $^{14}\text{CO}_2$ within incubation period of 25 days –
230 expressed as % of ^{14}C -substrate added to the soil). This approach enabled the raw data to be
231 analysed accounting for replication at the level of the experimental unit or site ($n=5$). For analysis we
232 used the \log_{10} of all variables other than pH.

233 **3 Results and discussion**

234 **3.1 General soil characteristics**

235 There were significant differences ($p < 0.05$) between sites in all of the soil properties measured. Soil
236 conductivity was highest at the natural high shore marsh, lowest at the agricultural site, and
237 intermediate at the restored high shore site (averages of 12.08, 0.14 and 4.40 mS cm^{-1} respectively,
238 Table 1). These large differences highlight the influence of seawater on both the natural and
239 restored sites, and of freshwater on the agricultural site. There was no significant difference
240 between the restored and natural low shore sites ($p = 0.582$) and all sites showed a decrease in
241 conductivity with depth. The sodium, potassium and calcium data, as would be expected, reflected
242 the differences observed in the conductivity data – they were all significantly ($p \leq 0.009$) higher at
243 the natural high shore site and lowest at the agricultural site ($p \leq 0.010$, Table 1).

244 **3.2 Plant biomass and soil organic matter, C and N pools**

245 The restored high shore site was found to have approximately twice as much above ground plant
246 biomass compared to the natural high shore site ($p = 0.037$, Fig. 2) due to it being dominated by a
247 monoculture of *P. maritima*. On the other hand, there was 16 times more below ground plant
248 biomass in the natural high shore site than at any of the other sites sampled (average of 11.5 kg m^{-2}
249 compared to 0.7 kg m^{-2} for the other four sites; Fig. 2) due to the species-rich vegetation consisting
250 of long lived perennials with woody tap roots. This translated into the natural high shore site soil
251 having significantly greater ($p < 0.05$) organic matter content (and therefore less mineral material)
252 than all other sites at all three depths (average of 21.8% compared to an average of 5.2% for the
253 other four sites, Table 1); this appears consistent with data collected from created *Spartina*
254 *alterniflora* marshes along the North Carolina coast (Craft, 2000), which showed that macro-organic
255 matter (MOM) content increased with age of the created marsh. The agricultural site had
256 significantly lower ($p < 0.05$) soil organic matter content than all other sites at depth 0-10 cm ($p \leq$
257 0.004) but was not significantly different ($p < 0.05$) to the two restored sites at depths 10-20 cm and
258 20-30 cm. This suggests that even after 15 years of inundation, soils below 10 cm depth retain

259 properties characteristic of the agricultural soil – an idea also supported by Craft (2000) in
260 constructed *Spartina alterniflora* saltmarshes in North Carolina. Spencer et al. (2008) found similar
261 evidence within a restored site 8 years after managed realignment, and hypothesized that the relic
262 land surface may have formed an aquaclude which prevents vertical soil water movement. This idea
263 is further supported in the current study by the pH and humic substances data, which both increased
264 with depth in the restored high shore site to values that were more comparable to the agricultural
265 site than the natural high shore site (supplementary Table 1).

266 The natural and restored low shore sites had the lowest soil carbon pool (average of 13.7
267 and 10.9 kg m⁻³ respectively, Fig. 2) and were not significantly different from each other (p = 0.260).
268 This is not surprising as the low shore sites are both inundated daily and are dominated by pioneer
269 annual species. In contrast, the natural high marsh site had a significantly greater soil carbon pool
270 than all other sites (p ≤ 0.016, average of 31.1 kg m⁻³), whilst the restored high shore and agricultural
271 sites had a similar soil carbon pool (p = 0.621) of 22.1 and 20.7 kg m⁻³ respectively (Fig. 2). This
272 suggests that there has been, at best, only a small overall increase in the soil carbon pool of the
273 restored high-shore site in the 15 years since managed realignment, and that the site is thus likely to
274 take many more years to accumulate an equivalent carbon store to the natural system.

275 Assuming that the restored high site previously resembled the agricultural site (see
276 methods), the estimated rate of carbon accumulated at the Tollesbury managed realignment site
277 was calculated to be 92.4 g C m⁻³ yr⁻¹, or 0.92 t C ha⁻¹ yr⁻¹. This is within the estimated UK saltmarsh
278 carbon sequestration range of 0.64 – 2.19 t C ha⁻¹ yr⁻¹ proposed by Cannell et al. (1999). We
279 therefore estimate that it would take approximately 100 years for the restored site to accumulate
280 the amount of carbon currently stored in the natural site. This is similar to the figure estimated by
281 Craft et al. (2003) of up to 70 years for the total organic carbon pool to become equivalent to natural
282 within a created *S. alterniflora* marsh. The 100 year estimate also corresponds that of Crooks et al.
283 (2002) for the length of time it could take vegetation in realignment sites to resemble that of natural
284 marshes, which (as noted above) has a substantial influence on the size of the soil organic matter

285 pool. It has been suggested, however, that the organic matter formed in constructed marshes
286 contains a greater proportion of labile organic compounds (Craft et al., 2003) which are turned over
287 more quickly by the microbial community and could result in constructed or restored wetlands being
288 less effective at sequestering carbon over time; this is considered further below.

289 The C/N ratio of the soil was significantly greater at the natural high shore site than all other
290 sampling sites ($p \leq 0.004$, Table 1). The restored high marsh site had slightly (albeit not significantly)
291 higher C/N than the agricultural site, which is again consistent with the interpretation that this site is
292 slowly transitioning towards the soil conditions currently observed at the natural marsh. This accords
293 with other terrestrial data that suggest it takes a very long time, even centuries, to reverse historic
294 enrichment of nutrient poor pools.

295 Measured total inorganic nitrogen at the agricultural site (2.46 mg kg^{-1} dry weight, 97.7% as
296 oxidised N, Table 1) was 2.5 times higher than at any of the other sites, and significantly higher ($p <$
297 0.05) at all sites except the natural high shore site. This is indicative of both nutrient enrichment
298 (wheat is cultivated at this site and is presumably fertilised) and aerobic conditions, both of which
299 favour nitrification and therefore increased nitrate versus ammonium concentrations. In contrast,
300 ammonium concentrations were at least four times higher at all other sites when compared to the
301 agricultural site, and ammonium was the dominant form of mineral N at the natural high, managed
302 high and managed low sites. These data suggest that managed realignment, which has led to the
303 reinstatement of anaerobic soil conditions that promote denitrification and leaching, has resulted in
304 rapid decreases in extractable nitrate levels. On the other hand, the presence of extractable
305 ammonium in all samples from all these sites ($0.27 - 0.92 \text{ mg kg}^{-1}$ dry weight, Table 1) suggests a
306 continued supply of mineral N via organic matter mineralisation in both natural and restored
307 conditions, consistent with the relatively low measured C/N ratios of soil organic matter (Table 1).
308 The fact that ammonium concentrations were not significantly higher in the restored high shore site
309 compared to the natural high site suggests that, despite soil C/N remaining lower in the restored

310 site, N mineralisation rates (and therefore nutrient N supply) may have returned to pre-agricultural
311 levels following managed realignment.

312 **3.3 Dynamics of organic matter cycling**

313 CH₄ and N₂O fluxes were near-zero for all sites and on all sampling occasions, with concentrations at
314 or close to ambient air concentrations (Table 2) – a result reflecting those of Livesley and Andrusiak
315 (2012) who found that CH₄ and N₂O fluxes were close to zero in temperate saltmarsh in south
316 eastern Australia. No significant between-site differences were observed. CH₄ production is known
317 to be inhibited by the presence of sulphate (e.g. from seawater), due to competition between
318 sulphate reducing bacteria and methanogens (Bartlett et al., 1987; Andrews et al., 2006).
319 Poffenbarger et al. (2011) proposed that the salinity regime required for methane flux to be
320 negligible was 18 µg l⁻¹ – well within the expected range of salinity at Tollesbury. However, a similar
321 study of a restored saltmarsh in the estuary of the River Torridge, Devon, UK, 6 months after
322 managed realignment, concluded that managed realignment could result in increased production of
323 N₂O, due the combination of high residual soil nitrogen levels from the agricultural site, and the
324 reinstatement of dry-wet cycles following tidal reconnection (Blackwell et al., 2010). The River
325 Torridge experiment was laboratory-based, and measured fluxes over simulated tidal cycles, which
326 may explain the difference in results (an average of 0.65 mg N₂O-N m⁻² h⁻¹ compared to zero in our
327 study). However, if tidal pumping – tidal forcing of seawater into the coastal aquifer – does in fact
328 flush CO₂ and other carbon forms out of re-flooded soils (an idea proposed by Kathilankal et al.,
329 2008) then our results should demonstrate maximum gaseous flux as they were taken within two
330 hours prior to high tide. Given the limited number of sampling occasions, we cannot draw clear
331 conclusions about the overall magnitude of N₂O flux at the Tollesbury sites, other than to note that
332 no evidence of a measurable flux was recorded at any of the sites on any of the sampling occasions.

333 Ecosystem respiration (R_{eco}) measurements using dark chambers indicated substantial CO₂
334 production rates, with the highest fluxes recorded for the restored low shore site, followed by the
335 natural low shore site (Fig. 3). Overall, between-site differences were significant ($p < 0.001$)

336 suggesting that both of the low shore sites were more microbially active, turning over carbon inputs
337 at a higher rate than either of the high shore sites. The natural high shore site had the lowest CO₂
338 flux but was not significantly lower than the restored high shore or agricultural site ($p \geq 0.3$)
339 suggesting that rates of carbon cycling among these three sites were similar. There was some
340 indication however of an inverse relationship between R_{eco} and C/N for these three sites, consistent
341 with higher rates of carbon turnover at the more nutrient-rich agricultural site. It is likely that lower
342 R_{eco} values at the agricultural site relative to the low shore sites are a consequence of reduced
343 substrate supply following the harvesting of the crop (see below). Based on our measurements, we
344 did not detect a clear influence of substrate quality on respiration rates, as suggested by Craft et al.
345 (2003), although we acknowledge that the dataset only covers a short time-period.

346 The total evolution of ¹⁴C-isotopically labelled C substrate as CO₂ (expressed as % of ¹⁴C-
347 substrate added to the soil) was highest at the agricultural site (Fig. 4). This result contrasts with the
348 low measured ecosystem respiration flux but, as noted above, the respiration measurements
349 occurred after crop harvesting at the site, when substrate inputs would have been very low. As well
350 as the direct production of CO₂ from plant respiration, the presence or absence of vegetation has
351 been shown to significantly affect microbial activity in soils via the supply of litter inputs and root
352 exudates for microbial respiration (Oburger and Jones, 2009). The observation that added substrate
353 was rapidly respired at the agricultural site indicates that potential rates of microbial carbon
354 turnover are higher than those observed at the time of sampling.

355 Among the natural and restored sites, mineralisation of ¹⁴C-labelled substrate was
356 consistently lower, and the restored high shore site was not significantly different to the natural high
357 shore site ($p = 0.107$, Fig. 4). As the added substrate was of high molecular weight carbon, results
358 should provide an insight into differences in carbon sequestration potential between sites, on the
359 basis that this material has the potential to be retained in the soil rather than respired, whereas low
360 molecular weight fractions would likely be utilised by the microbial community in all sites. The
361 results suggest an overall slower turnover rate, and therefore greater carbon storage potential, in

362 the natural site, when compared to the agricultural soil. The absence of a significant difference in
363 turnover rates between natural and restored high shore sites further suggests that, 15 years post-
364 restoration, rates of carbon sequestration at the Tollesbury managed realignment site have now
365 returned to those characteristic of the natural marsh.

366 At both low shore sites, measured substrate utilisation rates were higher than at the natural
367 and restored high shore (significantly different at $p < 0.05$ for all combinations other than the
368 restored high versus natural low sites, $p = 0.218$, Fig. 4), but lower than at the agricultural site.
369 Higher substrate utilisation rates at the low versus high shore natural and restored sites are
370 consistent with the ecosystem respiration measurements, again suggesting faster carbon turnover
371 rates at these sites. Higher extractable nitrate and phosphate, and lower soil C/N ratios, further
372 suggest that this rapid turnover is linked to higher nutrient availability, possibly due to tidal recharge
373 of nutrients.

374 **4 Conclusions**

375 In the UK, managed realignment is primarily undertaken for purposes of habitat and biodiversity
376 enhancement or restoration, or for coastal defence. Carbon and nutrient cycling are rarely
377 considered when these schemes are developed and monitored, let alone used as success criteria
378 (which currently only consider vegetation development). However, with a growing policy emphasis
379 on the wider ecosystem service implications of land-management, it is clear that enhanced carbon
380 sequestration could provide an additional benefit resulting from restoration, whilst reversion of the
381 nitrogen cycle to the low-nutrient levels characteristic of natural ecosystems may be a prerequisite
382 for full vegetation recovery. Our data suggest that managed realignment reduces nitrogen
383 mineralisation rates towards those of natural saltmarsh levels, but that soil C/N ratios remain well
384 below those of the natural site, suggesting that complete recovery to natural conditions may be far
385 slower. Similarly, the soil carbon pool of the restored site was more similar to the agricultural site
386 than the natural marsh, suggesting that there has been at best only a small overall increase in the
387 carbon pool of the restored high-shore site in the 15 years since managed realignment. On the other

388 hand, carbon mineralisation rates at the restored site were similar to the natural site, and lower
389 than the agricultural site, suggesting that the soil carbon pool of the restored site will ultimately
390 converge with that of the natural marsh. Our calculations predict that this will take approximately
391 100 years.

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Fig. 1. Experimental design at the Tollesbury managed realignment site, adjacent natural marshes and arable land of the Blackwater Estuary, south-east England (51°46'N, 0°51'E). Each circle represents one sampling location where three replicates were taken. Within replicate distances were 10 m, and between sampling location distances were in the order of 150 m. Open circles = high marsh, closed circles = low marsh and grey circles = agricultural.

Fig. 2. Above and below ground biomass and calculated soil carbon pool measured at all 5 sites. Values represent means \pm standard deviation. The site effect was evaluated using a linear mixed effects model, the p value of which is displayed. Significant differences between site means are denoted by different letters.

Fig. 3. Ecosystem respiration (R_{eco}) measured for all 5 sites. Values represent means \pm standard deviation. The site effect was evaluated using a linear mixed effects model, the p value of which is displayed. Significant differences between site means are denoted by different letters.

540 **Fig. 4.** Carbon mineralisation rates measured (as a % of total ¹⁴C-substrate added) for all 5 sites.
 541 Values represent means ± standard deviation. The site effect was evaluated using a linear mixed
 542 effects model on the final data points (total evolution within incubation period of 25 days –
 543 expressed as % of ¹⁴C-substrate added to the soil), the *p* value of which is displayed. Significant
 544 differences between site means are denoted by different letters.

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550 **Table 1.** Soil properties measured at all 5 sites. Site means (n = 18) are presented ± standard
 551 deviation. For bulk density n = 6. The site effect was evaluated using a linear mixed effects model.
 552 Significant differences (*p* < 0.05) between site means are denoted by different letters.

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	Agricultural		Restored		Natural		Restored		Natural	
			High		High		Low		Low	
pH	7.9 ± 0.5	<i>b</i>	7.2 ± 0.3	<i>c</i>	6.4 ± 0.2	<i>a</i>	7.6 ± 0.3	<i>bc</i>	7.8 ± 0.1	<i>b</i>
Soil conductivity (mS)	0.1 ± 0.1	<i>b</i>	4.4 ± 1.2	<i>c</i>	12.1 ± 2.7	<i>a</i>	5.9 ± 1.9	<i>c</i>	6.7 ± 0.6a	<i>b</i>
Total oxidised N (mg kg ⁻¹ dry weight)	2.41 ± 1.28	<i>b</i>	0.08 ± 1.00	<i>a</i>	0.07 ± 0.07	<i>a</i>	0.11 ± 0.16	<i>a</i>	0.37 ± 0.27	<i>c</i>
Ammonium (mg kg ⁻¹ dry weight)	0.06 ± 0.02	<i>b</i>	0.61 ± 0.58	<i>c</i>	0.92 ± 0.44	<i>a</i>	0.34 ± 0.25	<i>cd</i>	0.27 ± 0.19	<i>d</i>
Total inorganic N (mg kg ⁻¹ dry weight)	2.46 ± 1.28	<i>a</i>	0.69 ± 0.57	<i>c</i>	0.99 ± 0.47	<i>ab</i>	0.46 ± 0.29	<i>c</i>	0.64 ± 0.32	<i>bc</i>
Bulk Density (g cm ⁻³)	1.30 ± 0.13	<i>b</i>	0.84 ± 0.21	<i>c</i>	0.29 ± 0.03	<i>a</i>	0.42 ± 0.07	<i>a</i>	0.60 ± 0.03	<i>d</i>
Organic matter content (%)	3.8 ± 0.5	<i>b</i>	5.5 ± 1.2	<i>c</i>	21.8 ± 4.6	<i>a</i>	5.5 ± 1.3	<i>c</i>	6.0 ± 0.7	<i>c</i>
Humic substances (RAU cm ⁻¹)	0.90 ± 0.46	<i>b</i>	1.15 ± 0.85	<i>b</i>	0.33 ± 0.16	<i>a</i>	0.67 ± 1.07	<i>a</i>	0.15 ± 0.03	<i>c</i>
Sodium (g kg ⁻¹ dry weight)	0.02 ± 0.01	<i>b</i>	4.18 ± 1.57	<i>c</i>	39.49 ± 10.63	<i>a</i>	7.59 ± 4.63	<i>e</i>	10.31 ± 1.87	<i>d</i>
Potassium (g kg ⁻¹ dry weight)	0.02 ± 0.01	<i>b</i>	0.28 ± 0.09	<i>c</i>	1.75 ± 0.31	<i>a</i>	0.58 ± 0.29	<i>d</i>	0.64 ± 0.13	<i>d</i>
Calcium (g kg ⁻¹ dry weight)	0.07 ± 0.05	<i>b</i>	0.24 ± 0.09	<i>c</i>	3.54 ± 0.87	<i>a</i>	0.50 ± 0.31	<i>c</i>	0.63 ± 0.08	<i>c</i>
Phosphate (mg kg ⁻¹ dry weight)	3.42 ± 1.76	<i>b</i>	1.95 ± 0.35	<i>c</i>	1.02 ± 0.95	<i>a</i>	3.08 ± 1.14	<i>b</i>	3.38 ± 0.43	<i>b</i>

C (%)	1.7 ± 0.3 ^b	2.2 ± 0.4 ^b	9.7 ± 2.4 ^a	2.4 ± 0.7 ^b	2.2 ± 0.3 ^b
N (%)	0.17 ± 0.03 ^b	0.21 ± 0.03 ^c	0.72 ± 0.17 ^a	0.23 ± 0.05 ^c	0.25 ± 0.03 ^c
C/N ratio	10.0 ± 0.7 ^{bc}	10.5 ± 0.8 ^c	13.5 ± 0.7 ^a	10.0 ± 1.3 ^{bc}	8.7 ± 0.3 ^b

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559 **Table 2.** Organic matter cycling measurements at all 5 sites. Gas flux site means (n = 12) are
 560 presented \pm standard deviation. For carbon substrate mineralisation ($^{14}\text{CO}_2$ evolution) n = 6. The site
 561 effect was evaluated using a linear mixed effects model. Significant differences ($p < 0.05$) between
 562 site means are denoted by different letters. Non significant results are recorded as *ns* ($p > 0.05$).
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	Agricultural	Restored High	Natural High	Restored Low	Natural Low
CO_2 flux ($\text{mg m}^{-2} \text{h}^{-1}$)	69.8 ± 39.1 ^a	48.5 ± 57.5 ^a	18.7 ± 26.1 ^a	615.1 ± 200.4 ^c	264.6 ± 117.8 ^b
CH_4 flux ($\text{mg m}^{-2} \text{h}^{-1}$)	0.02 ± 0.12 ^{ns}	-0.01 ± 0.04 ^{ns}	-0.01 ± 0.03 ^{ns}	0.08 ± 0.08 ^{ns}	0.02 ± 0.02 ^{ns}
N_2O flux ($\text{mg m}^{-2} \text{h}^{-1}$)	-0.02 ± 0.04 ^{ns}	0.00 ± 0.06 ^{ns}	-0.02 ± 0.06 ^{ns}	0.00 ± 0.06 ^{ns}	-0.02 ± 0.06 ^{ns}
$^{14}\text{CO}_2$ evolution (% evolved of total ^{14}C -substrate added)	24.2 ± 1.2 ^b	14.6 ± 2.2 ^{ac}	12.3 ± 1.1 ^a	19.9 ± 2.8 ^d	16.3 ± 2.4 ^c

564 **Supplementary table 1.** Soil properties measured at all 5 sites by depth. Individual depth means by site (n = 6) are presented \pm standard deviation. Dry wt
 565 indicates dry weight.

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Depth (cm)	Agricultural			Restored High			Natural High			Restored Low			Natural Low		
	0-10	10-20	20-30	0-10	10-20	20-30	0-10	10-20	20-30	0-10	10-20	20-30	0-10	10-20	20-30
pH	7.6 \pm 0.3	7.8 \pm 0.5	8.3 \pm 0.2	7.0 \pm 0.1	7.1 \pm 0.2	7.4 \pm 0.3	6.4 \pm 0.2	6.4 \pm 0.2	6.5 \pm 0.3	7.4 \pm 0.3	7.6 \pm 0.3	7.7 \pm 0.2	7.8 \pm 0.1	7.8 \pm 0.1	7.8 \pm 0.2
Soil conductivity (mS)	0.2 \pm 0.1	0.1 \pm 0.1	0.1 \pm 0.0	5.6 \pm 0.7	4.4 \pm 0.8	3.2 \pm 0.8	15.0 \pm 1.1	11.8 \pm 1.5	9.4 \pm 1.4	8.3 \pm 0.7	5.4 \pm 0.6	4.1 \pm 0.5	7.1 \pm 0.4	6.8 \pm 0.6	6.2 \pm 0.4
Total oxidised N (mg kg ⁻¹ dry wt)	2.43 \pm 1.22	2.24 \pm 1.47	2.62 \pm 1.38	0.05 \pm 0.02	0.11 \pm 0.15	0.07 \pm 0.09	0.13 \pm 0.08	0.06 \pm 0.05	0.03 \pm 0.02	0.25 \pm 0.23	0.05 \pm 0.05	0.04 \pm 0.02	0.65 \pm 0.23	0.30 \pm 0.11	0.15 \pm 0.12
Ammonium (mg kg ⁻¹ dry wt)	0.07 \pm 0.03	0.05 \pm 0.02	0.05 \pm 0.02	0.38 \pm 0.33	0.33 \pm 0.18	1.13 \pm 0.71	0.86 \pm 0.27	0.98 \pm 0.62	0.92 \pm 0.44	0.45 \pm 0.20	0.18 \pm 0.05	0.40 \pm 0.35	0.34 \pm 0.15	0.17 \pm 0.10	0.30 \pm 0.27
Total inorganic N (mg kg ⁻¹ dry wt)	2.50 \pm 1.24	2.29 \pm 1.47	2.67 \pm 1.39	0.44 \pm 0.33	0.44 \pm 0.32	1.19 \pm 0.66	0.99 \pm 0.34	1.04 \pm 0.65	0.95 \pm 0.43	0.70 \pm 0.14	0.23 \pm 0.06	0.44 \pm 0.35	0.99 \pm 0.17	0.47 \pm 0.18	0.45 \pm 0.25
Organic matter content (%)	4.1 \pm 0.3	3.9 \pm 0.5	3.3 \pm 0.6	6.6 \pm 1.1	5.0 \pm 0.4	4.8 \pm 0.9	24.4 \pm 3.4	23.2 \pm 3.8	17.8 \pm 3.9	6.7 \pm 0.8	4.9 \pm 1.0	4.8 \pm 1.3	5.8 \pm 0.5	6.2 \pm 1.0	5.9 \pm 0.4
Humic substances (RAU cm ⁻¹)	0.89 \pm 0.36	1.04 \pm 0.67	0.69 \pm 0.10	0.51 \pm 0.49	1.00 \pm 0.70	1.92 \pm 0.72	0.45 \pm 0.16	0.30 \pm 0.12	0.23 \pm 0.12	0.16 \pm 0.04	0.38 \pm 0.16	1.47 \pm 1.63	0.17 \pm 0.01	0.14 \pm 0.02	0.14 \pm 0.03
Sodium (g kg ⁻¹ dry wt)	0.02 \pm 0.01	0.02 \pm 0.01	0.02 \pm 0.01	5.53 \pm 1.28	3.99 \pm 1.34	3.01 \pm 1.02	47.36 \pm 10.66	39.90 \pm 7.38	31.19 \pm 7.65	13.23 \pm 3.09	5.79 \pm 1.57	3.76 \pm 1.05	11.17 \pm 1.55	10.44 \pm 2.05	9.31 \pm 1.76
Potassium (g kg ⁻¹ dry wt)	0.03 \pm 0.01	0.02 \pm 0.01	0.01 \pm 0.00	0.36 \pm 0.06	0.26 \pm 0.07	0.21 \pm 0.06	1.92 \pm 0.35	1.66 \pm 0.26	1.65 \pm 0.27	0.88 \pm 0.28	0.49 \pm 0.13	0.36 \pm 0.10	0.66 \pm 0.04	0.65 \pm 0.16	0.62 \pm 0.16
Calcium	0.08 \pm 0.07	0.06 \pm 0.04	0.06 \pm 0.03	0.33 \pm 0.07	0.22 \pm 0.05	0.16 \pm 0.04	4.14 \pm 0.77	3.57 \pm 0.74	2.90 \pm 0.71	0.89 \pm 0.11	0.38 \pm 0.12	0.22 \pm 0.05	0.70 \pm 0.06	0.62 \pm 0.06	0.57 \pm 0.06

(g kg⁻¹ dry wt)

Phosphate	2.95 ± 0.69	4.30 ± 2.64	2.81 ± 0.77	2.05 ± 0.45	1.75 ± 0.09	2.05 ± 0.38	0.91 ± 0.57	1.10 ± 1.27	1.05 ± 1.05	4.40 ± 0.84	2.63 ± 0.56	2.21 ± 0.38	3.35 ± 0.40	3.53 ± 0.37	3.27 ± 0.55
(mg kg ⁻¹ dry wt)															
C	1.8 ± 0.3	1.8 ± 0.3	1.5 ± 0.5	2.6 ± 0.2	2.1 ± 0.3	1.9 ± 0.2	10.9 ± 1.1	11.2 ± 1.3	7.1 ± 2.1	2.7 ± 0.4	2.2 ± 0.6	2.2 ± 1.0	2.3 ± 0.1	2.2 ± 0.4	2.1 ± 0.4
N	0.18 ± 0.02	0.18 ± 0.02	0.15 ± 0.03	0.24 ± 0.02	0.20 ± 0.02	0.19 ± 0.01	0.81 ± 0.10	0.81 ± 0.11	0.54 ± 0.16	0.27 ± 0.04	0.22 ± 0.04	0.21 ± 0.06	0.26 ± 0.02	0.25 ± 0.03	0.24 ± 0.04
C:N	9.9 ± 0.5	10.1 ± 0.6	9.8 ± 1.3	11.0 ± 0.7	10.5 ± 0.8	10.0 ± 0.6	13.5 ± 0.7	13.8 ± 0.7	13.1 ± 0.6	9.8 ± 0.6	10.0 ± 1.2	10.1 ± 2.0	8.6 ± 0.3	8.9 ± 0.4	8.7 ± 0.3

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