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Publication Date

2023-09-01

DOI

10.1016/j.soilbio.2023.109078

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2 Chemical identity of carbon substrates drives differences in denitrification and N₂O
3 reduction within agricultural soils

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15 *Keywords:*

16 Denitrification; N₂O reduction; C substrate; C-to-N ratio; soil carbon

17

18 **Abstract**

19 Rates of nitrous oxide (N₂O) production from agricultural soils are highly variable across
20 space and time. Improving predictions of N₂O emissions will require improving our understanding of
21 the drivers of denitrification and the sources of variability in the rates of N₂O production between
22 soils and over time. While the amount of available carbon (C) is a known control on denitrification
23 and N₂O reduction, relatively little attention has been paid to the effect of the chemical identity of C
24 substrates on rates of denitrification and N₂O reduction. We investigated the effects of twelve
25 different C-substrate additions on the production and reduction of N₂O in five soils taken from two
26 distinct agricultural locations in Michigan under multiple land uses. We provided additions of
27 glucose, cellulose, N-acetyl-glucosamine, chitin, amino acids, protein, vanillyl alcohol, lignin, citrate,
28 succinate, methanol, and water in laboratory denitrification potential assays to determine the effects
29 of denitrifier C preference on denitrification rates. We found that amino acids, protein, and organic
30 acids stimulated the greatest rates of denitrification potential across all land uses. Similarly, we found
31 these same substrates caused the most N₂O reduction, resulting in the lowest net concentrations of
32 N₂O. Soils from agricultural rotations without cover crops had overall lower rates of denitrifier
33 activity, leading to less net N₂O production compared to soils from other land uses. In general, C-
34 utilization patterns were similar among all soils, and C-substrate identity had a much stronger effect
35 than land use. Here, we demonstrate that the chemical identity of available C gives rise to wide
36 variability in rates of denitrification and N₂O reduction.

37 **1. Introduction**

38 Nitrous oxide (N₂O) is a greenhouse gas with a global warming potential nearly 300 times
39 greater than that of CO₂ (IPCC, 2014). Nearly half of global N₂O emissions are anthropogenic, with
40 agriculture accounting for the largest share (Tian et al., 2020). Net emissions of N₂O are the result of
41 multiple microbially mediated processes, but denitrification is thought to be the predominant N₂O

42 generation pathway in agricultural systems (Opdyke et al., 2009; Liang and Robertson, 2021).
43 Denitrification is an anaerobic, respiratory metabolism where inorganic N species are reduced in a
44 stepwise manner within the electron transport chain to generate ATP through oxidative
45 phosphorylation, resulting in the production of N₂O and/or N₂. The balance of these two end products
46 determines the contribution of denitrification to greenhouse gas emissions.

47 There are multiple controls on the process of denitrification in general and on the end-product
48 ratio in particular (Firestone and Davidson, 1989). Even though the main drivers of denitrification are
49 known to include carbon (C), nitrogen (N), and O₂ availability, denitrification rates and net N₂O
50 emissions remain difficult to predict in the field and are subject to large spatial and temporal
51 variation. Much of this variation is due to the spatial distribution of the main drivers of denitrification
52 throughout the soil profile, giving rise to micro-scale variation and episodic fluxes in denitrification
53 rates (Groffman et al., 2009; Kuzyakov and Blagodatskaya, 2015). For example, anaerobic microsites
54 can exist within and between soil aggregates, promoting denitrification even within well-aerated soils
55 (Hojberg and Sorensen, 1993; Kravchenko et al., 2017; Schlüter et al., 2018). In addition, the
56 distribution of particulate C substrates throughout the soil creates zones of high N₂O production
57 (Parkin, 1987; K. Kim et al., 2020), but the effect of the chemical identity of these heterogeneously
58 distributed forms of C on rates of N₂O production and reduction has not been well described.

59 The importance of the quantity of available C seems obvious, with more available C yielding
60 more electrons and driving denitrification, but there are complex interactions that control how C
61 availability affects the balance of N₂O production and consumption. For instance, when C is limiting,
62 net N₂O production tends to be higher, due to a lower demand for terminal electron acceptors
63 (Pidello et al., 1996). Likewise, N₂O reduction has been shown to be inversely related to the
64 availability of alternative electron acceptors, such as O₂ or nitrate (NO₃⁻) (Firestone et al., 1980;
65 Miller et al., 2008; Senbayram et al., 2012). Therefore, by driving the consumption of terminal

66 electron acceptors, higher rates of C availability should stimulate more N₂O reduction and a lower
67 N₂O:N₂ ratio.

68 In addition to C quantity, one of the most important yet least explored factors determining
69 denitrification rates and N₂O production or consumption is the biochemistry of available C
70 compounds. Although the idea that the chemical identity of available C is a driver of denitrification
71 has been recognized for decades (e.g., de Catanzaro and Beauchamp, 1985), we still lack a clear
72 understanding of how substrate identity is tied to rates of denitrification, such as which chemical
73 characteristics are most important for denitrifiers. Multiple studies have come to widely different
74 conclusions on the effects of particular C substrates on denitrification. Some studies have described
75 glucose and other simple carbohydrates stimulating more denitrification than organic acids and
76 amino acids (Smith and Tiedje, 1979; Dendooven et al., 1996; Miller et al., 2008; Morley and Baggs,
77 2010), while others have found the opposite (Morley and Baggs, 2010; Morley et al., 2014).
78 Likewise, denitrifiers respond differently to whole plant residues compared to low-molecular-weight
79 C additions (de Catanzaro and Beauchamp, 1985; Senbayram et al., 2012; Giles et al., 2017). Much
80 of this inconsistency between studies comes from comparing experiments that were performed under
81 a variety of conditions, making it difficult to differentiate the effects of C chemistry from soil
82 characteristics and environmental or experimental factors, such as N availability and anaerobicity. By
83 comparing denitrifier C utilization between different soils within the same study, we can identify the
84 particular characteristics of C substrates that affect denitrification and whether C-based effects are
85 consistent across different soils and microbial communities.

86 The C compounds available in the soil possess various inherent characteristics that may
87 increase or decrease denitrification. For instance, the accessibility of C differs between monomeric
88 and polymeric compounds, with the degradation of polymeric residues by extracellular enzymes
89 often thought to be the rate-limiting step in the mineralization of complex C residues (Sinsabaugh,
90 1994). If depolymerization is limiting N₂O consumption by restricting the availability of C, then

91 additions of monomeric forms of C should result in lower net N₂O emissions than polymeric forms
92 because of the extra step involved in making C polymers accessible to denitrifiers. Moreover, the
93 C:N ratio of C substrates plays an important role in influencing denitrification and is also a good
94 predictor of N₂O production. As the C:N ratio narrows and the relative amount of N increases, N₂O
95 production tends to also increase (Huang et al., 2004; Millar and Baggs, 2004; Toma and Hatano,
96 2007). This is likely the result of low-C:N-ratio residues being a labile source of C since these
97 residues also tend to be easier to decompose. In addition to being a substrate for denitrification, N is
98 also a critical nutrient for microbial growth. Finally, the redox state of compounds can influence
99 denitrification, with more electron-rich, highly reduced substrates able to provide more electrons to
100 reduce more units of NO₃⁻ and therefore drive greater rates of denitrification. On the other hand,
101 more highly reduced substrates—such as simple carbohydrates—will often be available to organisms
102 with other anaerobic metabolisms, such as fermentation (Reddy et al., 1982; Pidello et al., 1996),
103 leading to competition that may reduce denitrifier access to such C sources but provide an ecological
104 opportunity for denitrifiers to specialize on more highly oxidized compounds such as organic acids.
105 Indeed, the organic acid succinate has been used as the C source in denitrifier isolation media
106 (Heylen et al., 2006). If competition with fermenters has influenced denitrifier C preference, then
107 compounds available more exclusively to denitrifiers, such as succinate or other organic acids,
108 should increase denitrification rates relative to simple sugars.

109 Denitrifier community composition and how different denitrifier species respond to different
110 C substrates is also likely to be a critical factor controlling N₂O production and reduction. Previous
111 research has demonstrated that individual denitrifier isolates possess their own C preferences and that
112 synthetic communities composed of denitrifiers with complementary C preferences can denitrify at
113 faster rates than other synthetic communities with overlapping resource niches (Salles et al., 2009,
114 2012). In addition, denitrifiers have differing capacities to carry out denitrification and N₂O
115 reduction. Community composition can be an important driver of N₂O emissions because denitrifier

116 communities vary across land uses, even within the same landscape (Cavigelli and Robertson, 2001;
117 Juhanson et al., 2017; Maul et al., 2019), and possess unique rates of denitrification and distinct
118 sensitivities to environmental factors (Cavigelli and Robertson, 2000; Krause et al., 2017; Maul et al.,
119 2019). These variations could extend to their use of C. In general, soil microbial communities from
120 separate land uses often differ in terms of C utilization. This is the basis for community-level
121 physiological profiling and popular techniques such as the Biolog plate assay (Garland and Mills,
122 1991). Underlying these patterns, differing legacies of C inputs between land-use histories can dictate
123 the C preference of soil microbial communities, giving rise to a home-field advantage where
124 microbes more quickly mineralize the C types they have historically been exposed to (Ayres et al.,
125 2009). In addition, extracellular enzyme activity has been linked to the quality and diversity of C-
126 input legacies, with rotational diversity and intercropping both increasing extracellular enzyme
127 activity (McDaniel et al., 2014; Curtright and Tiemann, 2021). Such land-use effects on enzyme
128 activity could potentially lead to differences in the amount of denitrification stimulated by polymeric
129 forms of C. Therefore, the C utilization profiles of denitrifiers may differ between land uses, with
130 higher overall rates of denitrification in soils with greater aboveground diversity. Such land use
131 effects could potentially account for discrepancies in C-utilization studies between denitrifiers from
132 different soils. However, how land use influences the C preference of denitrifiers has not yet been
133 examined.

134 Using lab incubations of soils from two agricultural field experiments with varying
135 management practices, we explore the interacting effects of land-use history, denitrifier community
136 structure, and C-compound quality and accessibility on N₂O production and consumption due to
137 denitrification. We hypothesized that in the same soils, the chemical identity of C inputs would result
138 in different levels of denitrification and N₂O reduction, specifically: (1) compounds with lower C:N
139 ratios will stimulate more denitrification, because of their ease of degradation; (2) monomeric
140 compounds will stimulate more gross N₂O production but lower net N₂O production than their paired

141 polymers, due to greater bioavailability of monomers; and (3) organic acids will stimulate more gross
142 N₂O production than reduced sugars, such as glucose, because organic acids will be preferentially
143 available to denitrifiers. Moreover, we hypothesized that (4) different land uses would lead to
144 denitrifier communities with distinct C-utilization profiles.

145 **2. Materials and Methods**

146 *2.1. Land Uses and Sampling*

147 We sampled soils from two locations in Michigan, USA. The first location was the W. K.
148 Kellogg Biological Station (KBS) (Hickory Corners, 42° 24' N, 85° 24' W), where we utilized field
149 treatments from the KBS Main Cropping System Experiment, a Long-Term Ecological Research site
150 established in 1989. Soils at this location are Typic Hapludalfs (fine-loamy, mixed, mesic; Table 1).
151 We sampled from three field treatments from this site: conventional agriculture, reduced-input
152 agricultural, and perennial switchgrass. The conventional and reduced-input agriculture treatments
153 are in a corn (*Zea mays*)-soybean (*Glycine max*)-wheat (*Triticum aestivum*) rotation with
154 conventional tillage and synthetic fertilizer inputs at locally recommended rates. The reduced-input
155 treatment receives a portion of its N inputs through winter leguminous cover crops after wheat
156 (*Trifolium pratense* L.). The reduced-input treatment also receives an annual ryegrass (*Lolium*
157 *multiflorum* Lam.) cover crop following corn. Both the conventional and reduced input treatments
158 receive herbicide applications to manage weeds. The perennial treatment previously contained
159 continuous alfalfa but was switched to continuous switchgrass (*Panicum virgatum* L.) in 2019.
160 Additional details relating to these soils can be found in Robertson and Hamilton (2015). Treatments
161 are organized in a randomized-block design, from which we sampled soils from four blocks. Soil
162 cores (1.9 cm diameter) were taken to a depth of 10 cm in April 2021, following a corn rotation and
163 prior to soybean planting.

164 The second set of soils was sampled from the Montcalm Research Center (MRC) (Montcalm,
165 43° 3' N, 85° 1' W). These soils are Oxyaquic Glossudalfs (fine, mixed, frigid; Table 1). We sampled
166 soils from two treatments of a field experiment established in 2015. Both treatments are in a potato
167 (*Solanum tuberosum*)-corn rotation and received the same conventional fertilizer and herbicide
168 applications, and they were both irrigated during the potato phase of the rotation. The two treatments
169 differed only in the use of cover crops; one treatment is seeded with a mixture of annual ryegrass and
170 hairy vetch (*Vicia villosa*), while the other treatment has no cover crops. Field treatments are
171 organized in a randomized-block design. We took soil cores to a depth of 10 cm in October 2020,
172 one week following potato harvest.

173 For each set of soil samples, soil cores were kept on ice in the field and brought back to the
174 lab for processing. Soils were sieved through a 2 mm mesh, and soil moisture content was assessed
175 gravimetrically. Soils were kept at 4°C until utilized in denitrification assays, within two weeks of
176 sampling.

177 2.2. Denitrification Assays

178 To assess denitrifier response to various C substrate additions, we modified a standard
179 denitrification enzyme activity assay protocol (Groffman et al., 1999). Specifically, 5 g of soil was
180 measured into 60 mL serum bottles. To each bottle, 5 mL of H₂O was added. After one hour, 10 mL
181 of KNO₃ solution (0.1 mg N mL⁻¹; bringing total solution volume to 15 mL) was added together with
182 one of 12 different C-substrate treatments to a final concentration of 4.4 mg C g⁻¹ soil (Table 2). The
183 different C substrates used were glucose, cellulose, N-acetyl-glucosamine (NAG), chitin, casamino
184 acid mix, soy protein isolate, vanillyl alcohol, lignin, citrate, succinate, and methanol. In addition, a
185 twelfth treatment contained no C addition. Casamino acid are a mixture of free amino acids produced
186 from an acid hydrolysis of casein protein; the acid hydrolysis removes most cystine and tryptophan.
187 Soluble forms of C were provided dissolved in the KNO₃ solution. Insoluble C treatments were

188 added in powder form together with an equivalent amount of KNO₃ solution to match the dissolved-
189 C additions. Jars were crimp capped with butyl-rubber septa. We then evacuated and flushed each jar
190 with UHP N₂ three times to atmospheric pressure to create an anaerobic atmosphere. Jars were
191 divided into two sets; one set received acetylene (C₂H₂) at 10% v/v and the other set received an
192 equivalent amount of N₂. C₂H₂ inhibits the enzyme responsible for the reduction of N₂O to N₂,
193 allowing for an estimation of gross versus net N₂O production (Yoshinari and Knowles, 1976).

194 Jars were kept on an orbital shaker at 150 rpm between gas sampling times, which were
195 determined based on earlier optimization studies that demonstrated nitrous oxide reductase (the
196 enzyme responsible for N₂O reduction) was fully induced after ~24 hours and that soil microbes had
197 not yet reached an exponential growth phase (Fig. S1). Headspace gas samples (3 mL) were injected
198 into pre-evacuated 12 mL vials at 24, 26, and 28 hours. The remaining volume of the gas tight vials
199 was filled with N₂. Following each gas sampling, the headspace removed from each jar was replaced
200 using either N₂ or a 90:10 mixture of N₂:C₂H₂. Gas samples were analyzed for N₂O and CO₂
201 concentrations on a TRACE 1310 (Thermo Fisher, USA) gas chromatograph equipped with an ECD
202 and TCD.

203 2.3. Statistics

204 Concentrations of N₂O in the jars that contained C₂H₂ were used for estimates of gross N₂O
205 production. Nitrous oxide concentrations in jars that did not receive C₂H₂ were used to estimate net
206 N₂O production. The proportional difference between gross and net N₂O production is often used to
207 scale N₂O emissions to total denitrification. When denitrification ends in the production of N₂O,
208 dN₂O will be high. Conversely, when N₂O is reduced fully to N₂, dN₂O will be low. We calculated
209 this value as dN₂O according to the following equation.

$$210 \quad dN_2O = \frac{N_2O_{net}}{N_2O_{gross}}$$

211 Statistical analyses were performed in R. Data from each study site, KBS and MRC, were
212 analyzed separately. For analysis, all concentration data were log transformed to achieve normality.
213 The *lme4* package (Bates et al., 2015) was utilized to create mixed-effects models with land-use,
214 substrate addition, and their interaction as fixed effects and field-treatment block as a random effect.
215 Marginal means were calculated using the *emmeans* package (Lenth et al., 2019). Using the *lmerTest*
216 package (Kuznetsova et al., 2017), type III tests for fixed effects were performed using the Kenward-
217 Roger method for calculating the denominator degrees of freedom. When fixed effects were found to
218 be significant, mean comparisons between substrate treatments within fields and between fields
219 within substrate were performed using Fisher's LSD at a Type I error rate of 0.05.

220 3. Results

221 3.1. C-Substrate Effects on Potential Gross N₂O Production

222 C substrate treatments had large effects on gross N₂O production in soils from all land uses
223 across both locations (Table 3). We found that amino acids and protein stimulated the most gross
224 N₂O production in each soil, regardless of land management, with up to 42 μg N₂O-N g⁻¹ soil d⁻¹
225 being produced from the amino-acid-amended soils from the KBS reduced input treatment (Fig. 1A).
226 Organic acids also consistently stimulated some of the largest amounts of gross N₂O production
227 (citrate produced 20 μg N₂O-N g⁻¹ soil d⁻¹ in reduced-input soils), followed by glucose (13.4 μg N₂O-
228 N g⁻¹ soil d⁻¹) and NAG (8.7 μg N₂O-N g⁻¹ soil d⁻¹). Vanillyl alcohol, lignin, and methanol had the
229 lowest rates of gross N₂O production. However, these substrates still stimulated significantly more
230 gross N₂O production than the no-C additions, which resulted in between 2.3 and 4.8 μg N₂O-N g⁻¹
231 soil d⁻¹ among the land uses at KBS and 2.4 and 2.9 μg N₂O-N g⁻¹ soil d⁻¹ in the MRC soils.

232 The monomeric compounds inconsistently stimulated more gross N₂O production than their
233 polymeric counterparts. Within each land use at KBS, glucose and amino acids stimulated about
234 twice as much gross N₂O production compared to cellulose and protein, respectively (Fig. 1A).

235 However, the denitrifier response to amino acids was stronger than to protein only in the KBS soils;
236 in the MRC soil with no cover crops, protein stimulated approximately 75% more gross N₂O
237 production than amino acids (Fig. 1B). Vanillyl alcohol never stimulated significantly more gross
238 N₂O production than lignin.

239 *3.2. C-Substrate Effects on Potential Net N₂O Production*

240 In each of the KBS soils, amino acid and protein additions resulted in the lowest rates of net
241 potential N₂O production (between 0.31 and 1.0 μg N₂O-N g⁻¹ soil d⁻¹), indicating the greatest N₂O
242 reduction (Fig. 2A). Organic acids also tended to have lower net N₂O production, but only the
243 succinate addition in the reduced-input treatment (0.59 μg N₂O-N g⁻¹ soil d⁻¹) was significantly
244 different from the no-C treatment (between 1.9 and 3.4 μg N₂O-N g⁻¹ soil d⁻¹). Meanwhile, in none of
245 the soils did any of the C additions result in net N₂O production significantly greater than that of
246 water alone, indicating that N₂O reduction kept pace with the N₂O production stimulated by each C
247 substrate.

248 Within MRC soils, proteins and citrate had the lowest net N₂O production with only 0.05 and
249 0.08 μg N₂O-N g⁻¹ soil d⁻¹ being produced by soils amended with protein in the cover cropped and
250 no-cover treatments, respectively (Fig. 2B). In contrast to the KBS soils, amino acids (~2.6 μg N₂O-
251 N g⁻¹ soil d⁻¹) did not result in lower net N₂O production than other substrates. In fact, amino acids
252 had significantly higher net N₂O production than water in the no-cover treatment (2.8 versus 1.4 μg
253 N₂O-N g⁻¹ soil d⁻¹). In the no-cover treatment at MRC, glucose had the highest amount of net N₂O
254 production (4.7 μg N₂O-N g⁻¹ soil d⁻¹), while cellulose, NAG, chitin, amino acids, and lignin also had
255 significantly higher net N₂O production compared to the water-only addition. In the cover crop
256 treatment, only glucose, cellulose, and lignin had significantly greater net N₂O production than the
257 water-only treatment (1.9 μg N₂O-N g⁻¹ soil d⁻¹).

258 The proportional difference between net and gross N₂O production, dN_2O , describes the
259 portion of total N₂O production that remains as N₂O following N₂O reduction. Across the KBS land
260 uses, proteins and amino acids had the lowest dN_2O (3.4–14%; Fig. 3A). Organic acids (7–24%) also
261 had a significantly lower dN_2O than other substrates in each of the KBS soils. Glucose and NAG had
262 an intermediate dN_2O . Within the conventional treatment, lignin had a low dN_2O comparable to that
263 of glucose and NAG. In addition, methanol-induced N₂O reduction was significantly greater than that
264 of water, but only in the conventional treatment. All other substrates had dN_2O values not
265 significantly different from that of water (~7%) in any land-use treatment at KBS. Protein and citrate
266 stimulated denitrification with the lowest dN_2O values in both treatments at MRC (Fig. 3B). Amino
267 acids and succinate also had low dN_2O values. In the cover crop treatment, NAG and chitin had
268 dN_2O values significantly lower than that of water. Within the conventional treatment, glucose,
269 vanillyl alcohol, and lignin had dN_2O values significantly higher than water.

270 *3.3. Land-Use Effects on Gross N₂O Production*

271 Among the KBS treatments, we observed that the conventional treatment had significantly
272 less potential N₂O production across most C substrate additions, with almost half as much N₂O
273 production in some additions (Fig. 1A). Interestingly, the reduced-input treatment tended to be more
274 similar in denitrifier C preferences to the perennial system than the conventional system, despite the
275 latter sharing a corn-soy-wheat rotation. Land-use differences were similar across most substrates.
276 Amino acids were the only substrate that did not have a significant land-use effect. At MRC, land-
277 use effects were not significant on gross N₂O production (Table 3).

278 *3.4. Land-Use Effects on Net N₂O Production*

279 At KBS, land use had a significant effect on net N₂O production, but this depended on the
280 substrate addition treatment (Table 3). The conventional-agriculture soils had significantly lower net
281 N₂O production than the other land uses in response to glucose, vanillyl alcohol, lignin, succinate,

282 and methanol (Fig. 2A). Protein resulted in higher net N₂O levels in the conventional-agriculture
283 soils compared to the other two treatments. The reduced-input and perennial systems tended to have
284 similar levels of net N₂O production across most substrates, but the reduced-input land use had
285 significantly lower net N₂O levels in response to succinate. When comparing dN₂O, a significantly
286 greater portion of N₂O remained when soils from the perennial management were amended with
287 glucose compared to soils under conventional management (Fig. 3A). Lignin stimulated a
288 significantly lower dN₂O in the conventional treatment compared to the other two land
289 managements. dN₂O was higher in the conventional treatment following protein addition and in the
290 perennial treatment following succinate, but these differences were not significant.

291 Among the MRC treatments, the cover cropped soils had higher net N₂O levels than the no-
292 cover-crop treatment and this effect did not depend on substrate additions (Table 3). Comparing
293 dN₂O, there were no significant land-use effects.

294 **4. Discussion**

295 Our results support the general hypothesis that the chemical identity of available C affects
296 rates of N₂O production and reduction. Although preliminary data indicated that exponential growth
297 did not occur after 24 hours in unamended soils, it appears that microbial growth may have been
298 stimulated with the addition of some of the substrates in the KBS soils (Fig. S2). In the glucose,
299 amino acid, protein, citrate, and succinate treatments, the rate of CO₂ production over 26 hours was
300 significantly greater than the rate of CO₂ production in the first 6 hours. The denitrification
301 stimulated by these substrates could therefore be caused by an increase in the number of denitrifiers
302 as well as, or instead of, greater activity of the extant denitrifier community. Stimulating growth is
303 one potential mechanism behind the substrate differences we observed; however, it is unlikely to
304 account for all the substrate differences. For instance, although substrate addition did not appear to
305 stimulate growth in MRC soils after 26 hours, denitrifiers from KBS and MRC soils still

306 demonstrated similar C-utilization characteristics with respect to N₂O production and reduction. In
307 addition, differences in the N₂O end-product ratio do not fully match the patterns in apparent
308 stimulated growth. Specifically, while glucose addition appeared to stimulate microbial growth in
309 KBS soils, it did not reduce dN₂O as much as other substrates.

310 By providing forms of C with particular chemical characteristics, we tested specific
311 hypotheses designed to help identify the various physiological and ecological mechanisms that may
312 account for the wide differences in denitrification rates that we observed between substrates. Below,
313 we discuss a handful of such mechanisms and whether they are supported by our data. We note at the
314 outset that the conditions of these assays were not representative of those in the field, and the
315 conclusions are therefore somewhat limited in their scope.

316 *4.1. C-to-N Ratio of Available Substrates*

317 Amino acids and proteins tended to stimulate the most N₂O production, with nearly ten times
318 more N₂O production than the no-C addition and a five-fold increase compared to glucose in the soils
319 from KBS (Fig. 1). In addition to C, amino acids supply varying levels of N, and in the field this
320 additional N could ultimately lead to increased denitrification since N availability is known to be a
321 primary control on denitrification (Wallenstein et al., 2006). For instance, among different plant-
322 residue additions, those with greater amounts of N tend to stimulate more N₂O production by
323 alleviating N limitation and providing the requirements for denitrification (de Catanzaro and
324 Beauchamp, 1985; Aulakh et al., 1991; Huang et al., 2004). However, in our experiments, all C
325 substrate additions also received non-limiting quantities of NO₃⁻, ensuring adequate amounts of N to
326 serve as electron acceptors. Further, the organic N of amino acids is in a reduced form and would
327 first need to be removed from the amino acid and then oxidized through either autotrophic or
328 heterotrophic nitrification to NO₃⁻ before being used as an electron acceptor in denitrification (Tiedje
329 et al., 1983). Since most nitrification pathways utilize ammonium monooxygenase, which requires

330 oxygen, the oxidation of amino-acid N to NO_3^- seems unlikely in our anaerobic assays (Martikainen,
331 2022). Moreover, among the C substrates we added was NAG, which also contains additional
332 organic N. But despite this additional N, we never observed NAG to stimulate more denitrification
333 than glucose. Therefore, the increased denitrification from amino acids and proteins is not due simply
334 to the provision of additional N to fuel denitrification.

335 While N availability *per se* might not explain the differences we observe among substrates,
336 compounds with more N tend to be more easily incorporated into biomass. Amino acids are the
337 building blocks of proteins, and environmental amino acids are rapidly recycled into new biomass
338 (Geisseler et al., 2009, 2010). Similarly, proteins are broken down into peptides and individual amino
339 acids, which are quickly utilized by soil microbes (Payne, 1976; Hill et al., 2011). A high availability
340 of these biochemical building blocks could stimulate growth as well as result in faster production of
341 denitrification enzymes and other metabolic machinery, leading to an increase in microbial activity.
342 The degree of C substrate incorporation into biomass—so called, C-use efficiency—can be measured
343 using stable-isotope-labelled C additions. However, when directly compared to glucose and other C
344 compounds, amino acids have tended to show lower C-use efficiencies (Brant et al., 2006; Frey et al.,
345 2013), suggesting they are not typically preferred for biomass any more than other substrates.
346 Nevertheless, past experiments measuring the relative C-use efficiency of amino acids have been
347 carried out under aerobic conditions, quite distinct from the anaerobic conditions in our experiment.
348 Since we did not track the C-use efficiency of denitrifiers in this study, we are unable to determine
349 whether direct incorporation of C substrates in new biomass is driving substrate differences in
350 denitrification rates.

351 Interestingly, although amino acids and protein stimulated the greatest amount of
352 denitrification, they also induced the most reduction of N_2O , resulting in the lowest dN_2O values.
353 While somewhat paradoxical, this indicates that these substrates stimulated N_2O reduction just as
354 much, or more than, N_2O production. Previous studies have reported on the effect of the amino acids

355 cysteine (Morley et al., 2014) and glutamate (Giles et al., 2017) on the denitrification end-product
356 ratio, with the former resulting in less N₂O reduction than glucose and the latter having no significant
357 difference. Our study is notable insofar as it demonstrates such high denitrification efficiency with an
358 amino acid mixture and proteins. The availability of substrates, such as amino acids and proteins, to
359 be incorporated into new biomass may be particularly important for N₂O reduction. If these
360 substrates also stimulated the greatest amount of microbial growth (Fig. S2), it could suggest that fast
361 growth may favor N₂O reduction as a trait of copiotrophic denitrifiers. When comparing different
362 denitrifying taxa, others have also found that denitrifiers with the greatest affinities for N₂O reduction
363 also happened to have the highest growth rates (Conthe et al., 2018; Qi et al., 2022). Nevertheless,
364 the available research linking N₂O reduction to other microbial traits such as growth rate is quite
365 limited, and we are hesitant to draw any premature conclusions regarding the ecological significance
366 of N₂O reduction to life-history strategies without additional study.

367 *4.2. Competition and Niche Partitioning*

368 We hypothesized that resource partitioning would result in some C compounds, namely
369 organic acids, being utilized more readily by denitrifiers than reduced sugars, which would be the
370 object of competition with other anaerobes. Evidence for niche partitioning between denitrifiers and
371 fermenters is hard to observe directly (Stevens et al., 1998). Paul and Beauchamp (1989) observed
372 the production and consumption of fermentation byproducts and suggested that syntrophies between
373 denitrifiers and fermenters improved the energy yield of available substrates. We found that citrate
374 did stimulate more denitrification than glucose and NAG (Fig. 1), and it also resulted in a greater
375 portion of N₂O being reduced (Fig. 3), supporting the possibility of niche differentiation between
376 denitrifiers and fermenters. However, succinate did not result in more denitrification or N₂O
377 reduction. Although denitrifiers did appear to prefer citrate over glucose, this did not reflect a

378 universal preference for organic acids. We are thus unable to conclude whether this C preference
379 emerged via resource partitioning with fermenters or by other ecological mechanisms.

380 Resource partitioning among denitrifier populations could also account for the particularly
381 high rates of denitrification following the addition of amino acids and protein. Rather than being a
382 single source of C, these additions were a mixture of different amino acids, each potentially available
383 to a different set of denitrifiers. These different forms of C can allow for niche partitioning and
384 complementarity in resource utilization, allowing more microbes to be metabolically active at the
385 same time (Goldfarb et al., 2011; Baran et al., 2015). Niche partitioning has been shown to increase
386 overall denitrification rates in synthetic communities of denitrifiers with complementary substrate
387 usage (Salles et al., 2009). Comparing the effects of C substrates within mixtures is, of course, more
388 representative of the soil environment (Henry et al., 2008) and demonstrates an important aspect of
389 the effect that C substrate identity has on denitrification rates. The single-substrate additions of our
390 assays were of course not fully representative of how denitrifiers would encounter C substrates in the
391 field.

392 *4.3. Substrate Bioavailability*

393 The bioavailability of C substrates will determine the rate of their utilization, with polymeric
394 forms of C being relatively less bioavailable than dissolved monomeric forms (Sinsabaugh, 1994).
395 To determine how this affected denitrification rates, we included pairs of polymers and their
396 composite monomers among our C addition treatments. At KBS, glucose and NAG tended to
397 stimulate more denitrification than their polymeric counterparts, cellulose and chitin, although the
398 difference between NAG and chitin was only statistically significant in one soil (Fig. 1A). The effect
399 of monomers on dN₂O values was more consistent, with monomers stimulating more N₂O reduction
400 relative to total denitrification (Fig. 3A). A greater supply of ready electron donors (monomers)
401 increases demand for terminal electron acceptors, thereby driving the reduction of N₂O to N₂ and

402 reducing dN₂O (Senbayram et al., 2012). While previous studies have demonstrated dN₂O to be
403 inversely related to C availability (Beauchamp et al., 1989; Weier et al., 1993; Miller et al., 2008;
404 Qin et al., 2017), they did not address the role of depolymerization in making substrates available.
405 Even though we provided all substrates in the same C-normalized quantity, monomeric substrates
406 were immediately available for rapid uptake and metabolism, while the bioavailability of polymeric
407 substrates was determined by the rate of extracellular depolymerization. In our study, this resulted in
408 a greater amount of N₂O being consumed when more biologically available monomers were
409 supplied.

410 The difference between monomers and polymers was not universal across all substrates and
411 land uses. Vanillyl alcohol never stimulated more denitrification or N₂O reduction than lignin. The
412 constituent monomers of lignin, such as vanillyl alcohol, are phenolic compounds that require highly
413 specialized biochemical pathways for their anaerobic degradation (Rabus, 2005). Although some
414 denitrifiers are capable of utilizing phenolic compounds as their sole C source, these microbes are
415 often fastidious and slow growing (van Schie and Young, 1998), so the lack of a large response of
416 denitrifiers to lignin and vanillyl alcohol in our study is not wholly unexpected. Among the soils
417 from KBS, the difference between monomers and polymers was greatest with amino acids and
418 proteins (Fig. 1A), but interestingly, proteins stimulated more gross N₂O production than amino acids
419 at MRC (Fig. 1B). Peptidase degradation of proteins was clearly not a rate limiting step in these soils,
420 and it was interesting to observe proteins stimulating more denitrification than amino acids. Proteins
421 can be broken down into peptide fragments, which may be more efficiently taken into cells than
422 individual amino acids (Matthews and Payne, 1980; Geisseler et al., 2010), potentially resulting in a
423 more rapid supply of C. Accordingly, within the rumen, peptides have been found to be utilized by
424 microbes more readily than amino acids (Wallace, 1996). The apparent preference for proteins over
425 amino acids could also be due to the specific composition of the two mixtures. The amino acid
426 addition was derived from a digestion of casein protein, whereas the protein addition was an isolation

427 of soy protein. While these contained similar levels of total amino acids overall, the specific
428 distribution and amounts of each can differ between the two. For example, tryptophan and cystine are
429 often lost in the acid hydrolysis step of prepared amino acid mixtures. The specific composition of
430 the protein mixture may have contained essential amino acids that were limiting to the denitrifiers in
431 the MRC soils or for which these microbes had a strong affinity (Wallace, 1996; Kajikawa et al.,
432 2002; Liu et al., 2020).

433 *4.4. Land Use and Soil Microbial Communities*

434 To determine the effect of land-use legacies on denitrifier C utilization we utilized different
435 field treatments from two locations in Michigan. Previous research has demonstrated dissimilar
436 denitrifier communities among soils at the KBS site that were functionally distinct with respect to the
437 sensitivity of nitrous oxide reductase to oxygen (Cavigelli and Robertson, 2001) (though only one of
438 the treatments utilized here was included in that research). Across both locations, we found that the
439 land uses that did not have cover crops tended to have the least potential for N₂O production across
440 most C substrate additions (Fig. 1). In general, cover cropping has been shown to increase overall
441 microbial activity (Tiemann et al., 2015; N. Kim et al., 2020). Therefore, the lower denitrification
442 potential in the land uses that did not include cover crops may be due to an overall reduction in
443 microbial activity, rather than an altered response of denitrifiers to substrate additions. It was also
444 interesting to note that, at KBS, denitrifier C utilization in the reduced-input treatment tended to be
445 more similar to C-utilization patterns in the perennial system rather than the conventional treatment,
446 which had the same crop rotation as reduced-input but without cover crops. This suggests that the
447 amount of time that plants cover the soil may be a stronger determinant of land-use effects rather
448 than specific plant composition or diversity, *per se* (Garland et al., 2021). Moreover, the N dynamics
449 of these soils have been extensively characterized previously (Millar and Robertson, 2015); despite
450 fewer fertilizer inputs, the reduced-input and switchgrass treatments have similar inorganic N levels

451 to the conventional treatment throughout the year, due to mineralization of cover crop residues and
452 soil organic matter. While the overall amount of available N remains similar, the form and rate of
453 release of this N may produce land-use differences between the treatments.

454 Among the treatments at each location, we did not find a strong effect of land use on the C-
455 utilization profiles of denitrifiers. The denitrifier communities from the soils we used in our
456 experiment appear to be functionally equivalent in their C preference. While representing distinct
457 management strategies, each of the land uses in this experiment were agricultural systems. Other
458 agricultural management factors, such as weed management or aboveground biomass removal, may
459 be more influential in determining denitrifier C-utilization profiles than the chemical composition of
460 plant residue inputs. For instance, previous studies have found that soil type is more important than
461 plant identity in determining denitrification rates (Graf et al., 2016). In another comparison of land
462 use on denitrifier communities, edaphic factors, such as pH and soil organic C, were identified as
463 primary drivers of differences in denitrification rates (Krause et al., 2017). While these land uses at
464 KBS have previously been shown to have differences in C content (Grandy and Robertson, 2007),
465 these may be too slight to affect the C preference of denitrifiers. Similar studies on a broader range of
466 soils and land uses will be required before drawing general conclusions as to the universality of
467 denitrifier C preference. It is also possible, and likely, that differentiating C preferences between
468 denitrifier communities manifest in the utilization of C compounds that were not included in this
469 study.

470 Microbial community effects were strongest in the qualitative differences in N₂O reduction
471 between the two sites. At KBS, amino acids and proteins resulted in significantly less net N₂O
472 production compared to the other substrates. On the other hand, at MRC amino acids did not result in
473 any less net N₂O production, while proteins stimulated almost complete removal of N₂O from the
474 microcosms. Microbial communities have widely different capacities for N₂O reduction depending
475 on the abundance and type of N₂O reducers present. In particular, communities that have more

476 organisms with *nosZ-II* may have a greater ability to consume N₂O than those with *nosZ-I* (Graf et
477 al., 2014; Jones et al., 2014; Yoon et al., 2016). Therefore, the addition of substrates stimulating
478 *nosZ-II* organisms could result in lower net production of N₂O. At least one study has shown that
479 *nosZ-II* organisms are more responsive to changes in C availability than those with *nosZ-I* (Assémien
480 et al., 2019); however, other studies have shown that both clades of *nosZ* are equally responsive to C
481 availability (Domeignoz-Horta et al., 2015, 2018; Juhanson et al., 2017). How these two groups of
482 organisms respond to different forms of C deserves further study.

483 **5. Conclusion**

484 In this study, we demonstrate how the chemical identity of C inputs influences N₂O
485 production and reduction in different agricultural soils. To resolve apparently divergent patterns in
486 denitrifier C utilization between different studies, we compared a set of twelve C-addition treatments
487 over five soils under identical assay conditions. We found that amino acids, proteins, and organic
488 acids consistently stimulated the most denitrification and N₂O reduction among the substrates
489 examined here. While soils from distinct land uses had differing overall rates of denitrification, C-
490 utilization profiles were largely similar between soils, suggesting denitrifier C preferences may be
491 widely held between microbial communities, at least within agricultural soils. The bioavailability of
492 C substrates appears to be a significant driver in denitrification and N₂O reduction, with labile
493 substrates stimulating greater activity than polymeric and recalcitrant C additions. Nevertheless, this
494 pattern was not universal across all the substrates tested. Given the heterogeneous distribution of
495 different forms of C throughout the soil profile, substrate effects likely contribute to the spatial and
496 temporal variability of N₂O production within soils. Determining whether these denitrifier C
497 preferences also occur under field conditions may provide opportunities to reduce emissions of N₂O
498 from soils by controlling soil C inputs.

499 **Acknowledgements**

500 This work was part of the MMPRNT project, which is funded by the U.S. Department of
501 Energy, Office of Science, Office of Biological and Environmental Research [award number
502 DESC0014108]. Additional funding was also provided by USDA-NIFA [award number 2017-67013-
503 26257] and the Foundation for Food and Agriculture Research [award number 430834]. Support for
504 this research was also provided by the NSF Long-term Ecological Research Program (DEB 2224712)
505 at the Kellogg Biological Station and by Michigan State University AgBioResearch.

506

507 **Tables**

508 **Table 1: Edaphic factors of study sites**

	Total C (%) ^c	pH	CEC (cmol kg ⁻¹)
Kellogg Biological Station ^a			
Conventional Row Crop	0.91 ± 0.08	6.12 ± 0.03	8.13 ± 0.23
Reduced Input Row Crop	1.09 ± 0.05	6.28 ± 0.01	7.95 ± 0.11
Perennial Switchgrass	1.42 ± 0.06	6.38 ± 0.02	7.95 ± 0.15
Montcalm Research Center ^b			
Maize-Potato with No Cover	0.87 ± 0.03	6.5	10
Maize-Potato with Vetch and Rye	1.15 ± 0.19		

509
 510 Total soil carbon, pH, and cation exchange capacity of the soil utilized in the study. ^a Soil data for
 511 Kellogg Biological Station was obtained from Robertson and Simmons (2020). ^b Cation exchange
 512 capacity and pH for soils sampled from Montcalm Research Center were obtained from the Web Soil
 513 Survey (USDA NRCS). ^c Soil carbon data for KBS was obtained from Grandy and Robertson (2007).
 514
 515

Table 2: Characteristics of added carbon substrates

Substrate	Formula	C:N Ratio	NOSC	Utilization	Degradation
Glucose	$C_6H_{12}O_6$	No N	0	Short-term energy storage. Monomer of cellulose	Widely used substrate in cellular respiration and fermentation reactions
Cellulose	$(C_6H_{10}O_5)_n$	No N	0	Polymer of glucose. Structural component of plant cell walls	Requires specialized enzymes to depolymerize extracellularly
N-acetylglucosamine (NAG)	$C_8H_{15}NO_6$	8:1	0	Monomeric subunit composing the cell walls of bacteria (peptidoglycan) and fungi (chitin)	Initial degradation steps require specialized enzymes to remove the acetyl and amino groups
Chitin	$(C_8H_{13}O_5N)_n$	8:1	0	Polymer of NAG. Structural component of fungal cell walls and insect exoskeletons	Requires specialized enzymes to depolymerize extracellularly
Amino acids		~4:1 (3:2 to 9:1)	-0.1	Monomeric subunit of proteins	Can be recycled into new proteins or catabolized via cellular respiration and fermentation depending on the specific microbe and amino acid
Protein		~4:1	-0.1	Polymer of amino acids. Multiple uses in cellular functioning and structure	Requires specialized enzymes to depolymerize extracellularly
Vanillyl alcohol	$C_8H_{10}O_3$	No N	-0.5	Monomeric subunit of lignin	Aromatic ring structure requires specialized pathways to break down. Often requires an oxidase
Lignin		No N	-0.5	Highly stable aromatic polymer, structural component of plant cell walls	Requires specialized enzymes to depolymerize extracellularly
Citrate	$C_6H_8O_7$	No N	+1	Intermediate of central metabolism	Intermediate in central metabolism for nearly all microbes, but use as a C source is restricted by the ability to transport into the cell. Not a typical fermentation substrate but there are some specialized pathways
Succinate	$C_4H_6O_4$	No N	+0.5	Intermediate of central metabolism	Intermediate in central metabolism for nearly all microbes, but use as a C source is restricted by the ability to transport into the cell. Not a typical fermentation substrate but there are some specialized pathways

Methanol

CH₃OH

No N

-2

Waste product

Energy dense but requires specialized pathways to be utilized

Characteristics of added C substrates, including the C:N molar ratio, the nominal oxidation state of C (NOSC), how the compounds are commonly utilized by microorganisms, and how they can be degraded. NOSC is related to the amount of energy that is attainable from a C substrate. More energy-dense substrates will be more reduced and have a more negative NOSC, while more oxidized compounds will have a more positive NOSC.

Table 3: Type III ANOVA table of fixed effects

	Gross N ₂ O		Net N ₂ O		dN ₂ O	
	F	P	F	P	F	P
Kellogg Biological Station						
Substrate	93.9	< 0.001	5.5	< 0.001	33.4	< 0.001
Land Use	89.1	< 0.001	3.5	0.100	0.7	0.546
Substrate × Land Use	2.0	0.013	3.5	< 0.001	1.4	0.140
Montcalm Research Center						
Substrate	207.0	< 0.001	85.3	< 0.001	70.7	< 0.001
Land Use	6.0	0.092	12.1	0.040	0.5	0.520
Substrate × Land Use	1.9	0.081	1.9	0.074	2.1	0.048

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Figure 1: Differences in gross N₂O production between substrate additions at KBS (A) and MRC (B)

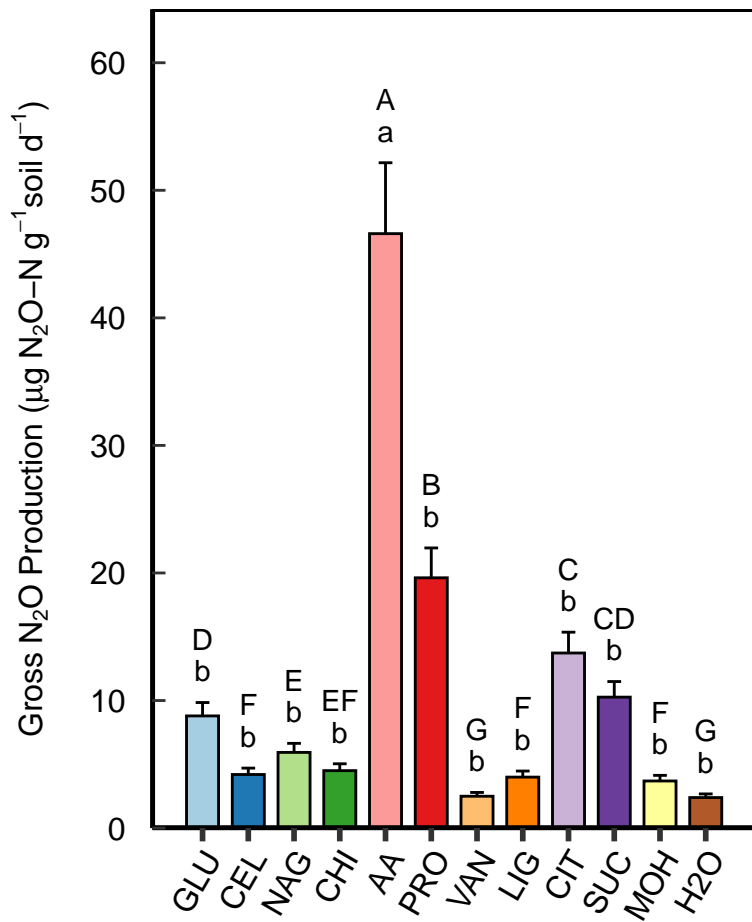
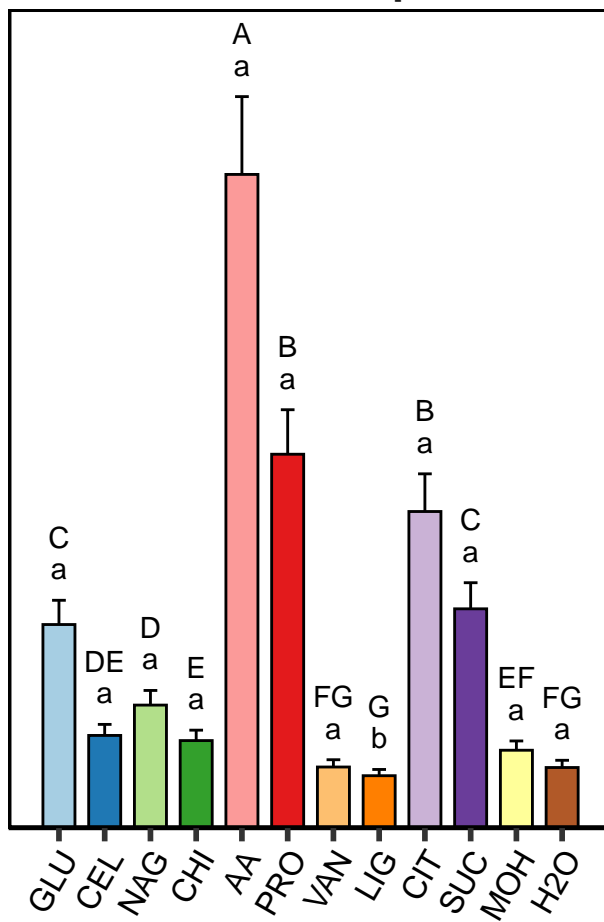
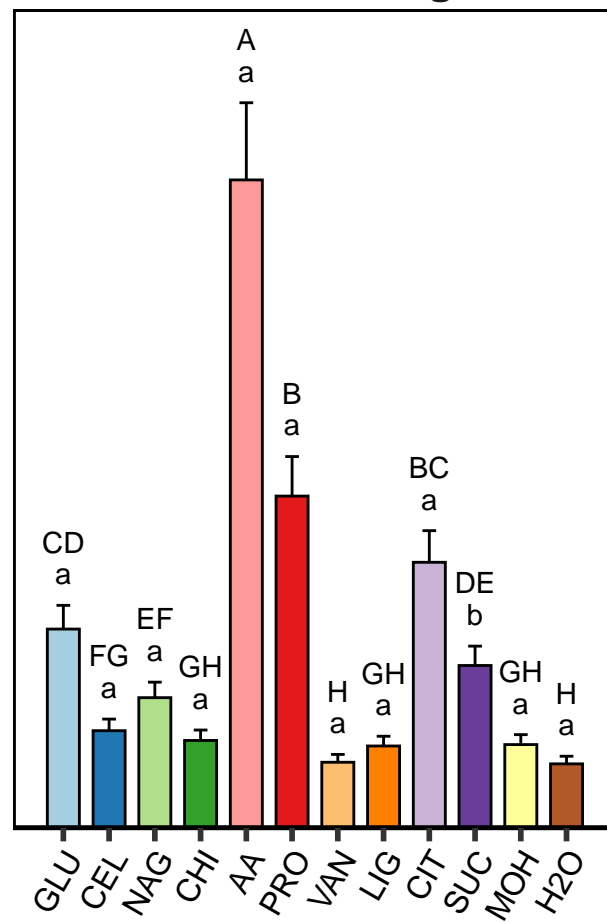
Cumulative production of N₂O in treatments receiving acetylene. Acetylene inhibits N₂O reduction; thus, N₂O concentrations represent total gross production of N₂O following the addition of glucose (GLU), cellulose (CEL), N-acetyl-glucosamine (NAG), chitin (CHI), casamino acids (AA), soy protein isolate (PRO), vanillyl alcohol (VAN), lignin (LIG), citrate (CIT), succinate (SUC), methanol (MOH), or no C addition (H₂O). Means are shown with error bars representing one standard error (n = 4). Capital letters indicate significantly different means between substrates within each land use; lowercase letters indicate significantly different means between land uses for each substrate ($\alpha = 0.05$).

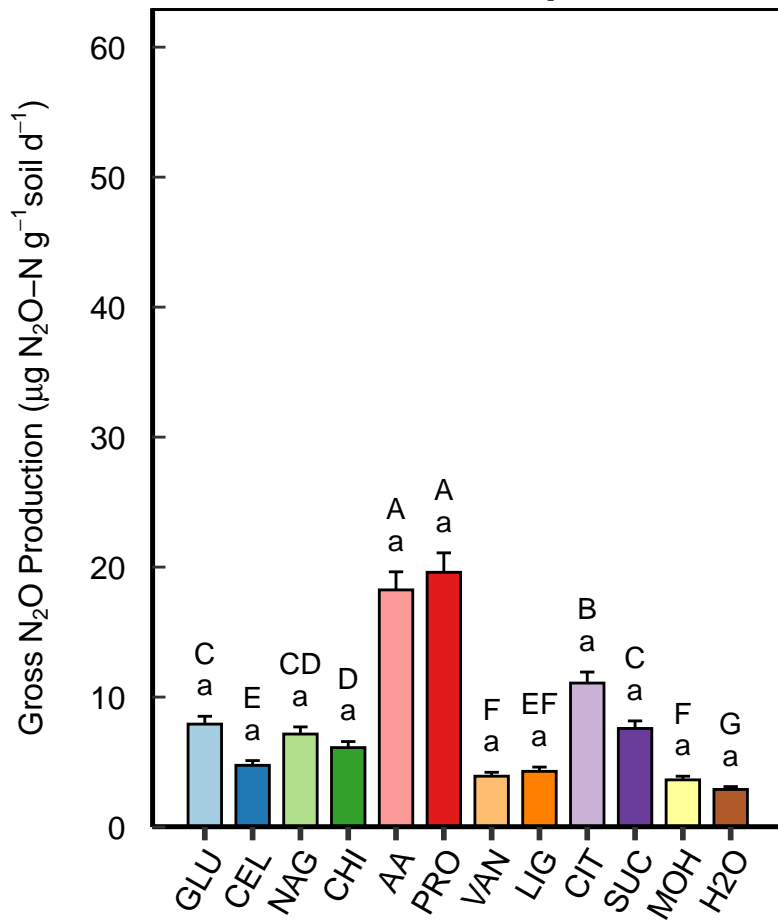
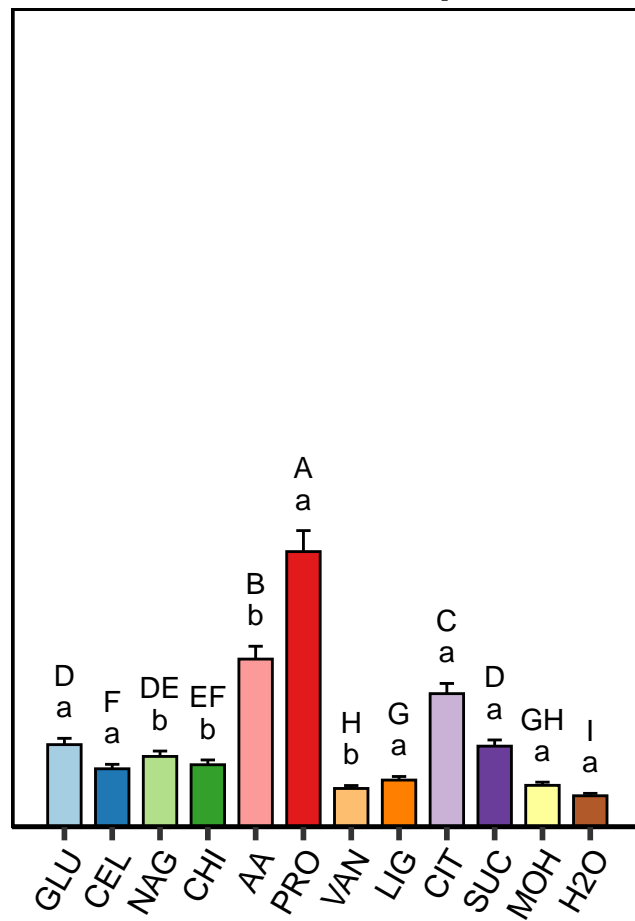
Figure 2: Differences in net N₂O production between substrate additions at KBS (A) and MRC (B)

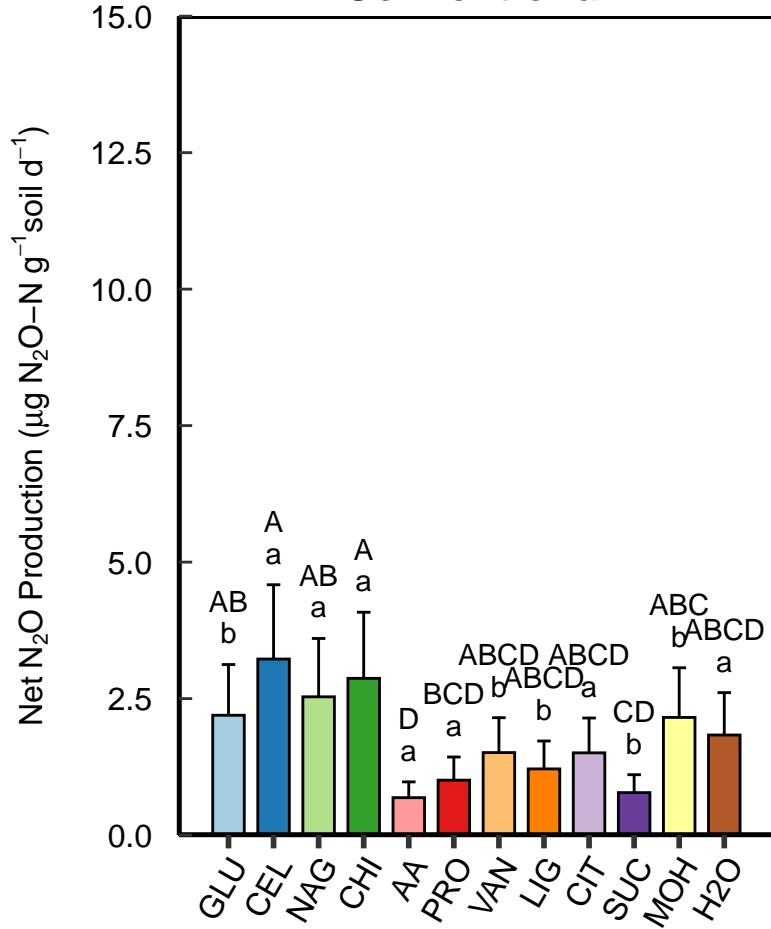
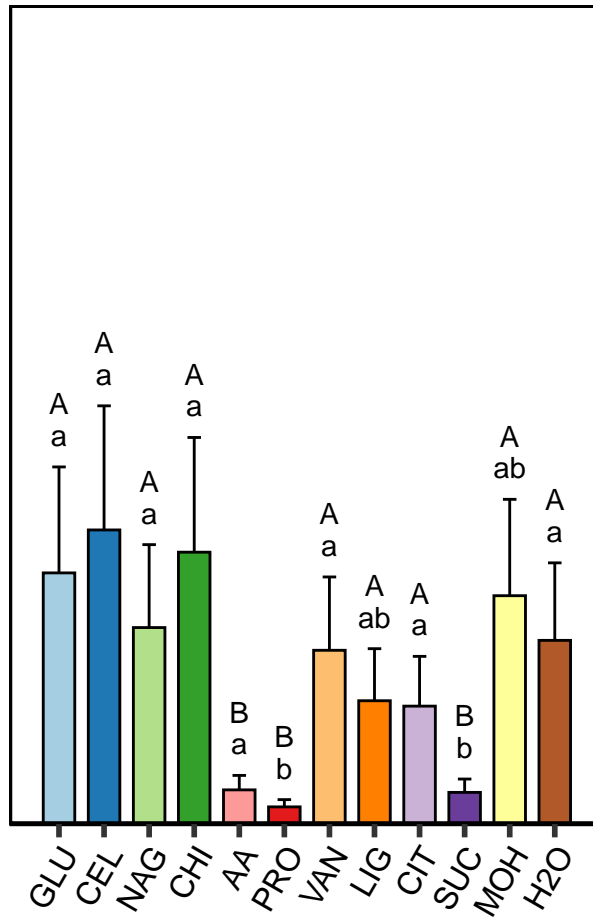
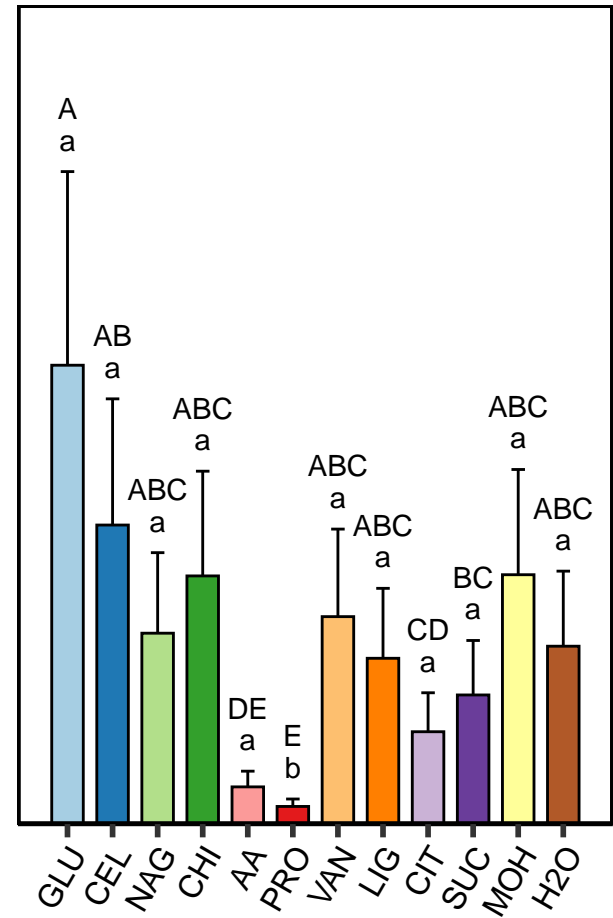
Cumulative production of N₂O in treatments that did not receive acetylene. N₂O values reflect the balance of N₂O production and N₂O reduction following the addition of glucose (GLU), cellulose (CEL), N-acetyl-glucosamine (NAG), chitin (CHI), casamino acids (AA), soy protein isolate (PRO), vanillyl alcohol (VAN), lignin (LIG), citrate (CIT), succinate (SUC), methanol (MOH), or no C addition (H₂O). Means are shown with error bars representing one standard error (n = 4). Capital letters indicate significantly different means between substrates within each land use; lowercase letters indicate significantly different means between land uses for each substrate ($\alpha = 0.05$).

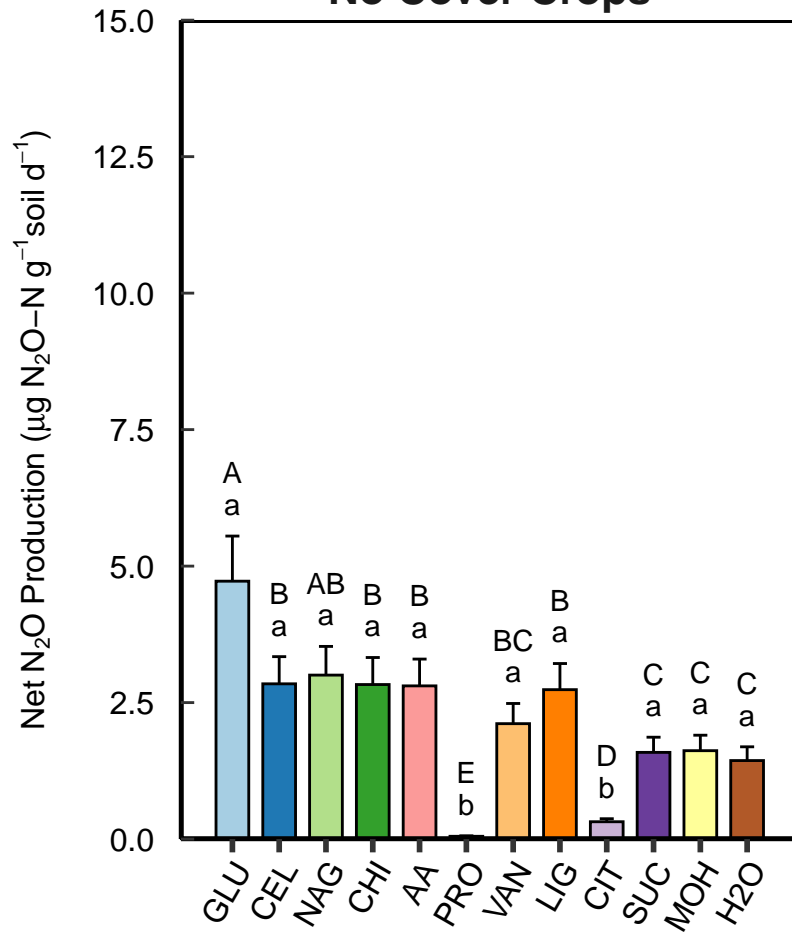
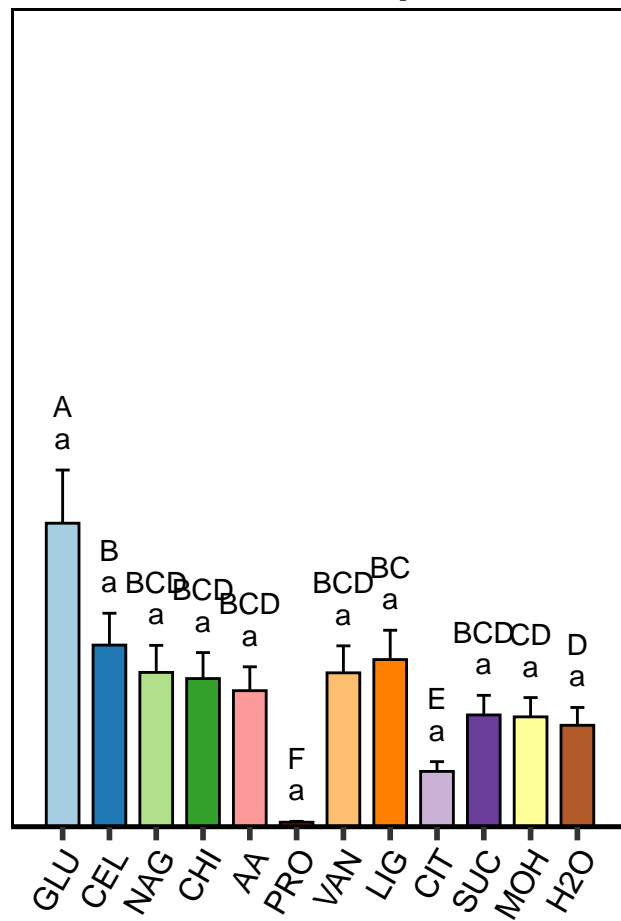
Figure 3: Portion of total N₂O production remaining as N₂O between substrate additions at KBS (A) and MRC (B)

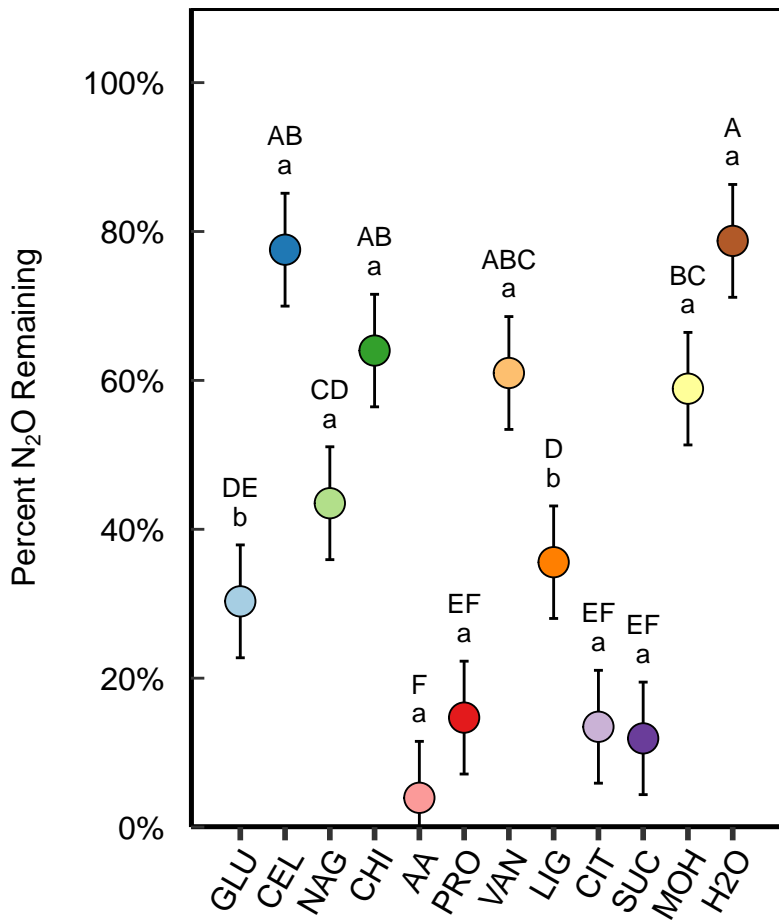
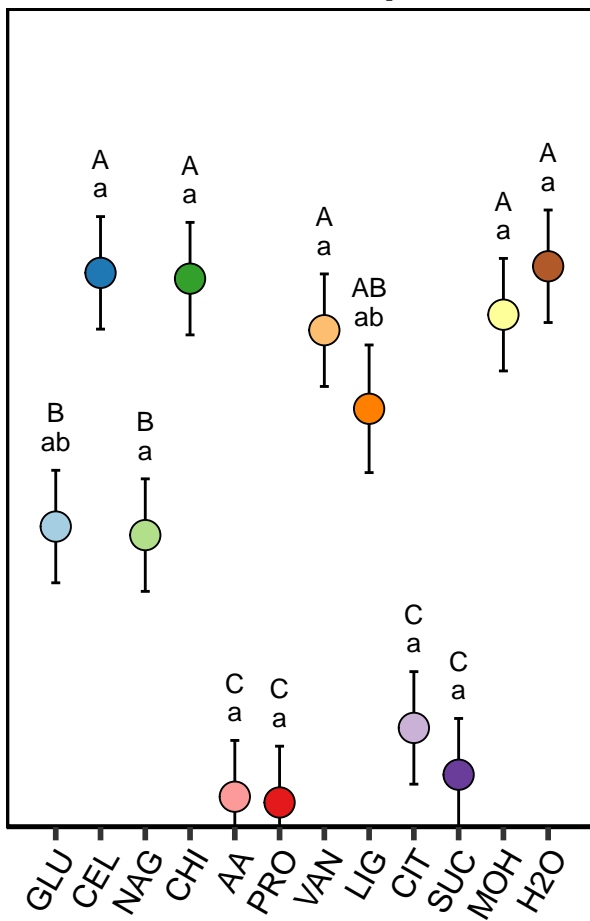
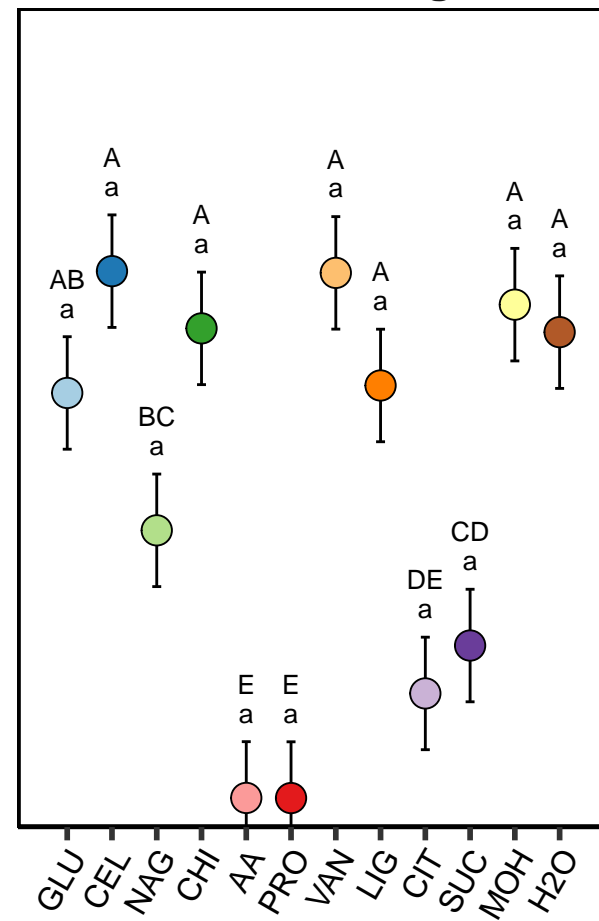
The portion of N₂O remaining (dN₂O) scales net production of N₂O to total gross production of N₂O. Higher values indicate less N₂O reduction occurred relative to total denitrification, while values close to zero indicate near complete reduction of all N₂O produced. Values are means of treatments receiving glucose (GLU), cellulose (CEL), N-acetyl-glucosamine (NAG), chitin (CHI), casamino acids (AA), soy protein isolate (PRO), vanillyl alcohol (VAN), lignin (LIG), citrate (CIT), succinate (SUC), methanol (MOH), or no C addition (H₂O). Error bars represent one standard error (n = 4). Capital letters indicate significantly different means between substrates within each land use; lowercase letters indicate significantly different means between land uses at each location for each substrate ($\alpha = 0.05$).

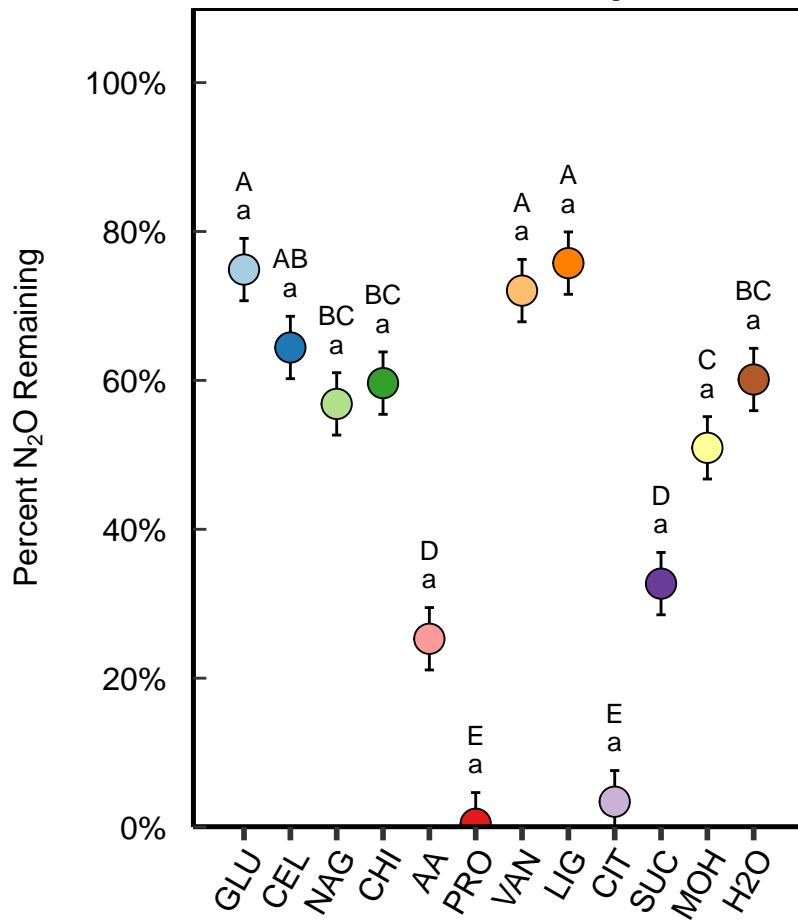
A**Conventional****Reduced Input****Perennial Switchgrass**

B**Cover Crops****No Cover Crops**

A**Conventional****Reduced Input****Perennial Switchgrass**

B**No Cover Crops****Cover Crops**

A**Conventional****Reduced Input****Perennial Switchgrass**

B**No Cover Crops****Cover Crops**