

# Bison and cattle grazing increase soil nitrogen cycling in a tallgrass prairie ecosystem

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Abstract Nitrogen (N) is a necessary element of soil fertility and a limiting nutrient in tallgrass prairie but grazers like bison and cattle can also recycle N. Bison and cattle impact the nitrogen (N) cycle by digesting forage that is consumed, and recycled back to the soil in a more available forms stimulating soil microbial N cycling activities. Yet we do not know how both grazers comparatively affect N cycling in tallgrass prairie. Thus, we investigated if bison and cattle had similar impacts on N cycling in annually burned tallgrass prairie relative to ungrazed conditions over a 3-year period (2020-2022) at the Konza Prairie Biological Station. We examined: soil pH, soil water content, mineralized N, nitrification potential, denitrification potential and extracellular enzyme assays. Interannual variability in precipitation controlled soil water and N cycling microbial activities but grazing effects had a stronger influence on N cycling. We found significant differences and increased soil pH, nitrification and denitrification potential and less N limitation in bison vs cattle grazed soils where bison grazed soils

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University of Kansas, Kansas Biological Survey & Center for Ecological Research, Lawrence, KS, USA exhibited faster N cycling. Differences between the grazers may be attributed to the different management of bison and cattle as both can impact N cycling. Overall, these data provide some evidence that bison and cattle affect N cycling differently at this study site, and improve the ecological understanding of grazer impacts on N cycling dynamics within the tall-grass prairie ecosystem.

**Keywords** Nitrogen · Tallgrass prairie · Bison · Cattle · Soil

# Introduction

Nitrogen (N) is a limiting nutrient in many terrestrial ecosystems, including tallgrass prairies (Blair 1997; Schlesinger and Bernhardt 2020). In this ecosystem, frequent fire volatilizes N from plant litter, slowing the accumulation of soil organic N; therefore, fire maintains conditions in which N-limited plants and soil microbes rapidly assimilate and immobilize, and effectively retain, soil available inorganic N (Dodds et al. 1996; Dell and Rice 2005; Dell et al. 2005). In addition, tallgrass prairies were historically grazed by large mammalian herbivores, which often enhance soil N cycling rates, N heterogeneity, and soil fertility (Hobbs 1996; Frank and Evans 1997; Blair et al. 1998; Knapp et al. 1999; Bakker et al. 2003). Bison grazing can increase soil N cycling rates in areas managed with annual fire to levels equivalent to areas

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experiencing infrequent fire (Groffman et al. 1993; Johnson and Matchett 2001).

Plains bison, also known as the American Buffalo (Bison bison), are keystone herbivores that once ranged across the whole North American continent (Knapp et al. 1999; Lott 2002; Anderson 2006; List et al. 2007). However, in the 1880s, extermination through settler colonization (Dunbar-Ortiz 2014) decimated bison populations which numbered in the millions to less than 1000 individuals, driving societal collapse and negative economic impacts on Native American communities that remain until this day (Hornaday, 1913; Flores 1991; Shaw 1995; Lott 2002; Feir et al. 2021). Domesticated cattle (Bos taurus) now outnumber bison by an estimated 950 million individuals (Samson et al. 2004; Kohl et al. 2013). In the 1980s and 90 s, ecological researchers recognized bison as major ecosystem drivers in maintaining Great Plains grasslands and their nutrient cycles (Hulbert 1986; Vinton et al. 1993; Ojima et al. 1994; Coppedge et al. 1998a, b; Woodmansee and Duncan 1980; Risser and Parton 1982; Blair 1997). All things considered, replacement of the keystone bison with cattle raises concerns of whether both animals occupy the same functional roles on the landscape (Allred et al. 2011; Kohl et al. 2013), mirroring global concerns and studies on the alteration of nutrient cycling following the replacement of native megaherbivores with domesticated cattle in managed rangeland (Enquist et al. 2020; Abraham et al. 2023; Roy et al. 2023).

Cattle and bison have certain redundant roles in ecosystem N-cycling function, but also differ physiologically in potentially influential ways. They both graze similar grasses in tallgrass prairie (Allred et al. 2011) and excrete dung and urine which in turn increases bioavailable N for soil microorganisms and plants (McNaughton 1983; Detling 1988; Schlesinger and Hartley 1992; Anderson 2006). However, bison are hardier and tolerant of extreme hot and cold weather temperatures, enabling them to travel and spend more time grazing away from streams on upland prairie (Christopherson et al. 1978; Allred et al. 2011; Larson et al. 2013; McMillan et al. 2021; McMillan et al. 2022), while cattle are less weather hardy and tend to travel infrequently by comparison choosing to spend more time near riparian areas (Kohl et al. 2013; McMillan et al. 2021). Therefore, the distribution of N across the landscape by bison and cattle depends on decisions to travel and forage and drink water (Plumb and Dodd 1993; Augustine and Frank 2001; Raynor et al. 2021). Many comprehensive studies in tallgrass prairie focus on aboveground plant responses to grazing by bison and cattle, with less emphasis on soil microbial functions that allow N to become available for forage regrowth (Plumb and Dodd 1993; Coppedge and Shaw 1997; Coppedge et al. 1998a, b; Towne et al. 2005; McMillan et al. 2011; McMillan et al. 2019; Ratajczak et al. 2022). To our knowledge, no studies have directly assessed whether bison and cattle similarly influence N cycling in tallgrass prairie soils.

Therefore, we investigated soil microbial N cycling activities in annually burned tallgrass prairie, in bison grazed, cattle grazed, and ungrazed areas, focusing on microbially mediated N cycling transformations. We predicted that all soil N cycling rates would be higher, and that soil microbial N limitation would be lower, in grazed relative to ungrazed treatments, and that bison and cattle would have similar magnitudes of influence on soil N cycling. In addition, we considered that the influence of grazing on soil N cycling rates may vary due to differences in soil water availability stemming from precipitation variability, which is a primary control over ecosystem and N cycling dynamics in tallgrass prairie and other grasslands around the world, with higher water generally promoting more plant production and faster N cycling (Groffman et al. 1993; Broderick et al. 2022; Chen et al. 2022). To assess the predictions, we sampled upland soils in annually burned watershed-scale experimental grazing treatments at the Konza Prairie Biological Station (KPBS), each summer from 2020 through 2022, and measured resin-bound inorganic N (a proxy for the amount of mineralized N available for plant and soil microbial uptake through a growing season), nitrification potential rates, denitrification potential rates, denitrification enzyme activity rates, and hydrolytic extracellular enzyme activity rates (which were also used to calculate an index of soil microbial N limitation).

## Methods

Study site and sampling design

The KPBS is a 3487-ha tallgrass prairie preserve located in the Flint Hills region of northeastern

Kansas near Manhattan, KS, USA (39° 05' N, 96° 35' W). KPBS is situated on one of the last remaining tracts of tallgrass prairie, was established as a research station in 1971, and became host to a Long-Term Ecological Research (LTER) project in 1980. KPBS maintains watershed scale treatments of differing fire intervals. Bison were reintroduced to a subset of these experimental watersheds between 1987 and 1992, and cattle were introduced to another subset of watersheds in the 1990s. Bison are stocked at 0.4 ha per animal unit month (AUM: the forage required to feed a 454 kg animal or its equivalent for 1 month), or 0.98 acres per animal, reproduce on-site, and are present year-round (Blair 2023); while cattle graze annually as cow-calf pairs between May 1 to October 1 and are stocked at 0.7 ha per AUM or 1.7 acres per cow-calf pair (Olson 2023), but for a shorter period of the year (April through October). Dominant plants of this area include Sorghastrum nutans, Andropogon gerardii, Schizacryum scoparium, Panicum virgatum, Amorpha canescens and Rhus glabra. Mean annual precipitation (MAP) at this site is 899 mm and mean annual temperature (MAT) is 12.5 °C.

For this study, research was restricted to upland soils, to control for variability in soil type. Samples were collected on the Florence-Benfield complex soil map unit (Clayey-skeletal, smectitic, mesic Udic Argiustolls and Fine, mixed, superactive, mesic Udertic Argiustolls), which is widespread across the Flint Hills of Kansas. Soil sampling was undertaken once late in each summer growing season from 2020 to 2022. These years spanned a range of above-average rainfall (2020) to well below average (2021) and slightly below average (2022) (Fig. 1). We sampled along four 10-m transects, parallel to long-term plant sampling transects in each experimental watershed, in two bison grazed (N1A and N1B), two cattle grazed (C1A and C1B), and two ungrazed (1D and SpB) watersheds, all of which are burned annually. We acknowledge that landscape-scale replication of grazing treatment is low in this study; however, the field experimental design provides good standardization of soil type, which often affects baseline levels of microbial N-cycling (Zeglin et al. 2007).

Each transect covered six sampling points at 0, 0.1, 0.5, 1, 5, and 10 m, from which 2-cm diameter mineral soil samples were collected with an Oak-field corer (Oakfield, WI, USA), to a depth of up to 15 cm. Sample locations at 0, 1, 5, and 10 m were



**Fig. 1** Cumulative precipitation on KPBS in the 2020, 2021, and 2022 growing seasons. Historic (30-year) average precipitation is denoted by a blue line overlaying the gray one standard deviation boundaries. Accumulated precipitation is denoted by the red line for each respective year, and dashed vertical lines indicate soil sampling dates for the year

geolocated using the WGS84 datum with a Garmin GPSMAP 64x (Garmin, Olathe, KS, USA). Samples were taken using sterile technique, i.e., while wearing nitrile gloves, and by washing the corer in ethanol between each sample. Samples were stored in a cooler on ice and transported to the lab, where all samples were aseptically sieved using a No. 4 (4 mm) sieve, to remove rocks and plant roots while largely retaining soil aggregate structure. A portion of each soil sample was frozen and stored at -20 °C before soil physical analysis, and the remaining fresh soil was stored at 4 °C for no more than 48 h before measuring N cycling activity potential rates.

Soil physical characteristics and N-cycling rates

Soil water content was measured gravimetrically by drying soil at 105 °C for 24 h. Soil pH was measured from a 1:3 slurry of field-moist soil and DI H<sub>2</sub>O, on samples collected in 2021 and 2022. Soil available N was measured using ion exchange resin bags installed from June to September (Baer and Blair 2008; Nieland et al. 2021) in 2021 and 2022. Resin bag sorbed  $NH_4^+$ –N and  $NO_3^-$ –N was quantified using a modified indophenol method and VCl<sub>3</sub>/Griess reagent method (Hood-Nowotny et al. 2010), respectively, and measured spectrophotometrically with a Filtermax F5 Multimode Microplate Reader (Molecular Devices, San Jose, CA, USA).

We measured rate potentials of two microbiallymediated N transformation processes, nitrification

and denitrification, and we measured denitrification potential in two ways, to learn about different limiting factors on soil N cycling. Nitrification is the process of oxidation of ammonium to nitrate, and is an important consideration in soil fertility since fewer types of plant can readily assimilate both ammonium and nitrate, while nitrate is more easily lost from the soil due to leaching or denitrification. Denitrification is the process of reduction of nitrate to dinitrogen gas or nitrous oxide, atmospheric gases that are not useful to plants. Nitrification and denitrification processes bring energy to different types of specialist microorganisms, and are thus limited by the abundance of those organisms, as well as the availability of ammonium and oxygen (for nitrification), and the availability of nitrate, dissolved organic carbon, and anoxic conditions (for denitrification). We assayed both processes in lab incubations with saturating levels of substrates and optimal levels of oxygen, effectively measuring an index of maximum microbial potential for nitrification and denitrification, and also measured denitrification rates in assays with no substrate added to understand process rates under ambient soil conditions.

Nitrification potential (NP) rates were measured in an aerobic soil slurry amended with saturating concentrations (250  $\mu$ M) of NH<sub>4</sub><sup>+</sup>-N, and shaken at 120 rpm. After 0.25 and 24 h, 1 ml of each soil slurry sample was transferred into a 1.5 ml tube, centrifuged at 15,000 rpm, and the supernatant was frozen at -20° C until measuring NO<sub>3</sub><sup>-</sup>-N as described above (Taylor et al. 2010). The resulting increase in NO<sub>3</sub><sup>-</sup>-N over time, due to ammonia oxidation by soil microorganisms, was used to calculate the maximum nitrification potential rate of each soil sample.

Denitrification potential activity (DNP) and denitrification enzyme activity (DEA) were measured in parallel (Groffman et al. 2009; Nieland et al. 2021). Both are estimates of the reduction of  $NO_3^-$ –N to N<sub>2</sub>O–N in an aerobic soil slurry in the presence of acetylene, which prevents the transformation of N<sub>2</sub>O–N to N<sub>2</sub>–N. DEA is defined as the maximum enzymatic potential at which denitrification can occur over 1 h, measured with the addition of both glucose and KNO<sub>3</sub>, which provides the optimal resources necessary for bacterial denitrification. In contrast, DNP assays were not amended, reflecting denitrification rates attainable under levels of nitrate and carbon availability in the soil sample, and measured over a 4-h period. The production of  $N_2O-N$  used to calculate DNP and DEA was measured using a Shimadzu 2014 GC analyzer (Shimadzu Scientific Instruments, Inc., Columbia, MD, USA).

Soil extracellular enzyme activity (EEA) and microbial N limitation

The enzymatic hydrolysis of amino acids and amino sugars from soil organic matter controls soil fertility by limiting the rate of net N mineralization from biologically inaccessible soil N into forms that can be assimilated by plants and microorganisms (Schimel and Weintraub 2003; Sinsabaugh et al. 2009). The expression of these enzymes is generally regulated by product suppression, i.e., if soil N availability is higher, then N demand is lower, and fewer enzymes to produce available N are synthesized relative to synthesis of enzymes catalyzing hydrolysis of bioavailable C or P (Allison and Vitousek 2005; Nieland et al. 2024).

For this study, we measured the activity potential of two common N acquiring enzymes, β-Nacetylglucosaminidase (NAG; EC 3.2.1.14, 4-MUB-N-acetyl-β-D-glucosaminide) leucyl and aminopeptidase (LAP; EC 3.4.11.1, L-leucine-7-amido-4-MC), as well as one carbon acquiring enzyme, β-glucosidase (βG; EC 3.2.1.21, 4-MUB-β-D-glucoside). Hydrolytic enzyme activity rates were measured using fluorometric substrates (methylumbelliferone (MUB) for NAG and BG, and methylcoumarin (MC) for LAP). Soil samples were thawed and 1 g of each soil sample was added to a solution of 100 ml 50 mM sodium acetate buffer (pH 5), forming a slurry. We combined 200 µl of soil slurry and 50 µl of the target substrate in 96 well assay plates, with six analytical replicates and triplicate quench standards per sample and replicate blanks, negative controls, and 200 µM reference standards. Assays for NAG were incubated for 3.5 h, LAP for 16 h, and βG for 2 h. After incubations, reactions were halted with the addition of 10 µl of 0.5 M NaOH, raising the pH to > 8. Fluorescence of hydrolyzed substrate was measured at excitation/emission of 360/450 nm with a Filtermax F5 Multimode Microplate Reader (Molecular Devices, San Jose, CA, USA). Finally, we calculated indices of soil microbial N limitation:  $(\ln(\beta G))/$  $(\ln(NAG + LAP))$ , which decreases under conditions of higher N- than C-limitation, and ((ln(NAG+LAP)/ ln(Phos)), which increases under conditions of higher N- than P-limitation (Sinsabaugh and Shah 2012).

# Data analysis

All statistical analysis was done using the R programing language in the R studio interface for statistical analysis (R Core Team 2022). While sampling was performed using a log-distance design to assess spatial heterogeneity patterns, this structure was surprisingly weak, so we proceeded with a standard statistical approach. To test for the direct and interactive effects of grazing treatment and year on soil characteristics and N cycling rates, we used two-way analysis of variance (ANOVA) models, with post-hoc Tukey's honest significant difference (HSD) tests for pairwise comparisons of within-group differences. Coefficient of correlation  $(R^2)$  was used to assess linear relationship strength between soil characteristics, N cycling rates, and EEAs. For each variable, diagnostic Q-Q plots and histograms were used to assess assumptions of statistical normality; if these assumptions were not met, a square root or natural log transformation was used to shift the data distribution to better satisfy normality assumptions. Statistical results with a P-value of < 0.05 are reported in the text.

# Results

Soil water content, pH, and available N

Soil gravimetric water content and pH varied with both grazing treatment and year, independently of one another (Table 1). Mean soil water content at the time of sampling was highest in 2020 and lowest in 2022, and was also higher in ungrazed soils than in bison or cattle grazed soils (Fig. 2). Soil pH was measured higher in 2022 than 2021, and was also consistently higher in bison grazed soils than in cattle grazed or ungrazed soils (Table 1, Fig. 2). Resin-sorbed N responses to grazing and year were also independent of one another (Table 1): In 2022, resin-sorbed NH4+-N, NO3-N, total inorganic N, and  $NO_3^--N:NH_4^+-N$  were all higher than in 2021 (Fig. 3); also, bison grazed treatments had higher resin-sorbed nitrate and total inorganic N than ungrazed and cattle treatments, but resin-sorbed  $NH_4^+$ -N and the ratio of  $NO_3^-$ -N:NH<sub>4</sub><sup>+</sup>-N did not respond to grazing (Fig. 3).

Nitrification and denitrification potentials

Soil nitrification potential (NP) responded to grazing and year independently, such that NP was lower in 2022 than in 2020 or 2021, and was also

Table 1Two-way ANOVAresults (F statistic and Pvalues) showing the directand interactive effectsof grazing and samplingyear on soil GWC andpH, resin-bound N,N-cycling potential rates,and extracellular enzymeactivities	Response variable	Graze F, P	Year F, P	Graze * Year F, P	
	GWC <sup>Sqrt</sup>	16.0, < 0.001	83.1, < 0.001	1.17, 0.32	
	pH	22.3, < 0.001	6.52, 0.012	2.36, 0.10	
	$NH_4^+ - N^{\ln}$	1.32, 0.27	262.2, < 0.001	1.18, 0.31	
	NO <sub>3</sub> <sup>-</sup> -N <sup>ln</sup>	12.8, < 0.001	24.9, < 0.001	2.37, 0.10	
	$NO_{3}^{-}-N+NH_{4}^{+}-N^{ln}$	13.1, < 0.001	34.3, < 0.001	2.16, 0.12	
	$NO_{3}^{-}-N:NH_{4}^{+}-N^{ln}$	2.49, 0.086	84.2, < 0.001	0.37, 0.69	
	NP <sup>In</sup>	20.4, < 0.001	6.36, 0.002	0.37, 0.83	
	$DNP^{ln}$	2.29, 0.11	36.8, < 0.001	1.35, 0.25	
	DEA <sup>sqrt</sup>	13.7, < 0.001	28.6, < 0.001	1.35, < 0.001	
	$\mathrm{BG}^{\mathrm{ln}}$	2.20, 0.11	62.5, < 0.001	0.67, 0.62	
	CBH <sup>ln</sup>	3.03, 0.050	47.6, < 0.001	1.72, 0.15	
Bolded values indicate	NAG <sup>ln</sup>	5.70, 0.004	60.1, < 0.001	2.72, 0.030	
P<0.05	LAP <sup>In</sup>	14.5, < 0.001	138.2, < 0.001	9.41, < 0.001	
Sqrt denotes square root transformation and In denotes natural log transformation	Phos <sup>ln</sup>	2.21, 0.11	47.4, < 0.001	2.37, 0.053	
	lnBG:ln(NAG+LAP)	11.0, < 0.001	4.03, 0.019	2.93, 0.021	
	ln(NAG + LAP):ln(Phos)	6.90, 0.001	1.89, 0.15	3.23, 0.013	





Fig. 2 A Soil gravimetric water content over a 3-year period in grazed and ungrazed soils and **B** soil pH over a 2-year period. Tukey's HSD post-hoc results are shown with different letters indicating years (top) or grazing treatments (x-axis) that differed from each other at P < 0.05

consistently higher in bison grazed soils than cattle grazed or ungrazed soils (Table 1; Fig. 4). Soil denitrification potential (DNP) did not respond to grazing treatment, but was higher in 2020 than in 2021 or 2022 (Table 1; Fig. 4). Notably, DNP rates were only 14% (on average) of the denitrification enzyme activity (DEA) when detectable, and were below detectable limits in 2022 and in cattle and ungrazed treatments in 2021. DEA responses to grazing varied interannually (Table 1). In 2020, soil DEA in bison grazed treatments was greater than in both cattle grazed and ungrazed treatments; in 2021, DEA was higher in cattle grazed than in ungrazed soils, and intermediate in bison grazed soils; and in 2022, DEA was higher in bison grazed than in ungrazed soils, and intermediate in cattle grazed soils (Fig. 4).



**Fig. 3** A NH<sub>4</sub><sup>+</sup>–N, **B** NO<sub>3</sub><sup>-</sup>–N, **C** NO<sub>3</sub><sup>-</sup>–N+NH<sub>4</sub><sup>+</sup>–N, and **D** NO<sub>3</sub><sup>-</sup>–N: NH<sub>4</sub><sup>+</sup>–N sorbed to resin bags through the summer growing season in two study years. Tukey's HSD post-hoc results are shown with different letters indicating years (top) or grazing treatments (center) that differed from each other at P < 0.05

Soil extracellular enzyme activities and relative N limitation

Extracellular enzyme activity potentials varied interannually, tending to be highest in 2022 and lowest in 2020 (Table 1; Fig. 6). Only N-acquiring enzyme activities (NAG and LAP) and the indices of N demand relative to C demand (lnBG:ln(NAG+LAP)) and relative to P demand  $(\ln(NAG + LAP):\ln Phos)$ responded to grazing treatment (Table 1; Figs. 5, 6). In 2020, the ungrazed treatment soils were most N-limited relative to C and P, cattle grazed soils were least, and bison grazed soils were intermediate; in 2021, bison grazed soils were less N limited relative to C and P than either cattle grazed or ungrazed soils; and in 2022, bison grazed soils were less N limited relative to C than either cattle grazed or ungrazed soils, and less N limited relative to P than ungrazed soils (Fig. 5).

# Correlations

Many linear model correlations among soil characteristics and N cycling parameters were statistically



Fig. 4 A NP, B DNP, and C DEA in soils from different grazing treatments over 3 years. Tukey's HSD post-hoc results are shown with different letters indicating years (top) or grazing treatments (center) that differed from each other at P < 0.05

significant, but none had an  $R^2$  value higher than 0.15 (Table 2). Soil water content was positively correlated with DNP and negatively correlated with DEA, and DNP and DEA were negatively correlated with one another. Soil pH was positively



Fig. 5 A ln(BG):ln(NAG:LAP) and B ln(NAG:LAP):ln(Phos), indicators of relative soil microbial demand for C and N, and N and P, respectively, in soils from different grazing treatments over 3 years. Tukey's HSD post-hoc results are shown with different letters indicating grazing treatments that differed from each other at P < 0.05 level each year

correlated with resin-sorbed  $NO_3^--N$ , NP, and lnBG:ln(NAG+LAP). Resin-sorbed  $NH_4^+-N$  was positively correlated with resin-sorbed  $NO_3^--N$ , but negatively correlated with NP. Microbial N-limitation index values (lnBG:ln(NAG+LAP)), which are higher in less N-limited conditions, were positively correlated with both NP and DNP.

# Discussion

We wanted to learn whether bison and cattle influence soil microbial N cycling activities in tallgrass prairie similarly, and investigated this question over a three-year period. Our results show that bison and cattle grazing have qualitatively similar but



**Fig. 6** A Beta glucosidase, **B** cellobiohydrolase, **C** N-acetyl glucosaminidase, **D** leucine aminopeptidase, and **E** acid phosphatase enzyme activity rates (nmol substrate released  $g^{-1}$  dry soil  $h^{-1}$ ) through the summer growing season in 3 sampling

years. Tukey's HSD post-hoc results are shown with different letters indicating years or grazing treatment units within year that differed at the P < 0.05 level

 Table 2
 Correlation test results (r statistic and P values) among soil GWC and pH, N-cycling rates, and the microbial N limitation index

r, P	GWC <sup>Sqrt</sup>	pН	resin-NH4 <sup>+</sup> -N <sup>ln</sup>	resin-NO <sub>3</sub> <sup>-</sup> -N <sup>ln</sup>	NP <sup>ln</sup>	DNP <sup>ln</sup>	DEA <sup>sqrt</sup>
рН	- 0.06, 0.44						
resin-NH4 <sup>+</sup> -N <sup>ln</sup>	- 0.17, 0.12	0.01, 0.96					
resin-NO <sub>3</sub> <sup>-</sup> –N <sup>ln</sup>	- 0.14, 0.20	0.27, 0.013	0.35, < 0.001				
NP <sup>ln</sup>	- 0.02, 0.80	0.36, 0.002	- 0.27, 0.031	0.15, 0.25			
DNP <sup>ln</sup>	0.23, 0.005	0.15, 0.20	- 0.24, 0.050	0.11, 0.35	0.14, 0.09		
DEA <sup>sqrt</sup>	- 0.34, < 0.001	0.09, 0.47	- 0.02, 0.89	0.03, 0.81	0.13, 0.11	- 0.24, 0.004	
lnBG:ln(NAG+LAP)	0.02, 0.74	0.28, 0.001	0.08, 0.46	0.12, 0.30	0.24, 0.004	0.39, < 0.001	0.02, 0.86

Bolded values indicate P<0.05

Sqrt denotes square root transformation and In denotes natural log transformation

quantitatively different effects: Bison and cattle both tended to increase N availability and N cycling activities in comparison to ungrazed soils, but the influence of bison tended to be stronger than that of cattle. Specifically, soil pH, resin-sorbed nitrate and nitrification potentials (NP) were consistently highest in bisongrazed soils, and microbial N-limitation was lowest in bison-grazed soils in two of the three sampling years. Also, despite interannual variability in cattle versus bison responses, microbial N-limitation was always highest, and denitrification enzyme activity (DEA) was always lowest, in ungrazed soils relative to both cattle- and bison-grazed soils. However, the magnitude of temporal variation was stronger than grazing effects for soil water content, resin-sorbed N, nitrification potential (NP), and denitrification potentials (DNP). Resin-sorbed N was lowest in the driest year of the study, and both NP and DNP were highest, while DEA was lowest, in the wettest year of the study. Finally, while coarse relationships among measured variables support mechanistic discussion points at the interannual and grazing treatment scale, correlations were not strong enough to suggest predictive relationships among soil water content, pH, and microbial N-cycling variables at the soil sample scale.

# Interannual variation in soil water and microbial N cycling

Soil water content at the time of sampling was higher in 2020 than in 2021 and 2022, while summer precipitation was within the historic 95% confidence interval range in 2020 and 2022, but not in 2021, which was notably dry (Figs. 1, 2). Levels of  $NO_3^--N$ ,  $NH_4^+-N$ , and  $NH_4^+ - N + NO_3^- - N$  sorbed to resin bags coincided with this variability in water, such that with less than average precipitation there was less mineralized inorganic N (Fig. 3). In 2020, both NP and DNP were highest in magnitude, suggesting that wetter soil conditions supported higher mineralization and mobilization of the ammonium and nitrate substrates driving these two microbial metabolic activities (Fig. 4). Furthermore, extracellular enzyme activities (particularly N-acquiring activities) were lowest, indicating greater product availability (particularly of soluble nitrogenous compounds) and lower investment into enzyme production acquisition (Sinsabaugh and Follstad Shah 2012; Burns et al. 2013), during the wet year (Figs. 5, 6). The resin-sorbed N and microbial N-cycling datasets support the conclusion that interannually, N availability and N-cycling rates are positively associated with seasonal precipitation and soil water content.

The process of denitrification has complex controls, including limitation by nitrate availability, C availability, or low anoxia (Wallenstein et al. 2006; Robertson and Groffman 2014), and because DNP rates never reached DEA rate potential levels (Fig. 4), at least one of these factors must have limited the process. Anoxic conditions would have been highest in the wettest year, when oxygen diffusion into the soil pore space was most restricted. While we did not measure soil C availability, at the landscape level, growing season precipitation has a stronger effect than grazing on annual forage growth measured as aboveground net primary productivity (ANPP), as well as on root production (Johnson and Matchett 2001; Fay et al. 2003), so C availability from plant production should also have been highest in the wettest year. However, DNP rates remained only 9% (on average) of DEA rate potentials in the 2020 sampling year. Further, while DEA rate potentials did not decrease during the dry year of 2021, DNP dropped substantially, and was only detectable in bison-grazed soils where N availability was highest. Based on these observations, denitrification in this system is likely more limited by nitrate than by water, anoxia, or carbon, and only conditionally high, in agreement with conclusions made using the in situ amended core incubation technique by Groffman et al. (1993). The lack of recovery of DNP in 2022, despite a wetter summer overall, could be related to the long period of time following a precipitation event preceding the sampling time (Fig. 1), since as soils dry, soluble nutrient availability also declines, until rewetting stimulates pulses of microbial activity (Schimel 2018).

# Grazing and N availability

Mineralization of N from soil organic matter is a primary microbial mechanism that makes N available for plant and microbial uptake, and is controlled by microbial enzyme activity (Tabatabai et al. 2010). Resin bags are an index, not an in situ measure of mineralization, yet they can provide a reliable indicator of N mineralized during the growing season (Baer and Blair 2008; Nieland et al. 2021). Further, ratios of extracellular enzyme activities are indicative of relative microbial investment in N acquisition (Sinsabaugh and Follstad Shah 2012). N availability was higher in bison-grazed soils than ungrazed soils using both of these indicators in 2021 and 2022 (Figs. 3, 5). It is plausible that higher water availability was connected to the weaker bison effect on soil microbial N limitation in 2020, if wetter conditions supported greater inorganic N mineralization and lower N limitation overall, but unfortunately, we do not have resinsorbed N data from 2020 to corroborate this interpretation. Still, the evidence points to increased N availability in bison treatments for microbial immobilization and plant assimilation, relative to both cattle grazed and ungrazed treatment soils.

Despite the general bison grazing effect on N availability, specific N-cycling enzyme activities did not all respond the same. For example, in cattle and ungrazed treatments, polypeptide-decomposing

(LAP) activity supported higher N demand in 2021, while microbial cell wall-decomposing (NAG) activity did in 2022, suggesting that different components of the soil organic N pool were microbial N sources in each year (Fig. 6). Also, nitrate-N rather than ammonium-N responded to grazing treatment, suggesting that either ammonium uptake was lower, or nitrification was higher, in bison grazed soils (Fig. 3). Notably, despite interannual variation, bison grazed soils maintained a higher (less acidic) pH than cattle grazed soils (Fig. 2). Soil pH broadly constrains soil chemical transformations and the microbial enzymatic activities which drive N cycling, such that different N sources may support soil microbial N demand under different pH conditions due to changes in available N and microbial enzyme production (Sharpley 1991; Zeglin et al. 2007; Sinsabaugh et al. 2008; Nannipieri et al. 2018; Barber et al. 2023). In this study, soil pH was likely affected by bison grazing activity through urine and dung inputs, which add alkalinity to the soil (Somda et al. 1997; Hong et al. 2021).

## Grazing and nitrification potential

Both NP and resin-sorbed NO<sub>3</sub><sup>-</sup>-N were consistently higher in bison treatments (Figs. 3, 4), suggesting that under bison grazing, there is a higher likelihood of NO<sub>3</sub><sup>-</sup>-N becoming mobile in soil solution, and subsequently being taken up by plants or microorganisms, reduced by microorganisms and denitrified, or leached out of the soil. Overall, even though nitrification is a precursor for N loss through either nitrate leaching or denitrification, lower denitrification rates relative to nitrification rates suggest loss of soil N to the atmosphere in tallgrass prairie is a comparatively small factor in the N cycle at this site (Groffman et al. 1993; Blair et al. 1998). While nitrate leaching has not been constrained, we do know that stream water nitrate concentrations are low in this watershed despite the long-term grazing pressure (Dodds et al. 1996), and that local grasses can rapidly assimilate nitrate (Dell and Rice 2005); so while possible, nitrate production is not necessarily strongly tied to N leaching losses at the study site.

In addition to higher inorganic N availability overall, the consistently higher pH in bison grazed soils may boost NP through direct effects on ammonium availability. Chemically, soil pH controls the proportion of ammonia (NH<sub>3</sub>) in soil solution as ammonium  $(NH_4^+)$ , which in turn affects N availability for the process of nitrification (Kemmitt et al. 2006; Sahrawat 2008): With a higher pH, the nonprotonated form (ammonia) is favored, which supplies more of the substrate for ammonia monooxygenase, the rate limiting enzyme of nitrification found in both ammonia oxidizing bacteria and archaea (Nicol et al. 2008). This could help explain the positive relationships between pH and NP, and pH and resin-bound nitrate (Table 2). This finding is notable because nitrifier metabolism is the least functionally redundant in soils of all of the N-cycling processes measured in this study (Prosser and Nicol 2012) suggesting in turn that bison treatments could have higher abundance and/or a pH-specialized population of soil nitrifiers (Prosser and Nicol 2012), or a higher NP due to a more optimal soil pH.

## Grazing and denitrification potential

Denitrification, the process of NO<sub>3</sub><sup>-</sup>-N reduction to gaseous form, was measured in two ways: Under in situ soil N and C availability conditions (DNP) and with nitrate and DOC added to the assay to measure maximum denitrification enzyme activity (DEA). We found that DNP was much lower than DEA, especially in the drier years of 2021 and 2022 (Fig. 4). This indicates that while microbial biomass with enzymatic potential for denitrification exists, because either soil N or C substrate was limiting, little denitrification potential was realized. Only in bison grazed soils, where NP and nitrate availability was higher, was any DNP detected in 2021. In contrast, DEA was not different between bison and cattle treatments in 2021 or 2022, despite differences in NP and nitrate availability, while ungrazed treatments exhibited consistently lower NP, resin-sorbed nitrate, and DEA (Figs. 3, 4). This suggests that grazing intensity in general impacts soil DEA in some biologically similar way. Because many bacterial taxa carry the genetic potential to produce enzymes in the denitrification pathway (Nelson et al. 2016), whether or not conditions allow these enzymes to be used, general changes in the soil microbial community are more likely to affect DEA than the substrate-limited DNP (Wallenstein et al. 2006). At the same tallgrass prairie field site, bison dung is a microbial dispersal vector that increases soil microbial diversity and changes

microbial community composition (Hawkins and Zeglin 2022), and other investigations show that grazing by cattle, sheep, and goats can affect soil microbial composition (Clegg 2006; Eldridge et al. 2017; Wang et al. 2019). However, the redundancy of bison and cattle effects on the soil microbiome, and implications for denitrification, are not yet understood.

#### Grazing and grassland soil N cycling

Bison grazing substantially increased soil N-cycling rate potentials at this site in the North American Great Plains, and these elevated rates (NP of 0.05-0.5 µg N  $g^{-1}$  dry soil  $h^{-1}$ , DEA of 30–100 µg N kg<sup>-1</sup> dry soil  $h^{-1}$ ) were similar to those measured in other grazed grasslands worldwide. In Mongolia, grasslands that were grazed by sheep had an estimated NP of 0.5–1  $\mu$ g N g<sup>-1</sup> dry soil h<sup>-1</sup> and DEA of 400–700  $\mu$ g N kg<sup>-1</sup> dry soil h<sup>-1</sup> (Yingjin et al. 2022); in tropical savanna in the Ivory Coast, NP was 0-10 µg N  $g^{-1}$  dry soil  $h^{-1}$  and DEA was 0–100 µg N kg<sup>-1</sup> dry soil  $h^{-1}$  (Srikanthasamy et al. 2018). In cattle grazed Australian grasslands, NP was 0–0.5  $\mu$ g N g<sup>-1</sup> dry soil h<sup>-1</sup> and DEA was 20-40 µg N kg<sup>-1</sup> dry soil h<sup>-1</sup> (Mehnaz and Dijkstra 2016). Comparatively, our N cycling rates were close to those estimated in Australian grasslands but a bit lower than those estimated in Mongolia or West African savanna, suggesting that higher grazing intensity and more N recycling is occurring at those two sites and/or the physical and chemical soil properties and biological conditions for NP or DEA were different in the other grassland types. Historically, at the same tallgrass prairie site that we studied, but using different methods, rates were in a similar range as currently estimated: NP was 0.17–0.23  $\mu g$  N  $g^{-1}$  dry soil  $h^{-1}$  and DEA was 180–286  $\mu$ g N kg<sup>-1</sup> dry soil h<sup>-1</sup> (Groffman et al. 1993). Comparatively, given the three decades of additional grazing pressure between these two studies, it is somewhat surprising that rates are so similar. However, because variable precipitation mediates the magnitude of N-cycling, rates could be more strongly linked to the direct and indirect effects of soil water than to grazing intensity over time.

This tallgrass prairie region and world as a whole is predicted to have more variable climactic conditions in the future (IPCC 2022) impacting soil water and thus large grazers and microbial N cycling activities in this region and many others (Nippert et al. 2022; Abraham et al. 2023). Our nitrification potentials and denitrification estimates suggest losses of N from the soil are possible, and that higher losses are possible in bison-grazed areas. A recent metaanalysis of greenhouse gas emissions on grasslands showed that heavy grazing intensity did not increase N<sub>2</sub>O-N emissions, but instead overgrazing severely degraded rangeland habitat leading to soil runoff (Tang et al. 2019). This result suggests that management is the key to mitigating grazing animal effects on N loss from rangelands, specifically, maintaining proper stocking densities based on set carrying capacity (Holechek et al. 2011) and considering how the physical movement of animals controls N export and spatial heterogeneity (Coetsee et al. 2023).

Despite the clear enhancement of soil N-cycling rates, particularly nitrification potential, by bison grazing in this study, we do not know whether nitrate was subsequently leached out, assimilated by plants, or immobilized by microorganisms and retained in soil organic matter. Because the physical properties of these soils promote the retention of N in mineralassociated fractions (Soong and Cotrufo 2015) and the native biota conserve N tightly in both soil microbial biomass and plant tissue (Dell et al. 2005), elevated internal cycling rates may be coupled with N turnover through biotic pools, rather than linked to N losses from the ecosystem. Compared to ungrazed and annually burned tallgrass prairie, N cycling is not as "open" i.e., N recycling is slower and more closed in ungrazed tallgrass prairie even though there are larger inorganic N pools then grazed prairie (Connell et al. 2020). As such, ecosystem N retention may be high in all treatments of this study because they are burned annually (Dell et al. 2005). However, additional research is necessary to directly measure grazing effects on soil and ecosystem N retention.

#### Conclusion

We found that, despite interannual variability, bison impacted nitrification differently than cattle because of greater available soil inorganic N, higher soil pH and less N limitation. Also, annual variation in N cycling was apparent because of variability in summer precipitation and soil water availability controlling N cycling. As such, tracking soil characteristics and N cycling activities over time gives primary diagnostic information of N recycling and should strongly be considered in rangeland monitoring health assessments, as it is currently lacking (Pellant et al. 2020). However, due to concomitant differences in bison and cattle behavior (e.g., bison tendency to maintain grazing lawns) with differences in management (i.e., bison graze year-round while cattle are on pasture May-October), we cannot discern whether N cycling effects are inherent to the different grazers per se or to the human-grazer interaction. Human societies past and present around the world have always depended on large grazing animals, in particular the bison, and more recently cattle, in North America. Bison are tied to the existing cultural identity of tribal nations that have lived on grasslands for at tens of thousands of years (Kornfeld et al. 2016), and while in the past bison movement was not directly constrained by humans, bison numbers were orders of magnitude higher across their continental range. Thus, it is also difficult to speculate the extent to which soil fertility changed following the replacement of bison with cattle; however, our data suggest that a significant change was possible. Also knowing that bison grazing can dramatically shift grassland plant diversity (Ratajczak et al. 2022), it is further possible that the interactions between grazers, soils, and plants that maintained Great Plains grassland ecosystems pre-colonialization may have been different from what we understand today. Long term data on N cycling activities are sparse, and considering the data here in the context of grazer management and changing climate will provide useful information for future N cycling and budgeting in grassland ecosystems.

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**Data availability** The data generated and analyzed in this study are publicly available within the Environmental Data Initiative Repository (https://doi.org/https://doi.org/10.6073/pasta/34b4b622191f222c90e7d227291951c8).

#### Declarations

**Competing interests** The authors have no relevant financial or non-financial interests to disclose.

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# References

- Abraham AJ et al (2023) Understanding anthropogenic impacts on zoogeochemistry is essential for ecological restoration. Restor Ecol 31:e13778. https://doi.org/10.1111/rec.13778
- Allison SD, Vitousek PM (2005) Responses of extracellular enzymes to simple and complex nutrient inputs. Soil Biol Biochem 37:937–944. https://doi.org/10.1016/j.soilbio. 2004.09.014
- Allred BW, Fuhlendorf SD, Hamilton RG (2011) The role of herbivores in Great Plains conservation: comparative ecology of bison and cattle. Ecosphere 2:1–17. https://doi. org/10.1890/ES10-00152.1
- Anderson RC (2006) Evolution and origin of the central grassland of North America: climate, fire, and mammalian grazers. JTBS 133:626–647
- Augustine DJ, Frank DA (2001) Effects of migratory grazers on spatial heterogeneity of soil nitrogen properties in a grassland ecosystem. Ecology 82:3149–3162

- Baer SG, Blair JM (2008) Grassland establishment under varying resource availability: a test of positive and negative feedback. Ecology 89:1859–1871. https://doi.org/ 10.1890/07-0417.1
- Bakker C, Blair JM, Knapp AK (2003) Does resource availability, resource heterogeneity, or species turnover mediate changes in plant species richness in grazed grasslands? Oecologia 137:385–391. https://doi.org/10.1007/ s00442-003-1360-y
- Barber NA et al (2023) Restoration age and reintroduced bison may shape soil bacterial communities in restored tallgrass prairies. FEMS Microbiol Ecol. https://doi.org/ 10.1093/femsec/fiad007
- Blair J (2023) CBH01: Konza Prairie bison herd information. https://doi.org/10.6073/PASTA/4912DE8FAF25E93 D45551CB1891E7FAD
- Blair JM (1997) Fire, N availability, and plant response in grasslands: A test of the transient maxima hypothesis. Ecology 78:2359–2368
- Blair JM et al (1998) Terrestrial nutrient cycling in tallgrass prairie, Grassland dynamics: long-term ecological research in tallgrass prairie. Oxford University Press, New York, pp 222–243
- Broderick CM et al (2022) Climate segacy effects shape tallgrass prairie nitrogen cycling. J Geophy Res Biogeosci. https://doi.org/10.1029/2022JG006972
- Burns RG et al (2013) Soil enzymes in a changing environment: current knowledge and future directions. Soil Biol Biochem 58:216–234. https://doi.org/10.1016/j.soilbio. 2012.11.009
- Chen H et al (2022) Carbon and nitrogen cycling on the Qinghai-Tibetan Plateau. Nat Rev Earth Environ 3:701– 716. https://doi.org/10.1038/s43017-022-00344-2
- Christopherson RJ, Hudson RJ, Christophersen MK (1978) Seasonal energy expenditures and thermoregulatory responses of bison and cattle. Can J Anim Sci. https:// doi.org/10.4141/cjas79-077
- Clegg CD (2006) Impact of cattle grazing and inorganic fertilizer additions to managed grasslands on the microbial community composition of soils. Appl Soil Ecol 31:73– 82. https://doi.org/10.1016/j.apsoil.2005.04.003
- Coetsee C, Wigley BJ, Sankaran M et al (2023) Contrasting effects of grazing vs browsing herbivores determine changes in soil fertility in an East African Savanna. Ecosystems 26:161–173. https://doi.org/10.1007/ s10021-022-00748-7
- Connell RK, Nippert JB, Blair JM (2020) Three decades of divergent land use and plant community change alters soil C and N content in tallgrass prairie. J Geophys Res Biogeosci. https://doi.org/10.1029/2020JG005723
- Coppedge BR, Leslie DM, Shaw JH (1998a) Botanical composition of bison diets on tallgrass prairie in Oklahoma. J Range Manag 51:379–382. https://doi.org/10.2307/ 4003321
- Coppedge BR, Engle DM, Toepfer CS, Shaw JH (1998b) Effects of seasonal fire, bison grazing and climatic variation on tallgrass prairie vegetation. Plant Ecol 139:235–246. https://doi.org/10.1023/A:1009728104 508
- Coppedge BR, Shaw JH (1997) Effects of horning and rubbing behavior by bison (*Bison bison*) on woody vegetation in

a tallgrass prairie landscape. Am Midl Nat 138:189–196. https://doi.org/10.2307/2426665

- Dell CJ, Rice CW (2005) Short-term competition for ammonium and nitrate in tallgrass prairie. SSSAJ 69:371–377. https://doi.org/10.2136/sssaj2005.0371
- Dell CJ, Williams MA, Rice CW (2005) Partitioning of nitrogen over five growing seasons in tallgrass prairie. Ecology 86:1280–1287. https://doi.org/10.1890/03-0790
- Detling JK (1988) Grasslands and savannas: regulation of energy flow and nutrient cycling by herbivores. In: Pomeroy LR, Alberts JJ (eds) Concepts of ecosystem ecology. Springer, New York, pp 131–148
- Dodds WK et al (1996) Nitrogen transport from tallgrass prairie watersheds. J Environ Qual 25:973–981. https://doi. org/10.2134/jeq1996.00472425002500050007x
- Dunbar-Ortiz R (2014) An indigenous peoples' history of the United States. Beacon Press, Boston
- Eldridge DJ et al (2017) Competition drives the response of soil microbial diversity to increased grazing by vertebrate herbivores. Ecology 98:1922–1931. https://doi.org/10. 1002/ecy.1879
- Enquist BJ et al (2020) The megabiota are disproportionately important for biosphere functioning. Nat Commun 11:699. https://doi.org/10.1038/s41467-020-14369-y
- Fay PA et al (2003) Productivity responses to altered rainfall patterns in a C4-dominated grassland. Oecologia 137:245–251. https://doi.org/10.1007/s00442-003-1331-3
- Feir D, Gillezeau R, Jones MEC (2021) The slaughter of the bison and reversal of fortunes on the Great Plains. SSRN J. https://doi.org/10.2139/ssrn.4196271
- Flores D (1991) Bison ecology and bison diplomacy: The Southern plains from 1800 to 1850. J Am Hist 78:465. https://doi.org/10.2307/2079530
- Frank DA, Evans RD (1997) Effects of native grazers on grassland N cycling in Yellowstone National Par. Ecology 78:2238–2248
- Groffman PM et al (2009) Challenges to incorporating spatially and temporally explicit phenomena (hotspots and hot moments) in denitrification models. Biogeochemistry 93:49–77. https://doi.org/10.1007/s10533-008-9277-5
- Groffman PM, Rice CW, Tiedje JM (1993) Denitrification in a tallgrass prairie landscape. Ecology 74:855–862. https:// doi.org/10.2307/1940811
- Hawkins JH, Zeglin LH (2022) Microbial dispersal, including bison dung vectored dispersal, increases soil microbial diversity in a grassland ecosystem. Front Microbio 13:825193. https://doi.org/10.3389/fmicb.2022.825193
- Hobbs NT (1996) Modification of ecosystems by ungulates. J Wildl Manag 60:695–713. https://doi.org/10.2307/38023 68
- Holechek JL, Pieper RD, Herbel CH (2010) Range management: principles and practices, 6th edn. Pearson Education
- Hong J et al (2021) Significant soil acidification caused by grazing exclusion across China's grassland areas. Land Degrad Dev 32:535–545. https://doi.org/10.1002/ldr.3722
- Hood-Nowotny R et al (2010) Alternative methods for measuring inorganic, organic, and total dissolved nitrogen in soil. SSSAJ 74:1018–1027. https://doi.org/10.2136/sssaj2009. 0389
- Hornaday WT (1913) Our vanishing wild life; its extermination and preservation. C. Scribner's sons, New York

- Hulbert LC (1986) Fire effects on tallgrass prairie. Proceedings of the ninth North American Prairie conference. GK Clambey and RH Pemble, Fargo, pp 138-142
- IPCC (2022) Synthesis report of the IPCC, sixth assessment report (AR6). https://www.ipcc.ch/report/ar6/syr/
- Johnson LC, Matchett JR (2001) Fire and grazing regulate belowground processes in tallgrass prairie. Ecology 82:3377–3389. https://doi.org/10.2307/2680159
- Kemmitt SJ et al (2006) pH regulation of carbon and nitrogen dynamics in two agricultural soils. Soil Biol Biochem 38:898–911. https://doi.org/10.1016/j.soilbio.2005.08.006
- Knapp AK et al (1999) The keystone role of bison in North American tallgrass prairie: Bison increase habitat heterogeneity and alter a broad array of plant, community, and ecosystem processes. Bioscience 49:39–50. https://doi. org/10.1525/bisi.1999.49.1.39
- Kohl MT et al (2013) Bison versus cattle: are they ecologically synonymous? Rangel Ecol Manag 66:721–731. https:// doi.org/10.2111/REM-D-12-00113.1
- Kornfeld M, Frison GC, Larson ML (2016) Prehistoric huntergatherers of the High Plains and Rockies, 3rd edn. Routledge, London
- Larson DM et al (2013) Blazing and grazing: influences of fire and bison on tallgrass prairie stream water quality. Freshw Sci 32:779–791. https://doi.org/10.1899/12-118.1
- List R et al (2007) Historic distribution and challenges to bison recovery in the northern Chihuahuan desert. Conserv Biol 21:1487–1494. https://doi.org/10.1111/j.1523-1739.2007. 00810.x
- Lott DF (2002) American bison: a natural history. University of California Press, Berkeley
- McMillan BR, Pfeiffer KA, Kaufman DW (2011) Vegetation responses to an animal-generated disturbance (bison wallows) in tallgrass prairie. Am Midl Nat 165:60–73. https:// doi.org/10.1674/0003-0031-165.1.60
- McMillan NA et al (2019) Plant community responses to bison reintroduction on the Northern Great Plains, United States: a test of the keystone species concept. Restor Ecol 27:379–388. https://doi.org/10.1111/rec.12856
- McMillan NA et al (2021) Are bison movements dependent on season and time of day? Investigating movement across two complex grasslands. Ecosphere 12:e03317. https:// doi.org/10.1002/ecs2.3317
- McMillan NA et al (2022) Bison movements change with weather: implications for their continued conservation in the Anthropocene. Ecol Evol 12:e9586. https://doi.org/10. 1002/ece3.9586
- McNaughton SJ (1983) Compensatory plant growth as a response to herbivory. Oikos 40:329–336. https://doi.org/ 10.2307/3544305
- Mehnaz KR, Dijkstra FA (2016) Denitrification and associated N<sub>2</sub>O emissions are limited by phosphorus availability in a grassland soil. Geoderma 284:34–41. https://doi.org/10. 1016/j.geoderma.2016.08.011
- Nannipieri P et al (2018) Soil enzyme activity: a brief history and biochemistry as a basis for appropriate interpretations and meta-analysis. Biol Fertil Soils 54:11–19. https://doi. org/10.1007/s00374-017-1245-6
- Nelson MB, Martiny AC, Martiny JBH (2016) Global biogeography of microbial nitrogen-cycling traits in soil. PNAS 113:8033–8040. https://doi.org/10.1073/pnas.1601070113

- Nicol GW et al (2008) The influence of soil pH on the diversity, abundance and transcriptional activity of ammonia oxidizing archaea and bacteria. Environ Microbiol 10:2966–2978. https://doi.org/10.1111/j.1462-2920.2008. 01701.x
- Nieland MA et al (2021) Differential resilience of soil microbes and ecosystem functions following cessation of long-term fertilization. Ecosystems 24:2042–2060. https://doi.org/ 10.1007/s10021-021-00633-9
- Nieland MA et al (2024) Product-inhibition feedbacks, not microbial population level tradeoffs or soil pH, regulate decomposition potential under nutrient eutrophication. Soil Biol Biochem 189:109247. https://doi.org/10.1016/j. soilbio.2023.109247
- Nippert J et al (2022) Climate change in grassland ecosystems. In: Groskinsky B (ed) Climate actions: local applications and practical solutions, 1st edn. CRC Press, Boca Raton, pp 65–100
- Olson KC (2023) PBG06: Cattle grazing and cattle performance in the Patch-Burn Grazing experiment at Konza Prairie. https://doi.org/10.6073/PASTA/28C9AA49E9 42E3BD489A05E867F38ECC
- Ojima DS, Schimel DS, Parton WJ, Owensby CE (1994) Longand short-term effects of fire on nitrogen cycling in tallgrass prairie. Biogeochemistry 24:67–84. https://doi.org/ 10.1007/BF02390180
- Pellant M et al. (2020) Interpreting Indicators of Rangeland Health, Version 5: Bureau of Land Management Technical Reference 1734–6. Bureau of Land Management. https:// pubs.er.usgs.gov/publication/70215720
- Plumb GE, Dodd JL (1993) Foraging ecology of bison and cattle on a mixed prairie: implications for natural area management. Ecol Appl 3:631–643. https://doi.org/10.2307/ 1942096
- Prosser JI, Nicol GW (2012) Archaeal and bacterial ammoniaoxidisers in soil: the quest for niche specialisation and differentiation. Trends Microbiol 20:523–531. https://doi. org/10.1016/j.tim.2012.08.001
- Ratajczak Z et al (2022) Reintroducing bison results in longrunning and resilient increases in grassland diversity. PNAS 119:e2210433119. https://doi.org/10.1073/pnas. 2210433119
- Raynor EJ et al (2021) Cattle grazing distribution patterns related to topography across diverse rangeland ecosystems of North America. Range Ecol Manag 75:91–103. https:// doi.org/10.1016/j.rama.2020.12.002
- Risser PG, Parton WJ (1982) Ecosystem analysis of the tallgrass prairie: nitrogen cycle. Ecology 63:1342–1351. https://doi.org/10.2307/1938862
- Robertson GP, Groffman PM (2014) Nitrogen transformations. In: Paul EA (ed) Soil microbiology, ecology and biochemistry, 4th edn. Academic Press, San Diego, pp 341–364
- Roy S, Naidu DGT, Bagchi S (2023) Functional substitutability of native herbivores by livestock for soil carbon stock is mediated by microbial decomposers. Glob Change Biol 29:2141–2155. https://doi.org/10.1111/gcb.16600
- R Core Team (2022) R: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna
- Samson FB, Knopf FL, Ostlie WR (2004) Great Plains ecosystems: past, present, and future. Wildl Soc Bull 32:6–15

- Sahrawat KL (2008) Factors affecting nitrification in soils. Commun Soil Sci Plant Anal 39:1436–1446. https://doi. org/10.1080/00103620802004235
- Schimel JP (2018) Life in dry soils: effects of drought on soil microbial communities and processes. Annu Rev Ecol Evol Syst 49:409–432. https://doi.org/10.1146/annurevecolsys-110617-062614
- Schimel JP, Weintraub MN (2003) The implications of exoenzyme activity on microbial carbon and nitrogen limitation in soil: a theoretical model. Soil Biol Biochem 35:549– 563. https://doi.org/10.1016/S0038-0717(03)00015-4
- Schlesinger WH, Bernhardt ES (2020) Biogeochemistry: an analysis of global change, 4th edn. Academic Press, London
- Schlesinger WH, Hartley AE (1992) A global budget for atmospheric NH<sub>3</sub>. Biogeochemistry 15:191–211. https:// doi.org/10.1007/BF00002936
- Sharpley AN (1991) Effect of soil pH on cation and anion solubility. Commun Soil Sci Plant Anal 22:827–841. https:// doi.org/10.1080/00103629109368457
- Shaw H (1995) How many bison originally populated western rangelands? Rangelands 17:148–150
- Sinsabaugh R, Follstad Shah J (2012) Ecoenzymatic stoichiometry and ecological theory. Annu Rev Ecol Evol Syst 43:313–343. https://doi.org/10.1146/annurev-ecols ys-071112-124414
- Sinsabaugh RL, Hill BH, Follstad Shah JJ (2009) Ecoenzymatic stoichiometry of microbial organic nutrient acquisition in soil and sediment. Nature 462:795–798. https://doi. org/10.1038/nature08632
- Sinsabaugh RL et al (2008) Stoichiometry of soil enzyme activity at global scale: stoichiometry of soil enzyme activity. Ecol Lett 11:1252–1264. https://doi.org/10. 1111/j.1461-0248.2008.01245.x
- Soong JL, Cotrufo MF (2015) Annual burning of a tallgrass prairie inhibits C and N cycling in soil, increasing recalcitrant pyrogenic organic matter storage while reducing N availability. Glob Change Biol 21:2321–2333. https://doi. org/10.1111/gcb.12832
- Somda ZC, Powell JM, Batoino A (1997) Soil pH and nitrogen changes following cattle and sheep urine deposition. Commun Soil Sci Plant Anal 28:1253–1268. https://doi.org/10. 1080/00103629709369872
- Srikanthasamy T et al (2018) Contrasting effects of grasses and trees on microbial N-cycling in an African humid savanna. Soil Biol Biochem 117:153–163. https://doi.org/ 10.1016/j.soilbio.2017.11.016

- Tabatabai MA, Ekenler M, Senwo ZN (2010) Significance of enzyme activities in soil nitrogen mineralization. Commun Soil Sci Plant Anal 41:595–605. https://doi.org/10. 1080/00103620903531177
- Tang S et al (2019) Heavy grazing reduces grassland soil greenhouse gas fluxes: a global meta-analysis. Sci Total Environ 654:1218–1224. https://doi.org/10.1016/j.scito tenv.2018.11.082
- Taylor AE, Zeglin LH, Dooley S, Myrold DD, Bottomley PJ (2010) Evidence for different contributions of archaea and bacteria to the ammonia-oxidizing potential of diverse Oregon soils. AEM 76:7691–7698. https://doi.org/10. 1128/AEM.01324-10
- Towne EG, Hartnett DC, Cochran RC (2005) Vegetation trends in tallgrass prairie from bison and cattle grazing. Ecol Appl 15:1550–1559. https://doi.org/10.1890/04-1958
- Vinton MA, Hartnett DC, Finck EJ, Briggs JM (1993) Interactive effects of fire, bison (*Bison bison*) grazing and plant community composition in tallgrass prairie. Am Midl Nat 129:10–18. https://doi.org/10.2307/2426430
- Wallenstein MD et al (2006) Environmental controls on denitrifying communities and denitrification rates: insights from molecular methods. Ecol Appl 16:2143–2152
- Wang Z et al (2019) Impact of long-term grazing exclusion on soil microbial community composition and nutrient availability. Biol Fertil Soils 55:121–134. https://doi.org/10. 1007/s00374-018-013365
- Woodmansee RG, Duncan DA (1980) Nitrogen and phosphorus dynamics and budgets in annual grasslands. Ecology 61:893–904. https://doi.org/10.2307/1936759
- Yingjin S et al (2022) Short-term grazing rather than mowing stimulates N<sub>2</sub>O production potential through enhancing the bacterial pathway in semiarid grasslands. J Soils Seds 22:32–42. https://doi.org/10.1007/s11368-021-03051-5
- Zeglin LH et al (2007) Microbial responses to nitrogen addition in three contrasting grassland ecosystems. Oecologia 154:349–359. https://doi.org/10.1007/s00442-007-0836-6

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