

Glomalin and Soil Aggregation under Six Management Systems in the Northern Great Plains, USA

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ABSTRACT

The soil environment is linked to aboveground management including plant species composition, grazing intensity, levels of soil disturbance, residue management, and the length of time of a living plant is growing. Soil samples were collected under rangeland [native grass, rotational grazing (NGRG); tame grass, heavy grazing (TGRG); and tame grass, rotational grazing (TGHG)] and cropland [conventional till (CT); CT plus manure (CTM); and long term no till (NT)] systems. The rangeland systems were hypothesized to have higher glomalin content [measured as Bradford-reactive soil protein (BRSP)] and water stable aggregation (WSA) than the cropland systems. In addition, within both rangeland and cropland systems, BRSP and WSA were expected to decline with increased disturbance due to grazing or tillage and going from native to introduced plant species. Differences were detected for BRSP with NGRG and CTM having the highest values in range and cropland systems, respectively. However, the CTM system had higher BRSP values than one or both of the tame grass systems while the CT and NT systems had similar values. Correlation analysis showed strong relationships between all of the BRSP values and WSA.

Keywords: Glomalin; Arbuscular Mycorrhizal Fungi; Soil Aggregation; Grazing Systems; Crop Production Systems

1. Introduction

The symbiotic relationship between arbuscular mycorrhizal fungi (AMF) and about 90% of all vascular plants provides a direct linkage between above and below-ground activities [1,2]. Soil aggregation is a mechanism used to engineer the soil environment to maximize water and nutrient use efficiency [3]. Glomalin is a glycoprotein produced by AMF which is strongly linked to aggregation [4-6]. The aggregation process and glomalin production are both a carbon cost to AMF and may be hindered by soil disturbance through tillage [4-11] or overgrazing [12-20]. Therefore, the level of mycorrhizal dependency [1] of the host plant as well as photosynthetic activity may impact glomalin content and water stable aggregation (WSA) [2].

In rangeland systems, Liebig *et al.* [21] found that systems under heavy grazing intensity had higher soil C values than systems under moderate intensity. However, Klumpp *et al.* [19] and Ingram *et al.* [20] found that increases in grazing intensity decreased soil C. Unlike the Liebig *et al.* [21] study, these authors also examined mi-

crobial community and soil C dynamics under different grazing treatments and found that the rhizosphere environment was the driver behind changes in soil C [19,20]. These results are worth noting because often soil C levels are related to soil aggregation and glomalin concentrations [2, 22].

In cropland systems, no till (NT) treatments had higher soil organic C levels, glomalin concentrations, and more water stable aggregates than conventional till (CT) treatments [5-8,11]. These differences tended to increase with longer periods of time in NT [5,6]. Pikul *et al.* [4] found that particulate organic matter (POM) increased under NT while aggregate wettability decreased. Greater amounts of POM may support higher aggregate stability and water holding capacity while a decrease in aggregate wettability may negatively impact water infiltration and water holding capacity. In addition to C and N levels and aggregate stability, glomalin values also may be related to other AMF parameters such as spore counts, hyphal length or root colonization but these relationships are inconsistent [7,8,13,15,18,23].

Bradford-reactive soil protein (BRSP) and water stable aggregation (WSA) were measured at six sites near Platte, SD, USA. These sites included three rangelands [native grass, rotational grazing (NGRG); tame grass, heavy grazing (TGRG); and tame grass, rotational grazing (TGHG)] and three croplands [conventional till (CT); CT plus manure (CTM); and long term no till (NT)] systems. We hypothesized that the rangeland systems would have higher BRSP and WSA values than the cropland systems and values would increase across these systems in the following order: CT < CTM < NT < TGHG < TGRG < NGRG.

2. Materials and Methods

2.1. Site Description

Soil samples were collected on June 7, 2004 near Platte, SD, USA from under six management systems (NGRG, TGRG, TGHG, CT, CTM, and NT). The sites were selected to show a wide range of management scenarios on the same soil type (an Eakin silty clay loam or fine-silty, mixed, superactive, mesic Typic Argiustolls) and under the same climatic conditions (*i.e.* all six sites within a 10-km radius). The major grass species at the NGRG was big bluestem (*Andropogon gerardii*) with a small amount of switchgrass (*Panicum virgatum* L.). The grass species at the TGRG sites were predominantly brome grass (*Bromus inermis*) and crested wheatgrass (*Agropyron cristatum*) with small amounts of intermediate wheatgrass (*Thinopyrum intermedium*). The grass species at the TGHG was Kentucky bluegrass (*Poa pratensis*) with some brome grass. The cropping systems all had a corn-wheat rotation. Two samples were collected from each site at the 0 to 10 cm depth to create a bulk sample. A portion of the soil was sequentially passed through a series of screens to collect 1 to 2 mm aggregates.

2.2. Soil Analyses

Six 2-g subsamples were removed from both bulk soil and 1 to 2 mm aggregate samples and extracted for glomalin-related soil protein (GRSP). Soil was extracted with 50 mM sodium citrate, pH 8.0, for 15 to 26 1-hr cycles at 121°C [24] followed by extraction with 100 mM sodium pyrophosphate, pH 9.0, for four to nine 1-hr cycles at 121°C to remove a recalcitrant fraction of glomalin [2]. For both the citrate and the sodium pyrophosphate extractions, samples were centrifuged after each 1-hr extraction cycle to pellet the soil and collect the supernatant. All citrate or pyrophosphate extractions were combined, and the total volume measured.

The Bradford total protein assay was used to measure GRSP as Bradford reactive soil protein (BRSP) [2]. For the samples extracted with pyrophosphate, an aliquot equal in volume to the aliquot used for the unknowns of

100 mM pyrophosphate was added to the BSA standard to remove background cross-reactivity. Total protein values are calculated on a mg Bradford-reactive soil protein (BRSP) g⁻¹ soil or 1 to 2 mm aggregates basis. Both the bulk soil and 1 to 2 mm aggregates were corrected for coarse material in the sample using forced water to pass the extracted soil or aggregates through a 1 mm screen leaving the coarse fraction on the screen. Water stable aggregation (WSA) was measured on four subsamples of the 1 to 2 mm aggregates using methodology modified from Kemper and Rosenau [25] and described by Nichols and Wright [2].

2.3. Statistical Analysis

Means and standard error of the mean (SEM) were calculated for the BRSP and WSA. All means comparisons were made at the $\alpha \leq 0.05$ level using PROC MIXED (SAS software, ver. 9.1, SAS Institute, 2004) after the residuals met the assumptions for normality and homogeneity of variance. The correlations between BRSP, for both the citrate and pyrophosphate extractions and WSA were measured using PROC CORR after meeting the assumption for normal data distribution (SAS software, ver. 9.1, SAS Institute, 2004).

3. Results and Discussion

Concentrations of BRSP in both the bulk soil and 1 to 2 mm aggregate samples were similar for each of the individual management scenarios with the exception of the CT site which was lower in the bulk soil than in the 1 to 2 mm aggregates (**Table 1**). The three grazing systems had the highest glomalin concentrations with the NGRG system having the significantly highest values. Because most studies have shown that as grazing intensity increases, soil carbon levels and microbial activities decrease [19], it was unexpected that the TGRG site would have lower BRSP values than the TGHG system. However, a study by Liebig *et al.* [21] showed that a heavily grazed treatment had higher soil carbon levels than a moderately grazed treatment at sites in North Dakota which were in the same semi-arid ecoregion as the sites in this study but were approximately 560 km apart. These studies did indicate that the results were probably dependent upon if the plant species composition was dominated by native or tame grasses which impacts the intensity of the mycorrhizal relationship [1,13] and impacts rooting depth and rhizosphere dynamics [19-21].

In the crop production systems, the CTM site had the highest values while there was no difference between the CT and NT systems in the bulk soil but the CT site had significantly higher BRSP in the 1 to 2 mm aggregates. This was also unexpected since most studies have shown glomalin, or BRSP, increases with longer periods of NT [4-6]. The addition of manure has been shown to increase

Table 1. Values^a for glomalin extracted with sodium citrate (cit) followed by sodium pyrophosphate (pyro) from either bulk soil or 1 to 2 mm aggregates (aggreg) and measured as Bradford reactive soil protein (BRSP) on a mg protein g⁻¹ soil or aggregate weight basis and for water stable aggregation (WSA) at a% of total aggregates. Values are means^b ± SEs for each management system: native grass, rotational grazing (NGRG); tame grass, heavy grazing (TGRG); tame grass, rotational grazing (TGHG); long term no till (NT); conventional till (CT); and CT plus manure (CTM).

	NGRG	TGRG	TGHG	NT	CT	CTM
BRSP cit soil	9.555 ± 0.187 ^A	6.226 ± 0.249 ^B	6.895 ± 0.090 ^B	4.618 ± 0.152 ^D	4.354 ± 0.222 ^D	5.447 ± 0.128 ^C
BRSP pyro soil	0.700 ± 0.028 ^A	0.357 ± 0.014 ^C	0.379 ± 0.010 ^{BC}	0.252 ± 0.014 ^D	0.266 ± 0.012 ^D	0.439 ± 0.011 ^B
BRSP total soil	10.255 ± 0.166 ^A	6.583 ± 0.246 ^{BC}	7.274 ± 0.081 ^B	4.869 ± 0.161 ^D	4.619 ± 0.222 ^D	5.887 ± 0.120 ^C
BRSP cit aggreg	9.291 ± 0.113 ^A	6.423 ± 0.064 ^C	7.165 ± 0.092 ^B	5.093 ± 0.135 ^E	5.664 ± 0.146 ^D	6.128 ± 0.166 ^{CD}
BRSP pyro aggreg	0.575 ± 0.026 ^A	0.455 ± 0.021 ^B	0.301 ± 0.022 ^C	0.276 ± 0.013 ^C	0.325 ± 0.032 ^C	0.502 ± 0.021 ^{AB}
BRSP total aggreg	9.865 ± 0.090 ^A	6.878 ± 0.072 ^C	7.466 ± 0.096 ^B	5.369 ± 0.142 ^E	5.989 ± 0.152 ^D	6.630 ± 0.179 ^C
WSA	89.063 ± 1.028 ^A	83.125 ± 0.633 ^B	45.688 ± 0.806 ^C	24.875 ± 0.415 ^E	24.063 ± 0.544 ^E	42.063 ± 1.082 ^D

^aValues in a row which are followed by a different letter are significantly different at Prob < 0.05. ^bN equals six for the BRSP values and four for the WSA.

soil C, which may stimulate aggregation, and BRSP (Wright, personal communication). In addition, other organic compounds, such as humic substances and tannins may be co-extracted with glomalin and measured with the Bradford assay [2,26,27]. This might explain the CTM treatment values, but it was unclear in this study why the long term NT system did not have higher BRSP values than the CT site.

The authors speculate that the crops grown at the cropland sites might have impacted carbon allocation to AMF and AMF allocation to glomalin production and aggregation.

Despite these unexpected results from BRSP and WSA values, the Pearson Correlation Coefficients showed BRSP was strongly and significantly correlated WSA, $r > 0.7$ in almost all cases (**Table 2**). This is similar to many other studies where BRSP and WSA are correlated.

Native warm-season grasses, such as big bluestem and switchgrass, are all highly mycorrhizal while cool-season grasses are less dependent upon the mycorrhizal relationship [1,13]. Mycorrhizal fungi and glomalin production are sensitive to carbon exudates from the plant and soil disturbance. Additionally, AMF and glomalin are important in forming and stabilizing aggregates which play significant roles in water infiltration and water retention [5,6,14]. Finally, grazing pressure has been shown to decrease the amount of carbon fixed by photosynthesis and exuded from the roots [19,20]. For the most part, this did occur in this study but given the non-replicated design and the limited number of samples collected at each site, and the limited number of studies on glomalin and WSA in rangeland systems in the literature, more research is needed. Future studies will focus on the biological, chemical, and physical interactions occurring in the soil, particularly in the rhizosphere, which may be used to model above and belowground relationships. These models will help us understand the

linkage between levels of soil disturbance and the types and amounts of photosynthetically-derived carbon going belowground, especially as it relates to the plant-mycorrhizal relationship. Ongoing research projects in the northern Great Plains are measuring GRSP and WSA in other range and cropland studies. The range studies include systems with long-term (over 60 years) continuous management different levels of grazing intensity from no grazing to heavy grazing while the cropland studies incorporate perennial and cover crops which may be used as forages.

4. Conclusion

Similar to many other studies, glomalin and aggregate stability were strongly correlated in this study. However, the BRSP and WSA values did not increase from CT < CTM < NT < TGHG < TGRG < NGRG as expected. This was noted in particular in the cropland sites where CTM had the highest values and CT and NT were similar. The complex relationship between AMF and host plant especially as it relates to carbon allocation needs more studies to understand glomalin production and how it relates to aggregation.

5. Abbreviations and Acronyms

BRSP, Bradford reactive soil protein; C, carbon; CT, conventional till; CTM, conventional till plus manure; GRSP, glomalin-related soil protein; IR, infiltration rate; NGRG, native grass rotational grazing; NT, long-term no-till; TGHG, tame grass heavy grazing; TGRG, tame grass rotational grazing; WHC_s, water holding capacity; WSA, water stable aggregation.

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Table 2. Correlation coefficients^a for glomalin extracted with sodium citrate (cit) followed by sodium pyrophosphate (pyro) from either bulk soil or 1 to 2 mm aggregates (aggreg) and measured as Bradford reactive soil protein (BRSP) on a mg protein g⁻¹ soil or aggregate weight basis, for water stable aggregation (WSA) at a% of total aggregates, for the infiltration rate (IR) on a ml per hr basis for the first (1) and second (2) inches of water, and for the water holding capacity at saturation (WHC_s) for the first two inches of water.

	BRSP pyro soil	BRSP total soil	BRSP cit aggreg	BRSP pyro aggreg	BRSP total aggreg	WSA
BRSP cit soil	0.8684**	0.994***	0.9371***	0.5894*	0.9350***	0.7702**
BRSP pyro soil	1.000	0.8855***	0.8740***	0.7811**	0.8895***	0.7700**
BRSP total soil	0.8855***	1.0000	0.9409***	0.6083*	0.9400***	0.7700**
BRSP cit aggreg	0.8740**	0.9409***	1.0000	0.6362*	0.9983***	0.7893**
BRSP pyro aggreg	0.7811**	0.6083*	0.6362*	1.0000	0.6799*	0.7045*
BRSP total aggreg	0.8895***	0.9400***	0.9983***	0.6799*	1.0000	0.8032**

^aCorrelations are significant at the <0.05 (*), <0.01 (**), or <0.001 (***) levels.

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