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Evaluation of β -Glucosidase Activity as a Soil Quality Indicator for the Soil Management Assessment Framework

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The Soil Management Assessment Framework (SMAF) was developed to assess conservation effects on soil, and uses multiple soil quality indicator measurements to compare soil functioning. Our objective was to develop a SMAF-compatible scoring equation for soil β -glucosidase (BG) activity using published data sets representing different soils and management. The resulting equation was an S-shaped curve: $y = a/[1 + b\exp(-cx)]$, where x is the measured BG activity (mg p -nitrophenol [PNP] released kg^{-1} soil h^{-1}), a and b are constants, and c is a factor modified by soil classification, texture, and climate. Data from a study conducted near Mandan, ND were used to test the model for sensitivity to crop management systems. Soil organic C (SOC) content at the site measured 247 to 687 g kg^{-1} , while BG activity ranged from 33 to 675 $\text{mg kg}^{-1} \text{h}^{-1}$. Using SMAF, SOC indicator scores ranged from 0.25 to 0.73, while BG activity scores varied from 0.17 to 0.93. As the work progressed, it became apparent that when BG activity values were normalized to the SOC content, the resulting ratio could indicate C sequestration trends, with ratios of 10 to 17 $\text{g PNP kg}^{-1} \text{SOC h}^{-1}$ reflective of systems in equilibrium. Ratios >17 were mostly from recently altered management systems with SOC contents trending upward, while ratios <10 were generally from soils that were expected to continue to lose soil C. The application of a sensitive C cycling enzyme activity such as BG should improve the SMAF soil quality assessments for soil functions where soil metabolic activity or C-cycle enzyme activity play a role.

Abbreviations: BG, β -glucosidase; CEAP, Conservation Effects Assessment Project; CRP, Conservation Reserve Program; MBC, microbial biomass carbon; PNP, p -nitrophenol; SMAF, Soil Management Assessment Framework; SOC, soil organic carbon; SOM, soil organic matter.

Today our soils are expected to produce food, fiber, feed, and fuel while maintaining and protecting our air, water, and soil resources. Soil and water quality are inherently linked (National Research Council, 1993). Assessing soil quality on a watershed scale may be the link needed to demonstrate how agricultural management practices impact water quality in streams (National Research Council, 1993; Karlen et al., 2008). While the concept of a performance-based rating for soil is not new, it has most often been related to crop productivity (Cambardella et al., 2004). The soil quality concept has been broadened to include the soil's impact on the environment. It has been suggested that enhancing soil quality is critical for maintaining and improving water quality (Kennedy and Papendick, 1995). There continues to be a number of soil quality issues in the United States, including continued high rates of erosion, reductions in soil fertility and production, and exposure to chemical and heavy metal pollution (Karlen and Stott, 1994; Karlen et al., 2001; Doran, 2002; Andrews et al., 2004).

In 2003, the Conservation Effects Assessment Project (CEAP) was initiated to provide a scientific basis for a national assessment of conservation practices by the NRCS (Richardson et al., 2008). Initially, the primary thrust of CEAP was to assess the impact on water quality of implementing conservation practices within agricultural watersheds. In 2006, a study was initiated to assess the effects of these same conservation practices on soil quality within the USDA-ARS's 14 CEAP experimental watersheds.

For cropland, soil quality for a specific site can be affected by the interaction of many factors including tillage, crop rotation, and other management factors, as

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well as climate and soil type. Assessment tools are needed to evaluate the impact of management systems on critical soil functions related to soil quality, including nutrient cycling and water partitioning. Such tools need to be flexible with regard to the selection of soil functions to be assessed and indicators of the selected critical functions to be measured to ensure that the assessments are suitable for the management goals of interest. They also need to be sensitive to management changes, preferably within a year or two after implementation. For CEAP, the tool selected to help assess the impacts of management on soil was the SMAF (Andrews et al., 2004). A peer-reviewed NRCS report (Potter et al., 2006) described the use of the interpretation step of the SMAF for the national modeling portion of the CEAP project.

The SMAF provides site-specific interpretations for soil quality indicator results, and a beta version is now available (soil-quality.org/tools/smaf_intro.html; verified 19 Sept. 2009) or from us. The SMAF uses measured soil indicator data to assess management effects on soil functions using a three-step process that includes indicator selection, indicator interpretation, and integration into an index. Indicators used in the SMAF include soil physical, chemical, and biological characteristics that are management sensitive and therefore dynamic. Currently, the SMAF includes 10 indicators with scoring curves consisting of interpretation algorithms (some including logic functions). They are: aggregate stability, plant-available water holding capacity, bulk density, electrical conductivity, pH, Na adsorption ratio, soil P, SOC, microbial biomass C (MBC), and potentially mineralizable N (PMN). Interpretation algorithms that have been recently developed are water-filled pore space and Mehlich-extractable K (Wienhold et al., 2009). Nevertheless, the SMAF uses soil taxonomy as a foundation for assessment, allowing for the modification of many of the scoring indicator values based on soil suborder characteristics, and providing a contextual basis for indicator interpretation. Soil quality and its assessment is soil and site specific and depends on a variety of factors, including inherent soil characteristics, environmental influences such as climate, and human values such as intended land use, management goals, and environmental protection, all of which are considered (and can be manipulated by the user) in the SMAF tool.

Other soil quality assessment tools exist. The Soil Conditioning Index (SCI), which has been adopted by the NRCS, estimates the effects of crop management on SOC levels (NRCS, 2002). The SCI was designed to determine if SOC levels would increase, decrease, or remain stable under a given management system. When the SCI was compared with the SMAF SOC indicator (a more direct comparison than using the full suite of SMAF indicators), the SMAF SOC was more successful in separating the tested cropping systems (Zobeck et al., 2008). The AgroEcosystem Performance Assessment Tool (AEPAT) is a research-oriented index methodology that ranks agroecosystem performance among management practices for chosen functions and indicators (Liebig et al., 2004). There is general agreement between the AEPAT and the SMAF (Wienhold et al., 2006); however, the input requirements and intended uses of the two

tools are different, making a direct comparison difficult. There is also a farmer-oriented assessment tool: the Cornell Soil Health Test (Idowu et al., 2008). This assessment is narrowly tailored to assess only the crop production function of the soil.

Karlen et al. (2008) compared the SMAF with the SCI, the soil tillage intensity rating (STIR), and the N-leaching index that have been incorporated in the Revised Universal Soil Loss Equation, Version 2 (RUSLE2). The RUSLE2 estimates soil loss due to rill and interrill erosion caused by rainfall on cropland (USDA-ARS, 2005; Lightle, 2007). The STIR was developed to replace the soil disturbance rating used in the original SCI, and can function as a stand-alone rating to evaluate tillage and planting effects on factors other than ground cover and surface residue distribution. The N-leaching index is computed based on the soil hydrologic group and annual and winter rainfall (Pierce et al., 1991) and can be used to compare the potential for N leaching among various management systems. The SMAF soil quality index was significantly negatively correlated with soil loss (-0.46 , $P < 0.0001$) as calculated by RUSLE2 and the N-leaching index (-0.51 , $P < 0.0001$), significantly positively correlated with the SCI (0.29 , $P < 0.0001$), and showed no correlation with the STIR rating (0.08). The SMAF appeared to provide more information about the effects of management practices within the watershed examined (Karlen et al., 2008).

To increase the sensitivity of the SMAF to management impacts, the development of additional indicator scoring curves has been encouraged. Scoring curve development is a multistep process starting with the identification of an indicator, determining the type of relationship between the indicator and a specific soil function, identifying an appropriate mathematical equation(s) describing that relationship, and validating the scoring curve (Andrews et al., 2004; Wienhold et al., 2009). There are basically three types of relationships between scoring curves and soil function: (i) more is better (upper asymptotic sigmoid curve), (ii) less is better (lower asymptotic sigmoid curve), and (iii) having a midpoint optimum (Gaussian function) (Karlen and Stott, 1994; Andrews et al., 2004).

Currently the SMAF includes two microbial or biochemical indicators, PMN and MBC, both being represented by more-is-better curves (Andrews et al., 2004). The inclusion of PMN is based on its relation to nutrient availability and a theorized relationship between microbial activity and plant productivity (Hendrix et al., 1990; Sparling, 1997), while MCB is included based on its role as a readily available pool of C and N and an association with improved soil structural functioning (Elliott and Coleman, 1988; Hendrix et al., 1990). Neither of these parameters address microbial activity or the potential metabolic activity of the soil.

Soil enzymes mediate and catalyze a number of soil biochemical and nutrient-cycling processes involved in soil functions and are considered to be the most likely candidates for determining early responses to changes in soil management (Dick et al., 1996). Enzyme activities will increase as a response to increases in soil microbial populations and the resulting increase in enzyme synthesis, as microorganisms are the major source of enzymes in soil

(Tabatabai, 1994). Despite the short life-cycle of microorganisms, however, most enzymes continue to contribute to the metabolic capacity of the soil. Enzymes can be excreted by living cells or released by disintegrating cells to become free enzymes. In soil, free enzymes become adsorbed on organic and mineral constituents or complexed with humic substances or both. When soil is sampled to compare management impacts on soil functions, it provides a snapshot of the soil ecosystem as it exists at the time of sampling. Thus, while changes in enzymatic activities may be correlated with simultaneous changes in the soil microbial population, the shifts in activities are just as apt to reflect long-term fluctuations in microbial biomass and not necessarily the current population level.

Of the enzymes for which assay procedures exist for the soil environment, BG is one of the immobilized enzymes most often reported in the literature and has been suggested as an indicator of management effects (Bandick and Dick, 1999). The BG enzyme (EC 3.2.1.21; obsolete name: cellobiase) plays a major role in the degradation of soil organic matter and plant residues. It catalyzes the hydrolysis of β -D-glucopyranosides in the final, rate-limiting step in the degradation of cellulose, the most abundant polysaccharide in the earth, providing simple sugars for the soil microbial population. While no single enzyme activity can provide a full picture of soil metabolic functioning, BG has been shown to be sensitive to changes in soil and residue management as well as an early indicator of changes in SOC before these changes are reflected in total or organic C analyses (Miller and Dick, 1995; Deng and Tabatabai, 1996; Aon and Colaneri, 2001; Turner et al., 2002; Acosta-Martínez et al., 2003a; de la Horra et al., 2003; Roldán et al., 2005; Green et al., 2007). An increasing BG activity, which usually increases with increasing soil microbial biomass, would reflect on a soil's ability to break down plant residues and improve the availability of nutrients for subsequent crops. Within the structure of the SMAF, relative BG activity would relate to the following soil functions due to its importance in C cycling and providing simple sugars to support a diverse microbial population: nutrient cycling (for plant growth), biodiversity and habitat (within the soil and the plants and animals sustained by the soil), filtering and buffering (excess nutrients and toxic chemicals from the water), and physical stability and support (soil structure).

There has been some comparison of cropland activities to those found in native ecosystems and long-term (>10-yr) no-till, Conservation Reserve Program (CRP) land, and pastures. The activities of BG were generally two to five times greater in these systems than the typical conventional cropping systems, and one and a half to three times higher than conservation systems, with greater disparity occurring in the temperate climates (Bandick and Dick, 1999; Acosta-Martínez et al., 2003a, 2007b; de la Horra et al., 2003). Elevated levels of BG activity can be found within 1 to 3 yr of changing management from conventional to conservation or no-till practices (Roldán et al., 2005). Within the same soil type, BG activity is higher in more intensive cropping systems (Bandick and Dick, 1999; Acosta-Martínez et al., 2003a, 2007b), with systems that include a fallow period having among the lowest activity rates.

Large additions of organic material can initially result in two- to fourfold increases in BG activity, but subsequent additions do not sustain the high levels of activity (Martens et al., 1992). Soil structural improvement, as measured by decreased bulk density and increased infiltration rates, have resulted in enhanced levels of BG activity with time when compared with preamendment activities. In general, improved soil aggregation resulted in higher BG activity levels (Roldán et al., 2005).

For the CEAP soil quality assessment, BG activity will be measured at the 14 experimental watershed sites. To increase the usefulness of BG activity as a soil quality indicator for CEAP, a SMAF indicator curve is needed for this important biochemical measurement. The assessment of soil enzyme activity, including BG activity, is simple and has relatively low labor costs when compared with other soil biochemical analyses (Ndiaye et al., 2000; Acosta-Martínez et al., 2007a). Our objectives were to develop a set of algorithms to describe BG activity, as an indicator of soil metabolic activity, for use as a SMAF indicator and to provide an initial validation using published literature. An additional objective was assess BG activities in the same soil under different no-till cropping systems just before and 1 yr after implementation. The field experiment hypotheses were that the SMAF BG activity indicator would be sensitive to changes due to the different cropping systems.

MATERIALS AND METHODS

Soil Management Assessment Framework Curve Development for β -Glucosidase Activity

The basic procedure for SMAF curve development has been published previously (Andrews et al., 2004; Wienhold et al., 2009). A SMAF scoring curve consists of an algorithm or a logic statement (if, then, else) with multiple algorithms. The algorithms should be quantitative relationships between measured values of the indicator and a normalized, unitless score that represents the indicator's performance as part of a soil function(s). While it is assumed that the general shape of the relationship between a soil indicator and a soil function holds across ecosystems, the range for an indicator will vary among ecosystems. The variation from system to system results from differences in site-specific factors such as climate or inherent soil properties. Curve-fitting software is used to describe the general shape and help identify and calibrate site-specific coefficients to provide the appropriate contextual variation in the expected range.

β -Glucosidase activity has not been studied in relation to any of the soil functions described in the SMAF (Andrews et al., 2004), so we chose to relate BG activity to SOC because SOC content is a component of many of the soil functions outlined in the SMAF. The initial relationship between BG activity and soil C was determined by graphing the data and determining the general shape of the relationship (i.e., more is better, less is better, or local optimum). Once the relationship shape was defined, curve-fitting software (CurveExpert, version 1.3 shareware, available at curveexpert.webhop.net/; verified 19 Sept. 2009) was used to develop an algorithm that describes the relationship. CurveExpert compares the fit of the data to a library of available models, selects the model having the lowest root mean square error, and provides coefficient estimates for the model exhibiting the best fit.

Table 1. The soil organic matter factor classes used to determine the soil organic C indicator index value, as used in the Soil Management Assessment Framework (Andrews et al., 2004), presented in the first Conservation Effects Assessment Project report (Potter et al., 2006), and proposed for use in calculating the β -glucosidase indicator index value. Class 1 represents the suborders that tend to have the highest potential for sequestering SOC, while Class 4 has the lowest.

Class 1 (High)	Class 2	Class 3	Class 4 (Low)
Andisols	Andisols	Alfisols	Andisols
Aquands	Udands	Aqualfs†	Torrands
Gelisols	Ustands	Boralfs	Xerands
Histels	Inceptisols	Cryalfs	Aridisols
Turbels	Aquepts	Udalfst†	Argids
Histosols	Mollisols	Ustalfst†	Calcids
Fibrists	Albolls	Xeralfst	Cambids
Folists	Aquollst†	Andisols	Cryids
Hemists	Borolls	Cryands	Durids
Saprists	Cryolls	Vitrands	Gypsid
Oxisols	Rendolls	Entisols	Orthids
Aquoxs	Udolls†	Aquents	Salids
Spodosols	Ustolls†	Gelisols	Entisols
Aquods	Xerolls†	Orthels	Arents
	Oxisols	Inceptisols	Fluvents
	Udox	Andepts	Orthents†
	Spodosols	Anthrepts	Psamments
	Humods	Spodosols	Xerents
	Ultisols	Cryods	Inceptisols
	Aquults	Vertisols	Cryepts
	Humults†	Cryerts	Ochrepts
	Vertisols		Tropepts
	Aquertst		Udepts
	Xererts		Umbrepts
			Ustepts
			Xerepts
			Oxisols
			Orthox
			Perox
			Torroxx
			Ustox†
			Spodosols
			Orthods
			Ultisols
			Udults
			Ustults
			Xerults
			Vertisols
			Torrerts
			Usterts
			Uderts

† Soil suborders used in curve development.

‡ Soil suborders used in curve validation.

Once the basic algorithm has been developed, the impact of inherent soil differences must be considered. The inherent differences include characteristics such as soil classification, texture, and climate. These inherent characteristics might cause a shift in the expected range of the BG activity indicator or in the relationship between BG and the selected soil function (SOC). From the preponderance of the literature, we determined that the

factors most likely to affect the expected range of BG were inherent soil organic matter (determined using U.S. soil taxonomic suborders), soil texture, and climate (Tables 1–3). The determination of how these factors affect the expected range occurred during curve calibration.

Data Set Selection

The scientific literature was searched for a variety of data sets containing measurements of BG activity. Several criteria were used to determine if a data set should be used for the development or validation process. These included: (i) BG activity was reported as the release of PNP using methods based on Eivazi and Tabatabai (1988); (ii) the soil series or suborder was given; (iii) soil texture was determined; (iv) SOC content was measured or soil organic matter (SOM) content was reported; (v) the experimental location was specified or climatic information was included; and (vi) there were multiple treatments such as crop rotation, tillage practices, or site comparisons resulting in a range of BG activities. Data sets meeting these criteria were divided for use in development (D) or validation (V) to maximize the diversity within each group and to maximize overlap, particularly in the soil and climate conditions, between the two groups. The sets used in the developmental stage were chosen so that each grouping of soil orders, soil textures, and climates were represented. The remainder of the data sets were used for validation.

Field Experiment

The Mandan validation experiment was a subset of a larger project described by Tanaka et al. (2007). The crop \times crop-residue matrix experiment was located at the Area IV Soil Conservation District, USDA-ARS Northern Great Plains Research Laboratory's research farm approximately 7 km southwest of Mandan, ND (46°46'22" N, 100°57'9" W). The experiment was comprised of two sites (north and south), with soils classified as Temvik–Wilton silt loams (fine-silty, mixed, superactive, frigid Typic and Pachic Haplustolls). Before initiating the experiment, the sites were seeded to an oilseed sunflower (*Helianthus annuus* L.)–spring wheat (*Triticum aestivum* L.)–spring wheat crop sequence beginning in 1999 for the north site and 2000 for the south site. The inclusion of two sites for the experiment provided four site years of data. During the first year of the crop \times crop-residue matrix experiment (2002 and 2003 for the north and south sites, respectively), 10 crops were no-till seeded into spring wheat stubble: buckwheat (*Fagopyrum esculentum* Moench), canola (*Brassica napus* L.), chickpea (*Cicer arietinum* L.), corn (*Zea mays* L.), dry pea (*Pisum sativum* L.), grain sorghum [*Sorghum bicolor* (L.) Moench], lentil (*Lens culinaris* Medik.), oilseed sunflower, proso millet (*Panicum miliaceum* L.), and hard red spring wheat. In the second year, the same crops were no-till seeded perpendicular over the residue of the previous year's crop, resulting in a 10 \times 10 matrix with 100 treatment combinations. Treatments were replicated four times each year following a strip-block design. Cultural practices used during the experiment were similar to that of local no-till producers. Planting and harvesting operations were conducted based on locally optimal time periods for each crop. Additional details regarding management of the applied treatments can be found in Tanaka et al. (2007).

Soil samples were collected in the spring before planting and taken only from plots that were to be planted to corn in the second year. The first sampling occurred the spring after the preliminary 3-yr rotation of sunflower–spring wheat–spring wheat, while a second sampling

Table 2. The texture class used to determine the soil organic C (SOC) indicator index value, as presented in the Soil Management Assessment Framework (Andrews et al., 2004; Potter et al., 2006), and proposed for use in calculating the β -glucosidase indicator index value. Class 1 represents the texture groups that have the lowest intrinsic potential for sequestering SOC, while Class 5 has the highest. Texture groups are based on the work of Quisenberry et al. (1993).

Class	Texture group
1	sand loamy sand
2	sandy loam (with <8% clay) sandy loam (with clay >8%) sandy clay loam loam
3	silt loam silt
4	sandy clay clay loam silty clay loam silty clay
5	clay (<60%) clay (>60%)

occurred the spring after the initial 10 crops were harvested. Samples were collected from three replicates at each site where inherent soil conditions were most uniform based on inspection of a soil survey map (Replicates 1, 2, and 4 for the north site and Replicates 5, 7, and 8 for the south site). In each plot, 10 soil cores were collected from the 0- to 5-cm depth, bulked, stored in a plastic bag, and kept in cold storage at 4°C until processing. Samples were air dried and ground to pass a 2.0-mm sieve. Identifiable root material was removed during sieving.

The total SOC content was determined by dry combustion using a LECO CHN 2000 Analyzer (LECO Corp., St. Joseph, MI). There were no detectable carbonates, so the total C content was considered equal to the organic C content. The BG activity was determined using the method of Eivazi and Tabatabai (1988) and expressed as milligrams PNP released per kilogram of soil per hour of incubation.

RESULTS AND DISCUSSION

Assumptions for Algorithm Development

The activity of BG has not been studied in relation to the soil functions designated in the SMAF. Since BG activity plays a role in plant residue decomposition and SOC cycling, however, we chose SOC as a comparative factor since it is a component of several of the SMAF soil functions; nonetheless, this is an imperfect comparison. The BG activity is often significantly correlated to SOC within a given study, and accounts for 53 to 100% of the variation (r^2) in the observed BG activity (Bergstrom et al., 1998; Mullen et al., 1998; Bandick and Dick, 1999; Dumontet et al., 2001; Taylor et al., 2002; Acosta-Martínez et al., 2003b, 2004, 2007b; de la Horra et al., 2003; Roldán et al., 2005; Leon et al., 2006). There are many exceptions, however, with those studies showing no significant correlation (r^2 from 0.13 to 0.42) between SOC and BG activity within a soil type (Eivazi and Tabatabai, 1988; Ajwa et al., 1999; Bandick and Dick, 1999; Taylor et al., 2002; Acosta-Martínez et al., 2003a; Dodor and Tabatabai, 2005). Another exception is for Oxisols, either with native Cerrado vegetation or planted to corn (Green et al., 2007;

Table 3. The climate classes used to determine the soil organic C indicator index value, as presented in the Soil Management Assessment Framework (Andrews et al., 2004; Potter et al., 2006), and proposed for use in calculating the β -glucosidase indicator index value.

Class	Climate designation	Growing degree days	Precipitation
			mm
1	high/high	≥ 170	≥ 550
2	high/low	≥ 170	<550
3	low/high	<170	≥ 550
4	low/low	<170	<550

Green and Stott, unpublished data, 2008), where BG activities were significantly correlated with P content, the limiting nutrient in that region, but were poorly correlated with SOC content. Also, large additions of various organic amendments to soil have been shown to result in temporary two- to four-fold increases in BG activity, with only a minor shift in SOC content (Martens et al., 1992). Furthermore, those increases peaked and were not sustained, despite additional organic inputs.

In the SMAF, the MBC indicator is affected by season, while SOC is not. There appears to be little seasonal impact on BG activity (Ajwa et al., 1999; Bandick and Dick, 1999; Acosta-Martínez et al., 2004; Bastida et al., 2006), therefore seasonal affects were not considered to be important in developing these algorithms. Annual fluctuations in temperature and rainfall, which impact soil microbial biomass and plant growth, can impact BG levels within an ecosystem, however, and so should be considered when interpreting scores (Acosta-Martínez et al., 2004; Dodor and Tabatabai, 2005).

Since BG activity is generally related to SOC contents within a given study, especially in ecosystems where an equilibrium has been reached, e.g., native vegetation, we assumed that the same groupings of soils, textures, and climates used for the SOC indicator within the SMAF model (Andrews et al., 2004) could be applied to the BG indicator. Soil suborder (Table 1) groupings were based on the potential of an individual suborder to sequester C, and are referred to as SOM factor classes. Soil texture was grouped into five classes (Table 2) based on the work of Quisenberry et al. (1993). Climate designations were based on rainfall and temperature (Table 3). If climate data were missing from a data set within the United States, the Major Land Resource Area (NRCS, 2006) was used to assign a climate class.

Relationship between Soil Organic Carbon and β -Glucosidase Activity

When BG activity was compared collectively to SOC contents from all of the development and validation data set studies (Table 4), there was a poor correlation (Fig. 1). If it is assumed, however, that ecosystems with native vegetation, long-term pastures, and long-term CRP lands are relatively stable (no shift in SOC content with time, and plant and soil microbial biomass are at stable, sustainable levels), then it might be assumed that SOC contents and BG activity levels would be in equilibrium. When BG activity was normalized relative to SOC ($BG_n = \text{BG activity} / \text{SOC content}$), such ecosystems had BG_n values ranging from 10

Table 4. Summary of data sets used in the development and validation of the β -glucosidase indicator equation c-factor values (site-specific coefficients in algorithms) and scoring curves for use in the Soil Management Assessment Framework.

Reference	Data set ID†	Location	Soil identification	Crop and soil management	Range of values‡		
					SOC	BG	BG _n
					g kg ⁻¹	mg kg ⁻¹ h ⁻¹	g kg SOC ⁻¹ h ⁻¹
Acosta-Martínez et al. (2003b)	D01	Lubbock, TX	Paleustoll loam	cotton, wheat, sorghum in various rotations and tillage regimes	9.0–12.2	43–156	4.8–12.8
Dodor and Tabatabai (2005), Moore et al. (2000)	D02	Iowa	Hapludoll loam	corn, soybean, oat, meadow; conventional tillage	17.3–25.5	87–277	4.1–13.6
Eivazi and Tabatabai (1988)	D03	Iowa	Endoaquert silty clay, Hapludoll clay loam, Calciaquoll clay loam, Endoaquert silty clay loam, Endoaquoll silty clay loam	corn–soybean rotation; conventional tillage (original method development)	25.4–54.5	72–295	5.7–6.8
Dodor and Tabatabai (2005), Moore et al. (2000)	D04	Iowa	Hapludoll clay loam	corn, soybean, oat, meadow; various rotations; two fertilizer rates	22.7–44.3	89–272	3.2–8.7
Green et al. (2007)	D05	Sete Lagoas, MG, Brazil	Paleudalf fine sandy loam	corn–common bean rotation with varying tillage regimes; native vegetation	27.4–41.6	55–100	1.7–3.5
Roldán et al. (2005)	D06	Mexico	Aquert clay	native vegetation, long-term moldboard plow, recent conservation tillage	7.4–12.6	38–205	5.1–16.3
Acosta-Martínez et al. (2003b)	D07	Lubbock, TX	Paleustalf sandy loam, Paleustalf sandy clay loam	cotton, wheat, peanut, sorghum in various rotations and tillage regimes	1.4–9.6	11–131	4.0–42.0
Martens et al. (1992)	D08	Riverside, CA	Durixeralf sandy clay loam	fallow; heavy manure or plant residue additions	4.1–7.4	162–345	27.7–63.0
Acosta-Martínez et al. (2004)	D09	Gaines Co., TX	Paleustalf sand	cotton and peanut in varying 3-yr rotations; deep tilled; irrigation implemented	1.4–2.1	12.5–42	8.2–24.9
Green and Stott (unpublished data, 2008)	D10	Brazil	Ustox sandy clay, Ustox clay (<60% clay)	native Cerrado vegetation, several sites	19.2–31.6	24–70	0.8–2.3
Bandick and Dick (1999)	D11	Aurora, OR	Argixeroll silt loam	vegetables with legume or cereal cover crops or fallow; fescue grass seed production	13.7–16.5	53–81	3.4–5.8
Bandick and Dick (1999)	D12	Pendleton, OR	Haploxeroll silt loam	long-term winter wheat–summer fallow; fertility treatments; native pasture	9.0–15.2	42–202	4.3–13.3
Dumontet et al. (2001)	V01	Italy	Ustorthent clay loam	vetch, oat, wheat and fallow; conventional or reduced tillage	9.7–12.4	221–380	21.0–30.6
Acosta-Martínez et al. (2007b)	V02	Akron, CO	Paleustoll loam	low-, medium-, high-intensity cropping, pasture	5.5–10.1	42–156	7.1–15.5
de la Horra et al. (2003)	V03	Cordoba, Argentina	Argiudoll silty clay loam	native pasture; conventional and no-till corn	10.1–27.0	121–295	10.9–14.8
Acosta-Martínez et al. (2003a)	V04	Texas	Paleustalf loamy sand	continuous cotton, conventional tillage; cotton–wheat, conservation tillage; Conservation Reserve Program (CRP)	3.4–4.0	22–49	6.8–16.5
Acosta-Martínez et al. (2003a)	V05	Texas	Paleustalf sandy loam	continuous cotton, conventional tillage; cotton–wheat, conservation tillage; CRP, native rangeland	3.1–14.1	23–124	6.4–13.5
Acosta-Martínez et al. (2003a)	V06	Texas	Paleustoll sandy loam, Paleustoll sandy clay loam	continuous cotton, conventional tillage; cotton–wheat, conservation tillage; sunflower, CRP, native rangeland	3.2–18.7	31–231	8.2–21.5
Knight and Dick (2004)	V07	eastern Oregon	Xeroll silt loam	managed (wheat) vs. unmanaged (cemetery)	9.4–13.9	135–190	13.7–14.4
Knight and Dick (2004)	V08	western Oregon	Albaqualf silt loam	managed (annual rye grass) vs. unmanaged land (native grassland)	21–23	80–111	3.8–4.8

Table 4. (cont.)

Reference	Data set ID†	Location	Soil identification	Crop and soil management	Range of values‡		
					SOC	BG	BG _n
					g kg ⁻¹	mg kg ⁻¹ h ⁻¹	g kg SOC ⁻¹ h ⁻¹
Knight and Dick (2004)	V09	western Oregon	Palehumult silty clay loam	managed (Christmas tree) vs. unmanaged (Douglas-fir forest)	31–38	80–134	2.1–4.3
Bergstrom et al. (1998)	V10	Ontario	Hapludalf with various textures	corn–soybean–winter wheat; no-till vs. conventional tillage	19–23	109–139	4.7–7.0
Bergstrom et al. (1998)	V11	Ontario	Aquoll sandy loam	corn–soybean; no-till vs. conventional tillage	29–38	167–209	5.5–5.8
Bergstrom et al. (1998)	V12	Ontario	Hapludalf sandy loam, Hapludalf loam	corn, soybean, wheat; no-till vs. conventional tillage	17–37	92–209	2.6–6.1
Ajwa et al. (1999)	V13	Konza Prairie, Kansas	Argiustoll silty clay loam	native grass; burned vs. unburned; fertilized vs. unfertilized	30–35	21–41	0.7–1.3
Mullen et al. (1998)	V14	western Tennessee	Hapludalf silt loam	corn, no-till; fertilized vs. unfertilized; no cover crop vs. hairy vetch	10.0–16.9	58–172	5.7–11.7

† D, development data set; V, validation data set.

‡ SOC, soil organic C content; BG, β -glucosidase activity in mg *p*-nitrophenol released kg⁻¹ soil h⁻¹ incubation; BG_n, ratio of BG activity to SOC content.

to 17 g PNP released kg⁻¹ SOC h⁻¹ incubation (Tables 4 and 5) and appeared to be independent of soil type. Within the development and validation data sets (Table 4), there were 53 data points from 11 of the data sets that fell within this stable BG_n range, and they exhibited a significantly high correlation between SOC content and BG activity (Fig. 1) with an r^2 of 0.95. When developing the family of curves for the BG indicator scores, it was assumed that for data points in the development sets that were in the stable BG_n range, the BG indicator score would be about equal to the SOC indicator score as calculated by the SMAF (Andrews et al., 2004). Examples of these stable systems included native vegetation (Acosta-Martínez et al., 2003a; de la Horra et al., 2003; Roldán et al., 2005), long-term pasture or legume crop (Bandick and Dick, 1999; de la Horra et al., 2003; Acosta-Martínez et al., 2004, 2007b; Knight and Dick, 2004), and long-term conservation tillage or no-till management (Acosta-Martínez et al., 2003a,b). Systems that had ratios below this range (<10 g PNP released kg⁻¹ SOC h⁻¹ incubation) were generally considered to have a continual loss of SOC. These points comprised 67% of the 226 data points used in this study, with an r^2 of 0.67. The greater spread among the low-range data points was probably due to multiple factors other than SOC content limiting the BG activity. Systems that had moved to no-till or other conservation practices had either started or were anticipated to start accumulating SOC, based on earlier studies, and had increasing soil microbial activity and a high

(>17 g PNP released kg⁻¹ SOC h⁻¹ incubation) BG_n ratio (Martens et al., 1992; Dumontet et al., 2001; Acosta-Martínez et al., 2003a,b, 2004). Twenty-one data points fell in the high range (BG_n > 17 g PNP released kg⁻¹ SOC h⁻¹ incubation), with an r^2 of 0.87.

Curve Development

For developing the family of curves, 12 data sets were culled from eight published and one unpublished study (Table 4). Those data sets provided a range of soil orders, soil textures, climates, and management practices (Table 4). Two studies that included multiple sites with differing soil orders or textures were divided

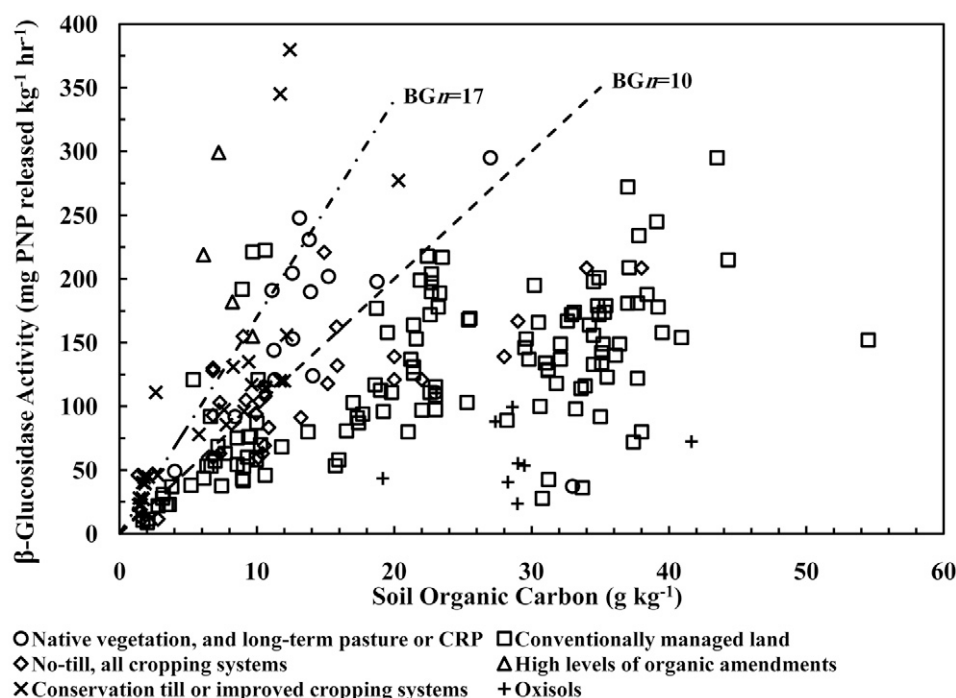


Fig. 1. Soil organic C (SOC) vs. β -glucosidase (BG) activity for the development and validation data sets used to determine site-specific coefficient (c factor) values for use in Eq. [1]. The data represent a broad range of soil and climatic conditions (Table 4). Lines represent normalized 8 pt-glucosidase activities (BG_n = BG activity/SOC content of 10 and 17 g *p*-nitrophenol [PNP] released kg⁻¹ SOC h⁻¹ incubation).

Table 5. Key endpoint samples from the data sets in Table 4 used in developing the Soil Management Assessment Framework indicator scoring curves for β -glucosidase activity.

Data set ID†	Points in data set‡	Class§	Management¶	SOC score#	BG activity†† mg kg ⁻¹ h ⁻¹	BC _n ‡‡ g kg ⁻¹ h ⁻¹	Target range§§	Final BG score¶¶
D01	13	2-2-2	continuous cotton, conventional tillage, irrigated	0.11	43	4.8	0.03–0.05	0.04
			wheat–cotton, reduced tillage, irrigated	0.19	156	12.8	0.15–0.25	0.19
			cotton with rye cover crop, no-till, irrigated	0.14	116	11.0	0.10–0.16	0.11
D02	26	2-2-3	corn–soybean, conventional tillage, 0 kg N	0.38	177	9.5	0.23–0.36	0.21
			corn–corn–oat–alfalfa, conventional tillage, 81.6 kg N	0.46	277	13.6	0.39–0.62	0.52
			corn–corn–oat–alfalfa, conventional tillage, 0 kg N	0.57	204	9.0	0.32–0.51	0.28
D03	5	2-4-3	corn–soybean, original methods development, Endoaquert silty clay	0.94	295	6.8	0.40–0.64	0.51
D04	44	2-4-3	continuous corn, no N fertilizer, Year 1	0.61	89	3.2	0.12–0.19	0.06
			corn–oat–meadow–meadow, no N fertilizer, Year 2	0.86	272	7.4	0.39–0.63	0.43
			corn–oat–meadow–meadow, 202 kg N ha ⁻¹ , Year 2	0.40	197	8.7	0.22–0.35	0.22
D05	4	2-4-1	corn–soybean, no-till	0.81	100	3.5	0.18–0.28	0.19
D06	5	2-5-1	native vegetation	0.16	205	16.3	0.16–0.26	0.18
			corn–common bean, moldboard plow	0.07	38	5.1	0.02–0.04	0.03
D07	17	3-2-2	continuous cotton, conventional tillage, irrigated	0.04	111	42	0.11–0.17	0.17
			wheat–cotton, reduced tillage	0.15	97	13	0.12–0.19	0.13
			cotton with wheat cover crop, no-till	0.03	46	34	0.06–0.10	0.05
			cotton with wheat cover crop, no-till, irrigated	0.04	12	4.1	0.01–0.02	0.03
D07	17	3-2-2	sorghum–cotton, reduced tillage	0.24	117	12.2	0.18–0.29	0.18
D08	5	3-2-2	additions of barley straw	0.09	345	63	0.34–0.55	0.96
D09	12	3-1-2	cotton–cotton–peanut, deep tilled, irrigated	0.03	12.5	8.0	0.02–0.03	0.03
			peanut–cotton–cotton, deep tilled, irrigated	0.03	18.5	13	0.03–0.04	0.03
			continuous peanut, deep tilled, irrigated	0.04	42	25	0.06–0.09	0.06
D10	2	4-4-1	native vegetation	1.00	24	0.8	0.05–0.08	0.06
			native vegetation	0.99	43	2.3	0.14–0.22	0.14
D11	4	2-3-3	vegetables with legume cover crop	0.22	81	4.9	0.07–0.11	0.06
			vegetables with cereal cover crop	0.16	80	5.8	0.06–0.09	0.06
			vegetables, winter fallow	0.20	53	3.4	0.04–0.07	0.04
D12	5	2-3-4	winter wheat, summer fallow, 90 kg N ha ⁻¹	0.09	46	0.03	0.02–0.04	0.03
			long-term pasture	0.17	202	13.3	0.14–0.23	0.18
V01	4	4-3-3	vetch, oat–wheat, reduced tillage	0.69	380	30.6	> 1	1.00
V02	8	2-4-4	medium-intensity cropping	0.05	88	15.5	0.05–0.08	0.06
			long-term pasture	0.10	156	15.5	0.10–0.16	0.12
			native grass	0.06	90	13.3	0.05–0.08	0.06
V03	3	2-4-3	native pasture	0.56	295	10.9	0.39–0.62	0.51
			conventional tillage, maize	0.09	121	12.0	0.06–0.10	0.09
V04	4	3-1-2	continuous cotton, dryland, conventional tillage	0.05	22	7.8	0.03–0.04	0.04
			cotton–wheat, dryland, conservation tillage	0.05	46	16.5	0.05–0.09	0.07
			Conservation Reserve Program (CRP) land	0.08	49	12.2	0.06–0.10	0.08

appropriately. The SOC contents of the data sets ranged from 1.4 to 54.5 g kg⁻¹, while BG activity ranged from 11 to 345 mg PNP released kg⁻¹ h⁻¹. An additional 14 data sets from eight publications were used for an initial validation. The SOC contents for the validation data sets ranged from 3.1 to 38.0 g kg⁻¹, while BG activity ranged from 21 to 380 mg PNP released kg⁻¹ h⁻¹, with BG_n values from 0.7 to 30.6 g PNP released kg⁻¹ SOC h⁻¹ incubation. For studies with multiple sampling depths, only surface samples were considered, and while most sampling depths were 0 to 5 cm (45% of the samples), some depths ranged to 20 cm.

The general equation for the curves was based on a more-is-better model and used the sigmoidal logistic model format:

$$\beta\text{-glucosidase score} = \frac{a}{1 + b \exp\left(\frac{-c \beta\text{-glucosidase activity}}{1000}\right)} \quad [1]$$

where the a is a constant equal to 1.01; b is a constant set to 48.4; and c is a factor that is equal to

$$c = c_1 c_2 + c_1 c_2 c_3 \quad [2]$$

where c_1 is the SOM class factor, c_2 is the texture class factor, and c_3 is the climate class factor (Tables 1–3).

A stepwise procedure was used to develop the site-specific c coefficient values, introducing only one new factor class at a time (listed in order in Tables 4 and 5). It was an iterative process, and not all points fit the model perfectly. We started with the data for semi-arid sandy soils from western Texas (Acosta-Martínez et al., 2003b), and since that was a study that compared two sites that represented two SOM classes (Table 4), it was split. The data set first considered was the Paleustoll loam (Data Set D01) that fell within the following three factor classes: SOM class 2, texture class 2, and climate class 2, with three representative points listed in Table 5. For those data

Table 5. (cont.)

Data set ID†	Points in data set‡	Class§	Management¶	SOC score#	BG activity†† mg kg ⁻¹ h ⁻¹	BG _n ‡‡ g kg ⁻¹ h ⁻¹	Target range§§	Final BG score¶¶
V05	5	3-2-2	cotton–cotton, dryland, conventional tillage	0.05	28	8.9	0.03–0.04	0.04
			wheat–cotton, dryland, conservation tillage	0.09	78	13.5	0.08–0.13	0.09
			CRP land	0.33	121	10.7	0.22–0.36	0.20
			native rangeland	0.52	124	8.8	0.29–0.46	0.20
V06	11	2-2-2	CRP land	0.16	144	12.8	0.13–0.21	0.16
			native rangeland	0.47	198	10.6	0.31–0.50	0.31
			wheat–cotton, dryland, conservation tillage	0.11	192	21.5	0.15–0.24	0.29
			sunflower, irrigated, conservation tillage	0.11	96	10.6	0.07–0.12	0.08
			native rangeland	0.16	191	17.2	0.17–0.28	0.29
V07	2	2-3-4	managed (winter wheat)	0.08	135	14.4	0.07–0.11	0.09
			unmanaged (cemetery)	0.14	190	13.7	0.12–0.20	0.16
V08	2	3-3-3	managed (annual rye grass)	0.70	80	3.8	0.17–0.27	0.08
			unmanaged (native grassland)	0.79	111	4.8	0.24–0.38	0.13
V09	2	2-4-3	managed (Christmas tree)	0.88	80	2.1	0.12–0.18	0.06
			unmanaged (Douglas-fir forest)	0.71	134	4.3	0.19–0.31	0.11
V10	4	3-4-3	corn–soybean–winter wheat, no-till	0.60	139	7.0	0.26–0.42	0.19
V11	2	2-2-3	corn–soybean, no-till	0.96	209	5.5	0.33–0.52	0.29
V12	8	3-2-3	corn–soybean–winter wheat, conventional tillage	1.00	181	4.9	0.30–0.49	0.38
			corn–soybean–winter wheat, no-till	0.99	209	6.1	0.38–0.61	0.51
			corn–soybean–winter wheat, no-till	0.96	139	5.0	0.30–0.48	0.22
			corn–soybean–winter wheat, conventional tillage	0.61	103	6.1	0.23–0.27	0.12
V13	4	2-4-3	native grassland, burned, fertilized	0.67	21	0.7	0.03–0.05	0.03
V14	12	3-3-1	corn, no-till, no fertilizer, no cover crop	0.25	58	5.7	0.09–0.14	0.12
			corn, no-till, 16 kg N ha ⁻¹ , no cover crop	0.48	92	6.6	0.20–0.31	0.30
			corn, no-till, no fertilizer, no cover crop	0.24	61	6.1	0.09–0.15	0.13
			corn, no-till, no fertilizer, vetch cover crop	0.53	172	11.7	0.39–0.62	0.86

† D, development data set; V, validation data set (see Table 4 for full identification).

‡ Number of points falling within the defined class.

§ Soil organic matter class (Table 1)–texture class (Table 2)–climate class (Table 3).

¶ Crop in bold is the crop planted when samples were taken, otherwise samples were taken while the first crop listed was planted.

Soil organic C (SOC) indicator score calculated by the Soil Management Assessment Framework (Potter et al., 2006).

†† BG, β -glucosidase activity in mg *p*-nitrophenol (PNP) released kg⁻¹ soil h⁻¹ incubation.

‡‡ BG_n, ratio of BG activity to SOC content, with final units of g PNP released kg⁻¹ SOC h⁻¹ incubation.

§§ The target range for the BG index was calculated as [(BG_n × SOC index)/either 10 or 16] with the range of 10–16 g PNP released kg⁻¹ SOC h⁻¹ representing the BG_n values where BG activity and SOC appear to be in equilibrium (Fig. 2).

¶¶ Using the endpoints and target range to set *c*-factor values (final values listed in Table 6), these are the final scores (using Eq. [1] and [2]) for the endpoints after several iterations.

points that had BG_n values that fell within the stable range, the BG indicator value for each point was assumed to be about equal to the SOC indicator value for that point. For those BG_n values falling below the stable range, e.g., those sample points where BG activity was lower per unit C than found in “stable” areas, the BG indicator values were expected to be lower than the SOC indicator values, and a target range of values was calculated based on the interpolation of the degree of difference between the actual BG_n and the stable BG_n range (10–17 g kg⁻¹ h⁻¹) and the SOC content (Table 5). Each sequential data set was selected based on introducing one new factor class at a time. There were no data sets representing SOM class 1 (Tables 1 and 4), so a value was assigned based on an interpolation from the other values in that class.

It was rare to find a data set that had BG_n values greater than the stable range. One example was D08 (Martens et al., 1992), where large amount of manures or plant residues had been added to the soil. The BG_n values at the end of the 31-mo study period

ranged from 28 to 63, therefore the highest BG activity was set as the near-maximum level of activity for that soil and climate type.

Another anomalous data set was D10, a study of native Cerrado soils, where the BG activity was highly correlated with soil P rather than SOC (Green and Stott, unpublished data, 2008). The BG activities were quite low, despite high SOC indicator values (Table 5), with BG_n values ranging from 0.8 to 2.3 g PNP released kg⁻¹ SOC h⁻¹ incubation (Table 4).

Once the *c*-factor values were set (Table 6), a family of related indicator equations and curves were generated (Fig. 2). The final results of calculating BG indicator values for the development data sets are summarized in Table 7. The correlations (*r*) between the SOC and BG activity indicator values were similar in most cases (± 0.1) to the correlations between the SOC contents and BG activities in a given study. The one exception was D04. This study (Dodor and Tabatabai, 2005) measured changes due to rotation phase (corn, soybean, oat [*Avena sativa* L.], or alfalfa [*Medicago sativa* L.] meadow in various combinations), in

Table 6. Site-specific coefficient (c-factor) values for the β -glucosidase activity indicator as modified by the soil organic matter class, based on soil orders and suborders (Table 1), the texture class (Table 2), and the climate class (Table 3). Values were determined using the data sets listed in Table 4.

Class	c factor
Soil organic matter, c_1	
1	0.9
2	2.9
3	3.8
4	5.8
Texture, c_2	
1	4.0
2	2.9
3	2.8
4	2.7
5	1.3
Climate, c_3	
1	2.10
2	0.85
3	0.70
4	0.45

which the meadow phase accrued C, but tillage in the corn phase resulted in the rapid loss of accrued C. Here a flush of C due to tilling and a spike in the microbial population occurred, with a lag before increases BG activity would be seen.

Validation with Published Data

Additional data sets meeting the same requirements as the development data sets were used for an initial validation of the proposed BG SMAF indicator. Some of the factor class combinations were the same as the development sets but from different studies, sites, and management systems, while others were unique combinations (Tables 4 and 5). Again, the correlations between

the SOC and BG activity indicator values for most of the data sets were about the same as those between the SOC contents and BG activities within a given data set (Table 7). There was one exception, V06, which had a large discrepancy between the two r values (observed vs. scored SOC and BG). This study (Acosta-Martínez et al., 2003a) compared conservation systems across four soil types, with one falling into the factor class combination 3-1-2 (V04), one into 3-2-2 (V05), and two into 2-2-2 (V06). The SOC indicator values were quite low; however, the BG activities were relatively high, with BG_n values ranging from 8.2 to 21.5 g PNP released kg^{-1} SOC h^{-1} incubation and a number of data points exceeding the stable range. The treatments varied among long-term CRP land (12 yr), native rangeland, and cropland under current management for 5 or more years. Apparently for these latter sites, the BG activities were in flux and increasing faster than the SOC contents. For all data sets, the BG indicators retained the sensitivity to changes in management seen in the original data.

Field Experiment

As part of a larger study (Tanaka et al., 2007), SOC and BG activities were measured in 10 cropping systems under no-till management in a semiarid environment (Table 8). The sites had been in a sunflower–spring wheat–spring wheat rotation before implementation. Samples were collected the spring before the experiment was implemented at each site, following spring wheat, and again the following spring after the 10 crops were harvested. Management systems were implemented at the north site 1 yr earlier than the south site, but otherwise the two sites had identical treatments. The SOC contents varied from 26.7 to 30.2 $g\ kg^{-1}$ at the north site and from 18.2 to 26.4 at the south site (Table 8). The BG activity varied from 385 to 687 mg PNP released kg^{-1} soil h^{-1}

incubation at the north site, and 248 to 531 mg PNP released kg^{-1} soil h^{-1} incubation at the south site (Table 8). There was no crop that had consistently higher or lower BG activities than the other crops. The north site overall had significantly higher SOC contents and BG activities than the south site. At both sites, the overall mean SOC contents and BG activities trended downward from the first to the second year, although individual plots might trend in either direction, depending on the crop. When BG activities were normalized to SOC, all the BG were 10 g PNP released kg^{-1} SOC h^{-1} , with many above 17. There was no distinct pattern with regard to crop type, but since all sites were in no-till, the surface layer may still be accruing C. For the north site, SOC contents explained 86% of the BG

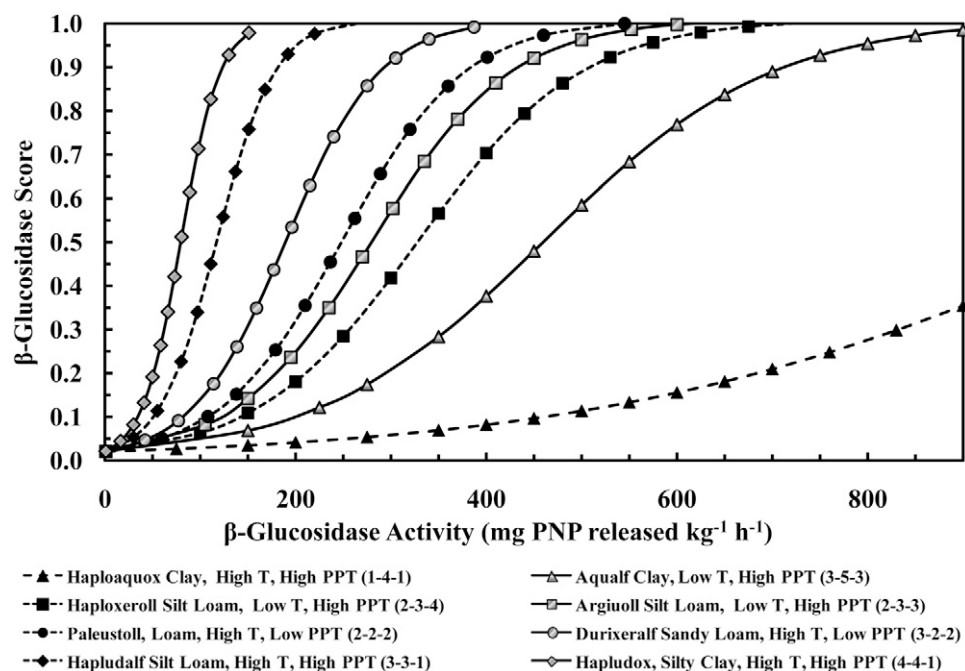


Fig. 2. Selection of curves based on Eq. [1] and [2] as well as the c-factor values listed in Table 6. The numbers in the legend represent the factor class designations in the following order: soil organic matter (based on soil taxonomy), texture, and climate (based on temperature [T] and precipitation [PPT]).

Table 7. Summary of the values used for the calculation of indicator scores using the Soil Management Assessment Framework (SMAF) for the soil organic C (SOC) content factor (Andrews et al., 2004) and Eq. [1] for the β -glucosidase activity (BG) factor.

Data set ID†	SOC vs. BG (<i>r</i>)‡	Class§	c factor for SOC scores¶	Range of SOC scores¶	Range of BG _n values#	c factor for BG scores††	Range of BG scores‡‡	SOC scores vs. BG scores (<i>r</i>)
D01	0.90	2-2-2	2.034	0.11–0.19	4.8–12.8	15.56	0.04–0.19	0.89
D02	0.40	2-2-3	1.841	0.33–0.69	4.1–13.6	14.30	0.07–0.52	0.31
D03	0.23	2-4-3	1.546	0.50–0.99	1.9–6.8	13.31	0.07–0.51	0.27
D04	0.37	2-4-3	1.546	0.40–0.95	3.2–8.7	13.31	0.06–0.43	0.98
D05	–0.27	2-4-1	1.872	0.77–0.98	1.7–3.5	24.27	0.07–0.19	–0.38
D06	0.91	2-5-1	1.783	0.07–0.16	5.1–16.3	11.69	0.03–0.18	0.90
D07	0.73	3-2-2	2.848	0.03–0.24	4.1–42.0	20.39	0.02–0.18	0.66
D08	0.03	3-2-2	2.848	0.06–0.14	27.7–63.0	20.39	0.36–0.96	0.02
D09	0.76	3-1-2	3.646	0.03–0.04	8.0–24.9	28.12	0.03–0.06	0.73
D10	–0.13	4-4-1	4.601	0.99–1.00	0.8–2.3	48.55	0.06–0.14	–0.07
D11	–0.31	2-3-3	1.620	0.16–0.22	3.4–5.8	13.80	0.04–0.06	–0.27
D12	0.90	2-3-4	1.535	0.07–0.17	4.3–13.3	11.77	0.03–0.18	0.94
V01	0.95	4-3-3	3.800	0.44–0.69	21.0–30.6	26.62	0.88–1.00	0.93
V02	0.73	2-4-4	1.744	0.05–0.10	7.1–15.5	12.19	0.03–0.12	0.80
V03	0.95	2-4-3	1.546	0.09–0.56	10.9–14.8	13.31	0.09–0.51	0.95
V04	0.37	3-1-2	3.646	0.05–0.08	6.8–16.5	28.12	0.04–0.08	0.43
V05	0.95	3-2-2	2.848	0.05–0.52	6.4–13.5	20.39	0.03–0.20	0.94
V06	0.83	2-2-2	2.034	0.04–0.47	8.0–21.5	15.56	0.03–0.43	0.68
V07	NA§§	2-3-4	1.535	0.08–0.14	13.7–14.4	11.77	0.09–0.16	NA
V08	NA	3-3-3	2.268	0.70–0.79	3.8–4.8	18.09	0.08–0.13	NA
V09	NA	2-4-3	1.546	0.71–0.88	2.1–4.3	13.31	0.06–0.11	NA
V10	–0.36	3-4-3	2.165	0.55–0.74	5.5–7.0	17.44	0.12–0.19	–0.33
V11	NA	2-3-3	1.841	0.81–0.96	5.5–5.8	14.30	0.18–0.29	NA
V12	0.58	3-2-3	2.577	0.61–1.00	2.6–6.1	18.73	0.10–0.51	0.49
V13	0.42	2-4-3	1.546	0.67–0.82	0.7–1.3	13.31	0.03–0.04	0.42
V14	0.84	3-3-1	2.745	0.25–0.67	5.7–11.7	32.98	0.12–0.86	0.83

† D, development data set; V, validation data set (see Table 4 for full identification).

‡ Coefficient of correlation of the observed values.

§ Soil organic matter class (Table 1)–texture class (Table 2)–climate class (Table 3).

¶ Calculated by SMAF (Andrews et al., 2004; Potter et al., 2006).

BG_n, BG normalized to the SOC content of the soil (BG/SOC), with final units of mg p-nitrophenol released kg^{–1} SOC h^{–1} incubation.

†† Calculated using Eq. [2] and values in Table 6.

‡‡ BG activity score as calculated by Eq. [1].

§§ NA, not applicable (data not available to make the calculation).

activity variation, while 82% of the variation in the south site values was explained, and when the sites were combined, 66% of the variation was explained by the SOC values.

Using the site-specific BG activity indicator developed here, the experimental site fell into the following factor classes: SOM class 2 ($c_1 = 2.9$), texture class 3 ($c_2 = 2.8$), and climate class 4 ($c_3 = 0.45$). The BG indicator scores ranged from 0.66 to 1.00 for the north site and 0.32 to 0.92 for the south site. When the 16 points from the stable group were added to the data found in the developmental and validation data sets in the same BG_n range (Table 8), 94% of the variation in BG activity could be explained by the SOC contents in the stable group. In the high-range group, the addition of the experimental data improved the comparison significantly ($r^2 = 0.91$).

CONCLUSIONS

Our evaluation of published data demonstrated the applicability of BG activity as a sensitive indicator of soil quality to be added to the SMAF. This will allow the SMAF to incorporate soil metabolic functioning in an overall assessment of soil quality. The BG activity has proven to be sensitive to a variety of dif-

ferent management regimes in several different climatic regions with various soil types and textures. This enzyme is an important indicator of the ability of a given soil ecosystem to degrade plant material and provide simple sugars for the microbial population. Soil enzyme activities, including BG, are generally simple, low-cost measurements to perform, especially compared with other biochemical measures.

Using data sets from a variety of native and natural ecosystems, a family of BG scoring curves to be used with the SMAF was developed. These initial curves may be modified as more data become available, especially maximum values that might be seen in native habitats. The scoring curves appropriately rank soil and crop management practices with regard to the observed BG activity in the field. Further validation of this indicator measurement, as part of a suite of indicators, will be part of the CEAP soil quality and other research experiments that utilize the SMAF to assess management impacts on soil quality. Including BG activity in the SMAF increases the sensitivity of this assessment tool, especially because measuring BG has the potential to detect changes due to shifting soil management sooner than measurements of SOC or other soil properties.

Table 8. Results from a field experiment conducted near Mandan, ND, and calculations of the Soil Management Assessment Framework scores for soil organic C (SOC) and β -glucosidase (BG) activity.

Crop†	BG‡	SOC§	SOC score¶	BG _n #	BG score††
	mg kg ⁻¹ h ⁻¹	g kg ⁻¹		g kg ⁻¹ h ⁻¹	
<u>North site, Year 1 (following spring wheat)</u>					
Buckwheat	505	28.6	0.62	17.7	0.90
Canola	600	29.0	0.63	20.7	0.97
Chickpea	564	29.4	0.65	19.2	0.95
Corn	599	29.1	0.64	20.5	0.97
Dry Pea	601	30.2	0.67	19.9	0.97
Lentil	576	26.4	0.53	21.8	0.96
Sorghum	618	32.1	0.73	19.3	0.98
Sunflower	687	29.0	0.63	23.7	1.00
Wheat	663	29.1	0.63	22.8	0.99
<u>North site, Year 2 (following crops listed below)</u>					
Buckwheat	486	27.0	0.56	18.0	0.87
Canola	426	28.9	0.63	14.7	0.76
Chickpea	466	29.3	0.64	15.9	0.84
Corn	420	30.1	0.67	13.9	0.75
Dry Pea	385	30.0	0.67	12.8	0.66
Lentil	416	26.7	0.55	15.6	0.74
Millet	440	27.1	0.56	16.2	0.79
Sorghum	479	29.2	0.64	16.4	0.86
Sunflower	587	28.5	0.61	20.6	0.96
Wheat	622	28.3	0.61	22.0	0.98
<u>South site, Year 1 (following spring wheat)</u>					
Buckwheat	279	21.8	0.36	12.8	0.36
Canola	319	20.2	0.31	15.8	0.47
Chickpea	439	25.4	0.50	17.3	0.79
Corn	384	20.2	0.31	19.1	0.66
Dry Pea	466	25.0	0.48	18.7	0.84
Lentil	438	23.1	0.41	19.0	0.79
Millet	339	22.5	0.39	15.1	0.53
Sorghum	338	19.0	0.27	17.8	0.53
Sunflower	376	23.0	0.41	16.3	0.64
Spring Wheat	531	26.4	0.53	20.1	0.92
<u>South site, Year 2 (following crops listed below)</u>					
Buckwheat	358	18.6	0.26	19.3	0.59
Canola	371	19.8	0.30	18.7	0.63
Chickpea	391	21.0	0.33	18.6	0.68
Corn	248	18.2	0.25	13.6	0.28
Dry Pea	395	21.7	0.36	18.2	0.69
Lentil	345	20.0	0.30	17.3	0.55
Millet	302	19.8	0.29	15.3	0.43
Sorghum	290	18.3	0.25	15.9	0.39
Sunflower	265	19.7	0.29	13.5	0.32
Spring Wheat	265	21.2	0.34	12.5	0.32

† Samples were collected in the spring before planting operations before (Year 1) and after (Year 2) the listed crops. At both sites, Year 1 reflected a baseline sampling that followed a 3-yr sunflower–spring wheat–spring wheat rotation. Sampling in Year 2 occurred the spring following the harvest of the 10 listed crops.

‡ β -glucosidase activity reported as mg p-nitrophenol released kg⁻¹ soil h⁻¹ incubation.

§ SOC content as g SOC kg⁻¹ soil.

¶ Calculated using the Soil Management Assessment Framework (Andrews et al., 2004).

BG normalized to the SOC content of the soil (BG/SOC), with final units of mg p-nitrophenol released kg⁻¹ SOC h⁻¹ incubation.

†† BG activity score as calculated by Eq. [1], using the values in Table 6 to calculate the c factor. For this site, $c_1 = 2.9$, $c_2 = 2.8$, $c_3 = 2.45$, and using Eq. [2], $c = 11.78$.

To fully test the concept that the BG_n ratio is a predictor of trends in C sequestration, studies that measure changes in enzyme activities for several years are needed. To test this indicator model, sites with high, near-maximum BG activities need to be sampled. Such sites would be native ecosystems or long-term pastures that have high SOC contents relative to the factor classes (Tables 1–3) representing the site.

The SMAF is a malleable model, as is the BG indicator developed here. It is hoped that others will use and modify the indicator and SMAF tool as necessary when new data become available. In the meantime, this indicator will allow soil enzymatic activity involved in C cycling to be assessed when using the SMAF ecosystem function analysis.

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