



## Image Analysis for Rapid Assessment and Quality-Based Sorting of Corn Stover

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Imaging in the visible spectrum is a low-cost tool that can be readily deployed for in-field or over-belt monitoring of biomass quality for bio-refining operations. Rapid image analysis coupled with innovative preprocessing may reduce the impacts of feedstock variability through identification of contaminants or other material attributes to guide selective sorting and quality management. Image analysis was employed to evaluate the quality of corn stover in red-green-blue (RGB) chromatic space. This study used controlled, bench-scale imaging as a proof-of-concept for rapid guality assessment of corn stover based on variations in material attributes, including chemical and physical attributes, that relate to biological degradation and soil contamination. Logistic regression-based classification algorithms were used to develop a method for biomass screening as a function of biological degradation or soil contamination. This study demonstrated the use of image analysis to extract features from RGB color space to investigate variations in critical material attributes from chemical composition of corn stover. Fourier transform infrared (FT-IR) suggested a correlation between red band intensity and biological degradation, while detailed surface texture analysis was found to distinguish among variations in ash. These insights offer promise for development of a rapid screening tool that could be deployed by farmers for in-field assessment of biomass quality or biorefinery operators for in-line sorting and process optimization.

Keywords: image analysis, rapid detection, corn stover, biological degradation, chemical composition, feedstock variability, FT-IR

## INTRODUCTION

The 2016 Billioin Tons Report (BT16) estimates that by 2040, more than 1 billion tons of biomass will be available to achieve a vision of a sustainable bioeconomy (US DOE, 2016). Lignocellulosic biomass has been considered a promising feedstock for biofuels production; therefore, tremendous research efforts have been made to enhance different aspects of the related processes (Cheah et al., 2020). Increases in fuel prices have challenged all countries around the world to develop their own biofuels from renewable resources such as lignocellulosic crops (Qureshi et al., 2010). *Zea mays* is a significant agricultural crop with potential as a biofuel feedstock due to its high carbohydrate content, low production cost, and high availability in the US corn belt (Li et al., 2020). Biomass variability

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originating from production and field conditions propagates with the feedstock through the value chain, posing a challenge to the emerging biorefinery industry (Ray et al., 2020). Variations in lignocellulosic biomass material and quality attributes are often overlooked when assessing feedstock value and pathways for conversion to fuels, chemicals, and products (Ray et al., 2020). Variations in cell wall composition, extractives, moisture content, inorganic species, and soil contaminants have been identified as critical factors affecting biomass quality, process uptime, and product yields (Ray et al., 2020; Sievers et al., 2020; Ding et al., 2021a). Hoover et al. (2019) developed several multiple regression models where five chemical characteristics could be used to estimate biochemical conversion performance. Using these models, an approach for a grading system was demonstrated that could be used to inform markets on the impacts of biomass variability. Hartley et al. (2020) used discrete event simulation that investigated feedstock quality on plant uptime and overall impact to biofuel cost through feedstock delivery cost. These works show how fundamental biomass information might enable real-time decision making on plant profitability and operability. Storage is an essential component of the biomass logistics supply chain, which can have significant impacts to the overall feedstock supply cost and operational reliability (Rentizelas, 2016). Many studies have established that the combination of living cell respiration, biological degradation, and thermo-chemical oxidative reactions bring about significant changes to critical biomass feedstock attributes and process efficiency during storage (Bose et al., 2020; Groenewold et al., 2020; Li et al., 2020). Changes that may occur in stored biomass feedstock include dry mass loss (up to 5.5% per month) (Thornqvist, 1985; Eisenbies et al., 2016), drying, decomposition, and energy content loss (Krigstin and Wetzel, 2016). In addition to capital and operational costs, it is imperative to consider the impact of storage method and format on chemical properties of biomass and overall process efficiency.

Image analysis techniques have been widely applied to measure lignocellulosic biomass features. Image analysis techniques for assessing crops and plant material involve extracting information from digital images of the lignocellulosic biomass (Yan et al., 2020). Unique features of the materials, such as shape, color, and surface texture, can be recognized. Recent research found that crop biomass is highly correlated with different crop parameters, such as leaf area index, crop height, and canopy volume (Lati et al., 2013; Bendig et al., 2014; Tilly et al., 2014), and is also correlated with different vegetation indices based on hyperspectral and red, green, blue (RGB) images (Gupta et al., 2000; Gitelson et al., 2003; Swain et al., 2010; Gitelson et al., 2014; Jannoura et al., 2015). Crop biomass yield has been estimated from spectral information from an unmanned aerial vehicle using standard RGB and multispectral or hyperspectral cameras (Jiang et al., 2019). Image analysis has also been used to estimate the distribution of plants in fields of a clover-grass mixture by using convolutional neural networks trained to predict semantic segmentation maps of clover, grass, and weeds in RGB images containing clover-grass mixtures (Skovsen et al., 2017). Further, Wang et al. found a strong positive correlation between the grayscale values of

biochar and its methylene blue and iodine adsorption capacity, and the Pearson's correlation coefficient range was 0.685–0.977 (Wang et al., 2015). In addition, surface texture can be measured directly from images taken with a laser profilometer (Chinga et al., 2007; Wagner and Horn, 2017), stereomicroscope (Mitra et al., 2014; Piselli et al., 2017), photographic scanner (Aguirre et al., 2018), or scanning electron microscope (Yan et al., 2020). The image analysis can also provide information on the surface roughness based on grayscale values of the image or height map (Chinga et al., 2007).

There are limited publications on visible, red-green-blue (RGB) analysis of biomass variability relevant to biorefineries and lack of such a study limits the ability to develop rapid screening tools for in-field assessment of biomass quality based on physical and chemical attributes. The novelty of this paper was to design and use controlled, bench-scale imaging, employing an off-the-shelf digital camera, as a proof-of-concept for rapid, quality-based assessment of corn stover in visible, redgreen-blue (RGB) space based on variability derived from soil contamination and biological degradation. Logistic regression classification algorithms were used to develop an image screening of biomass as a function of soil contamination and biological degradation. In addition, FTIR was used with a more detailed surface analysis to investigate variation in critical material attributes that arise from chemical composition. Finally, surface texture analysis of the same images distinguished among variable ash levels and degradation. The qualitative results presented in this study show promise for developing rapid screening tools to deploy in-field or in-line for rapid assessment of feedstock quality.

## MATERIALS AND METHODS

## **Corn Stover Bale Collection and Sample Preparation**

Two sets of corn stover samples were imaged and analyzed in this study. The first was a set of 216 core samples from 24 bales obtained from four fields in different central Iowa counties: Hamilton (4 bales), Hardin (6 bales), Story (6 bales), and Poweshiek (8 bales) described in Ray et al. (2020). Baling occurred between October 12 and 27, 2017, using an AGCO 2270XD large square baler, except for Poweshiek County, where a Heston 2270XD square baler was used. Preliminary screening for moisture and ash content was performed by taking three cores per bale to select the 24 bales used in this study. Selected bales were more thoroughly sampled by collecting nine cores per bale, illustrated in a previous study (Ray et al., 2020). Corn stover samples were dried at 40°C and milled with a 2 mm screen in a Thomas Model 4 Wiley Mill (Thomas Scientific, Swedesboro, NJ) for chemical composition analysis. Additional milling using a Retsch ZM200 (Haan, Germany) with a 0.2 mm screen was done to analyze inorganics. For surface analysis, the bales were size reduced through a Vermeer BG480 bale processor with a 75-mm screen, then a Bliss Hammermill with a 25-mm screen, and samples were collected and milled to pass a 2-mm screen for evaluation of surface properties.



FIGURE 1 | (A) Image workflow sample template with internal color standards and biomass samples; (B) Examples of the region of interest selection to measure and output RGB channel data.

The second set of 12 corn stover samples was selected to evaluate material attributes influenced by aging and degradation in storage operations. Bales were from Hardin County, IA (2 bales) and Story County, IA (3 bales) with harvest and baling information described for each county previously. Bale sections exhibiting visual evidence of biological degradation were selected

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for sampling as described in previous studies (Li et al., 2020; Groenewold et al., 2020; Bose et al., 2020). Samples were collected from bale flakes displaying variable extents of biological degradation-moderate biological degradation (medium brown coloration), severe biological degradation (dark brown to almost black), and mild-or negligible biological degradation (light brown). The two bales from Hardin County were size reduced using a Vermeer BG480 bale processer with a 75-mm screen, samples were collected following size reduction, and one sample per bale was visually identified as severely biologically degraded or mild/negligible biologically degraded. The three bales from Story County were manually deconstructed and dissected to collect samples from flakes of the bale that were observed to have mild/negligible biological degradation or moderate/severe biological degradation (Groenewold et al., 2020). For subsequent characterization, samples were milled to pass through a 2 mm and a 0.2 mm screen as described above.

#### **Image Processing and Analysis**

An imaging workflow was developed that uses a Panasonic Lumix G camera with a 88.9-cm square light shed and Metz Mecablitz 52 AF-1 digital flashes. Camera location and settings, as well as flash position, were kept consistent for all images. Each 2-mm sample was imaged using a standardized sample template with internal color standards in each sample photo and a consistent set of biomass samples in each photo (Figure 1A). The image processing workflow included post-processing in SilkyPix Developer Studio 8 SE and data extraction with ImageJ (https://imagej.nih.gov/ij/). Each photo's exposure and gray balance were set individually using the X-Rite ColorChecker Passport Photo 2 with color reference targets included in each image. Regions of interest were defined in each image, as displayed in Figure 1B. The digital photos were decomposed into red, green, and blue channel values ranging from 0 to 255 in relative intensity. The mean, median, standard deviation, minimum, and maximum for the red, green, and blue values range from the regions of interest for all samples were included in the data sets available in the Bioenergy Feedstock Library (Ding et al., 2021b). The image results were aligned with sample metadata, total inorganics, inorganic speciation, and chemical composition (NIRS predicted composition or wet chemical composition) (Ding et al., 2021b). In addition to analytical characterization, three independent observers inspected each sample to assist with positive classification of samples into qualitative sample categories: 1) clean and not degraded, 2) soil contaminated, or 3) degraded (Ding et al., 2021b). This is further discussed below.

#### **Chemical Analysis**

Inorganic speciation for Al, Ca, Fe, K, Mg, Mn, Na, P, Si, Ti, and S oxides were measured according to ASTM standards D3174, D3682, D6349 by a Huffman Hazen Laboratories in Golden, CO and represented on a % dry biomass basis. The chemical composition, including total inorganics, glucan, xylan, lignin, and total extractives, was measured using near-infrared spectroscopy (NIRS) predicted composition or wet chemical composition. For NIRS, predicted composition samples at 2 mm particle size were dried in a desiccator for a minimum of 72 h prior to NIRS analysis. Previous publications have described spectral analysis and NIRS calibration models in detail (Payne and Wolfrum, 2015; Ray et al., 2020). Wet chemical composition was performed in duplicate following the National Renewable Energy Laboratory's laboratory analytical procedures for standard biomass analysis (Sluiter et al., 2010), as described in a previous study (Hoover et al., 2019). The NIRS chemical composition predictive models were built on these same wet chemical procedures.

#### **Observational Dataset**

Three independent researchers performed an observational assessment of each sample and categorized sample quality on the basis of soil contamination and degradation due to biological heating. The surveyed researchers were highly experienced in biomass sampling, preparation, and analytical characterization for assessment of biomass quality, as well as identification of degraded samples that have undergone biological heating or biomass materials with soil contamination. Observational assessment consisted of a -visual (color and clear decay, soil contamination), tactile (texture from grit, soil, silt and/or sand entrapment, and fiber integrity), and odor (presence of astringent odors common to biological degradation) material inspection of each sample and recorded whether it was 1) not soil contaminated and not biologically degraded 2) soil contaminated, or 3) biologically degraded. Samples were assessed with a scale of 0-3 for soil contamination, and biological degradation with 0 = not observed, 1 = low, 2 =moderate, and 3 = high. The observational values were summed for each category for a minimum value of 0 (i.e., each observer selected 0) and a maximum of 9 (i.e., each observer selected 3) [(Ray et al., 2020) Supplementary Material]. Observational data are aggregated in the dataset (Ding et al., 2021b).

#### **Statistical Analysis**

JMP<sup>®</sup> Pro 16.0.0 was used to analyze the combined information from the image analysis, chemical analysis, and observational dataset as described above. Hierarchical cluster analysis using the Ward method was used to group the 222 samples with chemical composition data into four organic chemically distinct groups using glucan, xylan, lignin, and extractives contents (Ding et al., 2021b) along with four inorganic chemically distinct groups for the 191 samples in the dataset with inorganic speciation contents for 191 samples (Ding et al., 2021b). Principal component analysis (PCA) was also used for each organic and inorganic dataset to visualize and interpret the hierarchically defined clusters. Four linear regression models using least squares were generated relating 1) organic chemical components-glucan, xylan, lignin, a 2-way interaction between glucan and xylan, and a 3-way interaction between glucan, xylan, and lignin along with total inorganics-to the median red channel values (Ding et al., 2021b) as described in the image processing and analysis section, 2) the same organic chemical components along with SiO<sub>2</sub> and SO<sub>3</sub> inorganic constituents to the median red channel values 3) organic chemical components to the observed biological degradation dataset as described in the Observational

dataset section and 4) the observed biological degradation levels to median red channel value. For the development of these models, factors were retained if they significantly contributed to the response variable ( $p \le 0.05$ ) and were not highly correlated to other factors in the model except for cases when single factors were retained regardless of their significance if they were included in any interaction terms. Multicollinearity between model explanatory factors was determined based on the Variance Inflation Scores (VIF). A VIF score >10 was considered high multicollinearity between model factors. Nominal logistic fit regression models were used to develop predictive models for biological degradation and soil contamination levels, using the established hierarchical cluster groups as response variables and median values from the red, green, blue channels and 2-way interactions between these values. The dataset used for these models included the 222 samples for predicting the levels of biological degradation with organic chemical data available and 176 samples for predicting levels of soil contamination using samples with both organic chemical and inorganic speciation data available. Factors for each model were removed based on p-values greater than 0.05. The datasets were randomly split into 80% training and 20% validation for these predictive models.

# Fourier-Transform Infrared Spectroscopy Characterization

Corn stover samples for Sample Set 1 were milled using a Retsch ZM200 (Haan, Germany) to 0.2 mm, extractives were retained in the samples. Corn stover samples were extracted with 95:5 acetone/water on a Soxhlet apparatus (~70°C) to remove extractives for Sample Set 2. Corn stover samples for Sample Set 2 were milled in Retsch PM 100 mill fitted with one or two 50 ml ZrO<sub>2</sub> grinding jars and 10\*10 mm ball bearings 10 h to less or equal to 45  $\mu$ m (5 min mill with 10 min interval test). FTIR spectra of all corn stover samples were collected using a Bruker Vertex 70 FTIR spectrometer, equipped with a diamond ATR accessory (Bruker Corporation). Spectra were collected in the spectral range 4000-600 cm<sup>-1</sup>, using 64 scans and 1 cm<sup>-1</sup> resolution. Spectra were baseline corrected at 1840 cm<sup>-1</sup> and advanced diamond ATR fixed with the angle at 45°.

#### **Surface Texture Analysis**

Textural features of cropped images were quantified using the plugins SurfChar J 1q (Chinga et al., 2007), GLCM Texture Too (Haralick, 1979; Lan and Liu, 2018), and FracLac (Smith et al., 1996; Karperien et al., 2013). Before textural analysis, the original color corrected images were cropped to isolate a 900x900-pixel region-of-interest (ROI) centered on the sample cup. Then, the color information was discarded in the image of all subsequent texture analyses and used either 32-bit (SurfCharJ) or 8-bit (GLCM and FracLac) greyscale images. The SurfCharJ 1q package calculated an estimate for the root mean square deviation (Rq), arithmetical mean deviation (Ra), skewness (Rsk), Kurtosis (Rku), lowest valley (Rv), the highest peak (Rp), total profile height (Rt), the mean height of surface

profile (Rc), mean polar facet orientation (FPO), variation of the polar facet orientation (MFOV), the direction of azimuthal facets (FAD), mean resultant vector (MRV), and surface area (SA) from the image. The GLCM Texture Too plugin calculated angular second moment (ASM), inverse difference moment (IDM), contrast, entropy, homogeneity, variance, shade, prominence, inertia, and correlation. The FracLac plugin analyzed the fractal dimensions of the images. Local connected fractal dimension analysis was used, and the Dm output was collected. In total, 27 different surface texture parameters were calculated.

## **RESULTS AND DISCUSSION**

#### **Biological Degradation**

**Figure 2** shows the results and distribution of chemical compositions and inorganic speciation variability of all samples used for image analysis (detailed dataset in (Ding et al., 2021b). Total inorganics,  $SiO_2$ , and  $Al_2O_3$ , revealed substantial variation on a % dry matter basis. Both Si and Albased inorganic species are potential contributions from levels of soil contamination.

#### Image Analysis Biological Degradation

Hierarchical cluster analysis based on chemical components was used to glean key insights about sources of variability (i.e., degradation and soil contamination) that affect quality, with samples grouping into four distinct clusters observed through principal component analysis (PCA) (Figure 3). These chemical composition constituents vary by level of biological degradation (Andrews et al., 1999; Brand et al., 2011; Krigstin and Wetzel, 2016; Groenewold et al., 2020). The loading plot in Figure 3B shows how each of these chemical components is driving the observed cluster in the PCA score plot demonstrated in Figure 3A.

Figure 4A shows a linear relationship between the median red channel and the chemical properties of the samples and was used throughout the analyses. The red channel had the largest range of variability when compared to blue and green, high correlation to other factors used in regression analyses and was correlated to green and blue channels (Supplementary Figure S1, supporting information). Linear relationship factors were identified from chemical data using a stepwise technique as a function of their contributions toward explaining variability in the median red channel output, while minimizing multicollinearity between explanatory factors. For example, extractives and the interaction factor of (glucan x xylan) are highly correlated (Supplementary Figure **S1**, supporting information). Therefore, both were not necessary to explain the variability in the median red channel. The relationship between these chemical properties and the median red channel exceeds an R<sup>2</sup> value of 0.7. Interaction terms between glucan, xylan, and lignin content were included in this analysis. Table 1 displays the standardized coefficients and relative significance level for each chemical material attribute from linear relationships in Figure 4A. These coefficients suggest that xylan was the highest





contributor to explaining the variability seen in the red channel. At the same time, changes in glucan contribute the least in comparison to the other factors. This is consistent with prior work that demonstrated selective degradation of hemicellulose in response to biological heating during storage with cellulosic components involved to a lesser extent (Groenewold et al., 2020). Both factors indicate that as glucan and xylan contents decrease, the red channel signal decreases.

It should be noted that extractives were not used in the model due to the high correlation with glucan and xylan; however, as extractive content increases in the data set, the red channel decreases. Previous studies have demonstrated that extractives content increases with the extent of biological degradation (Groenewold et al., 2020; Li et al., 2020; Ray et al., 2020), consistent with reductions in hemicellulose and to a lesser extent, cellulosic components, as a function of biological



**TABLE 1** Chemical properties are used to form linear regressions for explaining the red-band and biological degradation/self-heating observation variability.

Chemical property	Red band coefficient <sup>a</sup>	Self-heating observation coefficient <sup>a</sup>
Glucan	0.10*	-0.26**
Xylan	0.44**	-0.49**
Lignin	0.25**	-0.18**
Glucan <i>x</i> Xylan	-0.26**	0.34**
Glucan x Xylan x Lignin	-0.15**	not significant
Total Inorganics	-0.35**	N/A

<sup>a</sup>Standardized coefficients.

\*Indicates significance 0.01 < p $\leq$  0.05.

\*\*Indicates highly significant p < 0.01.

degradation. The interaction terms between glucan and xylan suggest that with increases in both, the impact of xylan on the red channel output decreases. During degradation, xylan content, representing the hemicellulose fraction, is more reduced than glucan content (Qing and Wyman, 2011). This interaction term reflects the changing ratio between cellulose and hemicellulose. The same conclusion can be drawn from the interaction between glucan, xylan, and lignin, representing the changing ratios between these chemical attributes in response to degradation. Total inorganics also significantly contributed to changes in the median red channel response. This relationship is further discussed in the following section on soil contamination.

Linear regression analysis was used to relate organic components of glucan, xylan, and lignin to the severity of biological degradation. **Figure 4B** shows the resulting linear relationship, and the coefficients are shown in **Table 1**. The R<sup>2</sup> was 0.77 for this regression, similar to the regression formed between chemical properties and the red band in **Figure 4A**. This relationship also indicates that xylan is the highest contributor, suggesting that lower concentrations of xylan content correspond to higher biological degradation levels. The only organic-based significant factor not shared between

the two regressions (Figures 4A,B) was the interaction between glucan, xylan, and lignin (Table 1). This factor did not significantly contribute to explaining the biological degradation observations. Researcher observations of biological degradation less resolved (scale from 0-9 with each researcher selecting between 0-3) compared to the red channel range (88-161measured); therefore, observational data may not be able to capture the changes in glucanxylan-lignin property relationships, suggesting the basis for differing explanatory factors in Figures 4A,B. The similarities between the linear regressions formed between the red channel and chemical properties and observed levels of biological degradation and chemical properties suggest that the red channel from the image analysis identifies biological degradation in the samples. Figure 4C further corroborates this by showing the strong linear relationship between the red channel outputs and the biological degradation observations  $(R^2 = 0.79).$ 

The relationships observed between the chemical changes, image analysis of the red channel, and biological degradation observations generally align with the overarching hierarchical cluster grouping. Cluster 1 and 2 (**Figure 3A**) correspond with no or mild biological degradation, group 3 as moderate, and group 4 as the severely biological degradation samples.

#### Structural Properties of Biologically Degraded Corn Stover Biomass

To gain a more comprehensive understanding of the impacts of biological degradation on the structural properties of biomass, FT-IR was applied to characterize corn stover with different extents of degradation. Corn stover biomass was selected based on the degree of degradation classified as mild, moderate, and severe or scaled from 1 to 9. The detailed information for the red band values and chemical composition is listed in **Table 2**. The more severely biologically degraded biomass resulted in a lower red band value. **Figure 5** shows the FT-IR spectrum of selected corn stover biomass (Bales 5, 1, and

Sample	Degree of biological degradation	Red band	Glucan (%)	Xylan (%)	Lignin (%)	Total Inorganics (%)
Sample set 2, biologic	cally degraded corn stover <sup>a</sup>					
Bale 5 <sup>b</sup>	Mild	148	33.39	18.19	14.70	12.24
Bale 5 <sup>b</sup>	Moderate	110	32.23	13.08	18.69	8.81
Bale 5 <sup>b</sup>	Severe	106	30.23	10.51	20.41	10.43
Sample set 1, corn st	over <sup>c</sup>					
Bale 1 core 3 <sup>d</sup>	1	141	36.0	19.0	16.2	8.3
Bale 1 core 7 <sup>d</sup>	4	130	31.8	18.7	16.2	9.6
Bale 1 core 6 <sup>d</sup>	7	116	NP	NP	NP	NP
Bale 1 core 4 <sup>d</sup>	8	104	NP	NP	NP	NP
Bale 6 core 1 <sup>d</sup>	1	149	41.2	19.5	17.2	9.5
Bale 6 core 3 <sup>d</sup>	3	138	32.1	19.2	15.8	9.1
Bale 6 core 8 <sup>d</sup>	6	123	NP	18.5	15.0	10.4
Bale 6 core 7 <sup>d</sup>	8	108	31.9	10.1	23.8	12.9
Bale 6 core 6 <sup>d</sup>	9	92	NP	NP	NP	NP

TABLE 2 | Biological degradation properties, red band value, and chemical composition of corn stover for samples characterized using FT-IR.

<sup>a</sup>Obtained by wet chemistry.

<sup>b</sup>Extractives free for FT-IR, measurement, <45 μm.

<sup>c</sup>Obtained by NIR.

<sup>d</sup>Extractives retained for FT-IR, measurement, <0.2 mm.

NP, no prediction, samples fell outside of the NIRS prediction calibration.



6) (See image analysis dataset bale ID (Ding et al., 2021b)) and focuses on the region from 1800-800 cm<sup>-1</sup> reflecting the critical structural properties of biomass (Ma et al., 2018). Remarkable differences were observed in Bale 5. The C-O-C ether stretching at  $1030 \text{ cm}^{-1}$  and  $1242 \text{ cm}^{-1}$  decreased after biological degradation indicating the carbohydrates were hydrolyzed during biological degradation, which corresponded to the decrease of xylan. The absorbance of aromatic C-H stretching of lignin in 1510-1300 cm<sup>-1</sup> also decreased after degradation, suggesting that lignin condensation reactions happened during the biological degradation process. It was hypothesized that the biological oxidation reactions that occur during the degradation result in chromophore formation, such as conjugated carbonyl and quinone, leading to the darkened color of the biomass. In **Figure 5A**, the absorbance of unconjugated  $(1710 \text{ cm}^{-1})$  and conjugated  $(1651 \text{ cm}^{-1})$  carbonyl (C=O) increased and shifted with the reducing value of the red band, implying that lignin was oxidized during the biological degradation process supporting this hypothesis. **Figures 5B,C** show the FT-IR spectrum of the Sample Set 1 corn stover samples. Unlike the degraded samples, the extractives and soils were not removed, and the particle size was 0.2 mm, which is much larger than that used for the Sample Set 2 samples. The FT-IR spectra of the samples show similar results with the degraded samples. However, some samples, such as Bale 6 Cores1 and 8, have weaker absorbance caused by the particle size or soil contaminants. Also, the overall absorbance of Sample Set 1 is lower than the degraded samples (Sample Set 2). Therefore, the particle size of biomass and soil removal is essential for FT-IR ATR



FIGURE 6 | Principal component analysis of soil contamination (A) score plot and (B) loading plot for ash speciation represented on a % dry biomass basis.



FIGURE 7 | Linear regression of measured versus predicted red channel value explained by chemical composition and select inorganic species (SO<sub>3</sub> and SiO<sub>2</sub>) visualizing (A) ash clusters and (B) chemical clusters.

characterization. The FT-IR results demonstrated a potential correlation between the hydrolyzed carbohydrates and the condensed and oxidized lignin in biologically degraded corn stover. These results provide further insights into biopolymer degradation and modification mechanisms during storage, which could correlate to the red band intensity detected through image analysis.

#### Image Analysis of Soil Contamination

As shown in **Table 1** and **Figure 4A**, the total inorganics content significantly contributes to changes observed in the red channel, suggesting that image analysis could provide a valuable tool for

identifying levels of soil contamination. Hierarchical cluster analysis was used similarly on ash speciation data to identify distinct clusters based on inorganic features. Cluster 1 represents limited or no soil contamination, and cluster 4 represents high soil contamination. The PCA score plot (**Figure 6A**) displays these clusters. The variability in the 1<sup>st</sup> principal component suggests a relationship to inorganic species derived from soil contamination, including silicon and aluminum (Lacey et al., 2018). The second principal component relates to inorganic components that serve structural and physiological functions within the plant (i.e., biogenic ash), including sulfur, nitrogen, potassium, calcium, magnesium, and phosphorus (Thy et al.,

TABLE 3   Chemical properties, including select ash species (SO3 and SiO2), were
used to form linear regressions to explain the red-band variability.

Chemical property	Red band coefficient <sup>a</sup>		
Glucan	0.06		
Xylan	0.33**		
Lignin	0.22**		
Glucan x Xylan	-0.36**		
Glucan x Xylan x Lignin	-0.17**		
SiO <sub>2</sub>	-0.17**		
SO <sub>3</sub>	-0.30**		

<sup>a</sup>Standardized coefficients.

(\*) indicates significance 0.01 < p< 0.05.

\*\*indicates highly significant p < 0.01.

2013; Li et al., 2020). The inorganic clusters 1 and 2 vary primarily based on intrinsic inorganic species inherent to plant biomass, and clusters 3 and 4 show relationships to soil contaminant inorganic species (i.e., silicon and aluminum).

Figure 7A shows the resulting regression with the ash speciation clusters labeled, and Figure 7B shows the same regression with the chemical clusters, that were based on glucan, xylan, lignin, and extractives contents. Regression analyses highlight that for this dataset, the red channel relationships are being driven by the biological degradation severity, even with the highest ash sample in the data representing close to 62% total inorganics. The regression shown in Figure 7A includes SO<sub>3</sub> and SiO<sub>2</sub> content instead of total inorganics, as was included in the regression represented in Figure 4A, to represent the relationship of these intrinsic and extrinsic ash species to the image red channel intensity. Phosphorus and potassium oxides were not selected for the linear regression as representatives of intrinsic ash, as suggested by the principal component 2 in the principal component loadings plot (Figure 6B) as they did not significantly contribute to explaining the red variability. It should be noted that fewer samples in the dataset contained ash speciation compared to the organic composition data available (191 samples compared to 222 samples). Therefore, the coefficient changes cannot be directly compared to the model in Figure 4A; however, the general trends remain consistent. Both SO<sub>3</sub> (representative of intrinsic ash) and SiO<sub>2</sub> content (representative of soil contamination) significantly explain variability in the red channel (Table 3).

In summary, regression analyses associated with both biological degradation and soil contamination identified relationships among chemical components in corn stover, namely xylan, glucan, lignin, total inorganics, Si, and S, strongly correlated to the red band intensity detected through image analysis. The variations in organic chemical attributes xylan, glucan, and lignin were attributed to biological degradation. In contrast, variations in inorganic attributes, Si and S, corresponded to the extent of soil contamination and intrinsic inorganic content in the sample. Although a mechanistic understanding of the biological degradation (Groenewold et al., 2020; Ding et al., 2021a) and thermo-chemical oxidative reactions that alter biomass quality attributes during storage (Krigstin and Wetzel, 2016) is required to sort out confounding signals from the degradation and accumulation of inorganic species, qualitative results presented here show promise for developing rapid screening tools to deploy in-field or in-line for rapid assessment of quality (Ray et al., 2020).

#### **Screening Prediction Development**

The purpose of the regression analyses and relationships, as previously discussed, was to demonstrate that aspects of the images, the median red channel intensity specifically, could be quantitatively related to the chemical changes known to be impacted by biological degradation and soil contamination. Here the red channel along with the blue and green channels from the images was used to demonstrate potential screening methods for corn stover samples to predict the presence of biological degradation and soil contamination. For this demonstration, logistic regression was used to predict the hierarchical groups of chemical clusters representing biological degradation and inorganics clusters representing soil contamination considering the median red, blue and, green channel outputs and interactions between these color channels as predictors. For the organic chemical cluster logistic regression, the receiver operator characteristics (ROC) curve (Figures 8A,B) demonstrates the diagnostic ability of image analysis properties to correctly identify the organic chemical cluster, representing levels of biological degradation, that each sample belongs to by comparing the sensitivity, true identification rate, versus the specificity, falsepositive identification rate. These plots, along with the confusion matrix, give the actual number of samples predicted to be in each group versus their actual group (Table 4) for both the training and validation sets. The confusion matrix indicates that this model correctly identified the severely biologically degraded samples (cluster 4) 100% of the time with no false identifications. However, the identification accuracy was lower for cluster group 3, moderate levels of biological degradation, and reduced further for clusters 1 and 2, representing no and mild biological degradation. The validation sample results from this logistic regression follow the same trend. Samples with severe biological degradation (cluster 4) are accurately predicted 100% with no false positives and show progressive decreases in the sensitivity and specificity for clusters 3, 2, and 1. This preliminary model indicates that this approach could be employed to identify biomass samples exhibiting moderate to severe biological degradation, but requires expanded datasets and further refinement to distinguish between samples with mild or limited biological degradation and samples without degradation.

Linear regression analysis indicated that biological degradation was the primary factor driving variaitions in the red-channel intensity, with soil contamination contributing secondarily. Based on this knowledge, the proposed process for screening feedstock for soil contamination is to the first screen for evidence of moderate and severe biological degradation and then use



FIGURE 8 | Receiver operator characteristic (ROC) curves for the logistics regression training sets using image analysis data red, green, blue channels to predict the chemical cluster (associated with levels of biological degradation) (A) training set and (B) validation set and ash cluster (associated with levels of soil contamination) (C) training set and (D) validation set.

TABLE 4   Confusion matrix for chemical cluster logistics regression usin	ng red,
green, and blue channels image analysis data.	

Actual Training	Predicted				
	1	2	3	4	
1	31 (50%)	30	1	0	
2	20	62 (72%)	4	0	
3	3	6	16 (65%)	0	
4	0	0	0	5 (100%)	
Validation					
1	4 (25%)	12	0	0	
2	5	15 (71%)	1	0	
3	1	1	4 (67%)	0	
4	0	0	0	1 (100%)	

This table shows the actual number of samples predicted to be in each group versus their actual group and the true positive percentages for accurate prediction.  
 TABLE 5 | Confusion matrix for ash cluster logistics regressions using image analysis data red, green, and blue channels, and chemical cluster input.

Actual Training	Predicted				
	1	2	3	4	
1	47 (75%)	11	5	0	
2	14	39 (72%)	1	0	
3	7	4	12 (52%)	0	
4	0	0	0	1 (100%)	
Validation					
1	13 (81%)	2	0	1	
2	3	7 (58%)	0	2	
3	1	2	4 (57%)	0	
4	0	0	0	0 (.)	

This table shows the actual number of samples predicted to be in each group versus their actual group, along with the true positive percentages for accurately predicted.



parameters.

the outputs of the biological degradation prediction levels to screen for the additional presence of soil contamination. The resulting ROC curve and confusion matrices from this soil contamination logistics regression are shown in Figures 8C,D and Table 5. These results indicate enhanced diagnostic ability to predict higher levels of soil contamination (cluster group 4), representing a sample with 38% total ash and 28% SiO<sub>2</sub> content. As only one sample was available at this level, no validation samples were used for this cluster group. Cluster 1 samples representing the lowest ash samples with on average 10% ash content and 6% SiO<sub>2</sub> content was accurately 75 and 81% for the training and validation sets, respectively. The model had relatively equal power for identifying the lower levels of soil contamination representing average total ash 18 and 11% and SiO<sub>2</sub> contents of 11 and 6% for clusters 3 and 2, respectively. These results show the potential for the image analysis to be used as a screening tool for soil contamination after considering color changes due to biological degradation; however, a larger dataset with known validation samples are necessary to improve modelaccuracy.

#### **Surface Texture Analysis**

The image analysis based on color variability was used to identify biological degradation and soil contamination levels. In this research, image analysis of surface properties was also investigated to identify soil contamination, biological degradation, and additional properties impacting particle flow. For the samples with variable ash content, samples were selected with ash content of 5–10% (low ash) and 10–20% (high ash) (Sievers et al., 2020), surface roughness calculated as Rq (**Figure 9A**), and the GLCM parameter Shade (**Figure 9C**) appear to distinguish between high and low ash samples. The



pattern, however, was different between Rq and Shade, with the high ash sample measuring low Rq and higher Shade values. Rq is a measure of surface roughness that is positively correlated with inter-particle friction and could impede flowability. However, surface roughness is also correlated with hydrophobicity. Therefore, particles that trap less water should have better flowability. Shade characterizes the tendency of clustering of pixels as a measure of asymmetry. In our previous work using images of hammer-milled biomass, the asymmetric clustering that shade describes was positively related to poor conveyance (Gudavalli et al., 2020). The fractal analysis (**Figure 9B**) or the GLCM parameter ASM (**Figure 9D**) differed among the ash variable samples (Sample Set 1).

Mild, moderately, and severely degraded samples (Sample Set 2) were partly distinguished by three of the four texture parameters shown in **Figure 10**. The surface roughness (Rq, **Figure 10A**) reveals the most separation. Shade and ASM (**Figures 10C,D**) showed differences between mild and severe samples. These differences can be generated by how the samples fracture during milling, causing subtle differences in particle size and shape distributions—these differences in particles present as differences in the texture of biomass piles. For example, surface texture analysis could distinguish among variable ash levels or levels of degradation.

## CONCLUSION

This study used image analysis in visible, red-green-blue (RGB) chromatic space to evaluate the quality of corn stover across four Iowa counties representing a realistic supply shed in the US corn belt. Linear regression relationships with  $R^2>0.7$  were found between the red channel values from images of corn stover and changes in chemical properties resulting from biological degradation (xylan, glucan, and lignin) and soil contamination (Si and S). The FT-IR results demonstrated a potential correlation between the hydrolyzed carbohydrates and the condensed and oxidized lignin in biologically degraded corn stover, which could

correlate to the red band intensity detected through image analysis. In addition, surface texture analysis of for these same images was found to distinguish among variable levels of ash and degradation.

Further, logistic regression classification algorithms were used to develop an image analysis method for screening and classifying levels of biological degradation and soil contamination in corn stover. This prototype supports research and development that uses image analysis and other rapid characterization tools to further understand and describe corn stover and bioenergy feedstock quality during in-field or over-belt applications to support industrial operations for sorting biomass based on quality or presence of contaminants.

#### DATA AVAILABILITY STATEMENT

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found in the article/ **Supplementary Material**.

#### **AUTHOR CONTRIBUTIONS**

LD and K-TL contributed to investigation, methodology, formal analysis, and writing the original draft manuscript. AH and RE contributed to conceptualization, methodology, formal analysis, data curation, and writing—draft, reviewing and editing. JG and BD performed the surface texture analysis and contributed to writing the

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original draft, reviewing, and editing; BD contributed to funding acquisition. RC and BT were involved in investigation and formal analysis. JK contributed to conceptualization and editing. WS contributed to conceptualization. AR contributed to funding acquisition, supervision, conceptualization, writing—reviewing and editing.

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#### SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fenrg.2022.837698/full#supplementary-material

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