Spectral detection of nematodes in soybean at flowering growth stage using unmanned aerial vehicles

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ABSTRACT: Soybean is one of the main crop species grown in the world. However, there is a decline in productivity due to the various types of stress, including the nematodes Heterodera glycines and Pratylenchus brachyurus. The objectives were to determine the best spectral band for detecting H. glycines and P. brachyurus at the beginning of flowering (R1). Soil and root sampling was conducted at nine sampling sites in each of the five nematode-infested regions, totaling 45 sampling points. Flights were made at all regions using Phantom 4 Advanced, Sequoia and 14-band customized Sentera. For H. glycines, the red spectral band best explained the variability on soil and root nematode counts as well as the second stage of juveniles in soil. For P. brachyurus, Sentera RedEdge best explained the variability in root nematode counts and Sequoia NIR best explained soil juveniles. A multiple linear regression model using spectral data for detecting P. brachyurus and H. glycines improved R² compared to simple linear regressions. At flowering growth stage (R1), soybean spectral reflectance was associated with the number of H. glycines and P. brachyurus on soil and roots using low-cost and multispectral sensors.

Key words: Glycine max, drones, remote sensing, precision farming, integrated management.

INTRODUCTION

Soybean [Glycine max (L.) Merril] is one of the oldest crops cultivated by humans and is currently cultivated worldwide (HYMOWITZ, 1970). Soybean is used on grain-based foods for humans and animals, and as a raw material for biofuel production (ZHANG et al., 2017).

Various types of stressors affect Soybean causing yield losses, in particular the soybean cyst nematode Heterodera glycines and the root lesion nematode Pratylenchus brachyurus (NIBLACK, 2005; WRATHER & KOENNING, 2005; GOUART, 2008; WANG et al., 2015; PENG et al., 2016).

Heterodera glycines is a sedentary endoparasite that infects roots and may cause above-ground symptoms, including plant dwarfism, leaf chlorosis, early senescence, lower seed weight, and ultimately plant death (NIBLACK, 2005; ZHANG & SONG, 2017). Symptoms of attack of P. brachyurus are characterized by spots on leaves that often change in color. Roots infected by P. brachyurus have reddish-brown lesions progressing toward necrosis (DIAS et al., 2010). Irregular shapes of clustered plants can be observed after infection from H. glycines and P. brachyurus, but these patterns may be similar to the symptoms from water and nutritional

Received 03.31.20 Approved 10.23.20 Returned by the author 12.09.20 CR-2020-2803.183
stress (BLEVINS et al., 1995). Resistant soybean cultivars can prevent yield losses, but the integration of control methods is important keep resistance (TYLKA & MARK, 2002; CONCIBIDO et al., 2004; MITCHUM et al., 2007; NIBLACK et al., 2008).

In addition, the diagnosis of nematode-infected areas is costly because the precise estimation of population distribution requires a large number of soil samples across the field (MARTINS et al., 2017). One way of indirectly determine nematode damage in large areas and in short time intervals is the use of remote sensing. Symptoms on leaves, nematode low mobility in soil, and plant-infested clusters are argued to enable detection of nematodes by orbital and aerial imaging, facilitating the use of imaging applications on precision agriculture (HILLNHÜTTER et al., 2012). For nematode-induced stress, several sensor-based processes were tested using terrestrial and aerial platforms (HEATH et al., 2000; NUTTER et al., 2002; LAUDIEN, 2005; HILLNHÜTTER et al., 2012). Several authors T use of remote sensors on studies evaluating asymptomatic and symptomatic plant problems (MARCASSA et al., 2006; CHO et al., 2012; HUNT et al., 2013; OUMAR et al., 2013; ASHOURLOO et al., 2016).

Among the narrow spectral bands widely used in agriculture, hyperspectral bands are widely used to determine plant canopy characteristics, which can be caused to detect various types of stressors (GEBBERS & ADAMCHUK, 2010; MULLA, 2013; MAHLEIN et al., 2013). In short, hyperspectral remote sensing can be a valuable tool for early detection of plant disease, as the electromagnetic spectrum can detect imperceptible changes to the human eye (MAHLEIN et al., 2012; MARTINELLI et al., 2015). Spectral bands of sensors can record information from the electromagnetic spectrum within a pre-determined interval of wavelengths (BENDIG et al., 2014; JANNOURA et al., 2015). The central spectral band of the sensor typically contributes to most of the spectral information within the bandwidth. For better detection of pathogens, narrow band vegetation indices (IVs) are increasingly being used. Through them, it is possible to evaluate changes in vegetation at various scales, as well as to estimate the damage caused by pathogens (MARCUSI et al., 2010; MAHLEIN et al., 2013).

Unmanned Aerial Vehicles (UAVs), the miniaturization of spectral sensors, and new algorithms have enabled several applications of digital imaging for quick, high-precision cartographic products from large areas (COLOMINA & MOLINA, 2014). Using remote sensing for precise pathogen detection and mapping and localized management. Relatively rapid and early identification of the spatial distribution of a disease is possible using canopy-based reflectance methods (MOSHOU et al., 2004; QIN et al., 2009; SHAFRI & HAMDAN, 2009; SANKARAN et al., 2010). Few studies have explored the use of RGB, multispectral and hyperspectral images for detecting soybean cyst and root lesion nematode. NUTTER et al. (2002) used aerial remote sensing to quantify soybean cyst nematode population density reported and RE that 60% variability of the initial population can be explained by canopy reflectance. BAJWA et al. (2017) could distinguish a healthy soybean plant from another with cyst with high accuracy values.

The objectives of this study were to determine the best spectral band for detecting *H. glycines* and *P. brachyurus* using simple linear regression models and to determine a multiple linear regression model for their detection at the beginning of flowering (R1).

**MATERIALS AND METHODS**

**Study area**

The study area of approximately 330 hectares was located near the city of Rio Verde during the 2017/2018 growing season. This area had a dystrophic red Latosol, elevation of approximately 953 meters, slope ranging from flat (0 to 3%) to gentle undulating (3 to 8%), the average temperature of 22.3 °C and annual rainfall around 1600 millimeters (SOMA BRASIL, 2019) (Figure 1). Preliminary analysis determined that the experimental area had *H. glycines* from races 1, 3, and 6, *P. brachyurus* and *Helicotylenchus dihystera* (data not shown). Nematode species were identified under the microscope following methodology from COOLEN & D’HERDE (1972) by trained personnel from the Phytopathology Laboratory of the Federal Institute of Goiás, in the municipality of Rio Verde, State of Goiás. The races of *H. glycines* were determined using the method described in CORDEIRO et al. (2008).

**Materials**

The sensors used in this study included the Sequoia, Sentera and RGB on Phantom 4 Advanced. The Sequoia had four separate sensors: green at 550 nanometers (nm), red at 660 nm, red edge at 735 nm, and nir at 790 nm. Sentera had eight sensors from the traditional spectral bands found on common sensors: 446, 548, 586, 615, 650, 661, 775, and 825 nm. Sentera also had a NDVI sensor containing a 625 nm (red) and 850 nm (NIR) spectral bands. The NDRE
sensor contained another NIR band at 840 nm (NIR) and 720 nm (red edge).

Although, Sequoia had the RGB sensor, this sensor was not used because it had some noise in the image. The RGB sensor onboard of Phantom 4 Advanced was not specified by the manufacturer and was, therefore, set to red-Phantom, green-Phantom, and blue-Phantom. Sequoia sensor was boarded on a Phantom 4 Advanced. Sentera sensor was boarded on the Inspire 2. For each sampling site in the patches, an A4-size white sheet was used to find the sampling sites in the images. A Garmin etrex 20 was used to find the sampling sites in the 5 nematode-infested regions.

For preparing the flight plans for RGB-Phantom and Sequoia sensors, we used the DroneDeploy application. We used the Field Agent for flight definition for Sentera. Other materials such as hoe, properly identified plastic bags, sieves, among others, were used for collecting soil, root, and plants in the field and, consequently, nematode extraction. Pix4d software was used to create the orthorectified maps. Qgis software was used to extract pixel information from the sampling regions and to create maps. R software was used for statistical analysis (‘rumor’ package).

**Plant height, root biomass and nematode data**

A preliminary flight with the RGB sensor from the 2017/18 soybean crop was used to determine areas of interest that were potentially infested by nematodes. These areas were chosen to facilitate field sampling and increase the chances of finding higher degrees of severity (chlorosis). A total of 5 regions were selected for this research to have a significant number of samples for the elaboration of linear regressions.

Each region had 9 sampling sites. The first sampling site was at the center of the nematode-infested region. Sampling sites at 10, 20, 40, and 80 meters were determined in two directions from the center of the region (Figure 1). The directions were set so that the sites of one patch did not coincide with another patch. The sampling sites represented high, intermediate, and low (non-significant) populations of *H. glycines* and *P. brachyurus*.

The cultivar MONSOY 7198 was planted on October 10, 2018. This cultivar is considered an early group of maturation, and resistant to *H. glycines* races 1 and 3. Sampling and flights were performed on the same days during the R1 stage of soybean. This reproductive stage was chosen because of less...
interference of soil reflectance than that from the vegetative stage. At each sampling site, soil samples were taken at a depth from 0 to 20 cm in a “V” shape, and then the soil samples were mixed homogeneously. Roots were extracted to quantify the nematodes *H. glycines* and *P. brachyurus* and to measure the height, dry and green mass of the canopy of six plants, as well as the dry and green mass of six roots.

The extraction of *P. brachyurus* and *H. dihystera* in soil and root was done by the methods of JENKINS (1964), and COOLEN and D’HERDE (1972). For the second stage juveniles of *H. glycines* in soil and root, the adapted methodology of JENKINS (1964) and ALFENAS (2007) was used. For cysts (viable and non-viable) in soil, the methodology adapted by ARAÚJO (2009) was used and for females in the root, the method adapted by TIHOHOD (2000).

**Planning and operation of the flight**

The flights were made on the same days as the collection of agronomic and nematode data, so that they were performed whenever possible between 10h00 and 14h00, to minimize shadow effects generated by the position of the sun in relation to the objects.

In all flight plans, side and longitudinal overlaps were set to 80% to prevent problems during the creation of the orthomosaics. Our tests using several flight overlap indicated 80% as optimal for our drones and environmental conditions. Side overlap must prevent failures between consecutive lanes because of drift caused by high winds, drone inclinations, flight height variation, and terrain elevation. In addition, using a larger side overlaps avoids using the extreme edges of images whose radial lens distortion is often greater. Longitudinal overlap is important to provide supporting points by photo triangulation of images and stereoscopic.

The flight height for all sensors was set to 40 meters, resulting in a 1.2-cm pixel for the Phantom RGB sensor, 3.2-cm pixel for Sentera, and 4-cm pixel for Sequoia. It was not possible to obtain a cloud-free sky in all flights. However, the sampling regions were always cloud-free in the images.

**Flight processing and information extraction from orthomosaics**

Orthomosaic of the sampling areas was created using Pix4D software. It was necessary to extract some information associated with each image, such as camera calibration data and log file (coordinates of the center of each image and inclinations, ω, ϕ, k). As there was no calibration panel, digital number of the pixels were used in the analyses. Therefore, orthomosaic was imported into Qgis for using multiple spectral bands. The tool used for this was the raster calculator.

For each sampling site, a 1 m by 25 cm vector layer was created to extract the value of the corresponding pixels in each band of each sensor and obtain the reflectance average for further statistical analysis. A supervised classification removed pixels associated with soil and shadow background, leaving only pixels related to plants for the statistical analyses. The average of the pixels was obtained with the aid of the grid creation tools and statistics by zone.

**Statistical analysis**

Due to the large number of variables involved in this study and the search for a statistic that could best explain the behavior of the data, simple and multiple linear regressions were used. The objective of using the simple linear regression was to identify the best spectral bands for detecting the occurrence of soybean cyst nematode and root lesion nematode. In addition, the multiple regression identified the best combination of spectral bands for detecting nematodes. A forward model selection was conducted with all spectral bands and sensors to select a subset of them that would work together to characterize the variability on nematode densities. The most important spectral band was selected by the lowest P-value from the simple linear regression models. Other independent, non-redundant spectral bands were added sequentially (one-by-one) considering the lowest Mallows’s Cp value (HAIR et al., 2009; GIONGO et al., 2020). Residues from the models were visually inspected for normality, independence and homoscedastic. The data transformation was not needed. Simple and multiple regression models were plotted using digital number of soybean reflectance as independent variable (x) and number of nematodes as a response variable (y). Only models with significant P-values (α = 0.5%) are included in the tables.

**RESULTS AND DISCUSSION**

The dwarf plants and exposed soil within the nematode-infested regions were distinct from the other parts of the soybean field (Figure 2). This contrast offers an opportunity for growers to mapping nematodes using mathematical models and low-cost sensors such as that of Phantom 4 Advanced. The geographical information of nematode infestation
can support localized application of chemical or biological control products that can reduce the costs of controlling nematodes. *Pratylenchus brachyurus* was the most abundant nematode in the study with up to 126 nematodes in 100 cm³ of soil and up to 1439 nematodes in 10 grams of roots. The cultivar used in the study was race-resistant one and three; therefore, the population of *H. glycines* was considerably smaller in our evaluations, with the maximum number of viable cysts in the soil of 12, maximum number of non-viable cysts in the soil of 91; maximum number of juveniles of *H. glycines* of 125 in the soil and 13 in the roots. We also identified *H. glycines* (races one, three, and six) and *H. dihystera* nematodes in the samples. However, *H. dihystera* was excluded in the statistical analysis because it is an ectoparasite assumed to cause non-significant damage to soybean (MANSO et al., 1994). The resistant soybean cultivar partly explained the higher numbers of juveniles and cysts of *H. glycines* in the soil than that observed in the roots.

The spectral bands at 615 nm and red-Phantom were associated with *H. glycines* soybean cyst nematodes at all sampling regions (Table 1). The red-Phantom best explained the variability of data for juveniles and non-viable cysts (Table 1). Such a result is similar to the result of BAJWA et al. (2017) in soybean, referring to the study of simple correlation of bands with *H. glycines* and sudden death syndrome. Despite the low capability of explaining the variability on *H. glycines* data (R² = 0.197), our low-cost red-Phantom sensor detect spectral changes in soybean due to nematode injury similar to more expensive sensors (Table 1).

Juveniles of *H. glycines* from most of the sampling sites were not associated with visible symptoms and changes in plant spectral responses. Spectral bands associated with nematode injury were different between *P. brachyurus* and *H. glycines* (Table 1), indicating that spectral bands have different ability to detect plant responses to pathogens (SANKARAN et al., 2010). Because plant damage was directly associated with root nematodes rather than soil nematodes, the equations of great interest for simple linear regression for *H. glycines* are those using the red-Phantom and for root lesion nematodes, the Sentera RedEdge, as they have the best value of R² and p-value (Table 1).

For *H. glycines*, higher numbers of juveniles in roots increased the reflectance at visible spectral bands (Table 1). Increasing red reflectance may have indicated a lower biomass of shoots and roots, as well as a decrease in plant height (MARTINS et al., 2017; BAJWA et al., 2017; MAHLEIN et al., 2012). Changes in red reflectance may also be associated with chlorophyll content in the leaves (ALVES et
| Dependent nematode/agronomic variable | Independent variable (x) | Regression equation | P-value | R² |
|---------------------------------------|--------------------------|---------------------|---------|----|
| Non-viable cyst                       | 615 nm                   | Y = -43.4171 + 0.9169 × x | 0.032   | 0.081 |
|                                       | RedEdge - Sentera (720 nm) | Y = -25.9480 + 0.7000 × x | 0.041   | 0.072 |
|                                       | Red - NDRE (840 nm)       | Y = -3.4.2038 + 0.4268 × x | 0.034   | 0.079 |
|                                       | Red - Phantom             | Y = -87.5923 + 0.9258 × x | 0.003   | 0.162 |
| H. glycines in root                   | 615 nm                   | Y = -4.9.340 + 0.0827 × x | 0.021   | 0.096 |
|                                       | Red - Phantom             | Y = -9.0733 + 0.0850 × x | 0.001   | 0.197 |
| H. glycines in soil                   | 615 nm                   | Y = -55.8575 + 0.9485 × x | 0.006   | 0.138 |
|                                       | Red - Phantom             | Y = -93.7538 + 0.8870 × x | < 0.001 | 0.222 |
| P. brachyurus in root                 | RedEdge - Sentera (720 nm) | y² = 1286.9880 - 14.7590 × x | 0.007   | 0.137 |
|                                       | RedEdge - Sentera (735 nm) | y² = 1374.3350 - 8.2390 × x | 0.011   | 0.121 |
| P. brachyurus in soil                 | RedEdge - Sequoia (735 nm) | y² = 632.8905 + 0.0290 × x | 0.024   | 0.091 |
| Plant height                          | 615 nm                   | y² = 99.8122 - 0.0024 × x | 0.042   | 0.071 |
|                                       | Red - Sentera (650 nm)    | y² = 32.5222 - 0.1373 × x | 0.018   | 0.102 |
|                                       | Green - Sentera (548 nm)  | y² = 60.4020 - 0.3280 × x | < 0.001 | 0.422 |
|                                       | Blue - Sentera (446 nm)   | y² = 48.6091 - 0.2725 × x | 0.022   | 0.094 |
|                                       | Red - Phantom             | y² = 37.7422 - 0.1260 × x | 0.003   | 0.161 |
|                                       | RedEdge - Sequoia (735 nm) | y² = 12.9613 - 0.0003 × x | 0.005   | 0.146 |
| Shoot dry mass (MSA)                  | Air - Sequoia (790 nm)    | y² = 9.6545 - 0.0003 × x | 0.015   | 0.107 |
|                                       | 615 nm                   | y² = 3.9011 - 0.0203 × x | 0.045   | 0.068 |
|                                       | 586-nm                   | y² = 5.0940 - 0.0177 × x | 0.031   | 0.082 |
|                                       | Red - Sentera (650 nm)    | y² = 7.5385 - 0.0431 × x | < 0.001 | 0.245 |
|                                       | Green - Sentera (548 nm)  | y² = 6.3636 - 0.0259 × x | 0.015   | 0.109 |
|                                       | Blue - Sentera (446 nm)   | y² = 7.4970 - 0.0537 × x | 0.008   | 0.129 |
|                                       | Red - Phantom             | y² = 5.4740 - 0.0259 × x | < 0.001 | 0.236 |
|                                       | Green - Phantom           | y² = 5.3136 - 0.0160 × x | 0.013   | 0.113 |
|                                       | Blue - Phantom            | y² = 4.4609 - 0.0214 × x | < 0.001 | 0.210 |
|                                       | Green - Sequoia (550 nm)  | y² = 3.7115 - 0.0001 × x | 0.036   | 0.077 |
| Root dry mass (MSR)                   | 615 nm                   | y² = 10.1170 - 0.0634 × x | 0.023   | 0.093 |
|                                       | 586-nm                   | y² = 13.5265 - 0.0551 × x | 0.019   | 0.108 |
|                                       | Red - Sentera (650 nm)    | y² = 14.6932 - 0.0751 × x | 0.030   | 0.083 |
|                                       | Green - Sentera (548 nm)  | y² = 14.7055 - 0.0596 × x | 0.046   | 0.067 |
|                                       | Blue - Phantom            | y² = 9.2664 - 0.0367 × x | 0.048   | 0.066 |
|                                       | 586-nm                   | y² = 7.2453 - 0.0237 × x | 0.026   | 0.089 |
|                                       | Red - Sentera (650 nm)    | y² = 7.7008 - 0.0330 × x | 0.043   | 0.070 |
|                                       | Green - Sentera (548 nm)  | y² = 9.9035 - 0.0414 × x | 0.002   | 0.176 |
|                                       | Blue - Sentera (446 nm)   | y² = 10.1784 - 0.0688 × x | 0.009   | 0.125 |
|                                       | Red - Phantom             | y² = 6.6788 - 0.0249 × x | 0.011   | 0.119 |
|                                       | Green - Phantom           | y² = 8.2020 - 0.0264 × x | 0.002   | 0.179 |
|                                       | Blue - Phantom            | y² = 5.9044 - 0.0230 × x | 0.007   | 0.135 |
|                                       | Green - Sequoia (550 nm)  | y² = 5.6894 - 0.0001 × x | 0.008   | 0.130 |
| Shoot green mass (MVA)                | 615 nm                   | y² = 31.4041 - 0.2121 × x | 0.006   | 0.138 |
|                                       | Red - Sentera (650 nm)    | y² = 57.2280 - 0.3434 × x | < 0.001 | 0.253 |
|                                       | Blue - Sentera (446 nm)   | y² = 50.3016 - 0.3554 × x | 0.029   | 0.085 |
|                                       | Red - Phantom             | y² = 40.8630 - 0.2072 × x | < 0.001 | 0.246 |
| Root green mass (MVR)                 | 615 nm                   | y² = 26.3539 - 0.1318 × x | 0.011   | 0.120 |

For *P. brachyurus*, higher numbers of nematodes in roots increased reflectance at the red edge (Table 1). Stepwise multiple linear regression for detecting *P. brachyurus* and *H. glycines* improved R² in relation to simple linear regressions. Multi-band regression models also improved detection of injury from soybean aphids *Aphis glycines* (ALVES et al., 2018). For *P. brachyurus*, the most important spectral range was outside the visible spectral range. For *H. glycines*, the visible wavelengths were the most important spectral bands to estimate nematode injury on soybean (Table 2).
CONCLUSION

Soybean canopy reflectance from flowering growth stage (R1) was associated with the number of *H. glycines* and *P. brachyurus* on soil and roots. Therefore, low-cost sensors at the visible spectral range were sufficient to estimate *H. glycines*, but multispectral sensors were necessary to estimate *P. brachyurus*. Although, p-values indicated a significant association between soybean reflectance and nematode numbers, a better understanding of other factors affecting soybean canopy reflectance is still necessary to improve our models.

ACKNOWLEDGMENTS

This research was supported by equipment and laboratories from the Instituto Federal Goiano (IF) and the Polo de Inovação of the Rio Verde - Go campus. In addition, Coordenação de Aperfeiçoamento de Pessoal de Nível Superior (CAPES) financed the research grants. Thanks to the Federal Institute Goiano Rio Verde campus and Proppi for supporting the study and translation of the article.

DECLARATION OF CONFLICTS OF INTEREST

The authors declare no conflict of interest. The founding sponsors had no role in the design of the study; in the collection, analysis, or interpretation of data; in the writing of the manuscript, and in the decision to publish the results.

AUTHORS’ CONTRIBUTIONS

All authors contributed equally for the conception and writing of the manuscript. All authors critically revised the manuscript and approved of the final version.

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