



# Article A Method of Invasive Alien Plant Identification Based on Hyperspectral Images

Xi Qiao <sup>1,2,3,†</sup>, Xianghuan Liu <sup>1,2,†</sup>, Fukuan Wang <sup>1,2,†</sup>, Zhongyu Sun <sup>4</sup>, Long Yang <sup>4</sup>, Xuejiao Pu <sup>1</sup>, Yiqi Huang <sup>2</sup>, Shuangyin Liu <sup>3</sup> and Wanqiang Qian <sup>1,\*</sup>

- <sup>1</sup> Shenzhen Branch, Guangdong Laboratory of Lingnan Modern Agriculture, Genome Analysis Laboratory of the Ministry of Agriculture and Rural Affairs, Agricultural Genomics Institute at Shenzhen, Chinese Academy of Agricultural Sciences, Shenzhen 518120, China
- <sup>2</sup> College of Mechanical Engineering, Guangxi University, Nanning 530004, China
- <sup>3</sup> Guangzhou Key Laboratory of Agricultural Products Quality & Safety Traceability Information Technology, Zhongkai University of Agriculture and Engineering, Guangzhou 510225, China
- <sup>4</sup> Guangzhou Institute of Geography, Guangdong Academy of Sciences, Guangzhou 510070, China
- \* Correspondence: qianwanqiang@caas.cn
- + These authors contributed equally to this work.

Abstract: Invasive alien plants (IAPs) are considered to be one of the greatest threats to global biodiversity and ecosystems. Timely and accurate detection technology is needed to identify these invasive plants, helping to mitigate the damage to farmland, fruit trees and woodland. Hyperspectral technology has the potential to identify similar species. However, the challenge remains to simultaneously identify multiple invasive alien plants with similar colors based on image data. The spectral images were collected by a hyperspectral camera with a spectral range of 450–998 nm, and the raw spectra were extracted by Cubert software. First derivative (FD), Savitzky-Golay (SG) smoothing and standard normal variate (SNV) were used to preprocess the raw spectral data, respectively. Then, on the basis of preprocessing, principal component analysis (PCA) and ant colony optimization (ACO) were used for feature dimensionality reduction, and the reduced features were used as input variables for later modeling. Finally, a combination of both dimensionality reduction and non-dimensionality reduction is used for identification using support vector machines (SVM) and random forests (RF). In order to determine the optimal recognition model, a total of 18 combinations of different preprocessing methods, dimensionality reduction methods and classifiers were tested. The results showed that a combination of SG smoothing and SVM achieved a total accuracy (A) of 89.36%, an average accuracy (AA) of 89.39% and an average precision (AP) of 89.54% with a test time of 0.2639 s. In contrast, the combination of SG smoothing, the ACO, and SVM resulted in weaker performance in terms of A (86.76%), AA (86.99%) and AP (87.22%), but with less test time (0.0567 s). The SG-SVM and SG-ACO-SVM models should be selected considering accuracy and time cost, respectively, for recognition of the seven IAPs and background in the wild.

Keywords: hyperspectral analysis; invasive plants; data preprocessing; dimension reduction; classification

# 1. Introduction

Invasive alien plants (IAPs) have become one of the greatest threats to biodiversity, ecosystems and agriculture around the world, causing very large ecological and economic losses [1]. Furthermore, the threat caused by IAPs is growing due to the globalization of human activities and economics [2]. Various projects have been adopted to restrain and eliminate IAPs in recent years. During this time management has played a limited role and the threat is expanding [3,4]. More recently, accurate identification and monitoring, which can provide the detailed distribution of IAPs, has been considered an efficient and low-cost approach in IAP control [5].



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**Copyright:** © 2022 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). Traditionally, manual inspection has been the main method of monitoring for IAPs, which is considered an accurate but inefficient measure [6]. Moreover, various factors, such as the growth of plants, the inability to assess a large spatial coverage, the subjective judgment of scientists, and the inaccessible terrain have restricted monitoring [7]. Hyperspectral remote sensing is regarded as an effective monitoring method and has been successfully used in some IAP monitoring, such as for *Mikania micrantha* Kunth [8], *Spartina alterniflora* Loisel [9–11], *Psidium cattleianum* [12], *Lantana camara* L. [13], *Common milkweed* [14] and *Ailanthus altissima* [15]. These studies only identify single invasive plants and did not attempt to classify multiple species. However, the simultaneous identification of multiple IAPs based on imaging data remains challenging due to similar phenotypic colors. Hyperspectral images can provide rich spectral information and have the potential to simultaneously identify the colors of similar species [16,17]. This paper aims to identify multiple IAPs based on hyperspectral images.

Generally, the full-spectrum curve information of objects is extracted from hyperspectral images and used for identification through machine learning methods, such as support vector machine (SVM) and random forest (RF) [18]. There is no denying that capturing certain bands from hyperspectral images may result in the loss of important spectral information. However, seeking key information from the full spectral curve is indispensable since the hyperspectral image contains numerous invalid data, which will interfere with the identification and decrease the time-validity [19]. Furthermore, it is certain that some noise will occur in the wild and interfere with the hyperspectral information. Therefore, image preprocessing and the selection of critical spectral bands are essential [20].

Researchers have proposed a number of spectral preprocessing methods and have applied them in many studies. Sun et al. applied five preprocessing methods, including standard normal variate (SNV), multiple scattering correction, Savitzky–Golay (SG) smoothing, normalization and first derivative (FD), to pretreat the spectral data, and then a SVM model was used for the determination of moisture content in barley seeds [21]. Pang et al. proposed a method of detecting the vigor of variabilis seeds. In this research, using SG smoothing and second-order derivative (SD) preprocessing combined with least squares support vector machines (LS-SVM) achieved 98.81% accuracy in the test set [22]. Shen et al. explored an application of hyperspectral imaging technology to estimate the soil organic matter, no denoising, SG smoothing denoising and wavelet domain denoising were used to eliminate the noise in hyperspectral data [23].

Meanwhile, it is necessary to select the information that is valuable for the experiment, as hyperspectral data are always redundant and much information is useless [24]. The methods for selecting the information, which also means reducing the dimensionality of hyperspectral data, have been widely applied in many studies [25,26]. Principal component analysis (PCA) and ant colony optimization (ACO) have shown potential in their application for reducing dimensionality.

This research aims to explore the feasibility of classifying Mikania micrantha Kunth, *Sphagneticola calendulacea* (L.) Pruski, *Ageratum conyzoides* L., *Mimosa pudica* Linn., *Lantana camara* L., *Lpomoea cairica* (L.) Sweet, *Bidens pilosa* L. and background in the wild by hyperspectral data. Selecting a proper method for processing hyperspectral data and distinguishing the seven IAPs is a great challenge. Moreover, challenges are manifested in the variability of the hyperspectral data of IAPs in a complex field environment, the lack of prior knowledge and background interference.

To meet these challenges, hyperspectral preprocessing methods FD, SG smoothing, SNV and the dimensionality reduction algorithms PCA and ACO were applied for processing the hyperspectral data of IAPs and background. In the final classification, SVM and RF were used. Finally, preprocessing algorithms, dimensionality reduction algorithms and classifiers were randomly combined to study and explore an optimal identification method for IAPs in the field.

## 2. Materials and Methods

# 2.1. Sample Preparation

In this paper, we manually applied the high-speed imaging spectrograph S185 manufactured Cubert (Germany) to collect hyperspectral images of seven species of IAP in the wild. The spectrometer simultaneously recorded 138 bands in the spectral range of 450–950 nm, with a sampling interval of 4 nm. The collection sites were Jiulong town (23°22'29.5" N, 113°29'52.9" E; 23°22'13.4" N, 113°28'46.9" E) (collected *M. micrantha, S. calendulacea, A. conyzoides, M. pudica, L. camara* and *B. Pilosa*), Conghua district (23°28'25.8" N, 113°29'09.9" E) (collected *S. calendulacea, A. conyzoides, M. pudica, L. camara* and *L. cairica*), Guangzhou, China, and Sanjiao town (22°42'24.6" N, 113°25'41.7" E) (collected *M. micrantha, A. conyzoides, M. pudica, L. cairica* and *B. pilosa*), Zhongshan, China.

These seven IAPs pose a huge threat to native species and biodiversity. They are similar in color, phenotype, and appear together in small plots, which is why hyperspectral techniques were used to identify seven species simultaneously. The images were captured between 10 am and 3 pm around solar noon on between 21 and 25 November 2020, when the weather was cloudy. This heightened the chances of suitable weather conditions for the measurements [27,28]. Before image acquisition, dark reference (by closing the camera lens) and white reference (using a white plate) images were collected to calibrate the spectrometer according to the following equation [29]:

$$I_{C} = (I_{R} - I_{D}) / (I_{W} - I_{D})$$
(1)

where  $I_C$  is the calibrated image,  $I_R$  is the raw image,  $I_W$  is the white reference, and  $I_D$  is the dark reference.

Then, the spectrometer's lens was aimed directly at the surface of the invasive plant and manually focused to the middle of the leaves. Eighty images of vegetation were collected using the hyperspectral camera and were used in this study. Partial samples of seven invasive plants are shown in Figure 1. The images contain leaves, flowers, stalks, other plants and the no-plant cover area. Among them, flowers, stalks and the no-plant cover area belong to background. The leaves and background feature points were extracted manually from the original images using Cubeware software (approximately 43 per image, 38 of which are leaves feature points), as shown in Figure 2A; the spectral curves are shown in Figure 2B and saved in ASCII format. In this work, 3359 raw spectral data samples (*M. micrantha*: 428, background: 409, *S. calendulacea*: 428, *A. conyzoides*: 425, *M. pudica*: 419, *L. camara*: 408, *L. cairica*: 418, *B. pilosa*: 424) were collected, including a random partition a training set (2400 samples, each category has 300 samples) and testing set (959 samples).

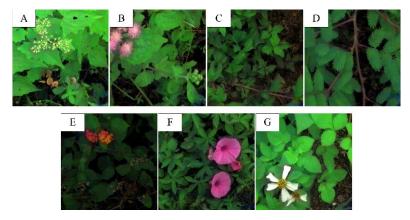
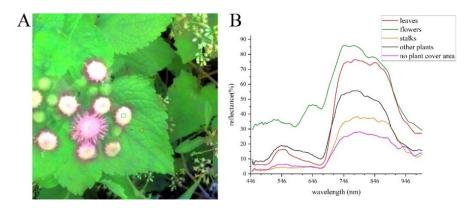


Figure 1. Partial samples of seven invasive plants and the calibration plate. (A) *Mikania micrantha* Kunth, (B) *Sphagneticola calendulacea* (L.) Pruski, (C) *Ageratum conyzoides* L., (D) *Mimosa pudica* Linn., (E) *Lantana camara* L., (F) *Ipomoea cairica* (L.) Sweet, (G) *Bidens pilosa* L.



**Figure 2.** Extraction of raw spectral data. (**A**) Pixels of hyperspectral images used for raw spectra extraction, (**B**) raw spectral data.

# 2.2. Preprocessing Methods

Hyperspectral data have a large amount of interference redundancy information, which seriously affects the accuracy of the identification. Studies have shown that the interference of noise, astigmatism and baseline drift caused by human factors and background can be reduced by preprocessing [30]. At present, preprocessing methods are relatively mature and can be divided into derivative algorithms, smoothing algorithms, and light scattering correction algorithms. FD, SG smoothing and SNV were used in this paper.

## 2.2.1. First Derivative

First derivatives (FD) can reduce hyperspectral mutual interference and reduce noise in the analysis of hyperspectral data [31], and they are often used for baseline correction and background elimination in hyperspectral analysis [32]. The formula for spectral reflectance for the FD at wavelength point *k* can be solved as follows:

$$x_{i,FD} = \frac{x_i - x_{i+k}}{k} \tag{2}$$

In the above formula, *k* is the window width, k = 9.

# 2.2.2. Savitzky-Golay Smoothing

The smoothing methods include moving window smoothing and Savitzky–Golay (SG) convolution smoothing [33]. Compared with the moving window smoothing method, SG convolution smoothing can achieve a better denoising effect because moving window smoothing is limited by the number of windows. If the number of windows is too large or too small, the denoising effect will be affected. The two smoothing methods are similar, but the difference is that SG convolution smoothing focuses on the calculation of the central value, while moving window smoothing focuses on the number of windows. That is, SG smoothing is an algorithm based on polynomial smoothing, which smooths the spectral curve through moving windows combined with least squares fitting. Its calculation formula is as follows:

$$Y_{k,SG} = \frac{1}{N} \sum_{i=-m}^{m} (Y_{k+i}A_i)$$
(3)

where  $A_i$  is the smoothing coefficient,  $Y_{k,SG}$  is the smoothed value of k points,  $Y_{k+i}$  is the unsmoothed value of k, i is the number of smoothed data points from this point to the left, m refers to the number of smoothed data points from this point to the right, and N is the normalization factor.

## 2.2.3. Standard Normal Variable Transformation

The standard normal variate (SNV) transformation is mainly used to eliminate the influence of solid particle size, surface scattering and optical path changes on the reflected spectrum [34]. SNV is the same as the standardized algorithm, except that the standardized algorithm addresses a set of spectra (based on the column of the spectral array), while SNV addresses a spectrum (based on the row of the spectrum) [35].

$$x_{snv} = (x - \omega) / \sqrt{\frac{\sum_{k=1}^{m} (x_k - \omega)^2}{(m - 1)}}$$
(4)

on the type of  $\omega = \frac{\sum_{k=1}^{m} x_k}{m}$ , *m* is the number of wavelength points.

# 2.3. Dimension Reduction

The raw and preprocessing spectral data are 138-dimensional, and a high number of dimensions may lead to low efficiency of hyperspectral modeling and poor model performance. Therefore, it is necessary to reduce the dimensionality of hyperspectral data and retain most of the dataset information. Many studies have shown that the dimensionality reduction processing of hyperspectral data can not only shorten the time but can also improve the recognition efficiency to some extent [36]. In this paper, PCA and ACO were used to reduce the dimensionality of hyperspectral data of invasive plants.

## 2.3.1. Principal Component Analysis

PCA is a dimensionality reduction method that transforms multidimensional data features into a few comprehensive features and uses variable information represented by a few variable features [37,38]. The PCA dimension reduction transformation process is as follows: suppose that the n-dimensional preprocessing result variable  $(x_1, x_2, x_3, \dots, x_n)$  to perform the linear transformation of Formula (5) on variable  $x_i$  ( $i = 1, 2, 3, \dots, n$ ) to obtain the comprehensive variable  $(y_1, y_2, y_3, \dots, y_n)$  is,

$$\begin{cases} y_1 = a_{11}x_1 + a_{12}x_2 + a_{13}x_3 + \dots + a_{1n}x_n \\ y_2 = a_{21}x_1 + a_{22}x_2 + a_{23}x_3 + \dots + a_{2n}x_n \\ \vdots \\ y_n = a_{n1}x_1 + a_{n2}x_2 + a_{n3}x_3 + \dots + a_{nn}x_n \end{cases}$$
(5)

Equation (5) linear transformation satisfies the following conditions:

(1)  $y_1, y_2, y_3, \cdots, y_n$  are linearly independent;

(2)  $a_{i1}^2 + a_{i2}^2 + a_{i3}^2 \cdots + a_{in}^2 = 1$ ,  $(i = 1, 2, 3, \cdots, n)$ ;

(3) the composite variables are arranged according to their variances.

The process of PCA is based on the contribution rate of a single principal component and the cumulative contribution rate. When the cumulative contribution rate of the first m principal components reaches 85%, the first *m* principal components can represent most of the information of the n-dimensional data to achieve dimensionality reduction. The calculation formulas of the principal component contribution rate and cumulative contribution rate are as follows:

$$\eta = \frac{\lambda_i}{\sum_{i=1}^n \lambda_i} \tag{6}$$

$$\rho = \frac{\sum_{k=1}^{m} \lambda_k}{\sum_{i=1}^{n} \lambda_i} \ (m = 1, 2, 3, \cdots, n)$$
(7)

In Formulas (6) and (7),  $\lambda_i$  is the principal component contrast.

#### 2.3.2. Ant Colony Optimization

ACO is a probabilistic dimension reduction method for spectral feature selection of hyperspectral data [38]. Based on ACO, the main idea is to feature the selection problem

of modeling for solving the minimum cost path search graph, thus creating a node of the search space and design procedures to find the manners of solution. ACO states the path that the ant chooses of the choice probability of the feature subset, based on the pheromone and heuristic information to compute the ant's choice. In this paper, the Fisher criterion is used to calculate heuristic information, and the ant's transition from one node to another node is determined by probability. Assume that the transfer probability  $P_{ij}^k$  of ant k from node i to node j at time t is calculated by the following formula:

$$P_{ij}^{k}(t) = \begin{cases} \frac{\tau_{ij}^{\alpha}(t)\gamma_{ij}^{\beta}(t)}{\sum_{s \in allowed_{k}}\tau_{is}^{\alpha}(t)\gamma_{is}^{\beta}(t)}, j \in allowed_{k} \\ 0, else \end{cases}$$
(8)

When the ant completes a cycle, the pheromone on each path is calculated as follows:

$$\tau_{ij}(t+n) = (1+\rho)\tau_{ij} + \Delta\tau_{ij} \tag{9}$$

 $\tau_{ij}$  refers to the pheromone concentration on the path from node *i* to node *j* at time *t*, and  $\gamma_{ij}$  is the heuristic function, which refers to the expectation of ant *k* choosing to remove node *j* from node *i*. The calculation formula is as follows:

$$\gamma_{ij} = 1/d_{ij} \tag{10}$$

where  $d_{ij}$  is the distance from node *i* to node *j*. The value of  $allowed_k = 1, 2, ..., n - 1, n$  is the total number of nodes,  $allowed_k$  represents the nodes that ant *k* can select next,  $\alpha$ ,  $\beta$  is the weight parameter, which plays a role in the importance of pheromones and heuristics,  $\rho$  is the pheromone evaporation coefficient, and  $\Delta \tau_{ij}$  is the pheromone concentration increment on the path from node *i* to node *j*.

#### 2.4. Identification Models

The typical algorithms based on hyperspectral image data classification include the neighborhood algorithm, SVM, RF, and neural network. Domestic and foreign researchers have accomplished many achievements on invasive plants based on hyperspectral data, most of which are based on the spatial distribution [39], area extraction [40], and species identification of invasive plants [41]. According to the characteristics of the samples, SVM and RF were selected to identify seven invasive plants.

SVM is a classical machine learning algorithm based on traditional theory; the core content of this is to transform the object of our study into a high-dimensional feature space by nonlinear transformation and to construct a linear decision function in the high-dimensional space to realize the nonlinear decision function of the original space [42]. In this paper, the kernel function selected by SVM classifier is the radical basis function (RBF). For each SVM model, a grid search combined with a 5-fold cross validation method was used to optimize parameters. The search interval of the penalty coefficient and kernel parameter was set as  $2^{-5} \sim 2^5$ , the step size was 0.1.

RF is a combination of classification trees that implements randomization [43]. In the 1980s, Breiman et al. invented the classification tree algorithm [44]. It is calculated by repeating dichotomous data. In the same year, Breiman merged classification trees into random forests. RF generate multiple classification trees by randomizing the data and variables, and then obtaining statistical results. The main parameters of RF are the number of trees and the number of nodes. In this paper, the number of trees was set to 1000, and the number of nodes was set to 10 for all of the RF models.

## 2.5. Performance Evaluation

The performance of all of the combined models was evaluated based on four statistical parameters, namely, accuracy (A), average accuracy (AA), average precision (AP) and test

time, which were often used to evaluate the performance of the classification models and calculated from Equations (11) and (12) [45].

$$Accuracy = \sum_{i=1}^{8} TP_i / N \times 100\%$$
(11)

Average Precision = 
$$\sum_{i=1}^{8} \left(\frac{TP_i}{TP_i + FP_i}\right)/8$$
 (12)

where *TP* is the true positive sample, *TN* is the true negative sample, *FN* is the false negative sample, *FP* is the false positive sample, *N* is the total sample number, *i* is the number of categories.

All of the aforementioned steps were coded and developed in the environment of MATLAB R2020a (The Math Works Inc., Natick, MA, USA). The GPU of the PC is a NVIDIA GTX 3060.

#### 3. Results

#### 3.1. Preprocessing

Raw and preprocessing spectral data of seven invasive plants and background are shown in Figure 3A–D. In Figure 3A, M. micrantha has the highest reflectivity in the wavelength range of approximately 700 to 900 nm, followed by B. pilosa, M. pudica, L. cairica, A. conyzoides, S. calendulacea and L. camara. In the range of approximately 450 to 680 nm, the background reflectance is relatively scattered, while in the range of approximately 700 nm to 998 nm, the background overlaps with the spectral reflectance of seven invasive species, and the overlap region between species was large. Therefore, it is challenging to classify seven invasive species simultaneously. To facilitate the identification of raw spectral data, three processing methods were used to reduce data noise and highlight the distribution patterns of reflectivity with wavelength. The original spectral data processed by SG smoothing are shown in Figure 3C. Compared with Figure 3A, the curve of Figure 3C is relatively smooth, and small fluctuations in reflectance over the whole wavelength range are eliminated. The other two processing methods, FD and SNV, remove other noise from the original spectral data. Figure 3B presents the FD processing of direct difference analysis of the raw spectral data, that is, noise reduction is performed at both ends of the spectral band. Compared with Figure 3B, the raw spectral data are obviously interfered with by artificial and systematic noises, and the intraclass difference of spectral data after FD processing is less than that of the original spectral data. Figure 3D displays the preprocessing data of SNV, and the intraclass difference in spectral data is reduced.

In summary, all three preprocessing methods can eliminate part of the spectral noise. SG smoothing reduced the small range of spectral fluctuations, FD and SNV reduced the intra-class differences in species, but none of the processing methods significantly reduced the inter-class differences in species and background. Therefore, it is necessary to determine the inter-class differences of the seven species through subsequent treatments. To determine a more appropriate pretreatment method, the next step is to analyze the impact of each processing method combined with dimension reduction.

#### 3.2. Dimension Reduction by PCA

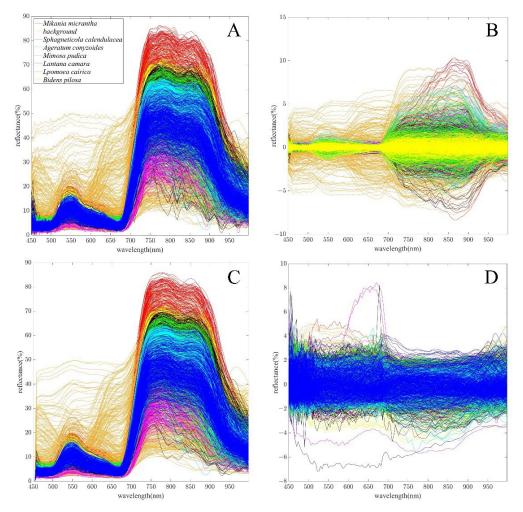
Based on the preprocessing results, dimension reduction analysis was conducted by PCA, as shown in Figure 4A,C,E, which are the point states drawn in horizontal (first principal component scores) and vertical (second principal component scores) coordinates. In this way, we can preliminarily judge the convergence degree of FD, SG smoothing and SNV. According to Figure 4A,C,E, it is obvious that FD-PCA has a better sample clustering effect, but *S. calendulacea, L. camara* and *B. pilosa* overlap clearly. In Figure 4C, the seven invasive plants are relatively scattered; that is, after SG-PCA treatment, the sample points of invasive plants have almost no similar convergence. In terms of clustering

performance, FD ranked first, SNV ranked second, and SG ranked third. The worst clustering performance of SG smoothing preprocessing is predictable because SG smoothing reduces noise but does not change details and overall trends. Therefore, more components need to be considered later.

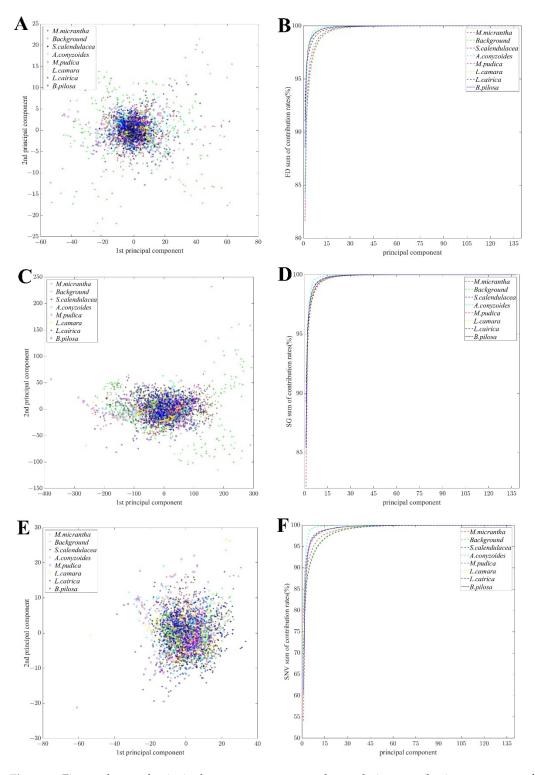
The cumulative contribution rate of the first 138 principal components were calculated as shown in Figure 4B,D,F. It is clear that at around the first 22 principal components, all curves tended to be smooth and the cumulative principal component contribution rate approaches 100%. Therefore, the principal component characteristics of the first 22 dimensions can fully represent the spectral information of 138 dimensions. In other words, after PCA, the 138-dimensional spectrum is reduced to 22 dimensions.

#### 3.3. Dimension Reduction by ACO

Based on the preprocessing results, feature selection was carried out by ACO, and the feature bands are shown in Table 1.



**Figure 3.** Raw spectral curve of 3359 samples (**A**), raw spectral curve, preprocessed by (**B**) first derivative, (**C**) SG smoothing, (**D**) standard normal variate.



**Figure 4.** First and second principal component scores and cumulative contribution rate curve of the principal components of 3359 samples based on (**A**,**B**) first derivative, (**C**,**D**) SG smoothing, (**E**,**F**) standard normal variable transformation.

Preprocessing Methods	Number of Feature Bands	Feature Bands (nm)
FD	28	450, 454, 458, 462, 466, 470, 474, 478, 482, 486, 494, 498, 502, 506, 510, 514, 518, 526, 530, 542, 598, 614, 738, 770, 834, 858, 934, 942,
SG smoothing	20	450, 454, 458, 462, 466, 470, 474, 478, 482, 486, 490, 494, 510, 518, 526, 638, 658, 702, 798, 914
SNV	25	450, 454, 458, 462, 466, 470, 474, 478, 482, 486, 494, 498, 502, 506, 510, 514, 518, 526, 530, 542, 578, 662, 774, 822, 950

Table 1. Feature bands.

The number of feature bands selected based on FD was 28, SNV was 25 and SG was 20. Table 1 shows that the feature bands selected based on the three preprocessing methods are concentrated in the range of 450–700 nm. That is, the feature wavelengths are concentrated in the visible range.

Wavelengths of 450, 454, 458, 462, 466, 470, 474, 478, 482, 486, 494, 510, 518 and 526 nm were common to all three preprocessing methods, indicating that these wavelengths were important features in distinguishing the seven species. Based on ACO, the extracted feature wavelengths were relatively continuous between 450–700 nm and relatively scattered between 700–998 nm. Furthermore, the selection of feature wavelengths is related to the quality of the preprocessing. After all, the number of feature wavelengths extracted based on SG is the lowest and the number based on FD is the highest.

## 3.4. Identification Performance Evaluation of Different Combination Models

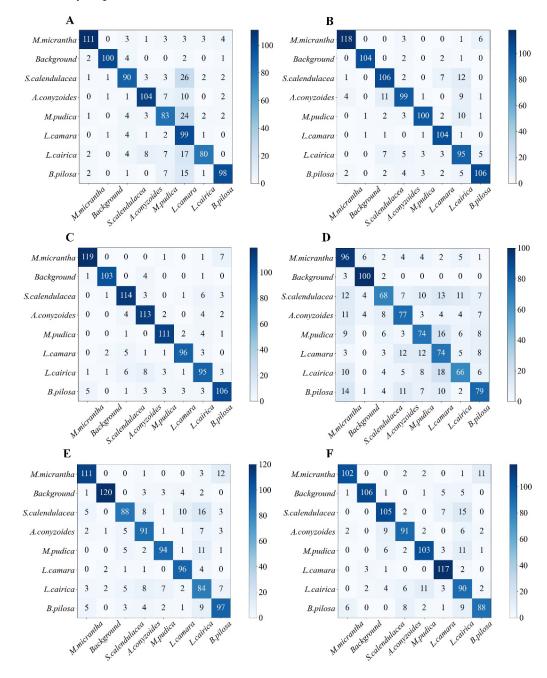
Among the 3359 samples, 2400 samples were used for model training and 959 samples were used for model testing. To explore the single species recognition rates and total recognition rates for each combined model, confusion matrices, single species recognition accuracy tables and total accuracy tables are presented, as shown in Figure 5, Tables 2 and 3 (Figure 5 and Table 2 only present the models with the highest accuracy under the same conditions).

Models	SNV-PCA-SVM	SG-ACO-SVM	SG-SVM	SNV-PCA-RF	SG-ACO-RF	SG-RF
M. micrantha	86.72%	92.19%	92.97%	80.00%	87.40%	86.44%
Background	91.74%	95.41%	94.50%	95.24%	90.23%	89.08%
S. calendulacea	70.31%	82.81%	89.06%	51.51%	67.18%	81.40%
A. conyzoides	83.20%	79.20%	90.40%	65.25%	81.99%	81.25%
M. pudica	69.75%	84.03%	93.28%	60.66%	82.46%	81.75%
L. camara	91.67%	96.30%	88.89%	63.25%	92.31%	95.12%
L. cairica	67.80%	80.51%	80.51%	56.41%	71.19%	76.27%
B. pilosa	79.03%	85.49%	85.49%	61.72%	80.17%	77.19%

Table 2. Recognition accuracy of combined model for single species.

SG-SVM showed outstanding advantages in the identification process among nearly all species. The SG-SVM identified 92.97%, 89.06%, 90.40%, 93.28%, 80.51% and 85.49% accuracy for *M. micrantha, S. calendulacea, A. conyzoides, M. pudica, L. cairica* and *B. Pilosa,* respectively, which are far better than those of SNV-PCA-SVM (86.72%, 70.31%, 83.20%, 69.75%, 67.80%, 79.03%), SNV-PCA-RF (80.00%, 51.51%, 65.25%, 60.66%, 56.41%, 61.72%). Meanwhile, the total accuracy (A), the average accuracy (AA), the average precision (AP) of SG-SVM are also excellent, reaching 89.36%, 89.39% and 89.54%, respectively. SG-ACO-SVM (86.76%, 86.99%, 87.22%) cannot be ignored. Compared to SG-SVM, the test time is also shorter, although A, AA and AP are all reduced. When the A of the model was reduced by about 3%, the cost time was nearly 5 times shorter. Moreover, for *L. camara* recognition,

SG-ACO-SVM (96.30%) is superior to SG-SVM (88.89%). SG-SVM and SG-ACO-SVM show advantages in recognition accuracy and time cost, respectively. In practice, when the total accuracy is relatively important, the SG-SVM model is selected, and when the test time cost is relatively important, the SG-ACO-SVM model is used.



**Figure 5.** Confusion matrices of the combined models, (**A**) SNV-PCA-SVM, (**B**) SG-ACO-SVM, (**C**) SG-SVM, (**D**) SNV-PCA-RF, (**E**) SG-ACO-RF, (**F**) SG-RF.

Models	A (%)	AA (%)	AP (%)	Test Time (s)
FD-PCA-SVM	77.89	78.13	79.82	0.09493
SG-PCA-SVM	72.99	73.37	75.63	0.09382
SNV-PCA-SVM	79.77	80.03	82.82	0.08942
FD-ACO-SVM	48.28	48.95	54.09	0.0766
SG-ACO-SVM	86.76	86.99	87.22	0.0567
SNV-ACO-SVM	67.67	68.16	67.76	0.0746
FD-SVM	67.26	67.66	67.70	0.3524
SG-SVM	89.36	89.39	89.54	0.2639
SNV-SVM	84.15	84.32	84.21	0.329
FD-PCA-RF	63.30	63.77	64.70	0.1031
SG-PCA-RF	64.44	64.42	64.36	0.0978
SNV-PCA-RF	66.11	66.76	66.75	0.1035
FD-ACO-RF	61.00	61.34	63.41	0.0841
SG-ACO-RF	81.44	81.61	81.74	0.0788
SNV-ACO-RF	67.36	68.08	68.01	0.0821
FD-RF	62.15	62.40	62.71	0.1407
SG-RF	83.63	83.56	84.15	0.0862
SNV-RF	71.01	71.23	71.11	0.1015

Table 3. The total accuracy, average accuracy and average precision of the 18 combined models.

#### 4. Discussion

Currently, manual monitoring is still the mainstay of invasive plant monitoring. However, this makes the monitoring of invasive species both laborious and time-consuming. Given these facts, unmanned aerial vehicle (UAV) with hyperspectral cameras can offer significant opportunities for invasive plant monitoring. Hyperspectral remote sensing, which combines the advantages of imaging and spectroscopy, has been successfully applied to invasive plant monitoring. Previous studies using hyperspectral imaging to monitor the spectral information of invasive plants at different time periods have been very effective. The results obtained in this study show that the spectral distribution of different invasive plants during the same period is different, but the differences are not significant. In general, the lower reflectance of invasive plants occurs in the visible part of the spectrum (400–700 nm), and the absorption of photosynthetic pigments in the region associated with it is largely concentrated in the blue (450-chlorophyll b) and red (650-chlorophyll a) regions.

The highest total accuracy of the combined model in this paper was only 89.39%. On the one hand, only traditional machine learning models are used, on the other hand, SVM and RF is mostly suitable for binary classification [46,47]. After all, SVM and RF can often achieve high accuracy in the application of binary classification [48]. Although the LIBSVM toolbox is capable of solving multiclassification problems [49], the total classification accuracy of samples is not high because the difference of spectral features between various samples is not big, or the difference of features between samples is further reduced after normalization. Dimensionality reduction reduces the training and testing time of the model, but inevitably leads to the loss of some important information, which is the main reason why the accuracy of dimensionality reduction models is lower than that of non-dimensionality reduction models. Of course, the small amount of data is also an important reason. Therefore, it is necessary to expand the datasets or try to use a new type of model to improve the accuracy and efficiency.

Recent studies have used hyperspectral imaging combined with deep learning methods to monitor invasive plants [50,51]. Meanwhile, deep learning is very popular in species classification [52,53], but it does not mean that the recognition effect of deep learning is superior to that of other algorithms. For example, in the recognition of grapevine species, the recognition effect of a SVM is not inferior to that of a convolutional neural network [54]. However, it cannot be denied that deep learning is a current trend in image recognition. To further improve the accuracy of recognition, it is necessary to adopt a deep learning algorithm. By studying the leaf spectra of the seven invasive plants, 18 combination models were able to distinguish seven invasive plants within a short time interval. By analyzing different combination models, it is found that the total accuracy of SG-SVM is the highest, followed by SG-ACO-SVM. It is worth noting that the spectrum of SG-ACO-SVM was only 20-dimensional, while the spectrum of SG-SVM was 138-dimensional. Although the total accuracy of the SG-ACO-SVM is slightly less, the testing time has been significantly shortened. Of course, the reduction in dimensionality does not necessarily indicate that the models will take less time to test. After all, the SG-PCA-RF and SNV-PCA-RF models are more time consuming than their non-dimensional reduction counterparts. However, the training time for the reduced dimension models is clearly lower than that for the non-dimensional reduction models, and this is true for all of the combined models. While the total accuracy is reduced, a satisfactory recognition accuracy is obtained in a shorter period of time. This is the purpose of using dimensionality reduction and the reason for comparing the 18 combined models.

In this paper, seven invasive plants were identified using 18 models, though promising, in-depth research is necessary before the technique can be applied to the field for accurate monitoring. One aspect that should be considered in future research is the spectral monitoring of invasive plants over their entire period. On the other hand is the consideration of the average spectra of more organs of the invasive plant, such as the flowers, stems, leaves, etc. of the plant. To achieve this, operational challenges have to be overcome. Such as the selection and collection of optimal conditions for hyperspectral images of the plant under test, the determination of the spectral distributions and correlated color temperatures of solar radiations, the calibration of black and white under different atmospheric conditions, the possibility of mounting a hyperspectral sensor on an unmanned aircraft, the calibration of the sensor, the collection, storage and analysis.

The main work of this research was to complete the identification of seven invasive species in four small fields. The image samples used were hyperspectral images taken with a handheld spectrometer. In practical applications, it is often necessary to identify a large range of invasive plants, which requires the installation of a hyperspectral imager on a UAV for image acquisition. Our study provides a reliable reference for hyperspectral image data processing and the classification of a variety of invasive plants.

# 5. Conclusions

Hyperspectral data of invasive plants in the wild were used as sample data in this work, and three groups of different results were obtained by different preprocessing methods (FD, SG, and SNV). Based on the three preprocessing results, nine groups of data were obtained by combining dimensionality reduction (PCA, ACO) and no dimensionality. The nine groups of data were combined with classification methods (SVM and RF) to establish different classification models to obtain the best recognition method for invasive plants.

The results show that the SG preprocessing method has a better preprocessing effect on hyperspectral data, and most values of the models obtained under this condition are optimal. Based on the preprocessing results, by comparing the results of dimensionality reduction and no dimensionality reduction, the method of dimension reduction may not improve the accuracy of recognition, but it can improve time efficiency. Compared with the RF classifier, the SVM classifier has a better recognition effect on invasive plants. In the dimension reduction method based on PCA and ACO, SVM model recognition based on the ACO dimension reduction method is better. Among the 18 models, the 2 best models are SG-ACO-SVM (AA, 86.99%, AP, 87.22%, TT, 0.0567) and SG-SVM (AA, 89.39%, AP, 89.54%, test time, 0.2639). **Author Contributions:** X.Q. and Y.H.: methodology, software, validation, writing. X.L.: methodology, software, validation, writing-original draft, writing-review. F.W.: software. Z.S. and L.Y.: project administration. X.P. and S.L.: project administration. W.Q.: project administration, Writing-original draft, Writing-review and editing. All authors contributed to the article and approved the submitted version. All authors have read and agreed to the published version of the manuscript.

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