# A PLANT RECOGNITION APPROACH USING HIGH RESOLUTION NETWORK

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**Abstract.** Plant species recognition plays an important role in agriculture, the pharmaceutical industry, and conservation. The traditional approaches may take days and have difficulties for non-experts. Several computer vision-based models have been proposed, which can partially assist and speed up the plant recognition process. Thanks to the development of data collection and computational systems, the models based on machine learning have considerably improved their performance in the last decades. In this paper, we present a model for plant recognition in Southeast Asia based on the high-resolution network. The evaluation is carried out on a public dataset consisting of 26 different species in Southeast Asia. It shows high accuracy in recognition.

Keywords. Plant classification; High-resolution network; Deep learning.

# 1. INTRODUCTION

The medicinal plant and conservation of natural resources have received increasing attention in recent years and depend greatly on identifying species. Species information can provide important characteristics. The manual methods were based on human intuition which is time-consuming, imprecise, and frustrating for non-experts due to the usage of words from the botanical major. The recognition methods based on machine learning and image processing techniques have been investigated by research communities. Digital image techniques may easily capture the leaf shape and venation of plants, which can provide crucial properties for recognition. For convenience of usage, the images could be received from the built-in camera in a mobile device. They have opened research directions in plant identification based on image processing techniques, computer vision, artificial intelligence, and machine learning.

One of the approaches based on computer vision for identifying the tree class from the leaf image was introduced by Oska JOS [1]. The different descriptors that describe the different features of leaves are compared. The dataset is from a joint project between Linkoping University and the Swedish Museum of Natural History. Gaston and O'Neill [2] proposed a method to identify plants and speed up the identification process by using computer vision and search engine. Wu S et al. [3] proposed an approach for leaf recognition by using a probabilistic neural network. Its performance was evaluated on a dataset of leaves collected from the campus of Nanjing University and the Sun Yat-Sen arboretum, Nanking. Most of

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them are common plants of the Yangtze Delta, China. How an extent combination of features can improve the performance of flower classification was investigated by ZA Nilsback [4, 5] The approaches were evaluated on a challenging database of flower images. In addition, a 103-class flower dataset was introduced. An approach for the classification of leaf images with a complicated background was proposed by Sonali Agrawala et al. [6]. This method consists of three steps including leaf segmentation from the complicated background image, feature extraction of morphological and texture, and leaf recognition by using multilevel classification. The proposed method could attain an accuracy of 94%.

Recently, following the advances in deep learning the performance of image recognizers has improved considerably [7–9]. In deep learning, the abstract and composite representation can be learned from each level. In the image recognition application, the raw input images are processed through layers to extract the features before being input into a recognizer. The success of deep learning models trained with supervision is typically contributed from large datasets of annotated images. For plant recognition, some datasets are available including PlantCLEL [10–15], Pl@ntNet [16], iNaturalist [17], etc., which allowed constructing challenges for classification training and evaluation. Several models have been proposed for plant recognition. However, they are not fully developed, and the datasets used in these studies are still heavily regional, and species oriented.

In this study, we investigate plant recognition by using the high-resolution neural network, which offers semantically richer and spatially more precise representation. The experiments are validated on the Bali26 dataset with 26 different plant species collected from the Southeast Asia region [18]. The proposed approach is promising with an accuracy of 100% on the test set.

This paper is the extended version of one presented at CIIA2022 [19]. The rest of the paper consists of three sections. The related works are presented in Section 2, Section 3 describes the model for plant recognition, the experiments are presented in Section 4, and finally, Section 5 is the conclusion.

## 2. RELATED WORKS

Traditional approaches for identifying species by biologists is to use field guides and dichotomous keys. The field guides utilize images and textual descriptions of unknown species. Dichotomous keys generate a decision tree based on the features of the organism. These approaches have limitations such as difficulty in searching or questioning people who a non-expertise is. These issues have been overcome by the electronic tools including online or handheld device applications [20–22].

Early methods for automatic recognition were based on the leaf shape [23]. Methods for classifying the images of chrysanthemum leaves were proposed by Mokhtarian and Abbasi [24]. Saitoh and Kaneko [25] introduced a method based on neural networks to identify wild-flowers by using their shape and color. A method based on shape descriptions and contexts was presented by Ling and Jacobs [26]. Wang et al. [27] introduced an approach based on the centroid-contour distance combined with more standard and global descriptions of shape. Felzenszwalb and Schwarts [28] proposed a hierarchical shape algorithm for plant classification. Another method based on the leaf shape was proposed by Peter N. Belhumeur et al. [29]. Most leaf recognition methods based on the shape of scanned leaves have the

limitation in the "in the wild" scenario due to the uniform background requirements [30].

Following the advances of artificial intelligence (AI) over the last years, the performance of AI-based recognizers for world flora has improved considerably. Several deep neural network (DNN) architectures have been developed. A broad range of convolutional neural networks (CNN) and transformer-based models are evaluated to validate the recognition capabilities of different feature extractors. The commonly used baselines include ResNet-50 [31], ResNet-101, Inception-v4, and Inception-ResNet-v2. The CNN-based models can identify 10,000 plant species from Europe and North America and 10,000 species from the Guiana Shield and the Amazonia with an accuracy of approximately 90 and 40%, respectively [30]. Mads Dyrmann et al. [32] proposed a method that relied on DNN. The experiments were validated on 22 species. The transfer learning from large-scale datasets to domain-specific datasets was introduced by Cui et al. [33]. Zheng et al. [34] proposed the trilinear attention sampling network that generates attention maps relying on inter-channel relationships. Malik et al. [35] proposed an approach based on Inception, MobileNet, and ResNet architectures. In 2017, Lasseck proposed a model that relied on three architectures including GoogLeNet, ResNet-152, and ResNeXt-101-64x4d. The model can attain the best accuracy of 88.5% on PlantCLEF2017 [36]. An approach based on Inception-ResNet-v2 and Inception-v4 was proposed by Sulc and Matas [37]. It can obtain the best accuracy of 88.4% on PlantCLEF2018.

In this paper, a method for plant recognition from the Southeast Asia region is developed. The model is based on deep learning architecture with high-resolution features.

## 3. THE MODEL FOR PLANT RECOGNITION

Deep convolutional neural networks CNNs have received considerable attention and can obtain state-of-the art results in several applications. Many network architectures have been proposed for object recognition, some reputed architectures are VGG [38], GoogLeNet [39], EfficientNet [40], AlexNet [41], and ResNet [31]. These architectures offer a low-resolution representation for image recognition. The medium-resolution representation can be handled by architectures including DeconvNet [42], SegNet [43], and U-Net [44]. In some applications, the high-resolution representation may give better performance. An architecture for maintaining the high-resolution representation in image processing was proposed by J Wang et al. [45]. It has reported promising results in problems such as object detection, classification, and semantic segmentation.



Figure 1: The overall architecture for plant recognition

The overall architecture for plant recognition based on a high-resolution representation is depicted in Figure 1. It consists of multiple stages with parallelly connected streams to maintain the resolution representations. Firstly, the input image is fed into a stem consisting

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of two convolutional blocks. Each convolutional block includes a stride-2  $3 \times 3$  convolution to decrease the resolution to 1/4, its output is passed into a batch normalization and a rectified linear unit. The output from the stem is then processed by the main body of a high-resolution network which consists of four stages, each stage has three main components including transition, multi-resolution convolution, and fusion as shown in Figure 2.



Figure 2: Components of a high-resolution network [45]

The main goal of the transition component is to add a high-to-low resolution. Let  $R_{ji}$  be the representation of the *i*-th resolution index at the *j*-th stage. The representation in the next stage is determined by

$$R_{j+1i} = \begin{cases} R_{ji} & \text{if } i \le j \\ f_{j+1}(R_{jj}) & \text{otherwise,} \end{cases}$$
(1)

where,  $f_{j+1}$  is a transitional function. In this study, the transitional function is corresponding to the downsampling of the input representation R through stride-2 3×3 convolutions. Note that we can downsample at 2× size by using multiple stride-2 3×3 convolutions. An example of transition from the third stage to the fourth stage is illustrated in Figure 3.



Figure 3: Transition component from the third stage to the fourth stage. The representation in the fourth stage is determined by  $R_{4i}=R_{3i}$  for  $i \leq 3$  and  $R_{44}=f_4(R_{33})$ , where  $f_4$  is implemented by downsampling using the stride-2 3×3 convolutions.

The multi-resolution consists of multi-streams of four blocks. Each block has the architecture as shown in Figure 4, which consists of  $3 \times 3$  and  $1 \times 1$  convolutions followed by batch normalization. The streams are parallelly connected and contain the resolutions from the previous stages. Therefore, it can maintain the high-resolution representations throughout the entire process to generate reliable high-resolution representations having high position sensitivity.



Figure 4: The architecture of a block in a stream

The main goal of the fusion component is to exchange data across representations with different resolutions. The output representation at the i-th index is computed by

$$R_i = \sum_j f_j(R_j), \qquad (2)$$

where  $R_j$  is the *j*-th representation and  $f_j$  is a fusion function. The selection of this function relies on the input and output resolution, which is expressed through indexes:

- (1) If j = i,  $f_j(R) = R$ ;
- (2) If j < i,  $f_{ji}(R)$  is a function to downsample R by using i j stride-2 3×3 convolutions;
- (3) If j > i,  $f_j(R)$  is a function to upsample R by using the bilinear interpolation followed by a 1×1 convolution for aligning the number of channels.

An illustration of aggregating data from different resolutions is depicted in Figure 5. In Figure 5a, the third representation is generated by  $R_3 = f_1(R_L) + f_2(R_2) + R_3 + f_4(R_4)$ , in which  $f_1$  is implemented by 2 stride-2 3×3 convolutions,  $f_2$  is implemented by one stride-2 3×3 convolution, and  $f_4$  is implemented by the bilinear interpolation followed by a 1×1 convolution. The Figure 5b is the second representation generated by aggregating data from resolutions 1,2,3, and 4 similar to the third representation except for the difference in the number of intermediate layers for downsampling from resolution 1 and upsampling from resolutions 3 and 4.

The architecture in our implementation consists of four stages. The first stage contains four residual blocks, each block consists of a bottleneck unit with a width of 64 followed by a  $3\times3$  convolution, which resizes the width of feature maps. The number of branches in the second, third, and fourth stages respectively, is two, three, and four. Each branch includes four residual blocks including two  $3\times3$  convolutions, batch normalization, and the rectified linear unit. The number of channels corresponding to the four resolutions are C, 2C, 4C, and 8C, respectively.

# 4. EXPERIMENTAL RESULTS

We evaluate the performance of plant recognition on the Bali26 dataset [46, 47] which consists of 26 categories of ethnobotanically significant seeds, fruits, plants, and trees. The



(a) Generating the third representation



(b) Generating the second representation





Papaya (971.jpg)

Coffeearabica (920.jpg)

Frangipani (976.jpg)

Figure 6: Typical images from the Bali26 dataset

number of images is from 1000 to 2500 for each category. This dataset was generated from late February to May 2020 [18, 46]. All images in the dataset are collected from the Island of Bali in the vicinity of Ubud, specifically in the villages of Penglipuran, Kerta, Jatiluwih, Buahan, Sekaan, and Bayung Gede. In comparison to the ImageNet dataset, the images in Bali26 have a higher information richness of the deliriously lush flora. The collection process was performed by a team of members collecting data in the wild, in a constrained location, and over a short period. The images with the size of  $1080 \times 1920$  in JPG format were captured by data collectors in the field using high-definition mobile phones. Some typical images are shown in Figure 6, and a summary of plant categories including common name, ethnobotany, and scientific name is given in Table 1.

	Common name	Fthnobotany	Scientific	
No.		Ethnobotany	name	
1	Bamboo	Cooked shoots could be added to vegetable soups	Gigantochloa	
			apus	
2	Banana	The core stems could be added to vegetable soups	Musa x paradisi-	
			aca	
3	Cacao	Powder of seeds powder can be used for inducing	Theobroma	
		relaxation	cacao	
4	Coffee ara- bica Dragon fruit	Powder of seeds can be used for inducing relaxation	Coffea canephora	
		Ding fruit goton fresh, Fruits can be used for me		
5		Ripe fruit eaten fresh; Fruits can be used for pro-	Hylocereus	
		Cessing of religious offerings	costaricensis	
6	Durian	Fruits can be used for processing or religious oner-	Durio zibethinus	
	Flophant foot	nhant fact		
7	Elephant loot	Tuberous roots can be used as staple food	nnorphophanus	
	Frangipani	The flowers can be used in religious offerings: The	Plumeria alba	
8		Flower juice can be applied to the skin for smallpox		
	Guava	Ripe Fruit can be used in religious offerings or	Psidium guajava	
9		eaten fresh.		
	Indonesian cinnamon	Decoction of leaves and barks can be used for hy-		
10		pertension, fever, heartburn, sore throat, cough,	Cinnamomum	
10		and to stimulate the appetite; Barks can be used	burmanii	
		as spices.		
11	Jackfruit	Leaf decoction can be used for diarrhea; Cooked	Artocarpus het-	
11		fruit and seeds can be added to vegetable soups.	erophyllus	
12	Lychee	Fruits can be used eaten fresh, for drying, or for	Litchi chinensis	
		religious offerings.		
13	Mango	Decoction of leaves can be used for hypertension	Mangifera indica	
		and diabetes		
14	Mangosteen	Ripe fruit can be eaten fresh, or used in religious	Garcinia man-	
14		offerings	gostana	

Table 1: A summary of plant categories from the Bali26 dataset [19]

15	Nilam	The leaves can be used for rheumatic, dysentery,	Pogostemon ca-		
		headache, and as diuretic	blin		
16	Papaya	Young leaves and cooked fruit can be added to veg-	Carico nonoro		
		etable soups	Carica papaya		
17	Passiflora	Ripe fruit can be eaten fresh	Passiflora edulis		
18	Sawo	Ripe Fruit can be used in religious offerings or	Manilkara za-		
		eaten fresh	pota		
19	Snake fruit	Ripe Fruit can be used in religious offerings or	Salacca zalacca		
		eaten fresh			
20	Star fruit	Juice of leaves can be used for heartburn; Cooked	Averrhoa caram-		
		fruit and leaves can be added to vegetable soups	bola		
21	Sugarplum	Boiled fruit is edible; Boiled inner stems can be			
		eaten as a staple food; Root decoction can be used	Arenga pinnata		
		for urolithiasis; Leaves can be used in religious of-			
		ferings			
22	Taro	Cooked leaves can be added to vegetable soups;	Colocasia escu-		
		Boiled tuberous roots can be eaten as a staple food.	lenta		
23	Vanilla	Fruit powder can be used as vanilla flavoring	Vanilla planifolia		
24	Water guava	Ding fruit can be got on fresh	Syzygium		
		Ripe fruit can be eaten fresh	aqueum		
25	White pep-	Seeds powder can be used as spice	Piper nigrum		
	per		- ipor ingruin		
26	Zodiac	Leaves can be used for mosquito repellent	Evodia sauve-		
		heaves can be abed for mosquito rependit	olens		

Table 2: The recognition results from the testing set

No.	Category	# Images	No.	Category	# Images
1	Cinnamon	136	14	Suweg	132
2	Dragon fruit	234	15	Sawo	136
3	Jack fruit	297	16	Banana	191
4	Mango	149	17	Mangosteen	180
5	Nilam	138	18	Durian	271
6	Papaya	183	19	Snake fruit	130
7	Passiflora	151	20	Bamboo	214
8	Vanilla	217	21	Coffee arabica	149
9	Water guava	205	22	Lychee	124
10	White pepper	157	23	Sugar palm	140
11	Star fruit	189	24	Frangipani	174
12	Taro	177	25	Guava	215
13	Zodiac	148	26	Cacao	187

The proposed scheme was implemented by using Python 3.7 programming language and PyTorch machine learning framework on Ubuntu 18.04 operating system. The hardware

configuration of CPU Intel® Xeon® Quad-core 2.9 GHz, RAM of 16GB, GPU GeForce GTX 2080ti with 16GB, and HDD of 1TB was used. The hyperparameters were selected based on try-and-error and previous publications. The Adam optimizer was used. The batch size, learning rate, momentum, and epochs were 32, 0.05, 0.9, and 50, respectively.

The model was pretrained on the ImageNet dataset, then trained on the Bali26 dataset. The performance is evaluated on the criterion of accuracy. The testing set corresponding to 26 categories is given in Table 2. The overall accuracy is highly impressive, which is 100% for the testing set.

The high-resolution input images can require the high-resolution network model which may result in more memory. There are several models for identifying plants, they are evaluated on different datasets. The DNN model proposed by Mads Dyrmann et al. (Dyr16) was evaluated on 22 species. The architectures based on GoogLeNet, ResNet-152, and ResNeXt-101-64×4d can attain the best accuracy of 88.5% on PlantCLEF2017. The model based on Inception-ResNet-v2 and Inception-v4 can attain the best accuracy of 88.4% on Plant-CLEF2018. In our study, most of the features in the dataset are fruits, leaves, and whole trees. The model based on the high-resolution architecture can attain an impressive accuracy of 100% on the test set of Bali26.

# 5. CONCLUSIONS

Identifying plant information plays an important role in not only agriculture but also medicine. Several models have been developed relying on image processing techniques and machine learning. They can attain promising results. However, they are validated on datasets collected from some specific regions. In this paper, an approach based on the high-resolution network is proposed and evaluated on a dataset consisting of 26 different species in Southeast Asia. The accuracy of the testing set is impressive. By training the model with more datasets for plant identification, we believe that the trained model can contribute to a variety of applications, especially in the pharmaceutical industry. In addition, plant identification may contribute to disease detection in plants to help solve agricultural problems.

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