

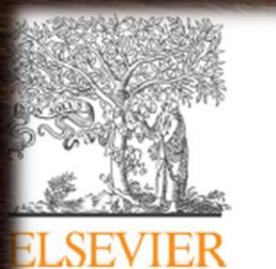
Seed-per-pod estimation for plant breeding using deep learning

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Paper



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Original papers

Seed-pod estimation for plant breeding using deep learning

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Fonte: Uzal *et. al.*, 2018.

Plant phenotyping



Fonte: Agriexpo, 2019.



Fonte: Embrapa, 2016.

Three major components of a soybean crop:

Pods per plant (PN)



Seeds per pod (SPP)



Seed size (SS)



Fonte: Technologytimes, 2014.

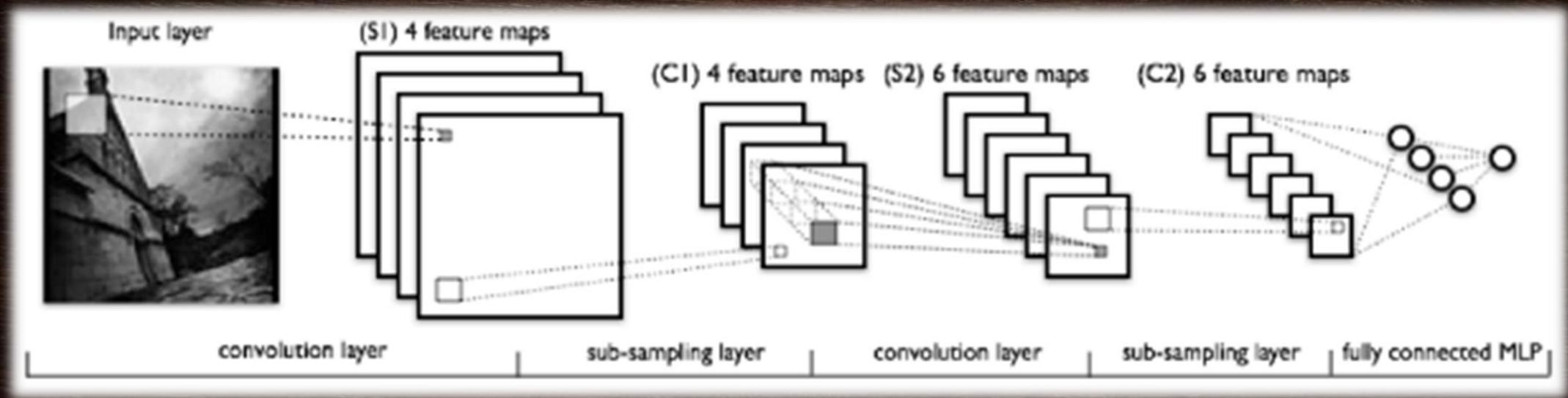
Fonte: Sistema Faep, 2018.

Fonte: Maissoja, 2018.

Classic procedure

- “Define and extract appropriate features for the problem at hand and then train a classifier” (UZAL, 2018)
- Three state-of-the-art classic classification methods were implemented:
 - SVM
 - Random Forest (RF)
 - Penalized Discriminant Analysis (PDA)

Convolutional neural networks



Fonte: Ravindra, S., 2017.

Data collection

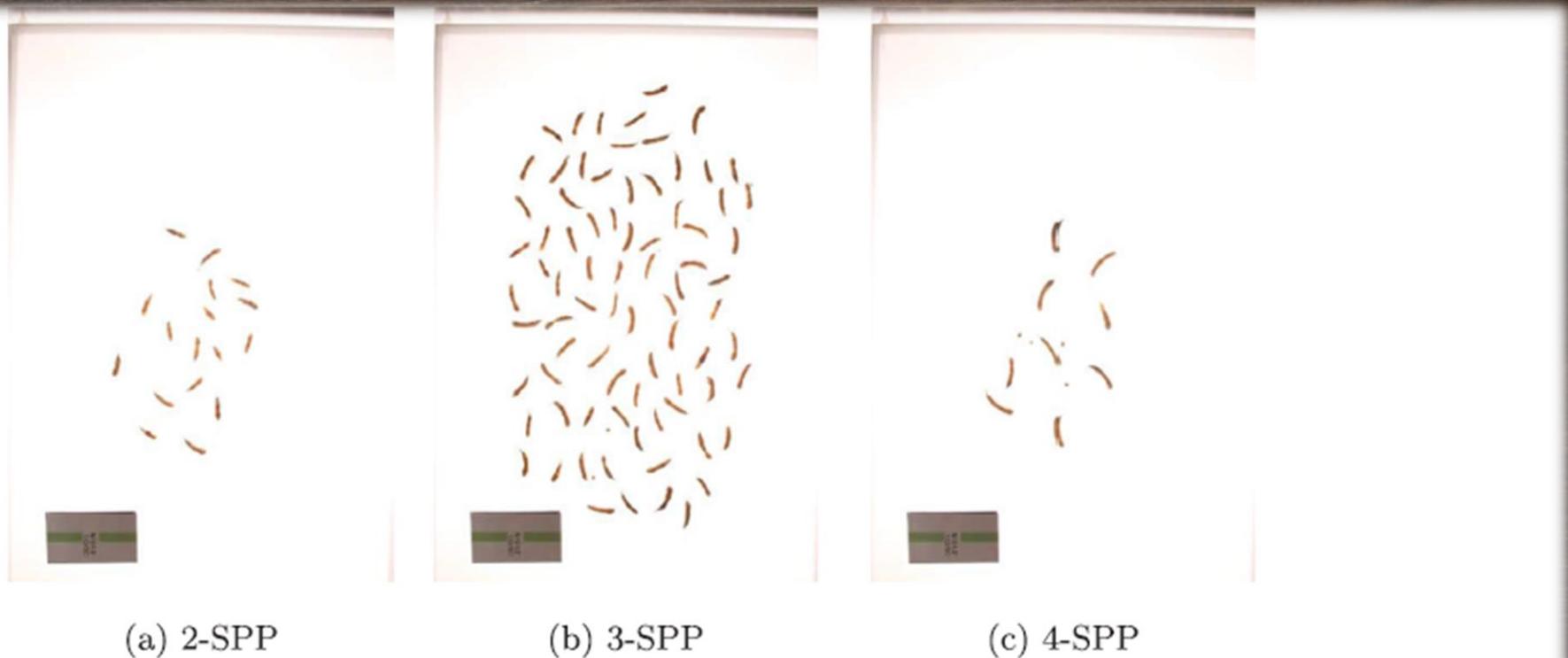
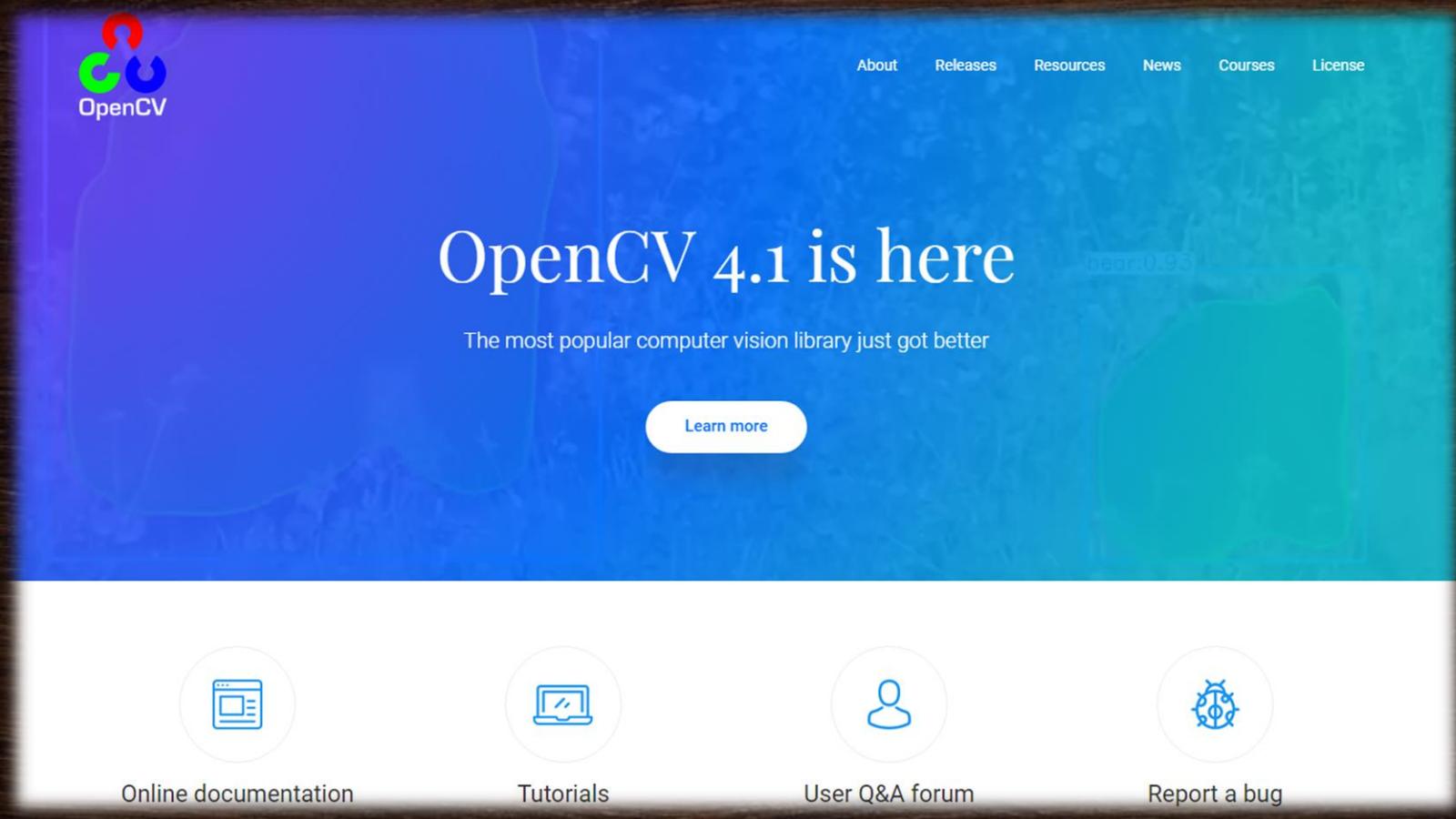


Fig. 1. Sample photographs of soybean pods used to build the dataset. Each pod is manually classified by an expert and photographed within its class group defined by the SPP number.

Fonte: Uzal *et. al.*, 2018.

Pods segmentation



Fonte: OpenCV, 2019.

Pods segmentation



Fig. 1. Sample photographs of soybean pods used to build the dataset. Each pod is manually classified by an expert and photographed within its class group defined by the SPP number.

Fonte: Uzal *et. al.*, 2018.

Pods segmentation

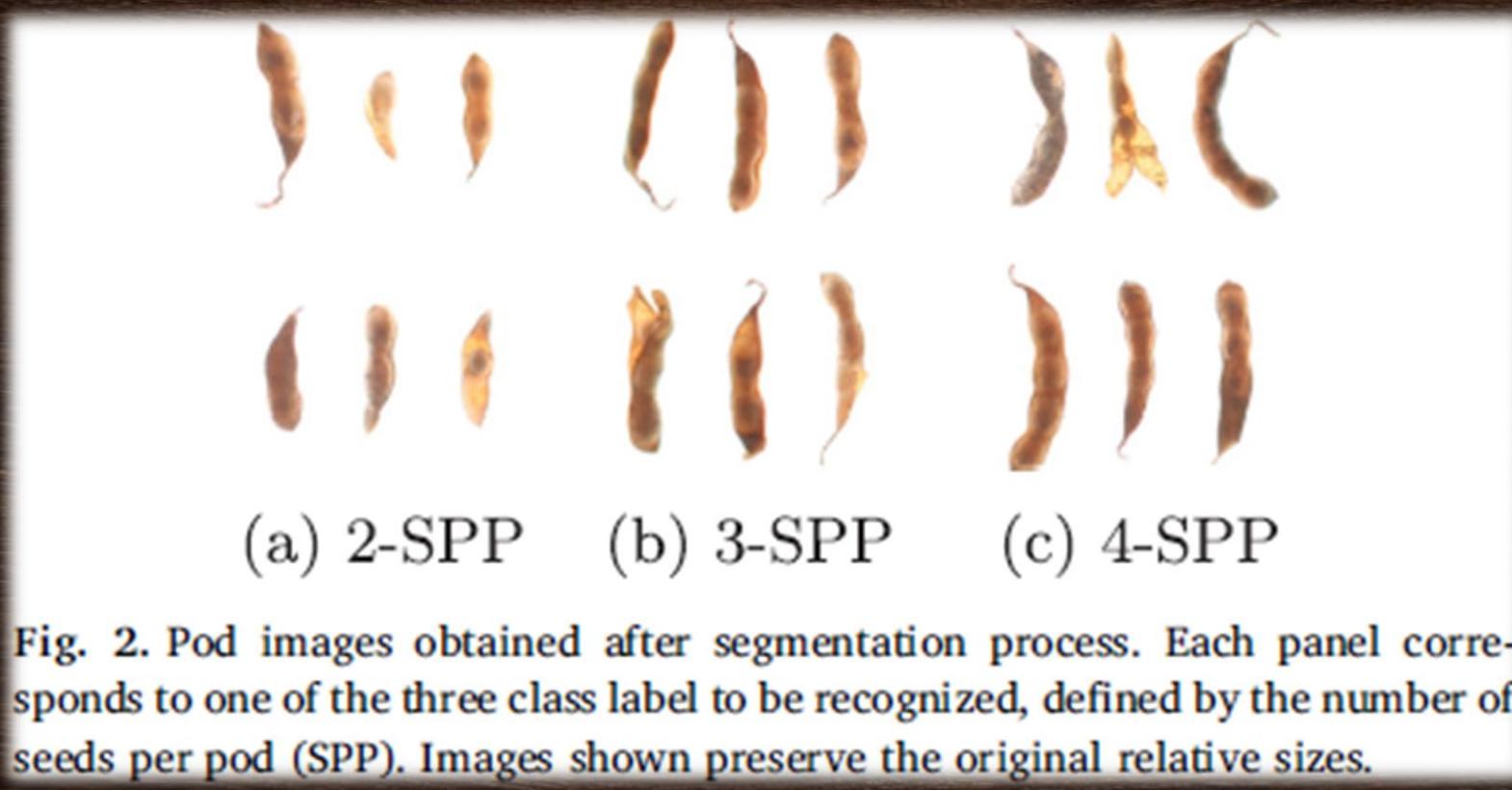


Fig. 2. Pod images obtained after segmentation process. Each panel corresponds to one of the three class label to be recognized, defined by the number of seeds per pod (SPP). Images shown preserve the original relative sizes.

Fonte: Uzal *et. al.*, 2018.

Pods segmentation

Table 1

Total number of examples corresponding to each class and season.

Class	Season 1	Season 2
2-SPP	811	3746
3-SPP	4598	5075
4-SPP	2444	1504
Total	7853	10325

Fonte: Uzal *et. al.*, 2018.

Preprocessing

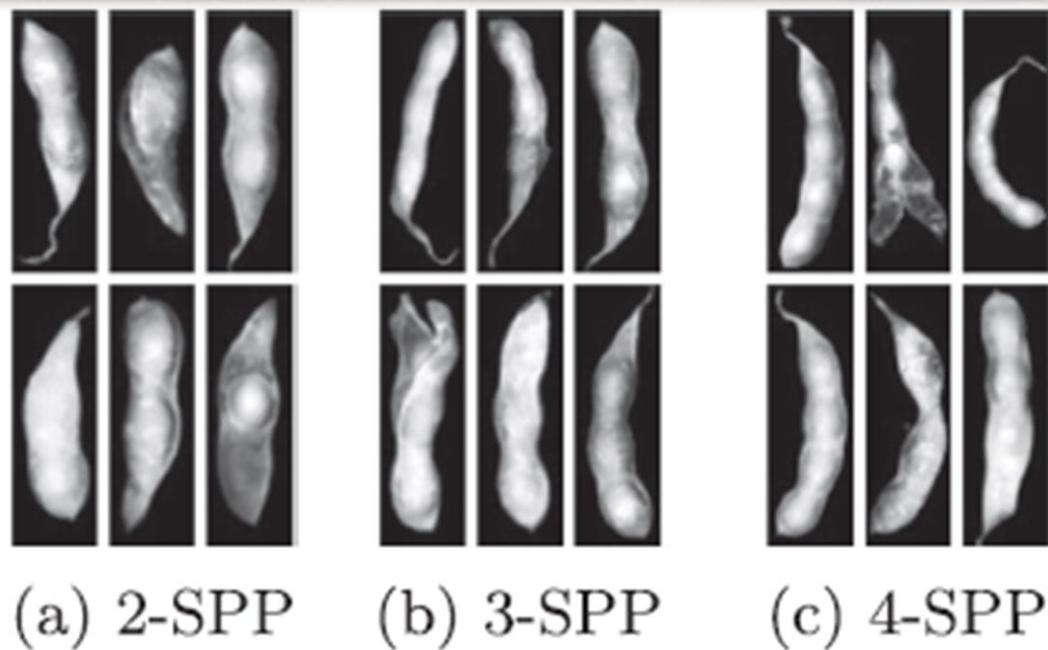


Fig. 3. Images obtained after preprocessing step. Samples are the same of Fig. 2.

Fonte: Uzal *et. al.*, 2018.

Data augmentation

The screenshot shows a red header bar with the Keras Documentation logo and a search bar. Below it is a sidebar with navigation links for Home, Why use Keras, GETTING STARTED, Guide to the Sequential model, Guide to the Functional API, FAQ, MODELS, LAYERS, and various layer types like About Keras models, Sequential, Model (functional API), etc. At the bottom of the sidebar are GitHub and navigation links for the previous and next pages. The main content area has a breadcrumb trail: Docs > Preprocessing > Image Preprocessing. It features a "Edit on GitHub" button and a large title "Image Preprocessing". Below that is a section for the "ImageDataGenerator class" with a "[source]" link. A code snippet shows the class definition:

```
keras.preprocessing.image.ImageDataGenerator(featurewise_center=False, samplewise_center=False, featurewise_std_normalization=False, samplewise_std_normalization=False, zca_epsilon=1e-06, zca_whitening=False, rotation_range=0, width_shift_range=0, height_shift_range=0, fill_mode='nearest', cval=0.0, horizontal_flip=False, vertical_flip=False, rescale=None, preprocessing_function=None, data_format=None)
```

. A description follows: "Generate batches of tensor image data with real-time data augmentation. The data will be looped over (in batches)." The "Arguments" section lists parameters with their descriptions:

- `featurewise_center`: Boolean. Set input mean to 0 over the dataset, feature-wise.
- `samplewise_center`: Boolean. Set each sample mean to 0.
- `featurewise_std_normalization`: Boolean. Divide inputs by std of the dataset, feature-wise.
- `samplewise_std_normalization`: Boolean. Divide each input by its std.
- `zca_epsilon`: epsilon for ZCA whitening. Default is `1e-6`.
- `zca_whitening`: Boolean. Apply ZCA whitening.
- `rotation_range`: Int. Degree range for random rotations.
- `width_shift_range`: Float, 1-D array-like or int
 - float: fraction of total width, if < 1, or pixels if ≥ 1 .
 - 1-D array-like: random elements from the array.
 - int: integer number of pixels from interval `(-width_shift_range, +width_shift_range)`
- With `width_shift_range=2` possible values are integers `[-1, 0, +1]`, same as with `width_shift_range=[-1, 0, +1]`, while with `width_shift_range=1.0` possible values are floats in the half-open interval `[-1.0, +1.0[`.
- `height_shift_range`: Float, 1-D array-like or int

Fonte: Keras, 2019.

Data augmentation

The screenshot shows the official website for GIMP (GNU Image Manipulation Program). The header features a dark green navigation bar with links for GIMP, DOWNLOAD, NEWS, ABOUT, DOCS, PARTICIPATE, TUTORIALS, and DONATE (with icons for PayPal, Credit Card, and Bitcoin). Below the header is a large banner with a blue gradient background showing a stylized cartoon dog holding a paintbrush. The word "GIMP" is written in large white letters, with "GNU IMAGE MANIPULATION PROGRAM" in smaller text below it. A red button with a download icon and "DOWNLOAD 2.10.10" is on the left, and a "RELEASE NOTES" button is on the right. The main content area has a white background. On the left, there's a section titled "The Free & Open Source Image Editor" with text about the software being free and open source, available for various operating systems, and suitable for graphic design, photography, illustration, and science. On the right, there's a "Recent News" section with three items: "GIMP 2.10.10 Released" (date: 2019-04-07), "GIMP and GEGL in 2018" (date: 2019-01-02), and "GIMP 2.10.8 Released" (date: 2018-11-08).

The Free & Open Source Image Editor

This is the official website of the GNU Image Manipulation Program (GIMP).

GIMP is a cross-platform image editor available for GNU/Linux, OS X, Windows and more operating systems. It is free software, you can change its source code and distribute your changes.

Whether you are a graphic designer, photographer, illustrator, or scientist, GIMP provides you with sophisticated tools to get your job done. You can further enhance your productivity with GIMP thanks to many customization options and 3rd party

Recent News

GIMP 2.10.10 Released
2019-04-07

GIMP and GEGL in 2018
2019-01-02

GIMP 2.10.8 Released
2018-11-08

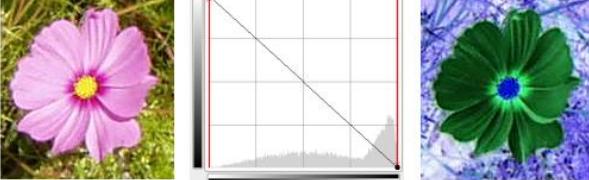
GIMP receives a \$100K donation
2018-08-30

Fonte: Gimp, 2019.

Data augmentation

8.11.3.2. Practical cases

Invert colors

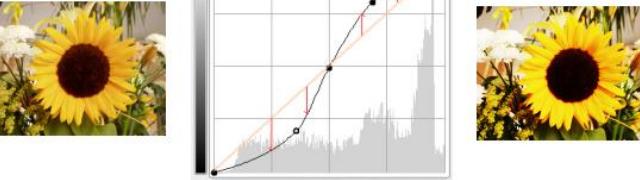


The image shows a pink flower on the left and its inverted color version on the right. The inverted version has a greenish-blue hue.

Inverted curve

Black is made White (fully colored / fully opaque). White is made black (black, fully transparent). All pixels adopt the complementary color. Why that? Because subtracting the channel values from 255 gives the complementary color.
For example: 19;197;248 a sky blue gives 255-19; 255-197; 255-248 = 236;58;7, a bright red.

Enhance contrast



The image shows a sunflower on the left and its enhanced contrast version on the right. The enhanced contrast version has more saturated colors and higher contrast.

Fonte: Gimp, 2019.

Feature extraction details

- Geometrical characteristics
 - Area
 - Perimeter
 - Major and minor axis length
- Shape features
 - Density
 - Elongation
 - Compactness
 - Rugosity
 - Axis ratio
- Etc

SVM implementation details

The screenshot shows the official scikit-learn website. At the top, there's a navigation bar with links for Home, Installation, Documentation, Examples, Google Custom Search, and a search icon. A "Fork me on GitHub" button is located in the top right corner. The main content area features a grid of nine small plots illustrating various machine learning models. Below this, there are six main sections: Classification, Regression, Clustering, Dimensionality reduction, Model selection, and Preprocessing. Each section provides a brief description, applications, algorithms, and examples. The "Classification" section includes a "Recent Models" gallery.

Classification
Identifying to which category an object belongs to.
Applications: Spam detection, Image recognition.
Algorithms: SVM, nearest neighbors, random forest, ...
— Examples

Regression
Predicting a continuous-valued attribute associated with an object.
Applications: Drug response, Stock prices.
Algorithms: SVR, ridge regression, Lasso, ...
— Examples

Clustering
Automatic grouping of similar objects into sets.
Applications: Customer segmentation, Grouping experiment outcomes
Algorithms: K-Means, spectral clustering, mean-shift, ...
— Examples

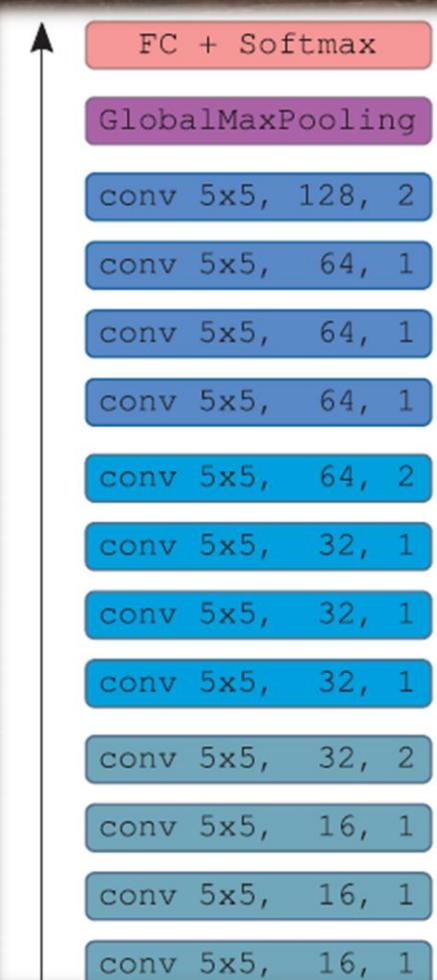
Dimensionality reduction
Reducing the number of random variables to consider.
Applications: Visualization, Increased efficiency
Algorithms: PCA, feature selection, non-negative matrix factorization.
— Examples

Model selection
Comparing, validating and choosing parameters and models.
Goal: Improved accuracy via parameter tuning
Modules: grid search, cross validation, metrics.
— Examples

Preprocessing
Feature extraction and normalization.
Application: Transforming input data such as text for use with machine learning algorithms.
Modules: preprocessing, feature extraction.
— Examples

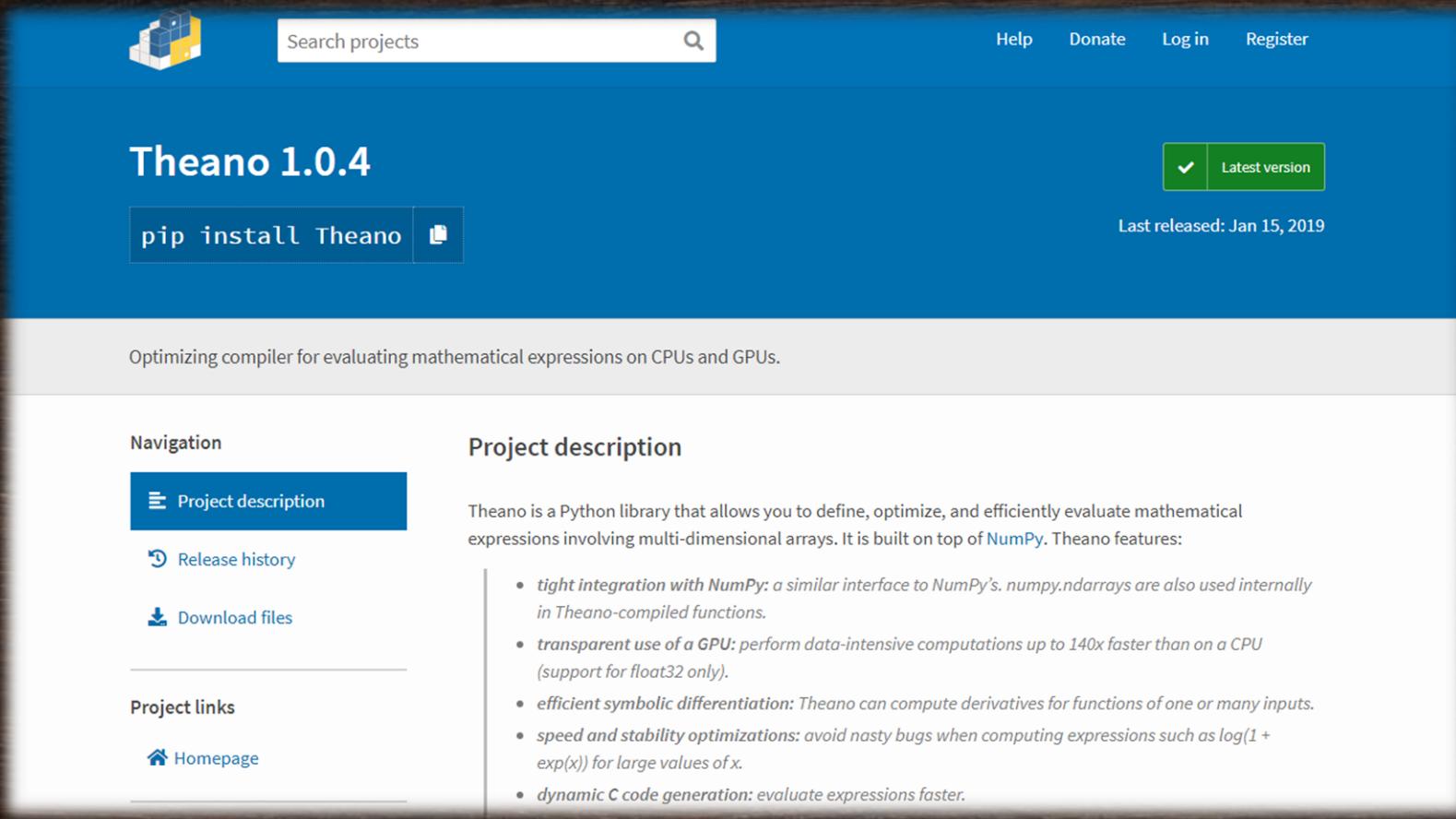
Fonte: Scikit-learn, 2019.

CNN implementation details



Fonte: Uzal *et. al.*, 2018.

CNN implementation details



Theano 1.0.4

pip install Theano

Last released: Jan 15, 2019

Optimizing compiler for evaluating mathematical expressions on CPUs and GPUs.

Navigation

- Project description**
- Release history
- Download files

Project links

- Homepage

Project description

Theano is a Python library that allows you to define, optimize, and efficiently evaluate mathematical expressions involving multi-dimensional arrays. It is built on top of NumPy. Theano features:

- tight integration with NumPy: a similar interface to NumPy's. numpy.ndarrays are also used internally in Theano-compiled functions.*
- transparent use of a GPU: perform data-intensive computations up to 140x faster than on a CPU (support for float32 only).*
- efficient symbolic differentiation: Theano can compute derivatives for functions of one or many inputs.*
- speed and stability optimizations: avoid nasty bugs when computing expressions such as $\log(1 + \exp(x))$ for large values of x .*
- dynamic C code generation: evaluate expressions faster.*

Fonte: Theano, 2018.

CNN implementation details

The screenshot shows the Lasagne documentation website. The top navigation bar includes a logo, the word "Lasagne", and a "latest" dropdown. A search bar labeled "Search docs" is present. The sidebar on the left lists various documentation sections: Installation, Tutorial, Layers, Creating custom layers, Development, and several sub-sections under lasagne.layers, lasagne.updates, lasagne.init, lasagne.nonlinearities, lasagne.objectives, lasagne.regularization, lasagne.random, and lasagne.utils. At the bottom of the sidebar is a "YOUR AD HERE" placeholder for an advertisement, which includes the text "Reach over 7 million devs each month when you advertise with Read the Docs." and "Sponsored - Ads served ethically". The main content area shows the "Welcome to Lasagne" page, which states that Lasagne is a lightweight library to build and train neural networks in Theano. It mentions that the project is still in progress and welcomes input. Below this is a "User Guide" section that explains how to install Lasagne, build neural networks, and contribute as a developer. The user guide also lists sub-sections such as Installation, Tutorial, Layers, and so on.

Docs » Welcome to Lasagne

Edit on GitHub

Welcome to Lasagne

Lasagne is a lightweight library to build and train neural networks in Theano.

Lasagne is a work in progress, input is welcome. The available documentation is limited for now. The project is on [GitHub](#).

User Guide

The Lasagne user guide explains how to install Lasagne, how to build and train neural networks using Lasagne, and how to contribute to the library as a developer.

- Installation
 - Prerequisites
 - Stable Lasagne release
 - Bleeding-edge version
 - Development installation
 - GPU support
 - Docker
- Tutorial
 - Before we start
 - Run the MNIST example
 - Understand the MNIST example
 - Where to go from here
- Layers
 - Creating a layer
 - Creating a network

Fonte: Lasagne, 2019.

Hyperparameter search

Table 2

Explored hyperparameters and selected values (see Section 4) for SVM, CNN and Data Augmentation.

Method	Hyperparameter	Range	Selected Valu	Description
SVM	C	[0.5, 1, 3, 5, 10, 50, 100, 200, 1000]	10	SVM C parameter
	gamma	[50, 20, 14, 10, 5, 2, 1, 0.5, 0.2, 0.1, 0.05]	10	Gaussian kernel gamma
CNN	blockSize	[1–4]	4	Layers per block
	nBlocks	[1–5]	3	Number of blocks
	widthFactor	[8, 16, 32, 48, 64, 96, 128]	16	Multiplicative factor for the number of maps
	log10lr	[-5.0 to -2.0]	-2.43	Learning rate (log10 scale)
	log10wd	[-5.0–0.0]	-1.10	Weight decay (log10 scale)
	batchSize	[16, 32, 48, 64, 96, 128]	128	Samples per minibatch
Data Augm.	zoom	[0.9–1.1]	0.97	Zoom range center
	zoomRange	[0.0–0.25]	0.18	Amplitude of zoom interval
	shear	[0.0–0.35]	0.14	Maximum shear angle (radians)
	wShift	[0.0–10.0]	2.8	Maximum horizontal shift (%)
	hShift	[0.0–10.0]	1.6	Maximum vertical shift (%)
	curves	[0.0–0.75]	0.58	Maximum curve strength
	rot	[0–20]	20	Maximum random rotation (degrees)

Fonte: Uzal *et. al.*, 2018.

Results

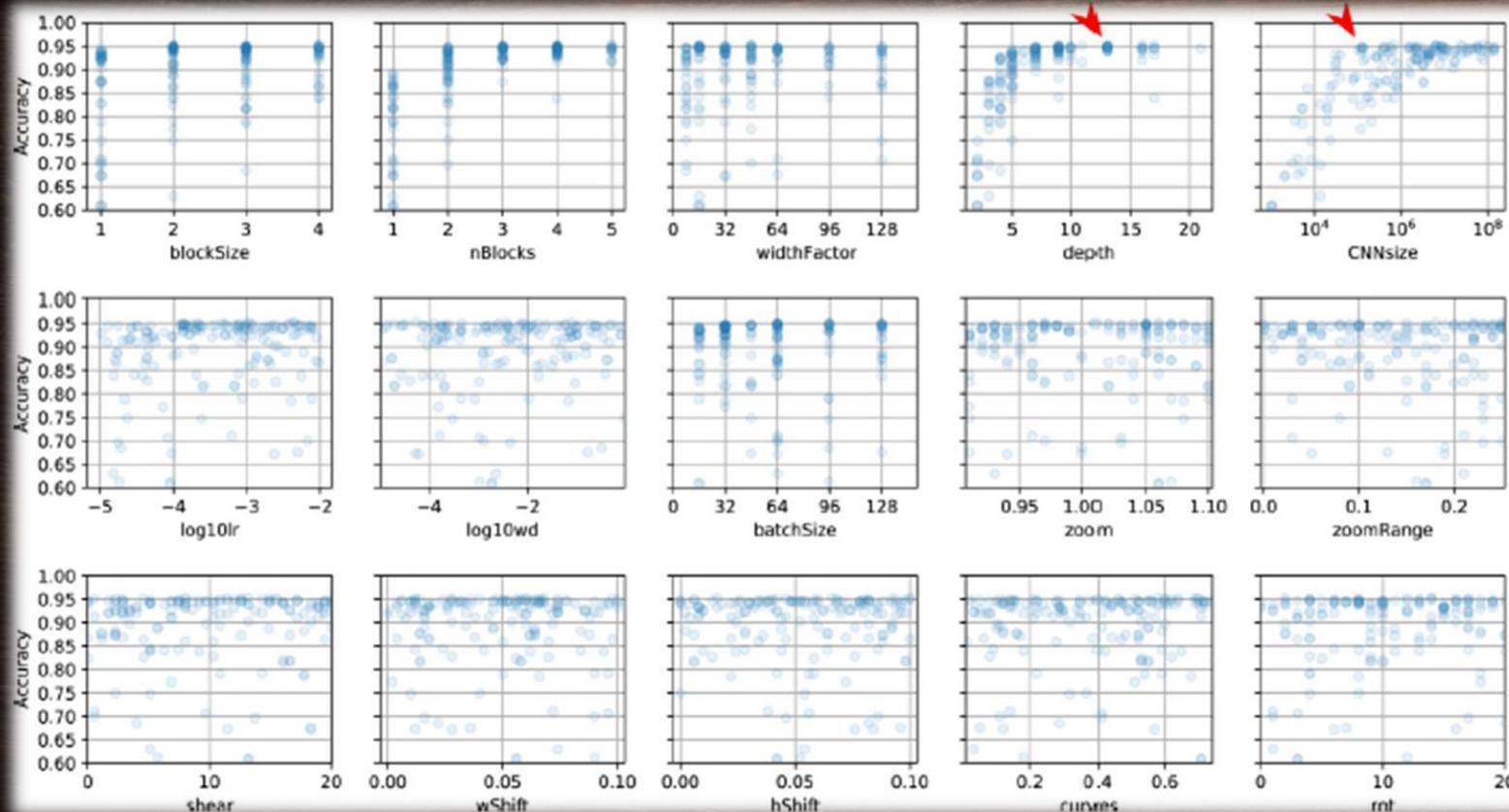
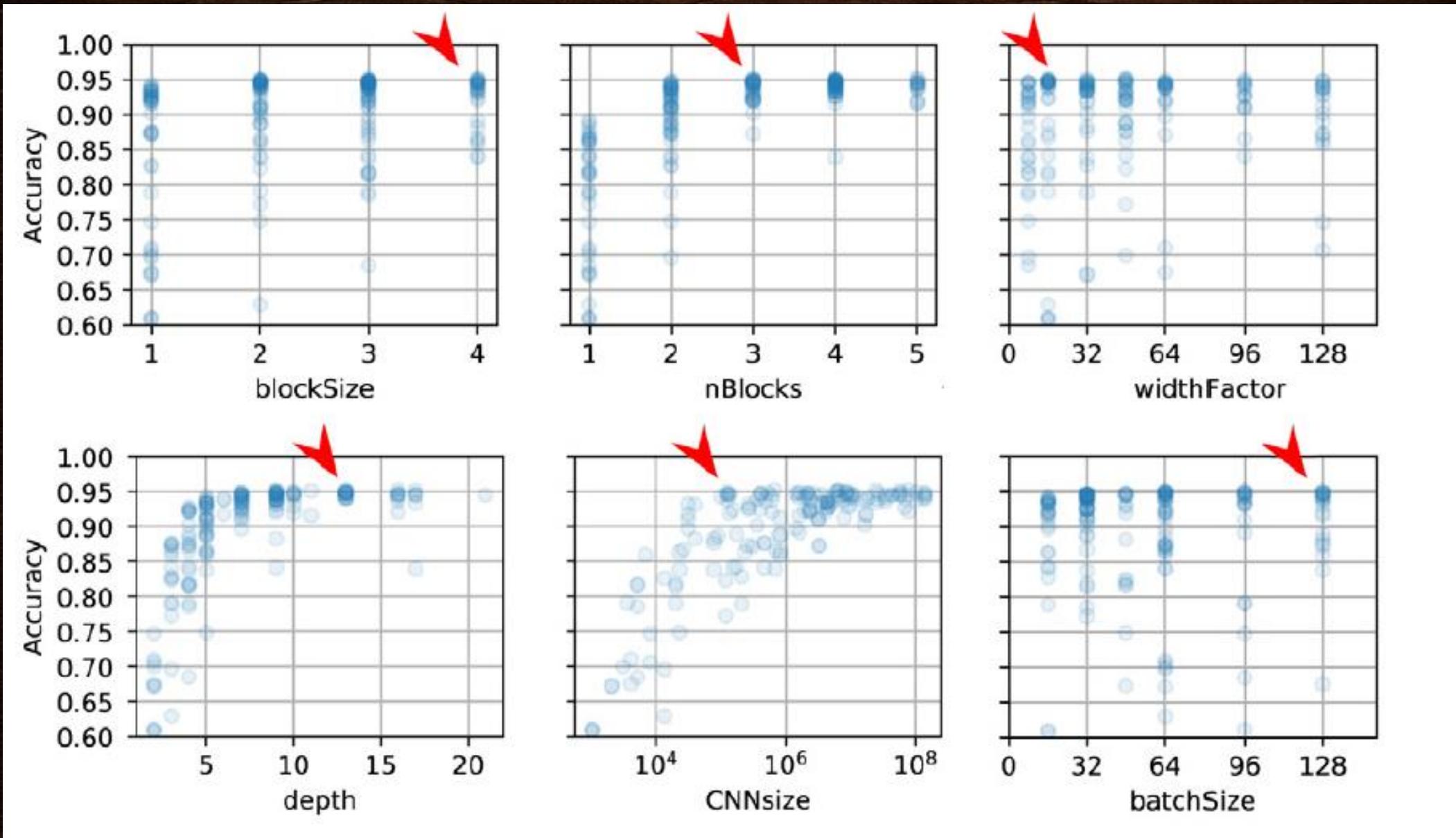


Fig. 5. Random search of hyperparameters for CNN training. The first row corresponds to model hyperparameters (defining network architecture). Variables `depth` and `CNNsize` are significative quantities derived from hyperparameters `blockSize`, `nBlocks`, and `widthFactor`. The rest of the panels corresponds to training algorithm and data augmentation hyperparameters (see Table 2 for details). Validation accuracy is almost insensitive to these training and data augmentation parameters. In order to reach high accuracies (above 90%), the only thing needed is to take a deep enough, high capacity network. Red arrows show the model selected which is a tradeoff between maximizing accuracy and minimizing model size.

Fonte: Uzal *et. al.*, 2018.

Relevant hyperparameters



Results

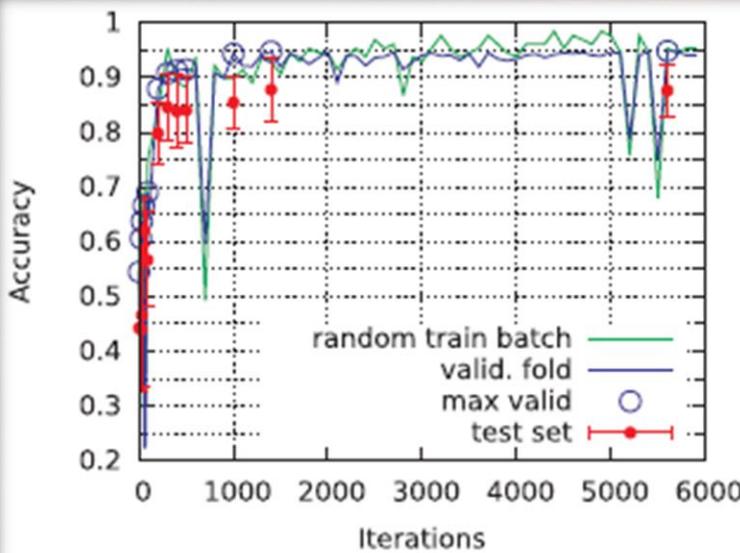
Table 3

Accuracy for different methods trained on Season 1 data and tested on Season 2 data. Mean and deviation for validation accuracies were computed with a group k-fold procedure over training data. Test accuracy mean and standard deviation were computed over test session groups and averaged over k-fold models.

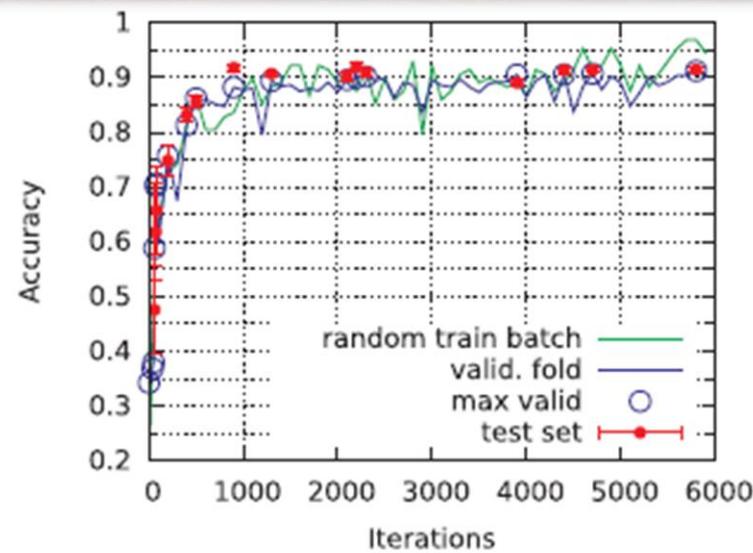
Method	Valid. Accuracy	Test Accuracy
Features + SVM	0.902 ± 0.022	0.504 ± 0.145
CNN without Data Augmentation	0.936 ± 0.009	0.827 ± 0.043
CNN with Data Augmentation	0.951 ± 0.005	0.862 ± 0.052

Fonte: Uzal *et. al.*, 2018.

Results



(a) Train set: Season 1. Test set: Season 2



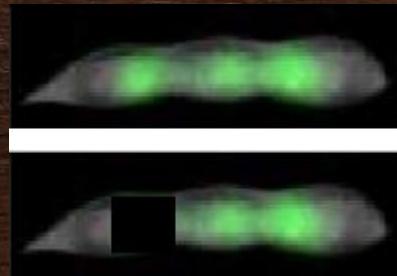
(b) Train set: Season 2. Test set: Season 1

Fonte: Uzal *et. al.*, 2018.

Fig. 6. Training curves for the two different seasons. Green lines: accuracy computed over a random minibatch during training. Blue lines: accuracy computed on the validation fold. Blue empty circles: new maximum in validation accuracy. Red dots: test accuracy. Season 2 dataset appears to contain more diverse and difficult examples which do not exist in the Season 1 version. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

Detection of important features

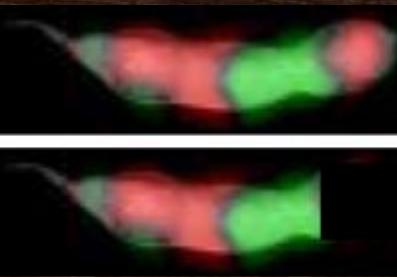
Original
Occluded



Seed per pod

3 (true and predicted)
2

Original
Occluded



Seed per pod

4 (prediction)
3 (true)

Fonte: Uzal *et. al.*, 2018.

Visualizing relevant patterns

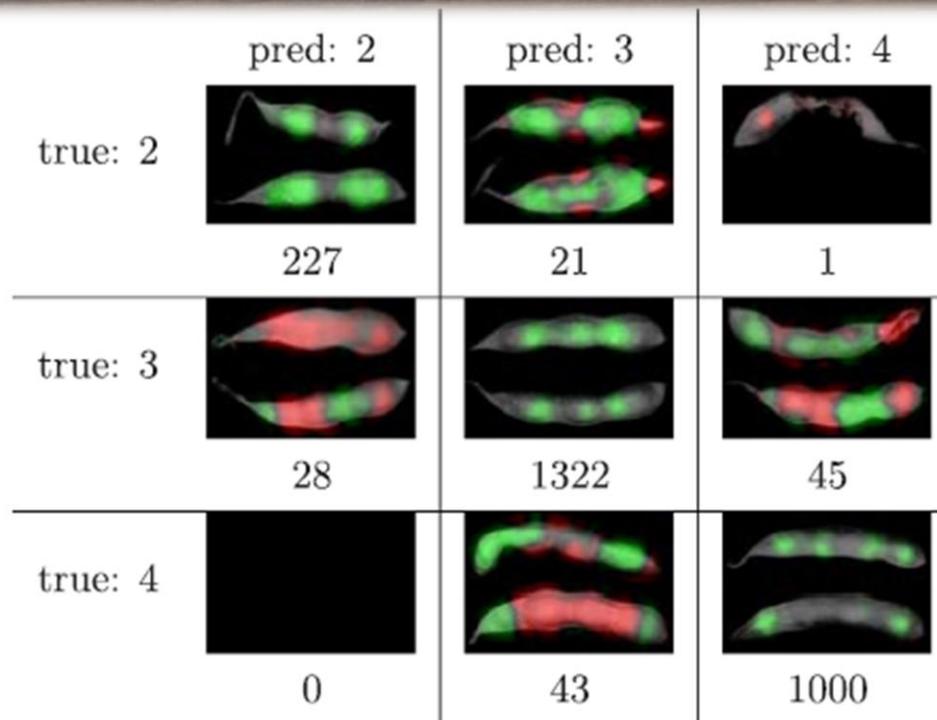
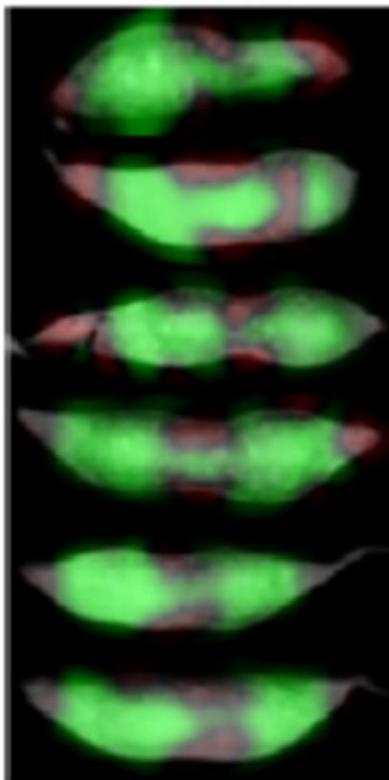


Fig. 7. Confusion matrix with representative samples visualization. Green (red) colored regions indicates regions of positive (negative) correlation with correct class CNN output probability obtained by occlusion experiments. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

Fonte: Uzal *et. al.*, 2018.

Visualizing relevant patterns

true: 2



true: 3

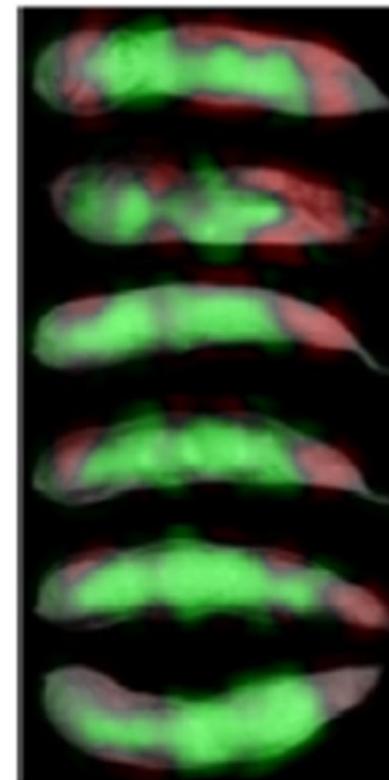


Fig. 8. Visualization of samples where the contour plays an important role. The colors indicate the same as in Fig. 7. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

Conclusion and future works

- CNNs have many hyperparameters, but high accuracy can be achieved without precise values
- CNNs outperformed classic approach
- CNNs learnt to detect each seed in the pod
- Detection in field

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- UZAL, L.C. *et. al.* Seed-per-pod estimation for plant breeding using deep learning. Computers and Electronics in Agriculture, v.150, p. 196-204, 2018.