Self-learning software to identify slipper orchids from pictures of flowers to simplify identification for customs officers

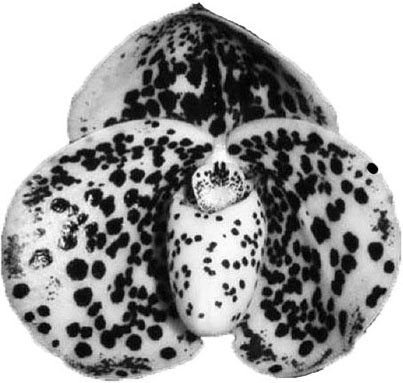


Figure 1 *Brachypetalum bellatulum* [1]

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# Abstract

Slipper orchids are very popular ornamentals among hobbyists and therefore highly endangered due to over collecting in the wild. To prevent them from going extinct all slipper orchids were placed on the international CITES list, which means that permits are required for trade all over the world. Some slipper orchid species are more endangered than others. This is reflected by different legislation as some species are placed on CITES Appendix 1 (highest level of protection) whereas others are placed on Appendix 2 (lower level of protection). Flowers of slipper orchids can be classified by characteristics like shape of the petals, the absence or presence of wards on the petals, and the colour of the lip. During this internship the concept of classifying slipper orchids to section level from pictures was tested. To test the possibilities of picture-based classification, pictures of well-identified orchids were collected. From these, different sets were created, one for training Artificial Neural Networks (ANNs) and one for testing them. The results show that some sections were harder to classify than others. Adding new pictures to the training set could improve the classification, but could also worsen the classification, depending on the quality of the new pictures and how well the new networks were trained. In general, better-trained networks, trained with more than a single picture, were able to correctly classify slipper orchid flowers, showing that visual recognition software can be used for identification of these orchids.

To make the classification software available to the outside world, a first concept of a website was developed. Users can upload a picture of a slipper orchid flower to the website. The website then runs the classification software using a pre-trained ANN. The result of this classification is then translated to a human readable result. This result is returned to the website. With this application, customs officers can check whether the permits of imported slipper orchids are correct. As compared to classification using a book, the website improves the identification by laymen, which will help customs officers to prevent illegal trade of these orchids.

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# Introduction

## Illegal trade in endangered orchids

There are thousands of different orchid species known all over the world [2]. None of these are allowed to be imported into the Netherlands without CITES permits. Since 1973, orchids are primarily protected by the Convention on International Trade in Endangered Species of Wild Flora and Fauna (CITES), which is signed by over 120 nations [3-4]. Despite this convention many orchids are traded illegally. To trade species that are protected by CITES, a licence or certificate is required.

It is difficult to monitor the illegal trade of orchids because some orchids look very similar to non-protected plants and so accurate identification can be very difficult. To improve identification, software that can identify orchids from pictures of tubers, leaves or flowers could be vital. This software would in particular aid customs officers and employees of nature conservation organizations involved in confiscating illegally-traded material. During this project, the focus was on slipper orchids and orchids from which *salep* is produced.

## Slipper orchids

In Europe and Asia the slipper orchids (Cypripedioideae) are widely distributed between sea level up to 2000 m altitude. They prefer to live in calcareous environments and are found in deciduous or mixed deciduous and coniferous woods. They grow best in light to deep shade. The slipper orchid is an herbaceous perennial plant species that has a long lifespan. It can grow up to 60 cm and each season the slipper orchid will produce new growths. Each stem of the orchid can contain 3 to 4 leaves that often have upwardly curved sides. The flower stalk can be one-flowered or two-flowered with leaf-like bracts. The sepals and petals are rarely green but commonly brightly coloured. These sepals and petals are also often twisted [5]. Slipper orchids are highly desired ornamentals, which is why they were placed on CITES Appendices, which means they are protected from unregulated export. Different levels of protection exist: the genera *Paphiopedilum* and *Phragmipedilum* and the species *Aerangis ellisii, Dendrobium cruentum, Laelia jongheana, Laelia lobata, Peristeria elata* and *Renanthera imschootiana* are placed on Appendix I (very strict control in trade) whereas all other species of ORCHIDACEAE are placed on Appendix II (less strict control in trade) [6].

## *Salep* orchids

Ground orchid bulbs of the Orchidoideae, also known as *salep,* are very popular in Turkey and adjacent countries in central Asia. They are used to produce ice cream in summer and hot drinks during winter. *Salep* is also used as medicine. In the early 1990s the trade of *salep* increased strongly. Official statistics from the Turkish State Institute of Statistics shows that the export between 1995 and 1999 was 282.000 kg annually. To achieve this amount of *salep* 9.825.000 – 19.650.000 bulbs are required. It is unknown if this information is related to pure *salep*, substitutes or mixtures. However, as this harvesting is unsustainable, laws have been established to protect these orchids. In Turkey there are three laws that protect them: the first law is the Turkish Forest Law, which regulates the use of non-wood forest products. In short, this law states that it is forbidden to collect and remove any form of forest vegetation. The second law, the Turkish Law of Natural Parks, states that “The production of forest products, hunting and disturbing the natural balance is prohibited.” Since collecting *salep* is classified as production of forest products, it is prohibited in all protected areas. The final law in Turkey is The Regulation on Collection, Production and Export of Bulbs of Wildflowers. As the title of this law suggests, this law regulates the production and the export of bulbs, roots and tubers of flowers. It also defines a list with species that may not be taken away from the wild for export [7]. The exact ingredients in *salep* cannot be identified without molecular identification tools, which makes it difficult to enforce these laws.

## How a web application can improve control in illegal orchid trade

To make it easier to follow the trade routes of orchid smuggling, a web application that can identify illegally traded orchids is desired. This application could be used on laptops or desktops as well as mobile devices by taking pictures of flowers, leaves or underground tubers and uploading the pictures to a website. A simple flowchart of the application is shown in figure 2. In this project the focus was on creating the website and integrating the identification application for pictures taken of flowers of slipper orchids. This application is currently under development at Naturalis.

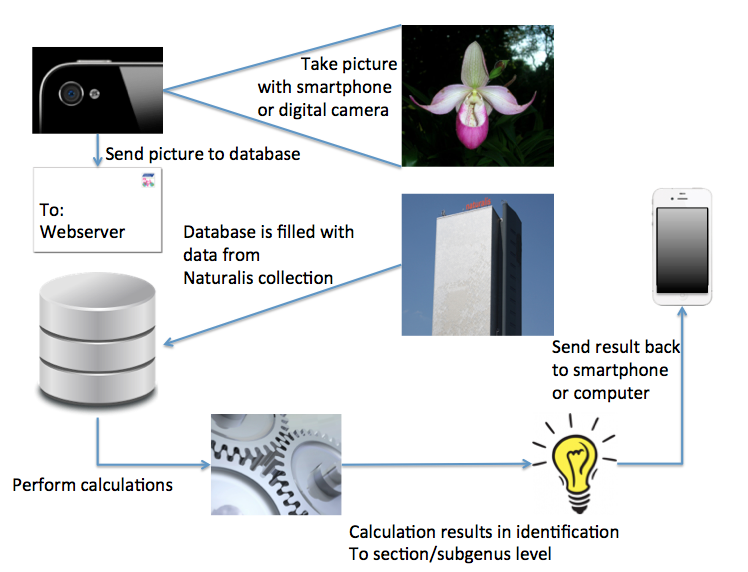


Figure 2 A simple flowchart of the application made during this project. Resources of the pictures: [8-15]

## Comparable software

There is software available that can identify a person using face recognition, for example the software KeyLemon [16]. This software could be used to unlock a computer.

Essentially the software takes a picture or series of pictures of your face. When it takes a series of pictures it is almost always required to move your head up and down and / or left and right. The software saves this picture or pictures. When you use the software to unlock your computer the software takes a picture or series of pictures of your face and compares this with the saved picture(s). If it finds a match you will be logged-in to your account. The existence of this kind of software demonstrates that it is possible to identify objects that look very similar to each other, such as two different persons. If this is possible, it would also be possible to identify orchids to section level. All sections have some very specific characteristics that are unique to that section.

Another example of comparable software is LeafSnap [17]. LeafSnap is an iPhone and android application that allows the user to take a picture of a leaf. After taking a picture the software turns the picture in a way that it can recognize the shape. The shape of the leaf will be used to create a list of possible trees.

These applications differ from the pipeline that was developing during this internship in several aspects. First of all, KeyLemon is made to recognize a human face and LeafSnap is made to recognize different leaves. The pipeline that was developed during this internship focused on slipper orchid flowers and *salep* orchid tubers. Secondly the pipeline will be available on a website, while the comparable software needs to be installed on a local machine (e.g. a computer or a smartphone). This makes it usable from different platforms. The comparable applications are specific for smartphones or computers, and also to the Operating System (OS) of the used device.

## Goal of this internship

The goal of this internship was to demonstrate that it is possible to classify slipper orchid flowers from pictures. When this concept was proved the classification software was trained very well. Once the classification software works sufficiently accurate, it will be available online. Customs officers can then use this software to check whether accompanying permits are correct when confiscating material. If this control is simplified, it will help prevent illegal trade in slipper orchids, which will prevent them from going extinct in the wild.

# Materials and Methods

## Website

During this internship a website was developed. Users of this website can upload a picture to the server. On the server the software will classify the orchid on the picture to section level. The processes behind this website, like clicking on a button, were written in Python 2.7 using the Django package. The layouts of the web pages are written in HTML, using CSS style sheets. There are two versions of every HTML file, one for computers and one for mobile devices. The different CSS style sheets, HTML files and python scripts can be found in appendices 1.2, 1.3 and 1.6. To make the website usable, a well-trained Artificial Neural Network (ANN) is required. To train this network, pictures of slipper orchid flowers were needed. These pictures were saved on a shared Flickr account on www.flickr.com, to make the pictures accessible to everyone who needs them.

## Flickr

To store the pictures in a safe place where they are accessible to whoever needs them, a shared Flickr account was created. Flickr is a website for saving and sharing pictures. Because the flower pictures are from a private collection of David Roberts, it is not allowed to freely share the pictures of the flowers, so the settings of the Flickr account were set to private. This means that only persons with the account name and password can access the pictures.

On Flickr it is possible to add metadata tags to the pictures. These tags were used later in the preparation process to save the pictures in the correct directory. To download the pictures and the metadata via the command line, a python script written by Hugo Haas, Offlickr.py, was modified and used (see appendix 1.5.4) [18].

## Training

### Preparation

Before training of the ANNs is possible, a preparation step is needed. First of all, pictures of the *salep* orchid tubers and the tubers of the look-a-likes were required. So pictures of *salep* orchid tubers and tubers of look-a-likes were taken at the Sylvius lab. It is required that the orientation is the same for all pictures. For instance, if the first picture of a tuber with appendices has the appendices on the right, all other tubers with appendices must have the appendices on the right as well. The user has to use the same orientation as the trainer. The background has to be one colour, like white or black, and this colour must be the same for every picture. The last requirement is that there is only one tuber on the picture.

The second step in this process is to download the pictures and metadata and search for the tags in the metadata. An example of the metadata can be found in supplementary 1.1. The blue square indicates the location of the tags. The next step is to convert the pictures from .jpg- to .png-format. This is needed for the training scripts, which work only with .png files. Finally the pictures will be placed into the correct directory using the tags. The tuber pictures were divided by shape and look-a-like or not, which resulted in six directories. Round, Spur and Oblong for the different *salep* orchid tubers, LRound, LSpur and LOblong for the different look-a-like tubers. The slipper orchid flowers were divided by section and within each section the pictures were divided by species. A section is a taxonomic level below a genus but above a species. Sections were created to obtain a better overview of species rich genera. Species in he same section share a common ancestor. In this project the classification of Chochai, who carried out molecular phylogenetic research to come up with an evolutionary based list of sections, is used [19]. Some sections contain many species, such as *P. Barbata* and *P. Paphiopedilum*. Other sections just contain a few species, such as *P. Pardalopetalum* and *P. Cochlopetalum*.

After separating the pictures of the tubers from the flowers, the pictures of the tubers will be segmented. This means that the background is made entirely white and the tuber is cut out of the picture. It was not necessary to split the flower pictures, because these pictures were already segmented.

### Slipper orchids

A total of 5 genera within the slipper orchids exist. The genera are *Cypripedium, Mexipedium Paphiopedilum, Phragmipedium* and *Selenipedium*. Due to time constraints, only the genus *Paphiopedilum* was used in this project. The sections of this genus are *Barbata, Brachypetalum, Chochlopetalum, Coryopedilmu, Paphiopedilum, Pardalopetalum* and *Parvisepalum*, which are based on molecular phylogenetic research [19]. The ANNs were trained on all these sections and tested on the four sections *Barbata, Brachypetalum, Coryopedilum* and *Parvisepalum* The ANNs were trained on more sections than they were tested with. This resulted in more realistic outcome, because the ANNs have to classify a new picture to all possible sections. If the ANNs were only trained on the sections used in the test runs, this could affect the outcome in a way that the results look much better because 3 of the 7 sections were missing.

The script that trained the network was already developed at Naturalis by Rutger Vos. This script accepts a parameter to specify the Desired Error. This is the maximum error rate that will be accepted. If the parameter is left empty the Desired Error will be 0.0001. When the error drops below this threshold the training is finished. To find the optimal Desired Error, seven networks were trained with a Desired Error in tenfold increments: the first network had a Desired Error of 0.0001, the second network had a Desired Error of 0.00001 and so on. The expectation was that a network with a lower Desired Error would classify a new picture better. To semi-automate the preparation for the training some scripts were written in python and bash. The workflow of these scripts can be found in figure 3. These scripts were for training the Artificial Neural Networks (ANNs), which is only done by the administrator of the website and not by the end-users of the website. The administrator has access to the shared Flickr account, so the manual check (and download) of the pictures that is required for one of the scripts, would not be a problem. Besides networks for classifying the slipper orchid flowers, ANNs for classifying *salep* orchid tubers were also created.

### *Salep* orchids

*Salep* orchid tubers are very popular in Turkey. These tubers are used to make ice creams during summer and hot drinks during winter. Due to a lack of information about the collected tubers, the ANNs could only be trained to recognize differences between *salep* orchid tubers and look-a-likes obtained from non orchid families. The tubers of look-a-likes that were used were from *Arum maculatum* (Araceae), *Asparagus officinalis* (Asparagaceae)*, Polygonatum verticillatum, Tulipa greigii,* and *T. sp.* (Liliaceae)[20]. Supplementary 2 contains pictures of these tubers.

Figure 3 Flowchart of training preparation scripts. Blue: preparation steps. Green: processing step to produce useful data for training the flower Artificial Neural Networks. Purple: Training of the Artificial Neural Networks

## Testing the networks

### Slipper orchids

After training the networks for the slipper orchid flowers, they were tested on pictures of four sections. The tested sections were *Barbata, Brachypetalum, Coryopedilum* and *Parvisepalum*. The networks were also tested with pictures of some primary hybrids of *Parvisepalum* and *Brachypetalum* with the aim of testing if it is also possible to correctly identify hybrids with the software. The assumption was that for primary hybrids a trained network would ideally list the sections of both parents as possible identification. However, some results of the first test round were very poor. To find the source of these results, pictures of some poorly classified species were checked for detection ability of the defining characteristics of the corresponding section. Table 1 holds a list with the characteristics per section based on Cribb P. [21]. During this research it was discovered that some species had only one picture in the train data. Therefore, extra pictures were added to the train data and the networks were trained again to find out if better results could be obtained after more extensive training. These networks were tested again with the same test sets.

Table 1 characteristics per section

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| *Characteristic* | *Barbata* | *Brachypetalum* | *Coryopedilum* | *Parvisepalum* |
| Petals | Spathulate | Broad-elliptic, pale yellow or white | Long, hanging down, spirally twisted, tapering, glandular at tip | Broad-elliptic,  Subcircular, Cream/yellow coloured |
| Spotted or warted? | Yes | Yes | Yes | No |
| Lip | Incurved side lobes | Small, ovoid with incurved margin | Incurved side lobes | Large, thin-textured |
| Staminode | Lunate | Ovate to transversely elliptic | Oblong | Large |

### *Salep* orchids

Due to time constraints, pictures of these orchids could not be investigated further during this internship.

## Testing the website

To test the website, a working ANN was required. After testing the ANNs, a working ANN was implemented on the website. This website was tested on receiving a picture, running the classify script using the given network and sending back a result. First this was tested for only one device. Once this was working correctly, the website was also tested to receive pictures from more than one device at the same time and send a result back. Receiving the pictures was working correctly, but sending back a result was not. The result of the first uploaded picture was send back to all devices. So this part of the website was made operable during this internship. The IP-address was used to save the uploaded picture in a directory. The name of this directory will be the IP-address. Only the picture in the directory with the correct IP-address will be classified, and its result is sent back to the website.

# Results

## Website

To make the software available for end-users a website was developed. A first design of this website was finished during this internship. It consists of a homepage, a page for uploading a picture, a page to show the picture is uploaded correctly, and a results page. On the homepage it is possible to choose to upload a picture or remove unused files from the server. The last option requires to log in with a valid account. A flowchart of this website is shown in figure 4. Figure 5 zooms in on the uploading part of the website. After selecting the upload option the user is forwarded to the upload page. On this page the user can select a picture to upload. On iPhones it is also possible to take a picture after tapping the “select file” button. The website will check whether the selected file is a picture. If it is not, the user stays on the upload page and a warning is issued. After uploading the file, the picture will be renamed. When these modifications are done the user is directed to the upload success page. Here the user can see the uploaded picture. The user can choose to see the results or cancel. If the user selects the result option, the classify.pl script will run to classify the picture (see appendix 1.4.1). A flowchart of the classification step of the website can be found in figure 6. The output of this program is a list with numbers. An example of this output can be found in supplementary 1.2. This list is sent to a python script, result.py, that translates this list to a readable result (see appendix 1.6.2). Figure 7 shows a flowchart of this script. The output of this script is sent back to the result page, where the user can view it. Supplementary 3 contains screenshots of this website.

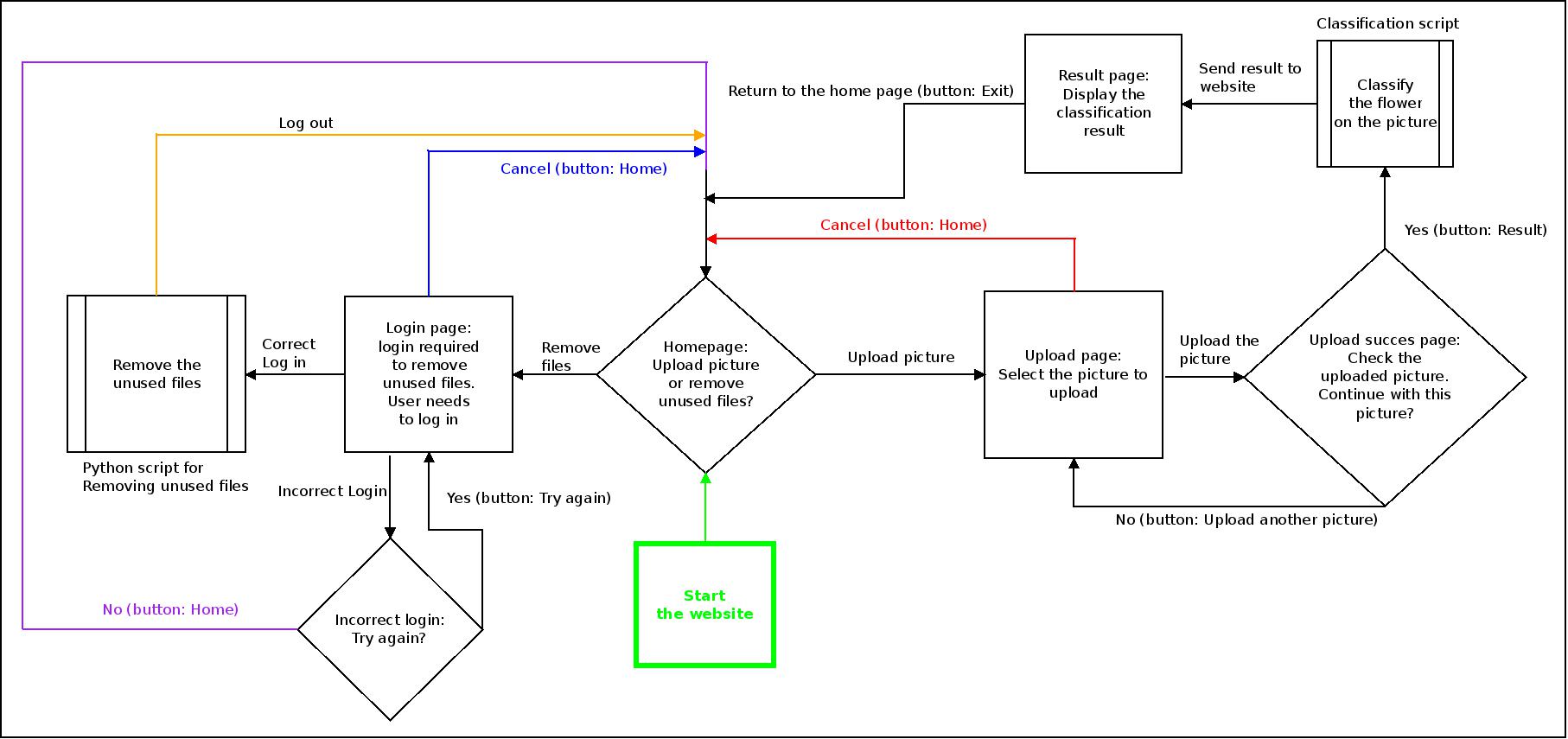


Figure 4 Flowchart of the website. Green: Start of the workflow, start the website.

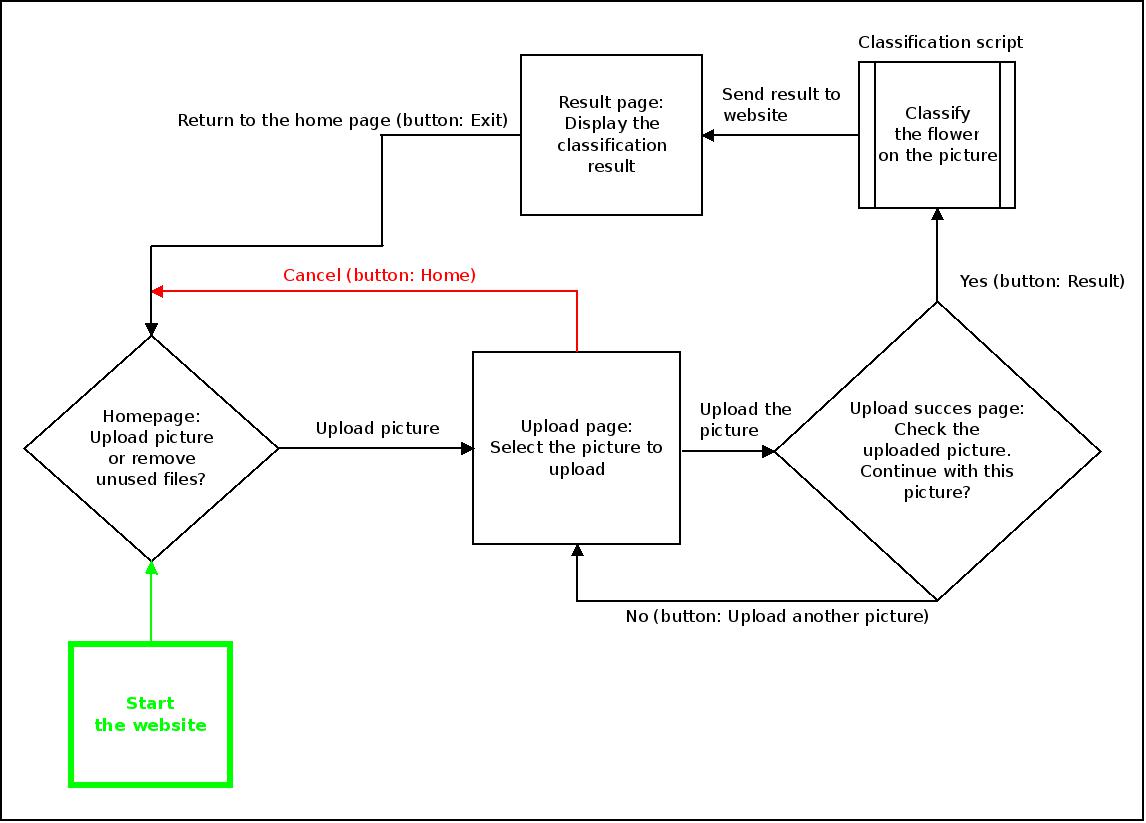


Figure 5 Flowchart of the website, zoomed in on uploading. Green: Start of the workflow, start the website.

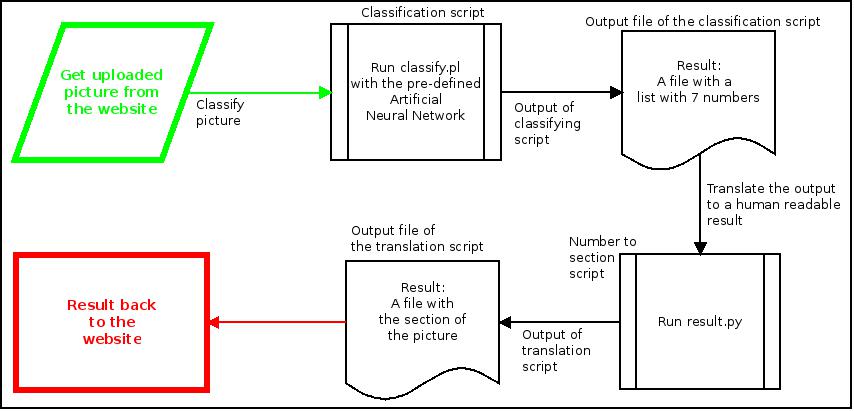


Figure 6 Flowchart of the classification step of the website. Green: Start of the workflow, get the picture, Red: End of the workflow, send the result.

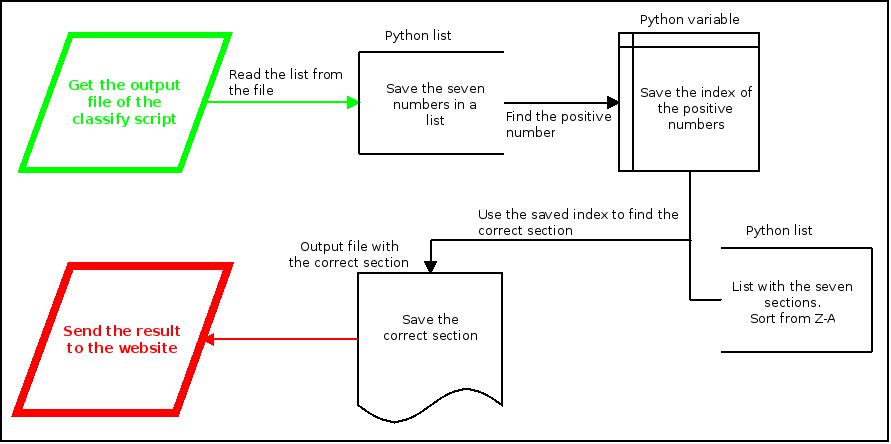


Figure 7 Flowchart of result.py. Green: Start of the workflow, get the input file. Red: End of the workflow, send the result.

## Preparation scripts

To semi-automate the preparation process, a bash script, training.sh, was developed (see appendix 1.1.3, and the first script in figure 3). Figure 8 contains a workflow of this script. The first step in this script was running Offlickr.py to download the pictures as .jpg and the metadata files as .xml. After downloading these files, the script will run another python script, get\_tags.py (see appendix 1.5.3), to get the original names and the tags from the metadata. A workflow of this script can be found in figure 9. This info will be saved in .txt files, using the id of the picture as name. For example, if the picture name is 123456789.jpg the tag file of this picture is 123456789\_tags.txt. At the end of this step the .xml files are removed. Example of different tag files can be found in supplements 1.3-1.5. The structure of these files is always the same: the first line is the original name of the picture, then an empty line, after that the shape for the tuber pictures or the section and species for the flower pictures. The last tag indicates whether the object is a slipper orchid flower or a *salep* orchid tuber. The knowledge of this structure can be used in the next step.

In the next step the pictures are divided between two directories, Flower and Tuber. Before the pictures and tags are moved to the correct directory, the pictures were converted from .jpg to .png. This conversion was required because the training scripts only work with png files. After converting the pictures, the .jpg files were not used anymore, so all .jpg files were removed.

After dividing the pictures between the directories Flower and Tuber, the separation goes further. First the Flower pictures are divided between the different slipper orchid sections and species. After this division there are some directories with the section names inside the Flower directory, and every section directory contains further species directories. After dividing the Flower pictures, the Tuber pictures are divided between shape and Look-a-Like or orchid. This step will produce six directories: LOblong, LSpur, LRound, Oblong, Spur and Round. All directories starting with an “L” are for the Look-a-Like tubers.

The last step in the preparation process is segmenting the pictures of the tubers, using a Perl script developed by Rutger Vos: splitter.pl [22] (see appendix 1.4.2). This script uses the Perl package Image::Magick to modify the picture so that only the tuber is on the picture, with a minimum background. With an option of Image::Magick the background is modified to be completely white (see the square in appendix 1.4.2). Figure S-11 in supplementary 2 shows a picture before and after segmentation. The pictures of the flowers were already segmented, so this step was not required for these pictures.

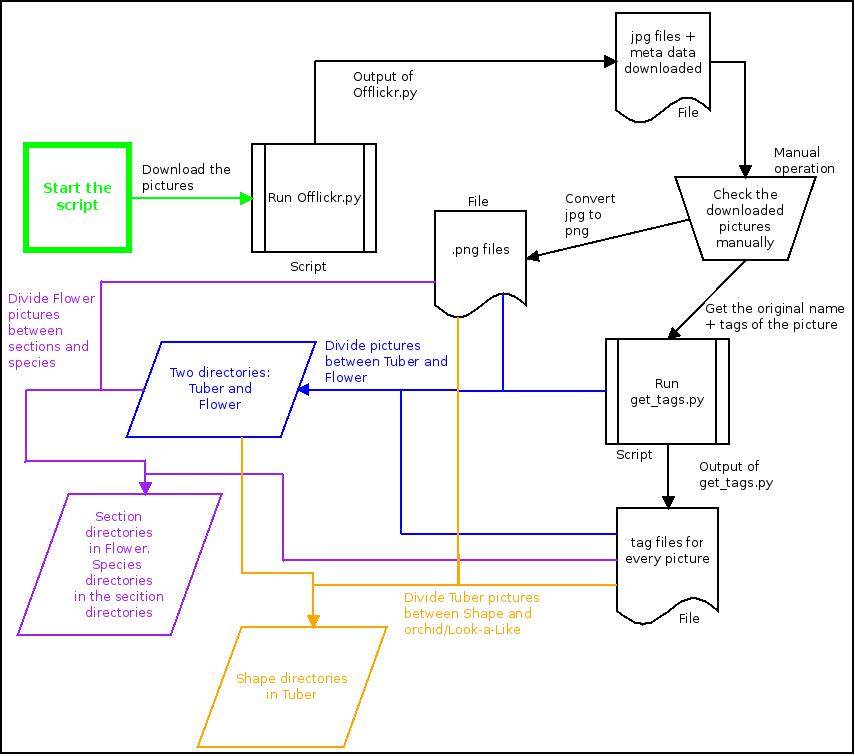


Figure 8 Flowchart of training.sh. The colour of the arrows indicates in which step they are involved.

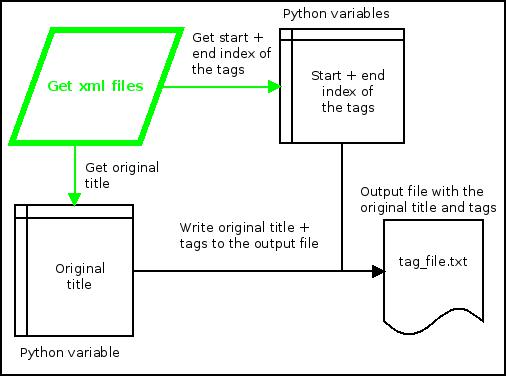


Figure 9 Flowchart of get\_tags.py. Green: Start of the workflow.

## Create training data

When the preparation script is finished the training data can be generated. To do this, a bash script, create\_traindata.sh, was developed (see appendix 1.1.1, and the second script in figure 3). This script runs two different Perl scripts developed by Rutger Vos. One of these scripts, traindata.pl, is for the tubers and the other, traindata2.pl, for the flowers (see appendices 1.4.4 and 1.4.5). The only difference between these two scripts is the accepted input files. The script for the tubers only accept files with a name in the format <number>,<number>.png, while the other script accepts files with a name that contains at least one number. These scripts create .tsv (tab separated value) files for every directory. So after running create\_traindata.sh there is a .tsv file for every species of the flower pictures and a tsv file for every shape of the tuber. These tsv files contain the intensities of Red, Blue and Green (RBG) of 50 vertical bins and 25 horizontal bins. This means that the classification is based on both colour and shape.

From this point the .tsv files of the tuber can be used for training the neural networks. The tsv files of the flowers need to be modified before using them for training the neural networks. To automate this modification process a bash script, modify\_flower\_data.sh, was developed that runs two python scripts (see appendix 1.1.2, the third script in figure 3 and appendices 1.5.1 and 1.5.1). The first python script, combine\_files.py, creates tsv files per section by combining the tsv files in that specific section. When all sections have a tsv file the second python script, add\_columns.py, is run. This script creates extra columns, one column per section. After creating the columns they are filled in. The guideline for this step is: the first section gets 1’s in the first column and -1’s in all other columns. The second section gets 1 in the second column and -1 in all other columns, and so on . The sections were sorted from A to Z. This convention was subsequently used in the classification step. This step produces a list with seven numbers, one for each possible section, sorted from Z to A. So when the last of these seven numbers is positive this means the pictures is classified as the first section and so on. After adding these columns the data can be used to train the neural networks.

## Neural networks

After completing these preparation process, a neural network can be trained. To train the network a Perl script written by Rutger Vos, trainai.pl [22], was used (see appendix 1.4.3). This script accepts a directory with tsv files, the number of categories (the number of extra columns), the output file, a Desired Error and the maximal number of epochs. The last one is the maximum number of rounds the network will be trained. When either the Desired Error or the number of epochs is reached, the training process will stop. Because the network is trained with the tsv files, it is trained to see patterns in the colour intensities of RGB. Now the network acts like it can see different colours and shapes.

During this project 14 ANNs were created for identifying slipper orchid flowers. Seven for the first test run and seven for the second test run. As mentioned earlier the Desired Error was incremented tenfold per network. The networks can be found on:

<https://github.com/naturalis/img-classify/tree/master/webapp/ann>

All ANNs with a name ending with 4x.ann were used for the first test run and all ANN’s with 5x.ann were used for the second test run. Due to time constraints, networks for tubers of *salep* orchids and look-a-likes were created, but not tested.

## First test run

In the first test run four of the seven sections of *Paphiopedilum*, i.e. *Barbata, Brachypetalum, Coryopedilum* and *Parvisepalum* were tested. The ANNs were also tested with a few pictures of hybrids between species of *Parvisepalum* and *Brachypetalum*. The results of these tests can be found in supplementary 4. The different networks are shown horizontally and the different pictures vertically. A green cell with the text ‘Correct’ means the network classified the picture to the correct section. A yellow cell with the text ‘Unknown’ means the network was unable to classify the picture to any section. A yellow cell with the text ‘Unknown <number>/<number>’ means the network was unable to select a section from two options. When this text is green, one of the options was the correct section. A red cell with the text ‘Incorrect <number>’ means the network classified the picture to a wrong section. The number indicates to which section the network erroneously classified the picture. Below, the obtained results will be discussed in more detail.

### Section *Barbata*

This section is characterized morphologically by spathulated, spotted and/or warted petals and incurved side lobes of the lip. Most of the species of the section *Barbata* were classified incorrectly. As seen in table 2 in Supplementary 4, most of the incorrectly classified pictures were classified as section *Paphiopedilum*. One explanation for this mistake could be that the species in both sections share several similar characteristics. The petals for instance have mostly the same shape. Another explanation is that both sections have many species so variation is high.

After characteristic pictures of three different species of the section *Barbata* were added to the test set, belonging to *P.* *argus, P. tonsum* and *P.* *wardii*, it turned out that even pictures with all characteristics of the section clearly visible were hard to classify correctly (see table 3 in supplementary 4). Like before, one of the possible explanations could be that species of the section *Barbata* share characteristics with the section *Paphiopedilum*. Another explanation could be that not enough train data of these species were added yet.

### Section *Brachypetalum*

This section is characterized morphologically by broad-elliptic petals with purple spots, a small and ovoid lip with an incurved margin and pale yellow or white flowers. The visual recognition software developed during this internship identified this section very well (see table 4 in supplementary 3). None of the pictures were classified incorrectly and only a few were classified as unknown. But even then, most of the time the correct section was one of the options. A possible explanation could be that this section contains few species only so variation is low.

### Section *Coryopedilum*

This section is characterized morphologically by long, tapering, spirally twisted and hanging petals with warts on the margins and glands at the tip and a lip with incurved side lobes. In this section one picture stands out, *P. rothschildianium*1. This picture was classified correctly with all networks (See table 5 in supplementary 4). So this is a good picture for classifying. Further, it is shown that the other two pictures of this species were much harder to identify. Possible explanations are that the first picture has more characteristics visible than the other two (i.e. spirally twisted petals and incurved side lobes) or that the resolution of the first picture is better than the other two.

### Section *Parvisepalum*

This section is characterized morphologically by broad-elliptic, subcircular cream/yellow coloured petals and a large staminode. This section also has a striking picture, *P. emersonii*2\_1. This picture was always incorrectly classified. Research on this picture shows that the petals are darker than the petals on the other pictures of this section. In addition, table 6 in supplementary 4 shows that *P.* *armeniacum* was the best recognized species. All test pictures of this species were classified correct by all networks. A possible explanation could be that this section contains few species only so variation is low.

### Hybrids of species from sections *Parvisepalum* and *Brachypetalum*

Besides testing the ANNs with pictures of the sections, the networks were also tested with pictures of hybrids of species of different sections. This was done to test if it was possible to identify one or both of the parental species. It could be hard to identify both parental species because the hybrid shares a part of its morphology with the mother and another part with the father. Since the ANNs were not trained with pictures of hybrids, the assumption was that only one parental section would be found by the neural networks. Table 7 in supplementary 4 shows that most of the time the pictures were indeed classified as one of the parental sections. In just one case, *armeniacum* x *concolor* using the network with Desired Error 0.00001, both parental sections were found.

## Characteristic research

Following the results of the first run, research on the characteristics of some species was done to understand why some species or sections were hard to classify. The results of this research can be found in supplementary 5. These results show that most of the test pictures do not have all section characteristics visible. The results indicate also that some characteristics were shared between different sections, like broad-elliptic petals in the sections *Brachypetalum* and *Parvisepalum*. This could be a reason why some species were classified very poorly. Essential characters necessary for correct identification turned out to be the shape of the petals, the presence or absence of wards on the petals, the colour of the flower, and the shape of the lip (incurved or not). Especially the colour was often distorted (i.e. white flowers looked yellow or yellow flowers looked green) which might explain some of the poor results obtained.

During this research it was also discovered that some species of the sections *Barbata* and *Coryopedilum* had only one picture in the train data. This could also be a reason why these species were classified poorly. Extra pictures of these species were added to the train data. This expanded train set was used to train the ANNs again. These new networks were tested again with the same test sets to find out if better classification results were obtained. The results of this experiment are described in more detail below.

## Second test run

To demonstrate that more pictures in the train data improves the ability of the ANNs to classify pictures, a second test run was done. The results of this run can be found in supplementary 6. The structure of these tables is the same as in supplementary 4. Some pictures (i.e. 50 out of 124; roughly 40%) were indeed classified better, showing that more training data pays of. Other pictures (i.e. 30 out of 124; roughly 25%) were classified worse than in the first test run. One explanation could be coincidence during training. When you train two networks with the same settings and the same train set, both networks will be different nonetheless. The cause of this phenomenon is that during the training the networks create connections. These connections will be different every time you train the network. To classify a new picture the connections in the network found at that moment are used to give a result. This means that different connections can result in a different classification. The remaining 44 pictures (roughly 35%) were classified as good as in the first test run.

### Section *Barbata*

After the second test, species belonging to section *Barbata*were still difficult to classify (See tables 12 and 13 in supplementary 6). The results do show that many of the pictures of this section were classified better (i.e. 25 out of 45; roughly 56%). It is striking that some pictures of species with additional pictures in the train set were classified worse, though, because the assumption is that more pictures in the train set will improve the classification. These concerned the species *P. argus*, *P. hennisianum, P. tonsum* and *P. wardii*. A possible explanation is the different connections in the ANN’s of the first test run and the second test run.

### Section *Brachypetalum*

*Brachypetalum* is one of the sections that did not get additional pictures in the train set. Nevertheless, there are some pictures that were better classified than before (See table 14 in supplementary 6). These concerned some pictures of *P. concolor* and *P. godefroyae*. But also in this section there were some pictures that were classified worse. These concerned other pictures of *P. concolor* and *P. godefroyae* and also some pictures of *P. niveum*. A possible explanation is again the different connections between the ANNs of the first test run and the second test run.

### Section *Coryopedilum*

This is a notable section. This section received additional pictures in the train data, but not of the species in the test data. Nevertheless, the pictures were much better classified than in the first run (See table 15 in supplementary 6). This is also the only section that has no longer incorrectly classified pictures. Still some pictures were classified as unknown. A possible explanation could be that the train data contains more variation after adding some pictures. The pictures in the test data looked more like the new variations, which made it possible to classify the picture to the correct section.

### Section *Parvisepalum*

After the first test run this section was one of the best-classified sections. Although there were no additional pictures of this section added, some pictures were classified better during the second test run (See table 16 in supplementary 6). Again, one of the possible explanations could be the different connections between the first ANNs and the second ANNs.

### Hybrids of species from section *Parvisepalum* and *Brachypetalum*

Most of the hybrids were classified correctly in the first test run. During the second test run it was seen that some pictures were classified better to one of the parental sections (See table 17 in supplementary 6). The picture that was classified to both parental sections in the first run, was classified to only one parental section in this run. Because it was only seen one time in the first test run that both parental sections were recognized, it could be coincidence. This could be an explanation to the results of the second test run. A possible explanation for the better results of some pictures could be again the different connections between the different ANNs.

# Discussion

## Tubers

Because it is hard to extract DNA from a dried tuber, most of the tubers of confiscated *salep* orchids could not yet be identified to species level during my internship. During this project they could therefore only be used to train the neural network to see differences between orchid tubers and lookalike tubers up to the family (i.e. orchid versus non orchid) level. The time necessary to test this was lacking, though, so this is not worked out completely. A new student will use the pictures that were made to further train the neural network to identify them to orchids versus lookalikes.

## Slipper orchids

Sections of slipper orchids that could be identified correctly included *Brachypetalum* (75-95% in the second test run), *Coryopedilum* (67-100% in the second test run), *Parvisepalum* (73-85% in the second test run) and primary hybrids of species of *Parvisepalum* and *Brachypetalum* (75-100% in the second test run). The only section that was less easy to identify was *Barbata* (24-41% in the second test run). The section *Barbata*has more characteristics in common with the section *Paphiopedilum* than the other sections, which might explain why pictures of the section *Barbata* were often classified as *Paphiopedilum*.

## Training networks

During training of the networks it was found that it was very hard to find good pictures of flowers of some species. When this fact is combined with the fact that some species had few pictures in the train set, this could be a reason why these species were poorly classified. This assumption is supported by the fact that when more pictures were added to the train set, identification improved. This was especially found for *P. hennisianum, P. tonsum, P. violascens, P. glanduliferum, P. wilhelminiae* and *P. rothschildianium*.

# Conclusions

## Classifying slipper orchids from pictures is possible

Although some species were very hard to classify, the results show that it is possible for most species to be classified to section level correctly using a picture of a flower. Some species need more or better pictures in the train set, but the concept of classifying slipper orchids from a picture of a flower has been demonstrated during this project.

## Differences between well classifiable vs hard classifiable sections of slipper orchids

The only section of those tested that remained hard to classify was the section *Barbata*. Most of the floral characteristics of this section are shared with the section *Paphiopedilum*. All other tested sections have unique characteristics. This might explain why some of the pictures of the section *Barbata* were classified as *Paphiopedilum*.

After adding pictures of this section to the training set the classification improved a little bit. Still many pictures were classified as *Paphiopedilum*. This supports the assumption that the shared characteristics make it hard to classify pictures of this section correctly.

# Suggestions for follow-ups

To make this software available for the outside world, the website needs to become online.

Before the website will be made online, the ANNs need to be tested with the sections *P. Cochlopetalum, P. Paphiopedilum* and *P. Pardalopetalum*. Depending on the results of these tests, the ANNs need to be trained again with more/better train data of one or more of these sections. The ANNs have to be re-trained with more/better train data of the section *Barbata* anyway.

When the website is online, customs officers can use this website to classify orchid flowers. To make the website more useful for custom officers, the software needs to be expended by classifying orchid tubers and orchid leafs.

# Acknowledgements

I would like to thank the following people for helping me: Serrano Pereira for segmenting some pictures and test some of my scripts, David Roberts for providing slipper orchid flower pictures, Rogier van Vught for providing some look-a-like tubers and Benjamin Versteeg for helping me out with Django.

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[22] Rutger Vos, Bioinformaticist at Naturalis Biodiversity Center

[23] Photographs by Patrick Wijntjes

# Supplements

## Input and output files

### Example of an xml file

<photo id="12342126885" secret="ca74c114fa" server="7451" farm="8" dateuploaded="1391690138" isfavorite="0" license="0" safety\_level="2" rotation="0" originalsecret="29fddd19e8" originalformat="jpg" views="0" media="photo">

<owner nsid="113733456@N06" username="patrick\_naturalis" realname="Patrick Wijntjes" location="" iconserver="0" iconfarm="0" path\_alias=""/>

<title>charlesworthii5</title>

<description/>

<visibility ispublic="0" isfriend="0" isfamily="0"/>

<dates posted="1391690138" taken="2012-08-29 17:13:09" takengranularity="0" lastupdate="1391690248"/>

<permissions permcomment="0" permaddmeta="0"/>

<editability cancomment="1" canaddmeta="1"/>

<publiceditability cancomment="0" canaddmeta="0"/>

<usage candownload="1" canblog="1" canprint="1" canshare="0"/>

<comments>0</comments>

<notes/>

<people haspeople="0"/>

<tags>

<tag id="113688134-12342126885-10993197" author="113733456@N06" raw="genus:Paphiopedilum" machine\_tag="0">genuspaphiopedilum</tag>

<tag id="113688134-12342126885-188931923" author="113733456@N06" raw="species:chariesworthii" machine\_tag="0">specieschariesworthii</tag>

<tag id="113688134-12342126885-535" author="113733456@N06" raw="Flower" machine\_tag="0">flower</tag>

</tags>

<urls>

<url type="photopage">http://www.flickr.com/photos/113733456@N06/12342126885/</url>

</urls>

</photo>

### Example of an output file of classify.pl

Entry: color-100,451.png

$VAR1 = [

'-0.456072704368231',

'-0.973874677375054',

'-1',

'-0.999999808175595',

'0.990980698747592'

];

Entry: color-129,334.png

$VAR1 = [

'-0.811027796569903',

'-0.999812464339958',

'0.99572448600037',

'-0.996400768699541',

'-1'

];

Entry: color-141,711.png

$VAR1 = [

'0.971726551475096',

'-0.999312299486709',

'-0.981400879260904',

'-0.99960776872456',

'-1'

];

Entry: color-222,648.png

$VAR1 = [

'0.996738096627912',

'-0.919456120335819',

'-0.999987677705016',

'-0.999994486624424',

'-0.99999973650751'

];

Entry: color-447,1931.png

$VAR1 = [

'0.935067461526289',

'-0.999821434281431',

'-0.999766410047794',

'-0.996420240471305',

'-1'

];

Entry: color-46,379.png

$VAR1 = [

'0.997483774166035',

'-0.994043337286478',

'-0.998988899850576',

'-0.999999999293947',

'-0.99999999995774'

];

Entry: color-50,380.png

$VAR1 = [

'0.940404321606217',

'-0.758403411429261',

'-0.999999840072036',

'-0.999999955862603',

'-0.999999983567608'

];

Entry: color-56,651.png

$VAR1 = [

'-0.897130009835269',

'0.994303714419604',

'-0.999999951703297',

'-0.999999993987055',

'-1'

];

Entry: color-64,737.png

$VAR1 = [

'0.93481407861017',

'0.345711139511172',

'-0.999999999999977',

'-0.999999985064081',

'-1'

];

Entry: color-7,390.png

$VAR1 = [

'-0.999853813787961',

'-0.303503446288619',

'-0.999984006201254',

'-0.999999238570459',

'-1'

];

### Tag file of a flower

bellatulum12 phot

genus:Brachypetalum

species:bellatulum

Flower

### Tag file of a *salep* tuber

TEH-1.1\_5

Orchid\_spur

Orchid

Spur

genus:Dactylorhiza

species:incarnata

Tuber

### Tag file of a Look-a-Like tuber

tulipa red riding hood3

Look-a-Like\_round

Look-a-Like

Round

genus:Tulipa

species:greigii

Tuber

## Photographs of *salep* orchid look-a-likes [23]



Figure S-1 *Arum maculatum* [23]



Figure S-2 *Asparagus officinalis* [23]



Figure S-3 *Polygonatum verticillatum* [23



Figure S-4 *Tulipa greigii* [23]



Figure S-5 *Tulipa sp.* [23]



Figure S-6 A: A picture of a tuber, before segmenting. B: A picture of the same tuber after segmenting [23].

## Screenshots of the classification website

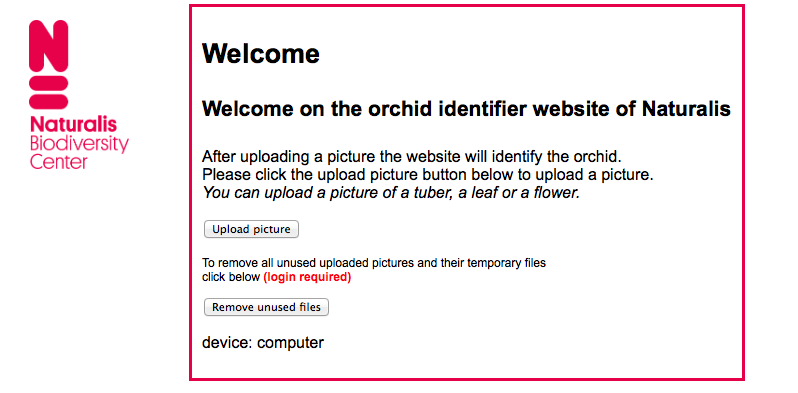


Figure S-1 Homepage of the classification website

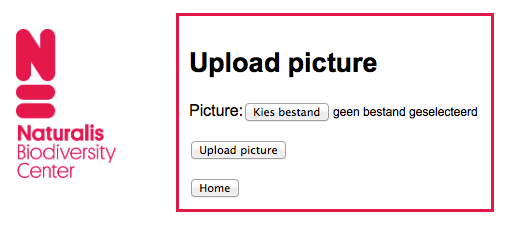


Figure S-2 Upload page of the classification website

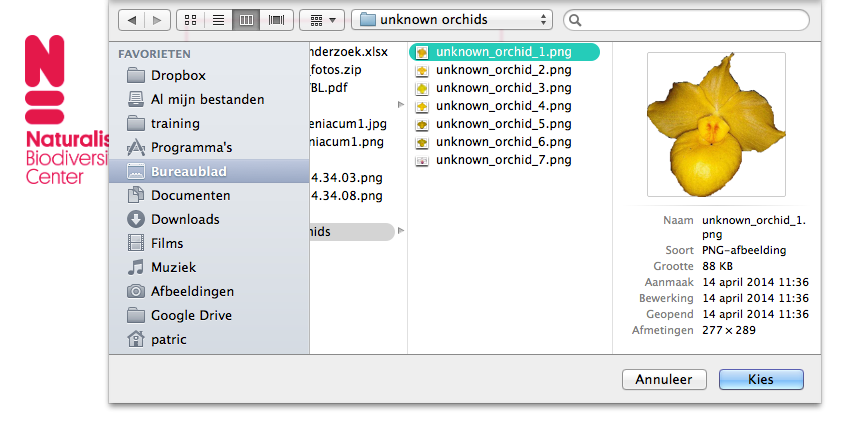


Figure S-3 Screen to select the picture to be classified

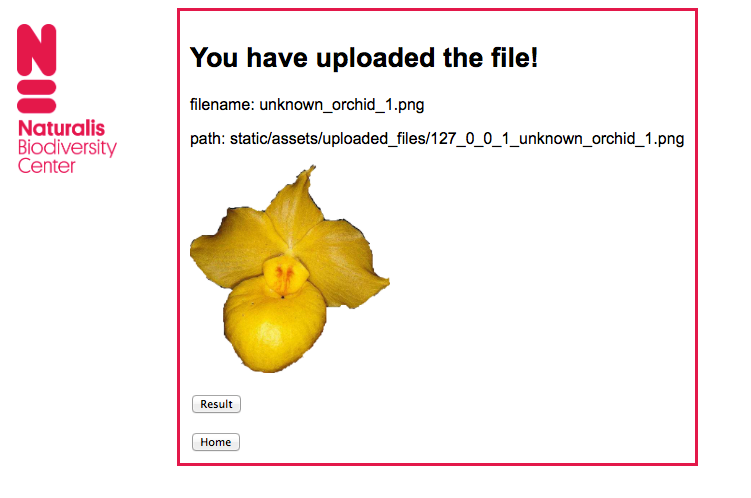


Figure S-4 page to check if the picture is uploaded correctly

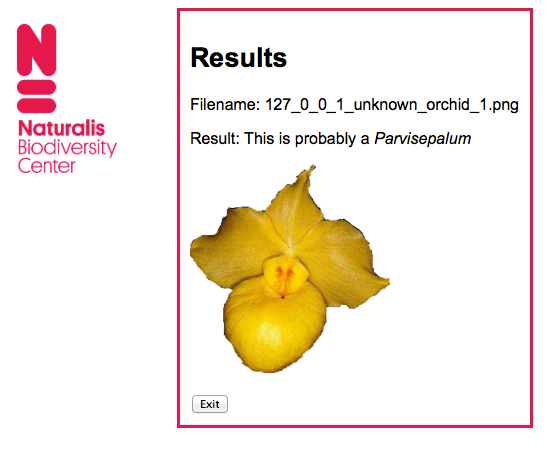


Figure S-5 Result page of the classification website

## Results of first test run

Table 2 Results of first test of *Paphiopedilum* section *Barbata*. Red: incorrectly classified. Green: Correctly classified. Yellow background and brown text: Unable to classify (to one section). Yellow background and green text: Unable to classify to one section, but the correct one is in the options.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  |  | ***Paphiopedilum* sect. *Barbata* run1\_2** | | | |  |  |
|  |  |  |  |  |  |  |  |
| **ANN** | **41** | **42** | **43** | **44** | **45** | **46** | **47** |
| **Desired Error** | 1,00E-04 | 1,00E-05 | 1,00E-06 | 1,00E-07 | 1,00E-08 | 1,00E-09 | 1,00E-10 |
| **Time to classify (s)** | 16 | 15 | 15 | 15 | 14 | 15 | 14 |
| **Pictures** |  |  |  |  |  |  |  |
| acmodontum1 | Incorrect 1 | Incorrect 1 | Unknown 7/1 | Incorrect 1 | Incorrect 1 | Incorrect 1 | Incorrect 1 |
| appletonianum1 | Incorrect 3 | Unknown | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 6 | Incorrect 6 |
| argus1 | Unknown 7/3 | Unknown | Correct | Unknown 7/3 | Unknown 7/3 | Correct | Unknown 3/7 |
| argus2\_2 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| barbatum1 | Incorrect 1 (low) | Unknown 7/3 | Unknown 3/1(low) | Incorrect 3 | Unknown 7/3 | Incorrect 3 | Unknown |
| barbatum2\_2 | Incorrect 3 | Incorrect 3(low) | Unknown | Unknown | Incorrect 3 | Unknown | Incorrect 3 |
| bougainvilleanum1\* | Unknown 7/1(low) | Incorrect 3 | Unknown | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 |
| bullenianum1 | Unknown 3/1(low) | Incorrect 3 | Incorrect 1 | Unknown 1/3 | Incorrect 3 | Incorrect 1 | Unknown |
| callosum1 | Incorrect 6 | Unknown 6/3(low) | Unknown | Incorrect 1(low) | Incorrect 1 | Incorrect 6 | Incorrect 6 |
| callosum2 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 1 | Unknown | Incorrect 3 |
| callosum3 | Unknown | Incorrect 3 | Incorrect 2 | Unknown | Incorrect 3 | Unknown | Correct |
| dayanum1 | Unknown | Unknown | Unknown | Unknown | Correct | Incorrect 6 | Unknown |
| dayanum2\_3 | Unknown 7/2 | Correct | Unknown 7/3 | Unknown | Incorrect 6 | Correct | Unknown |
| dayanum2 | Incorrect 3 | Incorrect 3 | Unknown | Unknown 3/1(low) | Unknown | Incorrect 1 | Unknown |
| fowliei1\* | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Unknown | Correct | Correct |
| fowliei2\* | Unknown | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 |
| hennisianum1 | Unknown 3/1 | Incorrect 1 | Incorrect 3 | Incorrect 1 | Incorrect 1 | Incorrect 3 | Unknown 1/7 |
| hennisianum2\_1 | Unknown 3/7(low) | Incorrect 3 | Correct | Unknown 7/3(low) | Unknown 7/3 | Correct | Unknown |
| hennisianum2 | Incorrect 3 | Correct(low) | Incorrect 3 | Unknown | Incorrect 3 | Incorrect 5 | Unknown |
| hookerae1 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3(low) | Incorrect 3 | Incorrect 3 |
| hookerae2\_4 | Correct | Correct | Incorrect 3 | Correct | Correct | Correct | Correct |
| javanicum1\* | Correct | Correct | Correct | Unknown 1/2/7(low) | Correct | Correct | Correct |
| javanicum2\* | Unknown 7/1 | Unknown | Unknown 7/1 | Incorrect 1 | Unknown | Correct | Unknown |
| lawrenceanum1 | Correct | Unknown | Unknown | Incorrect 5 | Unknown | Unknown | Unknown |
| mastersianum1 | Incorrect 3 | Incorrect 3 | Unknown 3/1(low) | Incorrect 3 | Incorrect 3 | Unknown 3/7 | Correct |
| mastersianum2\_5 | Correct | Correct | Correct | Correct | Correct | Correct | Incorrect 5 |
| papuanum1\* | Correct | Incorrect 1 | Incorrect 1 | Incorrect 1 | Correct | Unknown | Correct |
| purpuratum1 | Incorrect 1 | Incorrect 1 | Incorrect 6 | Incorrect 1 | Unknown 1/3 | Incorrect 1(low) | Unknown 1/3 |
| purpuratum2\_1 | Unknown 7/3 | Correct | Unknown 3/7 | Correct | Correct | Unknown 7/3 | Correct |
| purpuratum2 | Incorrect 1 | Incorrect 1 | Unknown 1/7 | Incorrect 1 | Incorrect 1 | Incorrect 1 | Unknown 1/7 |
| schoseri1\* | Incorrect 1 | Unknown | Unknown | Unknown | Unknown | Incorrect 6(low) | Unknown |
| sukhakulii1 | Incorrect 1 | Unknown | Incorrect 1 | Correct | Unknown | Incorrect 3 | Unknown |
| superbiens1§ | Unknown 3/7 | Correct | Incorrect 3 | Correct | Incorrect 3(low) | Unknown 7/3 | Incorrect 3 |
| tonsum1 | Correct | Unknown 7/1 | Correct | Correct | Incorrect 1 | Correct | Correct |
| tonsum2\_2 | Correct | Correct | Correct | Correct | Unknown | Incorrect 3 | Incorrect 3 |
| tonsum2 | Correct | Correct | Unknown 3/7 | Correct | Unknown | Correct | Incorrect 3 |
| tonsum3 | Correct | Correct | Unknown | Unknown | Unknown | Correct | Unknown |
| urbanianum1 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3(low) | Unknown 1/7 | Correct | Unknown 3/7 |
| urbanianum2\_3 | Unknown | Correct | Correct | Unknown | Correct | Correct | Unknown |
| violascens1 | Incorrect 1 | Unknown 7/1 | Incorrect 3 | Correct | Incorrect 3 | Incorrect 3 | Unknown 3/1 |
| violascens2 | Unknown | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Unknown 3/7 | Incorrect 3 |
| wardii1 | Incorrect 1 (low) | Unknown 3/1 | Unknown 7/1 | Incorrect 1 | Incorrect 1 | Incorrect 1 | Unknown |
| wardii2\_1 | Correct | Correct | Unknown 7/1 | Correct | Unknown 5/1 | Unknown 7/5 | Incorrect 5 |
| wentworthianum1\* | Incorrect 1 | Unknown | Incorrect 4 | Incorrect 1 | Incorrect 1 | Incorrect 1(low) | Unknown |
| wentworthianum2\* | Unknown | Incorrect 3 | Incorrect 4 | Unknown | Unknown | Unknown | Unknown |
| **Pictures** | **41** | **42** | **43** | **44** | **45** | **46** | **47** |
|  |  |  |  |  |  |  |  |
|  |  |  | 1:*Parvisepalum* | 2:*Pardalopetalum* | 3:*Paphiopedilum* | 4:*Coryopedilum* | 5: *Cochlopetalum* |
|  | \*= not in train set / on dear.smartweb.tw | | 6:*Brachypetalum* | 7:*Barbata* |  |  |  |
|  | §=only var. Curtisii in train set | | The order in Unknown x/y is <best\_sore>/<lower\_score>. No numbers? All scores are negative | | | | |

Table 3 Results of first test of extra pictures of *Paphiopedilum* section *Barbata*. Red: incorrect classified. Green: Correct classified. Yellow background and brown text: Unable to classify (to one section). Yellow background and green text: Unable to classify to one section, but the correct one is in the options.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  |  |  | ***Paphiopedilum* sect. *Barbata* run1\_2** | |  |  |  |
|  |  |  |  |  |  |  |  |
| **ANN** | **41** | **42** | **43** | **44** | **45** | **46** | **47** |
| **Desired Error** | 1,00E-04 | 1,00E-05 | 1,00E-06 | 1,00E-07 | 1,00E-08 | 1,00E-09 | 1,00E-10 |
| **Time to classify (s)** | 2 | 2 | 2 | 2 | 2 | 1 | 2 |
| **Picture** |  |  |  |  |  |  |  |
| paph\_argus | Correct | Unknown | Correct | Unknown 2/7 | Correct | Correct | Correct |
| Paptonsum\_blaine | Correct | Unknown | Unknown | Correct | Unknown | Unknown | Unknown |
| Paphtonsum\_elle | Unknown | Incorrect 1 | Incorrect 3 | Unknown | Incorrect 3 | Incorrect 1 | Unknown |
| Paphwardii | Incorrect 3 | Incorrect 3 | Correct | Incorrect 3 | Incorrect 3 | Incorrect 3 | Unknown |
|  |  |  |  |  |  |  |  |
|  |  |  | Legend Incorrect/Unknown | | |  |  |
|  |  |  |  |  |  |  |  |
|  |  |  |  | 1 | *Parvisepalum* | 2 | *Pardalopetalum* |
|  |  |  |  | 3 | *Paphiopedilum* | 4 | *Coryopedilum* |
|  |  |  |  | 5 | *Cochlopetalum* | 6 | *Brachypetalum* |
|  |  |  |  | 7 | *Barbata* |  |  |
|  |  |  | The order in Unknown x/y is <best\_score>/<lower\_score>. No numbers? All scores are negative! | | | | |

Table 4 Results of first test of *Paphiopedilum* section *Brachypetalum.* Red: incorrect classified. Green: Correct classified. Yellow background and brown text: Unable to classify (to one section). Yellow background and green text: Unable to classify to one section, but the correct one is in the options.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  |  | ***Paphiopedilum* sect. *Brachypetalum* run1** | | | |  |  |
|  |  |  |  |  |  |  |  |
| **ANN** | **41** | **42** | **43** | **44** | **45** | **46** | **47** |
| **Desired Error** | 1,00E-04 | 1,00E-05 | 1,00E-06 | 1,00E-07 | 1,00E-08 | 1,00E-09 | 1,00E-10 |
| **Time to classify (s)** | 8 | 7 | 7 | 7 | 7 | 7 | 8 |
| **Pictures** |  |  |  |  |  |  |  |
| bellatum1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| concolor1 | Correct | Unknown 6/1 | Correct | Correct | Unknown | Correct | Correct |
| concolor2\_2 | Correct | Correct | Unknown 6/1 | Correct | Unknown 6/1 | Correct | Correct |
| concolor2\_4 | Correct | Correct | Correct | Correct | Correct | Unknown 6/3 | Correct |
| concolor2\_6 | Correct | Unknown 6/1 | Unknown 6/1(low) | Correct | Unknown 6/1 | Correct | Correct |
| concolor2 | Correct | Correct | Correct | Correct | Unknown 6/3 | Unknown | Correct |
| godefroyae1 | Unknown 6/1(low) | Unknown | Unknown 6/7 | Unknown 6/1 | Unknown 7/6 | Correct | Correct |
| godefroyae2\_11 | Unknown 6/3 | Correct | Unknown 6/3 | Correct | Correct | Correct | Correct |
| godefroyae2\_13 | Correct | Unknown | Unknown 1/3 | Correct | Correct | Correct | Correct(low) |
| godefroyae2\_15 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| godefroyae2\_1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| godefroyae2\_3 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| godefroyae2\_5 | Correct | Correct | Correct | Correct | Correct | Correct | Unknown 6/3 |
| godefroyae2\_7 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| godefroyae2\_9 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| godefroyae2 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| niveum1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| niveum2\_4 | Unknown 6/2(low) | Correct | Correct | Correct | Correct | Correct | Correct |
| niveum2\_5 | Correct | Correct | Correct | Correct | Correct | Unknown | Correct |
| niveum2\_7 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
|  |  |  | 1:*Parvisepalum* | 2:*Pardalopetalum* | 3:*Paphiopedilum* | 4:*Coryopedilum* | 5: *Cochlopetalum* |
|  |  |  | 6:*Brachypetalum* | 7:*Barbata* |  |  |  |
|  |  |  | The order in Unknown x/y is <best\_score>/<lower\_score>. No numbers? All scores are negative! | | | | |

Table 5 Results of first test of *Paphiopedilum* section *Coryopedilum*. Red: incorrect classified. Green: Correct classified. Yellow background and brown text: Unable to classify (to one section). Yellow background and green text: Unable to classify to one section, but the correct one is in the options.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  |  | ***Paphiopedilum* sect. *Coryopedilum* run1** | | | |  |  |
|  |  |  |  |  |  |  |  |
| **ANN** | **41** | **42** | **43** | **44** | **45** | **46** | **47** |
| **Desired Error** | 1,00E-04 | 1,00E-05 | 1,00E-06 | 1,00E-07 | 1,00E-08 | 1,00E-09 | 1,00E-10 |
| **Time to classify (s)** | 4 | 3 | 3 | 3 | 3 | 3 | 3 |
| **Pictures** |  |  |  |  |  |  |  |
| glanduliferum3 | Incorrect 3 | Unknown | Unknown | Incorrect 3 | Incorrect 3 | Unknown | Unknown |
| rothschildianium1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| wilhelminiae1 | Unknown | Unknown 4/3 | Unknown | Unknown | Correct | Correct | Correct |
| philippinense1 | Correct | Correct | Unknown | Incorrect 1 | Unknown | Unknown | Unknown |
| rothschildianium2 | Unknown 4/5 | Unknown | Incorrect 1 | Correct | Correct | Correct | Correct |
| rothschildianium3 | Correct | Correct | Incorrect 1(low) | Incorrect 1(low) | Unknown | Unknown 3/4 | Correct |
|  |  |  |  |  |  |  |  |
|  |  |  |  |  |  |  |  |
|  |  |  | Legend Incorrect/Unknown | |  |  |  |
|  |  |  |  |  |  |  |  |
|  |  |  |  | 1 | *Parvisepalum* | 2 | *Pardalopetalum* |
|  |  |  |  | 3 | *Paphiopedilum* | 4 | *Coryopedilum* |
|  |  |  |  | 5 | *Cochlopetalum* | 6 | *Brachypetalum* |
|  |  |  |  | 7 | *Barbata* |  |  |

Table 6 Results of first test of *Paphiopedilum* section *Parvisepalum*. Red: incorrect classified. Green: Correct classified. Yellow background and brown text: Unable to classify (to one section). Yellow background and green text: Unable to classify to one section, but the correct one is in the options.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  |  | ***Paphiopedilum* sect. *Parvisepalum* run1** | | | |  |  |
|  |  |  |  |  |  |  |  |
| **ANN** | **41** | **42** | **43** | **44** | **45** | **46** | **47** |
| **Desired Error** | 1,00E-04 | 1,00E-05 | 1,00E-06 | 1,00E-07 | 1,00E-08 | 1,00E-09 | 1,00E-10 |
| **Time to classify (s)** | 15 | 15 | 15 | 15 | 14 | 15 | 15 |
| **Pictures** |  |  |  |  |  |  |  |
| armeniacum1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| armeniacum2\_4 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| armeniacum2\_5 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| armeniacum2\_6 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| armeniacum2\_7 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| armeniacum2\_8 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| delenatii1 | Correct | Unknown | Unknown | Unknown | Correct | Correct | Correct(low) |
| delenatii2\_2 | Correct | Correct(low) | Correct | Correct | Correct | Correct | Unknown 1/7 |
| delenatii2\_4 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| delenatii2\_6 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| delenatii2 | Correct | Correct | Correct | Correct | Unknown 7/1 | Correct | Correct |
| emersonii1 | Unknown 3/1 | Unknown | Correct | Incorrect 3(low) | Correct | Correct | Unknown |
| emersonii2\_1 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Unknown | Incorrect 3 |
| emersonii2\_3 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| emersonii2\_5 | Correct | Correct | Unknown 1/3 | Correct(low) | Correct | Correct | Correct |
| malipoense1 | Correct | Correct | Unknown 3/1 | Unknown | Incorrect 3(low) | Correct | Incorrect 3 |
| malipoense2\_10 | Correct | Correct | Correct | Correct | Unknown 1/7 | Correct | Correct |
| malipoense2\_12 | Correct | Correct | Unknown 3/1 | Correct | Correct | Correct | Correct |
| malipoense2\_14 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| malipoense2\_16 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| malipoense2\_18 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| malipoense2\_2 | Correct | Correct | Correct | Correct | Correct | Unknown 1/3 | Correct |
| malipoense2\_4 | Unknown 1/7 | Correct | Correct | Correct | Correct | Correct | Correct |
| malipoense2\_6 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| malipoense2\_8 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| malipoense2 | Unknown | Unknown | Unknown | Correct | Correct | Unknown | Correct |
| malipoense3 | Correct | Unknown | Incorrect 7 | Unknown | Correct | Unknown | Correct |
| micranthum1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2\_11 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2\_13 | Correct | Correct | Unknown | Correct | Correct | Unknown | Correct |
| micranthum2\_15 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2\_17 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2\_19 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2\_1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct(low) |
| micranthum2\_21 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2\_3 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2\_5 | Correct | Correct | Incorrect 3 | Correct | Incorrect 3 | Unknown | Unknown |
| micranthum2\_7 | Correct (very low) | Correct | Unknown 7/1(low) | Correct | Incorrect 7 | Incorrect 5 | Unknown |
| micranthum2\_9 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2 | Correct | Unknown 1/6 | Correct | Correct | Correct | Correct | Correct |
| micranthum3 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| **Picture** | **41** | **42** | **43** | **44** | **45** | **46** | **47** |
|  |  |  |  |  |  |  |  |
|  |  |  |  |  |  |  |  |
|  |  |  |  |  |  |  |  |
|  |  |  | 1:*Parvisepalum* | 2:*Pardalopetalum* | 3:*Paphiopedilum* | 4:*Coryopedilum* | 5: *Cochlopetalum* |
|  |  |  | 6:*Brachypetalum* | 7:*Barbata* |  |  |  |
|  |  |  | The order in Un-known x/y is <best\_sore>/<lower\_score>. No numbers? All scores are negative | | | | |

Table 7 Results of first test of hybrids of *Paphiopedilum* sect. *Parvisepalum* and *Brachypetalum*. Red: incorrect classified. Green: Correct classification to one of the parental sections. Green and bold: Correct classification to both parental sections. Yellow background and brown text: Unable to classify.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  |  | ***Paphiopedilum* sect. *Parvisepalum* X *Brachypetalum* run1** | | | |  |  |
|  |  |  |  |  |  |  |  |
| **ANN** | **41** | **42** | **43** | **44** | **45** | **46** | **47** |
| **Desired Error** | 1,00E-04 | 1,00E-05 | 1,00E-06 | 1,00E-07 | 1,00E-08 | 1,00E-09 | 1,00E-10 |
| **Time to classify (s)** | 7 | 5 | 5 | 5 | 5 | 4 | 5 |
| **Pictures** |  |  |  |  |  |  |  |
| armeniacumXconcolor2 | Unknown | 1 | 1 | 1 | 1 | 1 | 1 |
| delenatiiXniveum2 | 1 | Unknown | Unknown | 6 | Unknown | 1 | Unknown |
| armeniacumXconcolor1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| armeniacumXconcolor6 | 6 | ***1&6*** | 1 | 1 | 1 | 1 | 1 |
| emersoniiXbellatulum | 3&1 | 1 | 1&7 | Unknown | Unknown | 3 | 1 |
| delenatiiXniveum1 | 6 | 6 | 6 | Unknown | 6(low) | 6 | 6 |
| armeniacumXconcolor5 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| armeniacumXconcolor4 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
|  |  |  |  |  |  |  |  |
|  |  |  |  |  |  |  |  |
|  |  |  |  |  |  |  |  |
|  |  |  | Legend |  |  |  |  |
|  |  |  |  | 1 | *Parvisepalum* | 2 | *Pardalopetalum* |
|  |  |  |  | 3 | *Paphiopedilum* | 4 | *Coryopedilum* |
|  |  |  |  | 5 | *Cochlopetalum* | 6 | *Brachypetalum* |
|  |  |  |  | 7 | *Barbata* |  |  |

## Results of characteristic research

Table 8 Characteristics o*f Paphiopedilum* section *Barbata*

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  |  | ***Paphiopedilum* sect. *Barbata*** | | |  |  |  |
|  |  |  |  |  |  |  |  |
| **Characteristics** | | **argus1\*** | **argus2\_2§** | **mastersianum1\*** | **mastersianum2\_5§** | **tonsum1\*$** | **tonsum2\_2§** |
|  | |  |  |  |  |  |  |
| spathulate petals | | **!** | ✓ | ✓ | ✓ | **!** | ✓ |
| spotted and/or warted petals | | ✓ | ✓ | 🗶 | ✓ | ✓ | ✓ |
| incurved side lobes | | ✓ | ✓ | ✓ | ✓ | ✓ | ✓ |
|  | Note | artificial white |  | artificial white |  | artificial white | artificial white |
|  |  | patches caused |  | patches caused |  | patches caused | patches caused |
|  |  | by scanning |  | by scanning |  | by scanning + | by scanning |
|  |  |  |  |  |  | Small pic. |  |
|  |  |  |  |  |  |  |  |
| **Characteristics** | | **tonsum2\*** | **tonsum3\*** | **wardii1\*** | **wardii2\_1§** |  |  |
|  | |  |  |  |  |  |  |
| spathulate petals | | ✓ | ✓ | 🗶 | ✓ |  |  |
| spotted and/or warted petals | | ✓ | ✓ | ✓ | ✓ |  |  |
| incurved side lobes | | ✓ | ✓ | ✓ | ✓ |  |  |
|  | Note | artificial white | artificial white | artificial white |  |  |  |
|  |  | patches caused | patches caused | patches caused |  |  |  |
|  |  | by scanning | by scanning | by scanning |  |  |  |
| ✓ | Visible |  |  |  |  |  |  |
| **!** | Difficult to see | |  |  |  |  |  |
| 🗶 | Invisible |  |  |  |  |  |  |
| \* | Picture scanned from Cribb, 1998 | | |  |  |  |  |
| § | Picture from dear.smartweb.tw | | |  |  |  |  |
| **Name** | Incorrectly identified to sectional level in max 1 of the 7 different networks | | | | |  |  |
| **Name** | Incorrectly identified to sectional level in max 3 of the 7 different networks | | | | |  |  |
| **Name** | Incorrectly identified to sectional level in more than 3 of the 7 networks | | | | |  |  |

Table 9 Characteristics of *Paphiopedilum* section *Brachypetalum*.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
|  |  | ***Paphiopedilum* sect. *Brachypetalum*** | | |  |  |
|  |  |  |  |  |  |  |
|  |  |  |  |  |  |  |
| **Characteristics** | | **concolor1\*** | **concolor2\_2§** | **concolor2\_4§** | **concolor2\_6§** | **concolor2\*** |
|  | |  |  |  |  |  |
| variously ornamented with purple spots | | ✓ | ✓ | ✓ | ✓ | ✓ |
| broad-elliptic petals | | ✓ | ✓ | ✓ | ✓ | **!** |
| small lip | | ✓ | ✓ | ✓ | ✓ | ✓ |
| ovoid lip | | ✓ | ✓ | ✓ | 🗶 | ✓ |
| lip with incurved margin | | ✓ | 🗶 | ✓ | ✓ | ✓ |
| pale yellow or white flower | | 🗶 | ✓ | ✓ | ✓ | ✓ |
|  | |  |  |  |  |  |
|  |  |  |  |  |  |  |
|  |  |  |  |  |  |  |
| ✓ | Visible |  |  |  |  |  |
| **!** | Difficult to see |  |  |  |  |  |
| 🗶 | Invisible |  |  |  |  |  |
| \* | Picture scanned from Cribb, 1998 | |  |  |  |  |
| § | Picture from dear.smartweb.tw | |  |  |  |  |
| **Name** | Incorrectly identified to sectional level in max 1 of the 7 different networks | | | |  |  |
| **Name** | Incorrectly identified to sectional level in max 3 of the 7 different networks | | | |  |  |
| **Name** | Incorrectly identified to sectional level in more than 3 of the 7 networks | | | |  |  |

Table 10 Characteristics of *Paphiopedilum* section *Coryopedilum*.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
|  |  | ***Paphiopedilum* sect. *Coryopedilum*** | | |
|  |  |  |  |  |
|  |  |  |  |  |
| **Characteristics** | | **rothschildianium1\*** | **rothschildianium2\*** | **rothschildianium3\*** |
|  | |  |  |  |
| Long petals | | ✓ | ✓ | ✓ |
| Petals hanging down | | ✓ | ✓ | ✓ |
| Tapering petals | | ✓ | ✓ | ✓ |
| Spirally twisted petals | | ✓ | ✓ | **!** |
| Warts on margins of petals | | 🗶 | 🗶 | 🗶 |
| Glandular at tip | | 🗶 | 🗶 | 🗶 |
| incurved side lobes | | ✓ | 🗶 | 🗶 |
|  |  |  |  |  |
|  |  |  |  |  |
|  |  |  |  |  |
| ✓ | Visible |  |  |  |
| **!** | Difficult to see | |  |  |
| 🗶 | Invisible |  |  |  |
| \* | Picture scanned from Cribb, 1998 | |  |  |
| § | Picture from dear.smartweb.tw | |  |  |
| **Name** | Incorrectly identified to sectional level in max 1 of the 7 different networks | | | |
| **Name** | Incorrectly identified to sectional level in max 3 of the 7 different networks | | | |
| **Name** | Incorrectly identified to sectional level in more than 3 of the 7 networks | | | |

Table 11 Characteristics of *Paphiopedilum* section *Parvisepalum*.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
|  |  | ***Paphiopedilum* sect. *Parvisepalum*** | | |  |  |
|  |  |  |  |  |  |  |
| **Characteristics** | | **delenatii1\*** | **delenatii2\_2§** | **delenatii2\_4§** | **delenatii2\_6§** | **delenatii2\*** |
|  | |  |  |  |  |  |
| broad-elliptic petals | | ✓ | ✓ | ✓ | ✓ | ✓ |
| subcircular petals | | ✓ | ✓ | ✓ | ✓ | **!** |
| large staminode | | ✓ | ✓ | ✓ | ✓ | ✓ |
| cream/yellow coloured petals | | 🗶 | ✓ | ✓ | ✓ | 🗶 |
|  | Note | artificial white |  |  |  | artificial brown |
|  |  | patches caused |  |  |  | patches caused |
|  |  | by scanning |  |  |  | by scanning |
|  |  |  |  |  |  |  |
| **Characteristics** | | **emersonii1\*** | **emersonii2\_1§** | **emersonii2\_3§** | **emersonii2\_5§** | **malipoense1\*** |
|  | |  |  |  |  |  |
| broad-elliptic petals | | ✓ | ✓ | ✓ | ✓ | ✓ |
| subcircular petals | | ✓ | ✓ | ✓ | ✓ | 🗶 |
| large staminode | | ✓ | ✓ | ✓ | ✓ | ✓ |
| cream/yellow coloured petals | | 🗶 | 🗶 | ✓ | 🗶 | ✓ |
|  | Note | artificial white | Flower dark |  | Flower grey/ |  |
|  |  | patches caused |  |  | blue like |  |
|  |  | by scanning |  |  |  |  |
|  |  |  |  |  |  |  |
| **Characteristics** | | **malipoense2\_10§** | **malipoense2\_12§** | **malipoense2\_14§** | **malipoense2\_8§** | **malipoense2\*** |
|  | |  |  |  |  |  |
| broad-elliptic petals | | ✓ | ✓ | ✓ | ✓ | **!** |
| subcircular petals | | ✓ | ✓ | ✓ | ✓ | 🗶 |
| large staminode | | ✓ | ✓ | ✓ | ✓ | ✓ |
| cream/yellow coloured petals | | ✓ | ✓ | ✓ | ✓ | 🗶 |
|  | Note |  |  |  |  | Not fully frontal |
| **Characteristics** | | **malipoense3\*** | **malipoense2\_16§** | **malipoense2\_18§** | **malipoense2\_2§** | **malipoense2\_4§** |
|  | |  |  |  |  |  |
| broad-elliptic petals | | **!** | ✓ | ✓ | ✓ | ✓ |
| subcircular petals | | ✗ | ✓ | ✓ | ✓ | ✓ |
| large staminode | | ✓ | ✓ | ✓ | ✓ | ✓ |
| cream/yellow coloured petals | | ✗ | ✓ | ✓ | ✓ | ✓ |
|  | Note | Not fully frontal |  |  |  |  |
|  |  |  |  |  |  |  |
| **Characteristics** | | **malipoense2\_6§** |  |  |  |  |
|  | |  |  |  |  |  |
| broad-elliptic petals | | ✓ |  |  |  |  |
| subcircular petals | | ✓ |  |  |  |  |
| large staminode | | ✓ |  |  |  |  |
| cream/yellow coloured petals | | ✓ |  |  |  |  |
|  |  |  |  |  |  |  |
|  |  |  |  |  |  |  |
| ✓ | Visible |  |  |  |  |  |
| **!** | Difficult to see |  |  |  |  |  |
| 🗶 | Invisible |  |  |  |  |  |
| \* | Picture scanned from Cribb, 1998 | |  |  |  |  |
| § | Picture from dear.smartweb.tw | |  |  |  |  |
| **Name** | Incorrectly identified to sectional level in max 1 of the 7 different networks | | | |  |  |
| **Name** | Incorrectly identified to sectional level in max 3 of the 7 different networks | | | |  |  |
| **Name** | Incorrectly identified to sectional level in more than 3 of the 7 networks | | | |  |  |

## Results of second test run

Table 12 Results of second test run of *Paphiopedilum* section *Barbata*. Red: incorrect classified. Green: Correct classified. Yellow background and brown text: Unable to classify (to one section). Yellow background and green text: Unable to classify to one section, but the correct one is in the options.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  |  | ***Paphiopedilum* sect. *Barbata* run2\_1** | | | |  |  |
|  |  |  |  |  |  |  |  |
| **ANN** | **51** | **52** | **53** | **54** | **55** | **56** | **57** |
| **Desired Error** | 1,00E-04 | 1,00E-05 | 1,00E-06 | 1,00E-07 | 1,00E-08 | 1,00E-09 | 1,00E-10 |
| **Time to classify (s)** | 17 | 15 | 14 | 15 | 16 | 14 | 14 |
| **Pictures** |  |  |  |  |  |  |  |
| acmodontum1 | Unknown 1/2 | Incorrect 1 | Incorrect 1 | Incorrect 1 | Unknown 7/1 | Incorrect 1 | Incorrect 1 |
| appletonianum1 | Incorrect 1 | Unknown | Incorrect 3(low) | Incorrect 6 | Unknown | Unknown | Incorrect 1 |
| argus1 | Unknown 7/3 | Correct | Correct | Unknown | Incorrect 3 | Correct | Unknown 7/3 |
| argus2\_2 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| barbatum1 | Unknown | Unknown | Incorrect 3 | Incorrect 3 | Incorrect 1 | Unknown 3/7 | Incorrect 3 |
| barbatum2\_2 | Unknown | Incorrect 5 | Unknown 7/3(low) | Unknown | Incorrect 4 | Incorrect 5 | Unknown |
| bougainvilleanum1\* | Unknown 7/3 | Incorrect 3 | Correct | Correct | Unknown 3/7 | Unknown 7/3 | Unknown 7/3 |
| bullenianum1 | Incorrect 1 | Incorrect 1(low) | Unknown | Unknown | Incorrect 1 | Incorrect 3 | Unknown |
| callosum1 | Unknown 6/3 | Incorrect 3 | Incorrect 6 | Unknown 3/6 | Incorrect 6 | Incorrect 6 | Unknown |
| callosum2 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 |
| callosum3 | Correct | Unknown | Unknown | Correct | Correct | Correct | Correct |
| dayanum1 | Unknown | Unknown | Unknown | Unknown | Incorrect 6 | Incorrect 6(low) | Unknown |
| dayanum2\_3 | Incorrect 4(low) | Unknown 1/7 | Unknown | Unknown | Unknown | Unknown 4/7 | Unknown |
| dayanum2 | Incorrect 3(low) | Incorrect 3 | Unknown 5/4 | Unknown | Unknown | Unknown | Incorrect 5 |
| fowliei1\* | Unknown low:3/7 | Unknown 3/7(low) | Correct | Incorrect 3 | Incorrect 3(low) | Incorrect 3 | Unknown |
| fowliei2\* | Incorrect 3 | Incorrect 3 | Incorrect 3(low) | Unknown | Incorrect 3 | Incorrect 3(low) | Incorrect 3 |
| hennisianum1 | Unknown | Incorrect 1(low) | Unknown | Unknown | Incorrect 1 | Incorrect 1 | Incorrect 1 |
| hennisianum2\_1 | Correct | Incorrect 3 | Correct | Correct | Unknown 7/3 | Correct | Correct |
| hennisianum2 | Incorrect 3(low) | Unknown | Unknown | Unknown | Unknown | Unknown | Incorrect 3 |
| hookerae1 | Incorrect 3 | Incorrect 3 | Unknown 3/7 | Incorrect 3 | Incorrect 3 | Unknown | Incorrect 3 |
| hookerae2\_4 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| javanicum1\* | Correct | Correct | Correct | Correct | Correct | Unknown 7/3 | Correct |
| javanicum2\* | Unknown | Incorrect 1 | Correct | Correct(low) | Unknown 7/1 | Unknown 7/1 | Incorrect 1(low) |
| lawrenceanum1 | Incorrect 4 | Unknown | Correct | Unknown | Unknown | Unknown | Unknown |
| mastersianum1 | Unknown | Correct | Unknown 7/3(low) | Correct | Correct | Correct | Correct(low) |
| mastersianum2\_5 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| papuanum1\* | Unknown 4/7/1(L) | Unknown | Unknown 7/6/1 | Unknown | Correct | Unknown 3/7(low) | Incorrect 5 |
| purpuratum1 | Incorrect 1 | Incorrect 1 | Incorrect 1 | Unknown 1/3(low) | Unknown | Incorrect 1 | Incorrect 1 |
| purpuratum2\_1 | Unknown 7/3 | Unknown 7/5 | Correct | Correct | Correct | Correct | Correct |
| purpuratum2 | Incorrect 1 | Unknown 7/1 | Unknown 1/7 | Unknown 1/7 | Unknown 7/1 | Incorrect 1 | Incorrect 1 |
| schoseri1\* | Unknown | Unknown | Unknown | Unknown | Incorrect 6 | Unknown | Incorrect 1 |
| sukhakulii1 | Incorrect 1 | Unknown 1/7(low) | Incorrect 3 | Incorrect 1 | Incorrect 4 | Incorrect 1 | Unknown |
| superbiens1§ | Unknown | Unknown 7/3 | Correct | Correct(low) | Unknown 3/7(L) | Unknown | Correct |
| tonsum1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| tonsum2\_2 | Correct | Correct | Incorrect 5 | Correct | Correct | Correct | Correct |
| tonsum2 | Unknown | Correct | Correct | Unknown | Correct | Correct | Correct |
| tonsum3 | Correct | Incorrect 3 | Correct | Correct | Unknown | Incorrect 3 | Unknown |
| urbanianum1 | Correct | Unknown 7/1(low) | Incorrect 3 | Unknown | Unknown | Unknown | Incorrect 3(low) |
| urbanianum2\_3 | Correct(low) | Correct | Correct | Correct(low) | Correct | Correct(low) | Correct |
| violascens1 | Unknown 3/7(low) | Incorrect 3 | Incorrect 3 | Unknown | Correct(low) | Unknown | Incorrect 3 |
| violascens2 | Unknown 7/3 | Incorrect 3(low) | Incorrect 3(low) | Correct | Unknown 3/7 | Unknown | Incorrect 3 |
| wardii1 | Incorrect 1 | Unknown 1/3(low) | Incorrect 1 | Incorrect 1 | Unknown 3/1 | Incorrect 1 | Unknown |
| wardii2\_1 | Correct | Correct | Correct | Correct | Correct | Correct | Unknown 1/7(low) |
| wentworthianum1\* | Correct | Unknown | Unknown | Correct(low) | Unknown | Unknown | Unknown |
| wentworthianum2\* | Unknown | Unknown | Correct | Unknown | Unknown | Unknown | Unknown |
| **Pictures** | **51** | **52** | **53** | **54** | **55** | **56** | **57** |
|  |  |  |  |  |  |  |  |
|  |  |  | 1:*Parvisepalum* | 2:*Pardalopetalum* | 3:*Paphiopedilum* | 4:*Coryopedilum* | 5: *Cochlopetalum* |
|  | \*= not in train set / on dear.smartweb.tw | | 6:*Brachypetalum* | 7:*Barbata* | L:low |  |  |
|  | §= only var. Curtisii in train set. | | The order in Unknown x/y is <best\_sore>/<lower\_score>. No numbers? All scores are negative | | | | |

Table 13 Results of second test of the extra pictures of *Paphiopedilum* section *Barbata*. Red: incorrect classified. Green: Correct classified. Yellow background and brown text: Unable to classify (to one section). Yellow background and green text: Unable to classify to one section, but the correct one is in the options.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  |  | ***Paphiopedilum* sect. *Barbata* run2\_2** | | | |  |  |
|  |  |  |  |  |  |  |  |
| **ANN** | **51** | **52** | **53** | **54** | **55** | **56** | **57** |
| **Desired Error** | 1,00E-04 | 1,00E-05 | 1,00E-06 | 1,00E-07 | 1,00E-08 | 1,00E-09 | 1,00E-10 |
| **Time to classify (s)** | 2 | 2 | 2 | 2 | 2 | 2 | 2 |
|  |  |  |  |  |  |  |  |
| paph\_argus | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| Paptonsum\_blaine | Incorrect 1(low) | Unknown | Unknown | Correct(low) | Correct | Correct | Correct |
| Paphtonsum\_elle | Incorrect 1 | Incorrect 3(low) | Unknown | Unknown | Unknown | Incorrect 1 | Incorrect 1(low) |
| Paphwardii | Incorrect 3 | Incorrect 3 | Unknown 3/6 | Incorrect 3 | Unknown 3/7 | Unknown | Incorrect 3 |
|  |  |  |  |  |  |  |  |
|  |  |  | Legend Incorrect/Unknown | |  |  |  |
|  |  |  |  |  |  |  |  |
|  |  |  |  | 1 | *Parvisepalum* | 2 | *Pardalopetalum* |
|  |  |  |  | 3 | *Paphiopedilum* | 4 | *Coryopedilum* |
|  |  |  |  | 5 | *Cochlopetalum* | 6 | *Brachypetalum* |
|  |  |  |  | 7 | *Barbata* |  |  |
|  |  |  | The order in Unknown x/y is <best\_score>/<lower\_score>/ No numbers? All scores are negative | | | | |

Table 14 Results of second test run of *Paphiopedilum* section *Brachypetalum*. Red: incorrect classified. Green: Correct classified. Yellow background and brown text: Unable to classify (to one section). Yellow background and green text: Unable to classify to one section, but the correct one is in the options.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  |  | ***Paphiopedilum* sect. *Brachypetalum* run2** | | |  |  |  |
|  |  |  |  |  |  |  |  |
| **ANN** | **51** | **52** | **53** | **54** | **55** | **56** | **57** |
| **Desired Error** | 1,00E-04 | 1,00E-05 | 1,00E-06 | 1,00E-07 | 1,00E-08 | 1,00E-09 | 1,00E-10 |
| **Time to classify (s)** | 7 | 7 | 8 | 7 | 7 | 6 | 7 |
| **Pictures** |  |  |  |  |  |  |  |
| bellatum1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| concolor1 | Correct | Correct | Correct | Unknown | Correct | Correct | Correct |
| concolor2\_2 | Correct | Unknown 6/1 | Unknown 6/1 | Correct | Unknown 6/1 | Unknown 6/1 | Correct |
| concolor2\_4 | Correct | Correct | Unknown 6/1 | Unknown 6/1 | Correct | Unknown 6/1 | Correct |
| concolor2\_6 | Correct | Correct | Unknown 6/1 | Correct | Unknown 6/1 | Unknown 6/1 | Correct |
| concolor2 | Correct | Correct | Correct | Correct | Unknown | Correct | Correct |
| godefroyae1 | Unknown | Unknown | Unknown 6/7 | Correct | Unknown 6/1 | Incorrect 1 | Incorrect 1 |
| godefroyae2\_11 | Correct | Unknown | Correct | Correct | Correct | Correct | Correct |
| godefroyae2\_13 | Correct | Unknown | Correct | Correct | Correct | Correct | Correct |
| godefroyae2\_15 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| godefroyae2\_1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| godefroyae2\_3 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| godefroyae2\_5 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| godefroyae2\_7 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| godefroyae2\_9 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| godefroyae2 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| niveum1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| niveum2\_4 | Unknown | Correct | Correct | Correct | Correct | Unknown 6/1 | Correct |
| niveum2\_5 | Correct | Correct | Unknown 6/3(low) | Unknown | Correct | Correct | Correct |
| niveum2\_7 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
|  |  |  |  | 1:Parvisepalum | 2:Pardalopetalum | 3:Paphiopedilum | 4:Coryopedilum |
|  |  |  |  | 5: Cochlopetalum | 6: Brachypetalum | 7:Barbata |  |
|  |  |  |  | The order in Unknown x/y is <best\_sore>/<lower\_score>. No numbers? All scores are negative! | | | |

Table 15 Results of second test of *Paphiopedilum* section *Coryopedilum*. Red: incorrect classified. Green: Correct classified. Yellow background and brown text: Unable to classify (to one section). Yellow background and green text: Unable to classify to one section, but the correct one is in the options.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  |  |  | ***Paphiopedilum* sect. *Coryopedilum* run2** | |  |  |  |
|  |  |  |  |  |  |  |  |
| **ANN** | **51** | **52** | **53** | **54** | **55** | **56** | **57** |
| **Desired Error** | 1,00E-04 | 1,00E-05 | 1,00E-06 | 1,00E-07 | 1,00E-08 | 1,00E-09 | 1,00E-10 |
| **Time to classify (s)** | 3 | 3 | 3 | 2 | 2 | 3 | 2 |
| **Picture** |  |  |  |  |  |  |  |
| glanduliferum3 | Correct | Correct | Unknown | Correct | Unknown 4/3 | Unknown | Correct |
| rothschildianium1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| wilhelminiae1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| philippinense1 | Correct | Unknown 4/1 | Correct | Correct | Unknown 4/6 | Correct | Correct |
| rothschildianium2 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| rothschildianium3 | Correct | Correct | Correct | Unknown | Correct | Correct | Correct |
|  |  |  |  |  |  |  |  |
|  |  |  | Legend Incorrect/Unknown | |  |  |  |
|  |  |  |  |  |  |  |  |
|  |  |  |  | 1 | *Parvisepalum* | 2 | *Pardalopetalum* |
|  |  |  |  | 3 | *Paphiopedilum* | 4 | *Coryopedilum* |
|  |  |  |  | 5 | *Cochlopetalum* | 6 | *Brachypetalum* |
|  |  |  |  | 7 | *Barbata* |  |  |
|  |  |  | The order in Unknown x/y is <best\_score>/<lower\_score>/ No numbers? All scores are negative | | | | |

Table 16 Results of second test of *Paphiopedilum* section *Parvisepalum*. Red: incorrect classified. Green: Correct classified. Yellow background and brown text: Unable to classify (to one section). Yellow background and green text: Unable to classify to one section, but the correct one is in the options.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  |  | ***Paphiopedilum* sect. *Parvisepalum* run2** | | | |  |  |
|  |  |  |  |  |  |  |  |
| **ANN** | **51** | **52** | **53** | **54** | **55** | **56** | **57** |
| **Desired Error** | 1,00E-04 | 1,00E-05 | 1,00E-06 | 1,00E-07 | 1,00E-08 | 1,00E-09 | 1,00E-10 |
| **Time to classify (s)** | 15 | 14 | 14 | 14 | 15 | 14 | 14 |
| **Picture** |  |  |  |  |  |  |  |
| armeniacum1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| armeniacum2\_4 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| armeniacum2\_5 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| armeniacum2\_6 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| armeniacum2\_7 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| armeniacum2\_8 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| delenatii1 | Unknown | Correct | Unknown | Correct | Correct | Correct | Unknown |
| delenatii2\_2 | Correct | Correct | Correct | Correct | Unknown | Correct | Unknown |
| delenatii2\_4 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| delenatii2\_6 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| delenatii2 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| emersonii1 | Unknown low:1/3 | Correct | Correct | Correct | Incorrect 3 | Incorrect 3 | Unknown |
| emersonii2\_1 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 | Incorrect 3 |
| emersonii2\_3 | Unknown 1/3(low) | Correct | Correct | Unknown | Unknown 3/1 | Unknown 1/3 | Correct |
| emersonii2\_5 | Unknown | Correct | Unknown | Unknown | Unknown | Unknown | Correct |
| malipoense1 | Correct | Unknown 1/3 | Unknown | Correct | Unknown 3/1(low) | Correct | Correct |
| malipoense2\_10 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| malipoense2\_12 | Correct | Correct | Correct | Correct | Incorrect 3 | Correct | Correct |
| malipoense2\_14 | Correct | Correct | Correct | Unknown | Correct | Correct | Correct |
| malipoense2\_16 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| malipoense2\_18 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| malipoense2\_2 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| malipoense2\_4 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| malipoense2\_6 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| malipoense2\_8 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| malipoense2 | Correct | Correct | Correct | Unknown | Correct | Correct | Correct |
| malipoense3 | Unknown | Unknown | Incorrect 7 | Unknown | Unknown | Incorrect 7 | Incorrect 7 |
| micranthum1 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2\_11 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2\_13 | Unknown | Unknown | Unknown | Correct | Unknown | Unknown | Unknown |
| micranthum2\_15 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2\_17 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2\_19 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2\_1 | Correct(low) | Incorrect 5 | Incorrect 6 | Correct | Correct | Correct | Unknown |
| micranthum2\_21 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2\_3 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2\_5 | Unknown | Incorrect 7 | Unknown | Correct | Unknown | Unknown | Unknown 3/1 |
| micranthum2\_7 | Correct | Correct | Correct | Correct | Correct | Correct | Correct(low) |
| micranthum2\_9 | Correct | Correct | Correct | Correct | Correct | Correct | Correct |
| micranthum2 | Correct | Correct | Incorrect 4 | Correct | Unknown | Correct | Correct |
| micranthum3 | Correct | Correct | Unknown 1/4 | Correct | Correct | Correct | Correct |
| **Picture** | **51** | **52** | **53** | **54** | **55** | **56** | **57** |
|  |  |  |  |  |  |  |  |
|  |  |  |  |  |  |  |  |
|  |  |  |  |  |  |  |  |
|  |  |  | 1:*Parvisepalum* | 2:*Pardalopetalum* | 3:*Paphiopedilum* | 4:*Coryopedilum* | 5: *Cochlopetalum* |
|  |  |  | 6:*Brachypetalum* | 7:*Barbata* |  |  |  |
|  |  |  | The order in Un-known x/y is <best\_sore>/<lower\_score>. No numbers? All scores are negative! | | | | |

Table 17 Results of second test of hybrids of *Paphiopedilum* sections *Parvisepalum* and *Brachypetalum*. Red: incorrect classified. Green: Correct classification to one of the parental sections. Green and bold: Correct classification to both parental sections. Yellow background and brown text: Unable to classify.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
|  |  | ***Paphiopedilum* sect. *Parvisepalum* X *Brachypetalum* run2** | | | |  |  |
|  |  |  |  |  |  |  |  |
| **ANN** | **51** | **52** | **53** | **54** | **55** | **56** | **57** |
| **Desired Error** | 1,00E-04 | 1,00E-05 | 1,00E-06 | 1,00E-07 | 1,00E-08 | 1,00E-09 | 1,00E-10 |
| **Time to classify (s)** | 5 | 4 | 4 | 4 | 4 | 4 | 3 |
| **Picture** |  |  |  |  |  |  |  |
| armeniacumXconcolor2 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| delenatiiXniveum2 | Unknown | Unknown | 1 | 2(low) | Unknown | 1 | 6 |
| armeniacumXconcolor1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| armeniacumXconcolor6 | 1 | 1 | 1 | 1 | 1 | 1 | 1(low) |
| emersoniiXbellatulum | 1 | 1&3 | 1&3 | 1 | Unknown | Unknown | Unknown |
| delenatiiXniveum1 | 6 | 6 | 6 | 6 | 6 | 6 | 6 |
| armeniacumXconcolor5 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| armeniacumXconcolor4 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
|  |  |  |  |  |  |  |  |
|  |  |  | Legend |  |  |  |  |
|  |  |  |  |  |  |  |  |
|  |  |  |  | 1 | *Parvisepalum* | 2 | *Pardalopetalum* |
|  |  |  |  | 3 | *Paphiopedilum* | 4 | *Coryopedilum* |
|  |  |  |  | 5 | *Cochlopetalum* | 6 | *Brachypetalum* |
|  |  |  |  | 7 | *Barbata* |  |  |

# Appendices

## Codes

### Bash

#### Create\_traindata.sh

#!/usr/bin/env bash

#Loop throug the directories

for directories in $(ls -d \*/)

do

directory=${directories%%/}

#The tuber pictures use another Perl script for training the neural network

#than the flower pictures. So check the name of the directory.

if [[ $directory == T\* ]]

#If it starts with T (=Tuber)

then

#Go into the directory

cd $directory

#Loop through the directories

for directories2 in $(ls -d \*/)

do

#Create variables

catagory=0

shape=${directories2%%/}

echo "run traindata.pl for $shape"

#Give the Look-a-like tubers catagory -1

if [[ $shape == L\* ]]

then

catagory=-1

#Give the salep tubers catagory 1

else

catagory=1

fi

#Run the traindata script and write the output to a tsv file.

#Give this file the name of the current directory

perl ../traindata.pl -d ./$shape -c $catagory > $shape.tsv

done

#After looping through the directories go out of the Tuber directory

cd ..

elif [[ $directory == F\* ]]

#If it starts with F (=Flower)

then

#Go into the directory

cd $directory

#Loop through the section directories

for section in $(ls -d \*/)

do

#Go into the section directory

cd $section

#Loop through the species directories

for species\_dir in $(ls -d \*/)

do

#Create variables

species=${species\_dir%%/}

echo "run traindata2.pl for $species"

#Run the traindata2 script and redirect the output to a tsv file.

#Give this file the name of the current directory

perl ../../traindata2.pl -d ./$species -c 0 > $species.tsv

done

#After loopging through the species directories go back to the Flower directory

cd ..

done

#After looping through the section directories go out of the Flower directory

cd ..

fi

done

#Clear the screen

clear

#Print a finish message

echo "Done"

echo "Please modify the flower data:"

echo "Run: ./modify\_flower\_data.sh"

#### modify\_flower\_data.sh

#!/usr/bin/env bash

#Go into the Flower directory

cd Flower

#Loop through the directories

for directory in $(ls -d \*/)

do

#Go into the directory

cd $directory

#Run combine\_files.py

echo "Run combine\_files.py"

python ../../combine\_files.py

#Go back to the Flower directory

cd ..

done

#Run add\_columns.py

echo "Run add\_columns.py"

python ../add\_columns.py

#Print a finising message

echo "Finished"

echo "To train a neural network use the folowing command:"

echo "perl trainai.pl -d <directory\_with\_traindata> -c <number\_of\_categories> -o <output>"

echo "To change the Desired Error add -t <desired\_error>"

echo "To change the number of epoch add -e <epochs>"

echo "To follow the run time add date; before and ;date after the command"

#### training.sh

#!/usr/bin/env bash

#Clear the screen.

clear

#===================================================================================================#

# Download pictures from Flickr #

#===================================================================================================#

#Download pictures from Flickr

python Offlickr.py -p -n -i 113733456@N06 -d .

#List all xml files and save it in a

ls | egrep xml > xml\_files.txt

#Get the tags for every picture

python get\_tags.py

#remove all xml files and the list of xml files saved in xml\_files.txt

rm \*.xml xml\_files.txt

#Clear the screen, ask the user to check manualy if al pictures are download correctly. If not, download them by hand.

clear

echo "Pictures downloaded"

echo "Please check the downloaded pictures."

echo "Download all broken pictures by hand:"

echo "log in on Flickr with the shared naturalis account and go to:"

echo "https://www.flickr.com/photos/113733456@N06/<picture\_id>/sizes/o/"

echo "Press enter to continue"

read

#===================================================================================================#

# Divide pictures between Flower and Tuber #

#===================================================================================================#

#Create the required directories

mkdir Flower Tuber Tuber/LOblong Tuber/LSpur Tuber/LRound Tuber/Oblong Tuber/Round Tuber/Spur

#Loop through every jpg file

for i in \*.jpg

do

name=(${i//./ }$0)

tags="$name""\_tags.txt"

#Conver from jpg to png

echo "Convert $i to $name.png"

convert $i $name.png

content=$(cat $tags)

echo "---------------------------------------------"

#Divide the pictiure and tags between Tuber and Flower

if [[ $content == \*Tuber\* ]]

then

mv $name.png Tuber

mv $tags Tuber

elif [[ $content == \*Flower\* ]]

then

mv $name.png Flower

mv $tags Flower

else

echo "No correct tag found"

echo "=================================================="

fi

done

#Remove the jpg that will not be used anymore

rm \*.jpg

#Clear the screen, print a message and wait for input from the user.

clear

echo "Pictures divided between flower and tuber"

echo "Press enter to continue"

read

#===================================================================================================#

# Divide the pictures in Flower between the different genera and the different species #

#===================================================================================================#

#go into the Flower directory

cd Flower

#Loop through the png files

for files in \*.png

do

name=(${files//./ }$0)

tags="$name""\_tags.txt"

#Create variables that will be used to create directories.

content=$(cat $tags)

sectioni=$(sed -n '3p' < $tags)

speciesi=$(sed -n '4p' < $tags)

section=(${sectioni##\*:})

species=(${speciesi##\*:})

#Create a direcory with the name of the section

mkdir $section

#Move the picure and tag file to the the correct directory

mv $files $section

mv $tags $section

#Go into the section directory and create a directory wiht the name of the species

cd $section

mkdir $species

#Move the picture and tag file to the correct directory

mv $files $species

mv $tags $species

#Go back to the Flower directory

cd ..

done

#After looping through the png files in the Flower directory go out of this directory

cd ..

#Clear the screen, print a message and wait for input from the user

clear

echo "Flower pictures divided"

echo "Press enter to continue"

read

#===================================================================================================#

# Devide the pictures in Tuber between shape(L-a-L) #

#===================================================================================================#

#Go into the Tuber Directory

cd Tuber

for files in \*.png

do

#Create the variables to divide the pictures.

name=(${files//./ }$0)

tags="$name""\_tags.txt"

content=$(cat $tags)

#Divide the pictures to the correct directory

if [[ $content == \*Look-a-Like\_round\* ]]

then

mv $name.png LRound/

mv $tags LRound/

elif [[ $content == \*Look-a-Like\_oblong\* ]]

then

mv $name.png LOblong/

mv $tags LOblong/

elif [[ $content == \*Look-a-Like\_spur\* ]]

then

mv $name.png LSpur/

mv $tags LSpur/

elif [[ $content == \*Round\* ]]

then

mv $name.png Round/

mv $tags Round/

elif [[ $content == \*Oblong\* ]]

then

mv $name.png Oblong/

mv $tags Oblong/

elif [[ $content == \*Spur\* ]]

then

mv $name.png Spur/

mv $tags Spur/

else

echo "No correct tag found"

fi

done

#After looping through the png files in the Tuber directory go out of this directory

cd ..

#Clear the screen, print a message and ask the user if he wants to split the Tuber pictures.

clear

echo "Tuber pictures divided"

echo "Press enter to continue"

read

#===================================================================================================#

# Split all pictures of Tuber #

#===================================================================================================#

#Go into the Tuber directory

cd Tuber

#Loop through all directories in this folder

for directories in $(ls -d \*/)

do

directory=${directories%%/}

#Set standard parameter values

size=10000

t=0.65

#Some directories requires other values for size or for t

if [[ $directory == \*LO\* ]]

then

size=50000

elif [[ $directory == \*LS\* ]]

then

size=50000

elif [[ $directory == R\* ]]

then

t=0.6

fi

#Loop through all pictures

for files in ./$directory/\*.png

do

#Split the picture, using the given parameter values

echo "Splitting $files"

perl ../splitter.pl -t $t -i $files

#Remove the noise pictures using the file size

for FILENAME in \*,\*.png

do

FILESIZE=$(stat -f%z $FILENAME)

if (( FILESIZE > size ))

then

mv $FILENAME ./$directory

else

rm $FILENAME

fi

done

done

done

#When this step is finished go out of the Tuber directory

cd ..

#Clear the screen and print a message

clear

echo "Tuber pictures splited"

#Print a finish message

echo "The program is finished"

echo "To create traindata"

echo "Please run ./create\_traindata.sh"

### CSS

#### computer.css

/\* This is the default lay-out for computers \*/

/\* Lay-out for the body \*/

body {

/\* Place the text in the middel of the page \*/

text-align: center;

}

/\* Lay-out for the page \*/

#page {

/\* the width \*/

width: 960px;

/\* place the text left \*/

text-align: left;

/\* clear automatically the area around the page \*/

margin: 10px auto 20px auto;

/\* Set the background color to white \*/

background-color:white;

}

/\* Lay-out for the logo \*/

#logo {

/\* Place the logo on the leftside of the page \*/

float: left;

/\* the width of the logo is 200 pixels \*/

width: 200px;

}

/\* Lay-out for sidebar, not used, but available for later \*/

/\*#sidebar {

float: right;

width: 200px;

border: 1px solid #000;

}\*/

/\* Lay-out for the content \*/

#content {

/\* place the content left, because logo is described first, the content will

be displayed right of the logo \*/

float: left;

/\* The width of this box will be automaticly changed to the content self \*/

width: auto;

/\* Place a border around the content

3 pixels widht with a pink-like color, The same color of the

Naturalis logo \*/

border: 3px solid #E3004A;

/\* Clear 10 pixels around the content, inside de border \*/

padding: 10px;

}

/\* Lay-out for the footer \*/

#footer {

/\* Place the footer on the bottom \*/

position:absolute;

bottom:10px;

}

/\* For all Font-settings below the Naturalis housestyle is aplied.

Which means that the Arial font is everywhere used \*/

/\* Font-settings of p \*/

p {

font-family: Arial;

font-size: 1em;

}

/\* Font-settings of h1 \*/

h1 {

font-family: Arial;

font-size: 1.9em;

}

/\* Font-settings of h2 \*/

h2 {

font-family: Arial;

font-size: 1.7em;

}

/\* Font-settings of h3 \*/

h3 {

font-family: Arial;

font-size: 1.5em;

}

/\* Font-settings of h4 \*/

h4 {

font-family: Arial;

font-size: 1.3em;

}

/\* id small, used to make text in p smaller than the settings for p \*/

#small {

font-size: 0.75em;

}

/\* Font-settings of form \*/

form {

font-family: Arial;

font-size: 1em;

}

/\* Font-settings of inut \*/

input {

font-family: Arial;

font-size: 0.75em;

}

/\* id button, for styling the individual buttons \*/

#button {

font-family: Arial;

font-size: 0.75em;

}

### mobile.css

/\* This is the default lay-out for mobile devices \*/

/\* Lay-out for the body \*/

body {

/\* Place a margin of 10 pixels around the text \*/

margin: 10px

}

/\* Lay-out for the logo \*/

#logo {

/\* Place the logo on the leftside of the page \*/

float: left;

}

/\* Lay-out for sidebar, not used, but available for later \*/

/\*#sidebar {

float: right;

width: 200px;

border: 1px solid #000;

}\*/

/\* Lay-out for the content \*/

#content {

/\* place the content left, because logo is described first, the content will

be displayed under the logo \*/

float: left;

/\* Place a border around the content

13 pixels widht with a pink-like color, The same color of the

Naturalis logo \*/

border: 13px solid #E3004A;

/\* Clear 10 pixels around the content, inside de border \*/

padding: 10px;

}

/\* Lay-out for the footer \*/

#footer {

/\* Place the footer on the bottom \*/

float:left;

bottom:10px;

}

/\* For all Font-settings below the Naturalis housestyle is aplied.

Which means that the Arial font is everywhere used \*/

/\* Font-settings of p \*/

p {

font-family: Arial;

font-size: 40px;

}

/\* Font-settings of h1 \*/

h1 {

font-family: Arial;

font-size: 120px;

}

/\* Font-settings of h2 \*/

h2 {

font-family: Arial;

font-size: 100px;

}

/\* Font-settings of h3 \*/

h3 {

font-family: Arial;

font-size: 75px;

}

/\* Font-settings of h4 \*/

h4 {

font-family: Arial;

font-size: 50px;

}

/\* id small, used to make text in p smaller than the settings for p \*/

#small {

font-size: 25px;

}

/\* Font-settings of form \*/

form {

font-family: Arial;

font-size: 30px;

}

/\* Font-settings of inut \*/

input {

font-family: Arial;

font-size: 40px;

}

/\* id button, for styling the individual buttons \*/

#button {

font-family: Arial;

font-size: 40px;

}

### HTML

#### computer\_invalid\_login.html

<!-- Use the computer template and override the content block -->

{% extends "computer.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Invalid login{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Display a header, telling the login details are incorrect -->

<h2>Your login details are invalid!</h2>

<!-- Button to login again -->

<form action="/accounts/login/">

<!-- The submit button with the text Try again inside -->

<input type="submit" value="Try again" id="button">

</form>

<p></p>

<!-- Button to go back to the welcome page -->

<form action="/welcome/">

<!-- The submit button with the text Home inside -->

<input type="submit" value="Home" id="button">

</form>

{% endblock %}

#### computer\_login.html

<!-- Use the computer template and override the content block -->

{% extends "computer.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Login{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Display a header -->

<h3>Login required to remove files!</h3>

<!-- Ask for the username and password -->

<form action="/accounts/auth/" method="post">{% csrf\_token %}

<label for="username">User name:</label>

<input type="text" name="username" value="" id="username">

<p></p>

<label for="password">Password:</label>

<input type="password" name="password" value="" id="pasword">

<p></p>

<input type="submit" value="login" id="button">

</form>

<p></p>

<!-- Button to go back to the home page -->

<form action="/welcome/">

<!-- The submit button with the text Home inside -->

<input type="submit" value="Home" id="button">

</form>

{% endblock %}

#### computer\_remove.html

<!-- Use the computer template and override the content block -->

{% extends "computer.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Files removed{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Display a header, telling the task has been completed -->

<h2>Removing complete!</h2>

<!-- Give detailed information -->

<p>All uploaded pictures and their results<BR>

are moved to the results directory</p>

<p>You have removed the following files:</p>

<p>{{uploads}}</p>

<p>{{temps}}</p>

<!-- Button to logout -->

<form action="/accounts/logout/">

<!-- The submit button with the text Upload picture inside -->

<input type="submit" value="Logout" id="button">

</form>

{% endblock %}

#### computer\_result.html

<!-- Use the computer template and override the content block -->

{% extends "computer.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Result{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Display a header to tell this is de result page -->

<h2>Results</h2>

<!-- Display the result -->

<p>Result: This is probably a <em>{{result}}</em> </p>

<!-- Display the picture, give it a link to view the picter on a new

webpage / tab -->

<p><a href="/static/assets/uploaded\_files/{{ip}}/{{filename}}"><img src="/static/assets/uploaded\_files/{{ip}}/{{filename}}" width="200"/></a></p>

<form action="/exit/">

<!-- The submit button with the text Upload picture inside -->

<input type="submit" value="Exit" id="button">

</form>

{% endblock %}

#### computer\_sorry.html

<!-- Use the computer template and override the content block -->

{% extends "computer.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Error{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Header -->

<h2>Sorry!</h2>

<!-- Explain the error and ask to try again -->

<p>During the process, your picture has been removed.<BR>

Please try again</p>

<!-- Button to try again -->

<form action="/upload/">

<!-- The submit button with the text Upload picture inside -->

<input type="submit" value="Upload picture" id="button">

</form>

<p></p>

<!-- Button to go back to the homepage -->

<form action="/welcome/">

<!-- The submit button with the text Home inside -->

<input type="submit" value="Home" id="button">

</form>

{% endblock %}

#### computer\_upload\_succes.html

<!-- Use the computer template and override the content block -->

{% extends "computer.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Upload succeeded{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Display a header, telling the file is uploaded -->

<h2>You have uploaded the file!</h2>

<!-- Display the picture, give it a link to view the picter on a new

webpage / tab -->

<p><a href="/{{path}}"><img src="/{{path}}" width="200"/></a></p>

<form action="/result/">

<!-- The submit button with the text Upload picture inside -->

<input type="submit" value="Result" id="button">

</form>

<!-- Display an empty line betwee both buttonts-->

<p></p>

<form action="/upload/">

<!-- The submit button with the text Upload another picture inside -->

<input type="submit" value="Upload another picture" id="button">

</form>

{% endblock %}

#### computer\_upload.html

<!-- Use the computer template and override the content block -->

{% extends "computer.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Upload picture{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Display a header to tell this is de upload page -->

<h2>Upload picture</h2>

<p style="color:red; font-size:0.9em"><em>{{message}}</em></p>

<!-- Use the form from orchid.views.upload to receive a picture -->

<form action="" method="post" style="{{style}}" enctype="multipart/form-data">{% csrf\_token %}

{{form}}

<p></p>

<!-- Place a submit button under the form, with the text Upload picture inside.

To place it under the form, it is required to display an empty line first-->

<input type="submit" value="Upload picture" id="button">

<!-- End of form -->

</form>

<p></p>

<!-- Place a button under the form to go back to the home screen -->

<form action="/welcome/">

<input type="submit" value="Home" id="button">

</form>

{% endblock %}

#### computer\_welcome.html

<!-- Use the computer template and override the content block -->

{% extends "computer.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Home{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Display a header, telling this is the welcome page -->

<h2>Welcome</h2

<!-- Tell the user what this webapplication will do -->

<h4>Welcome on the orchid identifier website of Naturalis</h4>

<p>After uploading a picture the website will identify the orchid.<BR>

Please click the upload picture button below to upload a picture.<BR>

<em>You can upload a picture of a tuber.</em></p>

<!-- Create a form which contains two buttons.

The first button is for uploading the file -->

<form action="/upload/">

<!-- The submit button with the text Upload picture inside -->

<input type="submit" value="Upload picture" id="button">

</form>

<!-- display a white line -->

<p></p>

<!-- Second button to remove all leftover files -->

<p id="small">To remove all unused uploaded pictures and their temporary files<BR>

click below <span style="color: red"><strong>(login required)</strong></span></p>

<form action="/admin/remove">

<input type="submit" value="Remove unused files" id="button">

</form>

{% endblock %}

#### computer.html

<!-- THIS IS THE STANDARD HTML FOR THE COMPUTER LAY-OUTS -->

<!-- Load in the static function -->

{% load static %}

<!-- declare the DOCTYPE and html language-->

<!DOCTYPE html>

<html lang="en">

<!-- Create the head -->

<head>

<!-- Create a title using a block -->

<!-- This title needs to be changed! The best way to change the title

is in the other html files witch extends this html -->

<title>{% block title %} My Base Template{% endblock %}</title>

<!-- Use the compyter.css stylesheet for all the lay-outs -->

<link rel="stylesheet" type="text/css" href="{% static "assets/css/computer.css" %}">

<!-- end of the head -->

</head>

<!-- Create the body -->

<body>

<!-- Create a page -->

<div id="page">

<!-- Create a logo block, using the logo lay-out from compyter.css -->

<div id="logo">

{% block logo %}

<!-- Place the Naturalis logo inside the logo block -->

<ul>

<a href="<http://www.naturalis.nl>"><img src="{% static "assets/images/Naturalis\_logo.png" %}" width="100"/></a>

</ul>

<!-- End of the logo block -->

{% endblock %}

</div>

<!-- Create a content block, using the content lay-out from computer.css -->

<div id="content">

<!-- Place a standard text in the content -->

<!-- This block will be override for all different html pages -->

{% block content %}This is the content of this block{% endblock %}

</div>

<!-- Create a footer block, using the footer lay-out from computer.css -->

<div id="footer">

<!-- Place some text in the footer block -->

{% block footer %}<p>&copy;2013-2014 Patrick Wijntjes</p>{% endblock %}

<!-- End of the page -->

</div>

<!-- End of the body and of the html -->

</body>

</html>

#### mobile\_invalid\_login.html

<!-- Use the mobile template and override the content block -->

{% extends "mobile.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Invalid login{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Display a header, telling the login details are incorrect -->

<h2>Your login details are invalid!</h2>

<!-- Button to login again -->

<form action="/accounts/login/">

<!-- The submit button with the text Try again inside -->

<input type="submit" value="Try again" id="button">

</form>

<p></p>

<!-- Button to go back to the welcome page -->

<form action="/welcome/">

<!-- The submit button with the text Home inside -->

<input type="submit" value="Home" id="button">

</form>

{% endblock %}

#### mobile\_login.html

<!-- Use the mobile template and override the content block -->

{% extends "mobile.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Login{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Display a header -->

<h3>Login required to remove files!</h3>

<!-- Ask for the username and password -->

<form action="/accounts/auth/" method="post">{% csrf\_token %}

<label for="username">User name:</label>

<input type="text" name="username" value="" id="username">

<p></p>

<label for="password">Password:</label>

<input type="password" name="password" value="" id="pasword">

<p></p>

<input type="submit" value="login" id="button">

</form>

<p></p>

<!-- Button to go back to the home page -->

<form action="/welcome/">

<!-- The submit button with the text Home inside -->

<input type="submit" value="Home" id="button">

</form>

{% endblock %}

#### mobile\_remove.html

<!-- Use the mobile template and override the content block -->

{% extends "mobile.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Files removed{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Display a header, telling the task has been completed -->

<h2>Removing complete!</h2>

<!-- Give detailed information -->

<p>All uploaded pictures and their results<BR>

are moved to the results directory</p>

<p>You have removed the following files:</p>

<p>{{uploads}}</p>

<p>{{temps}}</p>

<!-- Button to logout -->

<form action="/accounts/logout/">

<!-- The submit button with the text Upload picture inside -->

<input type="submit" value="Logout" id="button">

</form>

{% endblock %}

#### mobile\_result.html

<!-- Use the mobile template and override the content block -->

{% extends "mobile.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Result{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Display a header to tell this is de result page -->

<h2>Results</h2>

<!-- Display the result -->

<p>Result: This is probably a <em>{{result}}</em> </p>

<!-- Display the picture, give it a link to view the picter on a new

webpage / tab -->

<p><a href="/static/assets/uploaded\_files/{{ip}}/{{filename}}"><img src="/static/assets/uploaded\_files/{{ip}}/{{filename}}" width="200"/></a></p>

<form action="/exit/">

<!-- The submit button with the text Exit inside -->

<input type="submit" value="Exit" id="button">

</form>

{% endblock %}

#### mobile\_sorry.html

<!-- Use the mobile template and override the content block -->

{% extends "mobile.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Error{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Header -->

<h2>Sorry!</h2>

<!-- Explain the error and ask to try again -->

<p>During the process, your picture has been removed.<BR>

Please try again</p>

<!-- Button to try again -->

<form action="/upload/">

<!-- The submit button with the text Upload picture inside -->

<input type="submit" value="Upload picture" id="button">

</form>

<p></p>

<!-- Buttont to go back to the homepage -->

<form action="/welcome/">

<!-- The submit buttont with the thext Home inside -->

<input type="submit" value="Home" id="button">

</form>

{% endblock %}

#### mobile\_upload\_succes.html

<!-- Use the mobile template and override the content block -->

{% extends "mobile.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Upload succeeded{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Display a header, telling the file is uploaded -->

<h2>You have uploaded the file!</h2>

<!-- Display the picture, give it a link to view the picter on a new

webpage / tab -->

<p><a href="/{{path}}"><img src="/{{path}}" width="200"/></a></p>

<form action="/result/">

<!-- The submit button with the text Upload picture inside -->

<input type="submit" value="Result" id="button">

</form>

<!-- Display an empty line betwee both buttonts-->

<p style="font-size:10;"></p>

<form action="/upload/">

<!-- The submit button with the text Upload another picture inside -->

<input type="submit" value="Upload another picture" id="button">

</form>

{% endblock %}

### mobile\_upload.html

<!-- Use the mobile template and override the content block -->

{% extends "mobile.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Upload picture{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Display a header to tell this is de upload page -->

<h2>Upload picture</h2>

<!-- Font-size needs te be changed!!!!! -->

<p style="{{style}};"><em>{{message}}</em></p>

<!-- Use the form from orchid.views.upload to receive a picture -->

<form action="" method="post" style="{{style}}; font-size:60px" enctype="multipart/form-data">{% csrf\_token %}

{{form}}

<p></p>

<!-- Place a submit button under the form, with the text Upload picture inside.

To place it under the form, it is required to display an empty line first-->

<input type="submit" value="Upload picture" id="button">

<!-- End of form -->

</form>

<p></p>

<!-- Place a button under the form to go back to the home screen -->

<form action="/welcome/">

<input type="submit" value="Home" id="button">

</form>

{% endblock %}

### mobile\_welcome.html

<!-- Use the mobile template and override the content block -->

{% extends "mobile.html" %}

<!-- Override the title -->

{% block title %}Orchid Identifier - Home{% endblock %}

{% block content %}

<!-- All lines in this block will be diplayd in the content block -->

<!-- Display a header, telling this is the welcome page -->

<h2>Welcome</h2

<!-- Tell the user what this webapplication will do -->

<h4>Welcome on the orchid identifier website of Naturalis</h4>

<p>After uploading a picture the website will identify the orchid.<BR>

Please click the upload picture button below to upload a picture.<BR>

<em>You can upload a picture of a tuber.</em></p>

<!-- Create a form which contains two buttons.

The first button is for uploading the file -->

<form action="/upload/">

<!-- The submit button with the text Upload picture inside -->

<input type="submit" value="Upload picture" id="button">

</form>

<!-- display a white line -->

<p></p>

<!-- Second button to remove all leftover files -->

<p id="small">To remove all unused uploaded pictures and their temporary files<BR>

click below <span style="color: red"><strong>(login required)</strong></span></p>

<form action="/admin/remove">

<input type="submit" value="Remove unused files" id="button">

</form>

{% endblock %}

#### mobile.html

<!-- THIS IS THE STANDARD HTML FOR MOBILE DEVICES -->

<!-- Load in the static function -->

{% load static %}

<!-- declare the DOCTYPE and html language-->

<!DOCTYPE html>

<html lang="en">

<!-- Create the head -->

<head>

<!-- Create a title using a block -->

<!-- This title needs to be changed! The best way to change the title

is in the other html files witch extends this html -->

<title>{% block title %} My Base Template{% endblock %}</title>

<!-- Use the mobile.css stylesheet for all mobile lay-outs -->

<link rel="stylesheet" type="text/css" href="{% static "assets/css/mobile.css" %}">

<!-- end of the head -->

</head>

<!-- Create the body -->

<body>

<!-- Create a page -->

<div id="page">

<!-- Create a logo block, using the logo lay-out from mobile.css -->

<div id="logo">

{% block logo %}

<!-- Place the Naturalis logo inside the logo block -->

<ul>

<a href="<http://www.naturalis.nl>"><img src="{% static "assets/images/Naturalis\_logo.png" %}" width="250"/></a>

</ul>

<!-- End of the logo block -->

{% endblock %}

</div>

<!-- Create a content block, using the content lay-out from mobile.css -->

<div id="content">

<!-- Place a standard text in the content -->

<!-- This block will be override for all mobile html pages -->

{% block content %}This is the content of this block{% endblock %}

</div>

<!-- Create a footer block, using the footer lay-out from mobile.css -->

<div id="footer">

<!-- Place some text in the footer block -->

{% block footer %}<p>&copy;2013-2014 Patrick Wijntjes</p>{% endblock %}

<!-- End of the page -->

</div>

<!-- End of the body and of the html -->

</body>

</html>

### Perl

#### classify.pl [22]

#!/usr/bin/perl

use strict;

use warnings;

use Getopt::Long;

use Data::Dumper;

use AI::FANN ':all';

use Fingerprint 'make\_fingerprint';

use Bio::Phylo::Util::Logger ':levels';

# process command line arguments

my $verbosity = WARN;

my $resolution = 50;

my $dir;

my $ai;

GetOptions(

'verbose+' => \$verbosity,

'dir=s' => \$dir,

'ai=s' => \$ai,

'resolution=i' => \$resolution,

);

# instantiate helper objects

my $log = Bio::Phylo::Util::Logger->new(

'-level' => $verbosity,

'-class' => 'main',

);

$log->info("going to instantiate AI from file $ai");

my $ann = AI::FANN->new\_from\_file($ai);

# read from the directory

$log->info("going to classify images in dir $dir");

opendir my $dh, $dir or die $!;

while( my $entry = readdir $dh ) {

if ( $entry =~ /.png$/ ) {

$log->debug("going to classify $entry");

# analyse the input file

my @fingerprint = make\_fingerprint(

'file' => "$dir/$entry",

'resolution' => $resolution,

);

$log->debug("made fingerprint of file");

# do the classification

my $result = $ann->run(\@fingerprint);

print "Entry: $entry\n";

print Dumper($result);

}

}

#### splitter.pl [22]

#!/usr/bin/perl

use strict;

use warnings;

use Data::Dumper;

use Getopt::Long;

use Image::Magick;

use List::Util 'sum';

use Bio::Phylo::Util::Logger ':levels';

# will have deep recursions

no warnings 'recursion';

# process command line arguments

my $threshold = 0.7;

my $fuzzyness = 100; # pixels

my $verbosity = WARN;

my $infile;

GetOptions(

'threshold=f' => \$threshold,

'fuzzyness=i' => \$fuzzyness,

'verbose+' => \$verbosity,

'infile=s' => \$infile,

);

# instantiate helper objects

my $log = Bio::Phylo::Util::Logger->new(

'-level' => $verbosity,

'-class' => 'main',

);

my $img = Image::Magick->new;

my %seen;

my %area;

# read the image

$log->info("going to read image '$infile'");

my $msg = $img->Read($infile);

$log->warn($msg) if $msg;

# get width and height

my $width = $img->Get('columns');

my $height = $img->Get('rows');

$log->info("width: $width height: $height");

# iterate over all pixels

for my $x ( 0 .. $width ) {

for my $y ( 0 .. $height ) {

my $nucleus = "$x,$y";

recurse( 'x' => $x, 'y' => $y, 'nucleus' => $nucleus );

if ( $area{$nucleus} ) {

my $size = scalar @{ $area{$nucleus} };

if ( $size > $fuzzyness ) {

$log->info("found area of $size pixels around nucleus $nucleus");

}

}

}

}

# write large areas

for my $nucleus ( grep { scalar @{ $area{$\_} } > $fuzzyness } keys %area ) {

my ($min\_x) = sort { $a <=> $b } map { [ split(/,/, $\_) ]->[0] } @{ $area{$nucleus} };

my ($max\_x) = sort { $b <=> $a } map { [ split(/,/, $\_) ]->[0] } @{ $area{$nucleus} };

my ($min\_y) = sort { $a <=> $b } map { [ split(/,/, $\_) ]->[1] } @{ $area{$nucleus} };

my ($max\_y) = sort { $b <=> $a } map { [ split(/,/, $\_) ]->[1] } @{ $area{$nucleus} };

# compute new area

my $new\_width = $max\_x - $min\_x;

my $new\_height = $max\_y - $min\_y;

$log->info("going to write $nucleus to ${new\_width}x${new\_height} file");

# create new image, set dimensions, make white background

my $new\_img = Image::Magick->new( 'size' => "${new\_width}x${new\_height}" );

$msg = $new\_img->Read('xc:white');

$log->warn($msg) if $msg;

$log->info("instantiated new image");

# assign pixels

for my $x ( 0 .. $new\_width ) {

for my $y ( 0 .. $new\_height ) {

my $loc = ( $x + $min\_x ) . ',' . ( $y + $min\_y );

if ( $seen{$loc} ) {

$msg = $new\_img->SetPixel( 'x' => $x, 'y' => $y, 'color' => $seen{$loc} );

$log->warn($msg) if $msg;

}

}

}

$log->info("assigned new pixels");

# write image

$msg = $new\_img->Write("${nucleus}.png");

$log->warn($msg) if $msg;

$log->info("wrote image ${nucleus}.png");

}

sub recurse {

my %args = @\_;

# get sub args

my $nucleus = delete $args{nucleus};

my ( $x, $y ) = @args{qw(x y)};

# sample the focal pixel

my @pixel = $img->GetPixel(%args);

# if pixel is darker than threshold and not yet seen...

if ( sum(@pixel)/scalar(@pixel) < $threshold && ! $seen{"$x,$y"} ) {

$log->debug("$x,$y");

# store the pixel

$seen{"$x,$y"} = \@pixel;

# initialize area around current nucleus

$area{$nucleus} = [] if not $area{$nucleus};

# store id of the focal pixel

push @{ $area{$nucleus} }, "$x,$y";

# start growing the area

if ( $x > 0 ) {

recurse( 'x' => $x - 1, 'y' => $y, 'nucleus' => $nucleus );

}

if ( $y > 0 ) {

recurse( 'x' => $x, 'y' => $y - 1, 'nucleus' => $nucleus );

}

if ( $x < $width ) {

recurse( 'x' => $x + 1, 'y' => $y, 'nucleus' => $nucleus );

}

if ( $y < $height ) {

recurse( 'x' => $x, 'y' => $y + 1, 'nucleus' => $nucleus );

}

}

}

#### trainai.pl [22]

#!/usr/bin/perl

use strict;

use warnings;

use Getopt::Long;

use AI::FANN ':all';

use Bio::Phylo::Util::Logger ':levels';

# process command line arguments

my $verbosity = WARN;

my $epochs = 50000;

my $target = 0.0001;

my $datadir = 'data/traindata';

my $outfile = 'data/ai/butterbeetle.ann';

my $categories = 1;

GetOptions(

'verbose+' => \$verbosity,

'datadir=s' => \$datadir,

'epochs=i' => \$epochs,

'target=f' => \$target,

'outfile=s' => \$outfile,

'categories=i' => \$categories,

);

# instantiate helper objects

my $log = Bio::Phylo::Util::Logger->new(

'-level' => $verbosity,

'-class' => 'main',

);

# read the data files

my @interdigitated;

my $neurons;

$log->info("going to read traindata from $datadir");

opendir my $dh, $datadir or die $!;

while( my $entry = readdir $dh ) {

if ( $entry =~ /\.tsv$/ ) {

# read the table

$log->info("going to read $datadir/$entry");

open my $fh, '<', "$datadir/$entry" or die $!;

my @header;

LINE: while(<$fh>) {

chomp;

my @fields = split /\t/, $\_;

if ( not @header ) {

@header = @fields;

next LINE;

}

# first cell

my $file = shift @fields;

# last cells

my @categ;

for my $i ( 1 .. $categories ) {

push @categ, pop @fields;

}

# see AI::FANN docs for datastructure

push @interdigitated, \@fields, \@categ;

$neurons = scalar @fields; # +1 in hidden layer

$log->info("read fingerprint for $file (@categ)");

}

}

}

# create the training data struct

my $train = AI::FANN::TrainData->new(@interdigitated);

# create the AI

my $ann = AI::FANN->new\_standard( $neurons, $neurons + 1, $categories );

$ann->hidden\_activation\_function(FANN\_SIGMOID\_SYMMETRIC);

$ann->output\_activation\_function(FANN\_SIGMOID\_SYMMETRIC);

# train the AI

$ann->train\_on\_data( $train, $epochs, $epochs / 100, $target );

# save the result

$ann->save($outfile);

#### traindata.pl [22]

#!/usr/bin/perl

use strict;

use warnings;

use Getopt::Long;

use Fingerprint 'make\_fingerprint';

use Bio::Phylo::Util::Logger ':levels';

# process command line arguments

my $verbosity = WARN;

my $resolution = 50;

my $dir;

my $category;

GetOptions(

'category=i' => \$category,

'resolution=i' => \$resolution,

'dir=s' => \$dir,

'verbose+' => \$verbosity,

);

# instantiate helper objects

my $log = Bio::Phylo::Util::Logger->new(

'-level' => $verbosity,

'-class' => [ 'main', 'Fingerprint' ],

);

# print the header

print "image\t";

for my $axis ( qw(vert horiz) ) {

for my $color ( qw(red green blue) ) {

my $max = $axis eq 'horiz' ? $resolution / 2 : $resolution;

for my $i ( 1 .. $max ) {

print "${axis}.${color}.${i}\t";

}

}

}

print "category\n";

# start reading the images

$log->info("going to read images from $dir");

opendir my $dh, $dir or die $!;

while( my $entry = readdir $dh ) {

# only read png files created by splitter.pl

if ( $entry =~ /(\d+,\d+)\.png/ ) {

my $nucleus = $1;

my @row = ( $nucleus );

# read image

my $img = Image::Magick->new;

push @row, make\_fingerprint(

'file' => $dir . '/' . $entry,

'resolution' => $resolution,

);

$log->info("created fingerprint for $entry");

push @row, $category;

print join("\t", @row), "\n";

}

}

#### traindata2.pl [22]

#!/usr/bin/perl

use strict;

use warnings;

use Getopt::Long;

use Fingerprint 'make\_fingerprint';

use Bio::Phylo::Util::Logger ':levels';

# process command line arguments

my $verbosity = WARN;

my $resolution = 50;

my $dir;

my $category;

GetOptions(

'category=i' => \$category,

'resolution=i' => \$resolution,

'dir=s' => \$dir,

'verbose+' => \$verbosity,

);

# instantiate helper objects

my $log = Bio::Phylo::Util::Logger->new(

'-level' => $verbosity,

'-class' => [ 'main', 'Fingerprint' ],

);

# print the header

print "image\t";

for my $axis ( qw(vert horiz) ) {

for my $color ( qw(red green blue) ) {

my $max = $axis eq 'horiz' ? $resolution / 2 : $resolution;

for my $i ( 1 .. $max ) {

print "${axis}.${color}.${i}\t";

}

}

}

print "category\n";

# start reading the images

$log->info("going to read images from $dir");

opendir my $dh, $dir or die $!;

while( my $entry = readdir $dh ) {

# only read png files created by splitter.pl

if ( $entry =~ /(\d+,?\d\*)\.png/ ) {

my $nucleus = $1;

my @row = ( $nucleus );

# read image

my $img = Image::Magick->new;

push @row, make\_fingerprint(

'file' => $dir . '/' . $entry,

'resolution' => $resolution,

);

$log->info("created fingerprint for $entry");

push @row, $category;

print join("\t", @row), "\n";

}

}

### Python for training

#### Add\_columns.py

import os

#Collect all tsv files and save the names in files.txt

os.system("ls | egrep '.tsv' > files.txt")

#Function for creating the variables

def create\_variables():

#Read the names of the tsv files and save them as a list in python

files = open("files.txt", 'r').readlines()

#Create variables

number = len(files)

header = []

out\_list = []

counter = 0

maxi = 0

#Run the create\_output\_list function

create\_output\_list(number, header, out\_list ,files, counter, maxi)

#Function for creating the output lists

def create\_output\_list(number, header, out\_list, files, counter, maxi):

#add information to the output lists

for q in range(number):

out\_list.append(-1)

header.append("C%s"%(q+1))

#Run the change\_files function

change\_files(header, out\_list, number ,files, counter, maxi)

#Functin for changing the files. This function will add the columns to the files.

def change\_files(header, out\_list, number, files, counter, maxi):

#Loop through the list of files

for x in range(number):

#Create an output file

y = files[x].strip()

name = y.split(".")[0] + "\_new.tsv"

output = open(name, 'a')

#Read the content of the file, save it as a list

z = open(y, 'r').readlines()

#Loop through the list of content

for a in range(len(z)):

#Create variables, b is the content list, i is the output list

b = z[a].split("\t")

i = b[:-2]

#When a is 0, it is the header

if a == 0:

#Add the header list to the output list

for c in range(number):

i.append(header[c])

#Write the content of the output list to the output file

for d in i:

output.write(d.strip() + "\t")

#After looping through the output list write an enter to the output file.

output.write("\n")

#When a is not 0 it is a normal line

else:

#When the counter (=file number) is equal to x, column x gets an 1

if counter == x:

out\_list[x] = 1

#Otherwise it keeps a 0

else:

pass

#Add the list of -1's (and 1's for column x == counter) to the output list

for f in range(number):

i.append(out\_list[f])

#Write the contentn of the output list to the output file

for g in i:

output.write("%s\t" %(g))

#After looping through the output list write an entr to the output file.

output.write("\n")

#Search to the hights number of columns

if maxi < len(i):

maxi = len(i)

#Set the column with 1 back to 0 for the next file

out\_list[x] = -1

#Add 1 to the counter.

counter += 1

os.system("mv %s %s"%(y, y.split(".")[0]))

print "\nnumber of categories: ", counter

clean\_up(output)

def clean\_up(output):

#Close the output file

output.close()

#Remove the temporary file

os.system("rm files.txt")

#Run the create\_variables function

create\_variables()

#### Combine\_files.py

import os

#function for creating the variables

def create\_variables():

#Save all tsv files created with add\_columns.py in files.txt

os.system("ls | egrep '.tsv' > files.txt")

#Read the content of files.txt and save it as a list

files = open('files.txt', 'r').readlines()

#Create variables

number = len(files)

counter = 0

#Get the path working directory

paht = os.path.dirname(os.path.realpath("combine\_files.py"))

#The output file will named as the directory

name = paht.split("/")[-1] + ".tsv"

#print name

output = open(name, 'a')

#Run the process\_files function

process\_files(number, files, counter, output)

#Run the clean\_up function

clean\_up(output, name)

#Function for combining the files

def process\_files(number, files, counter, output):

#Loop through the files list

for file\_index in range(number):

file = files[file\_index].strip()

#Read the content of the file, save it as a list.

content = open(file, 'r').readlines()

#Loop through the lines

for row\_number in range(len(content)):

#create a list, every columns is an entry of the list

row = content[row\_number].split("\t")

#When counter is 0, this is the first file.

if counter == 0:

#The whole content of the file is written to the output file

for entry in row:

output.write(entry.strip() + "\t")

#After looping through the content, write an enter

output.write("\n")

#When counter isn't 0, it is not the first file

else:

#When row\_number is 0 it is the header

if row\_number == 0:

#The header will not be written to the output file, because it is already there

pass

#When row\_number isn't 0 it is not the header

else:

#The content will be written to the output file

for entry in row:

output.write(entry.strip() + "\t")

#After looping through the content, write an enter to the output file.

output.write("\n")

#Add 1 to the counter

counter += 1

#Function for removing temporary files.

def clean\_up(output, name):

#Close the output file

output.close()

#Remove the temporary files.txt file

os.system("rm files.txt")

#Move the output file out of the Flower directory

os.system("mv %s ../"%(name))

#Run the create variables function

create\_variables()

#### get\_tags.py

import os

#Function for getting all xml files

def get\_xml\_files():

#Open xml\_files.txt in read mode

files = open("xml\_files.txt", 'r')

#Create a list of all the xml files

infiles = files.readlines()

#Close the temporary file

files.close()

#Run the create\_tag\_files function

create\_tag\_files(infiles)

#Function for creating the tag files

def create\_tag\_files(infiles):

#Loop through the list

for file in infiles:

#Remove all enters at the back of the filename

infile = file.strip()

#Get the picture id, to save the tags with the same number as the picture

#Example the name of the meta file is 123456789.xml so the id is 123456789

number = infile.split(".")[0]

#Print a message

print "Collecting the tags of file %s"%(infile)

#Open the meta data file in read mode

open\_file = open(infile, 'r')

#Make a list of the meta data

read\_file = open\_file.readlines()

#Try to find the tags

try:

'''One line befor the first tag you can find "/t<tags>\n"

So the first tag will be the index of "/t<tags>\n" +1'''

#Get the index of the first tag

start = read\_file.index("\t<tags>\n") +1

'''One line after the last tag you can find "\t</tags>\n"

So the last tag will be the index of "/t</tags>\n" -1. Since a for-loop

loops from start to end, EXCLUDING the end, you use the index of "/t</tags>\n"'''

#Get the index of the end of the tags

end = read\_file.index("\t</tags>\n")

#Get the original name of the picture

title = read\_file[2].split(">")[1].split("<")[0]

#Save the output name (using the id of the picture)

out\_name = "%s\_tags.txt"%(number)

#Open the output file in write mode

output = open(out\_name, 'w')

print "The tags will be saved in %s"%(out\_name)

#Write the name of the picture and a white line to the output file

#The title will always be the first line of the output file

output.write("%s\n\n"%(title))

#Loop through the tags

for tag\_line in range(start, end):

#Write the text between <tag> and </tag> to the output file

output.write(read\_file[tag\_line].strip().split(" ")[4].split('"')[1])

#Write an enter to the output file

output.write("\n")

#When the loop ends, close the output file

output.close()

'''If there are no tags, a ValueError arise. Except this Error and print

a message that the file has no tags'''

except ValueError:

print "%s has no tags"%(infile)

#Close the output file

open\_file.close()

#break

#Run the get\_xml\_files function

get\_xml\_files()

#### Offlickr.py [18]

#!/usr/bin/python

# -\*- coding: utf-8 -\*-

# Offlickr

# Hugo Haas -- <mailto:hugo@larve.net> -- <http://larve.net/people/hugo/>

# Homepage: <http://larve.net/people/hugo/2005/12/offlickr/>

# License: GPLv2

#

# Daniel Drucker <[dmd@3e.org](mailto:dmd@3e.org)> contributed:

# \* wget patch

# \* backup of videos as well

# \* updated to Beej's Flickr API version 1.2 (required)

import sys

import libxml2

import urllib

import getopt

import time

import os.path

import threading

# Beej's Python Flickr API

# <http://beej.us/flickr/flickrapi/>

from flickrapi import FlickrAPI

import logging

\_\_version\_\_ = '0.22 - 2009-03-20'

maxTime = '9999999999'

# Gotten from Flickr

flickrAPIKey = '1391fcd0a9780b247cd6a101272acf71'

flickrSecret = 'fd221d0336de3b6d'

class Offlickr:

def \_\_init\_\_(

self,

key,

secret,

uid,

httplib=None,

dryrun=False,

verbose=False,

):

"""Instantiates an Offlickr object

An API key is needed, as well as an API secret and a user id."""

self.\_\_flickrAPIKey = key

self.\_\_flickrSecret = secret

self.\_\_httplib = httplib

# Get authentication token

# note we must explicitly select the xmlnode parser to be compatible with FlickrAPI 1.2

self.fapi = FlickrAPI(self.\_\_flickrAPIKey, self.\_\_flickrSecret,

format='xmlnode')

(token, frob) = self.fapi.get\_token\_part\_one()

if not token:

raw\_input('Press ENTER after you authorized this program')

self.fapi.get\_token\_part\_two((token, frob))

self.token = token

self.flickrUserId = uid

self.dryrun = dryrun

self.verbose = verbose

def \_\_testFailure(self, rsp):

"""Returns whether the previous call was successful"""

if rsp['stat'] == 'fail':

print 'Error!'

return True

else:

return False

def getPhotoList(self, dateLo, dateHi):

"""Returns a list of photo given a time frame"""

n = 0

flickr\_max = 500

photos = []

print 'Retrieving list of photos'

while True:

if self.verbose:

print 'Requesting a page...'

n = n + 1

rsp = self.fapi.photos\_search(

api\_key=self.\_\_flickrAPIKey,

auth\_token=self.token,

user\_id=self.flickrUserId,

per\_page=str(flickr\_max),

page=str(n),

min\_upload\_date=dateLo,

max\_upload\_date=dateHi,

#The next line is added by Patrick Wijntjes, 14-01-2014

privacy\_filter=5

)

if self.\_\_testFailure(rsp):

return None

if rsp.photos[0]['total'] == '0':

return None

photos += rsp.photos[0].photo

if self.verbose:

print ' %d photos so far' % len(photos)

if len(photos) >= int(rsp.photos[0]['total']):

break

return photos

def getGeotaggedPhotoList(self, dateLo, dateHi):

"""Returns a list of photo given a time frame"""

n = 0

flickr\_max = 500

photos = []

print 'Retrieving list of photos'

while True:

if self.verbose:

print 'Requesting a page...'

n = n + 1

rsp = \

self.fapi.photos\_getWithGeoData(api\_key=self.\_\_flickrAPIKey,

auth\_token=self.token, user\_id=self.flickrUserId,

per\_page=str(flickr\_max), page=str(n))

if self.\_\_testFailure(rsp):

return None

if rsp.photos[0]['total'] == '0':

return None

photos += rsp.photos[0].photo

if self.verbose:

print ' %d photos so far' % len(photos)

if len(photos) >= int(rsp.photos[0]['total']):

break

return photos

def getPhotoLocation(self, pid):

"""Returns a string containing location of a photo (in XML)"""

rsp = \

self.fapi.photos\_geo\_getLocation(api\_key=self.\_\_flickrAPIKey,

auth\_token=self.token, photo\_id=pid)

if self.\_\_testFailure(rsp):

return None

doc = libxml2.parseDoc(rsp.xml)

info = doc.xpathEval('/rsp/photo')[0].serialize()

doc.freeDoc()

return info

def getPhotoLocationPermission(self, pid):

"""Returns a string containing location permision for a photo (in XML)"""

rsp = \

self.fapi.photos\_geo\_getPerms(api\_key=self.\_\_flickrAPIKey,

auth\_token=self.token, photo\_id=pid)

if self.\_\_testFailure(rsp):

return None

doc = libxml2.parseDoc(rsp.xml)

info = doc.xpathEval('/rsp/perms')[0].serialize()

doc.freeDoc()

return info

def getPhotosetList(self):

"""Returns a list of photosets for a user"""

rsp = self.fapi.photosets\_getList(api\_key=self.\_\_flickrAPIKey,

auth\_token=self.token, user\_id=self.flickrUserId)

if self.\_\_testFailure(rsp):

return None

return rsp.photosets[0].photoset

def getPhotosetInfo(self, pid, method):

"""Returns a string containing information about a photoset (in XML)"""

rsp = method(api\_key=self.\_\_flickrAPIKey,

auth\_token=self.token, photoset\_id=pid)

if self.\_\_testFailure(rsp):

return None

doc = libxml2.parseDoc(rsp.xml)

info = doc.xpathEval('/rsp/photoset')[0].serialize()

doc.freeDoc()

return info

def getPhotoMetadata(self, pid):

"""Returns an array containing containing the photo metadata (as a string), and the format of the photo"""

if self.verbose:

print 'Requesting metadata for photo %s' % pid

rsp = self.fapi.photos\_getInfo(api\_key=self.\_\_flickrAPIKey,

auth\_token=self.token, photo\_id=pid)

if self.\_\_testFailure(rsp):

return None

doc = libxml2.parseDoc(rsp.xml)

metadata = doc.xpathEval('/rsp/photo')[0].serialize()

doc.freeDoc()

return [metadata, rsp.photo[0]['originalformat']]

def getPhotoComments(self, pid):

"""Returns an XML string containing the photo comments"""

if self.verbose:

print 'Requesting comments for photo %s' % pid

rsp = \

self.fapi.photos\_comments\_getList(api\_key=self.\_\_flickrAPIKey,

auth\_token=self.token, photo\_id=pid)

if self.\_\_testFailure(rsp):

return None

doc = libxml2.parseDoc(rsp.xml)

comments = doc.xpathEval('/rsp/comments')[0].serialize()

doc.freeDoc()

return comments

def getPhotoSizes(self, pid):

"""Returns a string with is a list of available sizes for a photo"""

rsp = self.fapi.photos\_getSizes(api\_key=self.\_\_flickrAPIKey,

auth\_token=self.token, photo\_id=pid)

if self.\_\_testFailure(rsp):

return None

return rsp

def getOriginalPhoto(self, pid):

"""Returns a URL which is the original photo, if it exists"""

source = None

rsp = self.getPhotoSizes(pid)

if rsp == None:

return None

for s in rsp.sizes[0].size:

if s['label'] == 'Original':

source = s['source']

for s in rsp.sizes[0].size:

if s['label'] == 'Video Original':

source = s['source']

return [source, s['label'] == 'Video Original']

def \_\_downloadReportHook(

self,

count,

blockSize,

totalSize,

):

if not self.\_\_verbose:

return

p = ((100 \* count) \* blockSize) / totalSize

if p > 100:

p = 100

print '\r %3d %%' % p,

sys.stdout.flush()

def downloadURL(

self,

url,

target,

filename,

verbose=False,

):

"""Saves a photo in a file"""

if self.dryrun:

return

self.\_\_verbose = verbose

tmpfile = '%s/%s.TMP' % (target, filename)

if self.\_\_httplib == 'wget':

cmd = 'wget -q -t 0 -T 120 -w 10 -c -O %s %s' % (tmpfile,

url)

os.system(cmd)

else:

urllib.urlretrieve(url, tmpfile,

reporthook=self.\_\_downloadReportHook)

os.rename(tmpfile, '%s/%s' % (target, filename))

def usage():

"""Command line interface usage"""

print 'Usage: Offlickr.py -i <flickr Id>'

print 'Backs up Flickr photos and metadata'

print 'Options:'

print '\t-f <date>\tbeginning of the date range'

print '\t\t\t(default: since you started using Flickr)'

print '\t-t <date>\tend of the date range'

print '\t\t\t(default: until now)'

print '\t-d <dir>\tdirectory for saving files (default: ./dst)'

print '\t-l <level>\tlevels of directory hashes (default: 0)'

print '\t-p\t\tback up photos in addition to photo metadata'

print '\t-n\t\tdo not redownload anything which has already been downloaded (only jpg checked)'

print '\t-o\t\toverwrite photo, even if it already exists'

print '\t-L\t\tback up human-readable photo locations and permissions to separate files'

print '\t-s\t\tback up all photosets (time range is ignored)'

print '\t-w\t\tuse wget instead of internal Python HTTP library'

print '\t-c <threads>\tnumber of threads to run to backup photos (default: 1)'

print '\t-v\t\tverbose output'

print '\t-N\t\tdry run'

print '\t-h\t\tthis help message'

print '\nDates are specified in seconds since the Epoch (00:00:00 UTC, January 1, 1970).'

print '\nVersion ' + \_\_version\_\_

def fileWrite(

dryrun,

directory,

filename,

string,

):

"""Write a string into a file"""

if dryrun:

return

if not os.access(directory, os.F\_OK):

os.makedirs(directory)

f = open(directory + '/' + filename, 'w')

f.write(string)

f.close()

print 'Written as', filename

class photoBackupThread(threading.Thread):

def \_\_init\_\_(

self,

sem,

i,

total,

id,

title,

offlickr,

target,

hash\_level,

getPhotos,

doNotRedownload,

overwritePhotos,

):

self.sem = sem

self.i = i

self.total = total

self.id = id

self.title = title

self.offlickr = offlickr

self.target = target

self.hash\_level = hash\_level

self.getPhotos = getPhotos

self.doNotRedownload = doNotRedownload

self.overwritePhotos = overwritePhotos

threading.Thread.\_\_init\_\_(self)

def run(self):

backupPhoto(

self.i,

self.total,

self.id,

self.title,

self.target,

self.hash\_level,

self.offlickr,

self.doNotRedownload,

self.getPhotos,

self.overwritePhotos,

)

self.sem.release()

def backupPhoto(

i,

total,

id,

title,

target,

hash\_level,

offlickr,

doNotRedownload,

getPhotos,

overwritePhotos,

):

print str(i) + '/' + str(total) + ': ' + id + ': '\

+ title.encode('utf-8')

td = target\_dir(target, hash\_level, id)

if doNotRedownload and os.path.isfile(td + '/' + id + '.xml')\

and os.path.isfile(td + '/' + id + '-comments.xml')\

and (not getPhotos or getPhotos and os.path.isfile(td + '/'

+ id + '.jpg')):

print 'Photo %s already downloaded; continuing' % id

return

# Get Metadata

metadataResults = offlickr.getPhotoMetadata(id)

if metadataResults == None:

print 'Failed!'

sys.exit(2)

metadata = metadataResults[0]

format = metadataResults[1]

t\_dir = target\_dir(target, hash\_level, id)

# Write metadata

fileWrite(offlickr.dryrun, t\_dir, id + '.xml', metadata)

#The following lines were commented out by Patrick Wijntjes, 14-01-2014

'''# Get comments

photoComments = offlickr.getPhotoComments(id)

fileWrite(offlickr.dryrun, t\_dir, id + '-comments.xml',

photoComments)'''

# Do we want the picture too?

if not getPhotos:

return

[source, isVideo] = offlickr.getOriginalPhoto(id)

if source == None:

print 'Oopsie, no photo found'

return

# if it's a Video, we cannot trust the format that getInfo told us.

# we have to make an extra round trip to grab the Content-Disposition

isPrivateFailure = False

if isVideo:

sourceconnection = urllib.urlopen(source)

try:

format = sourceconnection.headers['Content-Disposition'].split('.')[-1].rstrip('"')

except:

print 'warning: private videos cannot be backed up due to a Flickr bug'

format = 'privateVideofailure'

isPrivateFailure = True

filename = id + '.' + format

if os.path.isfile('%s/%s' % (t\_dir, filename))\

and not overwritePhotos:

print '%s already downloaded... continuing' % filename

return

if not isPrivateFailure:

print 'Retrieving ' + source + ' as ' + filename

offlickr.downloadURL(source, t\_dir, filename, verbose=True)

print 'Done downloading %s' % filename

def backupPhotos(

threads,

offlickr,

target,

hash\_level,

dateLo,

dateHi,

getPhotos,

doNotRedownload,

overwritePhotos,

):

"""Back photos up for a particular time range"""

if dateHi == maxTime:

t = time.time()

print 'For incremental backups, the current time is %.0f' % t

print "You can rerun the program with '-f %.0f'" % t

photos = offlickr.getPhotoList(dateLo, dateHi)

if photos == None:

print 'No photos found'

sys.exit(1)

total = len(photos)

print 'Backing up', total, 'photos'

if threads > 1:

concurrentThreads = threading.Semaphore(threads)

i = 0

for p in photos:

i = i + 1

pid = str(int(p['id'])) # Making sure we don't have weird things here

if threads > 1:

concurrentThreads.acquire()

downloader = photoBackupThread(

concurrentThreads,

i,

total,

pid,

p['title'],

offlickr,

target,

hash\_level,

getPhotos,

doNotRedownload,

overwritePhotos,

)

downloader.start()

else:

backupPhoto(

i,

total,

pid,

p['title'],

target,

hash\_level,

offlickr,

doNotRedownload,

getPhotos,

overwritePhotos,

)

def backupLocation(

threads,

offlickr,

target,

hash\_level,

dateLo,

dateHi,

doNotRedownload,

):

"""Back photo locations up for a particular time range"""

if dateHi == maxTime:

t = time.time()

print 'For incremental backups, the current time is %.0f' % t

print "You can rerun the program with '-f %.0f'" % t

photos = offlickr.getGeotaggedPhotoList(dateLo, dateHi)

if photos == None:

print 'No photos found'

sys.exit(1)

total = len(photos)

print 'Backing up', total, 'photo locations'

i = 0

for p in photos:

i = i + 1

pid = str(int(p['id'])) # Making sure we don't have weird things here

td = target\_dir(target, hash\_level, pid) + '/'

if doNotRedownload and os.path.isfile(td + pid + '-location.xml'

) and os.path.isfile(td + pid

+ '-location-permissions.xml'):

print pid + ': Already there'

continue

location = offlickr.getPhotoLocation(pid)

if location == None:

print 'Failed!'

else:

fileWrite(offlickr.dryrun, target\_dir(target, hash\_level,

pid), pid + '-location.xml', location)

locationPermission = offlickr.getPhotoLocationPermission(pid)

if locationPermission == None:

print 'Failed!'

else:

fileWrite(offlickr.dryrun, target\_dir(target, hash\_level,

pid), pid + '-location-permissions.xml',

locationPermission)

def backupPhotosets(offlickr, target, hash\_level):

"""Back photosets up"""

photosets = offlickr.getPhotosetList()

if photosets == None:

print 'No photosets found'

sys.exit(0)

total = len(photosets)

print 'Backing up', total, 'photosets'

i = 0

for p in photosets:

i = i + 1

pid = str(int(p['id'])) # Making sure we don't have weird things here

print str(i) + '/' + str(total) + ': ' + pid + ': '\

+ p.title[0].text.encode('utf-8')

# Get Metadata

info = offlickr.getPhotosetInfo(pid,

offlickr.fapi.photosets\_getInfo)

if info == None:

print 'Failed!'

else:

fileWrite(offlickr.dryrun, target\_dir(target, hash\_level,

pid), 'set\_' + pid + '\_info.xml', info)

photos = offlickr.getPhotosetInfo(pid,

offlickr.fapi.photosets\_getPhotos)

if photos == None:

print 'Failed!'

else:

fileWrite(offlickr.dryrun, target\_dir(target, hash\_level,

pid), 'set\_' + pid + '\_photos.xml', photos)

# Do we want the picture too?

def target\_dir(target, hash\_level, id):

dir = target

i = 1

while i <= hash\_level:

dir = dir + '/' + id[len(id) - i]

i = i + 1

return dir

def main():

"""Command-line interface"""

# Default options

flickrUserId = None

dateLo = '1'

dateHi = maxTime

getPhotos = False

overwritePhotos = False

doNotRedownload = False

target = 'dst'

photoLocations = False

photosets = False

verbose = False

threads = 1

httplib = None

hash\_level = 0

dryrun = False

# Parse command line

try:

(opts, args) = getopt.getopt(sys.argv[1:],

'hvponNLswf:t:d:i:c:l:', ['help'])

except getopt.GetoptError:

usage()

sys.exit(2)

for (o, a) in opts:

if o in ('-h', '--help'):

usage()

sys.exit(0)

if o == '-i':

flickrUserId = a

if o == '-p':

getPhotos = True

if o == '-o':

overwritePhotos = True

if o == '-n':

doNotRedownload = True

if o == '-L':

photoLocations = True

if o == '-s':

photosets = True

if o == '-f':

dateLo = a

if o == '-t':

dateHi = a

if o == '-d':

target = a

if o == '-w':

httplib = 'wget'

if o == '-c':

threads = int(a)

if o == '-l':

hash\_level = int(a)

if o == '-N':

dryrun = True

if o == '-v':

verbose = True

# Check that we have a user id specified

if flickrUserId == None:

print 'You need to specify a Flickr Id'

sys.exit(1)

# Check that the target directory exists

if not os.path.isdir(target):

print target + ' is not a directory; please fix that.'

sys.exit(1)

offlickr = Offlickr(

flickrAPIKey,

flickrSecret,

flickrUserId,

httplib,

dryrun,

verbose,

)

if photosets:

backupPhotosets(offlickr, target, hash\_level)

elif photoLocations:

backupLocation(

threads,

offlickr,

target,

hash\_level,

dateLo,

dateHi,

doNotRedownload,

)

else:

backupPhotos(

threads,

offlickr,

target,

hash\_level,

dateLo,

dateHi,

getPhotos,

doNotRedownload,

overwritePhotos,

)

if \_\_name\_\_ == '\_\_main\_\_':

main()

### Python for website

#### forms.py

# Import the required modules

from django import forms

from models import Orchid

# Class for uploading pictures

class UploadPictureForm(forms.ModelForm):

# The meta data

class Meta:

# The used model, Orchids in this case

model = Orchid

#### result.py

#Import the required module

import sys

#Function for creating the variables

def create\_variables():

#Save the ip-adress which is the second commandline argument

ip = sys.argv[1]

#Create a list with the possible sections.

#Sort them so form Z-A. The index of the section

#Now is corresponding the index of the list from the classifly.pl script

sections = ["Parvisepalum", "Pardalopetalum", "Paphiopedilum", "Coryopedilum", "Cochlopetalum", "Brachypetalum", "Barbata"]

#Open the output file from the classifly.pl script.

infile = open("%s\_out.txt"%(ip), 'r')

#Read in the content of the output file

file1 = infile.readlines()

#Create a variable for the index

index = -1

#Create an empty list. This list will be used

#To save the values of the output list form classify.pl

values = []

#Create a counter

counter = -1

#Create an output file

output = open("%s\_result.txt"%(ip) , 'w')

#Run the get\_section function, give it all the created variables.

get\_section(sections, file1, index, values, counter, output)

def get\_section(sections, file1, index, values, counter, output):

#Loop through the lines of the file

for line in file1:

#For every loop add 1 to the counter

counter += 1

#To accept error use a try-except

try:

#Find the lines which contains the values

if counter >= 1 and counter <= 7:

#Add these values to the values list

values.append(float(line.strip().strip(",").strip("'")))

#If an error occur, except this and continue

except:

continue

#Loop through the values

for number in values:

#When the value is bigger than 0, the picture

#Is classified to this section.

if number > 0:

#Get the index of this value

index = values.index(number)

#Otherwise

else:

#Go on

continue

#The result of the cassification script is the section

#Which index corresponds to the index of the positive value

result = sections[index]

#Write the result to this file and close the file

output.write("%s"%(result))

output.close()

#Run the create\_variables function to create the variables

create\_variables()

#### views.py

# import the required modules

from django.http import HttpResponseRedirect

from django.shortcuts import render\_to\_response

from forms import UploadPictureForm

from django.core.context\_processors import csrf

from django.contrib import auth

from time import time

from django.contrib.auth.decorators import login\_required

import os

# Function to get the used devise.

def get\_device( request ):

""" Redirect to the servers list. """

#Initiate the device variable

device = ""

#If the used device is in the list, the device is a mobile phone

'''I have test both html-styles on the iPad. The results shows that the iPad can

better show the computer style'''

if 'HTTP\_USER\_AGENT' in request.META and (

request.META['HTTP\_USER\_AGENT'].startswith( 'BlackBerry' ) or \

"Opera Mobi" in request.META.get('HTTP\_USER\_AGENT') or \

"Opera Mini" in request.META.get('HTTP\_USER\_AGENT') or \

"Windows CE" in request.META.get('HTTP\_USER\_AGENT') or \

"MIDP" in request.META.get('HTTP\_USER\_AGENT') or \

"Palm" in request.META.get('HTTP\_USER\_AGENT') or \

"NetFront" in request.META.get('HTTP\_USER\_AGENT') or \

"Nokia" in request.META.get('HTTP\_USER\_AGENT') or \

"Symbian" in request.META.get('HTTP\_USER\_AGENT') or \

"UP.Browser" in request.META.get('HTTP\_USER\_AGENT') or \

"UP.Link" in request.META.get('HTTP\_USER\_AGENT') or \

"WinWAP" in request.META.get('HTTP\_USER\_AGENT') or \

"Android" in request.META.get('HTTP\_USER\_AGENT') or \

"DoCoMo" in request.META.get('HTTP\_USER\_AGENT') or \

"KDDI-" in request.META.get('HTTP\_USER\_AGENT') or \

"Softbank" in request.META.get('HTTP\_USER\_AGENT') or \

"J-Phone" in request.META.get('HTTP\_USER\_AGENT') or \

"IEMobile" in request.META.get('HTTP\_USER\_AGENT') or \

"iPod" in request.META.get('HTTP\_USER\_AGENT') or \

"iPhone" in request.META.get('HTTP\_USER\_AGENT') ):

device = "mobile"

#Otherwise it is a computer.

else:

device = "computer"

#Return the device

return device

# Function to check if the uploaded file is a picture

def check\_upload(upload):

#Create a list with picture extentions

picture = ["jpg","png","jpeg"]

#Get the extension from the uploaded file

name = str(upload)

extension = name.lower().split(".")[-1]

#Check if the extension is in the picture list

if extension in picture:

#If it is, return true

return True

#Otherwise

else:

#Remove the file from the server

name = name.replace(" ","\ ")

os.system("rm static/uploaded\_files/%s"%(name))

#Retrun false

return False

# Welcome view (homepage)

def welcome(request):

#Get the used device, using the get\_device function

device = get\_device(request)

# Create the args dictionary and save the csrf in this dictonary

args = {}

args.update(csrf(request))

# Save the html name, with the used device

html = device+"\_welcome.html"

# Call the html, for the correct device, for de welcome page.

return render\_to\_response(html, args)

#Function to give the uploaded file a variable part in front of the filename

def processUpload(request, filename):

#To use paths spaces need to be replaced by \<space>

filename = str(filename).replace(" ","\ ")

#To make the new filename easier to acces repalce the spaces by "\_"

filename2 = str(filename).replace(" ","\_")

# Get the IP-adres of the computer

x\_forwarded\_for = request.META.get('HTTP\_X\_FORWARDED\_FOR')

if x\_forwarded\_for:

ip = x\_forwarded\_for.split(',')[0]

else:

ip = request.META.get('REMOTE\_ADDR')

# Replace the '.' in the ip-adres to '\_'

ip = ip.replace('.', '\_')

# Create an output file named <ip>.filename.txt

outfile = open('%s.filename.txt' %(ip), 'w')

# Place the variable part (the ip) in front of the filename of the uploaded file

os.system("mv static/uploaded\_files/%s static/uploaded\_files/%s\_%s"%(filename, ip, filename2))

# Write the new filename to the outputfile

outfile.write("%s\_%s" %(ip, filename2))

# Close the outputfile

outfile.close()

# The upload view (choice file and upload it)

def upload(request):

#Get the used device, using the get\_device function

device = get\_device(request)

# Get the IP-adres of the computer

x\_forwarded\_for = request.META.get('HTTP\_X\_FORWARDED\_FOR')

if x\_forwarded\_for:

ip = x\_forwarded\_for.split(',')[0]

else:

ip = request.META.get('REMOTE\_ADDR')

# Replace the '.' in the ip-adres to '\_'

ip = ip.replace('.', '\_')

#If the user uploaded a file that isn't a picture, a message will be displaid.

#Also the color of the upload button will be red.

#To make this posible, the variables need to be created befor the if statement.

message = ""

style = ""

# Check if the method is POST

if request.method == 'POST':

#If the method is POST give the message and style variables the

#correct values

message = "You didn't select a picture"

style = "color:red"

# Save the user input from the form

form = UploadPictureForm(request.POST, request.FILES)

# Check if the form is valid

if form.is\_valid():

# Save the form

form.save()

#Check if the uploaded file is a picture

is\_picture = check\_upload(request.FILES["picture"])

#When the uploaded file is a picture

if is\_picture:

# run the processUpload function to place the ip in front of the name of the uploaded file

processUpload(request, request.FILES["picture"])

''' save the filename and path in python variables

use the variable part (the ip) to create the path'''

filename = str(request.FILES["picture"]).replace(" ", "\_")

path = ("static/assets/uploaded\_files/%s\_%s" % (ip, filename))

# Create the args dictionary and save the csrf in this dictonary

args = {}

args.update(csrf(request))

# save the filename and path in the dictionary

args['filename'] = filename

args['path'] = path

# Save the html name, whit the used device

html = device+"\_upload\_succes.html"

# Call the upload\_succes html, for the correct device and give it the args dictonary

return render\_to\_response(html, args)

#When the uploaded file isn't a picture

else:

# Create the args dictionary and save the csrf in this dictonary

args = {}

args.update(csrf(request))

# Save the empty form, message and style in the dictionary

args['form'] = UploadPictureForm()

args['message'] = message

args['style'] = style

# Save the html name, with the used device

html = device+"\_upload.html"

# Call the upload html, for the correct device and give it the args dictionary

return render\_to\_response(html, args)

# When the method is not POST

else:

# Create a form to upload a picture

form = UploadPictureForm()

# Create the args dictionary and save the csrf in this dictonary

args = {}

args.update(csrf(request))

# Save the empty form, message and style in the dictionary

args['form'] = UploadPictureForm()

args['message'] = message

args['style'] = style

# Save the html name, with the used device

html = device+"\_upload.html"

# Call the upload html, for the correct device and give it the args dictionary

return render\_to\_response(html, args)

# The result view (to display the result of the analysis)

def result(request):

#Get the used device, using the get\_device function

device = get\_device(request)

try:

# Get the IP-adres of the computer

x\_forwarded\_for = request.META.get('HTTP\_X\_FORWARDED\_FOR')

if x\_forwarded\_for:

ip = x\_forwarded\_for.split(',')[0]

else:

ip = request.META.get('REMOTE\_ADDR')

# Replace the '.' in the ip-adres to '\_'

ip = ip.replace('.', '\_')

#Run converter.sh to convert jpg files to png

os.system("sh converter.sh %s"%(ip))

# Read in the filename from <ip>.filename.txt

infile = open('%s.filename.txt' %(ip), 'r')

filename = infile.read().strip()

# Close the infile

infile.close()

# Run the program to classify the orchid

os.system("python classify.py %s %s" % (filename, ip))

#After the previous step a list with numbers is created.

#Runt result.py to translate this list to a readable result.

os.system("python result.py %s" % (ip))

# Open the file with the result from the result.py program

result = open('%s\_result.txt' %(ip), 'r')

# Read in the result

read\_result = result.read()

# Close the file

result.close()

# Create the args dictionary and save the csrf in this dictonary

args = {}

args.update(csrf(request))

# Save the filename, the result and the ip in the args dictionary

args['filename'] = filename

args['result'] = read\_result

args['ip'] = ip

# Save the html name with the used device

html = device+"\_result.html"

# Call the result html, for the correct device, with the args dictionary

return render\_to\_response(html, args)

except IOError:

'''If an IOError occure, the picture is uploaded just when the administrator removed all

unused files. So the uploaded picture is also removed. Send the user to the sorry page,

which tells the user to try uploading again.'''

# Save the html name with the used device

html = device+"\_sorry.html"

# Go to the sorry html, for the correct device

return render\_to\_response(html)

# The exit view (to "close" the app and remove all created temporary files)

def exit(request):

# Get the IP-adres of the computer

x\_forwarded\_for = request.META.get('HTTP\_X\_FORWARDED\_FOR')

if x\_forwarded\_for:

ip = x\_forwarded\_for.split(',')[0]

else:

ip = request.META.get('REMOTE\_ADDR')

# Replace the '.' in the ip-adres to '\_'

ip = ip.replace('.', '\_')

''' Create the variable part for the filename using a timestamp.

replace all . into \_ to prevent errors for the extension '''

var\_part = str(time()).replace('.', '\_')

# Read in the filename from <ip>.filename.txt, save it and close the file

infile = open('%s.filename.txt' %(ip), 'r')

filename = infile.read().strip()

infile.close()

# Remove the temporary file <ip>.filename.txt

# Move the uploaded picture and its result to the result directory,

# Save it as timestamp\_ip.png, timestamp\_ip\_result.txt and timestamp\_ip\_section.txt

os.system("rm %s.filename.txt" %(ip))

os.system("mv static/uploaded\_files/%s/%s results/%s\_%s" %(ip, filename, var\_part, filename))

os.system("rm -r static/uploaded\_files/%s" %(ip))

os.system("mv %s\_out.txt results/%s\_%s\_result.txt" %(ip, var\_part, ip))

os.system("mv %s\_result.txt results/%s\_%s\_section.txt" %(ip, var\_part, ip))

# Go back to the welcome page

return HttpResponseRedirect('/welcome')

# To remove all leftover files, login is required

def login(request):

#Get the used device, using the get\_device function

device = get\_device(request)

# Create a dictionary and put the csrf in it

args = {}

args.update(csrf(request))

#Save the html name with the used device

html=device+"\_login.html"

#Go to the login html, for the correct device, give it the dictionary

return render\_to\_response(html, args)

# Function to check the username and password

def auth\_view(request):

# Get the username and password

username = request.POST.get('username', '')

password = request.POST.get('password', '')

''' If the username and password are incorrect user will be None

Otherwise it will be the user '''

user = auth.authenticate(username=username, password=password)

''' Go to the correct page (admin/remove for correct login, accounts/invalid for

invalid login)'''

if user is not None:

#Login the user

auth.login(request, user)

return HttpResponseRedirect('/admin/remove')

else:

return HttpResponseRedirect('/accounts/invalid')

# function for logout

def logout(request):

#Log the user out

auth.logout(request)

#Go back to the welcome page

return HttpResponseRedirect('/welcome/')

# Function for invalid login

def invalid\_login(request):

#Get the used device, using the get\_device function

device = get\_device(request)

# Go to the invalid login html, for the correct device

html = device+"\_invalid\_login.html"

return render\_to\_response(html)

@login\_required

#User need to be registreded. Even when the user is not active this user can login and remove the files.

def remove(request):

#Get the used device, using the get\_device function

device = get\_device(request)

# List all the files that will be removed using a command line command (ls)

'''Save the name(s) of the picture(s) that will be removed in uploads.txt and the

name(s) of the temporary file(s) in temps.txt'''

os.system("ls static/uploaded\_files > uploads.txt")

os.system("ls | egrep \*.filename.txt > temps.txt")

#Remove all the unused pictures and their temporary files

os.system("rm -r static/uploaded\_files/\*")

os.system("rm \*filename.txt")

#Read the content of the uploads.txt file and the temps.txt file and save the content in

# Python variables

uploads\_in = open("uploads.txt", 'r')

temps\_in = open("temps.txt", 'r')

uploads = uploads\_in.read()

temps = temps\_in.read()

# Create the args dictionary and save the csrf in this dictonary

args = {}

args.update(csrf(request))

#Save the list of the pictures that will be removed in the dictionary

args['uploads'] = uploads

#Save the list of the temporary files that will be removed in the dictionary

args['temps'] = temps

# Remove the text files wich contain the lists

os.system("rm uploads.txt temps.txt")

# Save the html name with the used device

html = device+"\_remove.html"

# Call the html, for the correct device, and give it the args directory

return render\_to\_response(html, args)