Application of Deep Learning techniques for the photo-identification of fish individuals

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Model implementation: code outline

The main file of our system for training is "main_training.py", whose function calls are outlined below. The optimization of hyperparameters in "bayesian_optimization_parameters.py" works essentially the same way, but without fixing the parameters to a value.

```
The following parameters are set: dataset_path, batch_size, load_dataset_again, use_augmentation,
params_augmentation, learning_rate, momentum, std_prob_threshold, input_shape, model_name,
tensorboard_log_path
Instantiating class...
siamese_network = SiameseNetwork(...)
                                          et process = 'train'
          Instantiating class...

Images_loader= Loader(...)

The following parameters are set image_height, image_width, images_dictionary, ray_names,
               train\_dictionary, \ source\_to\_evaluate, \ validation\_dictionary, \ evaluation\_dictionary,
               evaluable_rays
                    nages are loaded with load_dataset() and stored in images_dictionary
                use_augmentation is true then the dictionary of images is augmented...
   images_dictionary = ImageAugmentor(...).apply_augmentation()
                Split the dataset with divide_train_valid_eval() which, for each ray in evaluable_rays, takes n_eval images for test
set, and divides the remaining in 75%-25% for train and validation. All images of non evaluable rays are for the train set.
                                            summary_writer
                                                 with constructuct_siamese_architecture() as defined in the methodology, which al
The training starts when the following parameters are set: validate_each, number_of_train_iterations...
validation_accuracy = siamese_network.train_siamese_network()
                                 number of train iterations
               images, labels = convert_path_list_to_images_and_labels(images_path, is_one_shot_task =
                                              a list of two lists of length number_of_pairs in wh
ts or 0s depending on wheter the respective pair is a s
                      False)
                 train_loss, train_accuracy = model.train_on_batch(images, labels)
                validate_each
                 validation accuracy = images loader.few shots task()
For each ray in evaluable_rays:
                                    rin evaluable_rays
or each test_image of that ray:
    images, _ = get_one_shot_batch(current_ray = ray, test_image), which creates
    the batch of pairs of images; the test_image of the current_ray against an image of the
    support set of that ray and against an image of all the remaining evaluable rays (i.e., the labels are
                                             probabilities = model.predict_on_batch(images)
                                                                                                             th of its test_image
                           validation_accuracy
Once trained and validated, we make the predictions over the evaluation (test) set loading the weights of the best model
```

evaluation_accuracy = siamese_network.predict_after_train(), which predicts following the defined methodology
that resembles Ensemble Learning.

Figure 1: Training outline.

And the main file for prediction is "main_prediction.py", which shares classes with the previous scheme but whose outline goes as follows:

```
The following parameters are set: dataset_path, input_shape, model_name

We set the architecture of our net...
siamese_network = SiameseNetwork(...)

And load the model (i.e., the weights) with which we want to predict.

Then, we load in the dictionary images_to_predict the images of the unidentified individuals that we want to predict, which are those in folders containing "new_recapture".

Predicting with...

flag_new_individuals = siamese_network.load_and_predict(images_to_predict), which predicts following the defined methodology that resembles Ensemble Learning and creates the list flag_new_individuals that will contain the path of the individuals detected as new ones.

We store the information of new individuals in a .txt file...

store_new_individuals(dataset_path, flag_new_individuals)

Finally, when enough new individuals have been gathered, we process them to create new folders of identification...

siamese_network.process_new_individuals()

"Creating pairs to compare in a pairwise manner...\n "
    "Predicting same/different over the pair:"
    print(pair)
    images, _ = images_loader._convert_path_list_to_images_and_labels(pair, is_one_shot_task = True)
    probability = model.predict_on_batch(images)[0][0]

if probability = model.predict_on_batch(images)[0][0][0]
```

Figure 2: Prediction outline.