



# UNIVERSITÀ DI PISA

Artificial Intelligence and Data Engineering

Computational Intelligence and Deep Learning

## *Artist Identification with Convolutional Neural Networks*

Project Documentation

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# 1 — Introduction

Artist identification is traditionally performed by *art historians* and *curators* who have expertise and familiarity with different artists and styles of art. This is a complex and interesting problem for computers because identifying an artist does not just require object or face detection; artists can paint a wide variety of objects and scenes. Additionally, many artists from the same time period will have similar styles, and some such as **Pablo Picasso** (see figure 1) have painted in multiple styles and changed their style over time.



**Figure 1:** Both of these paintings were created by Pablo Picasso, but they have vastly different styles and content since they correspond to two different periods.

The aim of this project is to use Convolutional Neural Networks for the identification of an artist given a painting. In particular, the CNN networks will be modeled using multiple techniques: from scratch; via pretrained network; networks based on comparison with baseline input and using an ensemble network made of by the best classifiers found.

## 1.1 State of the Art

As mentioned, artist identification has primarily been tackled by humans. An example of that is the Artsy’s Art Genome Project <sup>1</sup>, which is led by experts who manually classify art. This strategy is not very scalable even if it is highly precise in the classification (the site is a marketplace of fine-arts for collects, you can find *Pisarro*, *Banksy* and other famous artists).

Most prior attempts to apply machine learning to this problem have been feature-based, aiming to identify what qualities most effectively distinguish artists and styles. Many generic image features have been used, including scale-invariant feature transforms (SIFT), histograms of oriented gradients (HOG), and more, but with the focus on *discriminating different style* in Fine-Art Painting<sup>2</sup>.

The first time the problem of artist identification was really tackled was with J. Jou and S. Agrawal<sup>3</sup> in 2011, they applied several multi-class classification techniques like Naïve Bayes, Linear Discriminant Analysis, Logistic Regression, K-Means and SVMs and achieve a maximum classification accuracy of 65% for an unknown painting across 5 artists. Later on, the problem of identifying artists was retackled by the *Rijksmuseum Challenge*<sup>4</sup>. The objective of the challenge was to predict the artist, type, material and creation year (each of them was a different challenge) of the 112,039 photographic <sup>5</sup> (containing different viewpoints of an artwork, and different types of them: sculptures, paintings, saucers, etc.) reproductions of the artworks exhibited

<sup>1</sup><https://www.artsy.net/categories>

<sup>2</sup>T. E. Lombardi. The classification of style in fine-art painting. ETD Collection for Pace University, 2005

<sup>3</sup>J. Jou and S. Agrawal. Artist identification for renaissance paintings.

<sup>4</sup>T. Mensink and J. van Gemert. The rijksmuseum challenge: Museum-centered visual recognition. 2014

<sup>5</sup>The dataset contains 6,629 artists in total, with high variation in the number of pieces per artist. For example, Rembrandt has 1,384 pieces, and Vermeer has only 4. There are 350 artists with more than 50 pieces, 180 artists have around 100, and 90 artists have 200 pieces.

in the Rijksmuseum in Amsterdam (the Netherlands). For the artist classification challenge, the paper said they read a test accuracy of about 60%. The year later, Saleh and Elgammal’s paper<sup>6</sup> was the first attempt to identify artists with a large and varied dataset, but still using generic features. The collection they used has images of 81,449 fine-art paintings from 1,119 artists ranging from fifteen centuries to contemporary artists, reaching an accuracy of 59%<sup>7</sup>.

More recent attempts are related to the *Painter by Numbers*, a **Playground Prediction Competition** by *Kaggle*<sup>8</sup>. This competition used a pairwise comparison scheme: participants had to create an algorithm which needs to examine two images and predict whether the two images are by the same artist or not. Thus, it is not our same objective, however it can be consider the first application of Deep Learning to the problem. The real deal was taken by Nitin Viswanathan<sup>9</sup> in 2017. Viswanathan, using the same dataset of the mentioned *Kaggle Challenge*, proposed the use of ResNet with transfer learning (he first held the weights of the base ResNet constant and updated only the fully-connected layer for a few epochs). This trained network reached a train accuracy of 0.973 and a test accuracy of 0.898.

## 1.2 Dataset

Unfortunately, the dataset provided by the *Kaggle Challenge* is too huge to be used in Colab, in fact it is about 60GB unbearable on the free version of Colab, which provides only about 30GB of disk. Stated that, we decided to use a different dataset<sup>10</sup> with only 2GB of data and about 8k unique images.

The data downloaded from Kaggle has the following directories and csv file:

```

/
├── images
│   └── images
│       ├── Albrecht_Durer
│       ├── Alfred_Sisley
│       ├── Amedeo_Modigliani
│       ├── Andrei_Rublev
│       ├── Andy_Warhol
│       ├── Camille_Pissarro
│       ├── Caravaggio
│       ├── Claude_Monet
│       ├── Diego_Rivera
│       ├── Diego_Velazquez
│       ├── Edgar_Degas
│       ├── Edouard_Manet
│       ├── Edvard_Munch
│       └── an many others (total of 50 different artists)
├── resized
└── artists.csv

```

The *resized* directory is not useful for our studies, hence we deleted it to save space on the disk. On the other hand, we first use the *csv* file to select only the artists with at least 200 pieces, this operation was done to reduce the number of classes to a number per which the ratio between the number of artists and images was reasonable for learning. Even done that, the dataset was still unbalanced, e.g. Van Gogh’s paintings are 877 against the 239 of Chagall’s,

<sup>6</sup>B. Saleh and A. M. Elgammal. Large-scale classification of fine-art paintings: Learning the right metric on the right feature. CoRR, abs/1505.00855, 2015

<sup>7</sup>In the paper they tried to use also CNN, but reaching only an accuracy of 33.62%

<sup>8</sup><https://www.kaggle.com/c/painter-by-numbers/data>

<sup>9</sup>Nitin Viswanathan, Artist Identification with Convolutional Neural Networks

<sup>10</sup><https://www.kaggle.com/ikarus777/best-artworks-of-all-time>

thus we consider to compute **class weights** in order to use them in the *fit function*:

$$\text{class\_weights} = \frac{\text{Total number of paintings considered}}{\text{Number of artists considered} \cdot \text{Number of paintings per author}}$$

Then, we modified the structure of the *images/images* directory in order to create two directories, **train** and **test**, containing 90% and 10% of the images from each different artist's directory respectively (considering only the artists with at least 200 paintings). The newly created directories have the same structured of *images/images*. This was done in *python* in this way:

```

1 import os
2 import numpy as np
3 import shutil
4
5 rootdir= '/content/images/images' #path of the original folder
6 classes = os.listdir(rootdir)
7
8 for i, c in enumerate(classes, start=1):
9     if c not in artists_top_name.tolist():
10         shutil.rmtree(rootdir + '/' + c)
11         continue
12     if not os.path.exists(rootdir + '/train/' + c):
13         os.makedirs(rootdir + '/train/' + c)
14     if not os.path.exists(rootdir + '/test/' + c):
15         os.makedirs(rootdir + '/test/' + c)
16
17     source = os.path.join(rootdir, c)
18     allFileNames = os.listdir(source)
19
20     np.random.shuffle(allFileNames)
21
22     test_ratio = 0.10
23     train_FileNames, test_FileNames = np.split(np.array(allFileNames),
24                                                [int(len(allFileNames)*
25                                                    (1 - test_ratio))])
26
27     train_FileNames = [source+'/' + name for name in train_FileNames.tolist()]
28     test_FileNames = [source+'/' + name for name in test_FileNames.tolist()]
29
30     for name in train_FileNames:
31         shutil.copy(name, rootdir + '/train/' + c)
32
33     for name in test_FileNames:
34         shutil.copy(name, rootdir + '/test/' + c)

```

After that we created the train/validation/test-sets using the *image\_dataset\_from\_directory* function provided by **Keras** in the following way:

```

1 import tensorflow as tf
2
3 training_images = tf.keras.preprocessing.image_dataset_from_directory(
4     TRAIN_DIR, labels='inferred', label_mode='categorical',
5     class_names=None, color_mode='rgb', batch_size=BATCH_SIZE,
6     image_size=(IMAGE_HEIGHT, IMAGE_WIDTH), shuffle=True, seed=RANDOM_SEED,
7     validation_split=VALIDATION_SPLIT, subset='training',
8     interpolation='bilinear', follow_links=False
9 )
10
11 val_images = tf.keras.preprocessing.image_dataset_from_directory(
12     TRAIN_DIR, labels='inferred', label_mode='categorical',
13     class_names=None, color_mode='rgb', batch_size=BATCH_SIZE,
14     image_size=(IMAGE_HEIGHT, IMAGE_WIDTH), shuffle=True, seed=RANDOM_SEED,
15     validation_split=VALIDATION_SPLIT, subset='validation',
16     interpolation='bilinear', follow_links=False
17 )

```

```
18
19 test_images = tf.keras.preprocessing.image_dataset_from_directory(
20     TEST_DIR, labels='inferred', label_mode='categorical',
21     class_names=None, color_mode='rgb', batch_size=BATCH_SIZE,
22     image_size=(IMAGE_HEIGHT, IMAGE_WIDTH), shuffle=True, seed=RANDOM_SEED,
23     interpolation='bilinear', follow_links=False
24 )
```

Where *VALIDATION\_SPLIT* is equal to 0.1.

Obtaining in this way:

- 3478 files for training (belonging to 11 classes).
- 386 files for validation (belonging to 11 classes).
- 438 files for testing (belonging to 11 classes).

Hence, we have a total of 4299 different pictures.

## 2 — General Information Useful for Training

In the following chapters we will make use of different strategies:

- Class Weights (already talked about)
- Data augmentation
- Regularization
- Dropout
- Multiple activation functions
- Multiple optimizers
- Genetic Algorithms

In order to allow a better and faster reading of the tests done, in the following paragraph the mentioned strategies are discussed.

### 2.1 Data Augmentation

Data augmentation takes the approach of generating more training data from existing training samples by augmenting the samples via a number of random transformations that yield believable-looking images. The goal is that, at training time, your model will never see the exact same picture twice. This helps expose the model to more aspects of the data so it can generalize better. In Keras, this can be done by adding a number of data augmentation layers at the start of your model. In our model, we included the following transformation:

```
1 data_augmentation = ks.Sequential(  
2     [  
3         layers.RandomFlip('horizontal'),  
4         layers.RandomRotation(0.1),  
5         layers.RandomZoom(0.2),  
6         layers.RandomHeight(0.1),  
7         layers.RandomWidth(0.1)  
8     ]  
9 )
```

### 2.2 Regularization

Regularization techniques are a set of best practices that actively impede the model's ability to fit perfectly to the training data, with the goal of making the model perform better during validation. This is called “regularizing” the model, because it tends to make the model simpler, more “regular”, its curve smoother, more “generic”; thus it is less specific to the training set and better able to generalize by more closely approximating the latent manifold of the data. A common way to mitigate overfitting is to put constraints on the complexity of a model by forcing its weights to take only small values, which makes the distribution of weight values more regular. This is called *weight regularization*, and it's done by adding to the loss function of the model a cost associated with having large weights. This cost comes in two flavors:

1. *L1 regularization*—The cost added is proportional to the absolute value of the weight coefficients (the L1 norm of the weights).



2. *L2 regularization*—The cost added is proportional to the square of the value of the weight coefficients (the L2 norm of the weights).
3. *L1-L2 regularization*—Combine L1 and L2.

## 2.3 Dropout

Dropout is one of the most effective and most commonly used regularization techniques for neural networks; it was developed by Geoff Hinton and his students at the University of Toronto. Dropout, applied to a layer, consists of randomly dropping out (setting to zero) a number of output features of the layer during training.

In `keras` can be set using the `layers.Dropout` function passing as parameter the *dropout rate*, in our case always 0.5.

## 2.4 Activation Functions

In the studies done in the following chapters we used three different activation functions:

- *ReLU*:  $\max(0, x)$
- *ELU*:  $\max(0.2x, x)$
- *Leaky ReLU*:  $f(x) = \begin{cases} x & x > 0 \\ \alpha(\exp(x) - 1) & x \leq 0 \end{cases}$

They will be useful in the genetic algorithm analysis done fore the *scratch architecture* and the *VGG16*

## 2.5 Optimizers

An optimizer is the mechanism through which the model will update itself based on the training data it sees, so as to improve its performance. In our project we make use of:

- *RMSprop*: the gist of RMSprop is to:
  - Maintain a moving (discounted) average of the square of gradients
  - Divide the gradient by the root of this average
  - It uses plain momentum, not Nesterov momentum.
- *Adam*: stochastic gradient descent method that is based on adaptive estimation of first-order and second-order moments.

## 2.6 Genetic Algorithms

Genetic algorithms are a family of search algorithms inspired by the principles of evolution in nature. By imitating the process of natural selection and reproduction, genetic algorithms can produce high-quality solutions for various problems involving search, optimization, and learning. At the same time, their analogy to natural evolution allows genetic algorithms to overcome some of the hurdles that are encountered by traditional search and optimization algorithms, especially for problems with a large number of parameters and complex mathematical representations. Thus, they come in handy for optimizing our networks. In order to make use of genetic algorithms we must decide some components, which are:

- Genotype
- Population

- Fitness Function
- Selection Algorithm
- Crossover Algorithm
- Mutation Algorithm
- Elitism

All of these components are implemented using the python library **deap**<sup>11</sup>.

### 2.6.1 Genotype

The *genotype* is a collection of genes that are grouped into chromosomes. In our specific case, our genes are bounded real-valued encoded and they represent three different parameters:

- *activation\_function*: bounded between 0, 1.999, 2.999. The integer part stands for ReLU (0), ELU (1), Leaky Relu (2);
- *optimizer*: bounded between 0 and 1.999. The integer part stands for rmsprop (0) and adam (1);
- *learning rate*: bounded between 0.001 and 0.1.

### 2.6.2 Population

At any point in time, genetic algorithms maintain a population of individuals (i.e., chromosomes)—a collection of candidate solutions for the problem at hand. Since for each individual we train a model and evaluate its performance, we decided to use only 20 individuals per generation to limit the computation.

### 2.6.3 Fitness Function

At each iteration of the algorithm, the individuals are evaluated using a fitness function (also called the target function). This is the function we seek to optimize or the problem we attempt to solve. In our problem, the fitness function is the the function which calculate the maximum validation accuracy reached by the model. Our objective is maximizing the validation accuracy.

### 2.6.4 Selection Algorithm

After calculating the fitness of every individual in the population, a selection process is used to determine which of the individuals in the population will get to reproduce and create the offspring that will form the next generation. The selection algorithm used by as is *tournament selection*. In each round of the tournament selection method, two individuals are randomly picked from the population, and the one with the highest fitness score wins and gets selected. We decided to select only two individuals since our population is small and selecting more could cause an abuse in exploitation.

### 2.6.5 Crossover Algorithm

To create a pair of new individuals, two parents are chosen from the current generation, and parts of their chromosomes are interchanged (crossed over) to create two new chromosomes representing the offspring. There exists different algorithms applicable to real-value encoded

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<sup>11</sup><https://deap.readthedocs.io/en/master/>

individuals, we decided to use the *Simulated Binary Bounded Crossover*, which is a bounded version of the Simulated Binary Crossover (SBX)<sup>12</sup>.

The idea behind the simulated binary crossover is to imitate the properties of the single-point crossover that is commonly used with binary-coded chromosomes. One of these properties is that the average of the parents' values is equal to that of the offsprings' values. When applying SBX, the two offspring are created from the two parents using the following formula:

$$offspring_1 = \frac{1}{2}[(1 + \beta)parent_1 + (1 - \beta)parent_2]$$

$$offspring_2 = \frac{1}{2}[(1 - \beta)parent_1 + (1 + \beta)parent_2]$$

Here,  $\beta$  is a random number referred to as the *spread factor*.

This formula has the following notable properties:

- The average of the two offspring is equal to that of the parents, regardless of the value of  $\beta$ .
- When the  $\beta$  value is 1, the offspring are duplicates of the parents.
- When the  $\beta$  value is smaller than 1, the offspring are closer to each other than the parents were.
- When the  $\beta$  value is larger than 1, the offspring are farther apart from each other than the parents were.

The probability to mate is set equal to 0.9.

### 2.6.6 Mutation Algorithm

The purpose of the mutation operator is to periodically and randomly refresh the population, introduce new patterns into the chromosomes, and encourage search in uncharted areas of the solution space. As mutation algorithm we decided to use the *Polynomial Bounded* method, which is a bounded mutation operator that uses a polynomial function for the probability distribution.

The probability to mutate is set equal to 0.5.

### 2.6.7 Elitism

While the average fitness of the genetic algorithm population generally increases as generations go by, it is possible at any point that the best individual(s) of the current generation will be lost. This is due to the selection, crossover, and mutation operators altering the individuals in the process of creating the next generation. In many cases, the loss is temporary as these individuals (or better individuals) will be re-introduced into the population in a future generation.

However, if we want to guarantee that the best individual(s) always make it to the next generation, we can apply the optional elitism strategy. This means that the top  $n$  individuals ( $n$  being a small, predefined parameter, in our case 5) are duplicated into the next generation before we fill the rest of the available spots with offspring that are created using selection, crossover, and mutation. The elite individuals that were duplicated are still eligible for the selection process so they can still be used as the parents of new individuals.

Elitism is made possible in our code thanks to the function *eaSimpleWithElitism*, which is a modification of the function *eaSimple* present in the **Deap** framework.

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<sup>12</sup><https://content.wolfram.com/uploads/sites/13/2018/02/09-2-2.pdf>

## 3 — CNN from Scratch

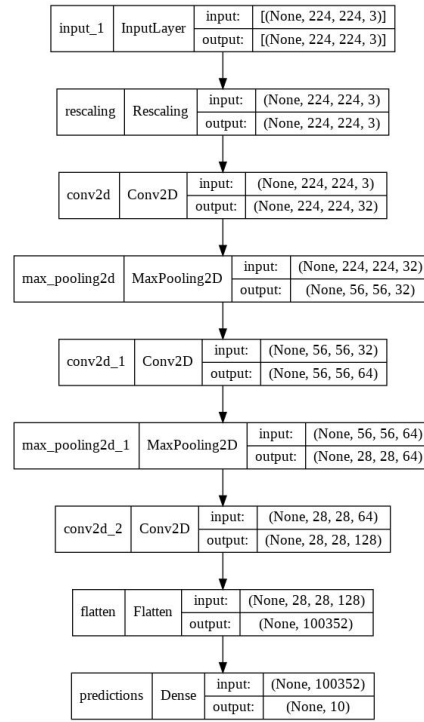
This chapter shows the results of the training of several custom architecture, that have been defined in order to solve the classification task. Starting from a very simple model, we start to analyze how to improve it and what modifications to apply in order to improve performance, taking into account mainly the accuracy on the validation test, but also considering other metrics like training time or number of parameters. The overall strategy is the following:

- test of different custom architectures defined from scratch and selection of the most promising one
- analysis of the level of fitting, try of different techniques to fight possible underfitting/overfitting
- evaluation of performance with the addition of a Batch Normalization layer
- hyperparameters optimization on the best model so far using a Genetic Algorithm(see Section 2.6)

The objective of the presented procedure is not the total exploration and exploitation of the search space, but it aims at finding good results in a reasonable time exploiting an ad-hoc heuristic search. The tested models are the following:<sup>13</sup>

### 3.1 Standard CNN

The first experiment has been conducted using a customized standard CNN that exploits Convolutional Layers and max Pooling to process input images. To start, we defined a very simple model, whose structure is reported in the image 2.



**Figure 2:** Customized Standard CNN Architecture

<sup>13</sup>all the models have been tested using data augmentation due to the lack of training samples. Some experiments without augmentation led to very quick overfitting and poor results, thus they have not been reported since considered as not interesting

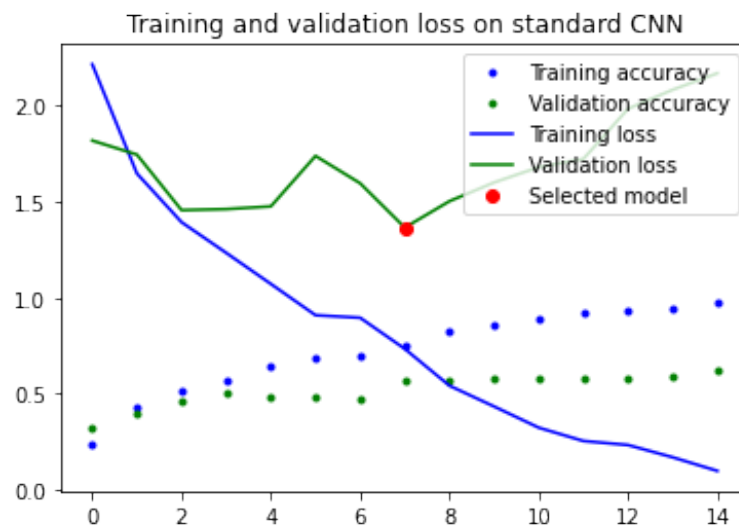
This model is trained using these default hyperparameters:

- optimizer: *ADAM*
- dropout rate :0.0
- learning rate: optimizer's default
- batch size: 128
- learning rate decay: none

In particular, we set a large value for batch size both because our main goal is to maximize the accuracy and also (as presented in the Introduction) to face off with the great variability of paintings with very dissimilar style but belonging to the same author. In this way, we increase the probability of a batch to be "complete", thus being representative of this variability.

The results obtained are the following:

StandardCNN				
Epoch stopped	Validation Accuracy	Testing Accuracy	Validation Loss	Testing Loss
15	0.5730	???	1,3656	???



**Figure 3:** Standard CNN: results

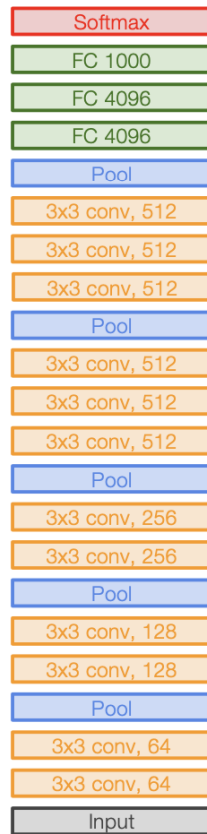
## 4 — Pre-Trained Models

This section describes the results obtained using different pre-trained architecture and strategies<sup>14</sup>. The pre-trained networks here tested are:

- VGG16
- ResNet50V2
- ResNet101V2
- InceptionV3

### 4.1 VGG16

VGG16<sup>4</sup> is a convolutional neural network model proposed by Simonyan et al., with several 3x3 convolutional layers in cascade occasionally interleaved with 2x2 max-pooling layers forming the so called *blocks*. Developed for the ILSVRC2014 challenge, it was able to achieve a top-5 accuracy of 92.7 on ImageNet.



**Figure 4:** VGG16 Architecture

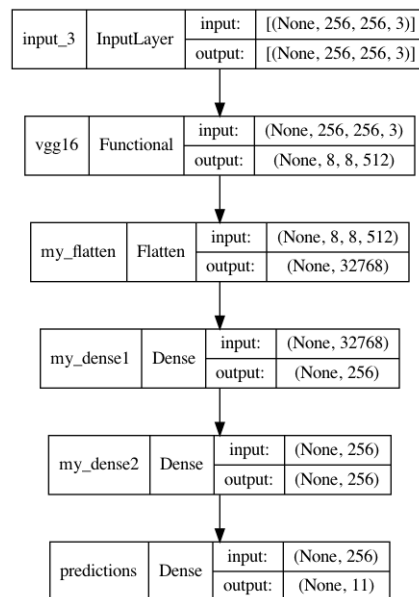
#### 4.1.1 Test 1: Classical VGG16 (Feature Extraction)

The original VGG16 comes with a couple of 4096 FC layers followed by 1000 softmax neurons, which is alright for ImageNet but definitely oversized for our purpose. Hence, the convolutional

---

<sup>14</sup>The data augmentation strategy is always used since we have very little data

base is left as it is, and the fully-connected block is replaced by the a shrunk version with only 256 neurons per layer, followed by our prediction layer made of 11 neurons<sup>5</sup>.

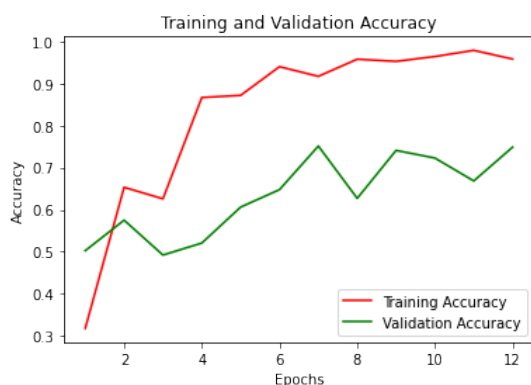


**Figure 5:** Our Feature Extraction Network

The result obtained, using RMSprop as optimizer, are:

Feature Extraction				
Epoch stopped	Validation Accuracy	Testing Accuracy	Validation Loss	Testing Loss
12	0.7409	0.7057	5.1959	5.7

The network begin to overfit very fast, hence some regularization methods are needed.



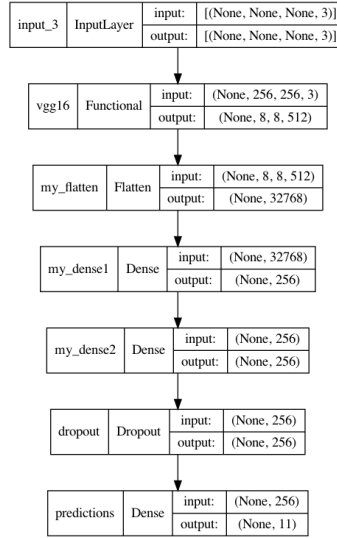
(a) Simple VGG16 Feature Extraction Accuracy



(b) Simple VGG16 Feature Extraction Loss

#### 4.1.2 Test 2: Adding Dropout to Test 1

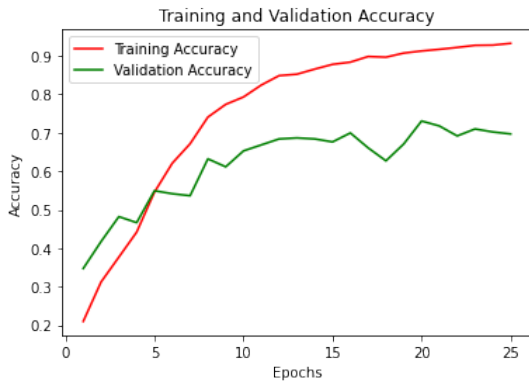
We have two possible positions to use the dropout layer in our network and they are after each 256-dense layer, but we decided to use just one layer at the end of the second 256-Dense layer (*my\_dense1*) as shown in Figure7. We didn't use a dropout layer between the two 256-dense layers, since this type of architecture led to worst performance, this mainly because we would have less units to fully train our topic-specific network.



**Figure 7:** Our Feature Extraction Network + Dropout

The result obtained, using RMSprop as optimizer, are:

Feature Extraction w/ dropout				
Epoch stopped	Validation Accuracy	Testing Accuracy	Validation Loss	Testing Loss
25	0.7306	0.7195	3.0616	3.0035



(a) Test 2 Accuracy



(b) Test 2 Loss

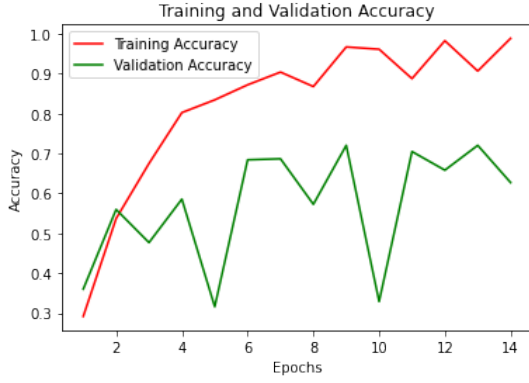
As expected, dropout mitigated the magnitude of overfitting, however our network perform slightly worst (now the validation accuracy is 0.73 and before was 0.74) than without the dropout layer.

#### 4.1.3 Test 3: Finetuning One Convolutional Layer

Using the model defined in test 1, the 3rd Conv2D layer in the 5th block is un-frozen and the network is trained. The result obtained using RMSprop as optimizer are the following:

Finetuning one convolutional layer				
Epoch stopped	Validation Accuracy	Testing Accuracy	Validation Loss	Testing Loss
28	0.7202	0.6253	2.3687	8.1807





(a) Test 3 Accuracy



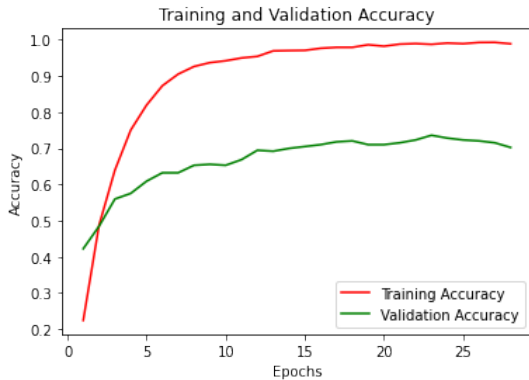
(b) Test 3 Loss

Again the performance are worst than in test one, but looking at the graphs it can be seen that not only our network overfitted very fast, but it forms also few fang-shaped changes in direction.

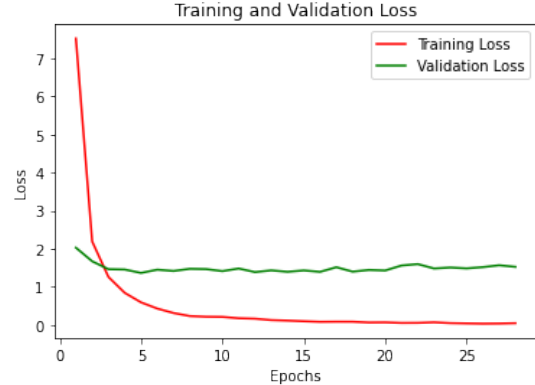
#### 4.1.4 Test 4: test 3 with dropout and different optimizer

To overcome the previous problems, the overfitting and the strange shape behavior, in this test we opt to use **Adam** as an optimizer, changing its default learning rate (i.e., 0.001) to 0.0001 in order to slowly learn and hoping to have a smoother accuracy and loss functions.

Finetuning one conv layer w/ dropout and Adam				
Epoch stopped	Validation Accuracy	Testing Accuracy	Validation Loss	Testing Loss
28	0.7358	0.7218	0.9871	1.1792



(a) Test 4 Accuracy



(b) Test 4 Loss

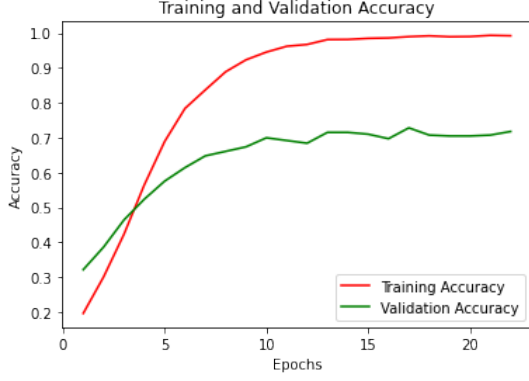
Adding dropout and a different optimizer with a little learning rate, we finally obtained what we were aiming. Anyway, the result is now comparable to the test 1, however now we are using a more complex network which is not good if the results are the same.

However we can still do something, the training accuracy increases very rapidly even if dropout is applied. To decrease this effect in test 6 we use weight regularization techniques.

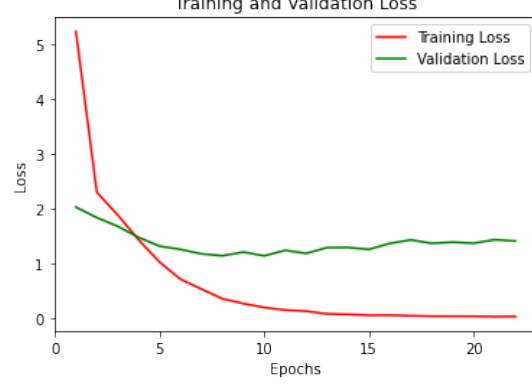
#### 4.1.5 Test 5: Finetuning Two Convolutional Layers

This test is basically test 4, but finetuning the last two convolutional layers of VGG16.

Finetuning two convolutional layers				
Epoch stopped	Validation Accuracy	Testing Accuracy	Validation Loss	Testing Loss
22	0.7280	0.7379	1.4253	1.2589



(a) Test 5 Accuracy



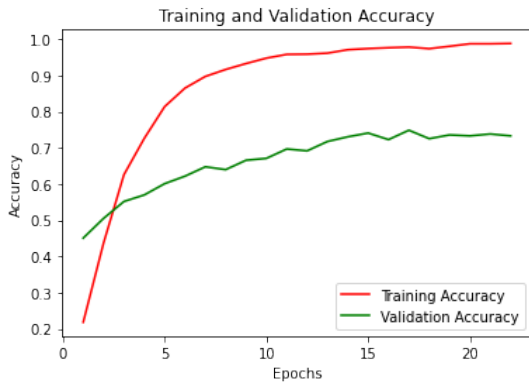
(b) Test 5 Loss

Following the approach used in the previous test we obtained also here more smoothed graphs, but the performance is little lower than before. The problem here is that we are propagating to the second layer in block 5 gradients that are not really improvements of the ones set by the *imagenet* default configuration.

#### 4.1.6 Test 6: Finetuning One Convolutional Layer and Weights Regularization

In this paragraph we exploited the conclusion mentioned in test 4, introducing here *L1\_L2 weight regularization* on the one convolutional layer finetuned network.

Finetuning one conv layer and weights regularization				
Epoch stopped	Validation Accuracy	Testing Accuracy	Validation Loss	Testing Loss
22	0.7487	0.7471	21.5590	9.8097



(a) Test 6 Accuracy



(b) Test 6 Loss

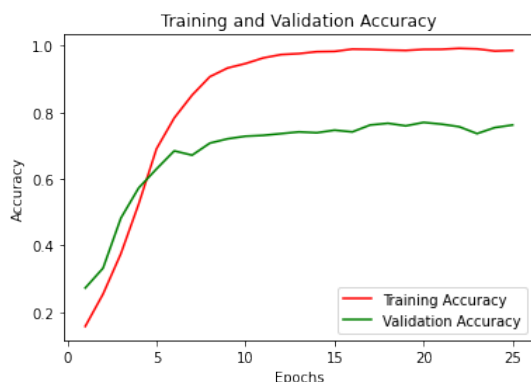
The shapes of the graphs are more or less the same of the two had in test 4, the only things that changes are the values of the losses which are here higher and more curvilinear due to the weight regularization approach used. Anyway, this heavy regularization helped us to surpass the result in test 1, not by much but it is an improvement that, maybe, can be exploited

finetuning more. Following this lead, in the next paragraph we use weight regularization with two convolutional layers finetuned.

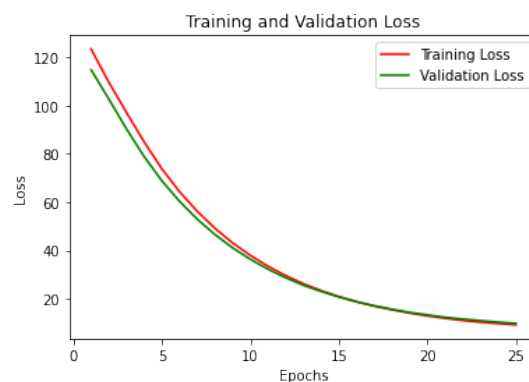
#### 4.1.7 Test 7: Finetuning Two Convolutional Layers and Weights Regularization

Following the good result obtained in test 6, in this paragraph we add, to the network used in test 5, *L1\_L2 weight regularization*.

Finetuning two conv layers and weights regularization				
Epoch stopped	Validation Accuracy	Testing Accuracy	Validation Loss	Testing Loss
25	0.7694	0.7494	15.6742	9.8097



(a) Test 7 Accuracy



(b) Test 7 Loss

Even if the increase the flexibility of our network finetuning more layers, as hoped, we achieved a better accuracy which is also the best obtained so far.

#### 4.1.8 Test 8: Genetic Algorithm for Hyper-parameters and Architecture Optimization

### 4.2 ResNet50V2

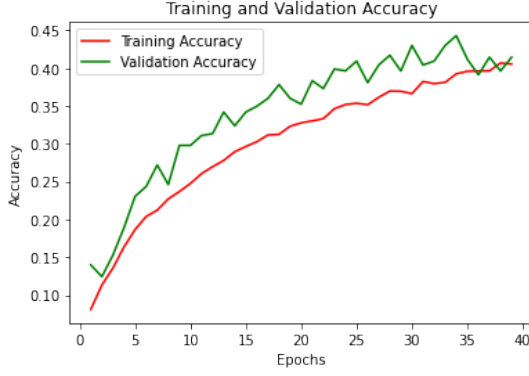
ResNet stands for Residual Network. It is an innovative neural network that was first introduced by Kaiming He, Xiangyu Zhang, Shaoqing Ren, and Jian Sun in their 2015 computer vision research paper titled "Deep Residual Learning for Image Recognition". This model was immensely successful, as can be ascertained from the fact that its ensemble won the top position at the ILSVRC 2015 classification competition with an error of only 3.57%. Additionally, it also came first in the ImageNet detection, ImageNet localization, COCO detection, and COCO segmentation in the ILSVRC & COCO competitions of 2015.

#### 4.2.1 Test 1: Classical ResNet50V2 (Feature Extraction)

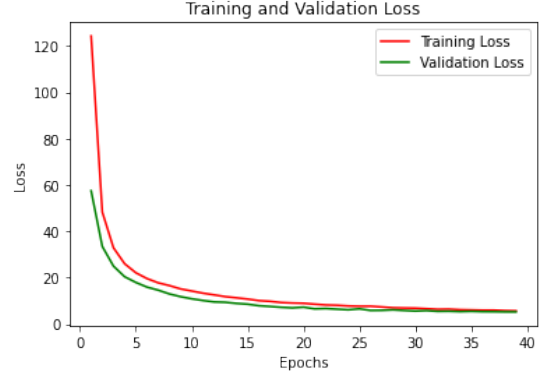
The original ResNet50 comes with a GlobalAveragePooling2D and a prediction layer soon after. In this test we used the same approach, resizing the prediction layer to the number of classes we have.

The results obtained after training are:

Feature Extraction				
Epoch stopped	Validation Accuracy	Testing Accuracy	Validation Loss	Testing Loss
39	0.4430	0.3241	5.3733	6.8513



(a) Simple ResNet50V2 Feature Extraction Accuracy



(b) Simple ResNet50V2 Feature Extraction Loss

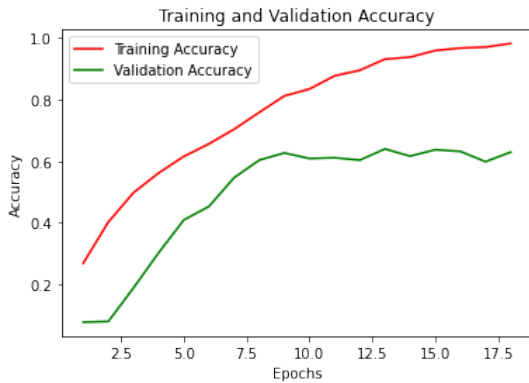
The network poorly performed compared with the results obtained using VGG16. This can be explained since the *imagenet dataset* is not specialized in distinguish paintings' artists, thus the frozen weights of our networks are very far to be the ones we need. To overcome this problem we can fine tune or add more layers after the *GlobalAveragePooling2D* layer.

Another problem can be noticed looking at the accuracy plot, the training accuracy is less than the validation accuracy. To understand this issue, it is important to consider the difference of the number of images in the training and validation set. The validation set is made of about few hundreds of pictures against the few thousands of the training set, the latter may involve, with greater probability, paintings being part of a different period of the artists (e.g., the example shown in Image1), which can lead to a worst accuracy.

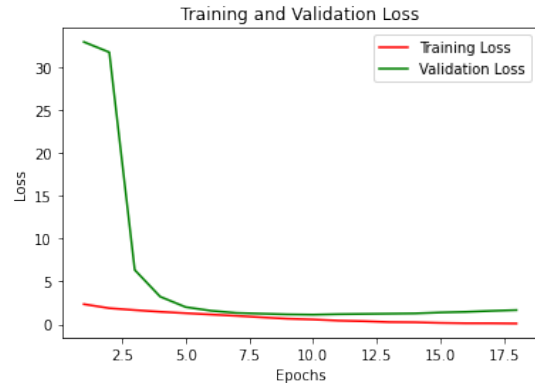
#### 4.2.2 Test 2: Finetuning 1 block

ResNet50V2 is made of multiple big blocks (i.e., 5) so called *conv* in the model. These blocks are made of sub-blocks connected by each other by an add layer which connects the processed input (e.g., processed by Conv2D, Padding, Pooling, BatchNormalization) and the residual input<sup>15</sup>. We finetuned considering this under-blocks, hence in this paragraph we finetuned the *conv5\_block3*.

Finetuning 1 block				
Epoch stopped	Validation Accuracy	Testing Accuracy	Validation Loss	Testing Loss
18	0.6399	0.5793	1.2453	1.7916



(a) ResNet50V2 Test 2 Accuracy



(b) ResNet50V2 Test 2 Loss

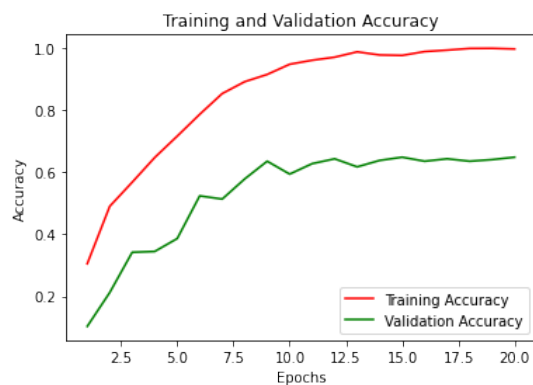
<sup>15</sup>Sometimes the residual is downsampled using a max pooling layer, this is done in order to match the actual size of the feature map obtained at that level of the network.

As expected, the network has definitely reach a more satisfactory result. Also the behaviors of the curves now is more reasonable.

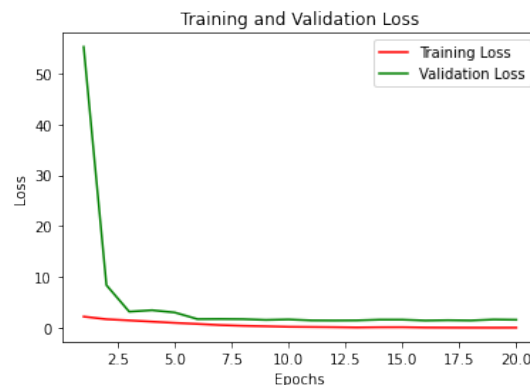
### 4.2.3 Test 3: Finetuning 2 blocks

Since finetuning a block led to better results, the next step we did was to tuning also the previous sub-block, thus starting to tuning from *conv5\_block2*.

Finetuning 2 blocks				
Epoch stopped	Validation Accuracy	Testing Accuracy	Validation Loss	Testing Loss
20	0.6477	0.6092	1.5918	1.7734



(a) ResNet50V2 Test 3 Accuracy



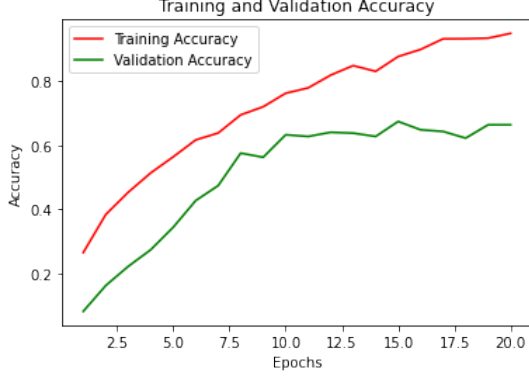
(b) ResNet50V2 Test 3 Loss

This results proves that our idea of finetuning was good since the network improved a little. Anyway, finetuning more can be risky since the computational power and time needed can be very high and becoming unbearable to the limits imposed by Colab. Thus, we tried to improved (without satisfaction, as the nexts to paragraphs describe) our network adding dense layers between our *GlobalAveragePooling2D* and our prediction layer, following the structure used in the VGG16 study.

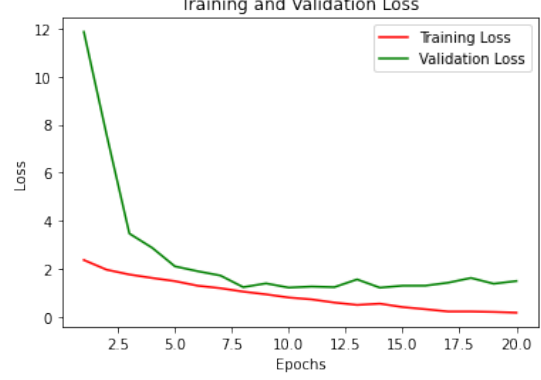
### 4.2.4 Test 4: Finetuning with One Block and Adding Two Dense layers

Here we took the same approach of test 3, but adding two dense layers followed by a dropout layer (as used in figure7).

Finetuning one block and dense layers				
Epoch stopped	Validation Accuracy	Testing Accuracy	Validation Loss	Testing Loss
20	0.6736	0.6368	1.6655	1.7734



(a) ResNet50V2 Test 4 Accuracy



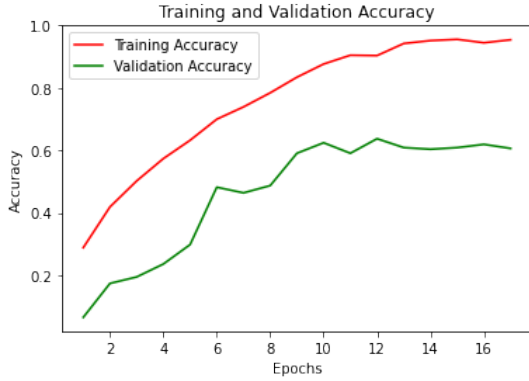
(b) ResNet50V2 Test 4 Loss

Enlarging in this way the network resulted in better performance. This can be because the dense layers before the softmax can better exploit the information of the *GlobalAveragePooling2D*, keeping the problem for more time to a higher dimensionally than the one used for prediction.

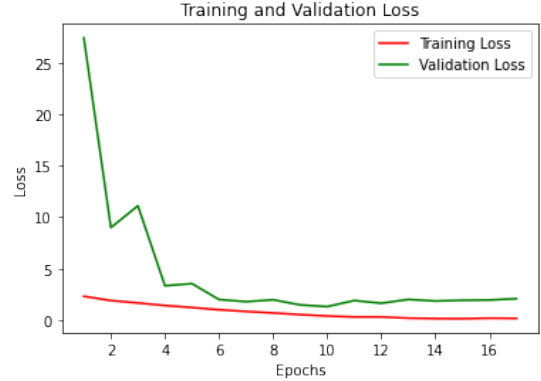
#### 4.2.5 Test 5: Finetuning with Two Blocks and Adding Two Dense layers

In this paragraph we did the same thing done in test 4, but finetuning two blocks as in test 3.

Finetuning two blocks and dense layers				
Epoch stopped	Validation Accuracy	Testing Accuracy	Validation Loss	Testing Loss
17	0.6373	0.5954	1.6404	2.1089



(a) ResNet50V2 Test 5 Accuracy



(b) ResNet50V2 Test 5 Loss

Unfortunately, finetuning this much with these additional layers led to worst performance. Perhaps, this is due to backpropagation, which changed the weights (which are 9M, against the 14M that are not trainable) to a not-optimal solution. Anyway, this result is also too poor compared to test 3, which without having the newly added dense layers still perform better.

### 4.3 ResNet101V2

ResNet101 was implemented as one of the model used in the ensemble method cited in the ResNet50V2 introduction. ResNet101 is an extension of ResNet50 that goes deeper reaching 101 layers in depth, simply adding more convolutional 'sub'-blocks in the blocks<sup>16</sup>. ResNet101

<sup>16</sup>Referring to the notation used before, we intend as a sub-block a block between two adds.

proved to be slightly more accurate than ResNet50.<sup>17</sup>

#### **4.3.1 Test 1: Classical ResNet101V2**

#### **4.3.2 Test 2: Completely Newly Output Layers Architecture**

#### **4.3.3 Test 3: Fine Tuning with One Layer**

#### **4.3.4 Test 4: Fine Tuning with Two Layers**

### **4.4 InceptionV3**

#### **4.4.1 Test 1: Classical ResNet101V2 with 50 classes**

#### **4.4.2 Test 2: Completely Newly Output Layers Architecture**

#### **4.4.3 Test 3: Fine Tuning with One Layer**

#### **4.4.4 Test 4: Fine Tuning with Two Layers**

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<sup>17</sup>It is not only additional network used in the ensemble method, an extension of it was also used, ResNet152. This model is the more accurate of the three, but because of its size we decide to limit our studies to the 101-layers version.