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Automatic underwater fish species classification with limited data using few-shot learning

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21 Abstract:

22 Underwater cameras are widely used to monitor marine biodiversity, and
23 the trend is increasing due to the availability of cheap action cameras.
24 The main bottleneck of video methods now resides in the manual
25 processing of images, a time-consuming task requiring trained experts.
26 Recently, several solutions based on Deep Learning (DL) have been
27 proposed to automatically process underwater videos. The main limitation
28 of such algorithms is that they require thousands of annotated images in
29 order to learn to discriminate classes (here species). This limitation
30 implies two issues: 1) the annotation of hundreds of common species
31 requires a lot of efforts 2) many species are too rare to gather enough
32 data to train a classic DL algorithm. Here, we propose to explore how
33 few-shot learning (FSL), an emerging research field, could overcome DL
34 limitations. Few-shot learning is based on the principle of training a Deep
35 Learning algorithm on “how to learn a new classification problem with
36 only few images”. In our case-study, we assess the robustness of FSL to
37 discriminate 20 coral reef fish species with a range of training databases
38 from 1 image per class to 30 images per class, and compare FSL to a
39 classic DL approach with thousands of images per class. We found that
40 FSL outperform classic DL approach in situations where annotated images
41 are limited, yet still providing good classification accuracy.

42

43 Keywords: few-shot learning, Deep learning, video, marine biodiversity

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Introduction

The world's ecosystems have entered an era of anthropogenic defaunation where human activities have triggered global decline in animal abundance, species range contraction and a new wave of species extinction [1]. This global change is threatening ecosystem services worldwide hence the stability of our food systems, economies, and health. Defaunation is more advanced in terrestrial and freshwater ecosystems than in the marine environment where it started centuries later. However, the pace of defaunation is accelerating in oceans mostly due to the advent of industrial fishing since a century ago [2]. Given this context of global changes rapidly affecting fish communities, it is imperative to monitor fish biodiversity over time, on a large scale and using non-destructive methods.

Fish biodiversity surveys in the marine environment are typically performed by divers. Although dive visual censuses provide a great deal of information on some shallow habitats, there are many limitations. First, divers are limited by depth and can hardly perform long dives to count fish below 30 m, ignoring mesophotic habitats and deeper ecosystems. Second, divers are limited by time and generally focus their 2-4 dives per day in the most speciose hard-substrate habitats, and ignore less rich and often immense adjacent soft-bottom habitats. Third, dive surveys provide data at a slow rate so that the compilation of global fish biodiversity database takes decades of efforts by multiple teams of highly skilled taxonomic divers (e.g. [3], [4]). This is a major restriction to the necessary temporal monitoring of global marine ecosystems, although a few time series exist in some countries¹ [5].

¹ AIMS, Long-Term Monitoring Program: Visual Census Fish Data (Great Barrier Reef)
<https://apps.aims.gov.au/metadata/view/5be0b340-4ade-11dc-8f56-00008a07204e>

Underwater videos (UV) are increasingly used [6] to overcome the limitations of diver-based surveys to quickly collect large amounts of data. For instance, more than 15,000 video stations were deployed in 58 countries in just three years for the first global assessment of the conservation status of reef sharks [7]. Furthermore, underwater video surveys can be performed in many habitats, with some example in shallow reefs [8], sandy lagoons [9], deep sea [10], and even in the pelagic ecosystem [11]. Deploying underwater video stations does not require expert taxonomists and is now quite inexpensive with the improvement of cheap action cameras since a few years. The bottleneck to analyse this data now resides in the manual processing of the videos. Indeed, manually extracting fish biodiversity and abundance data from raw videos requires unsustainable workload by highly trained taxonomic experts. Although this annotation work can be improved through citizen science [12]–[14], such time-consuming and expensive task cannot match the increasing size of datasets, up to 20,000 hours of videos for global surveys [7] and the necessary monitoring of global oceans over time.

As the demand for automatic methods to analyze underwater videos is rising, the latest generation of deep learning algorithms (DL), and in particular convolutional neural networks (CNNs) are increasingly used for species identification [14]–[17] and fish detection [18]–[21]. However, these algorithms require a large dataset of annotated images (thumbnails hereafter) in order to train a robust model, able to provide satisfying results. Therefore, this method still requires collecting an important image dataset manually annotated by experts. This is especially problematic in highly diverse faunas such as coral reef fish that encompass nearly 6500 species worldwide [22]. Furthermore, a universal pattern in species distribution, including fish communities, is that both rare and common species are found in every community, with the fraction of rare species

more important in rich ecosystems, such as coral reefs [23], [24]. It is therefore almost impossible to gather enough thumbnails of rare species to efficiently train a deep neural network in a “classic” way, which requires thousands of images per species [25]–[27]

There are two ways to tackle this problem of lack of data. The first one consists of directly addressing the data itself, through data augmentation [28]–[30]. The second option is to change the classification algorithm. Few-shots learning (FSL) algorithms [31], [32] are designed to compute a classification task (query, noted Q) with only a few thumbnails to train (Support Sets, noted SS), and it has been increasingly studied since 2017 [33]. Few-shots learning methods are divided into three main approaches. Metric-based methods are embedding both queries (Q) and support sets (SS), before assigning to the query a class, according to distances computed between Q and SS ([34]–[36]). The second approach consists of 1) training a model on a large database, and 2) adapt this model to a new task with few examples, while not forgetting the concepts learned previously [37], [38]. Finally, optimization-based methods are designed to adapt quickly to new tasks, hence able to learn a classification task with few examples [33], [39], [40]. Optimization-based algorithms showed promising results in deep learning few-shot classification [33], [41], [42]. Such methods propose to pre-train (or “meta-train”) a model with existing databases (e.g. MiniImageNet [43] , Ominglot [44]) on different tasks so it can adapt easily to a new one. For object identification, a task is defined by the classes the model has to discriminate. Once this model, called “meta-model” has been trained, it can then be tuned to operate on a new task with a very limited dataset, usually only 1-5 thumbnails per class.

In this study we propose to compare the efficiency of optimization-based few-shot learning and standard large dataset deep-learning methods to identify coral reef fish species on images. More specifically, we aim to determine how well a classic deep learning architecture trained with thousands of images [and the benefit of data augmentation](#) (hereafter DL) and FSL algorithms perform in situations where training thumbnail dataset is large or limited. To achieve this, we first trained a classic DL architecture built for image classification [45] on a large dataset of 69,169 thumbnails, and on a more limited dataset of 6,320 thumbnails for 20 coral reef fish species. Then, we trained a few-shots, optimization-based learning algorithm [39] on the exact same training datasets while varying the number of shots from 1 to 30. Finally, we compared the capacity of DL and FSL models to correctly identify species on an independent thumbnail dataset, and modelled the asymptotic relationship between classification accuracy and the number of thumbnails in the training datasets for both classic DL and FSL algorithms

Material and methods

Thumbnail datasets

We used three fish thumbnail datasets (*T0*, *T1*, and *T2*) extracted from 175 underwater videos recorded on reefs around Mayotte Island (Western Indian Ocean) using GoPro Hero 3+ and GoPro hero 4+ cameras with a resolution of 1920x1080 pixels. A thumbnail is defined as an image containing a single labelled fish belonging to one of the 20 most common fish species in the videos, and representing a broad range of sizes, colors, body orientations, and background (Supp. Fig. 1, Supp. Fig. 2).

T0 is composed of 69,169 thumbnails extracted from 130 videos, with a range of 1,134 to 7,345 thumbnails per species (Table 1). *T1* is composed of 6,320 thumbnails extracted from 20 videos with 40-1,436 images per species whereas *T2* is composed of 13,232 thumbnails extracted from 25 videos with 55-3,896 images per species. Thumbnails size originally ranged from 55x55 pixels to 500x450 pixels, but were resized to 84x84 pixels before being processed through FS and DL algorithms.

The datasets *T1* and *T2* correspond to two real scenarii where videos were recorded during two trips in the field of a week each.

The three thumbnails datasets are fully independent, as they were extracted from videos recorded at different sites, with different conditions (weather, lighting, depth, time of the day, seascape) and on different days.

To train our DL architecture, we applied data augmentation to *T0* and *T1*. For each natural thumbnails in *T0* and *T1*, we created 9 thumbnails through contrast augmentation or diminution, and horizontal flip. We then obtained augmented datasets composed of 691,690 (AT0) and 63,200

(AT1) images respectively [Supp. Table 1](#). Further details on thumbnail datasets and data augmentation are given in [46].

Table 1: Number of natural thumbnails extracted from the videos to build our three datasets

Family	Species	Training dataset $T0$	Training dataset $T1$	Test dataset $T2$
Acanthuridae	<i>Acanthurus leucosternon</i>	3,259	235	491
Acanthuridae	<i>Acanthurus lineatus</i>	1,008	114	864
Acanthuridae	<i>Naso brevirostris</i>	1,134	539	1932
Acanthuridae	<i>Naso elegans</i>	7,345	1,435	3,896
Acanthuridae	<i>Zebrasoma scopas</i>	4,970	48	579
Chaetodontidae	<i>Chaetodon auriga</i>	2,134	737	502
Chaetodontidae	<i>Chaetodon guttatissimus</i>	1,182	221	68
Chaetodontidae	<i>Chaetodon trifascialis</i>	5,234	41	630
Chaetodontidae	<i>Chaetodon trifasciatus</i>	4,421	71	82
Labridae	<i>Gomphosus caeruleus</i>	3,131	57	173
Labridae	<i>Halichoeres</i>	3,192	40	287

	<i>hortulanus</i>			
Labridae	<i>Thalassoma hardwicke</i>	4,951	181	275
Lethrinidae	<i>Monotaxis grandoculis</i>	3,893	797	1,422
Monacanthidae	<i>Oxymonacanthus longirostris</i>	2,553	54	55
Pomacentridae	<i>Abudefduf vaigiensis</i>	5,124	376	216
Pomacentridae	<i>Amblyglyphidodon indicus</i>	1,188	636	1,310
Pomacentridae	<i>Chromis opercularis</i>	1,525	81	93
Pomacentridae	<i>Chromis ternatensis</i>	3,640	300	156
Pomacentridae	<i>Pomacentrus sulfureus</i>	5,409	270	142
Zanclidae	<i>Zanclus cornutus</i>	3,876	86	59
TOTAL		69,169	6,320	13,232

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188 Experimental design

189 To compare classic deep-learning and few-shot algorithms in situations of
 190 large or small thumbnail datasets, [we led five experiments using datasets](#)
 191 [T0, T1, T2, AT0 and AT1 described in Supp. Table 1 :](#)

- 1) We trained a classic DL algorithms architecture with our biggest dataset *A70* as a baseline for the DL accuracy;
- 2) We trained the same DL architecture with the same hyper-parameters (e.g. model architecture and training process) but on a much more limited dataset (*A71*). Hyper-parameters are the parameters defining the architecture (number of layers, number and size of convolutions, connections between layers) and the training process of a Deep Model (learning rate, neurone activation, back-propagation computation).;
- 3) We trained the same DL architecture with limited datasets obtained by subsampling *T0* to 250 and 500 images per class (here after "species" when we are referring to our experiments), corresponding to 2500 and 5000 thumbnails in *A70*;
- 4) We pre-trained a FSL architecture on the 64 training classes of MiniImageNet (Supp. Fig. 3) and used *T0* to build support sets (*SS*) with 1, 5, 15 and 30 thumbnails for each fish species;
- 5) We pre-trained the same FSL architecture on MiniImageNet and used the more limited *T1* dataset to build support sets with 1, 5, 15 and 30 images per species.

We used ResNet 100 [45] as our classic deep-learning algorithm. Resnet is a convolutional neural network (CNN), a DL architecture which is able to both extract features from images and classify these images thanks to those features [47]. In order for a CNN to build an image classification model, the architecture is fed a large dataset, composed of pairs of labels and images. Using this dataset, the algorithms change their inner parameters in order to minimize the classification error, through a process called back-propagation. The ResNet architecture achieved the best results on ImageNet Large Scale Visual Recognition Competition (ILSVRC [43]) in 2015, considered the most challenging image

classification competition. It is still one of the best classification algorithms, while being easy to use and implement.

For the few-shot implementation, we used the Reptile algorithm [39]. Few-shot learning algorithms are specific DL algorithms, whose goal is to be able to fit a model with very few training images. The Reptile algorithm is based on the well-known MAML architecture [33], and more precisely on the first-order version of MAML [48]. The Reptile algorithm is based on the division of the training dataset into a number of tasks T_i , a task being a learning problem. Through repetitively changing the task during the first training phase (known as meta-training), this algorithm produces a quick learner, i.e. a learner that can quickly adapt to a new task with a small number of examples.

Here, the few-shot algorithms were tested on a classic *n*-ways *k*-shots procedure, *n* being the number of classes per support set, and *k* the number of images per class in the support set. For instance, a 5-ways 1-shots consists of training 5 classes with support sets composed of 1 image per classes (e.g. species). We set $n=5$ [34], [36], [40], [42], [49] and allowed *k* to vary between 1 shot and 30 shots for both experiments 4 and 5. We did not use data augmentation for FSL experiments for several reasons. First, the goal of FSL is to adapt quickly with a very limited number of images. Second, to have similar settings for method comparison. There were no data-augmentation in the original paper, so we reproduced that. It also allowed us to compare our results with those obtained on benchmarks. Third, the reason behind the use of raw data instead of augmented data in few-shot learning paper is that with very few training samples and few conditions, the risk of overfitting by using the same image modified multiple times is far greater than in classic approaches with important datasets with many conditions.

252

253 Model comparison

254 All the DL and FSL models were tested on the independent *T2* dataset.

255 First, we compared the results of experiments 1 and 2 in order to
256 estimate the decrease in performance of a classic ResNet DL architecture
257 when trained on a large dataset *AT0* (i.e. between 11,340 and 73,450
258 images per species after data augmentation, with an average of 3458
259 natural thumbnails per species) or trained on a more limited dataset *AT1*
260 (i.e. between 400 and 14,360 images per species after data
261 augmentation, with an average of 315 natural thumbnails per species).

262 Second, we compared the results of experiments 1 and 4 in order to
263 evaluate if the ResNet architecture outperforms the Reptile architecture in
264 a real-case situation where thumbnail dataset is large (*T0 and AT0*).

265 Finally, we compared the results of experiments 2 and 5 to determine
266 whether and to which extent a Reptile model performs better than a
267 ResNet model in a real-case situation where thumbnail dataset is limited
268 (*T1 and AT1*).

269

270

271 In order to better evaluate the performance of ResNet and Reptile
272 algorithms, we also modelled the relationship between model accuracy
273 and the number of thumbnails used to train the models. To achieve this,
274 we fitted the following asymptotic function to the results of experiments
275 1, 3 and 4 (obtained through training DL and FSL architectures on
276 datasets of various size obtained from *AT0* and *T0*):

277

$$278 \text{ Accuracy} = \text{Accuracy}_{\infty} \cdot (1 - \exp(-R \cdot N_{\text{image}})) \quad (\text{eq.1})$$

where $Accuracy_{\infty}$ is the asymptotic model accuracy when the number of thumbnails N_{image} is infinite, and R is the rate at which the asymptote is reached.

Equation 1 was fitted by non-linear mixed-effect modelling (NLME [50]) using species as a random effect. This method is widely used for fitting asymptotic processes. It allows estimating and comparing asymptotic accuracies of both FSL and DL algorithms, and the number of image to reach these asymptotic accuracies. The number of images required to reach the asymptotic accuracy was calculated as the number of images corresponding to an accuracy of 0.99 times the asymptotic value, meaning the asymptote was reached within 1%.

Results

The deep ResNet model trained on the large *AT0* dataset (3458 natural thumbnails in average per species) during the first experiment obtained a mean accuracy (i.e. percentage of correct classification) of 78.00% (standard deviation (SD) of 15.16%) on *T2* test-dataset (Table 2). With this model, accuracy varied among species between 54.14% (*Naso brevirostris*) and 99.07% (*Abudefduf vagiensis*). The same ResNet DL model trained on smaller *AT1* (315 natural thumbnails in average per species) during the second experiment showed highly degraded performance with a mean accuracy of only 42.21% (SD=24.95%). Among species variation ranged with this model from only 3.49% (*Chaetodon trifascialis*) to 85.86% (*Chaetodon auriga*).

The few-shot Reptile architecture trained on limited *T1* dataset during our fifth experiment obtained a mean accuracy of 32.04% for the 1-shot learning (SD= 12.70%) and 51.77% mean accuracy for the 30-shots learning (SD= 18.96%) (Table 2,). In this scenario of limited *T1* training dataset, the few-shot Reptile algorithm nearly equalled the ResNet DL model with only 5 shots (41.47% accuracy for 5-shots learning on *T1* vs 42.21% for DL on *AT1*), and performed better beyond 10 shots (45.92%

311 of accuracy on *T2* with 10-shots learning). A pairwise proportion test
312 showed a p-value <0.0001 , assessing that FSL was significantly better
313 than DL in this scenario beyond 10 shots (Supp. Table 3) accuracy of
314 Reptile models had a standard deviation from 12.70% with one-shot
315 learning, to 18.96% with 30-shots learning, indicating important variation
316 in accuracy among species. However, this standard deviation was smaller
317 than that of the ResNet algorithm trained on the same *AT1* limited
318 dataset (24.95%).

Figure 1: Relationship between the number of natural thumbnails per species for training and the accuracy of deep-learning and few-shot learning models. Non-linear mixed effects asymptotic model fit for (a) DL architecture at the fixed-effect level, and for FSL architecture at (b) the fixed-effect level and (c) the random-effect level. Grey areas represent 95% CI in fixed-effects estimates. Dotted lines represent the NLME estimate of the number of images per species required to reach the 99% asymptote value. We obtained similar magnitude with the 95% asymptote value, reached with 750 images for DL and 5 with FSL.

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330

331 The same few-shot Reptile architecture trained on subsets of T0 during
332 the fourth experiment obtained even better results than when trained on
333 T1, with a mean accuracy on T2 of 34.57% for 1-shot, 50.23% for 5-
334 shots, and up to 64.92% for 30-shots (Table 2).

335 Mixed-effects modelling (NLME) of T0 and AT0 experimental data showed
336 a clear pattern of asymptotic increase of accuracy with the number of
337 natural thumbnails for both Resnet and Reptile architectures (Figure 1).

338 NLME models included significant species random effect for both DL and
339 FSL (Log-likelihood tests, $P < 0.0001$).

340 The fixed-effect asymptotic value of accuracy was higher for ResNet
341 model ($Accuracy_{\infty} = 77.34\%$, 95% CI: 71.26-83.41%) than for Reptile
342 model ($Accuracy_{\infty} = 60.87\%$, 95% CI: 54.48-67.26%), illustrating higher
343 classification power of ResNet over Reptile when large numbers of
344 thumbnails are available. However, the slope of the asymptotic model
345 was two-orders of magnitude higher for Reptile (0.707, 95% CI: 0.559-
346 0.854) than for ResNet architecture (0.0040, 95% CI: 0.0032-0.0048),
347 illustrating the high capacity of Reptile FSL algorithm to learn from only a
348 few images. NLME modelling further showed that average asymptotic
349 accuracy was reached with only 7 natural thumbnails per species for
350 Reptile architecture, compared to 1153 natural thumbnails per species for
351 ResNet, confirming the strong power of Reptile method in situation of
352 limited thumbnail training dataset. However, model random effects
353 showed that some variation existed among species. For Reptile
354 architecture, asymptotic accuracy values ranged from 38.09%
355 (*Amblyglyphidodon indicus*) to 89.78% (*Pomacentrus sulfureus*), and was
356 reached with 4 to 16 training images per species. For DL architecture,
357 species asymptotes varied from 62.72% (*Monotaxis grandoculis*) to

358 96.81% (*Abudefduf vaigiensis*), and could be reached with 786-1776
359 thumbnails per species (*Supp. Table 4*).

360

361 *Table 2: Accuracy of our ResNet deep-learning (DL) and Reptile few-shots learning (FSL) models trained on T0 or*
362 *T1 thumbnails datasets for different number of shots. Accuracy is the % correct classification of models on T2 test*
363 *dataset. DL models were trained from T0 and T1 after data augmentation (AT0 and AT1).*

364

Image per species (on average)	DL		FSL					
	T0	T1	T1			T0		
	3458	315	1 shot	5 shots	30 shots	1 shot	5 shots	30 shots
<i>Abudefduf vaigiensis</i>	99.07	69.91	16.08	11.39	11.38	47.67	70.9	86.35
<i>Acanthurus leucosternon</i>	86.15	44.67	25.51	30.71	38.74	19.23	28.8	42.66
<i>Acanthurus lineatus</i>	59.72	20.37	39.86	56.04	72.50	32.93	61.01	72.02
<i>Amblyglyphidodon indicus</i>	58.78	60.78	25.75	26.74	32.86	28.26	32.55	40.64
<i>Chaetodon auriga</i>	87.05	85.86	18.16	25.68	36.56	27.8	35.18	53.20
<i>Chaetodon guttatissimus</i>	85.50	44.12	33.58	44.21	58.26	29.61	51.18	79.29
<i>Chaetodon trifascialis</i>	90.00	3.49	29.02	25.48	28.44	27.14	43.17	63.51
<i>Chaetodon trifasciatus</i>	87.80	28.05	38.73	50.63	66.72	32.41	51.07	70.63

<i>Chromis opercularis</i>	61.29	9.68	44.01	61.81	62.94	45.34	68.28	81.50
<i>Chromis ternatensis</i>	59.61	55.77	18.91	24.94	35.07	35.4	55.44	67.22
<i>Gomphosus caeruleus</i>	75.72	20.81	26.01	38.99	58.74	28.96	39.16	54.22
<i>Halichoeres hortulanus</i>	82.93	17.07	31.82	44.81	57.01	28.94	41.35	58.87
<i>Monotaxis grandoculis</i>	57.10	53.37	32.03	41.52	50.64	32.8	45.85	59.13
<i>Naso brevirostris</i>	54.14	68.60	47.06	54.26	64.66	54.47	58.08	61.00
<i>Naso elegans</i>	93.24	79.43	34.54	43.11	52.17	28.47	33.71	44.36
<i>Oxymonacanthus longirostris</i>	96.43	14.54	39.29	53.48		42.15	65.44	
					66.26			84.86
<i>Pomacentrus sulfureus</i>	90.14	61.97	70.90	88.18	93.93	65.53	86.21	90.00
<i>Thalassoma hardwicke</i>	90.90	51.64	25.64	44.4	67.96	26.72	45.7	70.60
<i>Zanclus cornutus</i>	81.36	40.68	18.33	31.28	44.44	26.08	40.82	62.72
<i>Zebrasoma scopas</i>	63.04	13.30	25.56	31.81	36.05	31.42	50.69	55.70
MEAN	78.00	42.21	32.04	41.47	51.77	34.57	50.23	64.92
SD	15.16	24.95	12.70	16.93	18.96	11.14	14.75	14.55

Discussion

Our experiments demonstrated that few-shot learning methods based on Reptile architecture can be effectively used to drastically reduce the number of annotated images for underwater fish identification. Accuracy levels obtained with few-shot learning algorithm trained with only five training images are close to those of a standard Deep Learning architecture such as ResNet trained with 400-14350 images per species. Further, FSL architecture trained with 10 images outperformed a ResNet 100 architecture trained with at least 400 images per species. This is a very promising result in situations where many species need to be identified from models trained with a few images, a typical characteristic in marine biodiversity applications.

However, the important standard deviation among the different trained species (18.96 SD on 30-shots) showed that few-shot algorithms may not be robust enough to discriminate among similar species showing only subtle differences. Nevertheless, in our 2nd experiment, our ResNet model achieved an accuracy under 40% for all the species with fewer training images than 1140 (after data augmentation, i.e. 114 natural images), and only 7 species were identified with an accuracy greater than 45%. These species were represented with a range of 2700-14350 images during the training phase. [We also show better results with the model trained on T0 than the model trained on T1. As expected, increasing the number of images per shot rely on better performances as well as increasing the per species images variability. However, in real conditions, few-shot learning is to be used in a context where very few images per classes are at disposal. Therefore, the dataset T1 corresponded more to a real use case scenario.](#)

Thus, there is a trade-off to make between accuracy and robustness on one hand, and the cost of video annotation by experts on the other.

Modeling the accuracy of neural networks using NLME allowed to understand the number of images per species required for the Few-shot and Deep architecture to reach 99% of their maximum potential accuracy. In our case study, there was a 150-fold factor between the average number of images required for a Deep Learning architecture (1153 images) and for a Few-shot architecture (7) to reach asymptotic accuracy. However, it is important to note that these numbers could vary according to the number and complexity of classes fed to the deep classifier.

In this work we used a Reptile FSL architecture. As the field of few-shot learning is quickly improving, new methods are proposed at a fast rate. While Reptile obtained a mean accuracy of 61.98% on the MiniImageNet dataset (the most used benchmark for few-shot learning methods) through a 5-shots learning, [51] recently achieved 80.51% of accuracy on the same dataset. Although further studies are required, we can reasonably assume that the improvements of FSL algorithms will further expand the possible use of few-shot learning for real-life use cases.

Applied to marine and coral reef ecology, such methods requiring few examples to fit a model on an identification task could be used for studies on species rarely seen on screen. A key characteristic of highly diverse ecosystems is that they are composed of few very common species and a large proportion of less-common and rare species. Hence, the important effort required to build databases with a sufficient number of images of all these rare species is the main bottleneck preventing the use of Deep Learning on a large number of species. The improvement of few-shot learning algorithms offers promises to build efficient identification models to automatically process images and videos to localise and identify rare fish species. [Such models could then be paired with more classic deep architectures, more efficient to identify abundant species with the leverage of important datasets.](#)

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<i>Abudefduf vaigiensis</i>	<i>Acanthurus leucosternon</i>	<i>Acanthurus lineatus</i>	<i>Amblyglyphidodon indicus</i>

<i>Chaetodon auriga</i>	<i>Chaetodon guttatissimus</i>	<i>Chaetodon trifascialis</i>	<i>Chaetodon trifasciatus</i>

<i>Chromis opercularis</i>	<i>Chromis ternatensis</i>	<i>Gomphosus caeruleus</i>	<i>Halichoeres hortulanus</i>

<i>Monotaxis grandoculis</i>	<i>Naso brevirostris</i>	<i>Naso elegans</i>	<i>Oxymonacanthus longirostris</i>

<i>Pomacentrus sulfureus</i>	<i>Thalassoma hardwicke</i>	<i>Zanclus cornutus</i>	<i>Zebrasoma scopas</i>

Supp. Fig. 1 : The 20 reef fish species considered in this study

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585 *Supp. Fig. 2: Diversity of individuals of the same species and of their*
586 *environments.*

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Supp. Fig. 3: Examples of classes' images in MiniImageNet

Supp. Table 1: Dataset usage during our experiments

Dataset name	Building	Number of annotations	Usage
T0	Human Annotation	69,169	Building of supports sets of 1,5,15 and 30 images for our fourth experiment
T1	Human Annotation	6,320	Building of supports sets or 1, 5, 15 and 30 images for our fifth experiment
T2	Human Annotation	13,232	testing dataset
AT0	Data augmentation applied on T0	691,690	DL training for our first experiment
AT1	Data augmentation applied on T1	63,200	DL training for our second experiment

Supp. Table 2: Mean accuracy obtained with FSL models trained with 1,5,10,15,20,25 and 30 images per species. All the images used for the supports set are from T1.

Species	Number of thumbnails in the Support set						
	1	5	10	15	20	25	30
<i>Abudefduf vaigiensis</i>	16.08	11.39	13.18	12.59	14.02	12.98	11.38
<i>Acanthurus leucosternon</i>	25.51	30.71	34.90	36.68	37.16	36.95	38.74
<i>Acanthurus lineatus</i>	39.86	56.04	63.52	66.21	70.02	70.93	72.50

<i>Amblyglyphidodon</i>							
<i>indicus</i>	25.75	26.74	28.06	28.80	30.24	32.20	32.86
<i>Chaetodo auriga</i>	18.16	25.68	30.67	32.63	33.78	35.42	36.56
<i>Chaetodon</i>							
<i>guttatissimus</i>	33.58	44.21	47.40	49.26	54.12	55.91	58.26
<i>Chaetodon trifascialis</i>	29.02	25.48	27.44	28.02	28.18	29.65	28.44
<i>Chaetodon</i>							
<i>trifasciatus</i>	38.73	50.63	53.47	60.17	63.19	64.38	66.72
<i>Chromis opercularis</i>	44.01	61.81	61.85	63.85	64.67	61.98	62.94
<i>Chromis ternatensis</i>	18.91	24.94	26.27	31.21	33.87	33.49	35.07
<i>Gomphosus</i>							
<i>caeruleus</i>	26.01	38.99	46.84	52.88	53.96	58.69	58.74
<i>Halichoeres</i>							
<i>hortulanus</i>	31.82	44.81	50.73	53.52	54.11	55.85	57.01
<i>Monotaxis</i>							
<i>grandoculis</i>	32.03	41.52	45.68	47.59	48.34	49.19	50.64
<i>Naso brevirostris</i>	47.06	54.26	61.01	59.46	59.30	62.40	64.66
<i>Naso elegans</i>	34.54	43.11	48.38	50.19	51.31	50.55	52.17
<i>Oxymonacanthus</i>							
<i>longirostris</i>	39.29	53.48	59.71	58.84	62.70	64.37	66.26
<i>Pomacentrus</i>							
<i>sulfureus</i>	70.90	88.18	90.92	93.08	92.67	94.13	93.93
<i>Thalassoma</i>							
<i>hardwicke</i>	25.64	44.40	57.80	62.03	64.97	67.86	67.96
<i>Zanclus cornutus</i>	18.33	31.28	37.27	41.71	41.70	42.95	44.44

<i>Zebrasoma scopas</i>	25.56	31.81	33.32	34.62	37.02	37.22	36.05
<hr/>							
Mean	32.04	41.47	45.92	48.17	49.77	50.86	51.77
SD	12.70	16.93	17.69	18.00	18.10	18.52	18.96
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Supp. Table 3: Probability values of the pairwise proportional test used to assess the significance of the difference between accuracies obtained through few-shot models with 5, 10, 15, 20, 25 and 30 shots and deep learning model trained on T1.

	DL	FS1	FS5	FS10	FS15	FS20	FS25
FS1	<2e-16						
FS5	0.28	<2e-16					
FS10	1.20E-08	<2e-16	3.80E-12				
FS15	<2e-16	<2e-16	<2e-16	0.0016			
FS20	<2e-16	<2e-16	<2e-16	4.10E-09	0.0379		
FS25	<2e-16	<2e-16	<2e-16	1.30E-14	8.90E-05	0.2362	
FS30	<2e-16	<2e-16	<2e-16	<2e-16	3.90E-08	0.0059	0.28

Supp. Table 4: Value of the asymptotic accuracy predicted by the NLME models, and number of natural images required for both Deep Learning architecture and Few-shot Learning architecture to reach 99% of this asymptote.

Species	Deep Learning		Few-Shot Learning	
	Number of images required to reach 99% of the asymptote.	Accuracy asymptote value	Number of images required to reach 99% of the asymptote.	Accuracy asymptote value
<i>Naso brevirostris</i>	1,506.47	67.43	5.04	38.10
<i>Monotaxis grandoculis</i>	1,769.71	62.72	7.59	38.14
<i>Amblyglyphidodon indicus</i>	1,766.55	62.89	5.34	41.76
<i>Chromis ternatensis</i>	1,492.65	67.75	6.16	46.55
<i>Acanthurus</i>	1,425.04	69.30	6.42	50.59

<i>lineatus</i>				
<i>Zebrasoma scopas</i>	1,776.17	62.74	5.92	54.94
<i>Gomphosus</i>				
<i>caeruleus</i>	1,341.80	71.51	6.90	53.15
<i>Chromis</i>				
<i>opercularis</i>	1,112.28	78.67	5.85	54.95
<i>Zanclus cornutus</i>	1,069.22	80.57	3.66	60.08
<i>Halichoeres</i>				
<i>hortulanus</i>	1,250.29	74.18	11.60	57.14
<i>Acanthurus</i>				
<i>leucosternon</i>	1,184.97	76.31	10.21	58.28
<i>Chaetodon</i>				
<i>guttatissimus</i>	868.64	90.93	6.46	64.95
<i>Chaetodon auriga</i>	1,053.06	81.31	14.97	65.94
<i>Chaetodon</i>				
<i>trifasciatus</i>	1,076.99	80.34	7.70	63.88
<i>Chaetodon</i>				
<i>trifascialis</i>	1,087.93	79.94	7.42	68.66
<i>Pomacentrus</i>				
<i>sulfureus</i>	960.99	85.63	15.41	72.98
<i>Thalassoma</i>				
<i>hardwicke</i>	994.27	84.00	5.86	78.42
<i>Naso elegans</i>	1,074.08	80.48	6.66	78.13
<i>Oxymonacanthus</i>				
<i>longirostris</i>	833.75	93.29	5.68	80.97
<i>Abudefduf</i>				
<i>vaigiensis</i>	785.95	96.82	4.12	89.78
Mean	1221.54	77.34	7.45	60.87

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