SEQUENTIAL LEARNING OF NEURAL NETWORKS FOR PREQUENTIAL MDL

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ABSTRACT

Minimum Description Length (MDL) provides a framework and an objective for principled model evaluation. It formalizes Occam’s Razor and can be applied to data from non-stationary sources. In the prequential formulation of MDL, the objective is to minimize the cumulative next-step log-loss when sequentially going through the data and using previous observations for parameter estimation. It thus closely resembles a continual- or online-learning problem. In this study, we evaluate approaches for computing prequential description lengths for image classification datasets with neural networks. Considering the computational cost, we find that online-learning with rehearsal has favorable performance compared to the previously widely used block-wise estimation. We propose forward-calibration to better align the models predictions with the empirical observations and introduce replay-streams, a minibatch incremental training technique to efficiently implement approximate random replay while avoiding large in-memory replay buffers. As a result, we present description lengths for a suite of image classification datasets that improve upon previously reported results by large margins.

1 INTRODUCTION

Within the field of deep learning, the paradigm of empirical risk minimization (ERM, Vapnik (1991)) together with model and hyper-parameter selection based on held-out data is the prevailing training and evaluation protocol. This approach has served the field well, supporting seamless scaling to large model and data set sizes. The core assumptions of ERM are: a) the existence of fixed but unknown distributions \( q(x) \) and \( q(y|x) \) that represent the data-generating process for the problem under consideration; b) the goal is to obtain a function \( \hat{y} = f(x, \theta^*) \) that minimizes some loss \( L(y, \hat{y}) \) in expectation over the data drawn from \( q \); and c) that we are given a set of (i.i.d.) samples from \( q \) to use as training and validation data. Its simplicity and well understood theoretical properties make ERM an attractive framework when developing learning machines.

However, sometimes we wish to employ deep learning techniques in situations where not all these basic assumptions hold. For example, if we do not assume a fixed data-generating distribution \( q \) we enter the realm of continual learning, life long learning or online learning. Multiple different terms are used for these scenarios because they operate under different constraints, and because there is some ambiguity about what problem exactly a learning machine is supposed to solve. A recent survey on continual- and life-long learning for example describes multiple, sometimes conflicting desiderata considered in the literature: forward-transfer, backward-transfer, avoiding catastrophic forgetting and maintaining plasticity (Hadsell et al., 2020).

Another situation in which the ERM framework is not necessarily the best approach is when minimizing the expected loss \( L \) is not the only, or maybe not even the primary objective of the learning machine. For example, recently deep-learning techniques have been used to aid structural and causal inference (Vowels et al., 2021). In these cases, we are more interested in model selection or some aspect of the learned parameters \( \theta^* \) than the generalization loss \( L \). Here we have little to gain from the generalization bounds provided by the ERM framework and in fact some of its properties can be harmful.

Independent of ERM, compression based approaches to inference and learning such as Minimum Description Length (Rissanen, 1984; Grunwald, 2004), Minimum Message Length (Wallace, 2005)
We propose two new techniques: forward-calibration with rehearsal. Throughout this empirical study, we consider the computational costs associated with these methods. As a result we present description lengths for a suite of popular image classification datasets that improve upon the previously reported results (Blier & Ollivier, 2018; Bornschein et al., 2020; Jin et al., 2021; Bornschein et al., 2021; Perez et al., 2021; Whitney et al., 2020). We investigate alternatives that are inspired by continuous-learning (CL) based methods. In particular, chunk-incremental and mini-batch incremental fine-tuning with rehearsal. Throughout this empirical study, we consider the computational costs associated with these methods.

Contributions. Previous work on computing prequential description lengths with neural networks relied on a block-wise (chunk-incremental) approximation: at some positions \( t \) a model is trained from random initialization to converge on data \( D_{<t} \) and then their prediction losses on the next intervals are combined (Blier & Ollivier, 2018; Bornschein et al., 2020; Jin et al., 2021; Bornschein et al., 2021; Perez et al., 2021; Whitney et al., 2020). We investigate alternatives that are inspired by continuous-learning (CL) based methods. In particular, chunk-incremental and mini-batch incremental fine-tuning with rehearsal. Throughout this empirical study, we consider the computational costs associated with these methods.

We propose two new techniques: forward-calibration and replay streams. Forward-calibration improves the results by making the model’s predictive uncertainty better match the observed distribution. Replay streams makes replay for mini-batch incremental learning more scalable by providing approximate random rehearsal while avoiding large in-memory replay-buffers. We identify exponential moving parameter averages, label-smoothing and weight-standardization as generally useful techniques. As a result we present description lengths for a suite of popular image classification datasets that improve upon the previously reported results (Blier & Ollivier, 2018; Bornschein et al., 2020) by large margins.

The motivation for studying prequential MDL stems from the desire to apply deep-learning techniques in situations violating the assumptions of ERM. In this work we concentrate on established i.i.d. datasets. While the motivation for choosing MDL over ERM is less clear in this scenario, it allows us to evaluate well known architectures and compare some results from ERM-based training, which has been the focus of the community for decades. We include additional experimental results on non-stationary data in Appendix C.7.

An example: model selection using ERM v.s. MDL. To showcase the properties of prequential MDL and its bias towards simpler models, we generate images by randomly drawing 3 MNIST examples and superimposing them as red(R), green(G) and blue(B) channels. The label is always
Figure 1: **Left:** Example images. **Center:** Best performing ERM-trained models on held-out data. There is no clear indication which conditioning should be preferred. **Right:** Log-loss and errors accumulated throughout the sequence. We observe that MDL-based model selection provides strong evidence that the model conditioned on the R-channel alone should be preferred. In all cases we randomly sampled 50 hyperparameters and use bootstrap-sampling to obtain 95% confidence intervals for the performance of the best performing model. See Figure 2 for detailed regret plots.

Figure 2: **Left:** Next-step neg. log-loss for the best performing models as a function of the position. The models are trained on all the previous examples. We observe that the models conditioned on the R, R and G, or R, G and B channels all rapidly improve their predictive performance when trained on more data. **Right:** The cumulative neg. log-loss (= description length) relative to the best performing model which is conditioned on the R channel only. During the first \( \approx 2k \) examples the models conditioned on irrelevant channels accumulate hundreds of excess nats description length, associated with the red channel; green and blue are distractors. We train a LeNet \cite{lecun1989backpropagation} convolutional neural network on 100k examples from this dataset; either within the ERM framework by drawing i.i.d. examples, training to convergence and validating on held-out data, or within the MDL framework by treating the data as a sequence, training online with replay buffer (see Section 2 for details) and recording the (cumulative) next-step prediction losses and errors. We compare 3 different models: a model conditioned only on the R channel, on the R and G, or on R, G and B channels. Figure 1 (middle) shows the validation-set performance of the best performing models after ERM training. Figure 1 (right) shows description lengths on the same data. It is evident that ERM does not give a clear indication which conditioning should be preferred. However, MDL strongly prefers the model conditioned only on the R-channel. Figure 2 shows a more detailed analysis of the three differently conditioned models in the MDL scenario. We observe that prequential MDL identifies the minimal conditioning because it takes the small-data performance of the models into account. Given sufficient training data the three alternatives perform almost indistinguishable (as seen from their almost parallel regret curves). When we treat the description lengths as evidence for Bayesian model selection to choose between \( M_R, M_{RG} \) and \( M_{RGB} \) we obtain

\[
p(M_R|D) = \frac{p(D|M_R)}{p(D|M_R) + p(D|M_{RG}) + p(D|M_{RGB})}.
\]

With the description length \(- \log p(D|M_R)\) about 300 nats smaller than the ones for \( M_{RG} \) and \( M_{RGB} \), we have \( 1: e^{-300} \) odds in favour of \( M_R \). The Appendix C.1 contains additional experiments with more architectures and the experimental details; demonstrating that this is a robust result regardless of architectures and optimization hyperparameters.

## 2 Online Learning Approaches to Prequential MDL estimation

In this section, we describe practical strategies for computing prequential description lengths with neural networks. Conceptually, computing the cumulative log-loss resembles a continual- or online-learning problem without constraints such as limiting access to previously observed data. In practice, however, some naive approaches such as retraining from scratch to convergence whenever a new example arrives are infeasible due to their high resource usage. In our study, we consider the compute requirements of each approach by counting the number of floating-point operations (FLOPs) until the learner reaches the end of the data sequence. We do not limit the learner’s ability to access or sample
We compare the following approaches to compute prequential description lengths:

- **Chunk Incremental / From-Scratch (CI/FS):** Variants of this approach have been used to compute the recently reported description lengths for deep-learning models (Blier & Ollivier, 2018; Bornschein et al., 2020; Jin et al., 2021; Bornschein et al., 2021; Perez et al., 2021). The data-sequence is partitioned into $K$ non-overlapping intervals; typically of increasing size (i.e. choosing exponentially spaced split points $\{s_k\}_{k=1}^K$, with $s_k \in [2, \ldots, n]$, $s_k<s_{k+1}$ and $s_K=n$). For each $k$, a neural network is randomly initialized and trained to convergence on all the examples before the split point $s_k$ and $s_{k+1}$. This corresponds to the block-wise approximation of the area under the curve: $\sum_{i=1}^n \log p(y_i|\hat{\theta}(D_{<s_k})) \approx \sum_{k=1}^{K-1} \sum_{j=s_k}^{s_{k+1}} \log p(y_j|\hat{\theta}(D_{<s_k}))$, where $\hat{\theta}(D)$ denotes parameters after training on data $D$. To ensure that the model produces calibrated predictions even when $D_{<s_k}$ is small, we use softmax temperature calibration (Guo et al., 2017): at each stage we split the data $D_{<s_k}$ into a 90% training and a 10% calibration data. Conceptually, we could perform post-calibration by first training the network to convergence and then, with all parameters frozen, replacing the output layer $\text{softmax}(h)$ with the calibrated output layer $\text{softmax}(\text{softplus}(\beta)h)$, where $\beta$ is a scalar parameter chosen to minimize the loss on calibration data. In practice, we optimize $\beta$ by gradient descent in parallel with the other model parameters. We alternate computing gradient steps for $\theta$, calculated from the training set and using the uncalibrated network (with final layer $\text{softmax}(h)$), with a gradient step on $\beta$, calculated from the validation set using the calibrated network (with final layer $\text{softmax}(\text{softplus}(\beta)h)$). This simple calibration procedure has proven to be surprisingly effective at avoiding overfitting symptoms when training large neural networks on small datasets (Guo et al., 2017; Bornschein et al., 2020).

- **Chunk Incremental / Continual Fine-tuning (CI/CF):** Similar to CI/FS, the sequence is split into increasingly larger chunks. But instead of training the model from scratch at each stage, the network is continuously fine-tuned on the now larger dataset $D_{<s_k}$. We use the same calibration strategy as for CI/FS. We expect to save compute by avoiding training from scratch. However, recent research suggests that, when first trained on data $D_A$ and then trained on data $D_A \cup D_B$, deep-learning previously observed data because storing and reading from large datasets is generally not considered a major technical bottleneck. However, in-memory (RAM) based implementations of replay-buffers suffer from limited capacity because data sets for training deep neural networks often exceed RAM sizes. Below we describe **replay-streams**, a simple approach to approximate random sampling for replay while only utilizing cheap sequential access to data on disk. With this approach, unlimited approximate random rehearsal from data on permanent storage becomes as easy as sampling from large datasets for ERM based optimization.

We compare the following approaches to compute prequential description lengths:

**Algorithm 1 Mini-batch Incremental Training with Replay Streams**

Require: data $(x_t, y_t)_{t=1}^T$; augmentation $Aug(\cdot)$; number of replay streams $K$; EMA decay $\alpha$

1. Initialize: parameters $\theta$; EMA parameters $\bar{\theta} = \theta$; softmax temperature calibration $\beta = 1$;
   - Replay positions $\{\rho_k = 1\}_{k=1}^K$; $L_{\text{preq}} = 0$
2. for $t = 1$ to $T$
   3. Compute next-step loss: $L_t := -\log p(y_t|x_t, \tilde{\theta}, \beta)$
   4. Update cumulative loss: $L_{\text{preq}} \leftarrow L_{\text{preq}} + L_t$
   5. Update temperature calibration: $\beta \leftarrow \beta - \nabla \beta \log p(y_t|x_t, \tilde{\theta}, \beta)$
   6. Update parameters: $\theta \leftarrow \theta - \nabla \theta \log p(y_t|x_t, \theta, 1.)$ with $\tilde{x}_t = Aug(x_t)$
   7. Update EMA parameters: $\bar{\theta} \leftarrow (1-\alpha)\bar{\theta} + \alpha \theta$
8. for $k = 1$ to $K$
   9. Get data from stream $k$: $x \leftarrow x_{\rho_k}$; $y \leftarrow y_{\rho_k}$, advance position $\rho_k \leftarrow \rho_k + 1$
10. Update parameters: $\theta \leftarrow \theta - \nabla \theta \log p(y|x, \theta, 1.)$ with $\tilde{x} = Aug(x)$
11. Update EMA parameters: $\bar{\theta} \leftarrow (1-\alpha)\bar{\theta} + \alpha \theta$
12. Maintain replay distribution: with probability $p_{\text{reset}}$ reset $\rho_k \leftarrow 1$ (see Equation (1))
13. end for
14. end for
15. return $L_{\text{preq}}$

In practice we perform the algorithm with mini-batches instead of individual examples and use gradient based optimizers such as AdamW (Loshchilov & Hutter, 2019) instead of plain SGD.
models struggle to converge to a solution that generalizes as well as training on $\mathcal{D}_A \cup \mathcal{D}_B$ from scratch; even when both $\mathcal{D}_A$ and $\mathcal{D}_B$ are sampled from the same underlying distribution. In [Ash & Adams (2020)] the authors observe that shrinking the model-parameters and adding some noise after training on $\mathcal{D}_A$ can improve the final generalization performance. We therefore run an ablation study and perform \textit{shrink \& perturb} operation whenever advancing to the next chunk $k$.

Mini-batch Incremental / Replay-Buffer (MI/RB) This approach uses continual online mini-batch gradient descent with an additional replay buffer to store previously seen examples. At each time $t$, the learner performs a number of learning steps on data in the replay buffer. This could be all or a subset of the data $\mathcal{D}_{\leq t}$ depending on the capacity of the replay. We propose \textit{forward-calibration} to optimize the calibration parameter $\beta$: each new batch of examples is first used for evaluation, then used to perform a single gradient step to optimize $\beta$, and finally used for training the parameters $\theta$ and placed into the replay-buffer. Calibration is computationally almost free: the forward-pass for the calibration gradient step can be shared with the evaluation forward-pass, and backward-propagation for $\beta$ ends right after the softmax layer. Appendix B.1 contains a more detailed description. If the replay buffer is limited by the memory capacity, we either use a FIFO replacement-strategy or reservoir-sampling (Vitter, 1985) to maintain a fixed maximum buffer size.

Mini-batch Incremental / Replay-Streams (MI/RS) Implementing large replay buffers can be technically challenging: in-memory (RAM) replay buffers have limited capacity and random access and sampling from permanent storage is often slow. We propose a simple yet effective alternative to storing replay data in-memory. We instead assume that the data is stored in its original order on permanent storage and we maintain $K$ replay streams (pointers into the sequence); each reading the data in-order from the sequence $(x_1, y_1) \ldots (x_T, y_T)$. We can think of a replay-stream as a file-handle, currently at position $\rho \in \ldots, T$. A read operation yields $(x_\rho, y_\rho)$ and increments the position to $\rho \leftarrow \rho + 1$. We denote the position of the $k$th replay-stream with $\rho_k$. Every time the learner steps forward from time $t$ to $t=t+1$ we also read a replay example from each of the $K$ replay-streams and perform gradient steps on them. By stochastically resetting individual streams with probability $p_{\text{reset}}$ to position $\rho \leftarrow 1$, we can influence the distribution of positions $\rho$. If we wish to use uniform replay of previous examples we reset with probability $p_{\text{reset}} = 1/t$. If instead we wish to prioritize recent examples and replay them according to $p_{\text{replay}}(\alpha)$, where $\alpha = t-i$ is the age of the example $(x_i, y_i)$ relative to the current position of the learner $t$, we use

$$p_{\text{reset}}(t) = \frac{\sum_{\alpha=1}^{t} p_{\text{replay}}(\alpha)}{\sum_{\alpha'=0}^{\infty} p_{\text{replay}}(\alpha')}$$

Note that $p_{\text{replay}}$ can be an unnormalized distribution and that by setting $p_{\text{replay}} \propto 1$ we recover $p_{\text{reset}} = 1/t$ for uniform replay. See Algorithm 1 for details.

3 Experiments

We empirically evaluate the approaches from Section 2 on a suite of models and image classification data sets. We use MNIST (LeCun et al., 2010), EMNIST, CIFAR-10, CIFAR-100 (Krizhevsky, 2009) and ImageNet (Russakovsky et al., 2015) and randomly shuffle each into a fixed sequence of examples. With MDL, we do not require validation or test set and could merge it with the training data. But because the data is known to be i.i.d. and we compare results with established ERM-based training, we instead maintain the standard test sets as held-out data. It must be emphasized that the primary objective throughout this study is to minimize the description length. All hyperparameters are tuned towards that goal. When we report test set performance it should be considered a sanity check. We evaluate the following architectures: MLPs, VGG (Simonyan & Zisserman, 2015), ResNets (He et al., 2016), WideResNets (Zagoruyko & Komodakis, 2016) and the transformer-based DeiT (Touvron et al., 2021) architecture. We refer to a variant of the VGG architecture with weight-standardized convolutions (Qiao et al., 2019) and with batch normalization (Ioffe & Szegedy, 2015) as VGG++.

For the bulk of our experiments, we use the AdamW (Loshchilov & Hutter, 2019) as optimizer; a series of control experiments however suggest that the quantitative results generalize to optimizers such as RMSProp and simple mini-batch SGD with momentum. We identified the following techniques that consistently improve the results and use them throughout this paper if not mentioned otherwise:
Figure 3: Shortest description lengths as function of compute resources consumed for different methods. For each method (in different colors) we run 250 (100 for ImageNet) independent experiments with randomly sampled hyperparameters. Each dot represents such an experiment and the solid lines show the Pareto-front of the best archived performance for different compute budgets. With the exception of CIFAR-100 in the large-FLOPs regime, the mini-batch incremental approach MI/RS results in the shortest description lengths.

Exponentially moving average (EMA) Similar to Polyak averaging [Polyak & Juditsky, 1992], EMA maintains a moving average of the model parameters $\theta$ and is used for next-step evaluations as well as when tuning the softmax temperature $\beta$.

Label smoothing [Szegedy et al., 2015] A reliable regularizer, label smoothing often helps for the continual fine-tuning methods such as MI/RB, MI/RS and CI/CF.

Weight standardization For the VGG architecture, weight standardized convolutions lead to significant improvements - bigger than for conventional ERM-based training. For other architectures such as ResNets, the results are not conclusive.

3.1 COMPARISON OF ONLINE LEARNING APPROACHES.

To compare the effectiveness of the different approaches listed in Section 2 we run a broad random hyperparameter sweep for EMNIST with LeNet, CIFAR-10 and CIFAR-100 with VGG++ and ImageNet with ResNet50. The hyperparameter intervals depend on the data and are detailed in Appendix B. We sample learning rate, EMA step size, batch size, weight decay; but crucially also number of epochs (or, correspondingly, number of replay streams for MI/RS) and an overall scaling of the model width (number of channels). We thus cover a large range of computational budgets, spanning about 2 orders of magnitude. When the capacity of the replay-buffer matches the total length of the sequence, MI/RB and MI/RS are equivalent. In these cases we omit MI/RB and instead use MI/RS to represent online-learning with unlimited rehearsal. A study of replay buffer v.s. replay streams can be found in Section 3.1. We show the overall Pareto front for these experiments in Figure 3. Mini-batch incremental learning obtains the lowest description lengths for a wide range of computational budgets. Only for very large compute budgets on CIFAR-100 we observe that the block-wise approximation with training from scratch (CI/FS) has a better results.

MDL for Deep Neural Networks. We apply MI/RS to a wider range of architectures. For most experiments we again use random hyperparameter sweeps, this time however without scaling the size
| Data               | Model     | MI/RS (Left half of the table) | ERM (Right half of the table) |
|-------------------|-----------|-------------------------------|------------------------------|
|                   |           | Cumulative                     | Test                         |
|                   |           | Loss | Errors | Loss | Error(%) | FLOPs | Loss  | Error(%) | FLOPs |
| MNIST             | Lenet     | 4.4k | 1362   | 0.03 | 1.0      | 1.4e13| 0.03  | 0.8      | 1.5e13|
| MNIST             | MnistNet  | 2.1k | 643    | 0.02 | 0.5      | 1.0e13| 0.02  | 0.5      | 3.2e14|
| CIFAR-10          | VGG++     | 22.1k| 6.6k   | 0.24 | 5.8      | 5.9e15| 0.21  | 6.9      | 8.0e15|
| CIFAR-100         | WRN-28-10 | 22.9k| 7.5k   | 0.23 | 7.4      | 3.7e16| 0.20  | 6.7      | 5.4e16|
| CIFAR-100         | VGG++     | 93.7k| 23.1k  | 1.14 | 32.0     | 8.5e15| 1.04  | 7.6      | 8.5e16|
| CIFAR-100         | WRN-28-10 | 87.0k| 22.3k  | 1.10 | 30.3     | 2.5e16| 1.08  | 28.3     | 4.3e16|
| ImageNet          | VGG++     | 2.55M| 549k   | 1.63 | 36.8     | 7.5e18| 1.25  | 29.9     | 7.0e17|
| ImageNet          | ResNet-34 | 2.35M| 507k   | 1.30 | 30.9     | 1.6e18| 1.25  | 29.9     | 7.0e17|
| ImageNet          | ResNet-50 | 2.21M| 496k   | 1.29 | 29.1     | 8.5e17| 1.15  | 28.7     | 1.9e18|
| ImageNet          | ResNet-50†| 1.93M| 431k   | 1.06 | 26.1     | 4.0e18| 1.10  | 26.6     | 4.6e18|
| ImageNet          | ResNet-101| 1.97M| 447k   | 1.20 | 28.2     | 3.1e18| 1.10  | 26.6     | 4.6e18|
| ImageNet          | DeiT-S    | 2.73M| 588k   | 1.40 | 31.7     | 1.2e18| 1.22  | 28.1     | 4.8e18|

**Prior Works:**
- MNIST (Blier & Ollivier, 2018) MnistNet 2.8k 0.5
- CIFAR-10 (Blier & Ollivier, 2018) ≈ VGG++ 31.4k 6.7
- ImageNet (Ioffinsen et al., 2020) ResNet-50 3.32M -

Table 1: **Left half of the table**: Suite of results when applying MI/RS for description length estimation. Cumulative loss in nats; cumulative error counts the number of prediction errors after reaching the end of the sequence. The test set metrics report the performance of the model selected by lowest description length. **Right half**: Results when using the same architecture and hyperparameter-sweep for conventional ERM training. † Manually tuned hyper parameters.

We present our results for the best run in each sweep and contrast them with previously reported description lengths from the literature in Table 1. We also show the test set performance of these models after they reached the end of training data. The rightmost columns show the test set performance when performing the same hyperparameter sweep for conventional ERM training on the full data set. Note that these runs do not use any kind of learning rate annealing, which is otherwise often employed to improve results.

Based on these experiments we plot a regret comparison for different architectures trained on ImageNet in Figure 4 (left). We observe that the additional depth and capacity of a ResNet101 is beneficial from the very beginning of the data-sequence. Figure 4 (right) shows the effect of scaling the number of channels of a VGG++ architecture on CIFAR-100 and it is evident that decreasing the model width has negative effects throughout the sequence and not just when the data gets larger.

![Figure 4: Left: Comparing architectures on ImageNet. Right: Scaling the size of VGG++.](image)

**Replay Buffer and Replay Streams** are equivalent for short data sequences, they differ when the length of a sequence exceeds the capacity of the replay buffer. In this section, we study the effects of limiting the replay buffer size. Figure 5 shows that reducing the capacity has a significant negative effect on the results. In addition, the strategy of which samples to keep in the buffer plays an important role. Using reservoir sampling instead of a FIFO replacement policy leads to further severely degraded results. This is maybe not surprising because with reservoir-sampling insertion to the buffer is stochastic; a fraction of the examples are thus never replayed at all. Appendix C.5 contains ablations for the number of replay-streams on CIFAR-100 and ImageNet.

**Ablations.** We provide ablations for using forward-calibration, label smoothing and weight standardization in Table 2. It is evident that each of these techniques helps obtaining better description lengths for the datasets and architectures under consideration. Regret plots for these ablations can be found in Appendix C.
Table 2: Description lengths (in nats) with various techniques added to the baseline model: forward-calibration (FC), label smoothing (LS) and weight standardization (WS).

| Data        | Model     | None       | +FC        | +LS        | +WS        | +ALL       |
|-------------|-----------|------------|------------|------------|------------|------------|
| CIFAR-10    | VGG++     | 32.4k      | 31.8k (-0.6k) | 32.1k (-0.3k) | 26.4k (-5.9k) | 23.3k (-9.0k) |
| CIFAR-100   | VGG++     | 103.4k     | 100.7k (-2.7k) | 102.8k (-0.6k) | 93.1k (-10.2k) | 89.1k (-14.2) |
| ImageNet    | ResNet50  | 2.33M      | 2.27M (-60.0k) | 2.28M (-50k)  | 2.21M (-120k)  |

Plasticity and the Warm-starting problem  We take a closer look at the large-FLOPs regime for CIFAR-100 where CI/FS obtained better results than MI/RS. Figure 6 (left) shows the regret comparison between the best runs for each training method. The split positions $s_k$ are clearly visible as kinks in the graph for CI/FS and CI/CF. For CI/CF the plot shows that right after fine-tuning, the predictive performance is approximately equal to MI/RS. For CI/FS we see that the regret relative to MI/RS decreases: it has better predictive performance. This effect persists even after running extensive additional hyperparameter sweeps. We conjecture that this is a signature of the warm-starting problem described in (Ash & Adams, 2020). Figure 6 (right) shows the Pareto front for the final test set accuracy of the best models (selected by their description length) over FLOPs. We observe a large gap between fine-tuning based methods and CI/FS: which is consistent with this hypothesis. Note that the last split-point $s_K = T$ implies that these models are effectively trained like in a conventional ERM paradigm. Appendix C shows test set pareto fronts for a range of datasets. Close inspection of various regret plots sometimes reveals the same signatures, suggesting that the effect is ubiquitous, but often much less pronounced. It is plausible that this also contributes to the gap in final test set performance in Table 1.

4 Discussion & Related Work

Compression based approaches to inference and learning, such as Solomonoff induction (Solomonoff, 1964), Minimum Message Length (Wallace, 2005) and Minimum Description Length (Rissanen, 1984; Grunwald, 2004), have been extensively studied. The Bayesian and variational Bayesian approaches to estimating code lengths have been popular and, at least outside the field of deep learning, also been very successful. This line of work includes methods such as AIC and BIC, as well as more sophisticated approximations to the posterior distribution over model parameters for many kinds of models, including neural networks (Hinton & Van Camp, 1993; MacKay, 2003). The description length depends crucially on the quality of these posterior approximations. Unfortunately, estimating the parameter posterior for modern neural networks is notoriously hard and an area of active research (Izmailov et al., 2021). Blier & Ollivier (2018) demonstrated that much shorter code lengths can be achieved by using the prequential approach with standard neural network architectures and optimizers. The prequential perspective with the block-wise approximation technique have been successively used in the context of image-classification (Bornschein et al., 2020), natural language processing (Voita & Titov, 2020; Jin et al., 2021; Perez et al., 2021) and causal structure discovery (Bornschein et al., 2021).
With our work, we aim to simplify and improve prequential code length estimation for neural networks with insights from the field of continuous learning (CL). For CL, fine-tuning with rehearsal is considered a strong baseline and large number of methods have been proposed to maintain its benefits while minimizing the size of the replay buffer (Mnih et al., 2013; Delange et al., 2021). We instead propose a method that makes it practical and relatively cheap to work with full rehearsal. Caccia et al. (2022) investigate very similar scenario to ours; largely phrased around how long a learner should wait and accumulate data when it is compute constraint; most experiments however consider the case without full rehearsal. He and Lin (He & Lin, 2020) discuss the close relationship between prequential coding and CL, however focus on compressive replay instead of accessing previous data. In general, the literature on CL methods and evaluation metrics is too broad to be adequately discussed here (Hadsell et al., 2020). However, we would not be surprised if some existing CL methods have the potential to vastly improve prequential MDL computation. In this sense, we see our work as a step of bringing CL and prequential MDL related works closer together.

Besides of taking inspiration from CL to improve description lengths, we believe that the MDL perspective can help guide future CL research: Much of the CL literature is rooted in an ERM-based perspective and thus considers distinct training, validation and test sets at different stages during learning. This has potentially played a role in the proliferation of many different evaluation metrics and training protocols used in CL research (Hadsell et al., 2020). This heterogeneity of the evaluation metrics makes it challenging to compare results. Forward-transfer is a commonly used evaluation metric and it is closely related to a prequential evaluation. However, it is often computed on separate held-out sets after training and does not take sample-efficiency into account as prequential MDL does. Recent work has embraced forward-validation in terms of average prediction-error on future training data as a evaluation metric, thus bringing it closer to a prequential MDL based evaluation but without discussing the connection (Cai et al., 2021; Lin et al., 2021). MDL has been conceived and analyzed as a theoretically well motivated objective when dealing with non-stationary data; without reference to the practical difficulties of actually performing parameter estimates. It is a challenging objective even for learners without constraints on compute and (replay-) memory.

5 Conclusions & Limitations

In this paper, we compare different approaches to computing prequential description lengths while considering their computational cost. We find that continuous-learning inspired mini-batch incremental training with full rehearsal is the preferred method over a wide range of computational budgets. We introduce two techniques that either improve results, or make it practical to apply full rehearsal even for very large data. As a result we report description lengths for a range of standard data sets that improve significantly over previously reported figures. Some of our results show that the warmstarting-effect can have a negative effect on the performance of continual fine-tuning. We also generally observe a gap in the final held-out performance after training models within the prequential framework vs. training models with the ERM-based approach. We are not aware of any technique to reliably mitigate the impact. It would be an important topic to tackle for future works. Furthermore, learning rate schedules such as cosine decay are popular and effective techniques for achieving better performance in ERM-based training. It is not obvious how to best translate them to the prequential learning scenario as the learner constantly receives new data.
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A Pitfalls of using test splits

Test set contamination is an increasingly grave unsolved problem that becomes more pressing as machine learning research moves from curated academic benchmark data to large user provided or web-scraped data. In the recent GPT-3 paper on large scale language modeling the authors dedicate multiple pages in the Appendix to discuss the issue (Brown et al., 2020). The paper details the method used to detect training-/test set duplicates and presents experiments and the conclusion that their particular model and evaluation approach is relatively insensitive to contamination. Indeed, some of the contamination was only discovered after the model had been trained, and the model could not be retrained with decontaminated data due to the prohibitive (computational) cost. This prompted some follow-up discussion on how validation and test sets should be created and decontaminated.

In (Søgaard et al., 2021) the authors show that depending on the exact splitting procedure significantly different results can be obtained. The discussion lays out arguments in favour of either using deliberately biased splits, or forward-validating on future data. This work refers to prior work (Gorman & Bedrick, 2019), which also described inconsistent results between different test sets, and between the random split, but at large came to a different conclusion.

To the best of our knowledge these issues have been primarily discussed in the NLP community. We mention them to point out that validation on held-out data can not blindly be regarded as the gold-standard for model evaluation when, at times, the procedure to create those splits is subject to intense discussion. Instead, careful consideration will be necessary in order to avoid misleading conclusions based on inappropriate validation and test sets.

Test set contamination is not only an increasingly important problem but also in a certain sense an unsolvable one, because it is ill-defined. At what threshold for example should a document that verbatimly cites a training-set document be removed from the test-set? When are two images scraped from the internet essentially the same?

On the other side: Test set contamination is no problem when using any of the MDL variants for model evaluation. For NML and Bayes this is true because they use model complexity and marginal predictions for regularization instead of held-out sets to avoid overfitting. But our limited ability to work with good parameter posterior estimates and the benign overfitting of overparametrized NNs make it largely non-applicable to deep learning. For prequential MDL it is true, because the model is evaluated and hence has to perform well already the first time a (potentially duplicate) data item appears.

For instance, assume the whole photo data base contains three or more (approximate) copies of each photo. If we randomly split off 10% as the test set, then the train set contains nearly all test set items. With held-out-validation a pure memorizer without any generalization capacity will perform nearly perfectly on the test set, but will fail miserably in practice on any newly taken photo. On the other hand, NML and marginal Bayes punish a memorizer due to its huge complexity, and prequential MDL due to the inability to predict (well) the first occurrence of each item. Test set evaluation is empirically sound for i.i.d. data, therefore the failure of this paradigm must be attributed to the non-iid nature of data with duplicates (Hutter, 2022).

For a more detailed discussion, see the FAQ in (Hutter, 2006/2020).

B Experimental Details

B.1 Forward Calibration

We train the neural network with standard cross-entropy loss and the categorical prediction-head \( p = \text{softmax}(h) \), where \( h \) is a vector of logits generated by the neural network. When using the model for evaluation on next-step data or on test set examples, we replace it with a calibrated prediction head \( p = \text{softmax}(\text{softplus}(\beta) h) \), where \( \beta \) is a scalar temperature parameter (Guo et al., 2017). We use forward-calibration to optimize the calibration parameter \( \beta \); each new batch of examples from the training sequence is used first for evaluation, then used to perform a single gradient step to optimize \( \beta \), and finally used for a training to optimizer \( \theta \) and placed into the replay-buffer. At the beginning of training we initialize \( \beta \) to \( \beta_0 = \log(\exp(1) - 1) \approx 0.5416 \), such that \( \text{softplus}(\beta_0) = 1 \). We use the same optimizer as for the model parameters \( \theta \), however scale the learning rate by a factor of \( \sqrt{K} \),
where $K$ is the number of training steps per evaluation step forward (i.e., number replay-streams plus one). Calibration is computationally almost free: the forward-pass for the calibration gradient step can be shared with the evaluation forward-pass, and backward-propagation for $\beta$ ends shortly after the softmax layer.

B.2 MNIST, EMNIST AND RGB-MNIST HYPERPARAMETERS

We run experiments with various model architectures: a) MLP with 1, 2 or 3 hidden layers, 512 units each, with dropout, and ReLU non-linearities, b) LeNet (LeCun et al., 1989) and c) the VGG inspired architecture used by (Blier & Ollivier, 2018), which we call MnistNet here.

We use the same hyperparameter and sampling intervals for all three model architectures. For the pareto front experiment we extend the number of replay-streams range to 10 . . . 100; otherwise we use the same intervals.

| Parameter                        | Distribution       | Values / Interval |
|----------------------------------|--------------------|-------------------|
| Number of replay-streams         | log-uniform        | 25 . . . 100       |
| Learning rate                    | log-uniform        | 1e-4 . . . 3e-3    |
| AdamW $\epsilon$                | log-uniform        | 1e-4 . . . 1       |
| EMA step size (Polyak averaging) | log-uniform        | 1e-3 . . . 1e-1    |
| Weight decay                     | log-uniform        | 1e-4 . . . 1.      |
| Batch size                       | fixed              | 32                |
| Label smoothing                  | fixed              | 0.001             |

B.3 CIFAR-10 AND CIFAR-100 WITH VGG++, AND WIDERESNET HYPERPARAMETERS

We use the same hyperparamater sampling intervals for all model architectures. Following Zagoruyko & Komodakis (2016) we use only minimal augmentations during training: Images will be horizontally flipped with a probability of 0.5. For the pareto-front plots we extend the number of epochs (= number of replay-streams) range to 10 . . . 100 and

| Parameter                        | Distribution       | Values / Interval |
|----------------------------------|--------------------|-------------------|
| Number of replay-streams         | log-uniform        | 25 . . . 100       |
| Learning rate                    | log-uniform        | 1e-4 . . . 3e-3    |
| AdamW $\epsilon$                | log-uniform        | 1e-4 . . . 1       |
| EMA step size (Polyak averaging) | log-uniform        | 1e-3 . . . 1e-1    |
| Weight decay                     | log-uniform        | 1e-4 . . . 1e-1    |
| Batch size                       | uniform            | 32, 64, 128       |
| Label smoothing                  | fixed              | 0.01              |

B.4 IMAGENET WITH VGG++ AND RESNETS HYPERPARAMETERS

We use randaugment for data augmentation and the same hyperparameter intervals for all experiments. For Pareto front experiments we additionally scale the architecture size (number of channels throughout) from $1/4 \times$ to $4 \times$ and extend the number of replay-streams interval to 10 . . . 100.

| Parameter                        | Distribution       | Values / Interval |
|----------------------------------|--------------------|-------------------|
| Number of replay-streams         | log-uniform        | 25 . . . 100       |
| Learning rate                    | log-uniform        | 1e-4 . . . 3e-3    |
| AdamW $\epsilon$                | log-uniform        | 1e-4 . . . 1       |
| EMA step size (Polyak averaging) | log-uniform        | 1e-3 . . . 1e-2    |
| Weight decay                     | log-uniform        | 1e-4 . . . 1.      |
| Batch size                       | fixed              | 512               |
| Label smoothing                  | fixed              | 0.01              |
B.5 CLOC with ResNets Hyperparameters

We use randaugment for data augmentation and the same hyperparameter intervals for all experiments.

| Parameter                        | Distribution | Values / Interval |
|----------------------------------|--------------|-------------------|
| Number of replay-streams        | log-uniform  | 10 . . . 25       |
| Learning rate                    | log-uniform  | 1e-4 . . . 3e-3   |
| EMA step size (Polyak averaging) | log-uniform  | 1e-3 . . . 1e-2   |
| Batch size                       | fixed        | 128               |
| Label smoothing                  | fixed        | 0.1               |

C Additional Empirical Results

C.1 RGB-MNIST Feature Selection

To illustrate that the model-selection property demonstrated in section 1 is a robust property of (prequential) MDL, we here present a suite of results on 3 different model architectures: 1) a MLP with 2 hidden layers, 512 units each; 2) the LeNet architecture from [LeCun et al., 1989]; and 3) the much higher capacity and better tuned convnet from [Blier & Ollivier, 2018].

In all cases we run the same hyperparameter sweep detailed in [Appendix B]. We show the regret plot for the best performing model in each sweep relative to the model conditioned only on the R channel. We see pMDL reliably and with confidence determines that the model conditioned on R only is the most appropriate one for this data; and we see that the small-data performance at the beginning of the sequence is crucial as the relative generalization performance becomes more similar as the length of the sequence increases:

MLP Regret Plot

LeNet Regret Plot

MnistNet Regret Plot
C.2 Pareto Fronts

**Left:** Description lengths as function of compute resources consumed for a selection of datasets and models. **Right:** Test set performance after reaching the end of the training data. We generally observe a gap between models that train from scratch (CI/FS) and continual finetuning methods. The gap is very pronounced for CIFAR-100. We do not observe a systematic gap between MI/RS and CI/CF.
C.3 Model Depth Ablation

We run ablations for different number of replay-streams (K in Algorithm 1) on CIFAR-100 and ImageNet. The results on ImageNet show that performing too much replay (choosing K too large) can be harmful.

C.4 Weight-Standardization Ablation

Besides of uniform replay for replay-stream training we experimented with alternative replay distributions. Exponentially decaying replay ($\rho_{\text{replay}}(a) = \lambda \exp(-\lambda a)$) is popular in continual-learning with reservoir sampled replay buffers, but lead to significant longer description lengths and suboptimal results in our experiments.

As an compromise between uniform replay and exponential decay we experimented with long-tailed Pareto distributions: $\rho_{\text{replay}}(a) = \frac{\alpha}{(a/\lambda)^{\alpha+1}} \cdot \frac{1}{\lambda^{\alpha+1}}$. The scale parameter $\lambda$ determines to what extend old examples are replayed and we use a fixed shape parameter $\alpha = \log_4 5 \approx 1.16$ for all out experiments:

C.5 Number of Replay Streams

C.6 Scale and Shape of the Replay Distribution

Besides of uniform replay for replay-stream training we experimented with alternative replay distributions. Exponentially decaying replay ($\rho_{\text{replay}}(a) = \lambda \exp(-\lambda a)$) is popular in continual-learning with reservoir sampled replay buffers, but lead to significant longer description lengths and suboptimal results in our experiments.

As an compromise between uniform replay and exponential decay we experimented with long-tailed Pareto distributions: $\rho_{\text{replay}}(a) = \frac{\alpha}{(a/\lambda)^{\alpha+1}} \cdot \frac{1}{\lambda^{\alpha+1}}$. The scale parameter $\lambda$ determines to what extend old examples are replayed and we use a fixed shape parameter $\alpha = \log_4 5 \approx 1.16$ for all out experiments:
Our experiments suggest that flat, almost uniform replay distributions lead to the shortest description lengths for the sequences from i.i.d. sources.

C.7 Non-stationary Sequence: Continual Geo-localization (CLOC)

In this study we concentrate on data sequences from stationary (i.i.d.) data sources to ensure we can compare our results with established ERM based training. However, one major benefit of MDL over ERM is that it is well defined for individual and non-stationary sequences. We here present empirical results on the CLOC data introduced in (Cai et al., 2021). CLOC is a large scale, chronologically sorted image classification sequence consisting of \( \approx 39M \) images with their geolocation as a categorical label. For \( \approx 5\% \) of the images we either received download errors, or the downloaded content could not be decoded as image data. In total we were able to obtain 37,093,769 labeled images. Figure 2 in (Cai et al., 2021) shows that the data is strongly non-stationary; and that traditional ERM training, which treats the data as i.i.d., results in a held-out error rate between 80 and 90%.

In (Cai et al., 2021) the authors reserve the first 2M (5\%) datapoints for pre-training the model statically before entering the online-phase, split-off 1\% of the data throughout for validation purposes and only evaluate the images from the first new album in each mini-batch instead of all images. Their online-learning task and metric are thus in spirit similar to the prequential MDL problem we are tackling here, in detail however sufficiently different that the quantitative results are not directly comparable. Following (Cai et al., 2021) we use a standard ResNet50 architecture and run experiments with replay buffers and replay streams. The hyperparameter sampling space is detailed in Appendix B.5. Note that we use a batch size of 128 because small batch-sizes allow the model to more quickly adapt to changes in the data distribution; and a relatively low number of 10-25 replay-steps / replay-streams to limit the total computational cost due to the size of the data.

We observe that heavily recency biased replay leads to significantly improved results. For replay streams we experimented with uniform replay, replay with exponentially decaying priority and with replay governed by a heavy tailed Pareto distribution (see C.6). From these the heavy tailed Pareto replay leads to the shortest description lengths. Even more than in the i.i.d. case, forward-calibration plays an crucial role to obtain good results.

Next-step performance  We plot the 5k rolling average next-step performance of the best performing model from our random hyperparameter sweep \((K=22)\) Pareto distributed replay streams with
a scale of 180). The model obtains a description length of 1.33G nats and a next-step error-rate of typically $60 \pm 10\%$.

**Ablations Left:** We show the description length regret for the best replay-buffer training run relative to the best replay-stream training run. Both experiments consumed about $5 \times 10^{18}$ FLOPs.

**Right:** Regret for replay-streams without forward calibration relative to replay streams with forward calibration: