Memory Replay with Data Compression for Continual Learning

Abstract

To continually learn a sequence of tasks, deep neural networks need to overcome catastrophic forgetting of the past. Memory replay of representative old training samples has been shown as an effective solution, and achieves the state-of-the-art (SOTA) performance. However, existing memory replay approaches are mainly built on a small memory buffer containing a few original data, which cannot fully characterize the old data distribution. In this work, we propose memory replay with data compression, which is an important baseline and a promising direction for continual learning. Data compression can largely reduce the storage of old training samples and thus increase their amount that can be stored in the memory buffer. We empirically validate that there is a trade-off between the quantity and quality of compressed data, which is highly nontrivial for the efficacy of memory replay. Then we propose a novel method based on determinantal point processes (DPPs) to efficiently determine an appropriate quality of data compression for memory replay. Across several benchmarks of class-incremental learning, using a naive technique of data compression with a properly selected quality can achieve the SOTA performance in a computation-efficient and plug-and-play way. We further demonstrate the advantages of our proposal in realistic continual learning of object detection for autonomous driving, where the incremental data are extremely large-scale with high storage cost.

1 Introduction

The ability to continually learn numerous tasks and infer them together is critical for deep neural networks (DNNs), which needs to mitigate *catastrophic forgetting* (McCloskey & Cohen, 1989) of the past. Memory replay of representative old training samples (referred to as *memory replay*) has been shown as an effective solution, and achieves the state-of-the-art (SOTA) performance (Hou et al., 2019). Existing memory replay approaches are mainly built on a small memory buffer containing a few original data, and try to construct and exploit it more effectively. However, due to the low storage efficiency of saving original data, this strategy of building memory buffer will lose a lot of information about the old data distribution. On the other hand, recent approaches in this direction attempted to improve the efficacy of memory replay by learning additional parameters (Liu et al., 2020b) or distilling old features (Hu et al., 2021), which will largely increase the computation cost.

Different from "artificial" memory replay in DNNs, a significant feature of biological memory is to encode the old experiences in a highly compressed form and replay them to overcome catastrophic forgetting (McClelland, 2013; Dragoi & Buzsáki, 2006; Davidson et al., 2009; Carr et al., 2011). Thus, a large amount of information about the past can be maintained in a small storage space as comprehensively as possible, and flexibly retrieved. Inspired by the compression feature of biological memory replay, we propose memory replay with data compression, which can largely increase the amount of old training samples that can be stored in the memory buffer by reducing their storage cost. As the operation of data compression only needs a small amount of extra computation, this strategy can both more effectively and more efficiently recover the old data distribution.

Given a limited storage space, data compression introduces an additional degree of freedom to explicitly balance the trade-off between quality and quantity for memory replay. With a properly selected quality, using a naive JPEG algorithm (Wallace, 1992) of data compression can achieve the SOTA performance of class-incremental learning with the lowest extra computation cost (Fig.

Figure 1: Averaged accuracy and computation cost on ImageNet-sub. Compared with the SOTA approaches, memory replay with data compression (ours) achieves the SOTA performance with the lowest extra computation cost (purple arrow), and can further boost their performance (gray arrow).

1, purple arrow) and can further improve the SOTA memory replay approaches in a *plug-and-play* way (Fig. 1, gray arrow). However, it is often computation-inefficient to empirically determine the compression quality[1]. We propose a novel method based on determinantal point processes (DPPs) to efficiently determine it without extra training. Further, we demonstrate the advantages of our proposal in realistic applications such as continual learning of object detection for autonomous driving, where the incremental data are extremely large-scale with high storage cost.

Our contributions include: (1) We present that memory replay with data compression is both an important baseline and a promising direction for continual learning; (2) We empirically validate that the trade-off between the quality and quantity of compressed data is highly nontrivial for memory replay, and provide a novel method to efficiently determine an appropriate quality of data compression; (3) Extensive experiments show that using a naive technique of data compression with a properly selected quality can substantially boost the efficacy of memory replay, and achieves the SOTA performance in a computation-efficient and plug-and-play way.

2 RELATED WORK

Continual learning needs to overcome catastrophic forgetting of the past when learning a new task. Regularization-based methods (Kirkpatrick et al., 2017; Chaudhry et al., 2018) approximated the importance of each parameter to the old tasks and selectively penalized its changes. Architecture-based methods (Rusu et al., 2016) allocated a dedicated parameter subspace for each task to prevent mutual interference. Replay-based methods (Rebuffi et al., 2017; Shin et al., 2017) approximated and recovered the old data distribution. In particular, memory replay of representative old training samples (referred to as *memory replay*) can generally achieve the best performance in class-incremental learning (Liu et al., 2021a; Hu et al., 2021) and in numerous other scenarios of continual learning, such as few-shot (Tao et al., 2020b), semi-supervised (Wang et al., 2021), and unsupervised continual learning (Khare et al., 2021), and continual learning of audio tasks (Ehret et al., 2020).

Existing works in memory replay are mainly built on a small memory buffer containing a few original data, and attempt to construct and exploit it more effectively. As the pioneer work, iCaRL (Rebuffi et al., 2017) proposed a general protocol of memory replay for continual learning. To better construct the memory buffer, Mnemonics (Liu et al., 2020c) parameterized the original data and made them optimizable, while TPCIL (Tao et al., 2020a) constructed an elastic Hebbian graph (EHG) optimized by competitive Hebbian learning. On the other hand, BiC (Wu et al., 2019), LUCIR (Hou et al., 2019), PODNet (Douillard et al., 2020), DDE (Hu et al., 2021) and AANets (Liu et al., 2021a) attempted to more effectively exploit the memory buffer, such as by mitigating the data imbalance between old and new classes (Hou et al., 2019; Wu et al., 2019; Hu et al., 2021).

In contrast to saving original data, numerous efforts attempted to remember the old data distribution more efficiently. One solution is based on continual learning of a generative model, which can be used to replay generated data (Shin et al., 2017; Wu et al., 2018) or compress old training data (Caccia et al., 2020). However, continual learning of such a generative model is extremely challenging, which limits its applications to relatively simple domains, and is generally computation-inefficient. Another solution is based on feature-replay: GFR (Liu et al., 2020a) replayed generated features by learning a feature generator from a feature extractor, but the feature extractor suffered from catastrophic

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[1]A naive grid search approach is to train continual learning processes with different qualities and choose the best one, resulting in huge computation cost.

forgetting since it was incrementally updated. REMIND (Hayes et al., 2020) saved the old features and reconstructed the synthesized features for replay, but it froze the majority of feature extractor after learning the initial phase, which limits the learning of representations for incremental tasks.

Data compression aims to improve the storage efficiency of a file, including lossless compression and lossy compression. Lossless compression needs to perfectly reconstruct the original data from the compressed data, which limits its compression rate (Shannon, 1948). In contrast, lossy compression can achieve a much higher compression rate by degrading the original data, so it has been broadly used in realistic applications. Representative hand-crafted approaches include JPEG (or JPG) (Wallace, 1992), which is the most commonly-used algorithm of lossy compression (Mentzer et al., 2020), WebP (Lian & Shilei, 2012) and JPEG2000 (Rabbani, 2002). On the other hand, neural compression approaches generally rely on optimizing Shannon's rate-distortion trade-off, through RNNs (Toderici et al., 2015; 2017), auto-encoders (Agustsson et al., 2017) and GANs (Mentzer et al., 2020).

3 Continual Learning Preliminaries

We consider a general setting of continual learning, which needs to incrementally learn numerous tasks $t = 1, 2, 3, \ldots$ from their task-specific training dataset $D_t = \{(x_{t,i}, y_{t,i})\}_{i=1}^{N_t}$, where $(x_{t,i}, y_{t,i})$ is a data-label pair and N_t is the number of such training samples. For classification tasks, the training samples of each task might be from one or several new classes, i.e., class-incremental learning. In continual learning, D_t is only available when learning task t, and all the tasks ever seen are evaluated at test time, which is known as the single-head evaluation (Chaudhry et al., 2018). Suppose a DNN with parameter θ has incrementally learned T tasks. When learning a new task, since the old training datasets $\bigcup_{t=1}^{T} D_t$ are unavailable, the DNN will adapt the learned parameters to fit D_{T+1}, and tend to catastrophically forget the old tasks (McCloskey & Cohen, 1989).

An effective solution of overcoming catastrophic forgetting is to select and store representative old training samples $D_t^{mb} = \{(x_{t,i}, y_{t,i})\}_{i=1}^{N_t^{mb}}$ in a small memory buffer (mb), and replay them when learning the new task. For classification tasks, mean-of-feature is a widely-used strategy to select D_t^{mb} (Rebuffi et al., 2017; Riemer et al., 2018; Hou et al., 2019; Douillard et al., 2020; Liu et al., 2020b; Hu et al., 2021). After learning each task, features of the training data can be obtained by the learned embedding function $F_\theta^e(\cdot)$, normalized under L_2-norm. In each class, several data points nearest to the mean-of-feature are selected into the memory buffer. Then, the training dataset of task $T + 1$ becomes $D_{T+1}' = D_{T+1} \bigcup D_{1:T}^{mb}$, including both new training samples D_{T+1} and replay samples $D_{1:T}^{mb} = \bigcup_{t=1}^{T} D_t^{mb}$, so as to prevent forgetting of the old tasks.

However, due to the limited storage space, only a few original training samples can be stored in the memory buffer, namely, $N_t^{mb} \ll N_t$. Although numerous memory replay approaches attempted to more effectively exploit the memory buffer, such as by alleviating the data imbalance between the old and the new training samples, this strategy of building memory buffer will lose a lot of information about the old data distribution. Several recent approaches attempted to improve the efficacy of memory replay through learning additional parameters (Liu et al., 2020b) or distilling old features (Hu et al., 2021), but resulting in huge computation cost.

4 Method

In this section, we first present memory replay with data compression for continual learning. Then, we empirically validate that there is a trade-off between the quality and quantity of compressed data, which is highly nontrivial for memory replay, and propose a novel method to determine it efficiently.

4.1 Memory Replay with Data Compression

Inspired by the biological memory replay that is in a highly compressed form (Carr et al., 2011), we propose an important baseline, that is, using data compression to increase the amount of old training samples that can be stored in the memory buffer, so as to more effectively recover the old data distribution. Data compression can be generally defined as a function $F_q^c(\cdot)$ of compressing the original data $x_{t,i}$ to $x_{q,t,i} = F_q^c(x_{t,i})$ with a controllable quality q. Due to the smaller storage cost of each $x_{q,t,i}$ than $x_{t,i}$, the memory buffer can maintain more old training samples for replay, namely,

Figure 2: Memory replay with data compression on ImageNet-sub. We make a grid search of the JPEG quality in $\{10, 25, 50, 75, 90\}$. The quality of 100 refers to the original data without compression.

$N_{q,t}^{mb} > N_t^{mb}$ for $D_{q,t}^{mb} = \{(x_{q,t,i}, y_{t,i})\}_{i=1}^{N_{q,t}^{mb}}$ in $D_{q,1:T}^{mb} = \bigcup_{t=1}^{T} D_{q,t}^{mb}$. For notation clarity, we will use D_q^{mb} to denote $D_{q,t}^{mb}$ without explicitly writing out its task label t, likewise for x_i, y_i, N_q^{mb}, N^{mb} and D. The compression rate can be defined as $r_q = N_q^{mb}/N^{mb} \propto N_q^{mb}$.

In analogy to the learning theory for supervised learning, we argue that continual learning will also benefit from replaying more compressed data, assuming that they approximately follow the original data distribution. However, the assumption is likely to be violated if the compression rate is too high. Intuitively, this leads to a trade-off between quality and quantity: if the storage space is limited, reducing the quality q of data compression will increase the quantity N_q^{mb} of compressed data that can be stored in the memory buffer, and vice versa.

Although lossless compression can perfectly reconstruct original data from compressed data, its compression rate is generally much lower than that of lossy compression (Shannon, 1948), so we mainly consider lossy compression for memory replay. Here we evaluate the proposed idea by compressing images with JPEG (Wallace, 1992), a naive but commonly-used lossy compression algorithm. JPEG can save images with a quality in the range of $[1, 100]$, where reducing the quality results in a smaller file size. Using a memory buffer equivalent to 20 original images per class (Hou et al., 2019), we make a grid search of the JPEG quality with representative memory replay approaches, such as LUCIR (Hou et al., 2019) and PODNet (Douillard et al., 2020). As shown in Fig. 2, memory replay of compressed data with an appropriate quality can substantially outperform that of original data. However, whether the quality is too large or too small, it will affect the performance. In particular, the quality that achieves the best performance varies with the memory replay approaches, but is consistent for different numbers of splits of incremental phases.

4.2 QUALITY-QUANTITY TRADE-OFF

Since the trade-off between quality q and quantity N_q^{mb} is highly nontrivial for memory replay, it needs to be carefully determined. Formally, after learning each task from its training dataset D, let's consider several compressed subsets $D_q^{mb} = \{(x_{q,i}, y_i)\}_{i=1}^{N_q^{mb}}$, where q is from a set of finite candidates $Q = \{q_1, q_2, q_3, ...\}$. Each D_q^{mb} is constructed by selecting a subset $D_q^{mb*} = \{(x_i, y_i)\}_{i=1}^{N_q^{mb}}$ of N_q^{mb} original training samples from D (following mean-of-feature or other principles). The size N_q^{mb} is determined as the maximum number such that the compressed version of D_q^{mb*} to a quality q can be stored in the memory buffer (see details in Appendix A). Thereby, a smaller q enables to save a larger N_q^{mb}, and vice versa. The objective is to select a compressed subset that can best represent D, namely, to determine an appropriate q for memory replay.

The effects of data compression to memory replay depend on the data compression approaches $F_q^c(\cdot)$, and the memory replay approaches for continual learning of the embedding function $F_\theta^e(\cdot)$. Since both of them are integrated to the feature embedding $f_{q,i} = F_\theta^e(F_q^c(x_i))$, we first calculate the feature matrix $M_q^c = [\bar{f}_{q,1}, \bar{f}_{q,2}, ..., \bar{f}_{q,N_q^{mb}}]$ of each compressed subset D_q^{mb}, where each column vector $\bar{f}_{q,i}$ is obtained by normalizing the feature embedding $f_{q,i}$ under L_2-norm to keep $||\bar{f}_{q,i}||_2 = 1$. Similarly, we obtain the feature matrix $M_q^* = [\bar{f}_1, \bar{f}_2, ..., \bar{f}_{N_q^{mb}}]$ of each original subset D_q^{mb*}. Then, we can analyze the quality-quantity trade-off in two aspects:

On the **empirical** side, in Fig. 3 we use t-SNE (Van der Maaten & Hinton, 2008) to visualize features of the original subset (light dot), which includes different amounts of original data, and its compressed subset (dark dot), which is obtained by compressing the original subset to just fit in the memory

Figure 3: t-SNE visualization of features of the original subset (light dots) and its compressed subset (dark dots) after learning 5-phase ImageNet-sub with LUCIR. From left to right, the quantity is increased from 37, 85 to 200, while the JPEG quality is reduced from 90, 50 to 10. We plot five classes out of the latest task and label them in different colors. The crossed area is out-of-distribution.

buffer. With the increase of quantity and the decrease of quality, the area of compressed subset is initially similar to that of original subset and expands synchronously. However, as a large number of low-quality compressed data occur out-of-distribution, the area of compressed subset becomes much larger than that of its original subset, where the performance also severely declines (see Fig. 2).

On the **theoretical** side, given a training dataset D, we aim to find the compressed subset D_q^{mb} that can best represent D by choosing an appropriate compression quality q from its range. To achieve this goal, we introduce $\mathcal{P}_q(D_q^{mb}|D)$ to characterize the conditional likelihood of selecting D_q^{mb} given input D under parameter q. Since the network parameter θ is fixed during compression, and the feature matrix $M_q^c = F_\theta^e(D_q^{mb})$, we rewrite $\mathcal{P}_q(D_q^{mb}|D)$ as $\mathcal{P}_q(M_q^c|D)$ equivalently. By focusing on maximum likelihood estimation (MLE), we propose to solve

$$\max_q \ \mathcal{P}_q(M_q^c|D). \tag{1}$$

We draw inspirations from determinantal point processes (DPPs), which are elegant probabilistic sampling models in data selection fields (Kulesza & Taskar, 2012). In particular, DPPs specify the probabilities for every possible subset with the volume spanned by all elements in this subset, using determinants. Depending on the nice properties of DPPs (e.g., high-quality diverse selection), our goal can be modeled as a DPP, i.e., a distribution over all subsets of D with cardinality N_q^{mb} (detailed in Appendix C.2). Formally, such a DPP formulates the probability $\mathcal{P}_q(M_q^c|D)$ as

$$\mathcal{P}_q(M_q^c|D) = \frac{\det(L_{M_q^c}(D; q, \theta))}{\sum_{|M|=N_q^{mb}} \det(L_M(D; q, \theta))}, \tag{2}$$

where $|M_q^c| = N_q^{mb}$, and $L(D; q, \theta)$ is a conditional DPP $|D| \times |D|$ kernel matrix that depends on the input D, parameters θ and q. $L_M(D; q, \theta)$ (resp., $L_{M_q^c}(D; q, \theta)$) is the submatrix sampled from $L(D; q, \theta)$ using indices from M (resp., M_q^c). The numerator defines the marginal probability of inclusion for the subset M_q^c, and the denominator serves as a normalizer to enforce the sum of $\mathcal{P}_q(M_q^c|D)$ for every possible M_q^c to 1. Generally, there are many ways to obtain a positive semi-definite kernel L. In this work, we employ the most widely-used dot product kernel function, where $L_{M_q^c} = M_q^{c\top} M_q^c$ and $L_M = M^\top M$.

However, due to the extremely high complexity of calculating the denominator in Eq. (2) (analyzed in Appendix C.1), it is difficult to optimize $\mathcal{P}_q(M_q^c|D)$. Alternatively, by introducing $\mathcal{P}_q(M_q^*|D)$ to characterize the conditional likelihood of selecting M_q^* given input D under parameter q, we propose a relaxed optimization program of Eq. (1), in which we (1) maximize $\mathcal{P}_q(M_q^*|D)$ since $\mathcal{P}_q(M_q^c|D) \leq \mathcal{P}_q(M_q^*|D)$ is always satisfied under lossy compression; and meanwhile, (2) constrain that $\mathcal{P}_q(M_q^c|D)$ is consistent with $\mathcal{P}_q(M_q^*|D)$. The relaxed program is solved as follows.

First, by formulating $\mathcal{P}_q(M_q^*|D)$ similarly as $\mathcal{P}_q(M_q^c|D)$ in Eq. (2) (detailed in Appendix C.2), we need to maximize

$$\mathcal{L}_1(q) = \mathcal{P}_q(M_q^*|D) = \frac{\det(L_{M_q^*}(D; \theta))}{\sum_{|M^*|=N_q^{mb}} \det(L_{M^*}(D; \theta))}, \tag{3}$$

where the conditional DPP kernel matrix $L(D; \theta)$ only depends on D and θ. For our task, $\mathcal{P}_q(M_q^*|D)$ monotonically increases with N_q^{mb}. Thus, optimizing $\mathcal{L}_1$ is converted into $\max_q \ N_q^{mb}$ equivalently, with significantly reduced complexity (detailed in Proposition 1 of Appendix C.3).

Second, to constrain that $\mathcal{P}_q(M_q^c|D)$ is consistent with $\mathcal{P}_q(M_q^*|D)$, we propose to minimize

$$\mathcal{L}_2(q) = \left| \frac{\mathcal{P}_q(M_q^c|D)}{\mathcal{P}_q(M_q^*|D)} - 1 \right| = \left| \frac{\det(M_q^{c\top} M_q^c)}{\det(M_q^{*\top} M_q^*)} Z_q - 1 \right| = \left| \left(\frac{\text{Vol}_q^c}{\text{Vol}_q^*} \right)^2 Z_q - 1 \right| = \left| R_q^2 Z_q - 1 \right|, \quad (4)$$

where $Z_q = \frac{\sum_{|M^*|=N_q^{mb}} \det(L_{M^*}(D;\theta))}{\sum_{|M|=N_q^{mb}} \det(L_M(D;q,\theta))}$. In particular, $\det(M_q^{*\top} M_q^*)$ has a geometric interpretation that it is equal to the squared volume spanned by M_q^* (Kulesza & Taskar, 2012), denoted as Vol_q^*, likewise for $\det(M_q^{c\top} M_q^c)$ with respect to Vol_q^c. Then we define $R_q = \frac{\text{Vol}_q^c}{\text{Vol}_q^*}$ as the ratio of the two feature volumes. To avoid computing Z_q, we can convert optimizing $\mathcal{L}_2$ into minimizing $|R_q - 1|$ equivalently, since both of them mean maximizing q (detailed in Proposition 2 of Appendix C.4).

Putting $\mathcal{L}_1$ and $\mathcal{L}_2$ together, our method is finally reformulated as

$$\begin{aligned} \max_q \; & g(q) \\ s.t., \; & q \in Q, \; |R_q - 1| < \epsilon, \end{aligned} \quad (5)$$

where ϵ is a small positive number to serve as the threshold of R_q. $g(\cdot) : \mathbb{R} \to \mathbb{N}$ represents the function that outputs the maximum number (i.e. N_q^{mb}) such that the compressed version of D_q^{mb*} to a quality q can be stored in the memory buffer. Note that we relax the original restriction of minimizing $|R_q - 1|$ by enforcing $|R_q - 1| < \epsilon$, since maximizing N_q^{mb} for $\mathcal{L}_1$ and maximizing q for $\mathcal{L}_2$ cannot be achieved simultaneously. Of note, R_q is calculated by normalizing the feature volume Vol_q^c with Vol_q^*, both of which depend on q (i.e., N_q^{mb}). Therefore, ϵ can largely mitigate the sensitivity of q to various domains and can be empirically set as a constant value (see Sec. 4.3).

Since the function $g(\cdot)$ in Eq. (5) is highly non-smooth, gradient-based methods are not applicable. Indeed, we solve it by selecting the best candidate in a finite-size set Q. Generally, the candidate values in Q can be equidistantly selected from the range of q, such as $[1, 100]$ for JPEG. More candidate values can determine a proper q more accurately, but the complexity will grow linearly. We found that selecting 5 candidate values is a good choice in our experiments. Once we solve Eq. (5), a good trade-off is achieved by reducing q as much as possible to obtain a larger N_q^{mb}, while keeping the feature volume Vol_q^c similar to Vol_q^*. This is consistent with our empirical analysis in Fig. 3.

4.3 VALIDATE OUR METHOD WITH GRID SEARCH RESULTS

In essence, the grid search described in Sec. 4.1 can be seen as a naive approach to determine the quality q, which is similar to selecting other hyperparameters for continual learning (Fig. 4, a). This naive strategy is achieved by learning each compressed subset D_q^{mb} together with the training dataset of incoming tasks for multiple epochs (Fig. 4, b), resulting in huge computation cost. In contrast, our method described in Sec. 4.2 can efficiently determine q, which only needs to calculate the feature volumes of each compressed subset D_q^{mb} and original subset D_q^{mb*}, without additional training (Fig. 4, c).

Figure 4: Determine an appropriate quality efficiently.

Now we validate the quality q determined by our method with the grid search results, where LUCIR and PODNet achieve the best performance on ImageNet-sub at the JPEG quality of 50 and 75, respectively (see Fig. 2). We present R_q in each incremental phase and the averaged R_q of all incremental phases for 5-phase split in Fig. 5 and for 10- and 25-phase splits in Appendix Fig.13. Based on the principle in Eq. (5) (we empirically set $\epsilon = 0.5$), it can be clearly seen that 50 and 75 are the qualities chosen for LUCIR and PODNet, respectively, since they are the smallest qualities

Figure 5: For 5-phase ImageNet-sub, we present R_q in each incremental phase with various qualities q, and the averaged R_q of all incremental phases.

that satisfy $|R_q - 1| < \epsilon$. Therefore, the quality determined by our method is consistent with the grid search results, but much more computation-efficient. Interestingly, for each quality q, whether $|R_q - 1| < \epsilon$ is generally consistent in each incremental phase and the average of all incremental phases. We further explore the scenarios where R_q might be more dynamic in Appendix D.4.

5 EXPERIMENT

In this section, we first evaluate memory replay with data compression in class-incremental learning of large-scale images. Then, we demonstrate the advantages of our proposal in realistic semi-supervised continual learning of large-scale object detection for autonomous driving. [2]

5.1 CLASS-INCREMENTAL LEARNING

Benchmark: We consider three benchmark datasets of large-scale images for continual learning. CUB-200-2011 (Wah et al., 2011) is a large-scale dataset including 200-class 11,788 colored images of birds with default format of JPG, split as 30 images per class for training while the rest for testing. ImageNet-full (Russakovsky et al., 2015) includes 1000-class large-scale natural images with default format of JPEG. ImageNet-sub (Hou et al., 2019) is a subset derived from ImageNet-full, consisting of randomly selected 100 classes of images. Following Hou et al. (2019), we randomly resize, crop and normalize the images to the size of 224×224, and randomly split a half of the classes as the initial phase while split the rest into 5, 10 and 25 incremental phases. We conduct single-head evaluation (Chaudhry et al., 2018) and report the averaged incremental accuracy (detailed in Appendix B.1). We further present the metric of averaged forgetting in Appendix E.3.

Implementation: The implementation is the same as representative memory replay approaches (Hou et al., 2019; Douillard et al., 2020), detailed in Appendix B.1. We mainly consider a memory buffer where the storage space is equal to 20 original images per class. We further discuss the effects of buffer size in Fig. 7, a fixed memory budget in Appendix E.4 and less compressed samples in Appendix E.5. For data compression, we apply JPEG (Wallace, 1992), a naive but commonly-used lossy compression algorithm, to compress images to a controllable quality in the range of $[1, 100]$.

Baseline: We evaluate representative memory replay approaches such as LwF (Li & Hoiem, 2017), iCaRL (Rebuffi et al., 2017), BiC (Wu et al., 2019), LUCIR (Hou et al., 2019), Mnemonics (Liu et al., 2020c), TPCIL (Tao et al., 2020a), PODNet (Douillard et al., 2020), DDE (Hu et al., 2021) and AANets (Liu et al., 2021a). In particular, AANets and DDE are the SOTA methods implemented on the backbones of LUCIR and PODNet, so we also implement ours on the two backbones. Since both AANets and DDE only release their official implementation on LUCIR, we further reproduce LUCIR w/ AANets and LUCIR w/ DDE for fair comparison.

Accuracy: We summarize the performance of the above baselines and memory replay with data compression (ours) in Table 1. For class-incremental learning, AANets and DDE are the SOTA approaches. Both of them are built on the backbones of LUCIR and PODNet. Using a naive JPEG algorithm of data compression with a properly selected quality, ours achieves the SOTA performance on the backbones of both LUCIR and PODNet, and can further boost the performance of AANets and DDE. For CUB-200-2011 with only a few training samples, all old training samples can be saved in the memory buffer with a high JPEG quality of 94. In contrast, for ImageNet-sub and ImageNet-full, only a part of old training samples can be selected, compressed and saved in the memory buffer,

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[2]All experiments are averaged by more than three runs with different random seeds.

Table 1: Averaged incremental accuracy (%) of classification tasks. The memory buffer size is equal to 20 original images per class. For CUB-200-2011, the performance of all baselines is reproduced from their officially-released code (if applicable), while for ImageNet-sub and ImageNet-full we present the reported performance of all baselines and the reproduced results of AANets and DDE, so as to make the comparison as fair as possible. The reproduced results might slightly vary from the reported results due to the usage of different random seeds. [1]With class-balance finetuning (Hu et al., 2021). [2]PODNet reproduced by Hu et al. (2021) underperforms that in Douillard et al. (2020). We present the ± standard deviation for all the reproduced results.

Method	CUB-200-2011			ImageNet-sub			ImageNet-full	
	5-phase	10-phase	25-phase	5-phase	10-phase	25-phase	5-phase	10-phase
LwF (Li & Hoiem, 2017)	$39.42_{\pm0.48}$	$38.53_{\pm0.96}$	$36.33_{\pm0.74}$	53.62	47.64	44.32	44.35	38.90
iCaRL (Rebuffi et al., 2017)	$39.49_{\pm0.58}$	$39.31_{\pm0.66}$	$38.77_{\pm0.73}$	65.44	59.88	52.97	51.50	46.89
BiC (Wu et al., 2019)	$45.29_{\pm0.88}$	$45.25_{\pm0.70}$	$45.17_{\pm0.27}$	70.07	64.96	57.73	62.65	58.72
Mnemonics (Liu et al., 2020c)	–	–	–	72.58	71.37	69.74	64.54	63.01
TPCIL (Tao et al., 2020a)	–	–	–	76.27	74.81	–	64.89	62.88
LUCIR (Hou et al., 2019)	$44.63_{\pm0.32}$	$45.58_{\pm0.28}$	$45.48_{\pm0.66}$	70.84	68.32	61.44	64.45	61.57
w/ AANets (Liu et al., 2021a)	–	–	–	72.55	69.22	67.60	64.94	62.39
w/ DDE (Hu et al., 2021)	–	–	–	72.34	70.20	–	67.51^{1}	65.77^{1}
w/ Ours	$46.68_{\pm0.60}$	$47.28_{\pm0.51}$	$48.01_{\pm0.72}$	$73.56_{\pm0.27}$	$72.70_{\pm0.47}$	$70.53_{\pm0.57}$	$67.53_{\pm0.08}^{1}$	$65.29_{\pm0.10}^{1}$
w/ AANets (Reproduced)	$46.87_{\pm0.66}$	$47.34_{\pm0.77}$	$47.35_{\pm0.95}$	$72.91_{\pm0.45}$	$71.93_{\pm0.52}$	$70.70_{\pm0.46}$	$63.37_{\pm0.26}$	$62.46_{\pm0.14}$
w/ AANets + *Ours*	$\mathbf{49.02}_{\pm1.07}$	$\mathbf{49.84}_{\pm0.87}$	$\mathbf{51.33}_{\pm1.42}$	$73.79_{\pm0.42}$	$73.73_{\pm0.37}$	$\mathbf{73.47}_{\pm0.35}$	$64.99_{\pm0.13}$	$63.04_{\pm0.11}$
w/ DDE (Reproduced)	$45.86_{\pm0.65}$	$46.48_{\pm0.69}$	$46.46_{\pm0.33}$	$73.04_{\pm0.36}$	$70.84_{\pm0.59}$	$66.61_{\pm0.68}$	$66.95_{\pm0.09}^{1}$	$65.21_{\pm0.05}^{1}$
w/ DDE + *Ours*	$47.16_{\pm0.60}$	$48.33_{\pm0.48}$	$48.37_{\pm0.34}$	$75.12_{\pm0.17}$	$73.39_{\pm0.29}$	$70.83_{\pm0.34}$	$67.90_{\pm0.05}^{1}$	$\mathbf{66.67}_{\pm0.23}^{1}$
PODNet (Douillard et al., 2020)	$44.92_{\pm0.31}$	$44.49_{\pm0.65}$	$43.79_{\pm0.44}$	75.54	74.33	68.31	66.95	64.13
w/ AANets (Liu et al., 2021a)	–	–	–	76.96	75.58	71.78	67.73	64.85
w/ DDE (Hu et al., 2021)	–	–	–	76.71	75.41	–	66.42^{2}	64.71
w/ Ours	$46.00_{\pm0.28}$	$46.09_{\pm0.37}$	$45.84_{\pm0.43}$	$\mathbf{78.08}_{\pm0.66}$	$\mathbf{76.02}_{\pm0.54}$	$72.72_{\pm0.74}$	$\mathbf{68.91}_{\pm0.16}$	$66.31_{\pm0.26}$

where our method can efficiently determine an appropriate quality of data compression for LUCIR and PODNet (see Sec. 4.3 and Appendix D.2), and for AANets and DDE (see Appendix D.3).

Computation Cost: Limiting the size of memory buffer is not only to save its storage cost, but also to save the computation cost of learning all old training samples again. Here we evaluate the computation cost of AANets, DDE and memory replay with data compression (ours) on LUCIR (see Fig. 6 for ImageNet-sub and Appendix Table 3 for CUB-200-2011). Both AANets and DDE require a huge amount of extra computation to improve the performance of continual learning, which is generally *several*

Figure 6: Averaged incremental accuracy (the column, left Y-axis) and computation cost (the triangle, right Y-axis) on ImageNet-sub. We run each baseline with one Tesla V100.

times that of LUCIR. In contrast, ours achieves competing or the largest performance improvement but only slightly increases the computation cost.

Figure 7: The effects of memory buffer size (equal to 10, 20, 40 and 80 original images per class) on ImageNet-sub. LUCIR (a) and PODNet (b) are reproduced from their officially-released codes.

Buffer Size: We further evaluate the effects of memory buffer size for memory replay with data compression (ours). As shown in Fig. 7, using the storage space equivalent to 10, 20, 40 and 80 original images per class, ours can consistently improve the performance of LUCIR and PODNet, where the improvement is generally more significant for more splits of incremental phases.

5.2 LARGE-SCALE OBJECT DETECTION

The advantages of memory replay with data compression are more significant in realistic scenarios such as autonomous driving, where the incremental data are extremely large-scale with huge storage cost. Here we evaluate continual learning on SODA10M (Han et al., 2021), a large-scale object detection benchmark for autonomous driving. SODA10M contains 10M unlabeled images and 20K labeled images of size 1920×1080 with default format of JPEG, which is much larger than the scale of ImageNet. The labeled images are split into 5K, 5K and 10K for training, validation and testing, respectively, annotating 6 classes of road objects for detection.

Figure 8: We present the grid search results (left), R_q in each incremental phase (middle), and the averaged R_q of all incremental phases (right) for SSCL on SODA10M. The quality determined by our method is 50, which indeed achieves the best performance in the grid search.

Since the autonomous driving data are typically partially-labeled, we follow the semi-supervised continual learning (SSCL) proposed by Wang et al. (2021) for object detection. Specifically, we randomly split 5 incremental phases containing 1K labeled data and 20K unlabeled data per phase, and use a memory buffer to replay labeled data. The memory buffer size is equal to 100 original images per phase. Following Han et al. (2021), we consider Pseudo Labeling and Unbiased Teacher (Liu et al., 2021b) for semi-supervised object detection (the implementation is detailed in Appendix B.2). Using the method described in Sec. 4.2 with the same threshold (i.e., $\epsilon = 0.5$), we can efficiently determine the compression quality for SSCL of object detection, and validate that the determined quality indeed achieves the best performance in grid search (see Fig. 8). Then, compared with memory replay of original data, memory replay with data compression (ours) can generally achieve *several times* of the performance improvement on finetuning, as shown in Table 2.

Table 2: Detection results with $\pm$ standard deviation (%) of semi-supervised continual learning on SODA10M. FT: finetuning. MR: memory replay.

	Method	AP	AP$_{50}$	AP$_{75}$
Pseudo Labeling	FT	40.36 $_{\pm 0.34}$	63.83 $_{\pm 0.35}$	43.82 $_{\pm 0.34}$
	MR	40.75 $_{\pm 0.30}$ / +0.39	65.11 $_{\pm 0.64}$ / +1.28	43.53 $_{\pm 0.20}$ / -0.29
	Ours	**41.50** $_{\pm 0.06}$ / **+1.14**	**65.36** $_{\pm 0.47}$ / **+1.53**	**44.95** $_{\pm 0.22}$ / **+1.13**
Unbiased Teacher	FT	42.88 $_{\pm 0.32}$	66.70 $_{\pm 0.59}$	45.99 $_{\pm 0.32}$
	MR	43.10 $_{\pm 0.06}$ / +0.22	66.88 $_{\pm 0.51}$ / +0.18	46.62 $_{\pm 0.02}$ / +0.63
	Ours	**43.72** $_{\pm 0.25}$ / **+0.84**	**67.80** $_{\pm 0.46}$ / **+1.10**	**47.36** $_{\pm 0.23}$ / **+1.37**

6 CONCLUSION

In this work, we propose memory replay with data compression, which is both an important baseline and a promising direction for continual learning. Using a naive technique of data compression with a properly selected quality, ours can achieve the SOTA performance in a computation-efficient and plug-and-play way. Since the compression quality is highly nontrivial for the efficacy of memory replay, we provide a novel method based on determinantal point processes (DPPs) to determine it efficiently, and validate our method in both class-incremental learning and semi-supervised continual learning of object detection. As data compression hardly affects the semantic information (detailed in Appendix A), a potential follow-up work is to regularize the differences between compressed data and original data, so as to further improve the performance of continual learning. In addition, our work inspires a new application of DPPs, which can provide a theoretical basis for memory replay in continual learning.

ETHIC STATEMENT

This work presents memory replay with data compression, which is an effective yet neglected baseline for continual learning. It can be used to reduce the storage requirement in memory replay and thus may facilitate large-scale applications of such methods to real-world problems (e.g., autonomous driving). As a fundamental research in machine learning, the negative consequences in the current stage are not obvious.

REPRODUCIBILITY STATEMENT

We ensure the reproducibility of our paper from three aspects. (1) Experiment: The implementation of our experiment described in Sec. 4.1, Sec. 4.3 and Sec. 5 is further detailed in Appendix B. (2) Code: Our code is included in supplementary materials. (3) Theory and Method: A complete proof of the theoretical results described in Sec. 4.2 is included in Appendix C.

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A   TRADE-OFF BETWEEN QUALITY AND QUANTITY

Given a limited memory buffer space, there is an intuitive trade-off between the quality q and quantity N_q^{mb} of compressed data. Here we plot the relation of q and N_q^{mb} for ImageNet and SODA10M used in our paper. As shown in Fig. 9, reducing the quality enables to save more compressed data in the memory buffer, and vice versa. Then, we present compressed images with various qualities in Fig. 10. Although reducing the quality tends to distort the compressed data and thus hurts the performance of continual learning, it can be clearly seen that the semantic information of such compressed data is hardly affected. There-

Figure 9: Using the same storage space as (a) 20 original images per class for ImageNet and (b) 50 original images per phase for SODA10M, we plot the relation between the quality and quantity of compressed data. The quality of 100 refers to the original data without compression.

fore, a potential follow-up work is to regularize the differences between compressed data and original data, so as to further improve the performance of memory replay.

Figure 10: Compressed images of ImageNet. (a) Each column contains compressed images of a specific JPEG quality. (b) Two exemplar images with different qualities.

B IMPLEMENTATION DETAIL

B.1 CLASS-INCREMENTAL LEARNING

Following the same implementation as Hou et al. (2019); Douillard et al. (2020); Tao et al. (2020a); Liu et al. (2021a); Hu et al. (2021), we train a ResNet-18 architecture for 90 epochs, with minibatch size of 128 and weight decay of 1×10^{-4}. We use a SGD optimizer with initial learning rate of 0.1, momentum of 0.9 and cosine annealing scheduling. We run each baseline with one Tesla V100, and measure the computation cost. We evaluate the performance of class-incremental learning with the averaged incremental accuracy (AIC) (Hou et al., 2019). After learning each phase, we can calculate the test accuracy on all classes learned so far in the previous t phases as A_t. Then we can calculate $AIC_t = \frac{1}{t} \sum_{i=1}^{t} A_i$.

To evaluate the effects of class similarity on the determined quality of data compression, we select similar or dissimilar superclasses from ImageNet, which includes 1000 classes ranked by their semantic similarity. Specifically, we select 10 adjacent classes as a superclass, and construct 10 adjacent (similar) or remote (dissimilar) superclasses. The index of classes selected in 10 similar superclasses is [1, 2, 3, ..., 100], while in 10 dissimilar superclasses is [1, 2, ..., 10, 101, 102, ..., 110, ..., 910].

B.2 LARGE-SCALE OBJECT DETECTION

For memory replay, we follow Wang et al. (2021) to randomly select the old labeled data and save them in the memory buffer, which indeed achieves competitive performance as analyzed by Chaudhry et al. (2018). We follow the implementation of Han et al. (2021) for semi-supervised object detection. Specifically, we use Faster-RCNN (Ren et al., 2015) with FPN (Lin et al., 2017) and ResNet-50 backbone as the object detection network for Pseudo Labeling (Han et al., 2021) and Unbiased Teacher (Liu et al., 2021b). For each incremental phase, we train the network for 10K iterations using the SGD optimizer with initial learning rate of 0.01, momentum of 0.9, and constant learning rate scheduler. The batch size of supervised and unsupervised data are both 16 images. For Pseudo Labeling, we first train a supervised model on the labeled set in the first 2K iterations. Then we evaluate the supervised model on the unlabeled set. A bounding box with a predicted score larger than 0.7 in the evaluation results is selected as a pseudo label, which would be used to train the supervised model in the later 8K iterations in each phase. For Unbiased Teacher, we follow the default setting and change the input size to comply with SODA10M. We run each baseline with 8 Tesla V100. The performance is evaluated by the metrics (AP, AP_{50}, AP_{75}) used in the prior works (Liu et al., 2021b; Han et al., 2021).

C THEORETICAL ANALYSIS

C.1 DETERMINANTAL POINT PROCESSES (DPPS) PRELIMINARIES

Arising in quantum physics and random matrix theory, determinantal point processes (DPPs) are elegant probabilistic models of global, negative correlations, and offer efficient algorithms for sampling, marginalization, conditioning, and other inference tasks.

A point process $\mathcal{P}$ on a ground set $\mathcal{V}$ is a probability measure on the power set 2^N, where $N = |\mathcal{V}|$ is the size of the ground set. Let B be a $D \times N$ matrix with $D \geq N$. In practice, the columns of B are vectors representing items in the set $\mathcal{V}$. Denote the columns of B by B_i for $i = 1, 2, \cdots, N$. A sample from $\mathcal{P}$ might be the empty set, the entirety of B, or anything in between. $\mathcal{P}$ is called a determinantal point process if, given a random subset Y drawn according to $\mathcal{P}$ (2^N possible instantiations for Y), we have for every $S \subseteq B$,

$$\mathcal{P}(Y = S) \propto \det(L_S), \tag{6}$$

where $S = \{B_{i_1}, B_{i_2}, \cdots, B_{i_{|S|}}\}$, $\mathcal{P}(Y = S)$ characterizes the likelihood of selecting the subset S from B. For some symmetric similarity kernel $L \in \mathbb{R}^{N \times N}$ (e.g., $L = B^T B$), where L_S is the similarity kernel of subset S (e.g., $L = S^T S$). That is, L_S is the submatrix sampled from L using indices from S. Since $\mathcal{P}$ is a probability measure, all principal minors $\det(L_S)$ of L must

be non-negative, and thus L itself must be positive semidefinite. These requirements turn out to be sufficient: any L, $0 \preceq L \preceq I$, defines a DPP.

L is often referred to as the marginal kernel since it contains all the information needed to compute the probability of any subset S being included in Y. Hence, the marginal probability of including one element B_i is $\mathcal{P}(B_i \in Y) = L_{ii}$, and two elements B_i and B_j is $\mathcal{P}(B_i, B_j \in Y) = L_{ii}L_{jj} - L_{ij}^2 = \mathcal{P}(B_i \in Y)\mathcal{P}(B_j \in Y) - L_{ij}^2$. A large value of L_{ij} reduces the likelihood of both elements to appear together in a diverse subset. This demonstrates why DPPs are "diversifying". Below is several important conclusions about DPPs related to our work (Kulesza & Taskar, 2012).

Geometric Interpretation. By focusing on L-ensembles in DPPs, determinants have an intuitive geometric interpretation. If $L = B^\top B$, then

$$\mathcal{P}(Y = S) \propto \det(L_S) = (\mathrm{Vol}(S))^2, \tag{7}$$

where the first two terms specify the marginal probabilities for every possible subset S, and the last term is the squared $|S|$-dimensional volume of the parallelepiped spanned by the items in S.

Quality vs. Diversity. While the entries of the DPP kernel L are not totally opaque in that they can be seen as measures of similarity— reflecting our primary qualitative characterization of DPPs as diversifying processes. In most practical situations, we want diversity to be balanced against some underlying preferences for different items in B. Kulesza & Taskar (2012) proposed a decomposition of the DPP that more directly illustrates the tension between diversity and a per-item measure of quality. Specifically, $L = B^\top B$ can be taken one step further, by writing each column B_i as the product of a quality term $q_i \geq 0$ and vector of normalized diversity features $f_i \in \mathbb{R}^D$, $\|f_i\| = 1$. Then, $L_{ij} = q_i f_i^\mathrm{T} f_j q_j$, where $f_i^\top f_j \in [-1, 1]$ is a signed measure of similarity between B_i and B_j. Let $\phi_{ij} = f_i^\top f_j = \frac{L_{ij}}{\sqrt{L_{ii}L_{jj}}}$, and $\phi = \{\phi_{ij}\}_{i,j=1}^N$. Then, depending on Eq. (6), we have

$$\mathcal{P}(Y = S) \propto \left(\prod_{B_{i_j} \in S} q_{i_j}^2 \right) \det(\phi_S), \tag{8}$$

where q_{i_j} can be seen as a quality score of an item B_{i_j} in B, and ϕ_S is the submatrix sampled from ϕ using indices from S.

Alternative Likelihood Formulas. In an L-ensemble DPP, the likelihood of a particular set $S \subseteq B$ is given by

$$\mathcal{P}_L(S) = \frac{\det(L_S)}{\sum_{S'} \det(L_{S'})} = \frac{\det(L_S)}{\det(L + I)}, \tag{9}$$

where S' is one of all 2^N possible subsets in B, and $L_{S'}$ is the submatrix sampled from L using indices from S'.

This expression has some nice intuitive properties in terms of volumes, and, ignoring the normalization in the denominator, takes a simple and concise form. However, as a ratio of determinants on matrices of differing dimension, it may not always be analytically convenient. Minors can be difficult to reason about directly, and ratios complicate calculations like derivatives. Such a high computation cost is just one key challenge in DPPs.

Conditional DPPs. A conditional DPP $\mathcal{P}(Y = S|B)$ is a conditional probabilistic model which assigns a probability to every possible subset $S \subseteq B$. The model takes the form of an L-ensemble:

$$\mathcal{P}(Y = S|B) \propto \det(L_S(B)), \tag{10}$$

where $L(B)$ is a positive semidefinite conditional $N \times N$ kernel matrix that depends on the input B.

k-DPPs. A k-DPP is obtained by conditioning a standard DPP on the event that the set Y has cardinality k, which is concerned only with the content of a random k-set (i.e., without the size k) (Kulesza & Taskar, 2012). Formally, the k-DPP $\mathcal{P}_L^k$ gives the probability of a particular set S as

$$\mathcal{P}_L^k(Y = S) = \frac{\det(L_S)}{\sum_{|S'|=k} \det(L_{S'})}, \tag{11}$$

where $|S| = k$ and L is a positive semidefinite kernel. Although numerous optimization algorithms have been proposed to solve DPPs problems, the high computation complexity about denominator cannot be avoided generally. On the one hand, computing denominator is a sum over $\binom{N}{k}$ terms. On the other hand, computing the determinant of each term through matrix decomposition is with $O(k^3)$ time. Just as Kulesza & Taskar (2012) claimed, sparse storage of larger matrices is possible for DPPs, but computing determinants remains prohibitively expensive unless the level of sparsity is extreme.

If the dot product kernel function is adopted to compute the kernel L in Eq. (11), similar to Geometric Interpretation of standard DPPs above, the probability $\mathcal{P}_L^k(Y = S)$ is further defined with volume sampling (Deshpande & Rademacher, 2010), i.e.,

$$\mathcal{P}_L^k(Y = S) = \frac{\det(L_S)}{\sum_{|S'|=k} \det(L_{S'})} \propto \det(L_S) = (k! \cdot (\text{Vol}(\text{Conv}(\bar{0} \cup S)))^2), \quad (12)$$

where $L = B^\top B$, $\text{Conv}(\cdot)$ denotes the convex hull, and $\text{Vol}(\cdot)$ is the k-dimensional volume of such a convex hull.

C.2 MODELING OUR CASE AS A CONDITIONAL N_q^{mb}-DPP

In this work, we aim to find the compressed subset D_q^{mb} that can best represent the training dataset D by choosing an appropriate compression quality q from its range. To achieve this goal, we introduce $\mathcal{P}_q(D_q^{mb}|D)$ to characterize the conditional likelihood of selecting D_q^{mb} given input D under parameter q. Since the network parameter θ is fixed during compression, and the feature matrix $M_q^c = F_\theta^e(D_q^{mb})$, we rewrite $\mathcal{P}_q(D_q^{mb}|D)$ as $\mathcal{P}_q(M_q^c|D)$ equivalently (the constructions of D_q^{mb} and M_q^c are detailed in Fig. 11). Depending on the nice properties of DPPs in Sec. C.1, we formulate our goal as a conditional DPP, where D is the associated ground set, and M_q^c is a desired subset in feature space. Then $\mathcal{P}_q(M_q^c|D)$ is a distribution over all subsets of D with cardinality N_q^{mb}, which is formally called a conditional N_q^{mb}-DPP, since $|M_q^c| = N_q^{mb}$ for any possible subset. Thus, such a DPP formulates the conditional probability $\mathcal{P}_q(M_q^c|D)$ as

Figure 11: Construction of D_q^{mb*}, D_q^{mb}, M_q^* and M_q^c. θ is fixed in this stage.

$$\mathcal{P}_q(M_q^c|D) = \frac{\det(L_{M_q^c}(D; q, \theta))}{\sum_{|M|=N_q^{mb}} \det(L_M(D; q, \theta))}, \quad (13)$$

where $L(D; q, \theta)$ is a conditional DPP $|D| \times |D|$ kernel matrix that depends on the input D parameterized in terms of parameters θ and q. $L_M(D; q, \theta)$ (resp., $L_{M_q^c}(D; q, \theta)$)) is the submatrix sampled from $L(D; q, \theta)$ using indices from M (resp., M_q^c).

Similarly, $\mathcal{P}_q(M_q^*|D)$ can be formulated by a conditional N_q^{mb}-DPP as

$$\mathcal{P}_q(M_q^*|D) = \frac{\det(L_{M_q^*}(D; \theta))}{\sum_{|M^*|=N_q^{mb}} \det(L_{M^*}(D; \theta))}, \quad (14)$$

where $|M_q^*| = N_q^{mb}$ (the constructions of M_q^* are detailed in Fig. 11). Unlike $L(D; q, \theta)$ above, the conditional DPP kernel matrix $L(D; \theta)$ only depends on D and θ without q, since q is fixed at its maximum, i.e., without compression. $L_{M^*}(D; \theta)$ (resp., $L_{M_q^*}(D; \theta)$) is the submatrix sampled from $L(D; \theta)$ using indices from M^* (resp., M_q^*).

Differences: Of note, two important differences between standard k-DPPs and our N_q^{mb}-DPP lie in the optimization variable and objective. For this work, we need to find an optimal set cardinality (i.e., N_q^{mb}), while it is fixed for standard k-DPPs. Although both of them can finally determine a desired subset with the maximum volume, our N_q^{mb}-DPP can uniquely determine it by N_q^{mb}, while standard k-DPPs obtain it by maximizing Eq. (11).

C.3 THE FIRST GOAL

First, we need to maximize

$$
\begin{aligned}
\mathcal{L}_1(q) = \mathcal{P}_q(M_q^*|D) &= \frac{\det(L_{M_q^*}(D;\theta))}{\sum_{|M^*|=N_q^{mb}} \det(L_{M^*}(D;\theta))} \\
&\propto \det(M_q^{*\top} M_q^*) = (\mathrm{Vol}_q^*)^2 \\
&= (N_q^{mb}! \cdot (\mathrm{Vol}(\mathrm{Conv}(\bar{0} \cup M_q^*)))^2),
\end{aligned}
\tag{15}
$$

where $|M_q^*| = N_q^{mb}$. $L(D;\theta)$ is a conditional DPP $|D| \times |D|$ kernel matrix that depends on the input D parameterized in terms of θ. $L_{M^*}(D;\theta)$ (resp., $L_{M_q^*}(D;\theta)$) is the submatrix sampled from $L(D;\theta)$ using indices from M^* (resp., M_q^*). The numerator defines the marginal probability of inclusion for the subset M_q^*, and the denominator serves as a normalizer to enforce the sum of $\mathcal{P}_q(M_q^*|D)$ for every possible M_q^* to 1. Here we employ the commonly-used dot product kernel function, where $L_{M_q^*} = M_q^{*\top} M_q^*$ and $L_{M^*} = M^{*\top} M^*$. Conv($\cdot$) denotes the convex hull, and Vol($\cdot$) is the N_q^{mb}-dimensional volume of such a convex hull.

Eq. (15) provides a geometric interpretation that the conditional probability $\mathcal{P}_q(M_q^*|D)$ is proportional to the squared volume of M_q^* with respect to N_q^{mb}, denoted as Vol_q^*. For our task, $\mathcal{P}_q(M_q^*|D)$ monotonically increases with N_q^{mb}. Thus, optimizing $\mathcal{L}_1$ is converted into $\max_q N_q^{mb}$ equivalently (detailed in Proposition 1 as below).

Proposition 1 *For any matrix $X = (\boldsymbol{x}_1, \boldsymbol{x}_2, \cdots, \boldsymbol{x}_n)$, where $\boldsymbol{x}_i \in \mathbb{R}^d$ is the i-th column of X, $d > n$ and $X^\top X \neq I$, define $\mathcal{M} = \{i_1, i_2, \cdots, i_m\}$ as a subset of $[n]$ containing m elements, and $X_{\mathcal{M}}$ as the submatrix sampled from X using indices from $\mathcal{M}$. Then for*

$$
P(\mathcal{M}) := \frac{\det\left(X_{\mathcal{M}}^\top X_{\mathcal{M}}\right)}{\sum_{|\mathcal{M}'|=m} \det\left(X_{\mathcal{M}'}^\top X_{\mathcal{M}'}\right)},
$$

we can ensure that the following two optimization programs

$$
\max_m P(\mathcal{M})
$$

and

$$
\max_m |\mathcal{M}|
$$

are equivalent.

Below is one critical lemma for the proof of Proposition 1.

Lemma 1 *For any matrix $X = (\boldsymbol{x}_1, \boldsymbol{x}_2, \cdots, \boldsymbol{x}_n)$, where $\boldsymbol{x}_i \in \mathbb{R}^d$ is the i-th column of X and $d > n$, define $\mathcal{M} = \{i_1, i_2, \cdots, i_m\}$ as a subset of $[n]$ containing m elements, and $X_{\mathcal{M}}$ as the submatrix sampled from X using indices from $\mathcal{M}$. Then for*

$$
P(\mathcal{M}) := \frac{\det\left(X_{\mathcal{M}}^\top X_{\mathcal{M}}\right)}{\sum_{|\mathcal{M}'|=m} \det\left(X_{\mathcal{M}'}^\top X_{\mathcal{M}'}\right)},
$$

we can ensure that

$$
[n] = argmax_{\mathcal{M}} P(\mathcal{M}).
$$

This means when $|\mathcal{M}| = n$, the probability $P(\mathcal{M})$ takes the maximum value.

Proof. From the definition, we have

$$
\begin{aligned}
0 \le P(\mathcal{M}) &= \frac{\det\left(X_{\mathcal{M}}^\top X_{\mathcal{M}}\right)}{\sum_{|\mathcal{M}'|=m} \det\left(X_{\mathcal{M}'}^\top X_{\mathcal{M}'}\right)} \\
&= \frac{\det\left(X_{\mathcal{M}}^\top X_{\mathcal{M}}\right)}{\det\left(X_{\mathcal{M}}^\top X_{\mathcal{M}}\right) + \sum_{\{|\mathcal{M}'|=m\} \cap \{\mathcal{M}' \neq \mathcal{M}\}} \det\left(X_{\mathcal{M}'}^\top X_{\mathcal{M}'}\right)} \le 1.
\end{aligned}
$$

If $|\mathcal{M}'| = m < n$, we have $\binom{n}{m} > 1$ possible options for subset $X_{\mathcal{M}'}$, where $\det\left(X_{\mathcal{M}'}^\top X_{\mathcal{M}'}\right) \geq 0$. Only if $|\mathcal{M}| = n$, then $\binom{n}{m} = 1$ and the probability $P(\mathcal{M})$ will definitely take the maximum value, i.e. $P(\mathcal{M}) = 1$. In addition, $X^\top X$ has full rank due to $d > n$, and then $\det(X^\top X) > 0$ always holds. That is, $P(\mathcal{M}) = 1$ can be guaranteed only when selecting all n items together. This means $[n] = \mathrm{argmax}_{\mathcal{M}} P(\mathcal{M})$. It completes the proof.

Proof of Proposition 1 From the definition in Deshpande & Rademacher (2010), we have

$$
\begin{aligned}
P(\mathcal{M}) &= \frac{\det\left(X_{\mathcal{M}}^\top X_{\mathcal{M}}\right)}{\sum_{|\mathcal{M}'|=m} \det\left(X_{\mathcal{M}'}^\top X_{\mathcal{M}'}\right)} \\
&\propto \det\left(X_{\mathcal{M}}^\top X_{\mathcal{M}}\right) \\
&= (m! \cdot (\mathrm{Vol}(\mathrm{Conv}(\bar{0} \cup \mathrm{X}_{\mathcal{M}})))^2),
\end{aligned}
\tag{16}
$$

where $\mathrm{Conv}(\cdot)$ denotes the convex hull, and $\mathrm{Vol}(\cdot)$ is the m-dimensional volume of such a convex hull. Here, from the monotonicity side, it can be easily verified that for $X_{\mathcal{M}} \cup \boldsymbol{x}_{i_{m+1}}$, where $\boldsymbol{x}_{i_{m+1}} \in \complement_X X_{\mathcal{M}}$, we have

$$
(m! \cdot (\mathrm{Vol}(\mathrm{Conv}(\bar{0} \cup \mathrm{X}_{\mathcal{M}}{}^*)))^2) < ((\mathrm{m}+1)! \cdot (\mathrm{Vol}(\mathrm{Conv}(\bar{0} \cup \mathrm{X}_{\mathcal{M}} \cup \mathrm{x}_{\mathrm{i}_{\mathrm{m}+1}})))^2),
$$

which means $P(\mathcal{M})$ is monotonically increasing with the increase of $|\mathcal{M}|$.

From the extremum side, by using Lemma 1, we can ensure that the following two optimization programs

$$
\max_m P(\mathcal{M})
$$

and

$$
\max_m |\mathcal{M}|
$$

have the same optimal value. Thus, maximizing $P(\mathcal{M})$ with respect to $|\mathcal{M}|$ can be converted into $\max_m |\mathcal{M}|$ equivalently.

C.4   THE SECOND GOAL

Second, we need to minimize

$$
\begin{aligned}
\mathcal{L}_2(q) &= \left| \frac{\mathcal{P}_q(M_q^c|D)}{\mathcal{P}_q(M_q^*|D)} - 1 \right| = \left| \frac{\det(L_{M_q^c}(D;q,\theta))}{\det(L_{M_q^*}(D;\theta))} Z_q - 1 \right| \\
&= \left| \frac{\det(M_q^{c\top} M_q^c)}{\det(M_q^{*\top} M_q^*)} Z_q - 1 \right| = \left| \left(\frac{\mathrm{Vol}_q^c}{\mathrm{Vol}_q^*}\right)^2 Z_q - 1 \right| = \left| R_q^2 Z_q - 1 \right|,
\end{aligned}
\tag{17}
$$

where $Z_q = \frac{\sum_{|M^*|=N_q^{mb}} \det(L_{M^*}(D;\theta))}{\sum_{|M|=N_q^{mb}} \det(L_M(D;q,\theta))}$. In particular, $\det(M_q^{*\top} M_q^*)$ has a geometric interpretation that it is equal to the squared volume spanned by M_q^* (Kulesza & Taskar, 2012), denoted as Vol_q^*, likewise for $\det(M_q^{c\top} M_q^c)$ with respect to Vol_q^c. Then we define $R_q = \frac{\mathrm{Vol}_q^c}{\mathrm{Vol}_q^*}$ as the ratio of the two feature volumes. To avoid computing Z_q, we can convert optimizing $\mathcal{L}_2$ into minimizing $|R_q - 1|$ equivalently, since both of them mean maximizing q (detailed in Proposition 2 as below).

Below we are going to introduce Proposition 2. We first introduce some necessary assumptions, then we present the theoretical guarantee of Proposition 2.

Assumption 1 *We make the following assumptions:*

1. ***Image dataset:*** *For any image dataset $D = (x_1, x_2, \cdots, x_n)$, where x_i is the i-th image in D, define $D_q = (x_1, x_2, \cdots, x_n)$ as a compressed image dataset with the compression quality q for D. The compression quality is bounded as $q \in Q$ (Q can be a finite/infinite set), in which the higher of q, the better of image quality.*

2. **Feature matrix:** *Denote* $X = (\boldsymbol{f}_1, \boldsymbol{f}_2, \cdots, \boldsymbol{f}_n) \in \mathcal{R}^{d \times n}$ *and* $X_q = (\boldsymbol{f}_1^q, \boldsymbol{f}_2^q, \cdots, \boldsymbol{f}_n^q) \in \mathcal{R}^{d \times n}$ *as the feature matrices of* D *and* D_q *with the same fixed feature extractor, respectively. Here,* $d > n$, $X^\top X \neq I$ *and* $X_q^\top X_q \neq I$.

3. **Selection principle:** *Define* $\mathcal{M} = \{i_1, i_2, \cdots, i_m\}$ *as a subset of* $[n]$ *containing* m *elements, and* $X_{\mathcal{M}}$ *as the submatrix sampled from* X *using indices from* $\mathcal{M}$. *Likewise, define* $\mathcal{M}_q = \{j_1, j_2, \cdots, j_m\}$ *as a subset of* $[n]$ *containing* m *elements, and* $X_{\mathcal{M}_q}$ *as the submatrix sampled from* X_q *using indices from* $\mathcal{M}_q$. *Both of them use the same selection principle.*

4. **Buffer size:** $g(\cdot) : \mathbb{R} \to \mathbb{N}$ *represents the function that outputs the maximum number (i.e.* m *) such that the compressed version of* D_q *to a quality* q *can be stored in the memory buffer. That is,* m *is uniquely determined by* q.

Proposition 2 *Under Assumption 1, for two probabilities*

$$P(\mathcal{M}) := \frac{\det\left(X_{\mathcal{M}}^\top X_{\mathcal{M}}\right)}{\sum_{|\mathcal{M}'|=m} \det\left(X_{\mathcal{M}'}^\top X_{\mathcal{M}'}\right)}$$

and

$$P(\mathcal{M}_q) := \frac{\det\left(X_{M_q}^\top X_{M_q}\right)}{\sum_{|\widetilde{\mathcal{M}}|=m} \det\left(X_{\widetilde{\mathcal{M}}}^\top X_{\widetilde{\mathcal{M}}}\right)},$$

we can ensure that the following two optimization programs

$$\min_q \left| \frac{\det\left(X_{\mathcal{M}_q}^\top X_{\mathcal{M}_q}\right)}{\det\left(X_{\mathcal{M}}^\top X_{\mathcal{M}}\right)} - 1 \right|$$

and

$$\min_q \left| \frac{P(\mathcal{M}_q)}{P(\mathcal{M})} - 1 \right|$$

are equivalent.

Below are two critical lemmas for the proof of Proposition 2.

Lemma 2 *Under Assumption 1, for a ratio*

$$R_q := \frac{\det\left(X_{\mathcal{M}_q}^\top X_{\mathcal{M}_q}\right)}{\det\left(X_{\mathcal{M}}^\top X_{\mathcal{M}}\right)},$$

we can ensure that

$$\sup Q = argmin_q \, |R_q - 1|.$$

This means when $q = \sup Q$ *(i.e, without image compression),* $|R_q - 1|$ *takes the minimum value.*

Proof. From the definition, we know $|R_q - 1| \geq 0$. To minimize $|R_q - 1|$, we need the ratio $R_q = 1$, i.e.,

$$\det\left(X_{\mathcal{M}_q}^\top X_{\mathcal{M}_q}\right) = \det\left(X_{\mathcal{M}}^\top X_{\mathcal{M}}\right). \tag{18}$$

Of note, both $X_{\mathcal{M}_q}$ and $X_{\mathcal{M}}$ are selected from the source dataset D with the same feature extractor, selection principle and set cardinality (i.e., m). That is, the only difference between them is image compression with respect to q.

By recurring to the Geometric Interpretation in k-DPPs (i.e., Eq. (12)), we have

$$\det\left(X_{\mathcal{M}_q}^\top X_{\mathcal{M}_q}\right) = (m! \cdot (\text{Vol}(\text{Conv}(\bar{0} \cup X_{\mathcal{M}_q})))^2)$$

and

$$\det\left(X_\mathcal{M}^\top X_\mathcal{M}\right) = \left(m! \cdot (\text{Vol}(\text{Conv}(\bar{0} \cup X_\mathcal{M})))^2\right).$$

Then, Eq. (18) holds only when $X_{\mathcal{M}_q}$ and $X_\mathcal{M}$ have the same convex hull. Due to great difficulty of analytically defining compression function, the two convex hull cannot be mathematically given. However, by conducting extensive experiments on this task (as shown in Fig. 3, Fig. 5, Fig. 8, Fig. 13 and Fig. 14), we find that the volume of $X_{\mathcal{M}_q}$ is larger than that of $X_\mathcal{M}$ for a specific m (i.e., q). Additionally, with q decreasing (i.e., m increasing), the volume of $X_{\mathcal{M}_q}$ is increasing more quickly. Thus, we have an empirical conclusion that the two volumes are the same only when $q = \sup Q$ (without image compression). It is reasonable since without image compression, the two selection problems about $X_{\mathcal{M}_q}$ and $X_\mathcal{M}$ are identical. This means when $q = \sup Q$, $|R_q - 1|$ takes the minimum value, which completes the proof.

Lemma 3 *Under Assumption 1, for*

$$P(\mathcal{M}) := \frac{\det\left(X_\mathcal{M}^\top X_\mathcal{M}\right)}{\sum_{|\mathcal{M}'|=m} \det\left(X_{\mathcal{M}'}^\top X_{\mathcal{M}'}\right)}$$

and

$$P(\mathcal{M}_q) := \frac{\det\left(X_{M_q}^\top X_{M_q}\right)}{\sum_{|\widetilde{\mathcal{M}}|=m} \det\left(X_{\widetilde{\mathcal{M}}}^\top X_{\widetilde{\mathcal{M}}}\right)},$$

we can ensure that

$$\sup Q = argmin_q \left|\frac{P(\mathcal{M}_q)}{P(\mathcal{M})} - 1\right|.$$

This means when $q = \sup Q$ (i.e, without image compression), $\left|\frac{P(\mathcal{M}_q)}{P(\mathcal{M})} - 1\right|$ takes the minimum value.

Proof. From the definition, we know $\left|\frac{P(\mathcal{M}_q)}{P(\mathcal{M})} - 1\right| \geq 0$. To minimize $\left|\frac{P(\mathcal{M}_q)}{P(\mathcal{M})} - 1\right|$, we need the ratio $\frac{P(\mathcal{M}_q)}{P(\mathcal{M})} = 1$. In fact,

$$\frac{P(\mathcal{M}_q)}{P(\mathcal{M})} = \frac{\det\left(X_{M_q}^\top X_{M_q}\right)}{\det\left(X_\mathcal{M}^\top X_\mathcal{M}\right)} \cdot \frac{\sum_{|\mathcal{M}'|=m} \det\left(X_{\mathcal{M}'}^\top X_{\mathcal{M}'}\right)}{\sum_{|\widetilde{\mathcal{M}}|=m} \det\left(X_{\widetilde{\mathcal{M}}}^\top X_{\widetilde{\mathcal{M}}}\right)} = R_q \cdot Z_q, \tag{19}$$

where $R_q = \frac{\det\left(X_{\mathcal{M}_q}^\top X_{\mathcal{M}_q}\right)}{\det\left(X_\mathcal{M}^\top X_\mathcal{M}\right)}$ and $Z_q = \frac{\sum_{|\mathcal{M}'|=m} \det\left(X_{\mathcal{M}'}^\top X_{\mathcal{M}'}\right)}{\sum_{|\widetilde{\mathcal{M}}|=m} \det\left(X_{\widetilde{\mathcal{M}}}^\top X_{\widetilde{\mathcal{M}}}\right)}$.

By using Lemma 2, we know that when $q = \sup Q$, $R_q = 1$ always holds. More importantly, $Z_q = 1$ also holds if $q = \sup Q$, since the numerator and denominator in Z_q will be equal without compression. That is, when $q = \sup Q$, the ratio $\frac{P(\mathcal{M}_q)}{P(\mathcal{M})} = 1$ and $\left|\frac{P(\mathcal{M}_q)}{P(\mathcal{M})} - 1\right|$ takes the minimum value. This completes the proof.

Proof of Proposition 2 From Lemma 2 and Lemma 3, we can ensure that under Assumption 1, the following two optimization programs

$$\min_q \left|\frac{\det\left(X_{\mathcal{M}_q}^\top X_{\mathcal{M}_q}\right)}{\det\left(X_\mathcal{M}^\top X_\mathcal{M}\right)} - 1\right|$$

and

$$\min_q \left|\frac{P(\mathcal{M}_q)}{P(\mathcal{M})} - 1\right|$$

have the same optimal value. Thus, maximizing $\left|\frac{P(\mathcal{M}_q)}{P(\mathcal{M})} - 1\right|$ with respect to q can be converted into $\min_q |R_q - 1|$ equivalently, where $R_q = \frac{\det\left(X_{\mathcal{M}_q}^\top X_{\mathcal{M}_q}\right)}{\det\left(X_\mathcal{M}^\top X_\mathcal{M}\right)}$.

D EMPIRICAL ANALYSIS

D.1 T-SNE VISUALIZATION OF NORMALIZED FEATURES

Figure 12: t-SNE visualization of features of different amounts of original data (dark dots) and all training data (light dots) for 5-phase ImageNet-sub with LUCIR. We randomly select five classes out of the latest task and label them in different colors.

To provide an empirical analysis of the quality-quantity trade-off and validate the theoretical interpretation, we use t-SNE (Van der Maaten & Hinton, 2008) to visualize features of all training data, the compressed subset M_q, and the original subset M_q^*. First, with the increase of quantity N_q^{mb}, the area of original data is expanded and can better cover the training data distribution, as shown in Fig. 12. This result is consistent with maximizing N_q^{mb} for $\mathcal{L}_1$. Second, with the decrease of quality q and increase of quantity N_q^{mb}, the compressed data tend to be distorted and thus become out-of-distribution, which has been discussed in the main text Fig. 3. This result is consistent with enforcing $|R_q - 1| < \epsilon$ for $\mathcal{L}_2$.

D.2 R_q FOR 5-, 10- AND 25-PHASE IMAGENET-SUB

Figure 13: We present R_q in each incremental phase with various qualities (left), and the averaged R_q of all incremental phases (right). From top to bottom are 5-, 10- and 25-phase ImageNet-Sub, respectively.

We present R_q in each incremental phase and the averaged R_q of all incremental phases for 5-, 10- and 25-phase ImageNet-sub in Fig.13. Based on the principle in Eq. (5) (we set $\epsilon = 0.5$ as the threshold of R_q), it can be clearly seen that 50 and 75 is a good quality for LUCIR and PODNet, respectively. Also, the determined quality is the same for different numbers of splits, which is consistent with the results of grid search in Fig. 2.

D.3 AANETS AND DDE

Figure 14: Determine the compression quality for AANets and DDE. We present R_q in each incremental phase (left), and the averaged R_q of all incremental phases (right) for 5-phase ImageNet-sub. The quality of 100 refers to the original data without compression.

Similar to LUCIR and PODNet, we apply the method described in Sec. 4.2 to determine the compression quality for AANets and DDE. Since both AANets and DDE only release their official implementation on LUCIR, here we focus on LUCIR w/ AANets and LUCIR w/ DDE. We present R_q in each incremental phase and the averaged R_q of all incremental phases for 5-phase ImageNet-sub in Fig. 14. The determined qualities for both LUCIR w/ AANets and LUCIR w/ DDE are consistent in different incremental phases, and are the same as that of LUCIR.

D.4 STATIC VS DYNAMIC QUALITY

For ImageNet-sub with randomly-split classes, whether $|R_q - 1| < \epsilon$ of each quality q is consistent among incremental phases and their average (see Fig. 13 and Fig. 14). Thus, the determined quality by our method usually serves as a static hyperparameter in continual learning, which has been validated by the results of gird search (see Fig. 2). Now, we wonder in which case the determined quality is dynamic, and whether using a dynamic quality is better than using a static one. Intuitively, the determined quality might be affected by the similarity of incremental classes, because the model learned to predict similar classes might be more sensitive to the image distortion caused by data compression.

Since the ImageNet classes are ranked by their semantic similarity, we select 10 adjacent classes as a superclass, and construct 10 adjacent (similar) or remote (dissimilar) superclasses (detailed in Appendix B.1). Similar to the 5-phase ImageNet-sub, we first learn 5 superlcasses as the initial phase, and then incrementally learn one superclass per phase with LUCIR. The results are presented in Fig. 15. For dissimilar superclasses, whether $|R_q - 1| < \epsilon$ is consistent in each incremental phase and their average, so the determined quality is stable at 50, validated by the results of grid search. For similar superclasses, the quality determined by the averaged R_q is 75, where using a stable quality of 75 indeed achieves the best performance in

Figure 15: Continual learning of similar or dissimilar superclasses with LUCIR. The quality of 100 refers to the original data without compression.

grid search. On the other hand, it can be clearly seen that $|R_{75} - 1| > \epsilon$ before phase 3 while $|R_{75} - 1| < \epsilon$ after it. Then we use the dynamically-determined quantity and quantity for each incremental phase. However, the dynamic quality and quantity result in severe data imbalance, so the performance is far lower than using a static quality of either 50, 75 or 90. A promising further work is to alleviate the data imbalance for the scenarios where the determined quality is highly dynamic.

E ADDITIONAL RESULT

E.1 COMPUTATION COST

In Table 3, we present the detailed results of computation cost and averaged incremental accuracy of LUCIR, LUCIR w/ AANets, LUCIR w/ DDE and LUCIR w/ Ours for CUB-200-2011 and ImageNet-sub.

Table 3: Comparison of computation cost and averaged incremental accuracy. We run each baseline with one Tesla V100.

| Methods | CUB-200-2011 | | | ImageNet-sub | | |
	5-phase	10-phase	25-phase	5-phase	10-phase	25-phase
LUCIR	1.97 h / 44.63%	3.60 h / 45.58%	8.03 h / 45.48%	8.65 h / 70.84%	10.47 h / 68.32%	15.63 h / 61.44%
LUCIR w/ AANets	3.52 h / 46.87%	5.56 h / 47.34%	14.34 h / 47.35%	21.94 h / 72.55%	40.58 h / 69.22%	91.26 h / 67.60%
LUCIR w/ DDE	7.28 h / 45.86%	13.09 h / 46.48%	31.02 h / 46.56%	55.36 h / 72.34%	61.81 h / 70.20%	79.42 h / 66.31%
LUCIR w/ *Ours*	2.70 h / 46.68%	5.00 h / 47.28%	11.00 h / 48.01%	9.09 h / 73.56%	11.27 h / 72.70%	18.83 h / 70.53%

E.2 BUFFER SIZE

In Table 4, we present the detailed results of the effects of memory buffer size.

Table 4: Averaged incremental accuracy (%) on ImageNet-sub. The storage space of the memory buffer is equal to 10, 20, 40 and 80 original images per class, respectively. The results of LUCIR and PODNet are reproduced from their officially-released codes.

	Buffer Size	10	20	40	80
	LUCIR	68.83	70.90	72.64	73.56
	w/ *Ours*	71.39 / +2.55	73.56 / +2.66	74.37 / +1.73	75.74 / +2.19
5-phase	PODNet	74.99	76.44	78.28	79.18
	w/ *Ours*	76.55 / +1.56	78.08 / +1.64	79.43 / +1.15	80.17 / +0.99
	LUCIR	64.64	68.31	70.84	72.56
	w/ *Ours*	68.13 / +3.49	72.70 / +4.39	74.11 / +3.28	76.09 / +3.54
10-phase	PODNet	70.73	73.91	76.63	78.26
	w/ *Ours*	72.78 / +2.05	76.02 / +2.11	77.82 / +1.19	79.27 / +1.01
	LUCIR	58.98	64.46	67.55	70.12
	w/ *Ours*	67.75 / +5.77	70.53 / +6.07	73.49 / +5.94	76.03 / +5.91
25-phase	PODNet	59.41	67.17	72.57	76.21
	w/ *Ours*	64.99 / +5.58	72.27 / +5.10	76.61 / +4.04	78.82 / +2.61

E.3 AVERAGED FORGETTING

In addition to averaged incremental accuracy, we evaluate the metric of averaged forgetting, which is calculated by averaging the test accuracy of each task minus its highest test accuracy achieved in continual learning (Caccia et al., 2020). In Table 5, we present the averaged forgetting of classes in the initial phase, which suffers from the most severe forgetting. It can be clearly seen that the averaged forgetting is largely alleviated by ours on each backbone.

Table 5: Averaged forgetting (%) of classes in the initial phase. The memory buffer size is equal to 20 original images per class. DDE (Hu et al., 2021) and AANets (Liu et al., 2021a) are reproduced from their officially-released code.

	CUB-200-2011			ImageNet-sub		
Method	5-phase	10-phase	25-phase	5-phase	10-phase	25-phase
LUCIR (Hou et al., 2019)	-2.86	-4.56	-4.72	-15.08	-17.32	-22.40
w/ *Ours*	-0.84	-1.08	-0.75	-14.11	-14.64	-17.96
w/ AANets (Reproduced)	-4.03	-6.58	-10.37	-11.78	-9.54	-12.86
w/ AANets + *Ours*	-1.40	-3.77	-6.82	-7.46	-8.13	-12.32
w/ DDE (Reproduced)	-1.41	-5.24	-3.91	-14.02	-17.68	-23.73
w/ DDE + *Ours*	-0.22	-0.97	-0.71	-11.24	-12.36	-17.80

E.4   FIXED MEMORY BUDGET

Here we evaluate memory replay with data compression (ours) under a fixed memory budget on ImageNet-sub. The storage space is equivalent to 2000 original images (20 original images per class × a total of 100 classes in ImageNet-sub), following a widely-used setting (Rebuffi et al., 2017; Wu et al., 2019). Under such a fixed memory budget, ours substantially boosts the performance of each backbone approach as shown in Table 6.

Table 6: Averaged incremental accuracy (%) under a fixed memory budget of 2000 original images.

	ImageNet-sub		
Method	5-phase	10-phase	25-phase
LUCIR (Hou et al., 2019)	71.87	69.37	65.72
w/ *Ours*	73.73	73.07	72.02
w/ DDE (Reproduced)	73.47	71.92	66.70
w/ DDE + *Ours*	75.35	73.93	71.17

E.5   LESS COMPRESSED SAMPLES

In Table 7, we present the results of LUCIR with different numbers of compressed samples on ImageNet-sub. Similar to the main text, we select the JPEG quality of 50 for LUCIR, where the storage space of 20 original images can save 85 such compressed images. Memory replay of 85 compressed images of quality 50 achieves a much better performance than that of 20 original images. When reducing the quantity from 85 to 70, 55 and 40, the accuracy will also decline. However, memory replay of 40 compressed images of quality 50 still outperforms that of 20 original images, where the average memory can be saved by 52.94%.

Table 7: Averaged incremental accuracy (%) of LUCIR on ImageNet-sub with different numbers of compressed data. "5-phase", "10-phase" and "25-phase" refer to the accuracy of 5-, 10- and 25-phase ImageNet-sub, respectively.

Quality	50	50	50	50	Original
Quantity	85	70	55	40	20
Total Storage	100%	82.35%	64.71%	47.06%	100%
5-phase	73.63	73.22	72.79	72.29	72.06
10-phase	72.65	72.01	71.35	70.16	68.59
25-phase	70.38	68.37	66.90	65.33	63.27