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Semi-supervised learning made simple with self-supervised clustering

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1 University of Trento  2 Inria†  3 SAP AI Research  4 Fondazione Bruno Kessler

Abstract

Self-supervised learning models have been shown to learn rich visual representations without requiring human annotations. However, in many real-world scenarios, labels are partially available, motivating a recent line of work on semi-supervised methods inspired by self-supervised principles. In this paper, we propose a conceptually simple yet empirically powerful approach to turn clustering-based self-supervised methods such as SwAV or DINO into semi-supervised learners. More precisely, we introduce a multi-task framework merging a supervised objective using ground-truth labels and a self-supervised objective relying on clustering assignments with a single cross-entropy loss. This approach may be interpreted as imposing the cluster centroids to be class prototypes. Despite its simplicity, we provide empirical evidence that our approach is highly effective and achieves state-of-the-art performance on CIFAR100 and ImageNet.

1. Introduction

In recent years, self-supervised learning became the dominant paradigm for unsupervised visual representation learning. In particular, much experimental evidence shows that augmentation-based self-supervision [3, 8–10, 14–16, 18, 21, 29, 32, 35, 71] can produce powerful representations of unlabeled data. Such models, although trained without supervision, can be naturally used for supervised downstream tasks via simple fine-tuning. However, the most suitable way to leverage self-supervision is perhaps by multi-tasking the self-supervised objective with a custom (possibly supervised) objective. Based on this idea, the community has worked on re-purposing self-supervised methods in other sub-fields of computer vision, as for instance in domain adaptation [25], novel class discovery [31, 77], continual learning [30] and semi-supervised learning [4, 13, 19, 72].

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Figure 1. Schematic illustration of the motivation behind the proposed semi-supervised framework. (a) Self-supervised clustering methods like SwAV [15] and DINO [16] compute cluster prototypes that are not necessarily well aligned with semantic categories, but they do not require labeled data. (b) Adding a linear classifier provides class prototypes, but the labeled (and unlabeled) samples are not always correctly separated. (c) Fine-tuning can help separating labeled data. (d) Our framework learns cluster prototypes that are aligned with class prototypes thus correctly separating both labeled and unlabeled data.

One of the areas that potentially benefits from the advancements in unsupervised representation learning is semi-supervised learning. This is mainly due to the fact that in semi-supervised learning it is crucial to efficiently extract the information in the unlabeled set to improve the
classification accuracy on the labeled classes. Indeed, several powerful semi-supervised methods [6, 13, 61, 72] were built upon this idea.

In the self-supervised learning landscape, arguably the most successful methods belong to the clustering-based family, such as DeepCluster v2 [14], SwAV [15] and DINO [16]. These methods learn representations by contrasting predicted cluster assignments of correlated views of the same image. To avoid collapsed solutions and group samples together, they use simple clustering-based pseudo-labeling algorithms such as k-means and Sinkhorn-Knopp to generate the assignments. A peculiar fact about this family of methods is the discretization of the feature space, that allows them to use techniques that were originally developed for supervised learning. Indeed, similar to supervised learning, the cross-entropy loss is adopted to compare the assignments, as they represent probability distributions over the set of clusters.

In this paper, we propose a new approach for semi-supervised learning based on the simple observation that clustering-based methods are amenable to be adapted to a semi-supervised learning setting: the cluster prototypes can be replaced with class prototypes learned with supervision and the same loss function can be used for both labeled and unlabeled data. In practice, semi-supervised learning can be achieved by multi-tasking the self-supervised and supervised objectives. This encourages the network to cluster unlabeled samples around the centroids of the classes in the feature space. By leveraging on these observations we propose a new framework for semi-supervised methods based on self-supervised clustering. We experiment with two instances of this framework: Suave and Daino, the semi-supervised counterparts of SwAV and DINO. These methods have several favorable properties: i) they are efficient at learning representations from unlabeled data since they are based on the top-performing self-supervised methods; ii) they extract relevant information for the semantic categories associated with the data thanks to the supervised conditioning; iii) they are easy to implement as they are based on the multi-tasking of two objectives. The motivation behind our proposal is also illustrated in Fig. 1. As shown in the figure, our multi-tasking approach enables to compute cluster centers that are aligned with class prototypes thus correctly separating both labeled and unlabeled data.

Our contributions can be summarized as follows:

- We propose a new framework for semi-supervised learning based on the multi-tasking of a supervised objective on the labeled data and a clustering-based self-supervised objective on the unlabeled samples;
- We experiment with two representatives of such framework: Suave and Daino, semi-supervised extensions of SwAV [15] and DINO [16]. These methods, while simple to implement, are powerful and efficient semi-supervised learners;
- Our methods outperform state-of-the-art approaches, often relying on multiple ad hoc components, both on common small scale (CIFAR100) and large scale (ImageNet) benchmarks, setting a new state-of-the-art on semi-supervised learning.

2. Related works

Self-supervised learning. The self-supervised literature has rapidly become extremely vast [38]. Excluding a recent trend involving denoising autoencoders [59] combined with vision transformers [27] like in MAE [34], the vast majority of existing methods are still relying on multiple views derived from each sample via data augmentation. These augmentation-based methods [3,8–10,14–16,18,21,29,32,35,71] are capable to learn rich representations during unsupervised pre-training, achieving performance comparable to their supervised counterparts when fine-tuned with the labels. Basically, these methods enforce correlated views of the same input to have coherent representations in latent space such that the model becomes invariant to the augmentations applied. This corresponds to maximizing the mutual information between views’ representations and, in the literature, it has been done using different loss functions.

Contrastive-based methods [8,18,35,37] define a loss based on noise-contrastive estimation [33] as instance discrimination [63] between positive (correlated) and negative views (the remaining samples in the mini-batch). A major drawback of these methods is that they require large mini-batches to have representative negative samples. To overcome this difficulty, consistency-based methods like [21,32] propose to maximize the cosine similarity of positive pairs without considering negatives, whereas redundancy-reduction-based methods [9,10,29,71] employ principled regularization terms to minimize features redundancy. For instance, in [71] a loss is introduced to minimize the cross-correlation between features of the positive pairs. Similarly, in [9,10] the features learning process minimizes the covariance alongside regulating the variance of the embeddings.

Clustering-based methods [3,14–16,45], instead, naturally discretize the latent space via clustering. They perform clustering either offline (using the whole dataset), as in DeepCluster v2 [14], PCL [45] and SeLa [3], or online (using mini-batches), as in SwAV [45] and DINO [16], and impose coherent clustering assignments of positive pairs through a cross-entropy. Noticeably, this loss contrasts targets (assignments) and predictions as in the standard supervised learning objective. Nonetheless, surprisingly, to the best of our knowledge there have been no previous attempts in the literature to exploit this favorable property in the semi-supervised scenario. Our work aims to fill this gap.
3. Clustering-based semi-supervised learning

Clustering-based self-supervision is typically adopted in unsupervised representation learning scenarios where no labeled data is available [14–16]. However, in this work, we aim to take advantage of the few annotations available in the semi-supervised setting to learn even better representations. Our main intuition is to replace the cluster centroids with class prototypes learned with supervision. In this way, unlabeled samples will be clustered around the class prototypes, guided by the self-supervised clustering-based objective. To this end, we jointly optimize a supervised loss on the labeled data and a self-supervised loss on the unlabeled data. It turns out that using the same loss function (cross-entropy) is a very good choice since it promotes synergy between the two objectives and eases up the implementation. In the following, we formalize self-supervised clustering in Section 3.1. Then, we describe the details of our novel semi-supervised learning framework in Section 3.2 and show its application through two popular self-supervised approaches in Section 3.3.

3.1. Clustering-based self-supervised learning

Given an unlabeled dataset \( D_u = \{x_u^1, \ldots, x_u^N\} \), two correlated views of the same input image, \((x, \hat{x})\), are generated via data augmentation and embedded through an encoder network \( f_0 \), composed of a backbone \( g \), a projector \( h \) and a set of prototypes (or centroids) \( p \) implemented as a bias-free linear layer. Performing the forward pass of the backbone and the projector produces two latent representations, \((h, \hat{h})\) for the two correlated views \((x, \hat{x})\), respectively. Subsequently, the cluster centroids \( p \) are used to produce two sets of logits \((\hat{p}, \hat{p})\) corresponding to each of the two latent representations. A softmax function \( \sigma \) can be applied to these logits to obtain probability distributions over the set of clusters, also referred to as cluster assignments.

In principle, the two assignments could be immediately compared in order to encourage the network to output similar predictions for similar inputs (correlated views). However, this may lead to degenerate solutions where all samples are assigned to the same cluster. To avoid collapse, simple clustering techniques are usually employed to embed priors into the cluster assignment process and regularize the training procedure. In practice, we compute:

\[ \hat{y} = \delta(\hat{p}, c, \epsilon), \]

where \( \delta \) is the clustering technique, which takes as input some predicted logits \( \hat{p} \), a context \( c \) and a temperature-like parameter \( \epsilon \) that usually regulates the entropy of the assignment \( \hat{y} \) (sometimes addressed as pseudo-label). The context \( c \) can be implemented in different flavors depending on the nature of \( \delta \). For instance, for offline clustering \( c \) is represented by the features of all the samples in the dataset, while
Pseudo-code for Suave and Daino.

```python
# x_l, x_u: labeled and unlabeled samples
# label: one-hot ground truth for x_l
def training_step(x_l, x_u, label):
    # correlated views
    x1_u, x2_u = aug(x_u), aug(x_u)
    # forward pass
    p_l = f(x_l)
    p1_u, p2_u = f(x1_u), f(x2_u)
    # compute assignments via clustering
    assign = delta(p2_u.detach())
    # concatenate predictions and targets
    pred = cat(p_l, p1_u)
    target = cat(label, assign)
    # compute loss
    loss = cross_entropy(pred, target)
    return loss
```

Figure 2. Left: overview of the proposed family of semi-supervised methods. Right: PyTorch-like pseudo-code for Suave and Daino.

for online clustering the context might contain the features of the current batch or simply a running mean of the overall distribution. The pseudo-label can be categorical (hard) or soft. Empirically soft cluster assignments were shown to yield superior performance [15].

The output of the clustering technique is then used as a target in the cross-entropy loss:

$$\ell(s, \hat{y}) = - \sum_{k=1}^{K} \hat{y}_k \log (s_k), \tag{2}$$

where $s = \sigma(p/\tau)$ has gone through softmax normalization with temperature $\tau$, and $K$ is the number of clusters. Note that a stop-gradient operation is performed, so that the gradient is not propagated through the pseudo-label. It is worth mentioning that the cross-entropy loss in Eq. 2 is asymmetric, but it can be made symmetric by swapping $x$ with $\hat{x}$.

Intuitively, this loss leverages cluster assignments as a proxy to minimize the distance between latent representations $(h, \hat{h})$ of augmented views of the same image. As a by-product, the objective also learns a set of cluster prototypes encoded in the last linear layer $p$, which represent semantic information in the latent space. However, these clusters have no guarantee to be aligned with the true semantic categories represented in the dataset. Nonetheless, the discretization of the feature space is particularly interesting in the context of semi-supervised learning, as described in the following.

### 3.2. Our semi-supervised learning framework

In the semi-supervised scenario, we assume having access to a partially labeled dataset, $D = D_u \cup D_l$, usually with $|D_l| \ll |D_u|$ and $D_l$ contains a number of $C$ known classes. We propose to exploit clustering-based self-supervised models described in Sec. 3.1 as a base to extract information from unlabeled data, and we extend them to take advantage of the labeled samples.

As mentioned, the main drawback of clustering-based self-supervised methods in the context of semi-supervised learning is that there are no guarantees that the stochastic optimization process will organize the clusters in the feature space according to the class labels. Indeed, they may be completely misaligned with the actual distribution of the classes. This also potentially hinders the effectiveness of the representations, as some prototypes may encode spurious correlations in the data. An ideal scenario is one where the clusters are centered on the actual class centroids such that the label can be propagated to the unlabeled samples by means of the clustering function. This will generate positive feedback that progressively transfers information from the labeled set into the unlabeled one, thus improving the feature representations learned by the network.

We propose to condition the cluster prototypes to encode the class information by resorting to multi-tasking of the self-supervised and supervised objectives. This can be achieved by optimizing the same loss function in Eq. 2 while replacing the pseudo-label with the ground-truth label when available:

$$\ell(s, \bar{y}) = \begin{cases} y, & x \in D_l \\ \hat{y}, & x \in D_u. \end{cases} \tag{3}$$

Since the linear layer $p$ that contains the prototypes is now shared between the two objectives, we set $K = C$ to have a matching label space, although in principle this is not a hard constraint (as shown in [31]). In a nutshell, in our framework we compute the forward pass for both labeled and unlabeled samples, concatenate the associated predictions and the targets and apply the cross-entropy loss simultaneously as described in Fig. 2. We empirically demonstrate in Sec. 5
that, despite its simplicity, our framework equipped with these design choices is a strong semi-supervised learner.

3.3. Suave and Daino

Our proposed framework described above can convert any clustering-based self-supervised method into a semi-supervised learner, without dropping, adding or replacing any architectural components and reusing the same loss function. We select two representative self-supervised methods to showcase our framework: SwAV [15] and DINO [16], whose semi-supervised extensions we name Suave and Daino. This choice is motivated by their superior representation learning capabilities and ease of use. In particular, both are online clustering methods, which means that the cluster assignments can be computed on-the-fly without accessing the whole dataset simultaneously. This is a great advantage, especially for large scale datasets. For these reasons, we discard offline clustering methods like DeepCluster [14] and PCL [45], while in principle our approach can be also applied to them.

Suave. SwAV [15], following [3], casts the pseudo-label assignment problem as an instance of the optimal transport problem and proposes the swapped prediction task where the assignment of a view is predicted from the representation of another view. Simply put, SwAV generates pseudo-labels such that each cluster is approximately equally represented in the current batch, preventing the network from falling into degenerate solutions. This is especially convenient as we only need the information in the current batch for clustering. In light of our proposed framework, reusing Eq. 1, in Suave the target for the first sample in the batch can be obtained as follows (the same reasoning can be trivially applied to the other samples in the batch):

$$\hat{y}_1 = \delta(\hat{p}_1, [\hat{p}_2, \ldots, \hat{p}_B], \epsilon)$$

where the context $c = [\hat{p}_2, \ldots, \hat{p}_B]$ contains all the logits in the batch except $\hat{p}_1$. Now we define $\hat{P} = [\hat{p}_1, \hat{p}_2, \ldots, \hat{p}_B]$ and $Y = [\hat{y}_1, \hat{y}_2, \ldots, \hat{y}_B]^T$, where $Y$ is the matrix that holds the unknown pseudo-labels of the whole batch. The clustering function $\delta$ will return the first column of $Y$ that is found by solving:

$$\hat{Y} = \max_{Y \in \Gamma} \text{Tr}(Y \hat{P}) + \epsilon H(Y),$$

where $\epsilon > 0$ is the temperature-like parameter mentioned in Eq. 1, $H$ is the entropy function, $\text{Tr}$ is the trace operator, and $\Gamma$ is the transportation polytope defined as:

$$\Gamma = \{ Y \in \mathbb{R}_{++}^{C \times B} \mid Y1_B = \frac{1}{C}1_C, Y^\top 1_C = \frac{1}{B}1_B \}.$$  

These constraints enforce that, on average, each cluster is selected $\frac{B}{C}$ times in each batch, automatically ensuring de-biased assignments. The solution to Eq. 5 is obtained using the Sinkhorn-Knopp algorithm [1, 23].

Daino. DINO [16] aims at further simplifying the pipeline described above. Instead of using optimal transport on the predictions of the current batch it uses two practical tricks to avoid collapse: a momentum encoder and a pseudo-labeling strategy based on centering and sharpening. A momentum encoder is a “slow” version of the encoder $f_{\theta}$ updated using exponential moving average (EMA). After each gradient step on $\theta$, the parameters $\phi$ of the momentum encoder $f_{\phi}$ are updated as follows:

$$\phi \leftarrow \eta \phi + (1 - \eta)\theta,$$

where $\eta$ is a rate parameter. The rationale behind this choice is that the momentum encoder serves as a teacher producing more stable representations throughout training, improving the optimization process. The teacher is used at every iteration to generate the logits $\tilde{p} = f_{\phi}(\tilde{x})$ which in turn are input to the clustering function $\delta$ in Eq. 1 to obtain the target assignment for Daino:

$$\hat{y} = \delta(\tilde{p}, \gamma, \epsilon) = \sigma \left( \frac{\tilde{p} - \gamma}{\epsilon} \right),$$

where $\sigma$ and $\epsilon$ are the softmax function and a temperature coefficient (here used to sharpen the distribution) respectively. The context $c = \gamma$ in this case is a centering vector that approximates and de-biases the overall distribution of the data over the clusters and is also updated using EMA:

$$\gamma \leftarrow \mu \gamma + (1 - \mu) \frac{1}{B} \sum_{b=0}^{B} \tilde{p}_b,$$

where $\mu$ adjusts the rate of the update. In brief, centering prevents one dimension to dominate but encourages high-entropy outputs, while sharpening does the opposite. Empirical evidence shows that this is enough to avoid collapse.

4. Implementation details

Architectures. For large-scale datasets (ImageNet), we adopt ResNet50 [36] and ViT-S/16 [27] backbones. For small-scale datasets (CIFAR100) we train a Wide ResNet (WRN-28-8) [70]. The convolutional backbone is followed by a projection head consisting of a multi-layer perceptron (with batch normalization in the hidden layers). We use 2 layers as in [6, 15]. After that, we perform L2-normalization and compute the predictions using a bias-free L2-normalized linear layer corresponding to the prototypes. We set the number of prototypes equal to the number of classes of the dataset at hand. Moreover, we perform an online linear evaluation at different depths of the network to identify which layer learns the best representations; we append a detached classification head on top of the backbone, the first and the second layer of the projection. We remark that such heads do not impact the efficiency of the training.
Semi-supervised pre-training. We pre-train our models using LARS [68] optimizer with linear warmup plus cosine learning rate schedule. Also, we adopt weight decay regularization. The backbone layers of the models can be initialized with self-supervised checkpoints of SwAV and DINO. At each training iteration, we sample a mini-batch composed of unlabeled images\(^1\) and labeled images. Note that we count a training epoch considering a full pass over the unlabeled dataset. We optimize the cross-entropy loss of Eq. 3, re-weighting the labeled and unlabeled terms by their frequency in the batch. Moreover, we soften the supervised targets with label smoothing to mitigate overfitting. For the pseudo-labeling, in Suave, we re-use the same parameters for the Sinkhorn-Knopp as in SwAV; in Daino, we inherit the hyperparameters for the centering and the momentum encoder, whereas we tune the sharpening coefficient. More details are in the supplementary material.

Data augmentation. Images are augmented differently based on whether they are unlabeled or labeled. For the unlabeled images, we follow the default self-supervised augmentations of SwAV and DINO, while for the labeled, we adopt lighter augmentation Inception-style (random crop and flip) [56] and color distortion (jittering and greyscale). Note that it is important not to over-distort the labeled images as they are needed to align the clusters and classes.

To boost the self-supervised feature learning, we employ the multi-crop [15] augmentation scheme for unlabeled images. From each input image, we derive two global views from larger and higher-resolution crops and multiple smaller views from tighter crops. This is common practice in self-supervised learning. Similar to SwAV and DINO, we compute a clustering assignment only for the global views and use them as targets for the smaller ones. All the multi-crop views are taken into account when weighting the loss.

Another augmentation technique we empirically found to be useful is based on the combination of CutMix [69] and MixUp [74]. We apply it to both unlabeled (global views only) and labeled images, but separately. For the unlabeled images, we interpolate the learned clustering assignments due to the lack of ground-truth labels. This augmentation allows for shifting decision boundaries to low-density regions of the data [12, 58]. We mix the whole batch at every iteration, but to not over-regularize, we concatenate the mixed images to the current batch, instead of substituting it.

Semi-supervised fine-tuning. Since self-supervised methods are trained with strong augmentations, it is preferable after pre-training to fine-tune them to slightly improve the performance. We discover that a semi-supervised fine-tuning recipe works better than the typical fully supervised fine-tuning e.g., [4]. In practice, we just keep training the model with the same objective (our semi-supervised clustering-based loss) for a few more epochs, while relaxing some of the stronger augmentations adopted during pre-training, i.e., disable multi-crop and color distortions.

5. Experiments

5.1. Experimental protocol

Datasets. We perform our experiments using two common datasets, i.e., CIFAR100 [41] and ImageNet-1k [26]. In the semi-supervised setting the training set of each of these datasets is split into two subsets, one labeled and one unlabeled. For CIFAR100, which is composed of 50K images equally distributed into 100 classes, we investigate three splits as in [55], retaining 4 (0.8%), 25 (5%), and 100 (20%) labels per class, resulting in total to 400, 2500, and 10000 labeled images, respectively. For ImageNet-1k (~1.3K images per class, 1K classes), we adopt the same two splits of [19] using 1% and 10% of the labels. In both datasets, we evaluate the performance of our method by computing top-1 accuracy on the respective validation/test sets.

Baselines. We compare our methods, Suave and Daino, with state of the art methods from the semi-supervised literature (see Section 2). In particular, we compare against hybrid consistency regularization methods like SimMatch [76] and ConMatch [40] (and others [46, 60, 73]), and methods that are derived from self-supervised approaches, like PAWS [6] and S4L-Rot [72]. We also compare with recent debiasing-based pseudo-labeling methods like DebiasPL [62]. For the sake of fairness, we leave out methods using larger architectures or pre-trained on larger datasets, e.g., DebiasPL with CLIP [52] and SimCLR v2 [19].

5.2. Results

First, we demonstrate the effectiveness of our semi-supervised framework in a small-scale dataset using the CIFAR100 benchmark. Then, we evaluate our models at large-scale on ImageNet-1k (see Section 5.2.1). Finally, we ablate the different components of our models (see Section 5.2.2).

5.2.1 Comparison with the state of the art

Results on CIFAR100. Table 1 shows a comparison between our methods and several semi-supervised approaches in the literature. In particular, we compare to consistency regularization semi-supervised methods (see Section 2) like ConMatch [40] and SimMatch [76], which are the strongest methods on this dataset. First, from the table we observe that both Suave and Daino achieve high performance in all the three splits (400, 2500, 10000). Daino obtains results comparable to the best competitors, while Suave outperforms all the baselines and beats the best methods CCSSL [67] and SimMatch [76] by +2.4%p, +1.3%p, +1.5%p in the three settings, respectively. Overall, these results clearly demonstrate how our clustering-based semi-
supervised learning methods achieves state-of-the-art performance without requiring any ad-hoc confidence thresholds for pseudo-labels as in most recent consistency regularization methods. Interestingly, comparing Suave and Daino with their self-supervised counterparts, SwAV and DINO, a remarkable improvement is achieved. In fact, SwAV and DINO obtain an accuracy of 64.9% and 66.8% [24], respectively, when linearly evaluated using 100% of the labels after self-supervised pre-training.

**Results on ImageNet-1k.** We also perform large scale experiments on ImageNet-1k considering the label splits of 1% and 10% as common in previous works [55, 76]. Due to limited computational resources, we primarily focus on Suave with a ResNet50 backbone, which enables us to compare with most of the related state-of-the-art methods. In addition, we also provide experimental evidence that our framework works well with a different clustering algorithm (Daino), backbone (ViT [27]), and a simpler training recipe (described in the supplementary material).

We compare against three families of methods, self-supervised-inspired semi-supervised approaches [6, 72], consistency regularization methods [13, 44, 50, 55, 64, 76], and debiasing-based methods [62], and present results in Table 3, Table 2, and Figure 3. To provide full context to our results, in Table 3, we also report the performance of self-supervised models when simply fine-tuned with labels. Suave and Daino obtain performances that are comparable with state-of-the-art methods on ImageNet-1k. In particular, when compared against related methods (e.g., DebiasPL, SimMatch, FixMatch-EMAN, CoMatch) with similar batch-size and number of epochs, Suave obtains the best score (+0.6%p) on the 10% setting and the third best score on the 1% setting, -1.0%p from the best baseline SimMatch [76]. Importantly, other approaches like PAWS [4] require larger batches to obtain comparable results. This behavior can be ascribed to the fact that PAWS adopts a k-nearest neighbor approach on the labeled instances to generate the assignment vectors (pseudo-labels), and thus it requires a sufficiently high number of labeled examples to well rep-
Table 4. Impact of our multi-task training strategy. Legend: UPT = “unsupervised pre-training”, MT = “multi-tasking” and SL = “same loss for all training phases”.

<table>
<thead>
<tr>
<th>Method</th>
<th>UPT</th>
<th>MT</th>
<th>SL</th>
<th>Epochs</th>
<th>Acc@1</th>
</tr>
</thead>
<tbody>
<tr>
<td>SwAV</td>
<td>✓</td>
<td>✓</td>
<td>✓</td>
<td>800</td>
<td>70.2</td>
</tr>
<tr>
<td>SwAV (repro)</td>
<td>✓</td>
<td>✓</td>
<td>✓</td>
<td>800</td>
<td>72.3</td>
</tr>
<tr>
<td>SwAV+CT [4]</td>
<td>✓</td>
<td>✓</td>
<td>✓</td>
<td>(400)30</td>
<td>70.8</td>
</tr>
<tr>
<td>Suave</td>
<td>✓</td>
<td>✓</td>
<td>✓</td>
<td>(800)100</td>
<td>75.0</td>
</tr>
</tbody>
</table>

Table 5. Impact of our semi-supervised fine-tuning strategy.

<table>
<thead>
<tr>
<th>Method</th>
<th>Labels (%)</th>
<th>Semi-supervised pre-training</th>
<th>Supervised fine-tuning</th>
<th>Semi-supervised fine-tuning</th>
</tr>
</thead>
<tbody>
<tr>
<td>Suave</td>
<td>1%</td>
<td>64.1</td>
<td>64.8</td>
<td>66.2</td>
</tr>
<tr>
<td></td>
<td>10%</td>
<td>73.4</td>
<td>74.8</td>
<td>75.0</td>
</tr>
</tbody>
</table>

Table 6. Online linear evaluation on the projector layers. We attach a linear layer (preceded by a stop-grad) after each projector layer and train it with labeled data.

<table>
<thead>
<tr>
<th>Method</th>
<th>Labels (%)</th>
<th>Backbone</th>
<th>Projection layer 1</th>
<th>Projection layer 2</th>
</tr>
</thead>
<tbody>
<tr>
<td>Suave</td>
<td>1%</td>
<td>65.2</td>
<td>66.2</td>
<td>65.1</td>
</tr>
<tr>
<td></td>
<td>10%</td>
<td>74.2</td>
<td>75.0</td>
<td>74.8</td>
</tr>
</tbody>
</table>

Figure 3. Comparison with related works on the training efficiency in terms of pre-training epochs (top) and size of the mini-batches (bottom). Plotted results refer to the ImageNet 10% split.

represent the class distributions. In contrast, Suave generates the assignments using the learnable cluster/class prototypes, which are a fixed number independent of the batch size.

Figure 3 (top) compares Suave with the best methods in Table 3, highlighting the fast convergence of our method. Performing 100 epochs of semi-supervised pre-training is enough to achieve comparable results to other related methods like FixMatch-EMAN, PAWS, and SimMatch, which require at least twice the semi-supervised epochs. It is worth noting that Suave naturally benefits from self-supervised initialization, as it basically shares the objective with the self-supervised counterpart SwAV. Indeed, by looking at Table 3 we can observe that better SwAV checkpoints lead to higher accuracy. At the same time, the difference between Suave (SwAV-100) and Suave (SwAV-800) is rather small, 1.4%p and 2.4%p on 10% and 1%, respectively.

5.2.2 Ablation study

Here, we assess the importance of the main components of our method: (i) the multi-task training objective, (ii) the adopted fine-tuning strategy, and (iii) the quality of representations at different depths of the network.

In Table 4, we demonstrate the importance of our multi-tasking strategy. Suave significantly outperforms vanilla SwAV, even in its improved version, i.e., SwAV (repro) obtained with the fine-tuning protocol borrowed from PAWS [6]. More importantly, when comparing SwAV to SwAV+CT (SwAV where the self-supervised loss is multi-tasked with a supervised contrastive loss [39]) we still observe that our method is far superior. This clearly indicates that it is better to exploit the available labels to condition the prototypes rather than using them to sample positive instances for contrastive training.

Table 5 provides evidence that fine-tuning Suave with our semi-supervised fine-tuning strategy (see Section 4) is more effective than adopting the classical fully supervised recipe. We believe that feeding the model with unlabeled data during the fine-tuning phase allows the model to better fit the real distribution. As shown in the table, the semi-supervised fine-tuning recipe enables a larger improvement over the performance after semi-supervised pre-training.

Finally, Table 6 shows how the quality of the learned representations changes at different depths of the projector. It turns out that regardless of the label split, we obtain the most discriminative representations at the first layer of the projection. This is consistent with what was observed by other works that use similar augmentations (e.g., PAWS [6]). We ascribe this behavior to the fact that the last layer tries to build as much invariance as possible to strong augmentations, which hurts the classification accuracy.

6. Conclusion

We have presented a novel approach for semi-supervised learning. Our framework leverages from clustering-based self-supervised methods and adopts a multi-task objective, combining a supervised loss with the unsupervised cross-entropy loss typically adopted for clustering assignments in [15, 16]. Despite its simplicity, we demonstrate that our approach is highly effective, setting a new state-of-the-art for semi-supervised learning on CIFAR-100 and ImageNet.

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