Scalable Bio-Inspired Training of Deep Neural Networks with FastHebb

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Abstract

Recent work on sample efficient training of Deep Neural Networks (DNNs) proposed a semi-supervised methodology based on biologically inspired Hebbian learning, combined with traditional backprop-based training. Promising results were achieved on various computer vision benchmarks, in scenarios of scarce labeled data availability. However, current Hebbian learning solutions can hardly address large-scale scenarios due to their demanding computational cost. In order to tackle this limitation, this contribution develops a novel solution by reformulating Hebbian learning rules in terms of matrix multiplications, which can be executed more efficiently on GPU. We experimentally show that the proposed approach, named FastHebb, accelerates training speed up to 70 times, allowing us to gracefully scale Hebbian learning expriments on large datasets and network architectures such as ImageNet and VGG.

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Keywords: Hebbian Learning, Deep Learning, Neural Networks, Biologically Inspired

1. Introduction

Recent efforts from the research community focused 2 on the development of biologically plausible alterna-3 tive to the backpropagation algorithms for Deep Neural Network (DNN) training. Biological constraints re-5 quire neurons to use only locally available information 6 to compute the weight updates, and neuroscientific observations suggest that synaptic plasticity follows the 8 Hebbian model [1, 2]. In simple terms, the weight up-9 date should be proportional to the input on the respec-10 tive synapse and the neuron output at a given point in 11 time. The study of biologically realistic learning models 12 is interesting both because they are well suited for neu-13 romorphic applications [3, 4], and for the perspective 14 to better understand the mechanisms behind biological 15 intelligence and use them to enhance current Artificial 16 Intelligence (AI) technologies. 17

Among the recently proposed bio-inspired learning approaches, Contrastive Hebbian Learning (CHL) [5] and Equilibrium Propagation (EP) [6] leverage recurrent architectures with Hebbian and anti-Hebbian

*Corresponding author Email addresses: gabriele.lagani@isti.cnr.it (Gabriele Lagani), fabrizio.falchi@isti.cnr.it (Fabrizio Falchi), claudio.gennaro@isti.cnr.it (Claudio Gennaro), hannes.fassold@joanneum.at (Hannes Fassold), giuseppe.amato@isti.cnr.it (Giuseppe Amato) phases, showing that the resulting update steps approximate backprop. More recently, the Forward-Forward (FF) approach has been proposed [7] for feedforward networks, which is also based on an alternation between two phases. While the approaches mentioned above focus on supervised learning solutions, a lot of attention on bio-inspired methods has converged on unsupervised learning. For example, the Similarity Matching criterion [8, 9, 10, 11] or the Hebbian PCA rule [12, 13] allow neurons to learn to extract the principal components from data. Similarly, Hebbian learning with Winner-Takes-All (WTA) competition allows neurons to find clusters in the data space [14, 15, 16, 17, 18, 19, 20]. This reveals interesting connections between the Hebbian theory of learning and data science.

In this work, we focus on a hybrid solution of unsupervised Hebbian learning and supervised backprop training, which are combined together in a semisupervised fashion. In fact, supervised training alone has the disadvantage of requiring numerous training samples to achieve high performances, but the latter are often expensive to gather, requiring a consistent manual effort. To circumvent this issue, a possible direction is to pre-train the model on a large amount of unlabeled data, with an unsupervised algorithm, and then fine-tune with supervision on a small labeled dataset [21, 22]. In this scenario, recent work has shown superior performance of Hebbian-based semi-supervised

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50 training, compared to other unsupervised methods for 102

51 pre-training, such as Variational Auto-Encoders (VAE) 103

 $_{52}$ [23], especially in scenarios where the available labeled $_{104}$

⁵³ data is scarce [13, 17]. Due to the difficulty of collect-⁵⁴ ing labeled data, these scenarios are of strong practical ⁵⁵ interest.

Despite their promising results, current Hebbian 108 learning solutions can hardly be used to address large-109 57 scale problems, due to their demanding computational 110 58 cost. In this perspective, the goal of our contribution is 59 to address the performance limitations of Hebbian algo-111 60 rithms. For this purpose, we developed a new Hebbian 112 61 learning solution, named *FastHebb*, which is designed 113 62 to better take advantage of GPU acceleration. This is 114 63 done in two steps. First, we notice that Hebbian learning 64 with mini-batch processing evolves in two stages, one 116 65 is the weight update computation for each sample in the 66 117 mini-batch, and the other is the aggregation of updates 67 over all the minibatch elements. These two phases can 118 68 be merged together with a significant speedup. Second, 119 69 the resulting Hebbian equations of synaptic updates can 120 70 121 translated in terms of matrix multiplications, which can 71 122 be executed very efficiently on GPU. 72

In order to provide an experimental evaluation of the 123 73 proposed method, we used established computer vision 124 74 benchmarks such as CIFAR10/100 [24], Tiny ImageNet 75 [25] and ImageNet [26]. Besides the image classifi-76 cation evaluation, we also studied the performance of 125 77 Hebbian neural features for Content-Based Image Re-78 trieval (CBIR). We considered sample efficient learn-79 126 ing scenarios, where label information is assumed to 80 be available only for a certain percentage of the data 128 81 used for training. Results confirm previous observation 129 82 about the superior performance of Hebbian-based semi-83 supervised approaches, compared to alternative solu-131 84 tions, especially in label-scarce learning regimes. More-132 85 over, the FastHebb solution exhibits a significant ac-133 87 celeration of training times, both compared to previous 134 Hebbian learning solutions, and compared to backprop-135 88 based alternatives. In particular, FastHebb achieves a 136 89 peak improvement in training up to 70 times faster than 137 90 corresponding Hebbian approaches not leveraging Fas-138 91 tHebb. This allowed to gracefully scale Hebbian al- 139 92 gorithms to large datasets, on the scale of ImageNet, 140 93 and architectures, on the scale of VGG [27]. Extending 141 94 Hebbian learning to other types of architectures, such 142 95 as residual networks [28] and transformers [29], in non-143 96 trivial and deserves to be explored in a separate work. 97 144

Some of the results on FastHebb were already presented in our recent conference publication [30]. However, those results were just preliminary and the aim of this paper is to significantly extend previous work. 148 Compared to [30], this paper performs a more comprehensive evaluation of the method by considering two types of test scenarios, image classification and CBIR, over four different computer vision benchmarks, including ImageNet. Moreover, we also extend the range of backbone architectures on which the approach is applied, pushing Hebbian learning to VGG-scale architectures for the first time, as far as we know.

In summary, our contribution is twofold:

- 1. A scalable solution for Hebbian synaptic updates is proposed;
- 2. Extensive evaluation of Hebbian algorithms is presented, including new experiments on largescale datasets (ImageNet) and architectures (VGG) which (to the best of our knowledge) have been out of reach for Hebbian algorithms so far.

Here is the structure of the following Sections: Section 2 illustrates related contributions; Section 3 describes the proposed FastHebb method more in detail; Section 4 delves into the details of our experimental scenarios in sample efficient and large-scale settings; in Section 5, the results of our experiments are described; conclusive remarks are presented in Section 6.

2. Background and related work

Some past contributions focused on addressing the biological plausibility problem of backpropagation by proposing solutions that can be shown to approximate backprop using Hebbian updates. Contrastive Hebbian Learning (CHL) [31] and Equilibrium Propagation (EP) [6] approaches do so by leveraging recurrent network architectures with two phases of activity. A free phase fixes the values of input neurons to represent a given sample, while output neurons and the remaining hidden units are left free. The recurrent dynamics lead the network to settle down into a steady state, where an anti-Hebbian update is performed. During the forced phase, the activations of output neurons are also fixed to a value closer to the desired target. Again, the recurrent dynamics will induce the network into another steady state, where a Hebbian update is performed. This combination of updates can be shown to approximate backprop at each neuron, using only local information. Another approach with strong biological support is Predictive Coding (PC) [32], in which a layer optimizes a local loss function that accounts for the error in predicting the next layer activations. Optimization is performed in a nested fashion. First, neuron activations are optimized to meet

the objective, which leads to the emergence of recur- 201 149 rent interactions among neurons, followed by optimiza- 202 150 tion of the weights. Again, the resulting updates can be 203 151 shown to match backprop updates using only local in- 204 152 formation [33]. The PC approach has been successfully 153 205 applied in different flavors to DNN training [34, 35, 36]. 154 206 More recently, the Forward-Forward (FF) approach has 155 207 been proposed [7]. This is based on standard feedfor-156 ward architectures, but using an input vector composed 209 157 of both sample and target. The approach alternates a 210 158 positive phase, where sample and the correct target are 211 159 provided to the network, which is required to maximize 212 160 its activations, and a negative phase, where the sample 213 161 is paired with a randomly generated target, and the net- 214 162 work is required to minimize its activations. In a pre-163 vious work [37], we have used a similar method for 164 training biologically realistic models of in vitro cultured 217 165 neural networks, where sample and target are provided 218 166 simultaneously to the network, and Hebbian plasticity 219 167 reinforces the connection between the two, so that at 220 168 test time, when a sample with no target is provided, the 221 169 network can recall the association. A weight normaliza-170 tion mechanism plays the role of the negative phase in 171 222 this case. 172

In addition to these attempts to model supervised 223 173 learning from a biologically plausible perspective, other 224 174 efforts have been focused on modeling bio-inspired un- 225 175 supervised learning mechanisms. Past works used Heb- 226 176 bian learning with WTA competition models to train 227 177 feature extractors in feedforward and/or convolutional 228 178 CNNs [14, 15, 16, 17, 18, 19, 20], showing impressive 179 convergence speed. In particular, a recent work [38] 229 180 showed that soft-WTA training of DNNs allows the net- 230 181 work to extract increasingly abstract representations, in 231 182 the same vein as backprop training, but at the cost of 232 183 using very wide layers. The authors also provide exper- 233 184 imental results with Hebbian learning on ImageNet, al-185 234 though only for a single training epoch. The method that 235 186 we propose allows us to run a full training session (20) 236 187 epochs or more) even on ImageNet scale. Miconi [39] 237 188 proposed translations of some Hebbian synaptic update 238 189 equations into optimizable objective functions, which 239 190 are more relatable to common frameworks for DL. An- 240 191 other line of research explored the Similarity Matching 241 192 objective as a possible direction to derive biologically 242 193 plausible neural models for principal subspace extrac-194 tion [8, 9, 10, 11], with extensions also to the supervised 244 195 end-to-end recurrent training case [40, 41]. 196

197 In our past contributions, we took a hybrid ap- 246 proach, and explored Hebbian WTA and PCA train-247 198 ing of DNNs in semi-supervised scenarios, using unsu-248 199 pervised Hebbian algorithms as a tool for pre-training 249 200

[12, 13, 42]. Experiments showed promising results, compared to backprop-based alternative methods, especially in scarce data learning scenarios. Due to the difficulty of gathering manually labeled data, these scenarios are of strong practical interest. Given the promising results obtained in previous works, in this contribution we further enhance previous solutions towards achieving higher efficiency and scalability to more complex scenarios.

Other works have explored semi-supervised approaches exploiting unsupervised pre-training with backprop-based auto-encoding architectures [43, 44, 21, 221. A different direction towards semi-supervised learning is instead based on pseudo-labeling or consistency-based methods [45, 46, 47, 48]. Since our approach belongs to the unsupervised pre-training category, we will focus our comparisons in this setting. However, it is worth mentioning that the other approaches are not mutually exclusive with unsupervised pre-training, and, indeed, these can be integrated together, as also suggested in Sec. 6

3. Speeding Up Hebbian learning with FastHebb

In this Section we present the FastHebb method. We start by introducing a convenient notation, that will be used to translate Hebbian synaptic update equations into the FastHebb formulation. Then, we illustrate the learning rules that are analyzed in this work, and we derive their FastHebb-enhanced counterpart.

3.1. A convenient notation

When working with common packages for DL, such as Pytorch, data are typically represented as tensors. In this context, a tensor is simply a data array with multiple dimensions. We wish to introduce a notation for tensors that is better suited for relating mathematical formalism with the corresponding implementation in DL packages. For example, a tensor has a number of dimensions, whose interpretation lies in the mind of the programmer (e.g. batch, channel, height, and width dimensions for images). Given a tensor, packages such as Pytorch allow us to transpose or permute any of its dimensions, which corresponds to reordering indexes. We can also unsqueeze singleton dimensions or squeeze them out, which correspond to adding or removing singleton indexes, i.e. indexes of dimension 1. Therefore, in essence, the notation that we introduce is motivated by a more straightforward mapping to the programming formalism for working with tensors.

We consider tensors, denoted by capital letters, followed by one index per dimension. The symbol used

to denote the index denotes its meaning. For example, 296 250 we can use an index p = 1...M to represent the *batch* 297 251 dimension, an index q = 1...Q to represent the *chan*- 298 252 *nel* dimension, and index m = 1...M to represent the 299 253 weight vector dimension. A 1 symbol used as an index 300 254 represents a singleton. For example, a neural feature 301 255 map at some layer can be denoted as $A_{p,q,1}$, which is 302 256 a tensor with one element for each mini-batch element 303 257 (dimension p) and for each neuron (dimension q). The 304 258 last dimension is a singleton because the output of each 259 neuron for each mini-batch element is a scalar. Simi-305 260 larly, a weight matrix can be denoted by $B_{1,q,m}$, because ³⁰⁶ 261 it does not extend along the mini-batch dimension, but 307 262 it has a number of channels (one per neuron), and each 308 263 corresponds to a weight vector. Concerning the input ³⁰⁹ 264 tensors, for example images, they have a mini-batch, ³¹⁰ 265 channel, height, and width dimensions. However, due to 266 the convolutional processing, a patch will be extracted 267 from each horizontal and vertical location of each im-268 age, which is treated, for the Hebbian learning purposes, 269 as a separate input. Therefore, our mini-batch is the col-270 lection of all patches extracted from all images. Each 271 patch is flattened into a vector, whose size corresponds 272 to the weight vector size of the next neural layer. On the 273 other hand, this tensor does not extend along the neu-274 ron dimension. Overall, our input tensors can be repre-275 sented as $C_{p,1,m}$. 276

This notation is convenient, because it allows us to 277 easily swap indexes, or squeeze and unsqueeze sin-278 gleton dimensions. If tensors have compatible dimen-279 313 sions, we can also perform element-wise operations (ad-280 ditions, multiplications, etc.). When a dimension is 281 a singleton, it automatically undergoes broadcasting, 282 i.e. the tensor is replicated along that direction until it 283 matches the corresponding dimension of the other ten-284 sor involved in the operation. 285

Matrix multiplication plays an important role in DNN 315 286 processing. We make the usage of matrix multipli- 316 287 cations explicit in our notation, by writing $bmm(\cdot, \cdot)$ 317 288 (which stands for batch matrix multiplication): 318 289

Notice that the matrix multiplication operation is 325 290 equivalent to taking the element-wise product between 326 291 292 tensors U and V, identifying the common index i and $_{327}$ summing over (or contracting) index k. Specifically, in-328 293 dex *i* represents a batch dimension, and the operation 329 294 is a batch matrix multiplication between *i* matrix pairs. 330 295

For each pair, the two matrices have indices (i, k) and (k, l), which are mapped to indices (j, l): $(j, k) \times (k, l) \rightarrow$ (j, l). The operation generalized to tensors with more that three dimensions as follows: all dimensions except the last two are considered as batch dimensions, while the last two dimensions represent rows and columns of the matrices being batch-multiplied. Batch dimensions must correspond between the two tensors, or be singleton (in which case, broadcasting takes place).

3.2. Hebbian synaptic updates: from computation to aggregation

We consider two types of Hebbian learning rules: Hebbian PCA (HPCA) and soft-Winner-Takes-All (SWTA). In this paper, we just give the definition of these learning rules, but the interested reader can find more details in [20, 19, 13, 12].

Given a layer of neurons whose activations are denoted by a_q (index q refers to the q-th neuron in the layer), whose weight vectors are denoted by \mathbf{b}_{q} , and whose input vector is denoted by c, the SWTA synaptic update equation is the following:

$$\Delta \mathbf{b}_q = \alpha \, s_q \, (\mathbf{c} - \mathbf{b}_q) \tag{2}$$

Where α stands for the learning rate and s_q is the softmax of the activations with temperature T [49]:

$$s_q = \frac{a_q/T}{\sum_k a_k/T} \tag{3}$$

Essentially, this modulates the update steps so that neurons with higher activations will also 'win' larger updates.

The HPCA learning rule, instead, is the following:

$$\Delta \mathbf{b}_q = \alpha \, a_q \, (\mathbf{c} - \sum_{k=1}^q a_k \mathbf{b}_k) \tag{4}$$

This rule can be shown to induce neurons to extract the principal components from data [50, 51].

When working with images and convolutional layers, these weight updates need to be computed for each patch extracted from a given image. However, due to the constraints of convolutional layers, neurons at different offsets need to maintain shared weights, hence they are bound to follow the same synaptic modifications. This can be achieved by aggregating the different weight updates, obtained from patches at different locations, into a unique update. Aggregation needs to be performed for all the images in a mini-batch as well. The overall two-phases approach is depicted in Fig. 1.

Aggregation is performed simply by averaging, in the HPCA case, or by a weighted average, where the weights are s_a coefficients, for SWTA.

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Figure 1: Two phases of weight update: update computation for each patch, followed by aggregation of several updates.

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331 3.3. From Hebbian synaptic updates to FastHebb

The Hebbian rules presented above can be rewritten in matrix form, including the aggregation step, using the notation outlined at the beginning of this Section:

$$\Delta B_{1,q,m} = \sum_{p} D_{p,q,1} \Delta B_{p,q,m} = \operatorname{bmm}(D_{q,1,p}, \Delta B_{q,p,m})$$
(5)

where $\Delta B_{p,q,m}$ is the collection of all weight updates that need to be aggregated, and $D_{p,q,1}$ is the tensor of coefficients for the aggregation.

Now that the two phases of weight computation and aggregation are merged together, we proceed differently depending on the specific learning rule.

FastHebb for SWTA. Let's rewrite the SWTA update rule as follows:

$$\Delta B_{1,q,m} = \alpha \sum_{p} D_{p,q,1} S_{p,q,1} \left(C_{p,1,m} - B_{1,q,m} \right)$$

$$= \alpha \sum_{p} (DS)_{p,q,1} (C - B)_{p,q,m}$$
(6)

$$= \alpha \operatorname{bmm}((DS)_{q,1,p}, (C-B)_{q,p,m})$$

340 Where $D_{p,q,1} = \frac{S_{p,q,1}}{\sum_p S_{p,q,1}}$.

The computational complexity required by this algorithm is O(PQM) in time. Moreover, if we wish to 359

exploit GPU parallelism, we need to keep a $P \times Q \times M$ tensor in memory, thus requiring O(PQM) space complexity as well, which can be prohibitive for large-scale scenarios.

However, it is possible to improve over these bounds by contracting the aggregation index p (which is typically the largest dimension) early:

$$\Delta B_{1,q,m} = \alpha \sum_{p} D_{p,q,1} S_{p,q,1} \left(C_{p,1,m} - B_{1,q,m} \right)$$

= $\alpha \sum_{p} (DS)_{p,q,1} C_{p,1,m} - \alpha \sum_{p} (DS)_{p,q,1} B_{1,q,m}$
= $\alpha \operatorname{bmm} \left((DS)_{1,q,m}, C_{1,q,m} \right) - \alpha \sum_{p} (DS)_{p,q,1} B_{1,q,m}$
= $\alpha \operatorname{bmm} \left((DS)_{1,q,p}, C_{1,p,m} \right) - \alpha E_{1,q,1} B_{1,q,m}$
(7)

Where $E_{1,q,1} = \sum_{p} (DS)_{p,q,1}$.

This requires only O(Q(P + M)) space complexity. Concerning the time complexity, this depends on the specific matrix multiplication algorithm adopted, but this can be made lower than O(PQM). This is the Fas-tHebb formulation for SWTA.

FastHebb for HPCA. Similarly to the SWTA case, we can rewrite the HPCA equation, together with the aggregation phase (in this case, the coefficient $D_{p,q,1}$ is just $\frac{1}{p}$), with the proposed notation:

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$$\Delta B_{1,q,m} = \alpha \frac{1}{P} \sum_{p} A_{p,q,1} \left(C_{p,1,m} - \sum_{q'=1}^{q} A_{p,q',1} B_{1,q',m} \right)$$

= $\alpha \frac{1}{P} \sum_{p} A_{p,q,1} \left(C_{p,1,m} - \sum_{q'=1}^{Q} T_{q,q'} A_{p,q',1} B_{1,q',m} \right)$
= $\alpha \frac{1}{P} \sum_{p} A_{p,q,1} F_{p,q,m}$
= $\alpha \frac{1}{P} \operatorname{bmm} \left(A_{q,1,p}, F_{q,p,m} \right)$ (8)

Where $F_{p,q,m} = (C_{p,1,m} - \sum_{q'=1}^{Q} T_{q,q'} A_{p,q',1} B_{1,q',m})$, and $T_{q,q'}$ is simply a lower-triangular matrix with all ones on and below the main diagonal and all zeros above.

The computation of the HPCA equation is slightly $_{374}$ more complex, requiring $O(PQ^2M)$ space and time, but this can be improved by reordering the sums: $_{376}$

$$\begin{split} \Delta B_{1,q,m} &= \alpha \, \frac{1}{P} \sum_{p} A_{p,q,1} \left(C_{p,1,m} - \sum_{q'=1}^{Q} T_{q,q'} A_{p,q',1} B_{1,q',m} \right) \\ &= \alpha \, \frac{1}{P} \sum_{p} A_{p,q,1} C_{p,1,m} \\ &- \alpha \, \frac{1}{P} \sum_{p} A_{p,q,1} \sum_{q'=1}^{Q} T_{q,q'} A_{p,q',1} B_{1,q',m} \\ &= \alpha \, \frac{1}{P} \text{bmm} \left(A_{1,q,p}, C_{1,p,m} \right) \\ &- \alpha \, \frac{1}{P} \sum_{q'=1}^{Q} \sum_{p} A_{p,q,1} A_{p,q',1} T_{q,q'} B_{1,q',m} \\ &= \alpha \, \frac{1}{P} \text{bmm} \left(A_{1,q,p}, C_{1,p,m} \right) \\ &- \alpha \, \frac{1}{P} \sum_{q'=1}^{Q} \text{bmm} \left(A_{1,q,p}, A_{1,p,q'} \right) T_{q,q'} B_{1,q',m} \\ &= \alpha \, \frac{1}{P} \text{bmm} \left(A_{1,q,p}, C_{1,p,m} \right) \\ &- \alpha \, \frac{1}{P} \sum_{q'=1}^{Q} G_{1,q,q'} B_{1,q',m} \\ &= \alpha \, \frac{1}{P} \text{bmm} \left(A_{1,q,p}, C_{1,p,m} \right) \\ &- \alpha \, \frac{1}{P} \sum_{q'=1}^{Q} G_{1,q,q'} B_{1,q',m} \\ &= \alpha \, \frac{1}{P} \text{bmm} \left(A_{1,q,p}, C_{1,p,m} \right) \\ &- \alpha \, \frac{1}{P} \sum_{q'=1}^{Q} \text{bmm} \left(A_{1,q,p}, C_{1,p,m} \right) \\ &- \alpha \, \frac{1}{P} \text{bmm} \left(A_{1,q,p}, C_{1,p,m} \right) \\ &- \alpha \, \frac{1}{P} \text{bmm} \left(G_{1,q,q'}, B_{1,q',m} \right) \end{split}$$

Where,
$$G_{1,q,q'} = \operatorname{bmm}(A_{1,q,p}, A_{1,p,q'}) T_{q,q'}$$
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	Layer 1	Layer 2	Layer 3	Lager 6	Layer 8	Layer 6
Paper	AL Conv. 54 No. 10 25 Parts 26 Parts Read	Ind Color, 128 Rel(J) Rel(J)	ALCON. 10 NO MACHINE 20 Machine 2	NJ Corn 256 NUCL RADI Very	Peter Rest Kenn Depot	Canadian Landare cere ciero

Figure 2: Backbone neural network model used for our experiments.



Figure 3: Bigger neural network model for ImageNet experiments.

The overall computation now has $O(Q^2 + QM)$ complexity in space, and up to $O(PQM + PQ^2 + Q^2M)$ in time. This is the FastHebb formulation for HPCA.

4. Evaluation scenario

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We evaluated the proposed methodology on a number of established computer vision benchmarks: CI-FAR10/100 [24], Tiny ImageNet [25], and ImageNet [26]. We performed an evaluation of Hebbian-based approaches in semi-supervised learning settings, on a backbone network model described in the following, compared to a Variational Auto-Encoder (VAE) [23, 21] baseline. In addition, we provide a FastHebb evaluation on VGG [27], to show the scalability of the proposed approach to large architectures. We evaluated the performance both in terms of classification accuracy, and in terms of training speedup achieved with FastHebb. We also provide an evaluation of Hebbian neural features for large-scale image retrieval tasks.

386 4.1. Neural network backbone for evaluation

In order to provide an evaluation for the proposed 387 approach, we need to define a suitable backbone net-388 work architecture for our experiments. For this purpose, 389 we need a network that presents the common architec-390 tural features of Convolutional Neural Networks (pool-391 ing and convolutional layers [52], batch normalization 392 [53], etc.). On the other hand, we need to exclude more 393 recent features such as residual connections [28] or at-394 tention layers [29], for Hebbian algorithms are not triv-395 ial to generalize to these cases, which deserve to be an-396 alvzed in a separate work. For a first experimentation 397 stage, we do not need to consider a very large model; it 398 is instead preferable to consider a more compact archi-399 tecture, which enables faster experimentation, and eas-400 ier analysis of deep features, also on a layer-by-layer 401 basis. It also makes reproducibility by other researchers 402 more accessible. Therefore, we opted for an AlexNet-403 inspired [52] architecture shown in Fig. 2, with 6 layers, 404 which is also consistent with previous works [20, 30]. 405 For larger-scale experiments on ImageNet we used an 406

extended version of the previous model with 10 layers, 455 407 shown in Fig. 3, as well as a VGG model [27]. 408

4.2. Semi-supervised training protocol for sample-409 efficient learning 410

We evaluated the proposed approach assuming a con-411 dition of scarcity of available labeled training data. We 412 define a sample efficiency regime as the percentage of 413 available labeled samples, over the total number of training samples. For each of the considered datasets, 415 467 we performed experiments in eight different sample ef-416 ficiency regimes: 1%, 2%, 3%, 4%, 5%, 10%, 25%, and 417 100%. 418

In order to take advantage of both labeled and un-419 labeled training samples, for each sample efficiency 420 regime, we followed a semi-supervised training proto-421 col in two phases: first, the network is pre-trained using 422 one of the proposed unsupervised Hebbian algorithms, 423 exploiting all the available training samples; second, 424 end-to-end fine-tuning is performed, using supervised 425 backprop training on a cross-entropy loss, and exploit-426 ing the labeled samples only. Finally, both the resulting 427 classification accuracy and the training time (in terms of 428 epoch duration, number of epochs, and total duration) 429 were recorded. 430

As a baseline for comparison, we used unsupervised 431 pre-training based on the Variational Auto-Encoder 432 (VAE) approach [54]. In this case, pre-training was 433 performed by using the deep layers (excluding the fi-434 nal classifier) of the proposed architectures as encoder, 435 mapping their output to 256 gaussian latent variables. 436 This was augmented with a another network branch, 437 acting as decoder, with a specular structure w.r.t. the en-438 coder (i.e. pooling layers replaced with unpooling, and convolutions with transpose convolutions), mapping the 440 latent variables to a decoded sample. The overall mod-441 487 els were trained in the encoding-decoding task, opti-442 mizing the β -VAE Variational Lower Bound [55], in an 443 end to end fashion, using all the available training sam-444 ples. At this point, the decoder was dropped, a linear 445 classifier was placed on top of the latent features, and 446 491 supervised backprop-based end-to-end fine tuning was 447 performed, using only the available labeled samples for 492 448 the given sample efficiency regime. Essentially, this is 493 449 the standard semi-supervised training approach based 450 494 451 on state-of-the-art VAE architectures [21]. Notice that 495 in this case, however, the pre-training phase, even if un-452 supervised, is still backprop-based, while Hebbian algo-497 453 rithms enable pre-training without requiring backprop. 498 454

4.3. Retrieval with neural features

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Deep features extracted from pre-trained networks were also used as vector descriptors for multimedia content indexing and retrieval [56, 57, 58]. The performance of the resulting feature representation was evaluated in Content-Based Image Retrieval (CBIR) tasks.

The CBIR systems architecture works as follows: in a first phase, feature representations are computed for all images in a given database, by extracting the deep representations from the convolutional part of the network. These feature representations are then mapped to a binary 256-dimensional descriptor which is then used for indexing the database images. This is done as in [57] by training another piece of network, with a 256 units hidden layer with tanh activations and a final classifier. This is trained in the classification task, so that the feature representation is mapped to the correct class, but passing through a compression stage into the desired 256 dimensional vector. The tanh activation is a "soft" proxy for the binarization operation, which doesn't block gradients from flowing backward during training. The 256 dimensional representation is then binarized by a thresholding operation: positive values are mapped to 1 and negative values are mapped to 0.

Test set images are used as sample queries: at test time, their 256-dimensional binary feature representation is computed as well, and the database images are ranked against the query based on the Hamming distance between feature representations. Retrieved images are considered to be a correct match if they belong to the same class as the query.

The evaluation measure used for the CBIR task is the Average Precision Score (APS) :

$$APS = \sum_{i=1}^{K} P_i \left(R_i - R_{i-1} \right)$$
(10)

where P_i is the precision at the *i*th retrieved item, R_i is the corresponding recall. This score is renormalized (so that its maximum value is always 1) and averaged over all the queries, thus obtaining the mean Average Precision (mAP).

4.4. Implementation details

The experiments, implemented in Pytorch, depend on a number of hyperparameters, whose search was pursued by Coordinate Descent (CD) [59], optimizing, for each dataset, the accuracy results of the trained models on the respective validation set. In the following, the resulting parameters and implementation details are illustrated.

All training sessions were performed over 20 epochs 548 499 (which were enough for the models to converge). Data 549 500 were processed in mini-batches of 64 samples each, 550 501 and each sample was an RGB image of 32 pixels in 551 502 height and width for the 6-layer CIFAR10/100 and Tiny 552 503 ImageNet network, 210 pixels for the 10-layer Ima-504 553 geNet network, and 224 pixles for VGG (specifically, 505 the VGG-11 model was used), pre-normalized to zero 506 mean and unit variance. 507

Concerning Hebbian pre-training, the learning rate 555 508 parameter was set to 10^{-3} . For ImageNet training, we ⁵⁵⁶ 509 also introduced an adaptive learning rate mechanism to 557 510 cope with the high variance of weight updates due to the 558 511 high dimensionality of the feature maps (causing insta-559 512 bility during training), which divides the learning rate 560 513 by the square root of the input size (this corresponds 514 to normalizing the output variance, assuming the inputs 515 are normalized). For SWTA training only, whitening 516 pre-processing was also necessary, as in [60, 24], al-564 517 though this step did not show any benefit on the other 565 518 approaches. SWTA uses 0.02 as inverse temperature pa-566 519 rameter 1/T. 567 520

Batch-norm layers used momentum 0.9. 521

Backprop-based training (i.e. both fine-tuning and 569 522 VAE pre-training) leveraged Stochastic Gradient De-570 523 scent (SGD) optimization with learning rate 10^{-3} , and ⁵⁷¹ 524 momentum 0.9, with Nesterov acceleration [61]. After 572 525 10 training epochs, learning rate was reduced by half 573 52 every 2 epochs until the end of the training session. The 574 527 best training epoch in terms of validation results was 575 528 then selected as final model (early stopping). 529

 β -VAE training used coefficient $\beta = 0.5$. 530

578 Supervised fine-tuning was regularized by dropout 531 with 0.5 rate, and L_2 weight decay with penalty equal 579 532 to $5 \cdot 10^{-2}$, 10^{-2} , $5 \cdot 10^{-3}$, $1 \cdot 10^{-3}$, for CIFAR10, CI-533 FAR100, Tiny ImageNet, ImageNet, respectively. 580 534

The implementation used Pytorch version 1.8.1 and 535 Python 3.7, with an Ubuntu 20.4 system running on 536 an I7 series 10700K Intel Processor, 32GB RAM, and 537 12GB NVidia Geforce 3060 GPU. 538

5. Results 539

The results of the experiments described in the pre-540 vious Section are illustrated hereafter. First, we re-541 port the recorded training speed, in terms of epoch du-542 ration, number of epochs for convergence, and total 543 544 duration, on CIFAR10/100, Tiny ImageNet, and ImageNet datasets, comparing VAE pre-training, ordinary 545 Hebbian learning, and FastHebb. Second, we report 546 the classification and retrieval results of the various ap-547

proaches in the label-scarcity scenarios described earlier. Finally, we report the results on the VGG architecture as well. The results were obtained from averaging five independent experiment iterations, and t-testing confirmed the observed differences to be statistically significant with p-values below 0.05.

5.1. Training speed analysis

Table 1 shows a comparison between the considered approaches in terms of computational performance of training, on the 10-layer (for ImageNet) and 6-layer (for the other datasets) architectures. The Table shows the single epoch duration, the number of epochs until convergence (measured as the point after which validation results stop improving), and the total training duration 1 . These results are specifically focused on the pre-training duration, while we observed no statistically significant difference in the duration of the successive fine-tuning phase for different pre-training approaches.

We can observe that FastHebb methods are significanly faster (up to 50 times for HPCA and HPCA-FH on ImageNet) than the traditional Hebbian counterparts, with an epoch duration becoming comparable to backprop-based VAE training. This enables Hebbian approaches to scale gracefully to complex datasets such as ImageNet, where the best speed-up by a factor of 50, in terms of epoch duration, is observed for HPCA. Moreover, the overall training duration of Hebbian approaches becomes faster (up to 5 times on ImageNet) than VAE, thanks to the lower number of epochs required to convergence. Among the Hebbian approaches, soft-WTA has lower time complexity, and it is in fact faster.

5.2. Label scarcity results

Table 2 illustrates the classification results, in terms of accuracy (top-1 for CIFAR10, and top-5 for the other datasets, since they contain many more classes), in various sample efficiency regimes, comparing the alternative approaches. Notice that in the results for HPCA and SWTA there is no difference between using FastHebb or not. In fact, despite the computational speedup, from the algebraic point of view FastHebb is equivalent to ordinary Hebbian learning, leading to the same results. Therefore, we show these results just once.

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¹For Hebbian approaches not using FastHebb, the training duration would be unfeasible to measure explicitly; instead, it was estimated by multiplying the single epoch duration by the number of epochs

Table 1: Analysis of algorithm performance on each dataset, for VAE, Hebbian PCA (HPCA), Hebbian PCA with FastHebb (HPCA-FH), soft-WTA (SWTA), and soft-WTA with FastHebb (SWTA-FH) methods, on the 10-layer (for ImageNet) and 6-layer (for the other datasets) networks.

Dataset	Method	Epoch Duration	Num. Epochs	Total Duration
	VAE	14s	17	3m 58s
	SWTA	4m 14s	1	4m 14s
CIFAR10	SWTA-FH	18s	1	18 s
	HPCA	6m 23s	12	1h 16m 36s
	HPCA-FH	19s	12	3m 48s
	VAE	15s	15	3m 45s
	SWTA	4m 16s	1	4m 16s
CIFAR100	SWTA-FH	18s	1	18 s
	HPCA	6m 25s	7	44m 55s
	HPCA-FH	19s	7	2m 13s
	VAE	33s	20	11m
	SWTA	9m 41s	1	9m 41s
Tiny ImageNet	SWTA-FH	41s	1	41s
	HPCA	14m 20s	14	3h 20m 40s
	HPCA-FH	43s	14	10m 2s
	VAE	2h 59m 19s	16	47h 49m 4s
	SWTA	105h 13m 24s	3	315h 40m 12s
ImageNet	SWTA-FH	3h 38m 6s	3	10h 54m 18s
	HPCA	155h 41m 39s	3	467h 4m 57s
	HPCA-FH	3h 39m 18s	3	10h 57m 54s

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The results show that, in conditions of label scarcity 616 591 (sample efficiency regimes below 4-5%), Hebbian ap-592 proaches perform significantly better than VAE. On the 593 other hand, it is only when far more labels are available 594 for the supervised fine-tuning phase that VAE-based 595 pre-training really kicks in. In these scenarios, however, 596 the performance of Hebbian approaches is comparable 597 or only slightly lower, but this is compensated by the 598 speedup in training time observed before. Comparing 599 HPCA and SWTA, it appears that the former performs 600 typically better. 601

5.3. Retrieval experiments 602

Table 3 shows the retrieval mAP results obtained 629 603 on the various dataset, for each of the considered ap-630 604 proaches, on the 10-layer (for ImageNet) and 6-layer 631 605 (for the other datasets) architectures. 606

This second scenario confirms the previous observa- 633 607 tions that, in conditions of extreme label scarcity (be- 634 608 low 10%), Hebbian-based neural features achieve better 635 609 results than VAE counterparts. Again, VAE-based pre-610 training improves in higher regimes, but, as observed 611 637 612 before, this is fairly compensated by the training time 638 advantage of Hebbian approaches. Comparing HPCA 613 and SWTA, also in this case it appears that the former 640 614 performs typically better. 615

5.4. Experiments on VGG

In Tab. 5, we report the training times required for the pre-training phase of VGG models using the different approaches considered so far. We do not consider VAEtype training of the VGG model, because that requires a large decoder, making the overall model very deep, which we found to be untrainable due to vanishing gradients [62, 63]. On the other hand, Hebbian pre-training was straightforward to apply in this case, as it requires no gradient backpropagation. Instead, as a baseline for comparison, we used Xavier initialization [64] (note that, since this is not properly a training method, it is not included in Tab. 5). In fact, it is known that appropriate initialization methods can achieve competitive results compared to end-to-end pre-training [64, 65].

Training times show once more the effectiveness of FastHebb methods in training large scale architectures, while using ordinary Hebbian learning would be unfeasible. In the best case, a speedup of almost 70 times is reached, comparing HPCA-FH with HPCA.

Finally, in Tab. 5, we report the results, both in terms of classification accuracy and retrieval mAP, achieved by training the VGG model in the semi-supervised task. We show the results achieved with Xavier initialization, HPCA pre-training, and SWTA pre-training.

When the scale of the architecture increases, it ap-

Regime	Method	CIFAR10	CIFAR100	Tiny ImageNet	ImageNet
1%	VAE	22.54	12.28	5.55	2.72
	SWTA	30.23	15.30	6.20	6.69
	HPCA	39.75	22.63	11.38	8.65
	VAE	26.78	15.25	6.74	6.14
2%	SWTA	36.59	20.76	8.56	11.52
	HPCA	45.51	30.83	15.71	13.64
	VAE	29.00	16.44	7.74	15.35
3%	SWTA	41.54	23.69	10.26	15.67
	HPCA	48.80	35.04	18.23	17.28
	VAE	31.15	17.89	8.45	23.97
4%	SWTA	45.31	26.91	11.52	19.95
	HPCA	51.28	38.89	20.55	20.39
	VAE	32.75	18.48	9.29	29.04
5%	SWTA	48.35	29.57	12.55	24.87
	HPCA	52.20	41.42	22.46	23.28
	VAE	45.67	23.80	13.51	43.73
10%	SWTA	58.00	38.26	16.70	41.54
	HPCA	57.35	48.93	28.13	34.27
	VAE	68.70	52.59	37.89	61.33
25%	SWTA	69.85	56.26	24.96	59.34
	HPCA	64.77	58.70	37.10	56.92
	VAE	85.23	79.97	60.23	76.84
100%	SWTA	85.37	79.80	54.94	76.10
	HPCA	84.38	74.42	53.96	77.28

Table 2: Accuracy results on each dataset (top-1 for CIFAR10, and top-5 for the other datasets, since they have many more classes), for the various approaches explored, on the 10-layer (for ImageNet) and 6-layer (for the other datasets) networks.

pears that SWTA approach improves over HPCA. Pre- 662 642 vious observations about Hebbian methods performing 663 643 better in low sample efficiency regimes (5% and below) 664 644 are confirmed. In particular, SWTA outperforms the 665 645 network with no pre-training by a margin up to 5 per- 666 646 cent points in accuracy, in the 1-2% sample efficiency 667 647 regimes. In terms of mAP, Hebbian pre-training is still 668 648 slightly superior, although the difference is not statisti- 669 649 cally significant. 650 670

6. Conclusions and future work 651

In this article, we have illustrated the FastHebb ap-652 proach for Hebbian learning, which leverages a matrix 653 multiplication formulation of Hebbian synaptic updates 654 to achieve higher efficiency. This makes Hebbian learn-655 ing more scalable, enabling teh use of Hebbian neural 677 656 features also on large datasets (ImageNet) and architec-657 tures (VGG), which (to the best of our knowledge) have 678 658 been computationally prohibitive for Hebbian learning 659 so far. Experimental scenarios of label scarcity show 660

promising results of Hebbian pre-training compared to 661 681 backprop-based alternatives such as VAE, considering classification accuracy, retrieval mAP, and training time.

Even though, in this paper, we have shown that it is possible to scale Hebbian training to large models such as VGG, further work needs to be done to adapt Hebbian approaches to more recent architectures, such as residual networks [28] or Transformers [29]. Moreover, additional performance improvements might come from the combination of Hebbian-based pre-training with pseudo-labeling and consistency-based semi-supervised methods [66, 48]. Finally, in line with recent efforts towards backprop-free learning (such as the Forward-Forward algorithm [7]), we plan to explore strategies to combine Hebbian approaches with local supervision signals.

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Table 3: mAP results on each dataset, for the various approaches explored, on the 10-layer (for ImageNet) and 6-layer (for the other datasets) networks.

Regime	Method	CIFAR10	CIFAR100	Tiny ImageNet	ImageNet
	VAE	21.90	6.10	3.18	0.95
1%	SWTA	28.29	8.16	3.80	1.94
	HPCA	37.39	15.39	6.61	4.88
	VAE	23.13	6.40	3.20	1.39
2%	SWTA	31.71	8.82	3.95	2.54
	HPCA	39.80	15.98	7.39	5.72
	VAE	24.27	6.36	3.52	2.02
3%	SWTA	34.54	9.13	4.16	3.21
	HPCA	41.64	16.40	7.57	6.56
	VAE	24.36	6.36	3.57	3.83
4%	SWTA	36.54	9.41	4.32	3.77
	HPCA	43.22	16.72	7.61	7.19
	VAE	24.65	6.49	3.57	5.49
5%	SWTA	38.69	9.66	4.39	4.19
	HPCA	44.92	17.10	7.79	7.75
	VAE	28.26	7.09	3.76	13.27
10%	SWTA	48.55	11.27	5.01	9.14
	HPCA	49.99	18.42	8.45	10.56
	VAE	62.30	13.69	7.59	24.50
25%	SWTA	63.95	16.37	6.98	20.75
	HPCA	58.81	21.49	10.13	21.24
	VAE	84.67	42.83	22.64	44.21
100%	SWTA	84.54	43.95	20.86	39.60
	HPCA	81.80	36.19	17.99	43.81

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Table 4: Comparison of ImageNet training times, for Hebbian PCA (HPCA), Hebbian PCA with FastHebb (HPCA-FH), soft-WTA (SWTA), and soft-WTA with FastHebb (SWTA-FH) methods, on VGG-11.

Dataset	Method	Epoch Duration	Num. Epochs	Total Duration
	SWTA	290h 44m	5	1454h
	SWTA-FH	7h 7m	5	35h 35m
ImageNet	HPCA	453h 32m	13	5896h
-	HPCA-FH	6h 25m	13	83h 25m

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Table 5: Accuracy results on ImageNet (top-5), and retrieval mean Average Precision (mAP) for the various approaches explored, on the VGG network.

Regime	Pre-train	Accuracy (%)	mAP (%)
	None	14.71	4.79
1%	SWTA	19.40	3.63
	HPCA	15.53	5.16
	None	26.88	6.42
2%	SWTA	31.91	5.18
	HPCA	27.24	6.70
	None	36.68	8.05
3%	SWTA	40.24	7.08
	HPCA	36.74	8.05
	None	44.01	9.34
4%	SWTA	46.98	8.95
	HPCA	43.69	9.68
	None	49,37	10.80
5%	SWTA	51.36	10.47
	HPCA	50.40	10.92
	None	65.61	17.44
10%	SWTA	65.09	17.53
	HPCA	65.49	17.65
	None	78.71	29.01
25%	SWTA	78.17	29.52
	HPCA	78.53	29.20
	None	90.03	50.58
100%	SWTA	88.00	49.93
	HPCA	88.54	50.06

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