

Investigating Hebbian Alternatives to Dense Associative Memory

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Abstract

Dense Associative Memories generalize traditional Hopfield networks while providing a substantially increased capacity. To do this, they show that by increasing the separation of similarity scores between query patterns and stored patterns. The cost of increased capacity is simplicity and biological plausibility. We propose to investigate the capacity characteristics of all Hebbian associative memories, to see if there exists any alternative to Dense Associative Memories which maintains the simplicity and appeal of Hopfield networks.

Humans are able to recognize and retrieve patterns of data using distorted, noisy, and partial patterns (Rumelhart et al., 1986). This capacity of human memory is known as *content-addressability*: patterns which are stored in memory are able to be “looked up” by themselves or their parts. Modeling this property is a classical task in computational cognitive and neuroscience (see Amari (1972), Little (1974), Marr (1971), Nakano (1972), and Stanley (1976)). The family of models which implement content-addressability are known as *associative memory models* (AMs). A recent revival of interest AMs in machine learning research, driven by their equivalence with “attention” layers in the transformer architecture (Ramsauer et al., 2021; Vaswani et al., 2023), has led to drastic advances in the storage capacity of AMs (Demircigil et al., 2017; Hu et al., 2024; Krotov & Hopfield, 2016).

The foundational model for modern associative memory research is the *Hopfield network* (J. J. Hopfield, 1984; J. Hopfield, 1982). Hopfield networks are single-layer neural networks with full, lateral connections. This means that each artificial neuron in the network is connected with every other neuron, except itself. Patterns that we wish to recall from the network are stored in the connection weights between every other unit. Hopfield networks learn new patterns to recall through a simple, biologically plausible learning rule called the *Activity Product Rule* (Haykin, 2009), which is a kind of Hebbian update rule (Hebb, 1949). The appeal of Hebbian update rules is that they are simple, in that they are totally defined by local interactions between layers in neural networks, and they are biologically plausible, having been observed in interactions between biological neurons (Bi & Poo, 1998; Markram et al., 1997; Rolls, 2013).

In spite of their simplicity and biological plausibility, Hopfield networks are inherently flawed. The number of patterns that Hopfield networks can store is pitifully small: with estimates in the range of 14% to 15% of the number of neurons in the network (Amit et al., 1987; J. Hopfield, 1982). In order to remedy the gap between the simplicity and plausibility of Hopfield networks and their child Dense Associative Memory models, we will theoretically derive and empirically test the storage capacities of all Hebbian update rules when used in single-layer neural networks with full lateral connections.

Discovering the limits of Hebbian alternatives to Dense Associative Memories is not only desirable because of simplicity and biological plausibility (while those are both laudable goals). For example, the study of Vector-Symbolic Architectures, used for representing high-level cognitive-tasks (Gayler, 1998; Kelly et al., 2013; Plate, 1995; Smolensky, 1990), requires an auto-associative memory “clean-up memory”. This means that the expressive capabilities of Vector-Symbolic Architectures are affected by the capacity of the associative memory used.

In the following we will discuss the problem in more depth. To do this, we will first lay out the theoretical motivation: beginning with an introduction to the literature in section 1, covering associative memories in subsection 1.1, Hopfield networks in subsection 1.2, their generalization with Dense Associative Memories in subsection 1.3, and Hebbian learning and learning rules in subsection 1.4. We will restate the problem of Hebbian Dense Associative Memories in section 2, and provide a work plan in subsection 2.1. Finally, we will discuss the timeline for completing this project in section 3.

1 Background

In order to understand the need for a Hebbian alternative to Dense Associative Memories, we must first familiarize ourselves with the literature up to this point. Associative Memory research has a wealth of literature and alternatives. In order to limit the scope of this research, we limit ourselves only to the family of models based on Hopfield networks (J. Hopfield, 1982).

1.1 Associative Memories

An associative memory, in particular an *auto*-associative memory, is a general structure that can be implemented by many mathematical and computational objects. Broadly speaking, an associative memory is a tuple of a set of objects \mathcal{X} , called the set of *traces* or *stored patterns*, and a (typically learned) identity map over the set of stored patterns, $T : \mathcal{X} \rightarrow \mathcal{X}$. More formally,

Definition 1.1.1 (Associative memories). An (auto) *associative memory* is a tuple $\langle \mathcal{X}, T \rangle$, where

- (a) \mathcal{X} is a set of objects, called the *pattern* set, or set of *traces*, typically high-dimensional vectors; and,

- (b) The *recall function*, T , maps from the set of traces back to the set of traces, such that $T(x) \approx x$, for all $x \in \mathcal{X}$, and $T(\bar{x}) \approx x$, where \bar{x} is a perturbed, masked, or degraded form of $x \in \mathcal{X}$.

We will be assuming throughout that the pattern set is the set of rows of the *pattern matrix* $\Xi = [\xi^1, \xi^2, \dots, \xi^N]$, where each *stored pattern* $\xi^i \in \{-1, 1\}^D$. It is sufficient to specify the pattern matrix, or the sequence of patterns, for an associative memory, and it should be assumed that the pattern set is just composed of the rows of the pattern matrix. Associative memories are *content-addressable*, as their recall function is able to recover desired associated patterns from partial information (Haykin, 2009; McClelland et al., 1986). This is in contrast to memory models which recall information based on arbitrary associations, like indices into memory locations.

While it is not essential to the most general definition of associative memories, we usually desire that the recall function be learned. Furthermore, associative memories have a maximum capacity of traces that they can store.

Definition 1.1.2 (Retrievability; Capacity). Following Bao et al. (2022), let us have an associative memory with patterns $\xi^1, \xi^2, \dots, \xi^N$, where each element ξ_j^i is -1 or 1 with equal likelihood. For any $\delta < 0.5$, we define the δ -perturbation of ξ^i , $\bar{\xi}^i$, as the D -dimensional vector ξ^i with each element flipped with a likelihood of δ . Then, we say that the set $\xi^1, \xi^2, \dots, \xi^N$ is (δ, ε) -retrievable if for every ξ^i , $i = 1, 2, \dots, N$ it is such that:

$$\mathbb{P}(T(\bar{\xi}^i) \neq \xi^i) < \varepsilon. \quad (1)$$

The *capacity* C of the associative memory is the maximum cardinality of the pattern set such that all patterns are (δ, ε) -retrievable.

In Krotov and Hopfield (2016), capacity is denoted by N^{\max} , where N is the first dimension of the pattern matrix. Associative memory capacity is affected by the (1) kind of data being stored, i.e. if the data has a high pairwise correlation, which leads to “cross-talk” (Kohonen, 1988), (2) the dimensionality of the data, and (3) the implementation of the recall function T . For example, if the associative memory is a neural network with tensor weights (Kelly et al., 2017), then the capacity scales with the number of the weights (Little & Shaw, 1978) (expanded further in subsection 1.3).

Before Hopfield networks, there were Correlation Matrix associative memories. So-called, because they relied on dot-product correlations for recall.

Example 1.1.1 (Correlation Matrix associative memory). *Let us have a pattern matrix $\Xi = [\xi^1, \xi^2, \dots, \xi^N]$ of vectors sampled from $\{1, -1\}^D$. The Correlation Matrix associative memory recall function is of the form:*

$$\begin{aligned} T(x) &= g \left(\sum_{\mu=1}^N \xi^\mu \left(\sum_{i=1}^D \xi_i^\mu x_i \right) \right) \\ &= g \left(\Xi^\top \Xi x \right), \end{aligned} \quad (2)$$

where g is an “activation function”, typically signum for binary vectors.

Example 1.1.2. Like above, let us have a pattern matrix $\Xi = [\xi^1, \xi^2, \dots, \xi^N]$ of vectors sampled from $\{1, -1\}^D$. Let our recall function T be:

$$T(x) = \xi^i, \text{ where } i = \arg \max_{i \in [1, N]} [\text{sim}(\xi^i, x)], \quad (3)$$

where sim is some similarity function (Kelly et al., 2017), e.g. the cosine similarity or Hamming distance.

Example 1.1.3 (Minerva2). *Minerva2* (Hintzman, 1984) is an associative memory used in cognitive science which is about as old as Hopfield networks. As per usual, we have binary patterns $\Xi = [\xi^1, \xi^2, \dots, \xi^N]$. The recall function is:

$$T(x) = \text{sgn} \left[\sum_{\mu=1}^N \xi^\mu \left(\frac{\sum_{i=1}^D \xi_i^\mu x_i}{\|\xi^\mu\| \|x\|} \right)^3 \right]. \quad (4)$$

1.2 Hopfield Networks

Hopfield networks (J. J. Hopfield, 1984; J. Hopfield, 1982) are an associative memory with deep connections to Example 1.1.1, except that they understand their recall function in terms of an *energy* function:

Definition 1.2.1 (Hopfield networks). Hopfield networks are single-layer neural networks of D computational units with full, lateral connections. The state of the network is described by the state vector σ at time t , denoted by $\sigma^{(t)}$. The *energy* of the network, $E(\sigma^{(t)})$ is:

$$E(\sigma^{(t)}) = -\frac{1}{2} \sum_{\mu=1}^N \left(\sum_{i=1}^D \xi_i^\mu \sigma_i^{(t)} \right)^2. \quad (5)$$

Let $\sigma[i := b]$ be the vector σ , with the i 'th component set to b . Then, update rule for the state of the network is:

$$\sigma_i^{(t+1)} = \text{sgn} \left[E(\sigma^{(t)}[i := -1]) - E(\sigma^{(t)}[i := 1]) \right]. \quad (6)$$

After setting the initial state at time $t = 0$, the network continuously updates random elements of the state vector depending upon whether the update lowers the energy of the network. The energy function, then, forms a “landscape”, where local minima in the space (ideally) correspond with desired patterns to be stored.

While the use of the energy function seems at odds with our definition of associative memories, we can align the two by considering a single-pass variant of the energy-minimization. It can be shown that updating every element i of the state vector $\sigma^{(t)}$ in a single pass (synchronously) corresponds with the update rule (Krotov et al., 2025):

$$T(\sigma) = \text{sgn} \left[\sum_{\mu} \xi^\mu \left(\sum_{i=1}^D \xi_i^\mu \sigma_i \right) \right], \quad (7)$$

which is just the update rule of Correlation Matrix associative memories (see, Example 1.1.1).

Hopfield networks unfortunately have a pitifully low storage capacity: the critical capacity C being around 14% to 15% of the number of neurons in the model (Amit et al., 1987; J. Hopfield, 1982). As a computational model of human memory this will not do. Likewise, this is unsuitable for machine learning tasks involving large datasets, e.g. large language models. Capacity problems led researchers to investigate how one can improve the capacity of Hopfield networks, while maintaining the simplicity and interpretability of energy minimization.

1.3 Dense Associative Memories

Dense Associative Memory models are a generalization of Hopfield networks that maintain energy minimization in recall, but have a dramatically increased storage capacity (Demircigil et al., 2017; Krotov & Hopfield, 2016). Instead of relying on full lateral connections between units in a single dimension, Dense Associative Memories either have multiple connections between computational units, or, increase the number of layers present in the model. Krotov and Hopfield (2021) notes that this is to be expected, given the information limits of single neurons.

Definition 1.3.1 (Dense Associative Memory). Given binary patterns $\xi^1, \xi^2, \dots, \xi^N$ sampled from $\{-1, 1\}^D$, a *Dense Associative Memory* is characterized by:

- (i) A D -dimensional *state vector* at time t , $\sigma^{(t)}$
- (ii) A *polynomial transmission function* F_n , where:

$$F_n(x) = \frac{1}{n}(x)^n; \quad (8)$$

- (iii) An *energy function* of the state of the network at time t ,

$$E(\sigma^{(t)}) = \sum_{\mu=1}^N F \left(\sum_{i=1}^D \xi_i^\mu \sigma_i^{(t)} \right), \quad (9)$$

- (iv) An update equation for the state of the network, such that:

$$\sigma_i^{(t+1)} = \text{sgn} \left[E(\sigma^{(t)})[i := -1] - E(\sigma^{(t)})[i := 1] \right]. \quad (10)$$

Like Hopfield networks, the state vector update rule is typically performed asynchronously. However, we can define a synchronous update rule which makes the Dense Associative Memory an Associative Memory with update rule:

$$T(\sigma) = \text{sgn} \left[\sum_{\mu=1}^N \xi^\mu F'_n \left(\sum_{i=1}^D \xi_i^\mu \sigma_i \right) \right], \quad (11)$$

where F'_n is the derivative of F_n . From this definition we immediately see how this framework generalizes Hopfield networks.

Example 1.3.1. *Hopfield networks in Definition 1.2.1 are Dense Associative Memories with polynomial transmission function F_2 .*

Example 1.3.2. *Minerva2 (from Example 1.1.3) are roughly Dense Associative Networks, assuming that all patterns $\xi^1, \xi^2, \dots, \xi^N$ and query patterns are normalized, and with polynomial transmission function F_4 .*

Knowing this, one can now see the relation between the state update rule of Hopfield networks in Definition 1.2.1 and Equation 7.

Dense Associative Memories improve the storage capacity of Hopfield networks in that the capacity C of a Dense Associative Memory, with polynomial function F_n , and dimensionality of patterns D is:

$$C \propto D^{n-1} \quad (12)$$

(Bao et al., 2022; Demircigil et al., 2017; Krotov & Hopfield, 2016). The increased capacity comes from the “steeper” energy basins around minima corresponding to patterns stored in the network, which is brought about by the exponentiation of the dot product similarity between the stored patterns and the query pattern (for discussion, see Kelly et al. (2017)). Memory capacity for Dense Associative Memories can also be increased by making more domain sensitive similarity functions (Millidge et al., 2022), e.g. with kernel estimation (Hu et al., 2024; Wu et al., 2024).

In spite of the enormous gains in capacity using non-linear transmission functions, Dense Associative Memories lose the ability to be interpreted as simple one-layer networks with lateral connections and trained via Hebbian update rules (Krotov & Hopfield, 2021; McAlister et al., 2025). On the one hand, this is to their benefit: a continuous variant of Dense Associative Memory called *Modern Hopfield Networks* has been shown to be equivalent to multi-head attention in Transformer neural networks (Ramsauer et al., 2021; Vaswani et al., 2023). Likewise, the framework behind Dense Associative Memories has been developed for modular neural network architectures (Krotov, 2021), and has been used to define an *Energy Based Transformer* (Hoover et al., 2023), as well as contribute to the interpretability of Diffusion Models (Pham et al., 2025).

But the cost is their simplicity. Correlation matrix memories trained with Hebbian update rules mimic observed phenomena in biological neural networks; meaning that they perform computations which are in principle achievable by biological systems.

1.4 Hebbian Learning Rules

Hopfield networks learn with the Activity Product Rule (Haykin, 2009). Talk about biological evidence for Hebbian learning rules. Stress simplicity.

The closest investigation to this proposed analysis is Lansner et al. (2025), which compares several extant Hebbian learning rules in the literature: namely,

the Willshaw learning rule, the Activity Product Rule, the Covariance Learning rule, the Presynaptic Covariance learning rule, the Bayesian Confidence Propagation learning rule, and the Bayesian Optimal Memory learning rule.

This work serves as an important comparison work between the theoretical results that we wish to derive.

2 Hebbian Dense Associative Memory

Outline of the project. Hopfield networks which, instead of manipulating a polynomial function, rather manipulate how patterns are stored to begin with.

2.1 Work Plan

Two main components of the project: theoretical and experimental. Theoretical involves proving some theorems about the minimum and maximum capacity of Hopfield networks with different Hebbian learning rules. Experimental work means implementing the different networks and testing them on a task. Suggest testing them on classification of degraded inputs.

3 Timeline

Theoretical work should be done by Christmas. Experimental work likewise requires only a couple of months. Data science certification requires emphasis here on large-scale projects. So, I would like to rigorously examine each model on a variety of tasks. One to two papers arises from this project: one talking about the theory of Hebbian DAM, and the other part including the experimental work. It would be better for a single paper, but if that runs too long, then experimental work can be a part 2.

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