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A Theory of Sequence Indexing and Working Memory in Recurrent Neural Networks

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To accommodate structured approaches of neural computation, we propose a class of recurrent neural networks for indexing and storing sequences of symbols or analog data vectors. These networks with randomized input weights and orthogonal recurrent weights implement coding principles previously described in vector symbolic architectures (VSA) and leverage properties of reservoir computing. In general, the storage in reservoir computing is lossy, and crosstalk noise limits the retrieval accuracy and information capacity. A novel theory to optimize memory performance in such networks is presented and compared with simulation experiments. The theory describes linear readout of analog data and readout with winner-take-all error correction of symbolic data as proposed in VSA models. We find that diverse VSA models from the literature have universal performance properties, which are superior to what previous analyses predicted. Further, we propose novel VSA models with the statistically optimal Wiener filter in the readout that exhibit much higher information capacity, in particular for storing analog data.

The theory we present also applies to memory buffers, networks with gradual forgetting, which can operate on infinite data streams without memory overflow. Interestingly, we find that different forgetting mechanisms, such as attenuating recurrent weights or neural nonlinearities, produce very similar behavior if the forgetting time constants are matched. Such models exhibit extensive capacity when their forgetting time constant is optimized for given noise conditions and network size.

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These results enable the design of new types of VSA models for the online processing of data streams.

1 Introduction

An important aspect of information processing is data representation. In order to access and process data, addresses or keys are required to provide a necessary context. To enable flexible contextual structure as required in cognitive reasoning, connectionist models have been proposed that represent data and keys in a high-dimensional vector space. Such models include holographic reduced representations (HRR; Plate, 1991, 2003) and hyperdimensional computing (HDC) (Gayler, 1998; Kanerva, 2009), and will be referred to here by the umbrella term *vector symbolic architectures* (VSA; see Gayler, 2003; section 4.1.1). VSA models have been shown to be able to solve challenging tasks of cognitive reasoning (Rinkus, 2012; Kleyko & Osipov, 2014; Gayler, 2003). VSA principles have been recently incorporated into standard neural networks for advanced machine learning tasks (Eliasmith et al., 2012), inductive reasoning (Rasmussen & Eliasmith, 2011), and processing of temporal structure (Graves, Wayne, & Danihelka, 2014; Graves et al., 2016; Danihelka, Wayne, Uria, Kalchbrenner, & Graves, 2016). Typically, VSA models offer at least two operations: one to produce key-value bindings (also referred to as role-filler pairs) and a superposition operation that forms a working memory state containing the indexed data structures. For example, to represent a time sequence of data in a VSA, individual data points are bound to time-stamp keys and the resulting key-value pairs superposed into a working memory state.

Here, we show that input sequences can be indexed and memorized according to various existing VSA models by recurrent neural networks (RNNs) that have randomized input weights and orthonormal recurrent weights of particular properties. Conversely, this class of networks has a straightforward computational interpretation: in each cycle, a new random key is generated, a key-value pair is formed with the new input, and the indexed input is integrated into the network state. In the VSA literature, this operation has been referred to as *trajectory association* (Plate, 1993). The memory in these networks follows principles previously described in reservoir computing. The idea of reservoir computing is that a neural network with fixed recurrent connectivity can exhibit a rich reservoir of dynamic internal states. An input sequence can selectively evoke these states so that an additional decoder network can extract the input history from the current network state. These models produce and retain neural representations of inputs on the fly, entirely without relying on previous synaptic learning as in standard models of neural memory networks (Caianiello, 1961; Little & Shaw, 1978; Hopfield, 1982; Schwenker, Sommer, & Palm, 1996; Sommer & Dayan, 1998). Models of reservoir computing include state-dependent

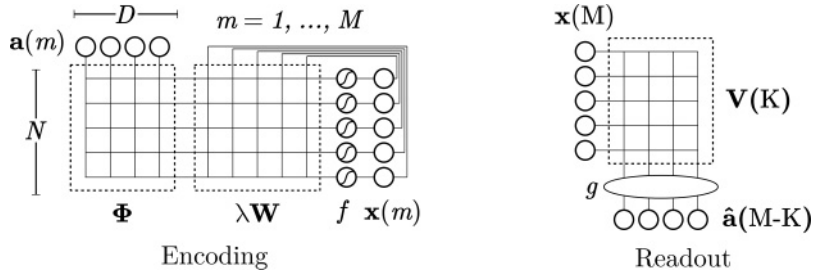

Figure 1: Network model investigated.

networks (Buonomano & Merzenich, 1995), echo-state networks (Jaeger, 2002; Lukoševičius & Jaeger, 2009), liquid-state machines (Maass, Natschläger, & Markram, 2002), and related network models of memory (White, Lee, & Sompolinsky, 2004; Ganguli, Huh, & Sompolinsky, 2008; Sussillo & Abbott, 2009). However, it is unclear how such reservoir models create representations that enable the selective readout of past input items. Leveraging the structured approach of VSAs to compute with distributed representations, we offer a novel framework for understanding reservoir computing.

2 Results

2.1 Indexing and Memorizing Sequences with Recurrent Networks. We investigate how a sequence of M input vectors of dimension D can be indexed by pseudo-random vectors and memorized by a recurrent network with N neurons (see Figure 1). The data vectors $\mathbf{a}(m) \in \mathbb{R}^D$ are fed into the network through a randomized, fixed input matrix $\mathbf{\Phi} \in \mathbb{R}^{N \times D}$. In the context of VSA, the input matrix corresponds to the codebook, and the matrix columns contain the set of high-dimensional random vector-symbols (hypervectors) used in the distributed computation scheme. In addition, the neurons might also experience some independent neuronal noise $\boldsymbol{\eta}(m) \in \mathbb{R}^N$ with $p(\eta_i(m)) \sim \mathcal{N}(0, \sigma_\eta^2)$. Further, feedback is provided through a matrix of recurrent weights $\lambda\mathbf{W} \in \mathbb{R}^{N \times N}$ where $\mathbf{W}$ is orthogonal and $0 < \lambda \leq 1$. The input sequence is encoded into a single network state $\mathbf{x}(M) \in \mathbb{R}^N$ by the recurrent neural network (RNN),

$$\mathbf{x}(m) = f(\lambda\mathbf{W}\mathbf{x}(m-1) + \mathbf{\Phi}\mathbf{a}(m) + \boldsymbol{\eta}(m)), \tag{2.1}$$

with $f(x)$ the component-wise neural activation function.

To estimate the input $\mathbf{a}(M-K)$ entered K steps ago from the network state, the readout is of the form

$$\hat{\mathbf{a}}(M-K) = g(\mathbf{V}(K)^\top \mathbf{x}(M)), \tag{2.2}$$

where $\mathbf{V}(K) \in \mathbb{R}^{N \times D}$ is a linear transform to select the input that occurred K time steps in the past (see Figure 1). In some models, the readout includes a nonlinearity $g(\mathbf{h})$ to produce the final output.

The effect of one iteration of equation 2.1 on the probability distribution of the network state $\mathbf{x}(m)$ is a Markov chain stochastic process, governed by the Chapman-Kolmogorov equation (Papoulis, 1984),

$$p(\mathbf{x}(m+1)|\mathbf{a}(m)) = \int p(\mathbf{x}(m+1)|\mathbf{x}(m), \mathbf{a}(m))\ p(\mathbf{x}(m))\ d\mathbf{x}(m), \tag{2.3}$$

with a transition kernel $p(\mathbf{x}(m+1)|\mathbf{x}(m), \mathbf{a}(m))$, which depends on all parameters and functions in equation 2.1. Thus, to analyze the memory performance in general, one has to iterate equation 2.3 to obtain the distribution of the network state.

2.1.1 Properties of the Matrices in the Encoding Network. The analysis simplifies considerably if the input and recurrent matrix satisfy certain conditions. Specifically, we investigate networks in which the input matrix $\mathbf{\Phi}$ has independent and identically distributed (i.i.d.) random entries and the recurrent weight matrix $\mathbf{W}$ is orthogonal with mixing properties and long cycle length. The assumed properties of the network weights guarantee the following independence conditions of the indexing keys, which will be essential in our analysis of the network performance:

- Code vectors $\mathbf{\Phi}_d$ are composed of *identically distributed* components,

$$p((\mathbf{\Phi}_d)_i) \sim p_{\mathbf{\Phi}}(x)\ \forall i, d \tag{2.4}$$

  where $p_{\mathbf{\Phi}}(x)$ is the distribution for a single component of a random code vector, and with $E_{\mathbf{\Phi}}(x)$, $V_{\mathbf{\Phi}}(x)$ being the mean and variance of $p_{\mathbf{\Phi}}(x)$, as typically defined by $E_{\mathbf{\Phi}}(\phi(x)) := \int \phi(x) p_{\mathbf{\Phi}}(x) dx$, $V_{\mathbf{\Phi}}(\phi(x)) := E_{\mathbf{\Phi}}(\phi(x)^2) - E_{\mathbf{\Phi}}(\phi(x))^2$, with $\phi(x)$ an arbitrary function.
- Components within a code vector and between code vectors are *independent*:

$$p\left((\mathbf{\Phi}_{d'})_i, (\mathbf{\Phi}_d)_j\right) = p((\mathbf{\Phi}_{d'})_i)\ p((\mathbf{\Phi}_d)_j)\ \forall j \neq i \vee d' \neq d. \tag{2.5}$$

- The recurrent weight matrix $\mathbf{W}$ is orthogonal and thus *preserves the mean and variance* of every component of a code vector:

$$\begin{aligned} E((\mathbf{W}\mathbf{\Phi}_d)_i) &= E((\mathbf{\Phi}_d)_i)\ \ \forall i, d, \\ \mathrm{Var}((\mathbf{W}\mathbf{\Phi}_d)_i) &= \mathrm{Var}((\mathbf{\Phi}_d)_i)\ \ \forall i, d. \end{aligned} \tag{2.6}$$

- The recurrent matrix preserves element-wise independence with a *large cycle time* (around the size of the reservoir):

$$p((\mathbf{W}^m \mathbf{\Phi}_d)_i, (\mathbf{\Phi}_d)_i) = p((\mathbf{W}^m \mathbf{\Phi}_d)_i)\ p((\mathbf{\Phi}_d)_i)\ \ \forall i, d; m = \{1, \ldots, O(N)\}. \tag{2.7}$$

The class of RNNs (see equation 2.1) in which the weights fulfill properties 2.4 to 2.7 contains the neural network implementations of various VSA models. Data encoding with such networks has a quite intuitive interpretation. For each input $a_d(m)$, a pseudo-random key vector is computed that indexes both the input dimension and location in the sequence, $\mathbf{W}^{M-m}\mathbf{\Phi}_d$. Each input $a_d(m)$ is multiplied with this key vector to form a new key-value pair, which is added to the memory vector $\mathbf{x}$. Each pseudo-random key defines a spatial pattern for how an input is distributed to the neurons of the network.

2.1.2 Types of Memories under Investigation. Reset memory versus memory buffer. In the case for finite input sequence length M, the network is reset to the zero vector before the first input arrives, and the iteration is stopped after the Mth input has been integrated. We refer to these models as *reset memories*. In the VSA literature, the superposition operation (Plate, 1991, 2003; Gallant & Okaywe, 2013) corresponds to a reset memory and, in particular, trajectory-association (Plate, 1993). In reservoir computing, the distributed shift register (DSR; White et al., 2004) can also be related to reset memories. In contrast, a *memory buffer* can track information from the past in a potentially infinite input stream ($M \to \infty$). Most models for reservoir computing are memory buffers (Jaeger, 2002; White et al., 2004; Ganguli et al., 2008). A memory buffer includes a mechanism for attenuating older information, which replaces the hard external reset in reset memories to avoid overload. The mechanisms of forgetting we will analyze here are contracting recurrent weights or neural nonlinearities. Our analysis links contracting weights (λ) and nonlinear activation functions (f) to the essential property of a memory buffer, the forgetting time constant, and we show how to optimize memory buffers to obtain extensive capacity.

Memories for symbols versus analog input sequences. The analysis considers data vectors $\mathbf{a}(m)$ that represent either symbolic or analog inputs. The superposition of discrete symbols in VSAs can be described by equation 2.1, where inputs $\mathbf{a}(m)$ are one-hot or zero vectors. A one-hot vector represents a symbol in an alphabet of size D. The readout of discrete symbols involves a nonlinear error correction for producing one-hot vectors as output, the winner-take-all operation $g(\mathbf{h}) = WTA(\mathbf{h})$. Typical models for reservoir computing (Jaeger, 2002; White et al., 2004) process one-dimensional analog input, and the readout is linear, $g(\mathbf{h}) = \mathbf{h}$ in equation 2.2. We derive the

information capacity for both uniform discrete symbols and gaussian analog inputs.

Readout by naive regression versus full minimum mean square error regression. Many models of reservoir computing use full optimal linear regression and set the linear transform in equation 2.2 to the Wiener filter $\mathbf{V}(K) = \mathbf{C}^{-1}\mathbf{A}(K)$, which produces the minimum mean square error (MMSE) estimate of the stored input data. Here, $\mathbf{A}(K) := \langle \mathbf{a}(M-K)\mathbf{x}(M)^\top \rangle \in \mathbb{R}^{N\times D}$ is the covariance between input and memory state, and $\mathbf{C} := \langle \mathbf{x}(M)\mathbf{x}(M)^\top \rangle \in \mathbb{R}^{N\times N}$ is the covariance matrix of the memory state. Obviously this readout requires inverting $\mathbf{C}$. In contrast, VSA models use $\mathbf{V}(K) = c^{-1}\langle \mathbf{a}(M-K)\mathbf{x}(M)^\top \rangle = c^{-1}\mathbf{W}^K\mathbf{\Phi}$, with $c = NE_{\mathbf{\Phi}}(x^2)$ a constant, which does not require matrix inversion. Thus, the readout in VSA models is computationally much simpler but can cause reduced readout quality. We show that the MMSE readout matrix can mitigate the crosstalk noise in VSA and improve readout quality in regimes where $MD \lesssim N$. This is particularly useful for the retrieval of analog input values, where the memory capacity exceeds many bits per neuron, limited only by neuronal noise.

2.2 Analysis of Memory Performance. After encoding an input sequence, the memory state $\mathbf{x}(M)$ contains information indexed with respect to the dimension $1, \ldots, D$ of the input vectors and with respect to the length dimension $1, \ldots, M$ of the sequence. The readout of a vector component, d, at a particular position of the sequence, $M-K$, begins with a linear dot product operation,

$$h_d(K) := \mathbf{V}_d(K)^\top \mathbf{x}(M), \tag{2.8}$$

where $\mathbf{V}_d(K)$ is the dth column vector of the decoding matrix $\mathbf{V}(K)$.

For readout of analog-valued input vectors, we use linear readout: $h_d(K) = \hat{a}_d(M-K) = a_d(M-K) + n_d$, where n_d is decoding noise resulting from crosstalk and neuronal noise. The signal-to-noise ratio, r, of the linear readout can then be defined as

$$r(K) := \frac{\sigma^2(a_d)}{\sigma^2(n_d)}, \tag{2.9}$$

where we suppressed the component index d and assume that the signal and noise properties are the same for all vector components.

For symbolic input, we will consider symbols from an alphabet of length D, which are represented by one-hot $\mathbf{a}$ vectors; that is, in each input vector, there is one component $a_{d'}$ with value 1 and all other a_d are 0. In this case, a multivariate threshold operation can be applied after the linear readout for error correction, the winner-take-all function: $\hat{\mathbf{a}}(M-K) = WTA(\mathbf{h}(K))$.

2.2.1 The Accuracy of Retrieving Discrete Inputs. For symbolic inputs, we will analyze the readout of two distinct types of input sequences. In the first type, a symbol is entered in every time step, and retrieval consists in classification to determine which symbol was added at a particular time. The second type of input sequence can contain gaps: some positions in the sequence can be empty. If most inputs in the sequence are empty, this type of input stream has been referred to as a sparse input sequence (Ganguli & Sompolinsky, 2010). The retrieval task is then detection: whether a symbol is present and, if so, reveal its identity.

For classification, if d' is the index of the hot component in $\mathbf{a}(M-K)$, then the readout with the winner-take-all operation is correct if in equation 2.8, $h_{d'}(K) > h_d(K)$ for all distracters $d \neq d'$. As we will see, under the independence conditions 2.4 to 2.7 and VSA readout, the h_d readout variables are the true inputs plus gaussian noise. The classification accuracy, p_{corr}, is

$$\begin{aligned} p_{corr}(K) &= p\left(h_{d'}(K) > h_d(K)\ \forall d \neq d'\right) \\ &= \int_{-\infty}^{\infty} p(h_{d'}(K) = h)\left[p(h_d(K) < h)\right]^{D-1} dh \\ &= \int_{-\infty}^{\infty} \mathcal{N}(h'; \mu(h_{d'}), \sigma^2(h_{d'}))\left[\int_{-\infty}^{h'} \mathcal{N}(h; \mu(h_d), \sigma^2(h_d))\, dh\right]^{D-1} dh' \\ &= \int_{-\infty}^{\infty} \mathcal{N}(h'; a_{d'}, \sigma^2(n_{d'}))\left[\int_{-\infty}^{h'} \mathcal{N}(h; a_d, \sigma^2(n_d))dh\right]^{D-1} dh'. \quad (2.10) \end{aligned}$$

For clarity in the notation of gaussian distributions, the argument variable is added: $p(x) \sim \mathcal{N}(x; \mu, \sigma^2)$.

The gaussian variables h and h' in equation 2.10 can be shifted and rescaled to yield

$$\begin{aligned} p_{corr}(K) &= \int_{-\infty}^{\infty} \frac{dh}{\sqrt{2\pi}}\, e^{-\frac{1}{2}h^2}\left[\Phi\left(\frac{\sigma(h_d)}{\sigma(h_{d'})}h - \frac{\mu(h_d) - \mu(h_{d'})}{\sigma(h_{d'})}\right)\right]^{D-1} \\ &= \int_{-\infty}^{\infty} \frac{dh}{\sqrt{2\pi}}\, e^{-\frac{1}{2}h^2}\left[\Phi\left(\frac{\sigma(n_d)}{\sigma(n_{d'})}h - \frac{a_d - a_{d'}}{\sigma(n_{d'})}\right)\right]^{D-1} \quad (2.11) \end{aligned}$$

where Φ is the normal cumulative density function.

Further simplification can be made when $\sigma(n_{d'}) \approx \sigma(n_d)$. The classification accuracy then becomes

$$p_{corr}(s(K)) = \int_{-\infty}^{\infty} \frac{dh}{\sqrt{2\pi}}\, e^{-\frac{1}{2}h^2}\left[\Phi\left(h + s(K)\right)\right]^{D-1}, \quad (2.12)$$

where the sensitivity for detecting the hot component d' from $\mathbf{h}(K)$ is defined:

$$s(K) := \frac{\mu(h_{d'}) - \mu(h_d)}{\sigma(h_d)} = \frac{a_{d'} - a_d}{\sigma(n_d)} = \frac{1}{\sigma(n_d)}. \tag{2.13}$$

For detection, the retrieval involves two steps: detecting whether an input item was integrated K time steps ago and identifying which symbol if one is detected. In this case, a rejection threshold, θ, is required, which governs the trade-off between the two error types: misses and false positives. If none of the components in $\mathbf{h}(K)$ exceed θ, then the readout will output that no item was stored. The detection accuracy is given by

$$\begin{aligned} & p^{\theta}_{corr}(s(K)) \\ &= p\left((h_{d'}(K) > h_d(K)\ \forall d \neq d') \wedge (h_{d'}(K) \geq \theta) | \mathbf{a}(M-K) = \delta_{d=d'}\right) p_s \\ &\quad + p\left((h_d(K) < \theta\ \forall d) | \mathbf{a}(M-K) = 0\right)(1-p_s), \end{aligned} \tag{2.14}$$

where p_s is the probably that $\mathbf{a}(m)$ is a nonzero signal. If the distribution of $\mathbf{h}(K)$ is close to gaussian, the two conditional probabilities of equation 2.14 can be computed as follows. The accuracy, given a nonzero input was applied, can be computed analogous to equation 2.12:

$$\begin{aligned} & p\left((h_{d'}(K) > h_d(K)\ \forall d \neq d') \wedge (h_{d'}(K) \geq \theta) | \mathbf{a}(M-K) = \delta_{d=d'}\right) \\ &= \int_{(\theta-1)s(K;Mp_s)}^{\infty} \frac{dh}{\sqrt{2\pi}} e^{-\frac{1}{2}h^2} \left[\Phi\left(h + s(K;Mp_s)\right)\right]^{D-1}. \end{aligned} \tag{2.15}$$

Note that equation 2.15 is of the same form as equation 2.12 but with different integration bounds. The second conditional probability in equation 2.14, for correctly detecting a zero input, can be computed by

$$p\left((h_d(K) < \theta\ \forall d) | \mathbf{a}(M-K) = 0\right) = \left[\Phi\left(\theta s(K;Mp_s)\right)\right]^{D}. \tag{2.16}$$

Special Cases

1. As a sanity check, consider the classification accuracy, equation 2.12, in the vanishing sensitivity regime, for $s \to 0$. The first factor in the integral, the gaussian, then becomes the inner derivative of the second factor, the cumulative gaussian raised to the $(D-1)$th power. With $s \to 0$, the integral can be solved analytically using the (inverse) chain rule to yield the correct chance value for the classification:

$$p_{corr}(s \to 0) = \frac{1}{D}\Phi(h)^D|_{-\infty}^{\infty} = \frac{1}{D}. \tag{2.17}$$

2. The case $D = 1$ makes sense for detection but not for classification retrieval. This case falls under classical signal detection theory (Peterson, Birdsall, & Fox, 1954), with s being the sensitivity index. The detection accuracy, equation 2.14, in this case becomes

$$p^{\theta}_{corr}(s(K; Mp_s); D = 1) = (1 - \Phi((\theta - 1)s(K; Mp_s)))\, p_s + \Phi(\theta s(K; Mp_s))\,(1 - p_s). \tag{2.18}$$

The threshold θ trades off miss and false alarm errors. Formulas 2.14 to 2.16 generalize signal detection theory to higher-dimensional signals (D).

3. Consider classification retrieval, equation 2.12, in the case $D = 2$. Since the (rescaled and translated) random variables $p(h_{d'}(K)) \sim \mathcal{N}(s, 1)$ and $p(h_d(K)) \sim \mathcal{N}(0, 1)$ (see Figure 2A) are uncorrelated, one can switch to a new gaussian variable representing their difference: $y := h_d(K) - h_{d'}(K)$ with $p(y) \sim \mathcal{N}(-s, 2)$ (see Figure 2B). Thus, for $D = 2$, one can compute equation 2.12 by just the normal cumulative density function (and avoiding the integration):

$$p_{corr}(s(K); D = 2) = p(y < 0) = \Phi\left(\frac{s(K)}{\sqrt{2}}\right). \tag{2.19}$$

The result, equation 2.19, is the special case $d = 1$ of table entry 10,010.8 in Owen's table of normal integrals (Owen, 1980).

In general, for $D > 2$ and nonzero sensitvity, the p_{corr} integral cannot be solved analytically, but can be numerically approximated to arbitrary precision (see Figure 14).

2.2.2 Accuracy in the High-Fidelity Regime. Next, we derive steps to approximate the accuracy in the regime of high-fidelity recall, following the rationale of previous analyses of VSA models (Plate, 2003; Gallant & Okaywe, 2013). This work showed that the accuracy of retrieval scales linearly with the number of neurons in the network (N). We will compare our analysis results with those of previous analyses and with simulation results.

We now try to apply to the case $D > 2$ what worked for $D = 2$ (see equation 2.19): that is, get rid of the integral in equation 2.12 by transforming to new variables $y_d = h_d - h_{d'}$ for each of the $D - 1$ distracters with $d \neq d'$. We can write $p(\mathbf{y}) \sim \mathcal{N}(-\mathbf{s}, \mathbf{\Sigma})$ with

$$\mathbf{s} = \begin{pmatrix} s \\ s \\ \dots \end{pmatrix} \in \mathbb{R}^{D-1}, \quad \mathbf{\Sigma} = \begin{pmatrix} 2 & 1 & 1 \\ 1 & 2 & \dots \\ 1 & \dots & 2 \end{pmatrix} \in \mathbb{R}^{(D-1)\times(D-1)}.$$

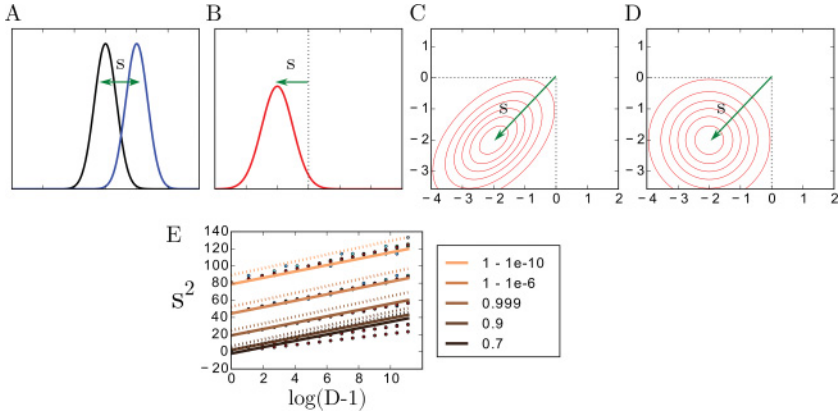

Figure 2: Approximating the retrieval accuracy in the high-fidelity regime. (A) The retrieval is correct when the value drawn from distribution $p(h_{d'})$ (blue) exceeds the values produced by $D-1$ draws from the distribution $p(h_d)$ (black). In the example, the sensitivity is $s = 2$. (B) When $D = 2$, the two distributions can be transformed into one distribution describing the difference of both quantities, $p(h_{d'} - h_d)$. (C) When $D > 2$, the $D-1$ random variables formed by such differences are correlated. Thus, in general, the multivariate cumulative gaussian integral, equation 2.20, cannot be factorized. The example shows the case $D = 3$, with the integration boundaries displayed by dashed lines. (D) However, for large s, that is, in the high-fidelity regime, the factorial approximation, equation 2.21, becomes quite accurate. The panel shows again the $D = 3$ example. (E) Linear relationship between the squared sensitivity and the logarithm of D. The numerically evaluated full theory (dots) coincides more precisely with the approximated linear theories (lines) when the accuracy is high (the accuracy is indicated by copper-colored lines; see the legend). The simpler linear theory (equation 2.24; dashed lines) matches the slope of the full theory but exhibits a small offset. The more elaborate linear theory (equation 2.25; solid lines) provides a quite accurate fit of the full theory for high accuracy values.

In analogy to the $D = 2$ case, equation 2.19, we can rewrite the result of equation 2.12 for $D > 2$ by

$$p_{corr} = p(y_d < 0 \; \forall d \neq d') = \Phi_{D-1}(\mathbf{s}, \mathbf{\Sigma}), \tag{2.20}$$

with multivariate cumulative gaussian, $\Phi_{D-1}(\mathbf{s}, \mathbf{\Sigma})$ (see the table entry $n0, 010.1$ in Owen's table of normal integrals; Owen, 1980).

The multivariate cumulative distribution, equation 2.20, would factorize, but only for uncorrelated variables, when the covariance matrix $\mathbf{\Sigma}$ is diagonal. The difficulty with $D > 2$ is that the multiple y_d variables are correlated: $\mathrm{Cov}(y_i, y_j) = E((y_i - s)(y_j - s)) = 1$. The positive uniform

off-diagonal entries in the covariance matrix means that the covariance ellipsoid of the y_d's is aligned with the $(1, 1, 1, \ldots)$ vector and thus also with the displacement vector $\mathbf{s}$ (see Figure 2C). In the high signal-to-noise regime, the integration boundaries are removed from the mean, and the exact shape of the distribution should not matter so much (see Figure 2D). Thus, the first step takes the factorized approximation (FA) to the multivariate gaussian to approximate p_{corr} in the high signal-to-noise regime:

$$p_{corr:\ FA} = \left[\Phi\left(\frac{s}{\sqrt{2}}\right)\right]^{D-1}. \tag{2.21}$$

Note that for $s \to 0$ in the low-fidelity regime, this approximation fails; the chance probability is $1/D$ when $s \to 0$ (see equation 2.17), but equation 2.21 yields 0.5^{D-1}, which is much too small for $D > 2$.

For an analytic expression, an approximate formula is needed for the one-dimensional cumulative gaussian, which is related to the complementary error function by $\Phi(x) = 1 - \frac{1}{2}\mathrm{erfc}(x/\sqrt{2})$. A well-known exponential upper bound on the complementary error function is the Chernoff-Rubin bound (CR; Chernoff, 1952). Later work (Jacobs, 1966; Hellman & Raviv, 1970) produced a tightened version of this bound: $\mathrm{erfc}(x) \leq B_{CR}(x) = e^{-x^2}$. Using $x = s/\sqrt{2}$, we obtain $B_{CR}(x/\sqrt{2}) = e^{-s^2/4}$, which can be inserted into equation 2.21 as the next step to yield an approximation of p_{corr}:

$$p_{corr:\ FA-CR} = \left[1 - \frac{1}{2}e^{-s^2/4}\right]^{D-1}. \tag{2.22}$$

With a final approximation step, using the local error expansion (LEE) $e^x = 1 + x + \ldots$ when x is near 0, we can set $x = -\frac{1}{2}e^{-s^2/4}$ and rewrite

$$p_{corr:\ FA-CR-LEE} = 1 - \frac{1}{2}(D-1)e^{-s^2/4}. \tag{2.23}$$

Solving for s^2 provides a simple law relating the sensitivity with the input dimension,

$$s^2 = 4\left[\ln(D-1) - \ln(2\epsilon)\right], \tag{2.24}$$

where $\epsilon := 1 - p_{corr}$.

The approximation, equation 2.24, is quite accurate (see Figure 2E, dashed lines) but not tight. Even if equation 2.21 was tight in the high-fidelity regime, there would still be a discrepancy because the CR bound is not tight. This problem of the CR bound has been noted for a long time, enticing efforts to derive tight bounds, usually involving more complicated multiterm expressions (e.g., Chiani, Dardari, & Simon, 2003). Quite

recently, Chang, Cosman, and Milstein (2011) studied one-term bounds of the complementary error function of the form $B(x;\alpha,\beta) := \alpha e^{-\beta x^2}$. First, they proved that there exists no parameter setting for tightening the original Chernoff-Rubin upper bound. Second, they reported a parameter range where the one-term expression becomes a lower bound: $\text{erfc}(x) \geq B(x;\alpha,\beta)$ for $x \geq 0$. The lower bound becomes the tightest with $\beta = 1.08$ and $\alpha = \sqrt{\frac{2e}{\pi}}\frac{\sqrt{\beta-1}}{\beta}$. This setting approximates the complementary error function as well as an eight-term expression derived in Chiani et al. (2003). Following Chang et al. (2011), we approximate the cumulative gaussian with the Chang bound (Ch) and follow the same FA and LEE steps to derive a tighter linear fit to the true numerically evaluated integral,

$$s^2 = \frac{4}{\beta}\left[\ln(D-1) - \ln(2\epsilon) + \ln\left(\sqrt{\frac{2e}{\pi}}\frac{\sqrt{\beta-1}}{\beta}\right)\right], \tag{2.25}$$

with $\beta = 1.08$. This law fits the full theory in the high-fidelity regime (see Figure 2E, solid lines), but it is not as accurate for smaller sensitivity values.

2.2.3 Information content and memory capacity. Memory capacity for symbolic input. The information content (Feinstein, 1954) is defined as the mutual information between the true sequence and the sequence retrieved from the superposition state $\mathbf{x}(M)$. The mutual information between the individual item that was stored K time steps ago ($a_{d'}$) and the item that was retrieved ($\hat{a}_d$) is given by

$$I_{item} = D_{KL}\left(p(\hat{a}_d, a_{d'}) \,||\, p(\hat{a}_d)p(a_{d'})\right) = \sum_{d}^{D}\sum_{d'}^{D} p(\hat{a}_d, a_{d'}) \log_2\left(\frac{p(\hat{a}_d, a_{d'})}{p(\hat{a}_d)p(a_{d'})}\right),$$

where $D_{KL}(p \,||\, q)$ is the Kullback-Leibler divergence (Kullback & Leibler, 1951).

For discrete input sequences, because the sequence items are chosen uniformly random from the set of D symbols, both the probability of a particular symbol as input and the probability of a particular symbol as the retrieved output are the same: $p(\hat{a}_d) = p(a_{d'}) = 1/D$. The $p_{corr}(s(K))$ integral evaluates the conditional probability that the output item is the same as the input item:

$$p(\hat{a}_{d'}|a_{d'}) = \frac{p(\hat{a}_{d'}, a_{d'})}{p(a_{d'})} = p_{corr}(s(K)).$$

To evaluate the $p(\hat{a}_d, a_{d'})$ $\forall d \neq d'$ terms, $p_{corr}(s(K))$ is needed to compute the probability of choosing the incorrect symbol given the true input. The

probability that the symbol is retrieved incorrectly is $1 - p_{corr}(s(K))$, and each of the $D - 1$ distracters is equally likely to be the incorrectly retrieved symbol, thus:

$$p(\hat{a}_d|a_{d'}) = \frac{p(\hat{a}_d, a_{d'})}{p(a_{d'})} = \frac{1 - p_{corr}(s(K))}{D - 1} \quad \forall\, d \neq d'.$$

Plugging these into the mutual information and simplifying,

$$\begin{aligned} I_{item}(p_{corr}(K)) &= p_{corr}(s(K)) \log_2 (p_{corr}(s(K))D) \\ &\quad + (1 - p_{corr}(s(K))) \log_2 \left(\frac{D}{D-1} (1 - p_{corr}(s(K))) \right) \\ &= D_{KL} \left(\mathcal{B}_{p_{corr}(s(K))} \,||\, \mathcal{B}_{\frac{1}{D}} \right), \end{aligned} \tag{2.26}$$

where $\mathcal{B}_p := \{p, 1 - p\}$ is the Bernoulli distribution. Note that the mutual information per item can be expressed as the Kullback-Leibler divergence between the actual recall accuracy p_{corr} and the recall accuracy achieved by chance, $1/D$.

The total mutual information is the sum of the information for each item in the full sequence:

$$I_{total} = \sum_{K=1}^{M} I_{item}(p_{corr}(s(K))) = \sum_{K=1}^{M} D_{KL} \left(\mathcal{B}_{p_{corr}(s(K))} \,||\, \mathcal{B}_{\frac{1}{D}} \right). \tag{2.27}$$

Note that if the accuracy is the same for all items, then $I_{total} = M\, I_{item}(p_{corr})$, and by setting $p_{corr} = 1$, one obtains the entire input information: $I_{stored} = M \log_2(D)$.

Memory capacity for analog input. For analog inputs, we can compute the information content if the components of input vectors are independent with gaussian distribution, $p(a_{d'}(m)) \sim \mathcal{N}(0, 1)$. In this case, distributions of the readout $p(\hat{a}_{d'})$ and the joint between input and readout $p(a_{d'}, \hat{a}_{d'})$ are also gaussian. Therefore, the correlation between $p(a_{d'})$ and $p(\hat{a}_{d'})$ is sufficient to compute the information (Gel'fand & Yaglom, 1957), with $I_{item} = -\frac{1}{2} \log_2(1 - \rho^2)$, where ρ is the correlation between the input and output. There is a simple relation between the signal correlation and the SNR r (9): $\rho = \sqrt{r/(r+1)}$, which gives the total information:

$$I_{total} = \frac{1}{2} \sum_{d'}^{D} \sum_{K}^{M} \log_2 (r(K) + 1). \tag{2.28}$$

The information content of a network is I_{total}/N in units bits per neuron. The memory capacity is then the maximum of I_{total}/N (27, 28) over all

parameter settings of the network. The network has extensive memory when I_{total}/N is positive, as N grows large.

2.3 VSA Indexing and Readout of Symbolic Input Sequences. In this section, we analyze the network model, equation 2.1, with linear neurons, $f(\mathbf{x}) = \mathbf{x}$ and without neuronal noise. After a reset to $\mathbf{x}(0) = 0$, the network receives a sequence of M discrete inputs. Each input is a one-hot vector, representing one of D symbols; we will show examples with alphabet size of $D = 27$, representing the 26 English letters and the space character. The readout, equation 2.2, involves the matrix $\mathbf{V}(K) = c^{-1}\langle \mathbf{a}(M-K)\mathbf{x}(M)^\top \rangle = c^{-1}\mathbf{W}^K\mathbf{\Phi}$ and the winner-take-all function, with $c = E_{\mathbf{\Phi}}(x^2)N$ a scaling constant. This setting is important because, as we will show in the next section, it can implement the working memory operation in various VSA models from the literature.

In this case, a sequence of inputs $\{\mathbf{a}(1), \ldots, \mathbf{a}(M)\}$ into the RNN, equation 2.1, produces the following memory vector:

$$\mathbf{x}(M) = \sum_{m=1}^{M} \mathbf{W}^{M-m}\mathbf{\Phi}\mathbf{a}(m). \tag{2.29}$$

Under the conditions 2.4 to 2.7, the linear part of the readout, equation 2.8, results in a sum of N independent random variables:

$$h_d(K) = \sum_{i=1}^{N} \left(\mathbf{V}_d(K)^\top \mathbf{x}(M)\right)_i = c^{-1}\sum_{i=1}^{N} (\mathbf{\Phi}_d)_i (\mathbf{W}^{-K}\mathbf{x}(M))_i = c^{-1}\sum_{i=1}^{N} z_{d,i}. \tag{2.30}$$

Note that under the conditions 2.4 to 2.7, each $z_{d,i}$ is independent, and thus h_d is a gaussian by the central limit theorem for large N. The mean and variance of h_d are given by $\mu(h_d) = c^{-1}N\mu(z_{d,i})$ and $\sigma^2(h_d) = c^{-1}N\sigma^2(z_{d,i})$.

The quantity $z_{d,i}$ in equation 2.30 can be written as

$$\begin{aligned} z_{d,i} &= (\mathbf{\Phi}_d)_i (\mathbf{W}^{-K}\mathbf{x}(M))_i \\ &= \begin{cases} (\mathbf{\Phi}_{d'})_i (\mathbf{\Phi}_{d'})_i + \displaystyle\sum_{m \neq (M-K)}^{M} (\mathbf{\Phi}_{d'})_i (\mathbf{W}^{M-K-m}\mathbf{\Phi}_{d'})_i & \text{if } d = d' \\ \displaystyle\sum_{m}^{M} (\mathbf{\Phi}_d)_i (\mathbf{W}^{M-K-m}\mathbf{\Phi}_{d'})_i & \text{otherwise} \end{cases}. \end{aligned} \tag{2.31}$$

Given the conditions 2.4 to 2.7, the moments of $z_{d,i}$ can be computed:

$$\mu(z_{d,i}) = \begin{cases} E_{\mathbf{\Phi}}(x^2) + (M-1)E_{\mathbf{\Phi}}(x)^2 & \text{if } d = d' \\ ME_{\mathbf{\Phi}}(x)^2 & \text{otherwise} \end{cases}, \tag{2.32}$$

$$\sigma^2(z_{d,i}) = \begin{cases} V_{\mathbf{\Phi}}(x^2) + (M-1)V_{\mathbf{\Phi}}(x)^2 & \text{if } d = d' \\ MV_{\mathbf{\Phi}}(x)^2 & \text{otherwise} \end{cases}, \tag{2.33}$$

with $E_{\mathbf{\Phi}}(x)$, $V_{\mathbf{\Phi}}(x)$ being the mean and variance of $p_{\mathbf{\Phi}}(x)$, the distribution of a component in the codebook $\mathbf{\Phi}$, as defined by equation 2.4.

Note that with linear neurons and unitary recurrent matrix, the argument K can be dropped because there is no recency effect and all items in the sequence can be retrieved with the same accuracy.

For networks with N large enough, $p(h_d(K)) \sim \mathcal{N}(c^{-1}N\mu(z_{d,i}), c^{-1}N\sigma^2(z_{d,i}))$. By inserting $\mu(h_d)$ and $\sigma(h_d)$ into equation 2.11, the accuracy then becomes

$$p_{corr} = \int_{-\infty}^{\infty} \frac{dh}{\sqrt{2\pi}} e^{-\frac{1}{2}h^2} \times \left[\Phi\left(\sqrt{\frac{M}{M - 1 + V_{\mathbf{\Phi}}(x^2)/V_{\mathbf{\Phi}}(x)^2}}\, h + \sqrt{\frac{N}{M - 1 + V_{\mathbf{\Phi}}(x^2)/V_{\mathbf{\Phi}}(x)^2}} \right) \right]^{D-1}. \tag{2.34}$$

Analogous to equation 2.12 for large M, the expression simplifies further to

$$p_{corr}(s) = \int_{-\infty}^{\infty} \frac{dh}{\sqrt{2\pi}} e^{-\frac{1}{2}h^2} \left[\Phi\left(h + s\right)\right]^{D-1} \quad \text{with } s = \sqrt{\frac{N}{M}}. \tag{2.35}$$

Interestingly, expression 2.35 is independent of the statistical moments of the coding vectors and thus applies to any distribution of coding vectors $p_{\mathbf{\Phi}}(x)$ equation 2.4. Since s is the ratio of N to M, it is easy to see that this network will have extensive capacity when s is held constant: $M = \beta N$:

$$\begin{aligned} \frac{I_{total}}{N} &= \frac{1}{N} \sum_{K=1}^{\beta N} I_{item}\left(p_{corr}\left(\sqrt{\frac{N}{\beta N}} \right) \right) \\ &= \beta I_{item}\left(p_{corr}\left(\sqrt{\frac{1}{\beta}} \right) \right) \\ &= const. \end{aligned} \tag{2.36}$$

However, it is a complex relationship between the parameter values that actually maximizes the mutual information. We will explore this in section 2.3.5. But first, in sections 2.3.1 to 2.3.4, we will show that $s = \sqrt{N/M}$, or a simple rescaling of it, describes the readout quality in many different VSA models.

2.3.1 VSA Models from the Literature. Many connectionist models from the literature can be directly mapped to equations 2.1 and 2.2 with the settings described at the beginning of section 2.3. In the following, we will describe various VSA models and the properties of the corresponding encoding matrix $\mathbf{\Phi}$ and recurrent weight matrix $\mathbf{W}$. We will determine the moments of the code vectors required in equation 2.34 to estimate the accuracy with in the general case (for small M). For large M values, we will show that all models perform similarly, and the accuracy can predicted by the universal sensitivity formula $s = \sqrt{N/M}$.

In hyperdimensional computing (HDC; Gayler, 1998; Kanerva, 2009), symbols are represented by N-dimensional random i.i.d. bipolar high-dimensional vectors (hypervectors), and referencing is performed by a permutation operation (see section 4.1.1). Thus, network 2.1 implements encoding according to HDC when the components of the encoding matrix $\mathbf{\Phi}$ are bipolar uniform random i.i.d. variables $+1$ or -1; that is, their distribution is a uniform Bernoulli distribution: $p_{\mathbf{\Phi}}(x) \sim \mathcal{B}_{0.5} : x \in \{-1, +1\}$, and $\mathbf{W}$ is a permutation matrix, a special case of a unitary matrix.

With these settings, we can compute the moments of $z_{d,i}$. We have $E_{\mathbf{\Phi}}(x^2) = 1$, $E_{\mathbf{\Phi}}(x) = 0$, $V_{\mathbf{\Phi}}(x^2) = 0$, and $V_{\mathbf{\Phi}}(x) = 1$, which can be inserted in equation 2.34 to compute the retrieval accuracy. For large M, the retrieval accuracy can be computed using equation 2.35. We implemented this model and compared multiple simulation experiments to the theory. The theory predicts the simulations precisely for all parameter settings of N, D, and M (see Figures 3A and 3B).

In holographic reduced representation (HRR) (Plate, 1993, 2003), symbols are represented by vectors drawn from a gaussian distribution with variance $1/N$: $p_{\mathbf{\Phi}}(x) \sim \mathcal{N}(0, 1/N)$. The binding operation is performed by circular convolution, and trajectory association can be implemented by binding each input symbol to successive convolutional powers of a random *key* vector, $\mathbf{w}$. According to Plate (1995), the circular convolution operation can be transformed into an equivalent matrix multiply for a fixed vector by forming the circulant matrix from the vector (i.e., $\mathbf{w} \circledast \mathbf{\Phi}_d = \mathbf{W}\mathbf{\Phi}_d$). This matrix has elements $W_{ij} = w_{(i-j)\%N}$ (where the subscripts on $\mathbf{w}$ a are interpreted modulo N). If $||\mathbf{w}|| = 1$, the corresponding matrix is unitary. Thus, HRR trajectory association can be implemented by an RNN with a recurrent circulant matrix and encoding matrix with entries drawn from a normal distribution. The analysis described for HDC carries over to HRR, and the error probabilities can be computed through the statistics of $z_{d,i}$, with $E_{\mathbf{\Phi}}(x) = 0$, $E_{\mathbf{\Phi}}(x^2) = 1/N$ giving $\mu(z_{d,i}) = (1/N)\delta_{d=d'}$, and with $V_{\mathbf{\Phi}}(x)^2 = 1/N$,

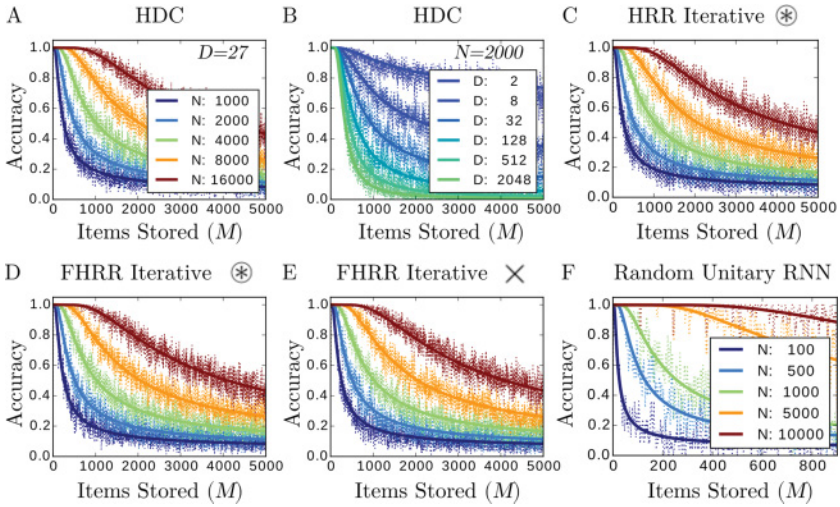

Figure 3: Classification retrieval accuracy: Theory and simulation experiments. The theory (solid lines) matches the simulation results (dashed lines) of the sequence recall task for a variety of VSA frameworks. Alphabet length in all panels except panel B is $D = 27$. (A) Accuracy of HDC code as a function of the number of stored items for different dimensions N of the hypervector. (B) Accuracy of HDC with different D and for constant $N = 2000$. (C) Accuracy of HRR code and circular convolution as binding mechanism. (D) Accuracy of FHRR code and circular convolution as the binding mechanism. (E) Accuracy of FHRR using multiply as the binding mechanism. (F) Accuracy achieved with random encoding and random unitary recurrent matrix also performs according to the same theory.

$V_{\mathbf{\Phi}}(x^2) = 2/N$ giving $\sigma^2(z_{d,i}) = (M + \delta_{d=d'})/N$. We compare simulations of HRR to the theoretical predictions in Figure 3C.

The *Fourier holographic reduced representation* (FHRR) (Plate, 2003) framework uses complex hypervectors as symbols, where components lay on the complex unit circle and have random phases: $(\mathbf{\Phi}_d)_i = e^{i\phi}$, with a phase angle drawn from the uniform distribution $p(\phi) \sim \mathcal{U}(0, 2\pi)$. The network implementation uses complex vectors of a dimension of $N/2$. Since each vector component is complex, there are N numbers to represent: one for the real part and one for the imaginary part (Danihelka et al., 2016). The first $N/2$ rows of the input matrix $\mathbf{\Phi}$ act on the real parts, and the second $N/2$ act on the imaginary part. Trajectory association can be performed with a random vector with $N/2$ complex elements acting as the key, raising the key to successive powers and binding this with each input sequentially. In FHRR, both element-wise multiply or circular convolution can be used as the binding operation, and trajectory association can be performed to encode the

letter sequence with either mechanism (see section 4.1.2 for further details). These are equivalent to an RNN with the diagonal of $\mathbf{W}$ as the key vector or as $\mathbf{W}$ being the circulant matrix of the key vector.

Given that each draw from $\mathcal{C}$ is a unitary complex number $z = (\cos(\phi), \sin(\phi))$ with $p(\phi) \sim \mathcal{U}(0, 2\pi)$, the statistics of $z_{d,i}$ are given by $E_{\mathbf{\Phi}}(x^2) = E(\cos^2(\phi)) = 1/2$, $[E_{\mathbf{\Phi}}(x)]^2 = E(\cos(\phi))^2 = 0$, giving $\mu(z_{d,i}) = \delta_{d=d'}/2$. For the variance, let $z_1 = (\cos(\phi_1), \sin(\phi_1))$ and $z_2 = (\cos(\phi_2), \sin(\phi_2))$. Then $z_1^\top z_2 = \cos(\phi_1)\cos(\phi_2) + \sin(\phi_1)\sin(\phi_2) = \cos(\phi_1 - \phi_2)$. Letting $\phi_* = \phi_1 - \phi_2$, it is easy to see that it also the case that $p(\phi_*) \sim \mathcal{U}(0, 2\pi)$. Therefore, $V_{\mathbf{\Phi}}(x)^2 = \mathrm{Var}(\cos(\phi_*))^2 = 1/4$ and $V_{\mathbf{\Phi}}(x^2) = 0$ giving $\sigma^2(z_{d,i}) = (M - \delta_{d=d'})/4$. Again we simulate such networks and compare to the theoretical results (see Figures 3D and 3E).

A random unitary matrix acting as a binding mechanism has also been proposed in the *matrix binding with additive terms* framework (MBAT) (Gallant & Okaywe, 2013). Our theory also applies to equivalent RNNs with random unitary recurrent matrices (created by QR decomposition of random gaussian matrix), with the same $s = \sqrt{N/M}$ (see Figure 3F). Picking an encoding matrix $\mathbf{\Phi}$ and unitary recurrent matrix $\mathbf{W}$ at random satisfies the required assumptions 2.4 to 2.7 with high probability when N is large.

2.3.2 Sparse Input Sequences. We next analyze detection retrieval of sparse input sequences, in which the input data vector $\mathbf{a}(m)$ is nonzero only with some probability p_s. The readout must first decide whether an input was present and determine its identity if present. With a random input matrix, linear neurons, and a unitary recurrent matrix, the sensitivity is $s = \sqrt{N/(Mp_s)}$. The crosstalk noise increments only when the input $\mathbf{a}(m)$ generates a one-hot vector. The threshold setting trades off hit and correct rejection accuracy (miss and false-positive error). We illustrate this in Figure 4A using equations 2.14 to 2.16 describing retrieval accuracy. The readout performance for sparse data sequences depends only on the product Mp_s. Thus, it appears possible to recover memory items with high sensitivity even for large sequence lengths $M > N$ if p_s is very small. However, our theoretical result requires assumptions 2.4 to 2.7, which break down for extremely long sequences. The result does not account for this breakdown and is optimistic for this scenario. Previous results that consider such extremely sparse input sequences have used the methods of compressed sensing and sparse inference (Ganguli & Sompolinsky, 2010; Charles, Yap, & Rozell, 2014; Charles, Yin, & Rozell, 2017), and show that recovering sparse input sequences with $M > N$ is possible.

2.3.3 Sparse Codebook. Often neural activity is characterized as sparse, and some VSA models use sparse codebooks. Several studies point to sparsity as an advantageous coding strategy for connectionist models (Rachkovskij, 2001). Sparse codes can be studied within our framework, equation 2.1, with a sparse input matrix—a random matrix in which

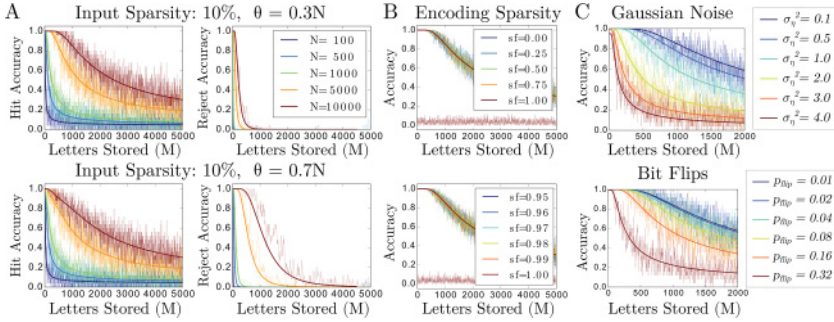

Figure 4: Detection retrieval accuracy, encoding sparsity and noise. (A) Retrieval of a sparse input sequence ($p_s = 0.9$, 10% chance for a zero vector). The hit and correct rejection performance for simulated networks (dashed lines) with different detection thresholds matches the theory (solid lines): a rejection is produced if $h_d < \theta \ \forall d$. (B) Performance is not affected by the level of encoding sparsity until catastrophic failing when all elements are 0. C. Simulations (dashed lines) match theory (solid lines) for networks corrupted by gaussian noise (top) and random bit-flips (bottom).

elements are zeroed out with a probability referred to as the *sparseness factor* (sf) Sparsity in the codebook affects both signal and the noise equally and cancels out to produce the same sensitivity, $s = \sqrt{N/M}$, as with a nonsparse input matrix. Thus, sparsity essentially has no effect on the capacity of the memory, up to the catastrophic point of sparsity where entire columns in the input matrix become zero (see Figure 4B).

2.3.4 Neuronal Noise. Here, we consider the case where each neuron experiences i.i.d. gaussian noise in each time step in addition to the data input. The effect of the noise depends on the ratio between noise variance and the variance of a component in the code vectors $V_{\mathbf{\Phi}}$. The sensitivity with neuronal noise is

$$s = \sqrt{\frac{N}{M(1 + \sigma_\eta^2 / V_{\mathbf{\Phi}})}}. \tag{2.37}$$

Thus, noise accumulation only scales s by a constant factor.

There are other ways to model noise in the network. For the case where there is only white noise added during the retrieval operation, it is easy to see that this noise will be added to the variance of $z_{d,i}$, giving $s = \sqrt{N/(M + \sigma_\eta^2 / V_{\mathbf{\Phi}})}$. If the noise was instead like a bit-flip in readout hypervector, with the probability of bit-flip p_f, then this gives $s = \sqrt{\frac{N(1-2p_f)^2}{M+2p_f}}$.

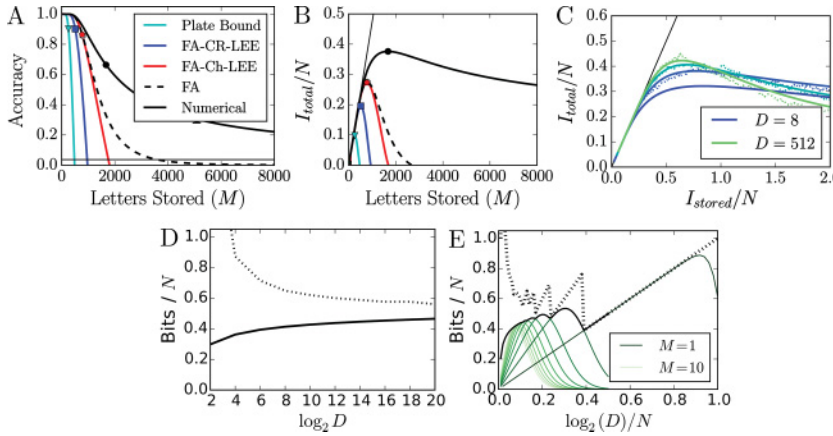

Figure 5: Information content and memory capacity. (A) Approximations of retrieval accuracy derived in section 2.2.2 and Plate (1993) are compared to the numerically evaluated accuracy (p_{corr}). The approximations underestimate the accuracy in the low-fidelity regime ($D = 27$, $N = 10{,}000$). (B) Total information content retrieved and memory capacity (solid points). High-fidelity retrieval recovers nearly all of the stored information (thin black line, $I_{stored} = M \log_2 D$, equation 2.27), but the true memory capacity is somewhat into the low-fidelity regime. (C) Retrieved information measured in simulations (dashed lines) compared to the predictions of the theory (solid lines). The memory capacity is dependent on D. (D) Memory capacity as a function of D (solid line) and information of the input sequence at retrieval maximum (I_{stored}, dashed). (E) Maximum information retrieved (solid black line) and total information stored (I_{stored}, dashed), where D is a significant fraction of 2^N ($N = 100$). The retrieved information for fixed sequence lengths $M = \{1, \ldots, 10\}$ are plotted (green lines of different shades). For $M = 1$, retrieved and stored information come close; with larger M, the gap grows.

Finally, with these derivations of s and equation 2.12, the empirical performance of simulated neural networks is predicted (see Figure 4C).

2.3.5 Memory Capacity of VSAs with Symbolic Input. The original estimate for the capacity of distributed random codes (Plate, 1993) considered a slightly different setup (see section 4.2.2) but follows similar ideas as the FA-CR-LEE high-fidelity approximation, equation 2.24, and we reformulated the Plate (1993) derivation to compare with our analysis. This work first showed that random indexing has linear extensive capacity and that the memory capacity is at least 0.16 bits per neuron. Figure 5A compares the approximations, equations 2.21, 2.23, and 2.24, with the full theory evaluated numerically, equation 2.12. These approximations are good in the

high-fidelity regime, where p_{corr} is near 1 but underestimate the performance in the low-fidelity regime.

With the relationship $s = \sqrt{N/M}$, the information contained in the activity state $\mathbf{x}(M)$ can be calculated. We compare the total information, equation 2.27, based on the approximations with the true information content determined by numeric evaluation of p_{corr} (see Figure 5B). In the linear scenario with unitary weight matrix, p_{corr} has no dependence on K, and so the total information in this case is simply $I_{total} = MI_{item}$, equation 2.27.

In the high-fidelity regime with $p_{corr} = 1 - \epsilon$ and small ϵ, we can estimate, with equation 2.24,

$$\frac{I_{total}}{N} = \frac{M}{N} D_{KL}\left(\mathcal{B}_{1-\epsilon} \,||\, \mathcal{B}_{\frac{1}{D}}\right) \approx \frac{M \log(D)}{N \log(2)} \approx \frac{\log(D)}{4 \log(2)(\log(D-1) - \log(2\epsilon))}. \tag{2.38}$$

We can see that for any fixed, finite D, the information per neuron depends on the admitted error ϵ and vanishes for $\epsilon \to 0$. If the alphabet size D is growing with N and for fixed small error ϵ, the asymptotic capacity value is $1/(4 \ln 2) = 0.36$. Our best high-fidelity approximation, equation 2.25, increases the total capacity bound above previous estimates to 0.39.

Results for the case of a finite moderate-sized alphabet size ($D = 27$) are shown in Figure 5B. The novel and most accurate high-fidelity approximation, equation 2.25, predicts 0.27 bits per neuron; the simpler high-fidelity approximations substantially underestimate the capacity.

Importantly, however, our full theory shows that the true information maximum lies outside the high-fidelity regime. The maximum capacity for $D = 27$ is nearly 0.4 bits per neuron (see Figure 5B, black circle). Thus, the achievable capacity is about four times larger than previous analysis suggested.

In a wide range of D, our full theory precisely predicts the empirically measured total information in simulations of the random sequence recall task (see Figure 5C). The total information per neuron scales linearly with the number of neurons, and the maximum amount of information per element that can be stored in the network is dependent on D. The memory capacity increases with D, reaching over 0.5 bits per neuron for large D (see Figure 5D, solid line).

Capacity without superposition. As the alphabet size, D, grows superlinear in N (approaching 2^N), one needs to reduce M in order to maximize the memory capacity (see Figure 5E, dashed line). The theory breaks down when there is no superposition, that is, when $M = 1$. This case is different because there is no cross-talk, but for very large D and randomly generated code vectors, collisions arise, another source of retrieval errors. Collisions are coincidental duplication of code vectors. The theory presented so far can describe effects of crosstalk but not of collisions.

For completeness, we briefly address the case of $M = 1$ and very large D. This case without cross talk shows that it is possible to achieve the theoretically optimal capacity of 1 bit per neuron and that crosstalk immediately limits the achievable capacity. If code vectors are drawn i.i.d. with a uniform Bernoulli distribution $p_{\Phi}(x) \sim \mathcal{B}_{0.5} : x \in \{-1, +1\}$, then the probability of accurately identifying the correct code word is

$$p_{corr}^{M=1} = \sum_{C} p_C/(C+1), \tag{2.39}$$

where p_C is the probability of a vector having collisions with C other vectors in the codebook of D vectors, which is given by the binomial distribution,

$$p_C = \binom{D}{C} q^C (1-q)^{D-C}, \tag{2.40}$$

where $q = 1/2^N$ is the likelihood of a pair of vectors colliding. The collisions reduce the accuracy $p_{corr}^{M=1}$ to $(1 - 1/e) \approx 0.63$ for $D = 2^N$ in the limit $N \to \infty$. However, this reduction does not affect the asymptotic capacity. It is $I_{total}/N \to 1$ bits per neuron as $N \to \infty$, the same as for a codebook without collisions (see section 4.4.3). The effects of collisions at finite sizes N can be seen in Figures 5E and 18A.

In the presence of superposition, that is, for $M > 1$, the crosstalk noise becomes immediately the limiting factor for memory capacity. This is shown in Figure 5E for a small network of 100 neurons. For $M = 2$, the memory capacity drops to around 0.5 bits per neuron and decreases to about 0.2 bits per neuron for large M values (see Figure 5E, black line). The capacity curves for fixed values of M (5E, green lines) show the effect of crosstalk noise, which increases as more items are superposed (as M increases). For $M = 1$ (see Figure 5E, dark green line), equations 2.39 and 2.27 can be evaluated as D grows to 2^N.

2.4 Indexed Memory Buffers with Symbolic Input. With the linear encoding network described in the previous section, there is no recency effect; the readout of the most recent input stored is just as the accurate as readout of the earliest input stored. In contrast, a network with a contracting recurrent weight matrix (see section 2.4.1) or nonlinear neural activation functions (see sections 2.4.2 and 2.4.3) will have a recency effect. In this section, the theory will be developed to describe memory in encoding networks (see equation 2.1) with recency effects. We juxtapose the performance of networks with recency effects in our two memory tasks, reset memory and memory buffer. We show that reset memories yield optimal capacity in the absence of any recency effect (Lim & Goldman, 2011). However, recency effects can avoid catastrophic forgetting when the input sequence is infinite.

Through the recency effect, sequence items presented further back in time will be attenuated and eventually forgotten. Thus, the recency effect is a soft substitution of resetting the memory activity before the input of interest is entered. Further, we show that contracting recurrent weights and saturating neural activation functions have very similar behavior if their forgetting time constants τ are aligned (see section 2.4.4). Finally, we optimize the parameters of memory buffers for high memory capacity and show that extensive capacity (Ganguli et al., 2008) is achievable even in the presence of neuronal noise by keeping the forgetting time constant proportional to N.

2.4.1 Linear Neurons with Contracting Recurrent Weights. Reset memory. Consider a network of linear neurons in which the attenuation factor $0 < \lambda < 1$ contracts the network activity in each time step. After a sequence of M input symbols has been applied, the variance of $z_{d,i}$ is $\left(\frac{1-\lambda^{2M}}{1-\lambda^2}\right) V_{\mathbf{\Phi}}$, and the signal decays exponentially with $\lambda^K E_{\mathbf{\Phi}}(x^2)$. The sensitivity for recalling the input that was added K time steps ago is

$$s(K) = \lambda^K \sqrt{\frac{N(1-\lambda^2)}{1-\lambda^{2M}}}. \tag{2.41}$$

Thus, the sensitivity decays exponentially as K increases, and the highest retrieval accuracy is from the most recent item stored in memory. The accuracy (see Figure 6A1) and information per item (see Figure 6B1) based on this formula for $s(K)$ show the interdependence between the total sequence length (M) and the look-back distance (K) in the history.

In equation 2.41, the sensitivity is monotonically increasing as λ increases, and thus to maximize it for the Kth element in history given a finite set of M stored tokens, we would want to maximize λ or have the weight matrix remain unitary with $\lambda = 1$.[1] The memory capacity is maximized as $\lambda \to 1$ when M is finite (see Figure 6C1) and as D grows large (see Figure 6D1), and there is no benefit of contracting weights in reset memories.

Memory buffer. For an infinite stream of inputs, $M \to \infty$, the setting $\lambda = 1$ results in catastrophic forgetting. However with $\lambda < 1$, the memory can operate even in this case because past signals fade away and make room for storing new inputs. These networks with $\lambda < 1$, $f(x) = x$, and $\mathbf{W}$ unitary have been denoted *normal networks* (White et al., 2004).

The value of λ affects both the signal and the crosstalk noise. For large M, the noise variance is bounded by $\frac{1}{1-\lambda^2}$, and the network reaches its saturated

[1] If we allowed λ to be larger than 1, then memories from the past would grow in magnitude exponentially. This would mean higher SNR for more distant memories at the cost of lower SNR for recent memories. This would cause the network to explode; however, normalization could be used.

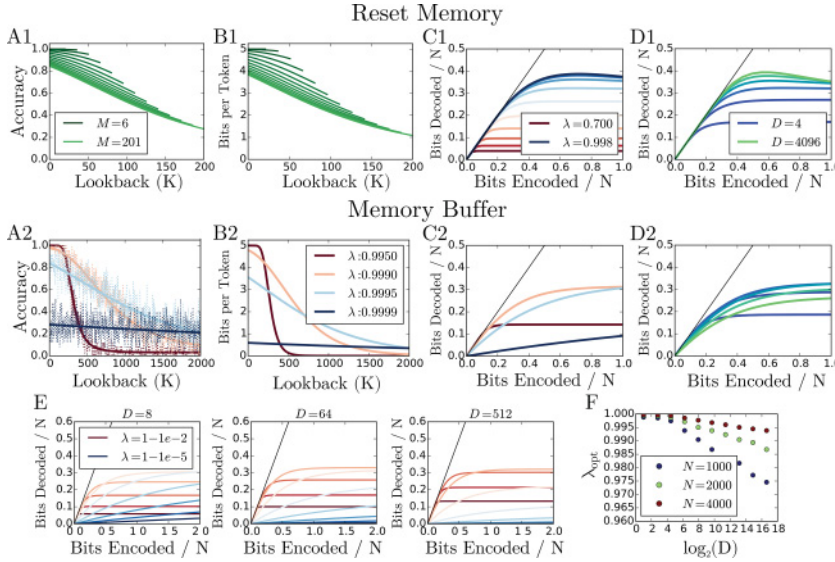

Figure 6: Linear network with contracting recurrent weights. (A1) Accuracy in networks with $\lambda < 1$. Multiple evaluations of p_{corr} are shown as a function of K for sequences of different lengths, M. ($\lambda = 0.996$, $N = 1000$). (B1) The information per item I_{item} also depends on K. (C1) The total retrieved information per neuron for different λ. The maximum is reached as λ approaches 1 when M is finite ($D = 64$; $N = 1000$). (D1) The retrieved information is maximized as D grows large ($\lambda = 0.988$). (A2) Accuracy in networks with $\lambda < 1$ as $M \to \infty$ ($N = 10{,}000$; $D = 32$). (B2) Information per item. (C2) Total information retrieved as a function of the total information stored for different λ. There is a λ that maximizes the information content for a given N and D ($D = 64$). (D2) Total information retrieved as a function of the total information stored for different D ($\lambda = 0.999$). Retrieved information is optimized by a particular combination of D and λ. (E) The total retrieved information per neuron versus the information stored per neuron for different D and λ over a wide range. As D increases, the information is maximized by decreasing λ. (F) Numerically determined λ_{opt} values that maximize the information content of the network with $M \to \infty$ for different N and D.

equilibrium state. The sensitivity for the Kth element back in time from the saturated state is

$$s(K) = \lambda^K \sqrt{N(1-\lambda^2)}. \tag{2.42}$$

The theory (see equations 2.42 and 2.12) predicts the performance of simulated networks with contracting recurrent weights that store a sequence of

symbols with $M \gg N$ (see Figure 6A2, solid lines) for different λ (see Figure 6A2, dashed lines). The information per item retrieved (see Figure 6B2) and the total information (see Figure 6C2) for different values of λ show the trade-off between fidelity and duration of storage. There is an ideal λ value that maximizes the memory capacity with $M \to \infty$ for given N and D. This ideal λ value differs depending on the alphabet size (D) (see Figure 6D2). For larger alphabets (meaning more bits per symbol), the inputs should be forgotten more quickly, and memorizing a shorter history optimizes the memory capacity (see Figure 6E). The values of λ that maximize the memory capacity were computed numerically; they drop with increasing N and D (see Figure 6F).

2.4.2 Neurons with Clipped-Linear Transfer Function. Squashing nonlinear neural activation functions induce a recency effect, similar to contracting recurrent weights. Consider equation 2.1 with $\lambda = 1$ and the clipped-linear activation function, $f(x) = f_\kappa(x)$, in which the absolute value of the activity of neurons is limited by κ:

$$f_\kappa(x) = \begin{cases} -\kappa & x \leq -\kappa \\ x & -\kappa < x < \kappa \\ \kappa & x \geq \kappa \end{cases} . \tag{2.43}$$

Clipping functions of this type with specific κ-values play a role in VSAs that constrain the activation of memory vectors, such as the binary-spatter code (Kanerva, 1996) or the binary sparse-distributed code (Rachkovskij, 2001).

We will analyze the HDC encoding scheme, a network with a bipolar random input matrix and the recurrent weights a permutation matrix. With this, the components of $\mathbf{x}$ will always assume integer values, and due to the clipping, the components will be confined to $\{-\kappa, -\kappa + 1, \ldots, \kappa\}$. As a consequence, $z_{d,i}$, defined as in equation 2.30, will also assume values limited to $\{-\kappa, \ldots, \kappa\}$. To compute s, we need to track the mean and variance of $z_{d,i}$. This requires iterating the Chapman-Kolmogorov equation 2.3. To do so, we introduce a vector $\mathbf{q}$ with $q_{\mathcal{J}(k)}(m) := p(z_{d,i}(m) = k)\ \forall k \in \{-\kappa, \ldots, \kappa\}$, which tracks the probability distribution of $z_{d,i}$. The probability of each of the integers from $\{-\kappa, \ldots, \kappa\}$ is enumerated in the $2\kappa + 1$ indices of the vector $\mathbf{q}$, and $\mathcal{J}(k) = k + \kappa$ is a bijective map from the values of $z_{d,i}$ to the indices of $\mathbf{q}$ with inverse $\mathcal{K} = \mathcal{J}^{-1}$. To compute the sensitivity of a particular recall, we need to track the distribution of $z_{d,i}$ with $\mathbf{q}$ before the item of interest is added, when the item of interest is added, and in the time steps after storing the item of interest. Note that κ is defined relative to the standard deviation of the codebook, $\kappa = \kappa^*/\sqrt{V_\Phi}$. A simple scaling can generalize the following analysis to account for codebooks with different variance.

Reset memory. At initialization $\mathbf{x}(0) = 0$, and so $q_j(0) = \delta_{\mathcal{K}(j)=0}$. For each time step that an input arrives in the sequence prior to the input of interest, a $+1$ or -1 will randomly add to $z_{d,i}$ up until the bounds induced by f_κ, and this can be tracked with the following diffusion of $\mathbf{q}$:

$$q_j(m+1) = \frac{1}{2}\begin{cases} q_j(m) + q_{j+1}(m) & \text{when } \mathcal{K}(j) = -\kappa \\ q_{j-1}(m) + q_j(m) & \text{when } \mathcal{K}(j) = \kappa \\ q_{j-1}(m) + q_{j+1}(m) & \text{otherwise.} \end{cases} \quad \forall\, m \neq M - K. \tag{2.44}$$

Once the vector of interest arrives at $m = M - K$, all entries in $z_{d,i}$ will have $+1$ added. This causes the probability distribution to skew:

$$q'_j(m+1) = \begin{cases} 0 & \text{when } \mathcal{K}(j) = -\kappa \\ q_j(m) + q_{j-1}(m) & \text{when } \mathcal{K}(j) = \kappa \\ q_{j-1}(m) & \text{otherwise} \end{cases} \quad m = M - K. \tag{2.45}$$

The $K - 1$ inputs following the input of interest will then again cause the probability distribution to diffuse further based on equation 2.44. Finally, $s(K)$ can be computed for this readout operation by calculating the mean and variance with $\mathbf{q}(M)$:

$$\mu(z_{d,i}) = \delta_{d=d'} \sum_{j=0}^{2\kappa} \mathcal{K}(j) q_j(M), \tag{2.46}$$

$$\sigma^2(z_{d,i}) = \sum_{j=0}^{2\kappa} (\mathcal{K}(j) - \mu(z_{d,i}))^2 q_j(M). \tag{2.47}$$

Memory buffer. For $M \to \infty$ the diffusion equation, 2.44, will reach a uniform equilibrium distribution with the values of $z_{d,i}$ uniformly distributed between $\{-\kappa, \ldots, \kappa\}$: $q_j(\infty) = 1/(2\kappa + 1)\ \forall j$. This means, as with contracting recurrent weights, the clipped-linear function bounds the noise variance of the saturated state. Here, the variance bound of $z_{d,i}$ is the variance of the uniform distribution, $((2\kappa + 1)^2 - 1)/12$. Thus, information can still be stored in the network even after being exposed to an infinite sequence of inputs. The sensitivity in the saturated state can be calculated with $M \to \infty$ by replacing in equation 2.45 $\mathbf{q}(m)$ with $\mathbf{q}(\infty)$, and then again using the diffusion equation 2.44 for the $K - 1$ items following the item of interest.

Figure 7 illustrates this analysis of the distribution of $z_{d,i}$. When the item of interest is added, the probability distribution is most skewed, and the signal degradation is relatively small. As more items are added later, the

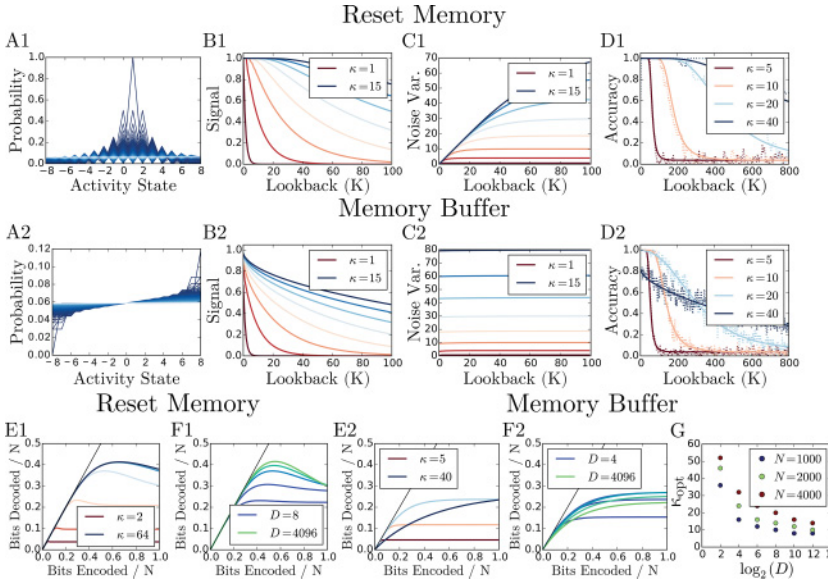

Figure 7: Capacity for neurons with clipped-linear transfer function. (A1) The probability distribution of $z_{d,i}$ for retrieval of the first sequence item, as the sequence length is increased. The distribution evolves according to equations 2.44 and 2.45; it begins at a delta function (dark blue) and diffuses to the uniform equilibrium distribution when M is large (light blue). (B1) The clipped-linear function causes the signal to degrade as more items are stored ($M = K$; $N = 5000$; $D = 27$). (C1) The variance of the distribution grows as more items are stored but is bounded. (D1) The accuracy theory fits empirical simulations, decoding the first input as more input symbols are stored (dashed lines; $M = K$; $N = 5000$; $D = 27$). (A2) The probability distribution of $\mathbf{z_{d,i}}$ for the memory buffer, that is, when $M \to \infty$. The most recent symbol encoded (dark blue) has the highest skew, and the distribution diffuses to the uniform equilibrium for readout further in the past (light blue). (B2) The signal is degraded from crosstalk and decays as a function of the look-back. (C2) The noise variance is already saturated and stays nearly constant. (D2) The accuracy exhibits a trade-off between fidelity and memory duration governed by κ. (E1) With reset memory, the information that can be decoded from the network reaches a maximum when κ is large ($D = 256$). (F1) The capacity increases with D ($\kappa = 20$). (E2) When $M \to \infty$, there is a trade-off between fidelity and memory duration; a particular κ value maximizes the retrieved information for a given D and N ($D = 256$). (F2) For a given memory duration ($\kappa = 20$), an intermediate D value maximizes the retrieved information. (G) The memory duration κ_{opt} that maximizes the retrieved information.

distribution diffuses to the uniform equilibrium and the signal decays to 0. The figure compares operation with the initial states corresponding to reset memory and memory buffer: empty (see Figures 7A1 to 7F1) and saturated (see Figures 7A2 to 7F2). Numerical optimization of memory capacity shows how κ_{opt} increases with N and decreases with D, the parameter when $M \to \infty$ (see Figure 7G).

2.4.3 Neurons with Squashing Nonlinear Transfer Function. The case when the neural transfer function $f(x)$ is a saturating or squashing function with $|f(x)|$ bounded by a constant also implies $z_{d,i}$ is bounded with $|z_{d,i}| \leq z_{\max}$. For any finite fixed error, one can choose an n large enough so that the distribution $p(z_{d,i} = k) = q_{\mathcal{J}(k)}$ can be approximated by discretizing the state space into $2n + 1$ equal bins in $\mathbf{q}$. Like the clipped-linear transfer function, one can construct a bijecteve map from values to indices and track $\mathbf{q}$ approximately using rounding to discretize the distribution, $\mathcal{J}(k) = \lfloor \frac{n}{z_{\max}}(k + z_{\max}) \rceil$, with inverse $\mathcal{K} = \mathcal{J}^{-1}$. The Kolmogorov equation, 2.3, simplifies in this discrete case and without neuronal noise to the following updates, one for encoding the signal, and one for encoding the distracters:

$$q_{j^*}(m+1) = \sum_{j=0}^{2n} \int dy \; p_{\mathbf{\Phi}}(y) q_j(m) \, \delta_{j^* = \mathcal{J}(f(\mathcal{K}(j)+y))} \quad \forall m \neq M - K. \tag{2.48}$$

The update for the signal, given at $m = M - K$, where we know that $\mathbf{\Phi}_{d'}$ was stored in the network is

$$q'_{j^*}(m+1) = \sum_{j=0}^{2n} \int dy \; p_{\mathbf{\Phi}}(y) q_j(m) \, \delta_{j^* = \mathcal{J}(f(\mathcal{K}(j)+y^2))} \quad m = M - K. \tag{2.49}$$

We illustrate our analysis for a network 2.1 with $\lambda = 1$ and $f(x) = \gamma \tanh(x/\gamma)$, where γ is a gain parameter. As in the previous section, the network implements HDC coding, so the codebook is a bipolar i.i.d. random matrix and the recurrent weights a permutation matrix. Our simulation experiments with this memory network examined both reset memories (see Figure 8, row 1) and memory buffers (see Figure 8, row 2). The iterative analysis (see equations 2.48 and 2.49) can be used to compute the sensitivity, and it predicts the experimentally observed readout accuracy very well. We find that a memory with the neural transfer function tanh possesses quite similar performance as a memory with the clipped-linear neural activation function.

2.4.4 Forgetting Time Constant and Extensive Capacity in Memory Buffers. We have analyzed different mechanisms of a recency effect in memory buffers, contracting weights, and squashing, nonlinear neural activation

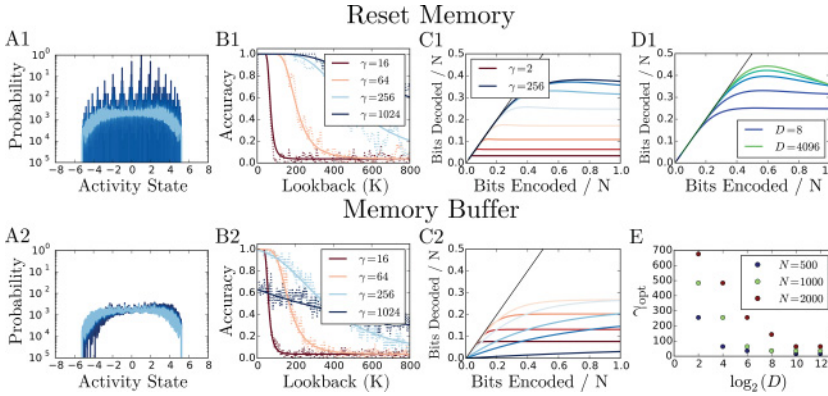

Figure 8: Capacity for neurons with saturating nonlinearity. (A1) The probability distribution of one term $z_{d,i}$ in the inner product used for retrieval of the first sequence item, as the sequence length is increased. The distribution begins as a delta function (dark blue) when only one input symbol is stored and approaches the equilibrium distribution when M is large (light blue). (B1) The accuracy theory (solid lines) correctly describes simulations (dashed lines) retrieving the first input as more inputs are stored ($M = K$; $N = 2000$; $D = 32$). (C1) Retrieved information as a function of the stored information. When M is finite, the maximum is reached for large γ ($D = 256$). (D1) The capacity increases as D increases ($\gamma = 64$). (A2) The probability distribution of $\mathbf{z_{d,i}}$ when a new item is entered at full equilibrium, that is, when $M \to \infty$. The distribution for the most recent input symbol possesses the highest skew (dark blue), and the distribution is closer to the uniform equilibrium (light blue) for input symbols encoded further back in the history. (B2) The accuracy exhibits a trade-off between fidelity and memory duration governed by γ. (C2) When M is large, there is a γ that maximizes the information content for a given D and N ($D = 256$). (E) Numerically computed γ_{opt} that maximizes the information content.

functions. Here we compare their properties and find parameters that optimize memory capacity.

For contracting weights, the forgetting time constant (τ) is defined from the exponential decay of the sensitivity λ^K in equation 2.42 by $\lambda = e^{-1/\tau}$:

$$\tau(\lambda) = -1/\log\lambda. \tag{2.50}$$

We compute the N that optimizes the capacity of the memory buffer for a desired time constant (see Figure 9A).

The forgetting time constants for nonlinear activation functions can be computed by equating the bound of the variance induced by the nonlinearity to the bound induced by contracting weights. For clipping, the noise variance is bounded by the variance of the uniform distribution,

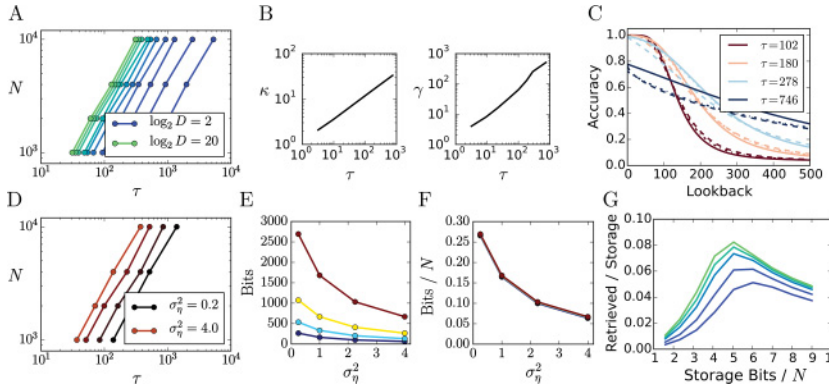

Figure 9: Buffer time constant and optimization. (A) The optimal N for given time constant τ and D. (B) The relationship between time constant and nonlinear activation parameters, κ and γ. (C) Accuracy comparison of contracting (solid), clipped-linear (dashed), and tanh (dotted) networks that share the same time constant, found from the bound on noise variance. (D) The optimal N and τ shifts in the presence of noise ($D = 32$). (E) The memory capacity decreases as noise increases ($N = [1K, 2K, 4K, 10K]$, blue to red). (F) When capacity is normalized by number of neurons, the curves in panel E collapse to a single curve, showing capacity per neuron to be constant with N and declining with neuronal noise. (G) Ratio between retrieved and stored information for the clipped HDC network. The ratio is optimized with four to five bits of resolution per element ($D = [8, 32, 256, 1024, 4096]$, dark to light).

$((2\kappa + 1)^2 - 1)/12$, which can be equated to the bound of contracting weights $1/(1 - \lambda^2)$. With equation 2.50, one obtains

$$\tau(\kappa) = \frac{-2}{\log\left(1 - \frac{3}{\kappa(\kappa+1)}\right)} \approx \frac{2}{3}\kappa^2 \approx \frac{2}{3}\frac{(\kappa^*)^2}{V_{\mathbf{\Phi}}}, \tag{2.51}$$

where the approximation holds for large κ.

Equating the bound of the noise variance to $1/(1 - \lambda^2)$ is a general technique to approximate the time constant for any nonlinear function with equation 2.50. For the tanh nonlinearity, we cannot compute analytically the forgetting time constant for a parameter value γ. Instead we use equations 2.48 and 2.49 to estimate its variance bound and equate it to $1/(1 - \lambda^2)$.

The relationships between τ, the clipping parameter κ (see equation 2.51), and the tanh gain parameter γ (numerically estimated) are not far from linear in logarithmic coordinates (see Figure 9B). When the forgetting time constants are matched, the accuracies of different recency mechanisms are not identical but quite similar (see Figure 9C).

With neuronal noise, the optimal forgetting time constant decreases with noise variance (see Figure 9D) and also the memorized information (see Figure 9E). Interestingly, the memory capacity is independent of N for a given noise variance (see Figure 9F), indicating that the memory buffer has extensive capacity. Since the effects of contracting weights and nonlinear neurons are similar, these networks can all achieve extensive memory in the presence of noise by keeping the time constant τ proportional to N.

With the clipped-linear activation function and HDC coding, for any finite setting of the clipping threshold κ, the number of bits required to represent or store the network state is finite and given by $I_{storage} = N \log_2(2\kappa + 1)$. One can now compute the ratio between the readout information and the information required to store the network state. In general, the amount of readout information increases with κ (see Figure 7E1) and with D (see Figure 7F1). However, as κ increases, so too do the number of bits required to represent the network. It turns out that there is an optimal ratio of about 0.08, achieved with neurons that represent about four to six bits, the exact value dependent on D (see Figure 9G).

2.5 VSA Indexing and Readout for Analog Input Vectors. The theory can be easily extended to the recall of coefficients of analog vectors. Rather than the input vector $\mathbf{a}(m)$ being a one-hot or zero vector, the input can be an arbitrary real vector, and we wish to store and retrieve a sequence of such analog vectors in the network. We can derive memory capacity under the assumption that the input vector is drawn independently from a normal distribution. In the following, the linear network with analog input is analyzed in two cases, operating as a reset memory (see section 2.5.1), and a memory buffer (see section 2.5.2).

2.5.1 Capacity of Reset Memories with Analog Inputs. A sequence of vectors with analog coefficients $\mathbf{a}(m)$ is encoded into the network state by equation 2.1 with a random input matrix $\mathbf{\Phi}$ and unitary recurrent weight matrix $\mathbf{W}$. We return to considering reset memories with linear neurons: $f(\mathbf{x}) = \mathbf{x}$. During the encoding, each coefficient is indexed with a pseudo-random key vector. To read out an individual coefficient, we use in equation 2.2 a linear readout function $g(\mathbf{h}) = \mathbf{h}$ and the readout matrix of VSA models: $\mathbf{V}(K) = c^{-1}\langle \mathbf{a}(M-K)\mathbf{x}(M)^\top \rangle = c^{-1}\mathbf{W}^K\mathbf{\Phi}$, with $c = NE_{\mathbf{\Phi}}(x^2)$. The readout variable can be decomposed into N contributions as in equation 2.30: $\hat{a}_{d'} = h_{d'} = \sum_i^N c^{-1} z_{d',i}$. For large enough N, $h_{d'}$ is distributed like a gaussian due to the central limit theorem. We can compute $z_{d',i}$ from equation 2.30:

$$\begin{aligned} z_{d',i} &= (\mathbf{\Phi}_{d'})_i (\mathbf{W}^{-K}\mathbf{x}(M))_i \\ &= (\mathbf{\Phi}_{d'})_i \left[(\mathbf{\Phi}_{d'})_i a_{d'}(M-K) \right] \end{aligned}$$

$$+ (\mathbf{\Phi}_{d'})_i \left[\sum_{d \neq d'}^{D} (\mathbf{\Phi}_d)_i a_d (M-K) + \sum_{m \neq (M-K)}^{M} \left(\mathbf{W}^{M-K-m} \left(\sum_{d}^{D} \mathbf{\Phi}_d a_d(m) \right) \right)_i \right]. \tag{2.52}$$

The signal and the noise term are on two lines in equation 2.52. In the expression $c^{-1} z_{d',i}$, the variance of the signal term is unity, and the resulting SNR is

$$r = \frac{\sigma^2(a_{d'})}{\sigma^2(n_{d'})} = \frac{N\sigma^2(a_{d'})}{\left(\sum_{d \neq d'} a_d^2 (M-K) + \sum_{m \neq (M-K)} \sum_d a_d^2(m) \right)} = \frac{N}{(MD-1)} \approx \frac{N}{MD}. \tag{2.53}$$

When neuronal noise is present, the SNR becomes

$$r = \frac{N}{MD} \left(\frac{1}{1 + \sigma_\eta^2 / (DV_{\mathbf{\Phi}})} \right). \tag{2.54}$$

If the input coefficients are all independent gaussian random variables, then the total information can be computed with equation 2.28:

$$\frac{I_{total}}{N} = \frac{MD}{2N} \log_2 (r+1) = \frac{\log_2 (r+1)}{2r} \left(\frac{1}{1 + \sigma_\eta^2 / (DV_{\mathbf{\Phi}})} \right). \tag{2.55}$$

Note that the memory capacity for analog input is a function of r. Thus, the memory capacity is extensive when MD is proportional to N. Without neuronal noise, the memory capacity depends on the product MD, and increasing either the sequence length or the alphabet size has the same effect on memory capacity.

We evaluated equation 2.55 with different parameters, and numerically optimized the memory capacity. We find linear extensive capacity (see Figure 10A1). Interestingly, unlike in the symbolic case, there is no catastrophic forgetting; the retrieved information content saturates to a maximum value as the sequence length M increases to infinity (see Figure 10B1). This means that in the limit of infinite sequence length, the information added by new data is perfectly cancelled by the information lost due to crosstalk (see Figure 10C1). The memory capacity is maximized for any large (finite) M or D compared to N, reaching $I_{total}/N = 1/(2 \log 2)$. This capacity bound can be easily derived analytically, since it is achieved for $r \to 0$:

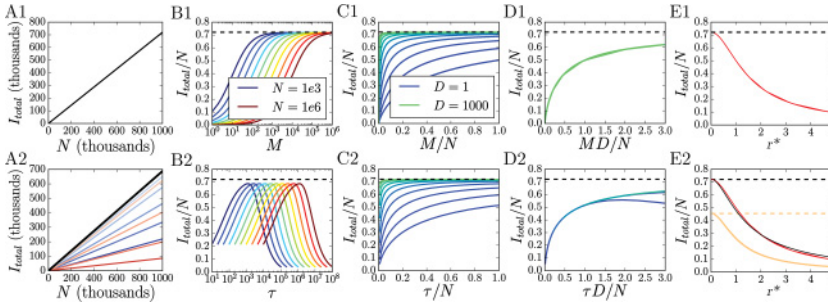

Figure 10: Memory capacity of gaussian inputs. (A1) Numeric evaluation shows extensive memory for finite gaussian input sequences. (B1) As M grows, the memory capacity saturates to the maximum. Note that each curve for different N and fixed $D = 10$ is similar. (C1) The family of curves in panel B1 reduces to a single curve with M/N for each D. As D grows, the memory capacity grows to the bound for any M. (D1) The family of curves in C1 reduces to a single curve with MD/N, which is directly related to $1/r$. Large MD (small r) reaches the memory capacity limit. (E1) The same function in D1 with r as the x-axis. (A2) Numeric evaluation shows extensive capacity whenever τ/N is held constant (colored lines), but a particular ratio results in the maximum (black line). (B2) Similar curves for different N and fixed $D = 10$ show an ideal τ that maximizes memory capacity. (C2) The curves in B2 reduce to a single curve for each D with the ratio τ/N. As D grows, the capacity is maximized. (D2) As τ and D grow large, the capacity saturates to the maximum. (E2) The information per neuron retained at high-fidelity I^* (copper) and the total mutual information per neuron (black) declines as the desired minimum r^* increases. The total mutual information of the memory buffer behaves similar to the total information in the linear case (compare black line to red line; red line same as in panel D1).

$$\frac{I_{total}}{N} = \frac{\log(r+1)}{2r\log 2}\left(\frac{1}{1+\sigma_\eta^2/(DV_\Phi)}\right)$$
$$= {}_{r\to 0}\frac{1}{2\log 2(1+\sigma_\eta^2/(DV_\Phi))} \approx \frac{0.72}{1+\sigma_\eta^2/(DV_\Phi)}. \tag{2.56}$$

However, the regime of optimal memory capacity with $r \to 0$ is not interesting for applications. The critical question is, what fraction of this capacity is available for a fixed desired level r^* of SNR? The answer is depicted in Figure 10D1 as a single curve. Because the memory capacity in the absence of neuronal noise depends only on the SNR, the curve describes all settings of the parameters N, M, and D. The capacity starts at the limit value, equation 2.56, and then decreases as r^* increases (see Figure 10E1). For instance, if the desired SNR is $r^* = 1$, one needs exactly as many neurons (N) as there are coefficients contained in the data sequence (MD) and achieves a memory capacity of 0.5 bits per neuron.

2.5.2 Capacity for Memory Buffer with Analog Input. Here, we analyze the memory buffer with linear neurons and contracting weights $\lambda < 1$ and an infinite sequence of analog input vectors, $M \to \infty$. To compensate for the signal decay, the linear readout contains a factor λ^{-K}:

$$\hat{\mathbf{a}}(M-K) = \mathbf{V}(K)^\top \mathbf{x}(M) = c^{-1}\lambda^{-K}\mathbf{\Phi}^\top \mathbf{W}^{-K}\mathbf{x}(M).$$

The readout then produces the original input, corrupted by gaussian noise, $\hat{a}_{d'} = a_{d'} + n_{d'}$, with the SNR:

$$r(K) = \lambda^{2K}\frac{N(1-\lambda^2)}{D(1-\lambda^{2M})}\left(\frac{1}{1+\sigma_\eta^2/(DV_{\mathbf{\Phi}})}\right). \tag{2.57}$$

The similarity between memory buffers with different mechanisms of forgetting still holds for the case of analog input. For buffers with nonlinear neurons, one can use the analysis of the forgetting time constants in section 2.4.4 and use equation 2.50 to determine the corresponding value of λ. These values can be used in equation 2.57 to compute the SNR of the readout.

If the input is independent gaussian random variables, the total information is

$$I_{total} = \frac{D}{2}\sum_{K}^{M}\log_2\left(r(K)+1\right). \tag{2.58}$$

Inserting equation 2.57 into equation 2.58, we obtain

$$I_{total} = \frac{D}{2}\log_2\left((-b_M q; q)_M\right), \tag{2.59}$$

where $q = \lambda^2$, $b_M := \frac{N(1-q)}{D(1+\sigma_\eta^2/(DV_{\mathbf{\Phi}}))(1-q^M)}$ and $(a; q)_M$ is the q-Pochhammer symbol (see section 4.3.1). The advantage of formulation 2.59 is that it is well defined and can be properly used for $M \to \infty$.

If one numerically optimizes the memory capacity using equation 2.58 for $MD \gg N$, as in the case of reset memories, one finds extensive capacity (see Figure 10A2). Extensive capacity is retained for any constant ratio τ/N, but there is a particular ratio that is the maximum (see Figure 10B2). A single curve for each value of D describes the memory capacity as a function of the ratio τ/N (see Figure 10C2). As D grows large and τ is optimized, the capacity saturates at the same asymptotic value as equation 2.56. By rescaling the x-axis with D, curves for different D become very similar (see Figure 10D2) but not identical; if τ is too large for a fixed D, then the information starts to decrease. These curves collapsed to the exact same curve in the reset memory (see Figure 10D1) because the effects of M and D are

fully interchangeable. However, the forgetting time constant introduces a distinction between the M and D parameters. The asymptotic memory capacity for the memory buffer, equation 2.59, yields the same numeric value as for the reset memory, when τ is optimized:

$$\frac{I_{total}}{N} = \frac{1}{2\log(2)(1+\sigma_\eta^2/(DV_\Phi))} \approx \frac{0.72}{1+\sigma_\eta^2/(DV_\Phi)}. \tag{2.60}$$

This bound is assumed when both τ, D become large enough (see section 4.3.1).

Again, the maximum memory capacity, equation 2.60, is not relevant for applications because the recall SNR for most memories is extremely low. To determine the usable capacity, we estimate $I^*(r^*) = \sum_{\{K:r(K)\geq r^*\}} I_{item}(K)$, the maximum information about past inputs that can be recalled with a given SNR r^* or better. The optimum is reached when as many inputs as possible can be read out with SNR greater than r^*. From the condition $r(K^*) = r^*$ and equation 2.57, one finds the optimal time constant (see section 4.3.2):

$$\frac{\tau_{opt}}{N} = \frac{2}{eDr^*(1+\sigma_\eta^2/(DV_\Phi))}. \tag{2.61}$$

The usable memory capacity for a given SNR threshold r^* is plotted as the copper line in Figure 10E2. Interestingly, the intercept of this curve for $r^* \to 0$ is lower than equation 2.60; the numeric capacity value can be computed (see section 4.3.2) as

$$\frac{I^*(r^* \to 0)}{N} = \frac{1-e^{-1}}{2\log(2)(1+\sigma_\eta^2/(DV_\Phi))} \approx \frac{0.46}{1+\sigma_\eta^2/(DV_\Phi)}. \tag{2.62}$$

The difference between the total capacity (see equation 2.60) and this result is the unusable fraction of information with $r(K) < r^*$.

If one counts usable and unusable information toward I_{total} in a network optimized for I^*, another interesting phenomenon occurs: I_{total} for a particular optimized r^* of the buffer memory is very similar to the capacity of memory with reset (see Figure 10E2, black line; red line for comparison is the same as in panel E1). In both cases, the information capacity drops very similarly as the required fidelity r^* is increased. Note that the meaning of r^* is different in both cases: with reset, it denotes the SNR for all memories, and for the buffer, it is the lowest accepted SNR. Although the total capacity is so similar for reset memory and buffer, the usable information is different. With a reset, all information is retrieved with exactly fidelity $r(K) = r^*$. For the memory buffer, the fraction depicted by the copper curve has fidelity $r(K) \geq r^*$, while inputs further back in history (beyond the critical value $r(K^*) = r^*$) still take up a significant fraction of memory but do

not count toward I^*. The buffer has lower capacity because its exponential decay of the input is only an imperfect substitute of a reset, leaving behind a sediment of unusable information.

2.6 Readout with the Minimum Mean Square Estimator. Thus far, we have analyzed readout mechanisms in equation 2.2 that were proposed in the VSA literature. In contrast, in the area of reservoir computing, a different readout has been studied, with a readout matrix determined by minimizing the mean square error between stored and recalled data. The readout is $\hat{\mathbf{a}}(M-K) = \mathbf{V}(K)^\top \mathbf{x}(M)$, with $\mathbf{V}(K)$ the minimum mean square estimator (MMSE; i.e., linear regression) or Wiener filter, given by $\mathbf{V}(K) = \mathbf{C}^{-1}\mathbf{A}(K)$. Here, $\mathbf{A}(K) := \left\langle \mathbf{a}(M-K)\mathbf{x}(M)^\top \right\rangle_R \in \mathbb{R}^{N\times D}$ is the empirical covariance between input and memory state over R training examples, and $\mathbf{C} := \left\langle \mathbf{x}(M)\mathbf{x}(M)^\top \right\rangle_R \in \mathbb{R}^{N\times N}$ is the covariance matrix of the memory state. In this section, we investigate this read-out method and compare its performance with traditional VSA read-out methods described in previous sections.

The MMSE readout matrix can be determined by solving a regression problem for an ensemble of training input sequences. It involves generating R synthetic input sequences of length M and encoding them in a neural network with one particular choice of input and recurrent matrix. This yields R copies of state vectors, each encoding one of the synthetic input sequences. The readout matrix is now determined by solving the regression problem between state vectors and synthetic input sequences. The particular choice of input and recurrent matrices does not significantly affect the following results as long as the VSA indexing assumptions 2.4 to 2.7 are satisfied. We show results for the input matrix being a gaussian random matrix and the recurrent weights being a permutation matrix.

2.6.1 Reset Memory with MMSE Readout. Symbolic inputs. The MMSE readout does indeed significantly increase the capacity of reset memories in the regime where $MD \lesssim N$ (see Figures 11A and 11B). However, as D grows larger, the performance of the MMSE readout falls back to the performance of VSA models. The comparison of the performances for symbolic input sequences is shown in Figures 11A and 11B.

To find out whether training both the input and decoding matrix, $\mathbf{\Phi}$ and $\mathbf{V}(K)$, has any advantages, we investigated the optimization of these matrices with backpropagation through time. A network of $N = 500$ linear neurons with fixed orthogonal recurrent matrix is fed a sequence of random input symbols and trained to recall the Kth item in the sequence history. The cross-entropy between the recalled distribution and the one-hot input distribution is the error function. During training, the winner-take-all function in equation 2.2 is replaced with softmax. The network is evaluated each time step as more and more input symbols are encoded. After 500 inputs are presented, the network is reset to 0 (see section 4.4.1 for further details).

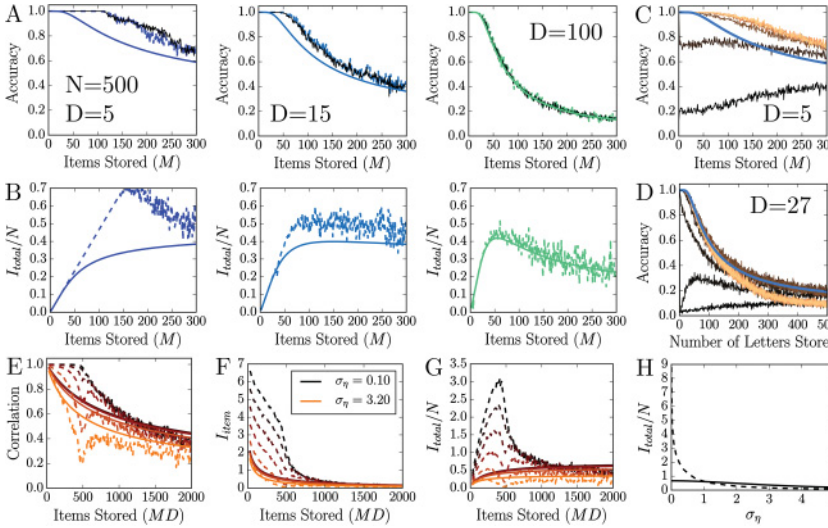

Figure 11: MMSE readout of reset memory networks. (A, B) The accuracy (A) and capacity (B) of VSA prescribed readout (solid lines) is compared to MMSE readout (dashed lines) for different D values. The empirical s measured from the MMSE readout is plugged into p_{corr}, equation 2.12, and matches the measured accuracy (black lines). (C, D) Accuracy of readout when both encoding and readout matrices are trained (black: early training iterations; copper: late training iterations; blue: VSA theory. After 500 inputs, the network is reset). Note that converged training in panel C (copper line) matches the dashed line in panel A. (E–H). MMSE training for analog inputs with neuronal noise utilizes the regime where $MD \lesssim N$. ($N = 500$ in all panels).

The readout accuracy with backpropagation learning improves successively with training (black to copper in Figures 11C and 11D), reaching the performance level of the MMSE readout (the copper line in Figure 11C matches the dashed MMSE line in Figure 11A). For larger D, the performance of backpropagation learning, MMSE readout, and VSA readout are equal (see Figure 11D). This convergence in performance shows that the simultaneous optimization of input and readout matrices (with backpropagation learning) yields no improvement over optimizing just the readout matrix for fixed input and recurrent matrix (with MMSE optimization).

Analog inputs. Compared to the case with discrete input, for analog input, the improvement achievable with MMSE has a similar pattern, but the magnitude of improvement is even more dramatic. When $MD \lesssim N$, MMSE can greatly diminish the crosstalk noise and increase memory capacity (see Figures 11E to 11G). This improvement is because the MMSE readout allows the network to fully utilize all N orthogonal degrees of freedom and store

nearly N analog numbers at high fidelity. The retrievable information per symbol is increased to many bits (see Figure 11F), and the memory capacity is limited only by neuronal noise (see Figure 11H).

Direct calculation of the readout matrix. The MMSE readout matrix for reset memory can also be computed without empirically solving the regression problem. The case where the number of training input sequences is sent to infinity $R \to \infty$ (see section 4.4.4) can be treated analytically. If $\mathbf{a}(m)$ has zero mean, the expected covariance matrix of the network states can be computed from $\mathbf{W}$ and $\mathbf{\Phi}$ as

$$\tilde{\mathbf{C}} = \langle \mathbf{x}(M)\mathbf{x}(M)^\top \rangle_\infty = M\sigma_\eta^2 \mathbf{I} + \sum_{k=1}^{M} \mathbf{W}^k \mathbf{\Phi}\mathbf{\Phi}^\top \mathbf{W}^{-k}. \tag{2.63}$$

An element of the expected covariance matrix is given by

$$\tilde{C}_{ij} = (\delta_{i=j})MDV_{\mathbf{\Phi}}(1 + \sigma_\eta^2/(DV_{\mathbf{\Phi}})) + (1 - \delta_{i=j}) \sum_{k=1}^{M} (\mathbf{W}^k)_i \mathbf{\Phi}\mathbf{\Phi}^\top (\mathbf{W}^{-k})_j. \tag{2.64}$$

Further, the covariance between inputs and memory states converges to the VSA readout: $\tilde{\mathbf{A}}(K) = \langle \mathbf{a}(M-K)\mathbf{x}(M)^\top \rangle_\infty = \mathbf{W}^K \mathbf{\Phi}$. Thus, the MMSE readout matrix is given by

$$\tilde{\mathbf{V}}(K) = \tilde{\mathbf{C}}^{-1} \mathbf{W}^K \mathbf{\Phi}. \tag{2.65}$$

Note from equation 2.65 that the MMSE readout involves VSA readout with an additional multiplication by the inverse covariance of the memory state. Thus, dimensions in the state space that are only weakly driven through the input and recurrent matrices are expanded to become fully useful in the decoding. The neuronal noise serves as ridge regularization of the regression, adding power in all dimensions. If the noise power σ_η^2 is positive, equation 2.65, is always well defined. However, without neuronal noise, the memory covariance matrix can become rank deficient and equation 2.65 undefined.

For symbolic input, this directly calculated linear filter matches the performance of the linear filter determined by linear regression from synthetic data. However for analog input, the filter determined by linear regression somewhat outperforms the directly calculated filter (see Figure 19).

2.6.2 Memory Buffer with MMSE Readout. Symbolic inputs. Our findings for the memory buffer are similar to our results for reset memories for both types of inputs. For discrete inputs and when D is small, the MMSE readout improves the performance of memory buffers in the regime where $MD \lesssim N$, significantly increasing retrieval accuracy and

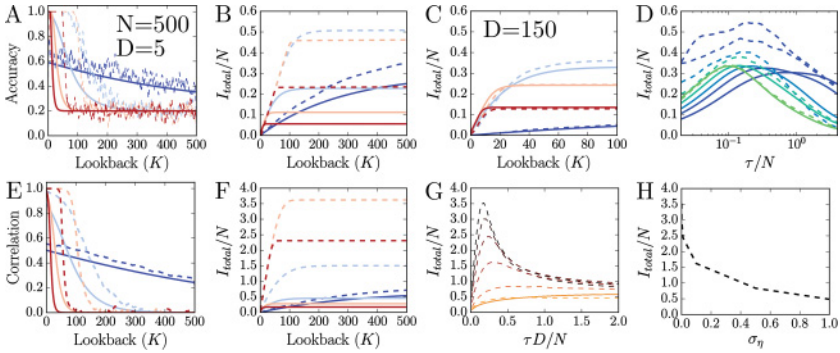

Figure 12: MMSE readout in memory buffer networks. (A–C) The accuracy (A) and capacity (B, C) of MMSE readout (dashed lines) are compared to VSA readout (solid lines) for memory buffers with discrete inputs. Four networks with $\tau = 0.8, 0.95, 0.99, 0.999$. (D) The capacity is computed as a function of the time constant for $D = 5, \ldots, 150$ (blue to green). For small D, the time constant can be used to enhance the readout accuracy and memory capacity, with optimal time constant between $(0.1 - 0.5)N$. (E, F) The correlation (E) and capacity (F) for analog inputs show that MMSE training can utilize nearly all N degrees of freedom with the right time constant ($N = 500$, $D = 5$). (G, H) The capacity is computed for different amounts of noise.

capacity (see Figures 12A to 12D). Optimizing the time constant is still required to maximize memory capacity (see Figure 12D), which occurs when τ is $(0.1 - 0.5)N$.

Analog inputs. For analog inputs, the time constant can be optimized to take advantage of the N orthogonal degrees of freedom and mitigate crosstalk noise (see Figures 12E to 12G). High-fidelity retrieval can be maintained for $K_{\max}$ items, with $K_{\max} D \lesssim N$. Many bits per item can be recovered this way with τ between $(0.1 - 0.5)N/D$, and the memory capacity is again limited only by neuronal noise (see Figure 12H).

Direct calculation of the readout matrix. The expected MMSE readout filter for the memory buffer is

$$\tilde{\mathbf{V}}(K) = \tilde{\mathbf{C}}^{-1} \lambda^K \mathbf{W}^K \mathbf{\Phi}. \tag{2.66}$$

For the case $D = 1$, this is the readout proposed in White et al. (2004).

The covariance matrix can be estimated for a given $\mathbf{W}$ and codebook $\mathbf{\Phi}$:

$$\tilde{\mathbf{C}} = \frac{\sigma_\eta^2}{1 - \lambda^2} \mathbf{I} + \sum_{k=1}^{\infty} \lambda^{2k} \mathbf{W}^k \mathbf{\Phi} \mathbf{\Phi}^\top \mathbf{W}^{-k}. \tag{2.67}$$

This can be simplified to a finite sum if the cycle time of the unitary matrix is known. Different unitary matrices can have different cycle times (random permutations have an expected cycle time of $0.62N$; Golomb, 1964), but we can pick a permutation matrix that has a maximum cycle time of N, s.t. $W^k = W^{k+N}$. This gives

$$\tilde{\mathbf{C}} = \frac{\sigma_\eta^2}{1-\lambda^2}\mathbf{I} + \sum_{k=1}^{N} \frac{\lambda^{2k}}{1-\lambda^{2N}} \mathbf{W}^k \mathbf{\Phi}\mathbf{\Phi}^\top \mathbf{W}^{-k}. \tag{2.68}$$

Here, the performance of MMSE with directly calculated readout matrix matches the performance with the readout matrix obtained by regression over synthetic data.

2.7 Examples of Storing and Retrieving Analog Inputs. To illustrate the analog theory and also show that the input distribution can be complex and nongaussian, we encoded a random ($12 \times 12 \times 3$) image patch into the vector $\mathbf{a}(m)$ each time step. Several networks of different sizes were used as memory buffers for the same image sequence, with the time constants held proportional to N. We empirically evaluated their performance. The retrieved images are shown as a function of N and look-back K for networks with VSA readout (see Figure 13A) and networks with full MMSE readout (see Figure 13B). The empirically measured SNR of the MMSE readout is greatly increased compared to the VSA readout (see Figure 13C).

One advantage of VSAs is that they can be used to form arbitrary composite representations and index data structures other than sequences. As a final example, we follow the procedure of Joshi, Halseth, and Kanerva (2016) that uses the HDC code and algebra to encode the statistics of letter trigrams. They show how this technique can be used to create an effective, simple, and low-cost classifier for identifying languages.

We performed the task of storing probabilities of letter n-grams by using the HDC algebra to create key vectors for each set of individual letters, bigrams, or trigrams from the base set of $D = 27$ tokens (26 letters and space). The text of *Alice in Wonderland* served as the input to accumulate statistics about n-grams. Important to note here is that a complex combination of multiplication and permutation is needed to create the composite n-gram representation from the base set of random vectors. For instance, the trigram *abc* is encoded as $\mathbf{x}_{abc} = \rho^2(\mathbf{\Phi}_a) \times \rho(\mathbf{\Phi}_b) \times \mathbf{\Phi}_c$, where ρ is the permutation operation. This encoding distinguishes the *abc* trigram from the trigram *bac* or any other n-gram that may share letters or differ only in letter order.

Unlike the sequence indexing presented before, this composite representation cannot be created by a fixed matrix multiply. However, the composite binding and permutation operations still follow the statistics of assumptions 2.4 to 2.7 and thus are still characterized by our theory. In fact,

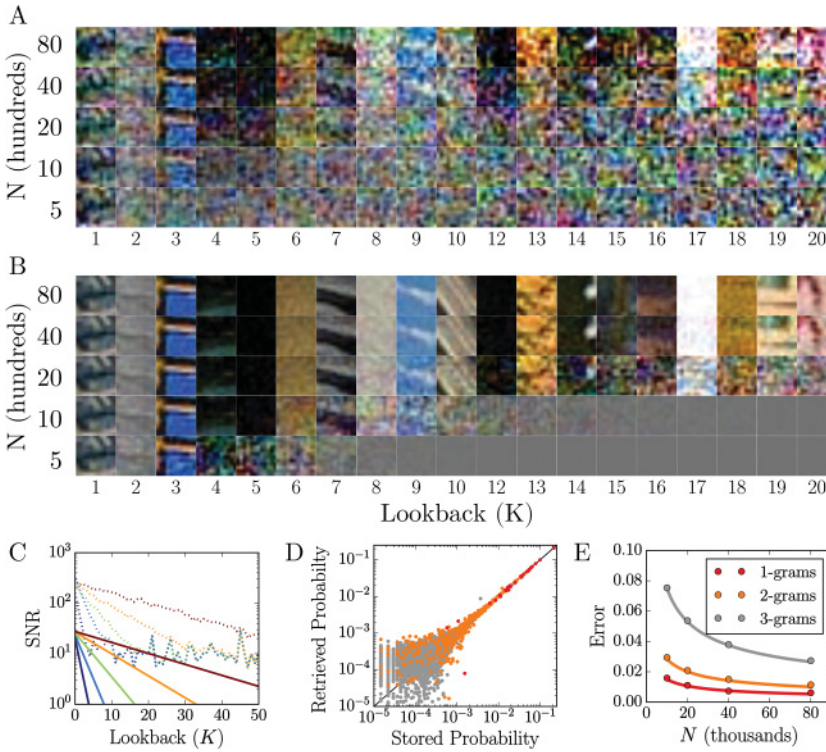

Figure 13: Analog coefficient storage and retrieval. (A, B) A long sequence of image patches was stored in networks with different N values, with τ proportional to N. The recent images were retrieved with VSA readout (A) and MMSE trained readout (B) and reconstructed for different look-back values, K. (C) The measured SNR of MMSE readout (dashed lines) is greatly increased compared to the SNR of VSA readout. (D) Indexing language n-grams with compositional binding operations has the same cross-talk properties as sequence indexing. (E) The measured readout error of language statistics (dots) matches the theory (lines).

all composite representations of discrete VSA base symbols ultimately follow these assumptions and are effectively a set of superposed random vectors. In terms of the neural network, the storage of n-gram statistics can be interpreted as the network accumulating the encoded n-gram vectors generated by external computations. The recurrent weight matrix is the identity, and thus the network counts up each n-gram vector that is given as input. The n-gram counts are indexed by the key vector made by the composite VSA operations and integrated into the memory state. We see that the

empirical performance of such an encoding behaves in the same qualitative manner and matches the theory (see Figures 13D and 13E).

3 Discussion

The ability of recurrent neural networks to memorize past inputs in the current activity state is the basis of working memory in artificial neural systems for signal processing (Jaeger & Haas, 2004), cognitive reasoning (Eliasmith et al., 2012), and machine learning (Graves et al., 2014). We try to dissect and understand what is often an intriguing but rather opaque property of recurrent neural networks. This is possible through the lens of vector-symbolic architectures (VSAs), a class of connectionist models proposed for structured computations with high-dimensional vector representations. Our work demonstrates that VSA sequence indexing can be mapped to recurrent neural networks with specific weight properties: the input weights random and the recurrent weights an orthogonal matrix. The computation of an iteration in such a network can now be concisely interpreted as generating a unique time stamp key and forming a key-value pair with the new input. These key-value bindings can be used to selectively retrieve data memorized by the network state. We were able to derive a theory describing the readout accuracy and information capacity achievable in such networks. Our results update and unify previous work on vector-based symbolic reasoning and contribute new results to the area of reservoir computing.

The theory includes two different forms of working memory. A reset memory operates like a tape recorder with start and stop buttons. The network state is initialized to zeros before the input of interest arrives and the sequence of inputs is finite. With the ability to reset, networks without forgetting reach optimal capacity (Lim & Goldman, 2011). Existing VSA models map onto networks that operate as reset memories. In contrast, a memory buffer can track recent inputs in an infinite stream without requiring external reset. We investigated palimpsest networks as memory buffers, which attenuate older information gradually by various mechanisms, such as contracting recurrent weights or saturating neural nonlinearities. We showed that there is one essential property of palimpsest memories: the forgetting time constant. We demonstrate how this constant is computed for different forgetting mechanisms and that the model performances are very similar if the forgetting time constant is the same. Further, we demonstrated that the time constant can be optimized for obtaining high and extensive information capacity, even in the presence of neuronal noise.

The theory analyzes memory networks for two different types of input data, symbolic or analog. Symbolic inputs, encoded by one-hot and binary input vectors, correspond to neural network models of vector-symbolic architectures (VSA) (Plate, 1994, 2003; Gallant & Okaywe, 2013). In these models, a naive linear readout is followed by a nonlinear winner-take-all

operation for additional error correction. The naive linear readout consists in projecting the memory state in the direction of the key vector associated with the wanted input. In addition to storing sequences, VSAs also allow the storage of other composite structures, such as unconnected sets of key-value pairs, trees, stacks, and *n*-grams, and our theory also extends to those (see Figures 13D and 13E).

The case of analog inputs has been considered before in reservoir computing (Jaeger, 2002; White et al., 2004; Ganguli et al., 2008). Some of these models have weight properties that fall outside the conditions of our theory. However, the regime described by the theory is particularly interesting. Previous work has shown that unitary recurrent weights optimize capacity (White et al., 2004). The readout in memories for analog data is typically linear. Going beyond the naive readout used in VSAs, these models use the Wiener filter providing the MMSE regression solution (White et al., 2004). The MMSE readout greatly improves performance when encoding independent analog values, as nearly the full N orthogonal degrees of freedom can be used with minimal cross-talk noise. In other regimes, however, the naive readout in VSAs has advantages, as it yields similar performance and is much easier to compute.[2]

3.1 Working Memory in Vector-Based Models of Cognitive Reasoning

3.1.1 Analysis of Existing VSA Models. We demonstrated how various VSA models from the literature can be mapped to equivalent reset memories with linear neurons and unitary recurrent matrix. This approach not only suggests concrete neural implementations of VSA models, but also allowed us to trace and highlight common properties of existing models and develop a common theory for analyzing them.

VSA models use the fact that random high-dimensional vectors are nearly orthogonal and that the composition operations (such as forming a sequence) preserve independence and generate pseudo-random vectors. The prerequisites of VSAs are formalized by conditions 2.4 to 2.7.

The previous analyses of specific VSA models (Plate, 1994, 2003; Gallant & Okaywe, 2013) were limited to the high-fidelity regime. Our theory yields more accurate estimates in this regime, revealing that the working memory performance is actually superior to what was previously thought. Specifically, we derived a capacity bound of 0.39 bits per neuron in the high-fidelity regime.

Our theory is not limited to the high-fidelity regime; it predicts the effects of cross talk and neuronal noise in all parameter regimes. In particular,

[2] Computing the inverse of the $R^{N\times N}$ matrix $\mathbf{C}$ scales with $O(N^3)$ and becomes unfeasible with N larger than a few thousand. For instance, the computation of $\mathbf{C}^{-1}$ took over 14 hours for the $N = 8000$ network shown in Figure 13B.

the theory revealed that different VSA frameworks from the literature have universal properties. For larger networks, recall accuracy is independent of the moments of a particular distribution of random codes, and therefore the sensitivity for recall is universally $s = \sqrt{N/M}$ (see equation 2.35). This finding explains that achieving large sensitivity for high-fidelity recall requires the number of memory items to be smaller than the number of neurons. These results can be used to design optimized VSA representations for particular hardware (Rahimi et al., 2017).

3.1.2 New VSA Models with Optimal Readout. VSA models use a linear readout that is suboptimal but fast to compute. Here, we asked how much optimal linear readout with the MMSE or Wiener filter can improve the performance of VSA models. VSA models are used to index both symbolic and analog input variables, and the input distribution has important consequences on the coding and capacity.

Symbols are represented by binary one-hot vectors and have an entropy of $M \log_2 D$. The MMSE readout can reduce crosstalk when the input can be represented with N orthogonal dimensions. However, the one-hot vectors still require MD dimensions to encode the input sequence, and the MMSE loses its advantage if D is large (see Figures 11A and 11B). Discrete input sequences that have $MD > N$ can still be stored with the randomized code vectors as the basis because the entropy is only $M \log_2 D < N$, but MMSE training does not much improve readout accuracy in this regime.

The analog input vectors we considered have independent entries. In this case, what matters for retrieval accuracy is the number MD of real numbers to store. Thus, the dimensions of the input vector, D, and the sequence length, M, contribute in the same way to the memory load. If and only if $MD \lesssim N$, the MMSE readout can remove crosstalk noise inherent in standard VSA models and can greatly increase the capacity to many bits per neuron. However, if $MD > N$, the performance of MMSE readout drops back to the performance of standard VSA readout.

3.1.3 New VSA Memory Buffers. Plate (1994) describes trajectory association, the mechanisms for indexing an input sequence with an RNN. Our analysis quantifies how recency effects, caused by contracting weights or nonlinear activations, influence recall accuracy. This analysis enabled us to construct memory buffers that are optimized to perform trajectory association in continuous data streams. For instance, consider an optimized digital implementation of a discrete memory buffer. We can use the HDC code framework, $p_{\Phi}(x) = \mathcal{B}_{0.5}$, $x \in \{-1, +1\}$, with clipping nonlinearity, and create a memory buffer where neurons only have integer activation states bounded by κ. Thus, the memory state can be represented by $N \log_2(\kappa)$ bits (see Figure 9G). The recurrent matrix is a simple permutation. This is an efficient way to utilize the coding scheme for a digital device that can continually encode a spatiotemporal input into an indexed, distributed memory

state. Readout neurons can be trained to recognize temporal input patterns (Kleyko, Frady, & Osipov, 2017). The complex vector representations used in FHRR could be useful for emerging computational hardware, such as optical computing or spiking neural hardware.

3.2 Contributions to the Field of Reservoir Computing

3.2.1 Memory Buffers Have Extensive Capacity with Optimized Forgetting Time Constant. Our theory captures working memory in neural networks that have contracting weights or saturating nonlinear neurons, both of which produce a recency effect. We derive for these diverse mechanisms the one single feature that is critical to the memory performance: the network's forgetting time constant. If the forgetting time constant is the same for networks with different forgetting mechanisms, the memory performance becomes very similar (see Figure 9). Putting the forgetting time constant in the center focus enables a unified view on a large body of literature investigating the scaling and capacity of recurrent neural networks (Jaeger, 2002; White et al., 2004; Ganguli et al., 2008; Hermans & Schrauwen, 2010; Wallace, Maei, & Latham, 2013).

Importantly, we found that memory buffers can possess extensive capacity in the presence of accumulating noise (see Figure 9E; enhanced by MMSE readout, Figure 12). To preserve extensive capacity with noise, the time constant (τ) has to be chosen appropriately for the given noise power (σ_η^2) and number of neurons (N). As N grows large, the time constant of the network must also increase proportionally (see Figure 9A). With this choice, the noise accumulation scales but does not destroy the linear relationship between N and the network capacity (see Figure 9D). It was first noted in White et al. (2004) that extensive capacity could be attained in normal networks (e.g., see the networks in section 2.4.1) by correctly setting the decay parameter in relation to N.

Since the dynamic range of the neurons determines the time constant of the recency effect, it must be optimized with N to achieve extensive capacity. Conversely, if the parameters related to the time constant are kept fixed as N grows large, the memory capacity does not scale proportionally to N. Thus, our result of extensive capacity is not in contradiction to Ganguli et al. (2008) where a subextensive capacity of $O(\sqrt{N})$ is derived for a fixed dynamic range (e.g., κ) in networks with nonnormal recurrent matrix. Our analysis confirms the subextensive capacity when the time constant is held fixed as N grows. With normal matrices, we see $O(\log N)$ scaling with fixed dynamic range. This logarithmic scaling matches results reported in Wallace et al. (2013).

We show that extensive capacity can be achieved by increasing the dynamic range of the neurons as N grows. Extensive capacity is achieved in this case, but only if the neuronal noise is fixed and not growing

proportionally to the dynamic range. If the noise assumptions are such that neuronal noise and dynamic range are proportional (Ganguli et al., 2008), then extensive capacity is not possible.

The time constant can also be increased by decreasing the variance of the codebook, V_Φ and keeping the dynamic range fixed (see equation 2.51). In the zero-noise case, optimizing the codebook variance can achieve extensive capacity with a fixed dynamic range. However, when fixed noise is present, reducing the variance also increases the impact of noise (see equation 2.37), which prohibits extensive capacity.

The analysis of the recency effect has implications for other models that use principles of reservoir computing to learn and generate complex output sequences (Sussillo & Abbott, 2009). The buffer time constant τ and its relationship to network size could be used for optimizing and understanding network limits and operational time-scales.

Previous studies have suggested that the RNN (see equation 2.1) can memorize sparse input sequences in which the sequence length is greater than the number of neurons, $M > N$, if sparse inference is used in the readout (Ganguli & Sompolinsky, 2010). Using the theory of compressed sensing, it has been shown for a fixed accuracy requirement that the number of required neurons scales as $N \propto Mp_s \log^\ell(M)$ (Charles et al., 2014, 2017). Our result for sparse input sequences, $N = s^2 Mp_s$, lacks the factor logarithmic with sequence length. One reason for this discrepancy is that our result requires conditions 2.4 to 2.7 to hold. For large enough M, condition 2.7 will inevitably fail to hold because unitary matrix powers eventually loop, and thus the sequence indexing keys for long sparse input sequences are not independent. Thus, compressed sensing theory might account for this gradual failing of producing independent time stamp keys, which reduces performance as the sequence length grows. Another reason for disagreement is that the compressed sensing readout requires sparse inference of the complete sequence history; it does not permit access to individual sequence entries. In contrast, the readout procedures presented in this article permit individual entries in the sequence to be retrieved separately.

3.2.2 Optimal Memory Capacity of Neural Reservoirs. White et al. (2004) proposed distributed shift register (DSR) models that store a one-dimensional temporal input sequence spatially in a delay line using a connectivity that avoids any signal mixing. This model shows how the full entropy of an RNN can be utilized and that capacity scales proportionally to N. The DSR performance is considered an upper bound of memory performance in reservoir computing. The DSR uses a constructed code for indexing and storing sequence history along each of the N neural dimensions, without any crosstalk. However, because of this construction, the network can break down catastrophically from small perturbations such as the removal of a neuron.

Table 1: Idealized Bounds on Memory Capacity.

	VSA		MMSE	
Capacity $\left(\frac{bits}{neuron}\right)$	Reset	Buffer	Reset	Buffer
Symbolic	≈ 0.5	≈ 0.3	1	1
Analog	0.72	0.46	∞	∞

White et al. (2004) also analyzed normal networks that can distribute the input sequence across the neurons and create a more robust memory. They used an annealed approximation for describing the readout performance for one-dimensional input sequences, which shares many characteristics with assumptions 2.4 to 2.7. Their approximation includes the MMSE readout and corresponds to analog memory buffer networks considered in section 2.6.2. This theory suggests much lower performance of normal networks compared to the DSR. We directly compare the White et al. (2004) theory to memory buffers with naive VSA readout and MMSE readout in section 4.4.2.

We show how VSAs can be used to perform indexing with random code vectors for inputs of arbitrary dimensions. Our theory precisely characterizes the nature of cross-talk noise, and we show how MMSE readout can remove much of this cross-talk noise in the regime where $MD \lesssim N$. We see that VSA indexing with MMSE readout outperforms the White et al. (2004) theory for normal networks and can reach the memory performance of the DSR. Compared to the constructed codes like the DSR, there is a small reduction of the capacity due to duplication of code vectors of random codes (see section 4.4.3), but this reduction becomes negligible for large network sizes. Thus, VSA encoding with MMSE readout can be distributed and robust while still retaining the full capacity of the DSR.

3.3 Survey of Memory Capacity across Models. Our results show that distributed representations indexed by VSA methods can be optimized for extensive capacity. We report the idealized, zero-noise memory capacity bounds found for different input streams and readouts in Table 1. The table reveals that the bounds are finite except for the case of MMSE readout with analog variables. The table summarizes the performance bounds of different models, although these bounds may not be achievable under realistic conditions.

4 Methods

4.1 Vector Symbolic Architectures. The different vector symbolic architectures described here share many fundamental properties, but they

Table 2: Summary of VSA Computing Frameworks.

VSA	Symbol Set ($p_{\mathbf{\Phi}}(x)$)	Binding	Permutation	Trajectory Association
HDC	$\mathcal{B}_{0.5} : x \in \{-1, +1\}$	$\times$	ρ	$\sum \rho^m(\mathbf{\Phi}_{d'})$
HRR	$\mathcal{N}(0, 1/N)$	$\circledast$	None	$\sum \mathbf{w}^m \circledast \mathbf{\Phi}_{d'}$
FHRR	$\mathcal{C} := e^{i\phi} : \phi \in \mathcal{U}(0, 2\pi)$	$\times$	$\circledast$	$\sum \mathbf{w}^m \times \mathbf{\Phi}_{d'}$ or $\sum \mathbf{w}^{\circledast m} \circledast \mathbf{\Phi}_{d'}$
MBAT	$\mathcal{N}(0, 1/N)$	Matrix	Matrix	$\sum \mathbf{W}^m \mathbf{\Phi}_{d'}$

Note: Each framework has its own set of symbols and operations on them for addition, multiplication, and a measure of similarity.

also have their unique flavors and potential advantages and disadvantages. Each framework utilizes random high-dimensional vectors (hypervectors) as the basis for representing symbols, but these vectors are drawn from different distributions (see Table 2). Further, different mechanisms are used to implement the key operations for vector computing: superposition, binding, permutation, and similarity.

4.1.1 Basics. The *similarity* operation transforms two hypervectors into a scalar that represents similarity or distance. In HDC, HRR, FHRR, and other frameworks, the similarity operation is the dot product of two hypervectors, while the Hamming distance is used in the frameworks that use only binary activations. The distance metric is inversely related to the similarity metric. When vectors are similar, their dot product will be very high or their Hamming distance will be close to 0. When vectors are orthogonal, their dot product is near 0 or their Hamming distance is near 0.5.

When the superposition ($+$) operation is applied to a pair of hypervectors, the result is a new hypervector that is similar to each one of the original pair. Consider HDC, given two hypervectors, $\mathbf{\Phi}_A, \mathbf{\Phi}_B$, which are independently chosen from $p_{\mathbf{\Phi}}(x) = \mathcal{B}_{0.5} : x \in \{-1, +1\}$ and thus have low similarity ($\mathbf{\Phi}_A^\top \mathbf{\Phi}_B = 0 + \textit{noise}$). Then the superposition of these vectors, $\mathbf{x} := \mathbf{\Phi}_A + \mathbf{\Phi}_B$, has high similarity to each of the original hypervectors (e.g. $\mathbf{\Phi}_A^\top \mathbf{x} = N + \textit{noise}$). In the linear VSA frameworks (Kanerva, 2009; Plate, 2003), we do not constrain the superposition operation to restrict the elements of the resulting vector to $\{-1, +1\}$, but we allow any rational value. However, other frameworks (Kanerva, 1996; Rachkovskij & Kussul, 2001) use clipping or majority rule to constrain the activations, typically to binary values.

The binding operation ($\times$) combines two hypervectors into a third hypervector ($\mathbf{x} := \mathbf{\Phi}_A \times \mathbf{\Phi}_B$) that has low similarity to the original pair (e.g., $\mathbf{\Phi}_A^\top \mathbf{x} = 0 + \textit{noise}$) and also maintains its basic statistical properties (i.e., it looks like a vector chosen from $p_{\mathbf{\Phi}}(x)$). In the HDC framework, the hypervectors are their own multiplicative self-inverses (e.g., $\mathbf{\Phi}_A \times \mathbf{\Phi}_A = \mathbf{1}$, where $\mathbf{1}$ is the binding identity), which means they can be dereferenced from the bound pair by the same operation (e.g., $\mathbf{\Phi}_A \times \mathbf{x} = \mathbf{\Phi}_B + \textit{noise}$). In the binary

frameworks, the binding operation is element-wise XOR, while in HRR and other frameworks, binding is implemented by circular convolution ($\circledast$).

In different VSA frameworks, these compositions are implemented by different operations. We note that all the binding operations can be mapped to a matrix multiply and the frameworks can be considered in the same neural network representation. The FHRR framework is the most generic of the VSAs and can utilize both multiply ($\times$) and circular convolution ($\circledast$) as a binding mechanism.

4.1.2 Implementation Details. The experiments are all implemented in Python as Jupyter notebooks using standard packages like numpy.

The experiments done with different VSA models use different implementations for binding, most of which can be captured by a matrix multiplication. However, for efficiency reasons, we implemented the permutation operation ρ and the circular convolution operation $\circledast$ with more efficient algorithms than the matrix multiplication. The permutation operation can be implemented with $O(N)$ complexity, using a circular shifting function (`np.roll`). Efficient circular convolution can be performed by fast Fourier transform, element-wise multiply in the Fourier domain, and inverse fast Fourier transform, with $O(N \log N)$ complexity.

To implement FHRR, we utilized a network of dimension N, where the first $N/2$ elements of the network are the real part and the second $N/2$ elements are the imaginary part. Binding through complex multiplication is implemented as

$$\mathbf{u} \times \mathbf{v} = \begin{bmatrix} \mathbf{u}_{real} \times \mathbf{v}_{real} - \mathbf{u}_{imaginary} \times \mathbf{v}_{imaginary} \\ \mathbf{u}_{real} \times \mathbf{v}_{imaginary} + \mathbf{u}_{imaginary} \times \mathbf{v}_{real} \end{bmatrix}.$$

The circular convolution operation can also be implemented in this framework, but with consideration that the pairs of numbers are permuted together. This can be implemented with a circulant matrix $\mathbf{W}$ with size $(N/2, N/2)$:

$$\mathbf{w} \circledast \mathbf{u} = \begin{bmatrix} \mathbf{W} & \mathbf{0} \\ \mathbf{0} & \mathbf{W} \end{bmatrix} \mathbf{u}.$$

The superposition ($+$) is the same, and similarity ($\cdot$) functions are defined for complex numbers as simply

$$\mathbf{u} \cdot \mathbf{v} = \mathbf{u}_{real} \cdot \mathbf{v}_{real} + \mathbf{u}_{imaginary} \cdot \mathbf{v}_{imaginary},$$

which is the real part of the conjugate dot product, $Re(\mathbf{u}^\top \mathbf{v}^*)$.

Either circular convolution or element-wise multiplication can be used to implement binding in FHRR, and trajectory association can be performed to encode the letter sequence with either operation:

$$\mathbf{x}(M) = \sum \mathbf{w}^{M-m} \times \mathbf{\Phi a}(m) \text{ or}$$

$$\mathbf{x}(M) = \sum \mathbf{w}^{\circledast(M-m)} \circledast \mathbf{\Phi a}(m),$$

where $\mathbf{w}^{\circledast m}$ means circular convolutional exponentiation (e.g., $\mathbf{w}^{\circledast 2} = \mathbf{w} \circledast \mathbf{w}$).

4.2 Accuracy of Retrieval from Superpositions

4.2.1 Comparison of Approximations for the High-Fidelity Regime. We compared each step of the high-fidelity approximation (see section 2.2.2) to the true numerically evaluated integral (see Figure 14) to understand which regimes of the approximations were valid (see Figure 15B).

We compare the CR bound and the Chang et al. (2011) approximation to the numerically evaluated normal cdf Φ and see that the CR lower bound does not get tight until multiple standard deviations into the very high-fidelity regime (see Figure 15A).

In Figures 15D and 15E, we see that while the approximations given are not strictly lower bounds, they are typically below the numerically evaluated accuracy. The Chang approximation can overestimate the performance, however, in the high-fidelity regime when D is large.

4.2.2 Previous Theories of the High-Fidelity Regime. The capacity theory derived here is similar to but slightly different from the analysis of Plate (2003), which builds on work done in Plate (1994). Plate (2003) frames the question: "What is the probability that I can correctly decode all M tokens stored, each of which is taken from the full set of D possibilities without replacement?" This is a slightly different problem, because this particular version of Plate's (2003) anlaysis does not use trajectory association to store copies of the same token in different addresses. Thus, M is always less than D, the M tokens are all unique, and there is a difference in the sampling of the tokens between our analysis frameworks.

Nonetheless, the analysis can be translated to a roughly equivalent framework given that D is relatively large compared to M. Plate (2003) derives the hit $p(h_{d'})$ and reject $p(h_d)$ distributions in the same manner as presented in our analysis, as well as uses a threshold to pose the probability problem:

$$p_{all-corr} = p(h_{d'} > \theta)^M p(h_d < \theta)^{D-M}. \tag{4.1}$$

```
from __future__ import division

import numpy as np

import scipy.special


def ncdf(z):

    return 0.5 * (1 + scipy.special.erf(z/2**0.5))


def p_correct_snr(M, N=10000, D=27, ares=2000):

    p = np.zeros((ares-1, len(M)))

    for iM,Mval in enumerate(M):

        s = (N / Mval)**0.5

        # span the Hit distribution up to 8 standard deviations

        av = np.linspace(s - 8, s + 8, ares)

        # the discretized gaussian of h_d'

        pdf_hdp = ncdf(av[1:])-ncdf(av[:-1])

        # the discretized cumulative gaussian of h_d

        cdf_hd = ncdf(np.mean(np.vstack((av[1:]+s, av[:-1]+s)),

            axis=0))

        p[:, iM] = pdf_hdp * cdf_h ** (D-1)

    return np.sum(p, axis=0) # integrate over av
```

Figure 14: Numeric algorithm for accuracy integral.

This can be interpreted as that the probability of reading all M tokens correctly ($p_{corr-all}$) is the probability that the dot product of the true token is larger than threshold for all M stored tokens ($p(h_{d'} > \theta)^M$) *and* that the dot product is below threshold for all $D - M$ remaining distracter tokens ($p(h_d < \theta)^{D-M}$).

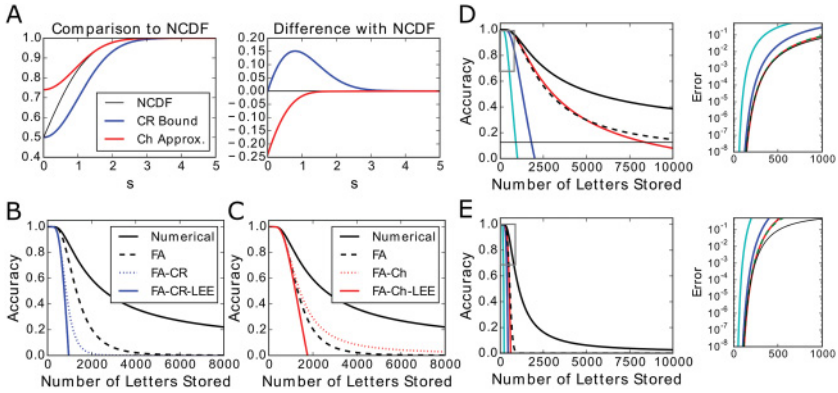

Figure 15: Comparison of different methods to approximate the retrieval accuracy. (A) The Chernoff-Rubin (CR; Chernoff, 1952) lower bound (blue) and the Chang et al. (2011) approximation (red) to compute the normalized cumulative density function (NCDF; black) analytically. The Chang et al. (2011) approximation becomes tight faster in the high-fidelity regime, but is not a lower bound. (B) Differences between the three methods of approximations and the numerically evaluated p_{corr} integral (black line). The factorial approximation (dashed black line) still requires numerical evaluation of the NCDF. Adding the CR lower bound (dashed blue) and the local error expansion, the high-fidelity regime can still be described well, but the low-fidelity regime cannot be captured. (C) Same as panel B, but using the Chang et al. (2011) approximation to the NCDF. (D) Accuracy curve and approximations for $D = 8$. (E) $D = 1024$. Right panels in panels D and E are zoom-ins into the high-fidelity regime (marked by gray box insets in the left panels).

In our framework, the probability of correctly reading out an individual symbol from the M items stored in memory is independent for all M items. This is given by equation 2.12; alter the equation to output the probability of reading all M tokens correctly, simply raise p_{corr} to te Mth power:

$$p_{all-corr} = \left[p_{corr}\right]^M = \left[\int_\theta^\infty \frac{dh}{\sqrt{2\pi}} e^{\frac{-h^2}{2}} \left[\Phi\left(h+s\right)\right]^{D-1}\right]^M. \tag{4.2}$$

In Figure 16, we compare our theory to Plate's by computing $p_{all-corr}$ given various different parameters of N, M, and D. We show that Plate's (2003) framework comparatively underestimates the capacity of hypervectors. There is slight discrepancy in our analysis frameworks because of how the tokens are decoded from memory. In our analysis framework, we take the maximum dot product as the decoded symbol, and there are instances that can be correctly classified that Plate's (2003) probability statement,

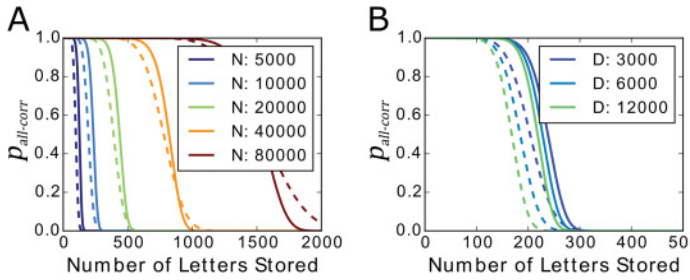

Figure 16: Comparison with the theory in Plate (2003). (A) Plate (2003) derived $p_{all-corr} = p_{corr}^{M}$, plotted in dashed lines for different values of N with D fixed at 4096. The new theory in solid lines. (B) Plate's (2003) theory in dashed lines with different values of D and fixed N. The new theory in solid lines.

equation 4.1, would consider incorrect. For instance, the true symbol and a distracter symbol can have dot products above threshold and the correct symbol can still be decoded as long as the true symbol's dot product is larger than the distracter. However, this scenario would be classified as incorrect by equation 4.1.

Plate next uses an approximation to derive a linear relationship describing the accuracy. Citing Abramowitz, Stegun, and Miller (1965), he writes:

$$erfc(x) < \frac{1}{x\sqrt{\pi}} e^{-x^2}.$$

This approximation allows Plate to estimate the linear relationship between N, M, $\log D$, and ϵ, arriving at

$$N < 8M \log\left(\frac{D}{\epsilon}\right).$$

The FA-CR-LEE approximation differs only by a factor of 2 because of the slightly different choice we made to approximate the cumulative gaussian as well as the different setup for the problem.

Subsequent work by Gallant and Okaywe (2013) proposed an alternative VSA framework that used a matrix as a binding mechanism. Based on their framework, they too, in analogous fashion to Plate (2003), derived an approximation to the capacity of vector symbols in superposition. Their derivation takes the high-fidelity factorial approximation as the starting point and uses e^{-x} as the bound on the tail of the normal distribution. This work is very similar to the derivation presented in this article, but we add more rigor and derive a tighter high-fidelity approximation utilizing

the Chernoff-Rubin bound and the updated approximation by Chang et al. (2011).

4.3 Derivations for Analog Gaussian Inputs

4.3.1 Analytic Capacity Bounds for Gaussian Analog Input. In section 2.5.1, we derived the memory capacity for gaussian inputs. Numerically, we showed that the equations suggest the memory capacity saturates to $1/(2\log 2)$ bits per neuron. It is possible to derive these capacity bounds analytically.

The total information is determined by $r(K)$, equation 2.57,

$$r(K) = \lambda^{2K} \frac{N(1-\lambda^2)}{D(1-\lambda^{2M})(1+\sigma_\eta^2/(DV_\Phi))},$$

and for gaussian variables, it can be computed as equation 2.58

$$I_{total} = \frac{D}{2} \sum_{K}^{M} \log_2 (r(K)+1).$$

Inserting equation 2.57 into 2.58, we obtain

$$I_{total} = \frac{D}{2} \sum_{K}^{M} \log_2 (r(K)+1) = \frac{D}{2} \log_2 \left(\prod_{K=1}^{M} (r(K)+1) \right). \tag{4.3}$$

The definition of the q-Pochhammer symbol or shifted factorial,

$$(a;q)_M := \prod_{K=0}^{M-1} (1-aq^K), \tag{4.4}$$

yields a more compact expression, equation 2.59,

$$I_{total} = \frac{D}{2} \log_2 ((-b_M q;q)_M)$$

with $q := \lambda^2$ and $b_M := \frac{N(1-q)}{D(1-q^M)(1+\sigma_\eta^2/(DV_\Phi))}$.

The approximation of the logarithm of the q-Pochhammer symbol for $|b_M q| < 1$ will now be useful (Zhang, 2013),

$$\log\left((b_M q;q)_\infty\right) = \frac{1}{2}\log(1-b_M q) - \frac{\tau}{2}Li_2(b_M q) - \frac{1}{6\tau}\frac{b_M q}{1-b_M q} + O(1/\tau^3)$$
$$= -\frac{\tau}{2}Li_2(b_M q) + \frac{1}{2}\log(1-b_M q) + O(1/\tau), \tag{4.5}$$

where $\tau := -2/\log(q)$ is the signal decay time constant for a given q and $Li_2(x)$ is the dilogarithm function. Note that for q close to one and large decay time constant, the first term in equation 4.5 becomes the leading term.

For the case $M \to \infty$ and any N, we can lead q so close to one that $b_\infty q = \epsilon$ becomes very small. This is accomplished by $q = 1 - \frac{\epsilon D(1+\sigma_\eta^2/(DV_\Phi))}{qN}$ and, equivalently, $\tau = \frac{2Nq}{\epsilon D(1+\sigma_\eta^2/(DV_\Phi))}$. With this setting, we can apply approximation 4.5 to compute the asymptotic information capacity. Neglecting nonleading terms,

$$\frac{I_{total}}{N} = \frac{D}{2N}\log_2\left((-b_\infty q;q)_\infty\right)$$
$$= -\frac{D\tau}{4N\log(2)}Li_2(\epsilon) = \frac{q}{2\log(2)(1+\sigma_\eta^2/(DV_\Phi))}$$
$$\approx_{q\to 1} \frac{0.72}{1+\sigma_\eta^2/(DV_\Phi)}. \tag{4.6}$$

Equation 4.6 uses the result for the polylogarithm: $\lim_{|z|\to 0} Li_s(z) = z$.

For the case of finite M and $D \propto N$, the identity $(a;q)_n = \frac{(a;q)_\infty}{(aq^n;q)_\infty}$ is useful in combination with approximation 4.5. We consider $D = N/\alpha$ with α, so that $b_M q = \frac{\alpha(1-q)q}{(1-q^M)(1+\sigma_\eta^2/(DV_\Phi))} = \epsilon$ becomes very small. Further, we set $\tau = -2/\log(q)$:

$$\frac{I_{total}}{N} = \frac{D}{2N\log(2)}\left[\log\left((-b_M q;q)_\infty\right) - \log\left((-b_M q^{M+1};q)_\infty\right)\right]$$
$$= \frac{\tau}{4\log(2)\alpha}\left[-Li_2(\epsilon) + Li_2\left(\epsilon q^M\right)\right] + O(\epsilon) + O(1/\tau)$$
$$= \frac{(1-q)q(1-q^M)\tau}{4\log(2)(1-q^M)(1+\sigma_\eta^2/(DV_\Phi))} + O(\epsilon) + O(1/\tau)$$
$$= \frac{1}{2\log(2)(1+\sigma_\eta^2/(DV_\Phi))} \times \frac{(1-q)q}{-\log(q))} + O(\epsilon) + O(1/\tau)$$
$$\approx_{q\to 1} \frac{0.72}{1+\sigma_\eta^2/(DV_\Phi)} \tag{4.7}$$

Thus, in both cases, we find the same asymptotic value for the information capacity.

4.3.2 Fixed Fidelity Retrieval Optimization for Memory Buffer. With the relation between r, K, and τ, we can find the τ_{opt} that maximizes K^* such that $r(K) \geq r^* \; \forall K \leq K^*$. Beginning with the SNR, equation 2.57,

$$r(K) = \lambda^{2K} \frac{N(1-\lambda^2)}{D(1+\sigma_\eta^2/(DV_{\mathbf{\Phi}}))} = e^{-2K/\tau} \frac{N(1-e^{-2/\tau})}{D(1+\sigma_\eta^2/(DV_{\mathbf{\Phi}}))}. \tag{4.8}$$

Setting $r(K^*) = r^*$ and solving for K^* gives

$$K^* = \frac{-\tau_{opt}}{2} \log\left(\frac{Dr^*(1+\sigma_\eta^2/(DV_{\mathbf{\Phi}}))}{N(1-e^{-2/\tau_{opt}})}\right). \tag{4.9}$$

Taking the derivative $dK^*/d\tau_{opt}$ and setting to 0,

$$-\frac{1}{2}\log\left(\frac{Dr^*(1+\sigma_\eta^2/(DV_{\mathbf{\Phi}}))}{N(1-e^{-2/\tau_{opt}})}\right) - \frac{e^{-2/\tau_{opt}}}{(1-e^{-2/\tau_{opt}})\tau_{opt}} = 0. \tag{4.10}$$

For moderately large τ_{opt}, the second term asymptotes to $\frac{1}{2}$, giving

$$\begin{aligned}
-1 &= \log\left(\frac{Dr^*(1+\sigma_\eta^2/(DV_{\mathbf{\Phi}}))}{N(1-e^{-2/\tau_{opt}})}\right), \\
e^{-1} &= \frac{Dr^*(1+\sigma_\eta^2/(DV_{\mathbf{\Phi}}))}{N(1-e^{-2/\tau_{opt}})}, \\
\tau_{opt} &= \frac{-2}{\log\left(1-\frac{eDr^*(1+\sigma_\eta^2/(DV_{\mathbf{\Phi}}))}{N}\right)}, \\
\frac{\tau_{opt}}{N} &= \frac{2}{eDr^*(1+\sigma_\eta^2/(DV_{\mathbf{\Phi}}))}.
\end{aligned} \tag{4.11}$$

From the first line of equations 4.11 and 4.9, it is easy to see that $K^* = \tau_{opt}/2$.

The information per neuron retrieved with a certain SNR criterion r^* is then given by

$$\frac{I^*(r^*)}{N} = \frac{D}{2N}\sum_{K=1}^{K^*} \log_2\left(r(K)+1\right) = \frac{D}{2N\log(2)} \log\left((b_\infty q; q)_{\tau_{opt}/2}\right) \tag{4.12}$$

with equation 4.11, $q = e^{-2/\tau_{opt}}$ and $b_\infty = er^*$.

For small r^* we can estimate

$$\frac{I^*(r^* \to 0)}{N} = \frac{D\tau_{opt}}{4N\log(2)}\left[Li_2(b_\infty q) - Li_2(b_\infty q^{\tau_{opt}/2+1})\right]$$
$$= \frac{D\tau_{opt} b_\infty}{4N\log(2)} q(1-q^{\tau_{opt}/2}) =_{r^*\to 0} \frac{1-q^{\tau_{opt}/2}}{2\log(2)(1+\sigma_\eta^2/(DV_{\mathbf{\Phi}}))}$$
$$= \frac{1-e^{-1}}{2\log(2)(1+\sigma_\eta^2/(DV_{\mathbf{\Phi}}))} \approx \frac{0.46}{1+\sigma_\eta^2/(DV_{\mathbf{\Phi}})}. \tag{4.13}$$

4.4 Capacity Results with MMSE Readout

4.4.1 Training Procedure for the Recurrent Neural Network. We used tensor flow to train a linear recurrent neural network at the sequence recall task. The parameter K could be altered to train the network to output the symbol given to it in the sequence K time steps in the history. However, larger K requires deeper backpropagation through time and becomes more expensive to compute and harder to learn. The training was based on minimizing the cross-entropy between $\mathbf{a}(m-K)$ and $\hat{\mathbf{a}}(m-K)$. The accuracy was monitored by comparing the maximum value of the output histogram with the maximum of the input histogram.

We initialized the network to have a random gaussian distributed encoding and decoding matrix ($\mathbf{\Phi}$, $\mathbf{V}(K)$) and a fixed random unitary recurrent weight matrix ($\mathbf{W}$). The random unitary matrix was formed by taking the unitary matrix from a QR decomposition of a random gaussian matrix. Such a matrix maintains the energy of the network, and with a streaming input, the energy of the network grows over time. After a fixed number of steps ($M = 500$), the recurrent network was reset, where the activation of each neuron was set to 0. This erases the history of the input. Only outputs K or more time steps after each reset were considered part of the energy function.

4.4.2 Comparison to DSR and Normal Networks in Reservoir Computing. White et al. (2004) describes the distributed shift register (DSR), a neural network that encodes an input sequence spatially by permuting the inputs down a chain of neurons. The final neuron in the chain, however, is not connected to any other postsynaptic neuron. This allows the network to store exactly N numbers. Since the last neuron is not connected, the network remembers exactly the last N most recent inputs, and the sequence length can be infinite. However, the network can be equated to our orthogonal networks with finite sequence length and a reset mechanism.

White et al. (2004) use a memory function that is the correlation between the input and decoded output to understand the information content

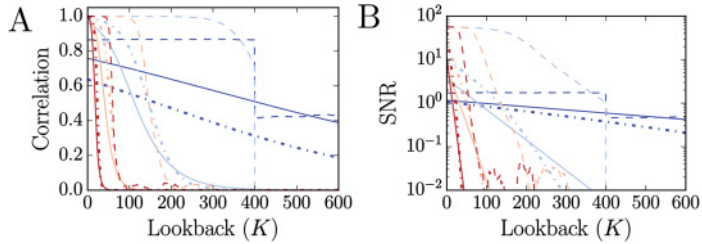

Figure 17: Memory performance in normal networks. Comparison of naive VSA readout (solid lines), approximation from White et al. (2004) (dot-dashed lines), and empirical performance (dashed lines) for analog memory buffers with different time constants, $\lambda^2 = [0.7, 0.95, 0.99, 0.999]$ red to blue, $N = 400$, $D = 1$. These parameters are taken from White et al. (2004).

of the DSR. This analysis extends to orthogonal networks with contracting weights, denoted *normal networks*. The correlation function for the DSR remains 1 until the look-back value K exceeds N, when the correlation function drops to 0. This has been taken as the upper limit on the information capacity for reservoir computing, and our results support this conclusion.

White et al. (2004) also derive an annealed approximation formula for the correlation function in normal networks. This approximation includes the MMSE readout and corresponds to analog memory buffer networks considered in section 2.6.2 (compare to Figure 12). We compare their theory to the naive VSA readout performance and the empirically measured performance of memory buffers with MMSE readout in Figure 17. The curves in Figure 17 match those in White et al. (2004, Figure 2; dot-dashed lines in Figure 17). These curves are given by

$$m(K) = \frac{\lambda^{2K} q}{1 + \lambda^{2K} q}, \tag{4.14}$$

where q satisfies

$$1 = N^{-1} \sum_{K=0}^{\infty} \frac{\lambda^{2K} q}{1 + \lambda^{2K} q} + \frac{\sigma_\eta^2 q}{1 + \lambda^2}. \tag{4.15}$$

These curves are generally more optimistic than the performance of naive VSA readout (see Figure 17, solid lines), except for larger time constants. However, the empirical performance of the MMSE readout (see Figure 17, dashed lines) still highly outperforms both naive VSA readout and the

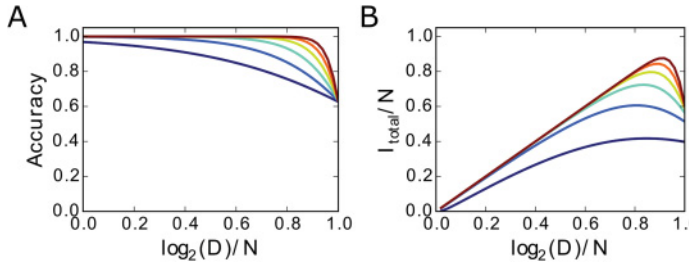

Figure 18: Finite size effects on information capacity in discrete DSRs with randomized codes. (A) The accuracy $p_{corr}^{M=1}$ with increasing N. (B) The retrieved information with increasing N.

White et al. (2004) theory. The MMSE readout performance approaches DSR-like performance, and nearly the full entropy of the neural space can be utilized if the time constant is appropriately optimized.

In our discrete framework, the discrete DSR would be considered a network with $D = 2$ (or with $D = 1$ and a detection threshold), and a sequence of binary inputs can be stored, with M increasing as high as N but not higher. Because of the way this representation is constructed, there is no interference noise, and the assumptions needed for our theory do not hold. However, the DSR can be reconsidered by focusing on the end result rather than the time evolution of the network. Ultimately, the DSR builds a binary representation for each possible binary input sequence and is equivalent to storing a single binary representation in the network. This is as if $M = 1$ and $D = 2^N$, where each possible input sequence corresponds to one of the D code vectors. Thus, we are able to apply our $M = 1$ analysis to understand the information capacity of DSRs and find that they are able to achieve the full capacity of binary neurons, i.e., 1 bit per neuron.

4.4.3 Randomized Vector Representations. In section 2.3.5, we compared the memory capacity of superpositions to the memory capacity of the $M = 1$ case as $D \to 2^N$. As D grows to a significant fraction of 2^N, the crosstalk from superpositions becomes overwhelming, and the memory capacity is maximized for $M = 1$. The retrieval errors are then due only to collisions between the randomized codevectors, and the accuracy $p_{corr}^{M=1}$ is given by equation 2.39. Figure 18A shows the accuracy for $M = 1$ as D grows to 2^N with a randomly constructed codebook for different (smaller) values of N; for large N, the numerical evaluation of equation 2.39 is difficult. As N grows, the accuracy remains perfect for an increasingly large fraction of the 2^N possible code vectors. However, at the point $D = 2^N$, the accuracy falls off to $(1 - 1/e)$, but this falloff is sharper as N grows larger. The information

retrieved from the network also grows closer and closer to 1 bit per neuron as N grows larger with $M = 1$ (see Figure 18B).

In Figure 18B, the capacity I_{total}/N of the randomly constructed codebook for $M = 1$ was computed with the equation we developed for superposed codes (see equation 2.27). However, the nature of the retrieval errors is different for $M = 1$; rather than crosstalk, collisions of code vectors are the error source. By performing an exhaustive analysis of the collision structure of a particular random codebook, the error correction can be limited to actual collisions, and the capacity of such a retrieval procedure is higher. The information transmitted when using the full knowledge of the collision structure is

$$I_{total} = \sum_C p_C \log_2 \left(\frac{p_C D}{C+1} \right). \tag{4.16}$$

For $D = 2^N$ and $N \to \infty$, the total information of a random vector symbol approaches 1 bit per neuron:

$$\lim_{N\to\infty} \frac{1}{N} \sum_C p_C \left(N + \log_2 \left(\frac{p_C}{C+1} \right) \right) \to 1. \tag{4.17}$$

It is an interesting and somewhat surprising result in the context of DSRs that a random codebook yields asymptotically, for large N, the same capacity as a codebook in which collisions are eliminated by construction (White et al., 2004). But it has to be emphasized that a retrieval procedure that uses the collision structure of the random codebook is necessary and advantageous only for the $M = 1$ case. For superpositions, even with just two code vectors ($M = 2$), the alphabet size D has to be drastically reduced to keep cross talk under control and the probability of collisions between random code vectors becomes negligible.

4.4.4 Capacity with Expected MMSE Readout. Further following White et al. (2004), we were able to compute $\tilde{\mathbf{C}}$, the expected covariance matrix of MMSE readout, without any training data (see section 2.6). White et al. (2004) focus on the memory buffer scenario where an infinite stream of inputs is given. They derive the result of MMSE readout in the buffer scenario, where they have an infinite input stream to act as training data. Their result is extended to higher D by equation 2.67. This matches the performance of empirically trained memory buffers for both discrete and analog inputs.

We derive an analogous equation for the expected covariance matrix of linear reset memories, equation 2.63. Computing $\mathbf{C}$ empirically requires training R parallel neural networks with the same connectivity but different

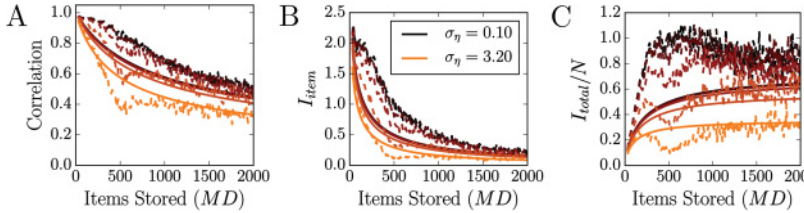

Figure 19: Performance of analog reset memory with expected covariance matrix. Compare to Figures 11E to 11H.

input sequences. The covariance matrix $\mathbf{C}$ is given by

$$\begin{aligned}\mathbf{C} &= \langle \mathbf{x}(M)\mathbf{x}(M)^\top \rangle_R \\ &= \frac{1}{R}\sum_r^R \left(\sum_{K_1}^M \mathbf{W}^{K_1}\mathbf{\Phi}\mathbf{a}(K_1;r) + \boldsymbol{\eta}(K_1;r)\right)\left(\sum_{K_2}^M \mathbf{W}^{K_2}\mathbf{\Phi}\mathbf{a}(K_2;r) + \boldsymbol{\eta}(K_2;r)\right)^\top \\ &= \sum_K^M \mathbf{W}^K\mathbf{\Phi}\left(\frac{1}{R}\sum_r^R \mathbf{a}(K;r)\mathbf{a}(K;r)^\top\right)\mathbf{\Phi}^\top\mathbf{W}^{-K} \\ &\quad + \sum_{K_1}^M\sum_{K_2\neq K_1}^M \mathbf{W}^{K_1}\mathbf{\Phi}\left(\frac{1}{R}\sum_r^R \mathbf{a}(K_1;r)\mathbf{a}(K_2;r)\right)\mathbf{\Phi}^\top\mathbf{W}^{-K_2} + M\sigma_\eta^2\mathbf{I}. \qquad (4.18)\end{aligned}$$

The covariance matrix is broken up into three parts: the diagonal term, the cross term, and the noise term. For $R \to \infty$, the diagonal term contains $\frac{1}{R}\sum_r^R \mathbf{a}(K;r)\mathbf{a}(K;r)^\top \to \mathbf{I}\sigma^2(\mathbf{a})$. The cross term converges to 0 if $\mu(\mathbf{a}) = 0$. This leaves equation 2.63:

$$\tilde{\mathbf{C}} = \langle \mathbf{x}(M)\mathbf{x}(M)^\top \rangle_\infty = M\sigma_\eta^2\mathbf{I} + \sum_{k=1}^M \mathbf{W}^k\mathbf{\Phi}\mathbf{\Phi}^\top\mathbf{W}^{-k}.$$

Using $\tilde{\mathbf{C}}$ in the readout does exceed the naive readout performance; however, it does not perform as well as an empirically trained MMSE network with finite R. Our simulations appear to be approaching the theory of equation 2.63, but second-order terms seem to play an important role in eliminating the cross-talk noise. These terms are small but still significant even for very large R. The performance of readout with the expected covariance matrix is shown in Figure 19 and can be compared to Figures 11E to 11H.

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