An Algorithmic Approach to Biological Information and Long-term Memory

John L. Pfaltz
Dept. of Computer Science,
University of Virginia

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Abstract

This paper presents two computable functions, \mathcal{R} and \mathcal{E} , that map networks into networks. If all cognition occurs as an active neural network, then it is thought that \mathcal{R} emulates long-term memory consolidation and \mathcal{E} models memory recall. These algorithms employ an intermediate form of network which is of considerable interest because it could be a model of biological information. It is a system of "chordless cycles" which can be shown to function like a vector space, and are thus ideal for the representation of data. It is thought that they could be the structural substrate of long-term memory; just as the double helix is the necessary substrate for genomic memory.

Such cycle systems are not just a mathematical construct; they exist in every cell of our bodies.

1 Introduction

There seems to be consensus that our sensations, ideas, and memories are really just active networks of neurons in our brains [24, 50]. And we have a good idea where in the brain specific kinds of mental activity occur, e.g. [36, 47] But, to our knowledge, no one has any idea as to what kinds of networks correspond to any specific sensation, concept or memory.

We know that neurons can stimulate other neurons by means of electric (or chemical) charges proceding along an axon to one, or more, synapses [49]. That would suggest that a directed, asymmetric network is a reasonable model. However, such an asymmetric network may best model neuronal behavior, but not neuronal state. Many neurons are interconnected by dendrites [32]. These are thought to be bi-directional, thus implementing symmetric relationships that may encode a state to be used by an active neuron.

Given this state of uncertainty, we have chosen to explore symmetric relationships, or graphs or networks, in this paper. Some of the mathematical results we present will be true

as well for asymmetric (directed) networks; some would require minor rewording; and some will no longer be true at all.

Regardless of whether our neural networks are essentially symmetric or asymmetric, it would appear that a mathematical treatment of networks, or graphs, or relationships, is a fruitful way to approach them. That we will do in this paper, parts of which are an expanded version of [46].

In Section 2, we clarify our interpretation of relationships and their visual representation as graphs or networks. We also introduce the concept of "closure". In Section 3, we describe a computational process, \mathcal{R} , which reduces any network to its unique, irreducible "trace". We will claim that this procedure appears to model the process of long-term memory "consolidation".

The irreducible trace of a network has unusual properties. In Section 4 it is shown to be a cycle system which behaves very much like a vector space under cycle composition. It seems ideal for representing biological properties.

The \mathcal{R} process is a well-defined function over the space of all finite networks in that for any network \mathcal{N} , \mathcal{R} yields a unique irreducible trace \mathcal{T} . Thus the inverse set, \mathcal{R}^{-1} , defines the abstract set of all networks that reduce to the same specific irreducible trace. In Section 5, we present a computational process which generates specific members within \mathcal{R}^{-1} . We will argue that this can model memory "recall" and "reconsolidation".

In Section 6 we present additional biological evidence to support our claims to model long-term memory consolidation and recall. Certain selected mathematical proofs are presented in an appendix, Section 7.

2 Sets, Relations, and Closure

Our computations are set based. The nature of the elements comprising the sets play no part, and can be quite arbitrary. So unlike most computational systems in which the variables will be int or float, our variables have type setid. We program using a set manipulation package in C++ with operators such as is_contained_in and union_of. Sets themselves are represented as extensible bit strings, so that the operators above are effectively of order O(1). There is no theoretical upper bound of these sets, but we have not tested it with sets of cardinality exceeding 50,000. A somewhat fuller description is given in [38]. All the following set-based operators and procedures have been implemented, and fully tested, using this system.

We use a standard set notation. A **set** S is comprised of elements $\{a, b, \ldots, y, z\}$ of unspecified type. The curly braces $\{\}$ indicate that these elements are regarded as a "set". Sets are denoted by upper case letters, e.g. X, Y; elements are always lower case, e.g. x, y. Sometimes we elide the commas, as in $Y = \{abc\}$.

If an element x is a member of the set X, we write $x \in X$. If a set X is contained in

another set Y, that is, $x \in X$ implies $x \in Y$ (here x is a variable running over all elements of X), we write $X \subseteq Y$. If the containment is **strict**, that is there exists $y \in Y$, $y \notin X$, we write $X \subset Y$. By $X \cup Y$ and $X \cap Y$ we mean the union and intersection (meet) of X and Y respectively.

One may have a "set of sets", which we call a **collection**, and denote with a caligraphic letter. Thus we may have $X \in \mathcal{C}$.

2.1 Relationships

Let N be any set. A **relation**, η , on N is a function, which given any subset $Y = \{y_1, y_2, \ldots, y_k\} \subseteq N$ returns the related set $Y.\eta = \{z_1, z_2, \ldots, z_n\} \subseteq N$. This is a bit unusual. It is more common to think of relations in terms of links, or edges, between elements, such as illustrated by the undirected graph, or network, of Figure 1. The relation η is sometimes regarded as a set of "edges" in a graph theoretic approach. But we prefer to define relations in terms of sets and functional operators. It provides an additional measure of generality which can be of value. We emphasize this set-based definition by using suffix notation, such as $Y.\eta$ to mean the set of elements $\{z\}$ that are related to Y by η . We call $Y.\eta$ "Y's **neighborhood**". In Figure 1, $\{a\}.\eta = \{a,d,f\}$, $\{d\}.\eta = \{a,b,d,f,g\}$, and



Figure 1: A small network illustrating neighborhood properties

 ${a,e}, \eta = {a,b,c,d,e,f,h}.$

If η has the property that

P1:
$$Y.\eta = \bigcup_{y \in Y} \{y\}.\eta$$
 (extensibility)

that is, $Y.\eta$ is the union of all the subsets $\{y\}.\eta$ for all $y \in Y$, we say that η is **extensible**, or graphically representable, so that Figure 1 is an accurate representation of η .

If the relationship is not extensible, then it constitutes a "hypergraph" [6, 18]. To more easily illustrate the concepts of this paper with graphs, we will assume that η is extensible; but unless explicitly noted none of the mathematical assertions require it. Moreover, we observe that for large, sparse relationships, matrix representations and operations are quite impractical [56].

In addition to extensibility, P1, a relationship η may also have any of the following 3 properties: that for all $X, Y \subseteq N$,

P2:
$$Y \subseteq Y.\eta$$
 (expansive or reflexive)¹

¹This is primarily for mathematical convenience.

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P3: X \subseteq Y implies X.\eta \subseteq Y.\eta (monotone)<sup>2</sup>
P4: X.\eta = Y implies Y.\eta = X (symmetric)<sup>3</sup>
```

The relation of Figure 1 is symmetric; its graph is **undirected**. By a **network**, $\mathcal{N} = (N, \eta)$, we mean a set N of nodes or elements, together with any relationship η . For this paper, we require that η satisfy the functional properties P2, P3 and P4.

2.2 Closure

The mathematical concept of "closure" plays a key role in our approach. In a discrete world, the interpretation of closed sets is somewhat different from the more traditional concepts encountered in classical point-set topology. Our view is that a closure operator, φ , is a set-valued function whose domains are also sets. If Y denotes any set, $Y.\varphi$ denotes its closure; that is it is the smallest closed set containing Y. Thus, like η , φ is a well-defined function mapping subsets, $X,Y \subseteq N$ of a given network into other subsets of N. More formally, φ is a **closure operator** that satisfies the following 3 closure axioms, for all sets X,Y:

```
C1: Y \subseteq Y.\varphi (expansive),
C2: X \subseteq Y implies X.\varphi \subseteq Y.\varphi (monotone), and
C3: Y.\varphi.\varphi = Y.\varphi (idempotent)
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Readily, any relationship operator, η , satisfying properties P2 and P3 is almost a closure operator. It has only to satisfy the idempotency axiom. But normally, $Y.\eta \subset Y.\eta.\eta$ since neighborhoods tend to grow.

An alternative definition of closure asserts that a collection $C = \{C_1, \ldots, C_n\}$ can be regarded as the closed sets of a superset N if and only if

C4: if
$$C_i, C_k in \mathcal{C}$$
 then $C_i \cap C_k \in \mathcal{C}$.

It is not difficult to prove that C4 implies C1, C2, and C3, and conversely.

We normally think of closure in terms of its operator definition. Because φ is expansive, C1, the superset N must be closed; by C4, if any two closed sets are disjoint, the empty set \emptyset must also be closed.

2.3 Neighborhood Closure

One important closure operator φ , called **neighborhood closure**, can be defined with respect to network relationships. We let

$$Y.\varphi = \bigcup_{z \in Y.\eta} \{\{z\}.\eta \subseteq Y.\eta\} \tag{1}$$

That is, if $z \in Y.\varphi$ then z is not related to any elements that are not already related to Y. Readily, since $\{a\}.\eta = \{adf\} \subseteq \{abdfg\} = \{d\}.\eta$ in Figure 1, $a \in \{d\}.\varphi$. Convince yourself

²Probably essential. If η is not monotone, we can prove very few mathematical results of interest.

³Unnecessary, relaxed in other papers such as [44, 45].

that $\{d\}.\varphi = \{adf\}$ and $\{h\}.\varphi = \{h\}$. It is not hard to show that φ , so defined with respect to η satisfies the closure axioms C1, C2 and C3 and that for all $\{y\}, \{y\} \subseteq \{y\}.\varphi \subseteq \{y\}.\eta$.

3 Irreducible Networks

Even though we are fairly sure that all mental activity, that is sensory apprehension, cognition, and ideation are creatures of our brain's neural system and we know what parts of the brain this activity is located [23, 29, 50], exactly which configurations of activated neurons might correspond to a particular experience or idea is totally unknown. The 206 interconnected nodes of Figure 2, some of which have been labeled with letters and integers, may

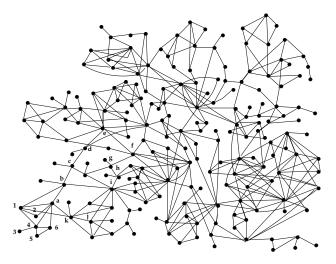


Figure 2: A moderately complex network whose nodes and links might model a neural configuration [7, 8].

be thought to schematically represent a momentary configuration of neurons in a mental process. Assuming this, we ask how could this mental experience be condensed into a more economical coded version, either for transmission or to be "remembered"?

If a singleton set $\{y\}$ is not closed, say $z \in \{y\}.\varphi$, then $\{z\}.\eta \subseteq \{y\}.\eta$, so z contributes little to understanding the structure of η in terms of closure. In the small network of Figure 1, $\{d\}.\varphi = \{a,d,f\}$ and $\{e\}.\varphi = \{c,e\}$. Removing the nodes a,f,c in this network will leave each of the singleton sets, $\{b\}$, $\{\}$, $\{e\}$, $\{g\}$ and $\{h\}$ as closed sets with respect to the η operator.

In Figure 2, $\{4\}$. $\varphi = \{3, 4, 5, 6\}$. Removing the nodes 3, 5, and 6, and their connections to 4, results in minimal information loss with respect to η as a whole. However, with these nodes gone, we now have $\{4\}$. $\eta = \{a, 1, 4\} \subseteq \{a, 1, 2, 4\} = \{a\}$. η . So the two elements, 1, 4,

and their connections can be removed as well. In fact the entire pendant substructure in the lower left corner, $\{1, 2, 3, 4, 5, 6\}$ on $\{a\}$, can be removed with no loss of global information. If it is removed $\{a\}.\varphi = \{a\}$, so $\{a\}$ is closed.

We say a network $\mathcal{N} = (N, \eta)$ is **irreducible** if every singleton set, $\{y\}$, is closed. That is, if for all $y \in N$, $\{y\}$.

If $\{y\}$ is not closed, only elements z in $\{y\}.\eta$ could possibly be in $\{y\}.\varphi$ so only those need be considered. If $\{z\}.\eta\subseteq\{y\}.\eta$ so that $\{z\}.\varphi\subseteq\{y\}.\varphi$, we say z is **subsumed** by y, or z **belongs** to y. We can remove z from N, together with all its connections, and add z to $\{y\}.\beta$, the set of all nodes belonging to $\{y\}$ which we call its β -set. Since, $y\in\{y\}.\beta$, its cardinality, or β -count, $|\{y\}.\beta| \ge 1$, a value we will use in Section 5. We use the pseudocode of Figure 3 to implement the process \mathcal{R} that reduces any network \mathcal{N} to its irreducible core, which is called its **trace**, $\mathcal{T}.^4$ This version of \mathcal{R} only records β -counts, not entire β -sets.

Figure 3: Reduction code, implementing \mathcal{R}

The irreducible trace, \mathcal{T} , of Figure 2 is shown in Figure 4. The trace is the dark network on the 83 elements with 234 bolder connections. Sets of subsumed nodes, or β -sets, have been encircled with dashed lines. The largest β -set, comprised of 12 nodes, is $\{y\}$. β in the upper right corner. The β -set $\{a\}$. β in the lower left corner consists of 7 nodes (including a itself). A total of 123 nodes were subsumed and eliminated.

In this particular network the outer loop of Figure 3 was executed 4 times with a 5th pass to verify that there remained no more reduceable nodes. The order in which individual nodes y are examined is arbitrary. One can create networks that require n = |N| iterations of this outer loop. So this process has a theoretical complexity of $O(n^2)$. However, in tests with rather complex networks of several thousand nodes, the maximal number of iterations

⁴In [43], this was called the "spine" of \mathcal{N} .

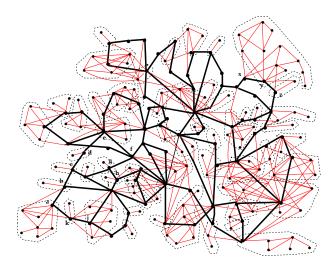


Figure 4: The irreducible trace \mathcal{T} of Figure 2

has never exceeded 7. Its effective complexity appears to be quite reasonable. Moreover, because of its local nature, the inner loop could be easily implemented in parallel.

We keep speaking of the function \mathcal{R} . It can be shown (see Section 7), that for any network \mathcal{N} , its irreducible trace, \mathcal{T} , is unique (up to isomorphism). Therefore, the pseudocode of Figure 3 does indeed embody a well-defined computational function which we denote by \mathcal{R} . Not only is \mathcal{R} a function, we can actually characterize its image, $\mathcal{T} = \mathcal{N}.\mathcal{R}$. When η is symmetric, if y is a node in $\mathcal{T} = \mathcal{N}.\mathcal{R}$, then y will be either: (a) an isolated node; (b) an element of a chordless cycle of length ≥ 4 ; or (c) an element in a path between two chordless cycles of length ≥ 4 (again see Section 7).

4 Chordless Cycles

The bold edges of the irreducible trace in Figure 4 constitute a system of chordless cycles. A **chordless cycle** is most easily visualized as a necklace of pearls (or beads). More formally, it is a sequence $\langle y_1, y_2, \ldots, y_n, y_1 \rangle$ where $y_{i\pm 1} \in \{y_i\}.\eta$, $y_1 \in \{y_n\}.\eta$ and $y_{i\pm k} \notin \{y_i\}.\eta$ if k > 1. In Figure 1 the 5-cycle $\langle b, d, g, h, e, b \rangle$ is chordless. In Figures 2 and 4 the 5-cycle $\langle a, b, i, j, k, a \rangle$ is also chordless, as is the 8-cycle $\langle b, c, d, e, f, g, h, i, b \rangle$. But the combined 11-cycle $\langle a, b, c, d, e, f, g, h, i, j, k, a \rangle$ is not. The link (b, i) is a chord.

Granovetter [26] called chordless cycles the "weak connections" of a social network. He felt they were the key to understanding the network structure as a whole. Even so, chordless cycle structures have been relatively unstudied, while "chordal graphs" (with no chordless cycles) have an exhaustive literature [35].

We can explore the mathematical properties of chordless networks a bit further. If some

collection, S, of subsets C_i of elements in S has the property that no set C_i is completely contained in another, then S^n , where n = |S|, constitutes a Sperner system.⁵ Such systems are so called after Emanuel Sperner who first described them [17]. With a little thought, we see that if C_i and C_k are chordless cycles, then C_i cannot contain C_k . Consequently, a system of chordless cycles $\{C_i\}$ constitutes a Sperner system with each cycle being a unique subset in this system. This property establishes a bijection between the set N of nodes and the set E of edges in any chordless cycle system. Consequently, each cycle C_i can be uniquely identified by its set of edges, which we denote by \bar{C}_i , or by its constituent nodes which we denote by \dot{C}_i .

This irreducible trace, \mathcal{T} , of chordless cycles preserves a number of important properties found in the original network, \mathcal{N} . First, it preserves the shortest path structure between retained nodes. Consequently, connectivity and the distances between nodes (as usually defined) are preserved. Further, "network centers", [5, 19, 20], whether with respect to distance or "betweenness", are preserved in the trace.

4.1 Cycle Composition

One can compose chordless cycles in an algebraic fashion. Consider the much smaller network of chordless cycles in Figure 5. Readily, it consists of 3 chordless cycles, \dot{C}_1 =

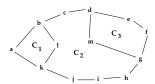


Figure 5: A small representative graph of chordless cycles.

 $\{ablk\}, \dot{C}_2 = \{bcdmghijkl\}$ and $\dot{C}_3 = \{defgm\}$ of lengths 4, 10 and 5 respectively. (We normally elide the commas when enumerating sets if no confusion is possible.)

But there are more than 3 chordless cycles in Figure 5. One is the cycle $C_4 = \dot{C}_4 = \{abcdmghijk\} = (\bar{C}_1 \cup \bar{C}_2) - (\bar{C}_1 \cap \bar{C}_2)$. We say C_m is the **composition** of C_i and C_k , denoted $C_i \circ C_k$, where

$$C_m = C_i \circ C_k = \dot{C}_m = (\bar{C}_i \cup \bar{C}_k) - (\bar{C}_i \cap \bar{C}_k)$$

$$\tag{2}$$

Not only is $C_4 = C_1 \circ C_2$ we have $C_5 = \{bcdefghijkl\} = (\bar{C}_2 \cup \bar{C}_3) - (\bar{C}_2 \cap \bar{C}_3)$ and $C_6 = C_3 \circ C_4 = C_1 \circ C_5 = C_1 \circ C_2 \circ C_3$. Because, $\bar{C}_1 \cap \bar{C}_3 = \emptyset$, the composition $C_1 \circ C_3$ is simply the two disjoint cycles, C_1 and C_3 ; that is $C_1 \cup C_3$. The disjoint union of chordless cycles must be treated mathematically as a chordless cycle itself.

⁵A collection of subsets $\{C_i\}$ is a Sperner system if $C_i \not\subset C_k$ for all $i \neq k$.

The **empty cycle**, C_{\emptyset} , in which $\dot{C}_{\emptyset} = \emptyset = \bar{C}_{\emptyset}$ functions as the identity element for the composition operator since for all i, $C_i \circ C_{\emptyset} = C_i$.

Table 1 completely details the composition operator over the 8 cycles in the cycle system of Figure 5. Here we are using $C_{1\cup 3}$ to denote $C_1\cup C_3$.

0	C_1	C_2	C_3	C_4	C_5	C_6	$C_{1\cup 3}$	C_{\emptyset}
C_1	C_{\emptyset}	C_4	$C_{1\cup 3}$	C_2	C_6	C_5	C_3	C_1
C_2	C_4	C_{\emptyset}	C_5	C_1	C_3	$C_{1\cup 3}$	C_6	C_2
C_3	$C_{1\cup 3}$	C_5	C_{\emptyset}	C_6	C_2	C_4	C_1	C_3
C_4	C_2	C_1	C_6	C_{\emptyset}	$C_{1\cup 3}$	C_3	C_5	C_4
C_5	C_6	C_3	C_2	$C_{1\cup 3}$	C_{\emptyset}	C_1	C_4	C_5
C_6	C_5	$C_{1\cup 3}$	C_4	C_3	C_1	C_{\emptyset}	C_2	C_6
$C_{1\cup 3}$	C_3	C_6	C_1	C_5	C_4	C_2	C_{\emptyset}	$C_{1\cup 3}$
C_{\emptyset}	C_1	C_2	C_3	C_4	C_5	C_6	$C_{1\cup 3}$	C_{\emptyset}

Table 1: Composition table for the cycle system of Figure 5.

It is not hard to show that for all i, k, m,

- (a) $C_i \circ C_k = C_k \circ C_i$,
- (b) $C_i \circ (C_k \circ C_m) = (C_i \circ C_k) \circ C_m$,
- (c) $C_k \circ C_k = C_{\emptyset}$, and
- (d) If $C_m = C_i \circ C_k$ then $C_k = C_i \circ C_m$.

Thus, a collection of chordless cycles constituting an irreducible network is an Abelian group of order 2.

4.2 Matroids

A matroid is a generalized vector space. It has basis elements, every set of which must have the same cardinality, which is said to be its *rank*. There is an abundance of literature on matroids, of which [3, 15, 59, 61] are just a sample.

A set $S = \{C_i\}$ of non-empty cycles is said to be **dependent** if there exists $C_m \in S$ such that $C_m = C_i \circ \ldots \circ C_k$, where $C_i \ldots, C_k \in S$. If S is not dependent, it is said to be **independent**. Independent sets are normally denoted by I. The non-empty cycles of an independent set are said to be **basic**. Any 3 of the non-empty cycles in Table 1 constitute an independent set.

We introduced closure operators in Section 2. Let $Y = \{C_1, \ldots, C_k\}$ be a set of cycles. By the **span** of Y, denoted $Y.\sigma$, we mean the set of all cycles $\{C_m\}$ such that $C_m = C_i \circ \ldots \circ C_j \circ \ldots \circ C_k$, where $C_j \in Y$ for all $j.^6$ If $C_m \notin Y$ then Y consists of a set of independent cycles. The spanning operator, σ is a closure operator over any set Y of cycles.

⁶In graph theory, the term "span" usually refers to a tree whose nodes include all $y \in N$. Since a tree has no cycles, it has no connection to our usage which is taken from the notion of spanning vector spaces.

A closure system is said to be a **matroid** if it satisfies the Steinitz-MacLane **exchange** axiom [27, 31, 61], that is:

if $x, y \notin Y \cdot \sigma$ and $y \in (Y \cup x) \cdot \sigma$ then $x \in (Y \cup y) \cdot \sigma$.

If σ satisfies the anti-exchange axiom [40], that is:

if $x, y \notin Y.\sigma$ and $y \in (Y \cup x).\sigma$ then $x \notin (Y \cup y).\sigma$

then the system is called an antimatroid.

Let \mathcal{C} be a cycle system and let σ be the spanning operator. We can show (see Section 7) that the system (\mathcal{C}, σ) satisfies the Steinitz-Maclane exchange axiom and is thus a matroid.

Since \mathcal{C} is a matroid, the cardinality of every maximal independent set is fixed and this number, r, is the **rank** of the system. The network of Figure 5 has rank 3.

A.I. and machine learning frequently use a vector space as their mathematical base. A feature vector denotes various properties of an individual element of the space. What we have shown is that individual cycles within a chordless cycle system have very similar properties.

We must note that a "cycle matroid" has been defined in the literature [61] in which any tree is independent and any subgraph containing a cycle (or circuit) is dependent. Thus the collection of all spanning trees constitutes the unique basis set for the matroid. This concept of a "cycle matroid" is very different from that described above.

4.3 Distribution of Chordless Cycles

The combinatorics of Sperner sets provides one mechinism for encoding information. The distribution of cycle lengths in a single irreducible network provides another. Counting the numbers of cycles of length k in a specific chordless network, \mathcal{N} , is not easy. Using the Sperner set property, the author has employed a brute force counting process that is limited by the size of \mathcal{N} which is described in [1]. Diane Castonguay, Elisangela Silva Dias et al. have developed more effective ways of counting [14, 30]. In [43], this distribution is called the signature of the network. The average cycle length is 23.4 in this network and the 6 longest chordless cycles have length 35.

4.4 Consolidation and the Memory Trace

The physical nature of human long-term memory is not at all a settled matter. We are fairly certain that the hippocampus of the brain is heavily involved [2, 22, 47]; but just how is not completely understood. One school of thought posits that long-term memories are recorded in some form of "memory trace", or "engram" [11, 52, 58, 48]. But, because no trace of these supposed "memory traces" has ever been physically detected (pun intended), others disbelieve this theory [12, 37]. We explore the possibility of that biological cycle systems constitute "memory traces" in Section 6.2.

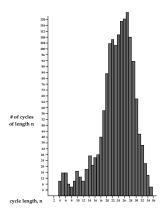


Figure 6: Distribution of chordless cycles in the irreducible trace of Figure 4.

There is more consensus that some form of processing which distinguishes long-term memory from short-term memory does occur. This process is commonly called **consolidation** [4, 33, 37]. We believe that \mathcal{R} is analogous to consolidation, and that chordless cycles, in some form, are analogous to the elusive "memory trace", whence our terminology.

5 Expanding Irreducible Networks

Since \mathcal{R} is a well-defined function mapping the space of all finite, symmetric networks into itself, one can consider $\mathcal{N}.\mathcal{R}^{-1}$, which is the collection of all networks \mathcal{N}_i such that $\mathcal{N}_i.\mathcal{R} = \mathcal{T} = \mathcal{N}.\mathcal{R}$. In this section we explore the nature of networks in this inverse set and how to generate them.

Two such networks, \mathcal{N}_i and \mathcal{N}_k , that have the same irreducible trace are said to be **structurally similar**. Readily, structural similarity is an equivalence relation. Even though \mathcal{N}_k may be similar to \mathcal{N}_i , they may have very different cardinalities. A network $\mathcal{N}_k = (N_k, \eta_k)$ is said to be **strongly similar** to $\mathcal{N}_i = (N_i, \eta_i)$ if $\mathcal{N}_k \mathcal{R} = \mathcal{N}_i \mathcal{R}$ and $|N_k| = |N_i|$.

The pseudocode below in Figure 7 describes a computational process \mathcal{E} that, given the trace \mathcal{T} of a network \mathcal{N} together with β -counts, randomly expands it to a strongly similar network $\mathcal{N}' = \mathcal{N}.\mathcal{R}.\mathcal{E}$. The process choose_random_in returns a random subset of its argument. Since $\{z\}.\eta = S \subseteq \{y\}.\eta$, the node z will be subsumed by (or belong to) y if reduced again ensuring that $\mathcal{N}.\mathcal{R}.\mathcal{E}.\mathcal{R} = \mathcal{N}.\mathcal{R}$. When a node $\{y\}$ is expanded, its β -count is decremented, and if > 1, part of the remainder may be added to the β -count of $\{z\}$. Consequently, by creating just as many new nodes as had belonged to any node $\{y\}$, we ensure that $|\mathcal{N}'| = |\mathcal{N}|$. This kind of \mathcal{E} process has been called an "expansion grammar" in [41]. The construction of \mathcal{E} , where $\{z\}.\eta \subseteq \{y\}.\eta$, assures us that $\mathcal{T}.\mathcal{E}.\mathcal{R}$ will be \mathcal{T} again.

```
for all {y} in N
    {
    while (|{y}.beta| > 1)
        {
        create new node z;
        S = choose_random_in ({y}.nbhd);
        {z}.nbhd = S;
        k = random_int(1, |{y}.beta|-1);
        |{y}.beta| = |{y}.beta| - k;
        |{z}.beta| = k;
        add {z} to N;
    }
}
```

Figure 7: Pseudocode for \mathcal{E} which generates strongly similar networks.

Consequently, for any network $\mathcal{N}' = \mathcal{T}.\mathcal{E}$, $\mathcal{N}' \in \mathcal{N}.\mathcal{R}^{-1}$, so \mathcal{N}' and \mathcal{N} are structurally similar.

Let \mathcal{N} be the network of Figure 2. The following Figure 8 shows a network \mathcal{N}' that was randomly expanded by \mathcal{E} , given the irreducible trace \mathcal{T} of Figure 4. The numbered nodes

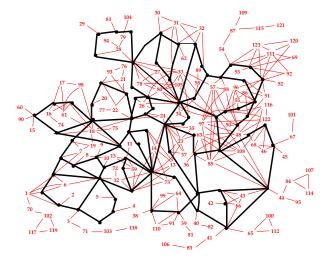


Figure 8: A reconstructed network $\mathcal{N}' = \mathcal{T}.\mathcal{E}$ in $\mathcal{N}.\mathcal{R}^{-1}$ that is strongly similar to \mathcal{N} of Figure 2

were randomly appended to the trace in numerical order and roughly correspond to the 123 nodes that were subsumed in Figure 4. They are numbered in the order that they were attached to the darker irreducible trace. \mathcal{N}' is strongly similar to the network \mathcal{N} of Figure

2 because $\mathcal{N}'.\mathcal{R} = \mathcal{T} = \mathcal{N}.\mathcal{R}$.

Such a semi-random "retrieval" process may be inappropriate in computer applications [45], but it seems to model biological recall rather well. It has been observed that the recall and reconstruction of our long-term memories is seldom exact [9, 10, 28]. Our memories often are confused with respect to detail, even when they are generally correct. Reconstruction of a network trace by \mathcal{E} has these very properties.

Given that for all networks \mathcal{N} , $\mathcal{N}.\mathcal{R}.\mathcal{E}.\mathcal{R} = \mathcal{N}.\mathcal{R} = \mathcal{T}$, it also supports the notion of "reconsolidation" which asserts than long-term memories are repeatedly recalled and re-written with no change, unless deliberately distorted in our (semi)conscious mind [37, 57].

6 Biological Memory

A computational model need not actually explain the behavior that it models. For example, the path of a thrown projectile has an excellent parabolic model. However, further study of this conic formulation contributes little to the understanding of either gravity or air resistence. By the same token, there need not be closure operators or chordless cycles involved in the performance of human memory, for the model to be valid. But, it would be a powerful verification of this model if we could demonstrate the existence of chordless cycle structures in a memory representation. We can't. Neither, to our knowledge, does anyone else know the actual structural format of our long-term memory.

Throughout this paper we have suggested parallels found in various memory studies. But, do these computational processes, \mathcal{R} and \mathcal{E} , really model biological memory? We just don't know. Are long-term memories really encoded as chordless cycles? In this section we offer a few more tantalizing clues which may, or may not, be significant.

6.1 Role of Closure

We employed "closure" as the basic mathematical concept in the preceding development. But, are instances of closure actually found in biological organisms? We offer two suggestive examples.

First, The visual pathway consists of layers of cells, beginning with the rods and cones of the retina passing stimuli toward the primary visual cortex. The neurological structure of this visual pathway is reasonably well understood, c.f. [24, 49, 54]. The individual functions of its layers are less well so.

Imagine that Figure 9 depicts a cross section of the retinal region. Dark cells denote visually excited cells. Although tightly packed, the actual neuronal structure is not as regular as this hexagonal grid; but this regularity plays no part in the process.

Let α be an existential operator defined as $Y.\alpha = black$ (excited), if and only if $\exists z \in Y.\eta$ where z is black (excited). Let β be the existential operator defined by $Y.\beta = white$ (quiesent), if and only if $\exists z \in Y.\eta$ such that z is white (quiesent). Figure 10(a) illustrates

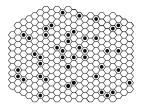


Figure 9: Excited cells in a cross section of the visual cortex.

the excited (small \times) neighbors of Figure 9. Figure 10(b) illustrates $Y.\alpha.\beta$ in which all

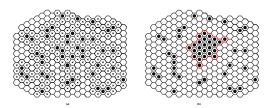


Figure 10: (a) $Y.\alpha$, excited cells, (b) $Y.\alpha.\beta$, remaining excited cells.

excited cells of Figure 10(a) that have at least one quiesent (white) neighbor become quiesent (white). The resulting central figure becomes evident; it is a closed object, because the pair of operators $(\alpha.\beta)$ is a closure operator. The pair $(\alpha.\beta)$ is idempotent because iterating them, as in $Y.(\alpha.\beta).(\alpha.\beta)$ yields no new black (excited) cells.

This two step operation can occur at the neural firing rate. It is an effective parallel process that was first proposed to eliminate salt and pepper noise in computer imagery [51]. Readily, such a "blob detection" capability would have evolutionary value. Does such a capacity exist? We don't know for sure. But, it is thought that the visual pathway is organized in an alternating manner to facilitate precisely this kind of two-step processing [54]. In any case, this example illustrates that this kind of all, or nothing, logic in which "for all" $(\forall x)$ can be interpreted as "there does not exist not x" $(\neg \exists \neg x)$ which is needed to implement the closure operator of (1) can be rendered in neural circuitry.

The second example is also "cognitive". In the development of "Knowledge Spaces" [16], Doignon and Falmagne call a coherent collection of facts or skills a "knowledge state". These are closed sets which are partially ordered by containment to form a lattice structure [40], which they call a "knowledge space". There is a considerable literature concerning closed knowledge "states" and knowledge "spaces". A somewhat similar approach to cognitive closure was presented in [44]. Closure operators can be an important aspect of cognitive behavior.

⁷Cord Hockemeyer, http://www.uni-graz.at/cord.hockemeyer/KST_Bibliographie/kst-bib.html, maintains a bibliography of over 400 related references.

6.2 Role of Chordless Cycles

Surprisingly, chordless cycles abound in all biological organisms as protein polymers.

One example, found in every cell of our bodies, is a 154 node phenylalaninic-glycinerepeat (nuclear pore protein), \mathcal{N} , which is shown in Figure 11.⁸ One can easily see the

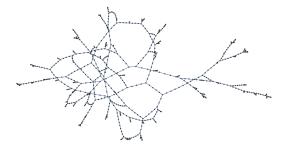


Figure 11: A 154 node protein polymer

chordless loops, with various linear tendrils attached to them. When these are removed by \mathcal{R} , there are 107 remaining elements involved in the chordless cycle structure. These protein polymers are thought to regulate transport of other proteins across the nuculear membrane [21, 39, 60].

The cyclic structure is even more apparent in the single 17-cycle with appendages shown in Figure 12. It depicts the HACN73 pheromone associated with civit cats. 9 Pheromones



Figure 12: The HACN73 pheromone.

are a basic biological signaling mechanism.

Readily, organisms with any form of memory, e.g. "movement toward light yields food", have survival benefit. Nature appears to reuse successful structures. If chordless cycles can successfully regulate one form of transport, it would not be surprising if evolutionary pressure led to their use in other control mechanisms.

⁸This network, \mathcal{N} , that we received from a lab at Johns Hopkins Univ. for inclusion in [1] was only identified as GrN2. We believe it is an natively unfolded phenylalanine-glycine (FG)-repeat [34].

⁹Courtesy of Alamy photos.

Moreover, there is experimental evidence that modification of protein polymers by means of phosphorylation [13, 25, 53, 55, 62] is involved in memory processes. It is not unreasonable to assume that phosphorylation processing corresponds to the removal and later replacement of tendrils on a cyclic substrate.

But is there reason to suspect that memory has any "structural" properties at all?

Perhaps the most important biological memory mechanism is our genetic memory which records the nature of our species. It is known to have a double helix structure which facilitates a near perfect recall. These coded sequences are subsequently "expressed" during development by an expansion process which might be similar to a non-random \mathcal{E} .

While the double helix facilitates a reliable read-only memory (ROM); the looser structure of chordless cycles appears to facilitate the encoding of eposidic information in a dynamic memory via a consolidation process such as \mathcal{R} .

Much of this section is speculation. But, both "closure" [44] and "chordless cycles" [45] would appear to have biological significance. The assertions of this paper have a solid mathematical base. As such, \mathcal{R} and \mathcal{E} provide useful examples within a category of *networks* that can formally model dynamic biological networks. If in addition, they actually model memory consolidation and recall as we suspect, that would be an additional bonus.

7 Appendix

Too much formal mathematics makes a paper hard to read. Yet, it is important to be able to check some of the statements made in the body of the paper. In this appendix we formally prove a representative selection of a few propositions tacitly found in the paper.

The order in which nodes, or more accurately the singleton subsets, of \mathcal{N} are encountered can alter which points are subsumed and subsequently deleted. Nevertheless, we show below that the reduced trace $\mathcal{T} = \mathcal{N}.\mathcal{R}$ will be unique, up to isomorphism.

Proposition 7.1 Let $\mathcal{T} = \mathcal{N}.\mathcal{R}$ and $\mathcal{T}' = \mathcal{N}.\mathcal{R}'$ be irreducible subsets of a finite network \mathcal{N} , then $\mathcal{T} \cong \mathcal{T}'$.

Proof: Let $y_0 \in \mathcal{T}$, $y_0 \notin \mathcal{T}'$. Then y_0 can be subsumed by some point y_1 in \mathcal{T}' and $y_1 \notin \mathcal{T}$ else because $y_0, \eta \subseteq y_1, \eta$ implies $y_0 \in \{y_1\}, \varphi$ and \mathcal{T} would not be irreducible.

Similarly, since $y_1 \in \mathcal{T}'$ and $y_1 \notin \mathcal{T}$, there exists $y_2 \in \mathcal{T}$ such that y_1 is subsumed by y_2 . So, $y_1.\eta \subseteq y_2.\eta$.

Now we have two possible cases; either $y_2 = y_0$, or not.

Suppose $y_2 = y_0$ (which is often the case), then $y_0.\eta \subseteq y_1.\eta$ and $y_1.\eta \subseteq y_2.\eta$ or $y_0.\eta = y_1.\eta$. Hence $i(y_0) = y_1$ is part of the desired isometry, i.

Now suppose $y_2 \neq y_0$. There exists $y_3 \neq y_1 \in \mathcal{T}'$ such that $y_2.\eta \subseteq y_3.\eta$, and so forth. Since \mathcal{T} is finite this construction must halt with some y_n . The points $\{y_0, y_1, y_2, \dots y_n\}$ constitute a complete graph Y_n with $\{y_i\}.\eta = Y_n.\eta$, for $i \in [0, n]$. In any reduction all $y_i \in Y_n$ reduce to a single point. All possibilities lead to mutually isomorphic maps.

In addition to $\mathcal{N}.\mathcal{R}$ being unique, we may observe that the transformation \mathcal{R} is functional because \mathcal{R} maps all subsets of N onto $N_{\mathcal{R}}$. So we can have $\{z\}.\mathcal{R} = \emptyset$, thus "deleting" z. Similarly, \mathcal{E} is functional because $\emptyset.\mathcal{E} = \{y\}$ provides for the inclusion of new elements. Both \mathcal{R} and \mathcal{E} are monotone, if we only modify its definition to be $X \subseteq Y$ implies $X.\mathcal{E} \subseteq Y.\mathcal{E}$, provided $X \neq \emptyset$.

The following proposition characterizes the structure of irreducible traces.

Proposition 7.2 Let \mathcal{N} be a finite symmetric network with $\mathcal{T} = \mathcal{N}.\mathcal{R}$ being its irreducible trace. If $y \in \mathcal{T}$ is not an isolated point then either

- (1) there exists a chordless k-cycle $C, k \geq 4$ such that $y \in C$, or
- (2) there exist chordless k-cycles C_1, C_2 each of length ≥ 4 with $x \in C_1$ $z \in C_2$ and y lies on a path from x to z.

Proof: (1) Let $y \in N_{\mathcal{T}}$. Since y is not isolated, we let $y = y_0$ with $y_1 \in y_0.\eta$, so $(y_0, y_1) \in E$. Since y_1 is not subsumed by y_0 , $\exists y_2 \in y_1.\eta, y_2 \notin y_0.\eta$, and since y_2 is not subsumed by y_1 , $\exists y_3 \in y_2.\eta$, $y_3 \notin y_1.\eta$. Since $y_2 \notin y_0.\eta$, $y_3 \neq y_0.\eta$.

Suppose $y_3 \in y_0.\eta$, then $< y_0, y_1, y_2, y_3, y_0 >$ constitutes a k-cycle $k \ge 4$, and we are done.

Suppose $y_3 \notin y_0.\eta$. We repeat the same path extension. $y_3.\eta \not\subseteq y_2.\eta$ implies $\exists y_4 \in y_3.\eta$, $y_4 \notin y_2.\eta$. If $y_4 \in y_0.\eta$ or $y_4 \in y_1.\eta$, we have the desired cycle. If not $\exists y_5,...$ and so forth. Because \mathcal{N} is finite, this path extension must terminate with $y_k \in y_i.\eta$, where $0 \leq i \leq n-3$, $n = |\mathcal{N}|$. Let $x = y_0, z = y_k$. (2) follows naturally.

Finally, we show that \mathcal{R} preserves the shortest paths between all elements of the trace, \mathcal{T} .

Proposition 7.3 Let $\sigma(x, z)$ denote a shortest path between x and z in \mathcal{N} . Then for all $y \neq x, z, \in \sigma(x, z)$, if y can be subsumed by y', then there exists a shortest path $\sigma'(x, z)$ through y'.

Proof: We may assume without loss of generality that y is adjacent to z in $\sigma(x, z)$. Let $\langle x, \ldots, x_n, y, z \rangle$ constitute $\sigma(x, z)$. If y is subsumed by y', then $y \cdot \eta = \{x_n, y, z\} \subseteq y' \cdot \eta$. So we have $\sigma'(x, z) = \langle x, \ldots, x_n, y', z \rangle$ of equal length. (Also proven in [42].)

In other words, z can be removed from \mathcal{N} with the certainty that if there was a path from some node x to z through y, there will still exist a path of equal length from x to z after y's removal.

Figure 13 visually illustrates the situation described in Proposition 7.3, which we call a **diamond**. There may, or may not, be a connection between y and y' as indicated by the dashed line. If there is, as assumed in Proposition 7.3, then either y' subsumes y or *vice* versa, depending on the order in which y and y' are encountered by \mathcal{R} . This provides one example of the isomorphism described in Proposition 7.1. If there is no connection between y and y' then we have two distinct paths between x and z of the same length.

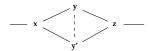


Figure 13: A network diamond

In the following proofs, σ denotes the spanning operator, not a shortest path.

Proposition 7.4 The spanning operator, σ is a closure operator over sets Y of cycles.

Proof: Readily, σ is expansive and monotone.

Let Y be a set of cycles $\{C_i\}$. Suppose $C_m \in Y.\sigma.\sigma$ implying that there exists some sequence $1 \le i \le k$ such that

$$C_m = C_1 \circ \ldots \circ C_i \circ \ldots \circ C_k \tag{3}$$

where $C_i \in Y.\sigma$, $1 \leq i \leq k$. Hence $C_i = C_{i_1} \circ \ldots \circ C_{i_n}$ where $C_{i_j} \in Y$. Thus, substituting into the sequence (3) for each i, we get $C_m = (C_{1_1} \circ \ldots \circ C_{1_n}) \circ (C_{2_1} \circ \ldots \circ C_{2_n})$ $\circ \ldots \circ (C_{k_1} \circ \ldots \circ C_{k_n})$ implying $C_m \in Y.\sigma$.

Proposition 7.5 Let C be a chordless cycle system and let σ be the spanning operator. The system (C, σ) satisfies the Steinitz-Maclane exchange axioms and is thus a matroid.

Proof: By Prop. 7.4, σ is a closure operator. Let $C_i, C_k \not\subseteq Y.\sigma$ where $Y = \{\dots, C_j, \dots\}$. Suppose $C_k \in (Y \cup C_i).\sigma$ implying that $C_k = C_i \circ (\dots C_j \dots) = C_i \circ C_m$ where $C_m \in Y.\sigma$. Consequently, since $C_i = C_k \circ C_m$ we have $C_i \in (Y \cup C_k).\sigma$.

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