

Transparallel processing by hyperstrings

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Human vision research aims at understanding the brain processes that enable us to see the world as a structured whole consisting of separate objects. To explain how humans organize a visual pattern, structural information theory starts from the idea that our visual system prefers the organization with the simplest descriptive code, that is, the code that captures a maximum of visual regularity. Empirically, structural information theory gained support from psychological data on a wide variety of perceptual phenomena, but theoretically, the computation of guaranteed simplest codes remained a troubling problem. Here, the graph-theoretical concept of “hyperstrings” is presented as a key to the solution of this problem. A hyperstring is a distributed data structure that allows a search for regularity in $O(2^N)$ strings as if only one string of length N were concerned. Thereby, hyperstrings enable transparallel processing, a previously uncharacterized form of processing that might also be a form of cognitive processing.

In the 1960s, Leeuwenberg (1) initiated structural information theory (SIT), which is a theory that aims at explaining how humans perceive visual patterns. A visual pattern can always be interpreted in many different ways, and SIT starts from the idea that the human visual system has a preference for the interpretation with the simplest descriptive code. In the 1950s, this idea had been proposed by Hochberg and McAlister (2), with an eye on Shannon’s work (3) as well as on early 20th century Gestalt psychology (ref. 4; see also ref. 5). To this idea, SIT adds a concrete visual coding language (see below), thus specifying the search space within which the simplest codes are to be found.

In interaction with empirical research, SIT developed into a competitive theory of visual structure. Leeuwenberg *et al.* (6–18) applied SIT to explain a variety of perceptual phenomena such as judged pattern complexity, pattern classification, neon effects, judged temporal order, assimilation and contrast, figure-ground organization, beauty, embeddedness, hierarchy, serial pattern segmentation and completion, and handedness. SIT started with a classification model, but nowadays it also contains comprehensive models of amodal completion (19, 20) and symmetry perception (21–24).

For object perception, SIT proposes an integration of viewpoint-independent and viewpoint-dependent factors quantified in terms of object complexities (19). A Bayesian translation of this integration, using precisions (i.e., probabilities $p = 2^{-c}$ derived from complexities c), suggests that fairly veridical vision in many worlds is a side effect of the preference for simplest interpretations (25). This idea, which challenges the traditional Helmholtzian idea that vision is highly veridical in only the one world in which we happen to live, is sustained by findings in the domain of algorithmic information theory (AIT), also known as the domain of Kolmogorov complexity or the domain of the minimal description-length (MDL) principle.

During the past 40 years, SIT and AIT showed similar developments. These developments, however, occurred in a different order, and until recently, SIT and AIT developed independently (see ref. 26 for an overview of AIT and ref. 25 for a comparison of SIT and AIT). Currently, noteworthy are the following two differences between SIT and AIT.

One difference applies to the complexity measurement. Unlike AIT, SIT takes account of the perceptually relevant distinction between structural and metrical information (27). For

example, the simplest codes of metrically different squares may have different algorithmic complexities in AIT but have the same structural complexity in SIT. By the same token, an AIT object class consists of objects with the same algorithmic complexity (ignoring structural differences), whereas an SIT object class consists of objects with the same structure (and hence with the same structural complexity) (25, 28). This might be a temporary difference, by the way. Since recently, AIT also seems to recognize the relevance of structures (29).

The other difference applies to the search space within which simplest codes are to be found. In both SIT and AIT, the simplest code of an object is to be obtained by “squeezing out” a maximum amount of regularity in a symbol string that represents a reconstruction recipe for the object; one might think of computer programs (binary strings) that produce certain output (an object). To formalize this idea, AIT did not focus on concrete coding languages that squeeze out specific regularities but instead provided (incomputable) definitions of randomness (30) to specify the result of squeezing out regularity. SIT, conversely, focused on a (computable) definition of “visual regularity,” which yielded a concrete coding language that squeezes out only transparent holographic regularities (for details, see ref. 31).

The transparent holographic character of these regularities has shown to be relevant in human symmetry perception (21–24). It also gave rise to the concept of “hyperstrings” that, in this article, is presented as a key to the computation of guaranteed simplest SIT codes. I begin by specifying the coding language of SIT and the related minimal-encoding problem.

SIT Coding Language

Basically, there are only three transparent holographic regularities, namely, iterations, symmetries, and alternations, which are described by, respectively, I-forms, S-forms, and A-forms (for short, ISA-forms), as given in the following definition of SIT’s coding language.

Definition 1: An SIT code \bar{X} of a string X is a string $t_1 t_2 \dots t_m$ such that $X = D(t_1) \dots D(t_m)$, where the decoding function $D: t \rightarrow D(t)$ takes one of the following forms:

I-form: $n^*(\bar{y}) \rightarrow yyy \dots y$ (n times y ; $n \geq 2$)

S-form: $S[(\bar{x}_1)(\bar{x}_2) \dots (\bar{x}_n), (\bar{p})] \rightarrow x_1 x_2 \dots x_n p x_n \dots x_2 x_1$ ($n \geq 1$)

A-form: $\langle (\bar{y}) \rangle / \langle (\bar{x}_1)(\bar{x}_2) \dots (\bar{x}_n) \rangle \rightarrow yx_1 yx_2 \dots yx_n$ ($n \geq 2$)

A-form: $\langle (\bar{x}_1)(\bar{x}_2) \dots (\bar{x}_n) \rangle / \langle (\bar{y}) \rangle \rightarrow x_1 y x_2 y \dots x_n y$ ($n \geq 2$)

Otherwise: $D(t) = t$ [1]

for strings y , p , and x_i ($i = 1, 2, \dots, n$). The code parts (\bar{y}) , (\bar{p}) , and (\bar{x}_i) are called “chunks”; the chunk (\bar{y}) in an I-form or A-form is called a “repeat”; the chunk (\bar{p}) in an S-form is called a “pivot,” which, as a limit case, may be empty; the chunk string $(\bar{x}_1)(\bar{x}_2) \dots (\bar{x}_n)$ in an S-form is called an “S-argument” consisting

Abbreviations: SIT, structural information theory; AIT, algorithmic information theory; MDL, minimal description length; I, iteration; S, symmetry; A, alternation; SPM, shortest-path method.

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of “S-chunks” (\bar{x}_i); and the chunk string ($\bar{x}_1\bar{x}_2\dots\bar{x}_n$) in an A-form is called an “A-argument” consisting of “A-chunks” (\bar{x}_i).

Hence, an SIT code may involve not only encodings of strings inside chunks [that is, from (y) into (\bar{y})] but also hierarchically recursive encodings of S-arguments or A-arguments ($\bar{x}_1\bar{x}_2\dots\bar{x}_n$) into ($\bar{x}_1\bar{x}_2\dots\bar{x}_n$). As I specify in the next section, this hierarchically recursive search for regularity creates the problem that, to compute simplest SIT codes, a superexponential amount of time seems to be required (see also ref. 32). The following sample of SIT codes of one and the same symbol string may give a gist of this problem.

- String: $X = abacadacbabacdacdad$
 Code 1: $\bar{X} = a b 2^*(acd) S[(a)(b), (a)] 2^*(cda) b$
 Code 2: $\bar{X} = \langle(aba)\rangle/\langle(cdacd)(bacdadacdad)\rangle$
 Code 3: $\bar{X} = \langle(S[(a), (b)])\rangle/\langle(S[(cd), (a)])(S[(b)(a)(cd), (a)])\rangle$
 Code 4: $\bar{X} = S[(ab)(acd)(acd)(ab)]$
 Code 5: $\bar{X} = S[S[(ab)((ab))((acd))]]$
 Code 6: $\bar{X} = 2^*(\langle(a)\rangle/\langle(b)(cd)(cd)(b)\rangle)$
 Code 7: $\bar{X} = 2^*(\langle(a)\rangle/\langle S[(b)((cd))]\rangle)$

[2]

Code 1 is a code with six code terms, namely, one S-form, two I-forms, and three symbols. Code 2 is an A-form with chunks containing strings that may be encoded as given in code 3. Code 4 is an S-form with an empty pivot and illustrates that, in general, S-forms describe broken symmetry (33); mirror symmetry then is the limit case in which every S-chunk contains only one symbol. Code 5 gives a hierarchically recursive encoding of the S-argument in code 4. Code 6 is an I-form with a repeat that has been encoded into an A-form with an A-argument that, in code 7, has been encoded hierarchically recursively into an S-form.

SIT’s Minimal-Encoding Problem

As said, the coding language of SIT specifies the search space within which simplest codes are to be found. To search this space for simplest codes, one of course needs a measure of code complexity, but this is a subordinate problem in this article. SIT has known complexity measures that were either empirically supported or theoretically plausible (28), but since about 1990, SIT uses a measure that is both (ref. 15; see also ref. 25). For any complexity measure, however, the question is whether one can ever be sure that a given code is indeed a simplest code. In other words, the fundamental problem of computing guaranteed simplest codes is to take account of all possible codes of a given string.

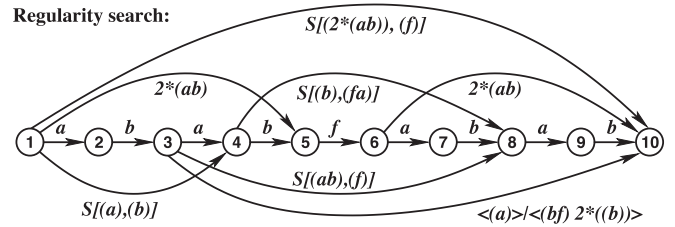
It is expedient to note that the SIT minimal-encoding problem differs from context-free grammar (CFG) problems such as finding the smallest CFG for any given string, for which fast approximation algorithms exist (e.g., see refs. 34 and 35). SIT starts from a particular CFG, namely, the coding language given in Definition 1, which was designed specifically to capture perceptually relevant structures in strings. The minimal-encoding problem of SIT then is to compute, for any given string, a guaranteed simplest code (i.e., no approximation) by means of the specific coding rules supplied by this perceptual coding language.

A part of SIT’s minimal-encoding problem can be solved as follows by means of Dijkstra’s (36) shortest-path method (SPM). Suppose that for every substring of a string of length N , one already has computed a simplest covering ISA-form, that is, a simplest substring code among those that consist of only one ISA-form. Then, Dijkstra’s $O(N^2)$ SPM can be applied to select a simplest code for the entire string from among the $O(2^N)$ codes that then still are possible (see Fig. 1; see ref. 37 for details on this application).

String:



Regularity search:



Shortest path method: $S[(2^*(ab)), (f)]$

Fig. 1. Suppose for the string $\mathcal{P} = ababfabab$ that the regularity search has yielded simplest covering ISA-forms for the substrings of \mathcal{P} (only a few of these ISA-forms are shown). Then, the SPM yields $S[(2^*(ab)), (f)]$ as the simplest code of \mathcal{P} . (Note: the number of string symbols in a code is taken to quantify its structural complexity.)

This, however, leaves open the much harder part of computing simplest covering ISA-forms for every substring. In general, as one may infer from Definition 1, a substring of length k can be encoded into $O(2^k)$ covering S-forms and $O(k2^k)$ covering A-forms. To pinpoint a simplest covering S-form or A-form, a simplest code for every one of the $O(2^k)$ S-arguments and $O(k2^k)$ A-arguments has to be computed as well, and so on, with $O(\log N)$ recursion steps. Hence, an algorithm that would process each and every S-argument and A-argument separately would require a superexponential $O(2^{N \log N})$ amount of computing time.

In the next sections, I show that the concept of hyperstrings provides a key to the solution of this daunting problem. First, I define and illustrate hyperstrings in a graph-theoretical setting (for an extensive course on graph theory, see ref. 38; for a brief course, see Appendix 1, which is published as supporting information on the PNAS web site). Then, I show that A-arguments and S-arguments group by nature into hyperstrings. Finally, I evaluate the fact that hyperstrings allow for what I call “transparent processing,” that is, they allow a search for regularity in $O(2^N)$ A-arguments or S-arguments as if only one A-argument or S-argument of length N were concerned.

Hyperstrings

The concept of hyperstrings is a generalization of the concept of strings. To identify context-free string structures, the only usable property is the identity of substrings. As I specify next, a generalization of this property holds for the hypersubstrings of a hyperstring.

Definition 2: A hyperstring is a simple semi-Hamiltonian directed acyclic graph (V, E) with a labeling of the edges in E such that for all vertices $i, j, p, q \in V$:

$$\text{either } \pi(i, j) = \pi(p, q) \text{ or } \pi(i, j) \cap \pi(p, q) = \emptyset, \quad [3]$$

where a substring set $\pi(v_1, v_2)$ is the set of label strings represented by the paths (v_1, \dots, v_2) in an edge-labeled directed acyclic graph. In a hyperstring, the subgraph formed by the vertices and edges in these paths (v_1, \dots, v_2) is called a “hypersubstring.”

It can easily be verified that a hyperstring is an st-digraph (i.e., a directed acyclic graph with only one source and only one sink) with only one Hamiltonian path from source to sink (i.e., a path that visits every vertex only once). The label string represented by this Hamiltonian path is what I call the “kernel” of the hyperstring.

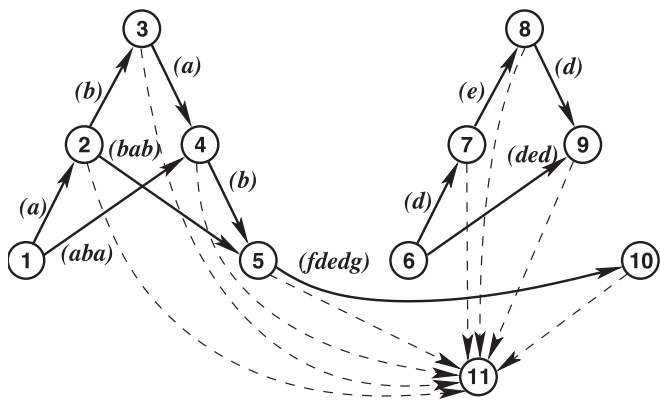


Fig. 5. The S-graph $\mathcal{S}(T)$ for the string $T = ababfdedgpfdedgbaba$, with two independent subgraphs. Dashed edges and bold edges represent pivots and S-chunks, respectively, in S-forms covering diafixes of T .

every substring $x_i \dots x_n p x_n \dots x_i$ of T can be covered by an S-form $S[(x_i \dots (x_{j-1}), (x_j \dots x_n p x_n \dots x_i)]$. The substrings $x_i \dots x_n p x_n \dots x_i$ are all centered around the midpoint of T and form what I next define to be “diafixes.” The notion of diafixes is convenient in the subsequent elaboration of how S-arguments group into hyperstrings.

Definition 4: A diafix of a string $T = s_1 s_2 \dots s_N$ is a substring $s_{i+1} \dots s_{N-i}$ ($0 \leq i < N/2$).

Definition 5: For a string $T = s_1 s_2 \dots s_N$, the S-graph $\mathcal{S}(T)$ is a simple directed acyclic graph (V, E) with $V = \{1, 2, \dots, \lfloor N/2 \rfloor + 2\}$ and, for all $1 \leq i < j < \lfloor N/2 \rfloor + 2$, edges (i, j) and $(j, \lfloor N/2 \rfloor + 2)$ labeled with, respectively, the chunk $(s_i \dots s_{j-1})$ and the possibly empty chunk $(s_j \dots s_{N-j+1})$ if and only if $s_i \dots s_{j-1} = s_{N-j+2} \dots s_{N-i+1}$.

Fig. 5 illustrates that an S-graph $\mathcal{S}(T)$ may contain several independent subgraphs and that every S-form covering a diafix of T is represented by a path in $\mathcal{S}(T)$. For instance, the path $(2, 5, 10, 11)$ represents the S-form $S[(bab)(fdedg), (p)]$ covering the diafix $babfdedgpfdedgbab$. Thus, in Definition 5, the edges $(j, \lfloor N/2 \rfloor + 2)$ represent all possible pivots in such S-forms, and the edges (i, j) represent all possible S-chunks in such S-forms.

Hence, without its pivot edges, an S-graph $\mathcal{S}(T)$ represents the S-arguments of all S-forms covering diafixes of T . As I establish next, these S-arguments group into hyperstrings.

Theorem 2. The S-graph $\mathcal{S}(T)$ for a string $T = s_1 s_2 \dots s_N$ consists of at most $\lfloor N/2 \rfloor + 2$ disconnected vertices and at most $\lfloor N/4 \rfloor$ independent subgraphs that, without the sink vertex $\lfloor N/2 \rfloor + 2$ and its incoming pivot edges, form one disconnected hyperstring each.

Proof: See Appendix 2.

For instance, Fig. 6 shows two S-graphs $\mathcal{S}(T_1)$ and $\mathcal{S}(T_2)$, which each consist of one independent subgraph that, without the pivot edges, forms a hyperstring. Although in Fig. 6 the two strings T_1 and T_2 are near identical, the substring sets $\pi(1, 5)$ and $\pi(6, 10)$ are identical for T_1 but disjoint for T_2 . This illustrates the crucial hyperstring property that substring sets are either completely identical or completely disjoint (see Definition 2).

The Computability of Simplest SIT Codes

Within AIT, guaranteed minimal encoding is not feasible, because without a definition of regularity, one can never be sure that one has extracted a maximum of regularity. Within SIT, regularity is defined as being constituted by transparent holographic configurations, but even then, guaranteed minimal encoding did not seem feasible: The required hierarchically recursive search for regularity seemed to imply that every

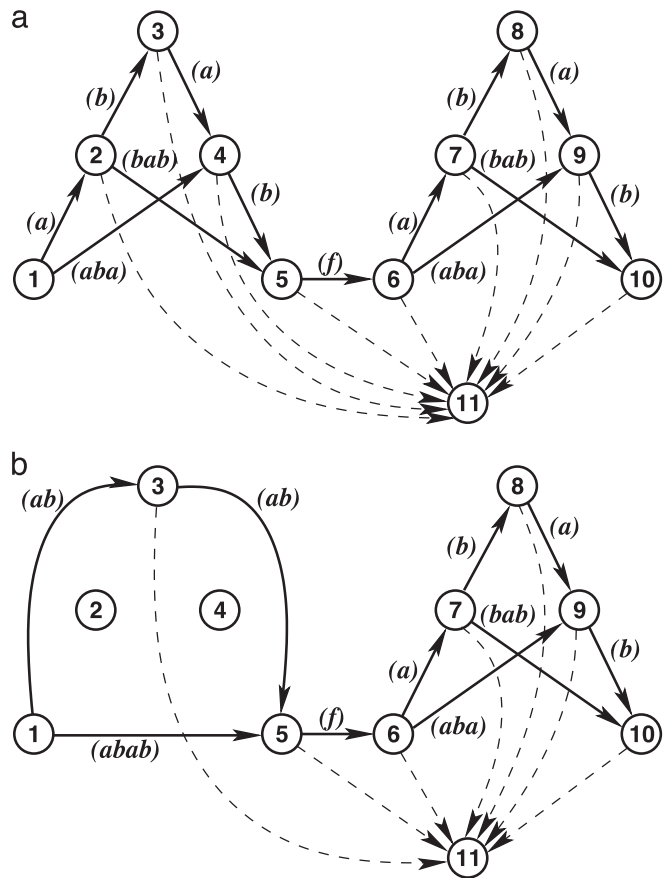


Fig. 6. (a) The S-graph for the string $T_1 = ababfababgbabafabab$, with, among others, identical substring sets $\pi(1, 5)$ and $\pi(6, 10)$. (b) The S-graph for the nearly identical string $T_2 = ababfababgbabafabab$, in which the substring sets $\pi(1, 5)$ and $\pi(6, 10)$ are disjoint.

S-argument and A-argument has to be processed separately, which would consume superexponential computing time.

The previous two sections, however, show that S-arguments and A-arguments group by nature into hyperstrings. More specifically, A-arguments group into independent hyperstrings, and S-arguments group into disconnected hyperstrings. As I discuss next, this paves the way for an encoding algorithm that determines guaranteed simplest SIT codes.

My quest for such an algorithm started in the mid-1980s, by developing an approximation algorithm (37). In the mid-1990s, I completed an algorithm that already used S-graphs and A-graphs without pseudo A-chunk edges (31). This algorithm is available on request; it determines guaranteed simplest SIT codes under the restriction that pseudo A-chunks do not occur (at the time, the need to include pseudo A-chunks was not yet evident). The upcoming upgrade (see below) removes this restriction by using hyperstrings as defined in this article. For the rest, the available algorithm deals with the hierarchically recursive search for regularity in A-arguments and S-arguments as outlined in the next overview of the upgrade.

Step 1: The search for simplest covering ISA-forms for the substrings of a string \mathcal{P} of length N can be embedded in an $O(N^3)$ all-pairs SPM (see ref. 39) that selects a simplest SIT code for every substring of \mathcal{P} , proceeding from small to large substrings, the largest substring being the entire string \mathcal{P} . That is, a simplest covering ISA-form for some substring of \mathcal{P} has to be computed only once all smaller substrings of \mathcal{P} have already been assigned a simplest code. This implies that the search for simplest covering

I-forms is “peanuts,” and that the hierarchically recursive search for regularity in A-arguments and S-arguments can start from A-forms $\langle(\bar{y})\rangle/\langle(\bar{x}_1)(\bar{x}_2)\dots(\bar{x}_m)\rangle$ and S-forms $S[(\bar{x}_1)(\bar{x}_2)\dots(\bar{x}_m), (\bar{p})]$ with simplest codes \bar{y} , \bar{p} , and \bar{x}_i for the smaller substrings inside the chunks.

Step 2: An A-graph $\mathcal{A}(T)$ or an S-graph $\mathcal{S}(T)$ for a substring $T = s_1s_2\dots s_M$ of \mathcal{P} represents $O(2^M)$ individual chunk strings, but it can be constructed in only $O(M^2)$ computing steps. For $\mathcal{A}(T)$, one only has to check for every substring of T whether this substring and the subsequent suffix have identical prefixes (see *Definition 3*). [In an $O(N^2)$ preprocess, every substring of \mathcal{P} can be assigned an integer that is the same for all and only all identical substrings.] Similarly, for $\mathcal{S}(T)$, one only has to check for every substring $s_i\dots s_{j-1}$ in the left-hand half of T whether it is identical to its symmetrical counterpart $s_{M-j+2}\dots s_{M-i+1}$ in the right-hand half of T (see *Definition 5*). Every edge in $\mathcal{A}(T)$ and $\mathcal{S}(T)$ can be given a complexity on the basis of the already processed content of the represented A-chunk, S-chunk, or pivot (see *Step 1*). At this point, pseudo A-chunks can be given an “infinite” complexity to prevent them from ending up as real A-chunks.

Step 3: Hierarchically recursively, the hypersubstrings in $\mathcal{A}(T)$ and $\mathcal{S}(T)$ can be processed starting with *Step 1*. That is, by *Definition 2*, a hyperstring with kernel length n can be conceived of as one string $\mathcal{H} = h_1h_2\dots h_n$ in which a substring $h_i\dots h_j$ stands for the substring set $\pi(i, j + 1)$ in the hyperstring. For instance, the hyperstring in Fig. 2 can be conceived of as a string $\mathcal{H} = h_1h_2\dots h_8$ in which, among others, substrings $h_1\dots h_3$ and $h_5\dots h_7$ are identical. This identity stands for the identity of the substring sets $\pi(1, 4)$ and $\pi(5, 8)$, which in one go captures the abc , ay , and xc identities in strings represented in the hyperstring. Inversely, one such substring identity already implies that the substring sets are identical. Hence, the ISA-forms in this single string \mathcal{H} account for all ISA-forms in all strings represented in the hyperstring, so this single string \mathcal{H} can be taken as the input of *Step 1*. In other words, *Step 1* can take the hyperstring as if it were only one string (namely, the hyperstring kernel) with, for various substrings, various *a priori* given alternatives that all but one will be dismissed during the all-pairs SPM (see Fig. 7).

Step 4: The argument of an A-form or S-form in a substring T of \mathcal{P} contains at most $\lfloor N/2 \rfloor$ chunks, so the recursion depth is at most $\log_2 N$. The recursion ends when *Step 2* yields A-graphs and S-graphs without identical substring sets. For these graphs and, backtracking, eventually for $\mathcal{A}(T)$ and $\mathcal{S}(T)$, the all-pairs SPM selects a simplest code for every hypersubstring, and by including repeats and pivots, respectively, it selects simplest covering A-forms and S-forms for, eventually, all suffixes and diafixes of T . These A-forms and S-forms are used in the all-pairs SPM that eventually selects a simplest code for \mathcal{P} .

As indicated in *Step 4*, the all-pairs SPM yields eventually simplest covering A-forms and S-forms for all suffixes and diafixes of T . Therefore, each pass, *Step 2* has to yield only $O(N)$ A-graph and S-graphs. Hence, *Step 2* yields $O(N^{\log N})$ such graphs during the entire hierarchically recursive search for regularity. Furthermore, because of the all-pairs SPM, the algorithm requires $O(N^3)$ computing steps per A-graph or S-graph. Hence, in total, it requires $O(N^{3+\log N})$ steps to compute a simplest SIT code for a string of length N .

It is true that this still implies a weakly exponential computing time, but it contrasts, in any case, very favorably with the uncomputability of a simplest AIT code and with the superexponential computing time that beforehand seemed to be required to compute a simplest SIT code. Furthermore, the weakly exponential factor $\log N$ is due to the number of hierarchical recursion steps in the worst case. Only few strings have a deep hierarchical structure, so in the average case, this weakly exponential factor hardly seems a problem.

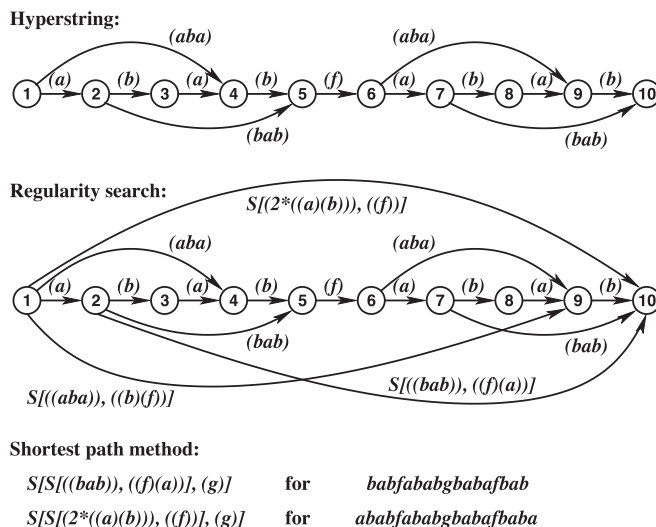


Fig. 7. For the hyperstring in the S-graph $\mathcal{S}(T_1)$ from Fig. 6a, with $T_1 = ababfababgbabafbaba$, the hierarchically recursive regularity search yields simplest covering ISA-forms for the hypersubstrings (only a few of these ISA-forms are shown). By including the pivots, the all-pairs SPM then yields simplest covering S-forms for the diafixes of T_1 (only a few of these S-forms are shown). Note that the number of string symbols in a code is taken to quantify its structural complexity, and for clarity the substrings *aba* and *bab* inside chunks are shown un-coded but should be read as $S[(a), (b)]$ and $S[(b), (a)]$, respectively.

Be that as it may, the central issue in this article is not this algorithm as a whole but rather the role of hyperstrings in it. As outlined in *Step 3*, hyperstrings imply that $O(2^N)$ S-arguments or A-arguments do not have to be searched serially or in parallel for regularity but can be processed as if only one S-argument or A-argument were concerned. I propose to call this a form of transparallel processing, which may be qualified as follows.

Transparallel Processing

To be clear, I do not use the term “transparallel” in the sense that Nelson (40) proposed it in the 1960s following Bush’s idea (41) to display related items in correspondence with the way humans think. Nelson used it in “transparallel displays,” which means so much as that connected items are shown together with their connections. Nowadays, such data structures are better known as “distributed representations,” and this is the term I use here as a leg up to what I call transparallel processing. To Nelson, a typical example of a distributed representation is a display showing two related stories side by side, with visible links between the corresponding parts. A more everyday example is a road map in which routes between places are not displayed separately but such that common parts are effectively displayed as common parts. Likewise, hyperstrings are distributed representations of strings.

A process that effectively exploits a distributed representation of items can be said to perform “distributed processing.” This is often taken to mean that the process is distributed over many processors, but here I take it to mean primarily that the items are processed simultaneously in the sense that every common part is processed only once. Different common parts can then be processed serially by one processor or in parallel by many processors. A classical example of serial distributed processing is Dijkstra’s SPM (ref. 36; see also Fig. 1). Generally, such a one-processor implementation can be converted into a many-processors implementation performing parallel distributed processing (see *Appendix 3*, which is published as supporting information on the PNAS web site).

Nowadays, distributed processing is a standard in many applications in computer science and in many models in cognitive science. For instance, in computer science, a deterministic finite automaton (DFA) is a distributed representation of the sentences in a regular language, which enables a quick serial distributed processing check on whether a given string is a sentence in this language (42). Furthermore, in cognitive science, distributed representations called networks are used in parallel distributed processing models of cognitive processes that, given certain input, select quickly a best matching item from among the items represented in the network (43). The items in the network could be words to be recognized in written or spoken language, for instance.

In the minimal-encoding algorithm outlined in the previous section, hyperstrings are subjected to all-pairs SPMs and, thereby, to distributed processing. In this respect, hyperstrings do not differ from deterministic finite automata and networks: These data structures all allow stored items to be processed simultaneously in the sense that every common part is processed only once. Hyperstrings, however, allow in addition for what I call transparallel processing, which, as I specify next, goes one step beyond distributed processing.

First, as said, in distributed processing, different common parts still have to be processed serially by one processor or in parallel by many processors. Second, in the minimal-encoding

algorithm, a hyperstring is subjected not only to an all-pairs SPM but also to a search for regularity in the strings represented in the hyperstring. During this regularity search, different common parts do not have to be processed serially by one processor or in parallel by many processors but can, by one processor, be processed as if only one part were concerned. As outlined in *Step 3* in the previous section, this form of processing is due to the hyperstring property that substring sets are either completely identical or completely disjoint (see *Definition 2*), and this is the form of processing I call transparallel processing.

Conclusions

In cognitive science, our brain is typically supposed to be attuned to relevant regularities in the world. Symmetry, for instance, is doubtlessly relevant: It is a regularity that is visible in the shape of virtually every living organism. In this article I showed that such visually relevant regularities lend themselves for transparallel processing. Hence, if our brain is indeed attuned to relevant regularities, then, just as distributed processing, transparallel processing might well be a form of cognitive processing.

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