

Oritatami systems assemble shapes no less complex than tile assembly model (aTAM)

Daria Pchelina (Дарья С Пчелина)

LIPN, Institut Galilée – Université Paris 13, France

Nicolas Schabanel

École Normale Supérieure de Lyon (LIP UMR5668 and IXXI, MC2), France

Shinnosuke Seki (関 新之助)

University of Electro-Communications, 1-5-1 Chofugaoka, Chofu, Tokyo 1828585, Japan

Guillaume Theyssier

Aix-Marseille Université, CNRS, I2M, Marseille, France

Abstract

Different models have been proposed to understand natural phenomena at the molecular scale from a computational point of view. Oritatami systems are a model of molecular co-transcriptional folding: the transcript (the “molecule”) folds as it is synthesized according to a local energy optimisation process, in a similar way to how actual biomolecules such as RNA fold into complex shapes and functions. We introduce a new model, called *turedo*, which is a self-avoiding Turing machine on the plane that evolves by marking visited positions and that can only move to unmarked positions. Any oritatami can be seen as a particular turedo. We show that any turedo with lookup radius 1 can conversely be simulated by an oritatami, using a universal bead type set. Our notion of simulation is strong enough to preserve the geometrical and dynamical features of these models up to a constant spatio-temporal rescaling (as in intrinsic simulation). As a consequence, turedo can be used as a readable oritatami “higher-level” programming language to build readily oritatami “smart robots”, using our explicit simulation result as a compiler.

As an application of our simulation result, we prove two new complexity results on the (infinite) limit configurations of oritatami systems (and radius-1 turedos), assembled from a finite seed configuration. First, we show that such limit configurations can embed any recursively enumerable set, and are thus exactly as complex as aTAM limit configurations. Second, we characterize the possible densities of occupied positions in such limit configurations: they are exactly the Π_2 -computable numbers between 0 and 1. We also show that all such limit densities can be produced by one single oritatami system, just by changing the finite seed configuration.

None of these results is implied by previous constructions of oritatami embedding tag systems or 1D cellular automata, which produce only computable limit configurations with constrained density.

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Supplementary Material Radius-1 Turedo to Delay-3 Oritatami compiler [23]: <https://hub.darcs.net/turedo2oritatami/turedo2oritatami/>; Oritatami simulator [22]: <http://perso.ens-lyon.fr/nicolas.schabanel/OSsimulator>.

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1 Introduction

A major trend of natural computing is the study of computational models inspired by molecular biology that are both theoretically rich and realistic enough to allow *in-vitro* implementations. Oritatami systems were introduced in [9, 10] to investigate the computational power of molecular co-transcriptional folding, in which an RNA sequence (transcript) folds upon itself into an intricate structure while being synthesized (transcribed). This phenomenon has proven programmable *in-vitro* by [12], in which Geary, Rothmund, and Andersen demonstrated how to encode a rectangular tile-like structure in a transcript and its folding pathway so that this transcript folds cotranscriptionally along the pathway into the encoded structure. This *RNA Origami* architecture has recently been highly automated by their software ROAD (RNA Origami Automated Design) [8]. ROAD extends the scale and functional diversity of RNA scaffolds, and is thus a promising direction for the design of RNA-based computation. DNA tile self-assembly did rely on the cellular automata theory to build up the abstract Tile Assembly model (aTAM) [24] which in turn allowed to develop experimental settings simple enough to be implemented *in vitro*, such as the Sierpinski triangle [21]. On the opposite, RNA origami was born first *in-vitro* and the oritatami system was created [11] to answer the lack of theoretical framework to design computations for cotranscriptional-based assembly systems. In this paper, we introduce the *turedo* model, implementable in oritatami, which, as opposed to oritatami, is simple enough to program, to wish for a design equivalent to the Sierpinski triangle experiment for cotranscription-based *in-vitro* systems.

An oritatami system consists of a “molecule” (the *transcript*) made of “beads” that attract each other. The molecule grows by one bead per step and, at each step, the δ most recently produced beads are free to move around to look for the position that maximizes the number of bonds they can make with each other (hence the folding is co-transcriptional). This process ends up self-assembling a shape incrementally. It is known from [11, 20] that oritatami systems are Turing universal. They can also build arbitrary shapes [4] modulo a small universal constant upscaling, as well as specific fractals [17]. However, oritatami systems remain notably challenging to design. Indeed, the only shapes that can be built by [11, 20] are space-time diagrams of cyclic tag-systems or 1D cellular automata; and [4] requires to hardcode the whole shape in the transcript. The new computational model introduced in this article (*turedo*) not only abstracts away the technical details of attraction rules and bead sequence of oritatami, but embraces the geometrical aspects of them, as opposed to the simulation of classical one-dimensional computational models. We demonstrate that *turedos* can be simulated up to upscaling by oritatami systems. Our simulation allows thus to take full advantage of *turedo* computations in building shapes, and can be used as a compiler to design powerful oritatami systems as demonstrated below.

Oritatami systems and Turedos. The classical model of Turing machines has already been considered in other settings than the one dimensional bi-infinite tape, in particular in higher dimensions [1]. A popular class of Turing machines in \mathbb{Z}^2 is that of turmites [16], which are free to move on the plane but do it by just looking at their current internal state and the tape content at their current position. In this paper, we introduce a somewhat orthogonal class of Turing machines on the plane, that we call *turedos*¹, which can look at the tape

¹ Inspired by the nicely coined terminology for turmites, as a reference to *toredo navalis* (shipworms) that would only grow self-avoiding tunnels in wood if they were infinite.

89 content around their position to decide their move (like in [1]), but are constrained to move
90 only in a self-avoiding way.

91 Both our models (oritatami and turedos) have two strong constraints: they are sequential
92 and self-avoiding (*i.e.* each position of the plane can only be visited once and becomes an
93 obstruction for future moves). They can be seen as the sequential counterpart of aTAM model
94 of self-assembly [19, 5] or freezing cellular automata [13, 2, 18]. But they are not just finite
95 state automata growing a self-avoiding path in a regular way. Their computational power is
96 in their ability to make moves depending on the configuration of neighboring positions.

97 Our main result is that oritatami can simulate turedos of lookup radius 1. Our notion
98 of simulation is strong enough to preserve the geometrical and dynamical features of these
99 models up to a constant spatio-temporal rescaling: the oritatami reproduces the whole
100 dynamics of the turedo using macro-cells and a constant spatio-temporal rescaling. This
101 definition is similar to intrinsic simulations developed for cellular automata [3] or self-assembly
102 tilings [5]. Theorem 1.1 is proved in section 3.

103 ► **Theorem 1.1** (Main result 1). *There is a universal bead type set \mathcal{B} such that for any*
104 *turedo \mathcal{T} of radius 1 with alphabet of size Q , there is a delay-3 oritatami system based on \mathcal{B}*
105 *with period $\Lambda = \Theta(Q^6 \log Q)$ which simulates intrinsically \mathcal{T} at space-scale $\Theta(Q^3 \sqrt{\log Q})$ and*
106 *time-scale Λ .*

107 **Complexity of limit configurations.** The Turing universality results in [11, 20] induce
108 undecidability results of the form: given an oritatami, a seed and a position, determining
109 whether the position will be visited is undecidable. However, these embeddings are such
110 that the obtained limit configurations are always computable because the space-time of the
111 simulated tag system (or cellular automaton) computation is progressively constructed in a
112 predictable way in a fixed region of oritatami's space. Precisely, in any limit configuration
113 c^∞ obtained this way, the map $z \mapsto c^\infty(z)$ is computable because there is a computable time
114 bound $\tau(z)$ such that if position z is not visited after $\tau(z)$ steps of the run, then it will never
115 be visited (see Lemma 4.1).

116 The first application of our simulation result is to prove that we can produce uncomputable
117 limit configurations from finite seeds with oritatami (section 4). This implies that there are
118 oritatami runs from finite seeds where there is no computable time bound $\tau(z)$ on the visit
119 time of position z .

120 Results on uncomputable limit configurations were already obtained in the model of
121 directed aTAM [15]. Nevertheless, the construction used takes full advantage of the massive
122 parallelism allowed in the aTAM model and *cannot* be translated into the turedo settings. Our
123 construction is actually simpler than that of [15] and shows that sequential self-avoiding models
124 can organize information in the plane in such a way that some regions allow 'uncomputable
125 comebacks'.

126 ► **Theorem 1.2** (Main result 2). *There exists a fixed oritatami with delay 3 and a fixed finite*
127 *seed σ such that the produced limit configuration c_σ^∞ is uncomputable as a map.*

128 The second application of our simulation result is about (upper) density of occupied
129 positions in the limit configurations obtained from finite seeds. Density is a natural geometrical
130 parameter to test the ability of our models to produce complex infinite self-avoiding paths from
131 finite seeds. We show that such densities are exactly the Π_2 -computable numbers between 0
132 and 1 (Theorem 5.3), where Π_2 -computable means being the limsup of a computable sequence
133 of rational numbers [25]. In particular turedos and oritatami can produce limit densities
134 which are not recursively approximable (*i.e.* not the limit of any computable sequence of

135 rational numbers). We actually show that the whole spectrum of density can be obtained
 136 in a single turedo by varying the seed (Theorem 5.3). Using our simulation framework, the
 137 following result is shown first for turedos and then for oritatami in section 5.

138 ► **Theorem 1.3** (Main result 3). *For any $\epsilon > 0$, there exists an oritatami of delay 3 such
 139 that for any Π_2 -computable number $d \in [0, 1 - \epsilon]$, there is a finite seed σ such that the limit
 140 configuration c_σ^∞ reached from it has density of occupied positions exactly d .*

141 Note that the densities that can be produced in the (directed) aTAM model or freezing
 142 cellular automata from finite initial configurations cannot be more complex (see Lemma 5.1).

143 The organization of the paper is as follows: we first present oritatami and turedo models
 144 and the notion of simulation (section 2); then, we establish our main simulation result
 145 (section 3) and its two applications (sections 4 and 5).

146 2 Definitions and Models

147 **Oritatami systems.** Oritatami systems are embedded in the triangular lattice $\mathbb{T} = (\mathbb{Z}^2, \sim)$,
 148 where $(x, y) \sim (u, v)$ if and only if $(u, v) \in \cup_{\epsilon=\pm 1} \{(x + \epsilon, y), (x, y + \epsilon), (x + \epsilon, y + \epsilon)\}$. Every
 149 position (x, y) in \mathbb{T} is mapped in the euclidean plane to $x \cdot \vec{e} + y \cdot \vec{sw}$ using the vector
 150 basis $\vec{e} = (1, 0)$ and $\vec{sw} = \text{RotateClockwise}(\vec{e}, 120^\circ) = (-\frac{1}{2}, -\frac{\sqrt{3}}{2})$. We will denote by
 151 $\vec{nw}, \vec{ne}, \vec{e}, \vec{se}, \vec{w}, \vec{sw}$ the six canonical unit vectors in \mathbb{T} . Let B be a finite set of *bead types*. A
 152 *configuration* c of a bead type sequence $p \in B^* \cup B^\mathbb{N}$ is a directed self-avoiding path $c_0c_1c_2\cdots$
 153 in \mathbb{T} , where for all integer i , the vertex c_i of c is labeled by p_i and refers to the *position* in \mathbb{T}
 154 of the $(i + 1)$ -th bead in the configuration. A *partial configuration* of p is a configuration of
 155 a prefix of p .

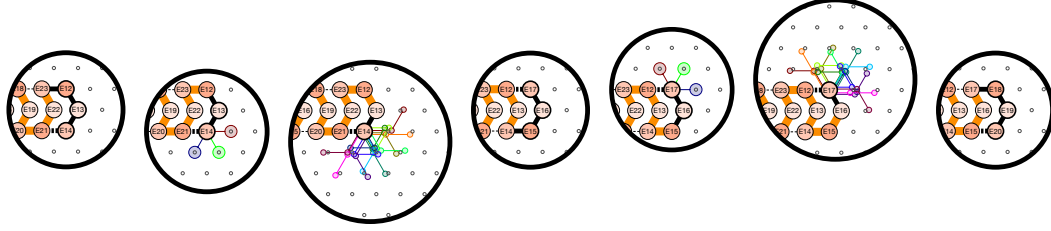
156 For any partial configuration c of some sequence p , an *elongation* of c by k beads (or
 157 *k-elongation*) is a partial configuration of p of length $|c| + k$ extending by k positions the
 158 self-avoiding path of c . We denote by \mathcal{C}_p the set of all partial configurations of p (the index
 159 p will be omitted whenever it is clear from the context). We denote by $c^{\triangleright k}$ the set of all
 160 k -elongations of a partial configuration c of sequence p .

161 An *oritatami system* $\mathcal{O} = (p, \heartsuit, \delta)$ is composed of (1) a (possibly infinite) bead type
 162 sequence p , called the *transcript*, (2) an *attraction rule*, which is a symmetric relation $\heartsuit \subseteq B^2$,
 163 and (3) a parameter δ called the *delay*. \mathcal{O} is said to be *periodic* if p is infinite and periodic.
 164 Periodicity ensures that the “program” p embedded in the oritatami system is finite (does not
 165 hardcode unbounded behavior) and at the same time allows arbitrarily long computation.

166 We say that two bead types a and b *attract* each other when $a \heartsuit b$. Furthermore, given
 167 a (partial) configuration c of a bead type sequence q , we say that there is a *bond* between
 168 two adjacent positions c_i and c_j of c in \mathbb{T} if $q_i \heartsuit q_j$ and $|i - j| > 1$. The *number of bonds* of
 169 configuration c of q is denoted by $H(c) = |\{(i, j) : c_i \sim c_j, j > i + 1, \text{ and } q_i \heartsuit q_j\}|$.

170 **Oritatami dynamics.** The folding of an oritatami system is controlled by the delay δ .
 171 Informally, the configuration grows from a *seed configuration* (the input), one bead at a time.
 172 This new bead adopts the position(s) that maximize(s) the potential number of bonds the
 173 configuration can make when elongated by δ beads in total. This dynamics is *oblivious* as it
 174 keeps no memory of the previously preferred positions [11].

Formally, given an Oritatami system $\mathcal{O} = (p, \heartsuit, \delta)$ and a *seed configuration* σ of a *seed
 bead type sequence* s , we denote by $\mathcal{C}_{\sigma,p}$ the set of all partial configurations of the sequence
 $s \cdot p$ elongating the seed configuration σ . The considered *dynamics* $\mathcal{D} : 2^{\mathcal{C}_{\sigma,p}} \rightarrow 2^{\mathcal{C}_{\sigma,p}}$ maps



■ **Figure 1 Oritatami model:** From left to right, the growth from bead E12 to bead E18 of a self-supported oritatami glider with delay $\delta = 3$, transcript $p = E12 \dots E23$ and rule $\{E12 \heartsuit E17, E14 \heartsuit E21, E18 \heartsuit E23, E20 \heartsuit E15\}$. At each step, the set of nascent paths and maximizing the number of bonds is shown. The nascent beads are highlighted in bold black. The nascent paths are drawn in bold black until the last bond made and ends in colors when their tail is free to move (i.e., is not bounded by any bond).

every subset S of partial configurations of length ℓ elongating σ of the sequence $s \cdot p$ to the subset $\mathcal{D}(S)$ of partial configurations of length $\ell + 1$ of $s \cdot p$ as follows:

$$\mathcal{D}(S) = \bigcup_{c \in S} \arg \max_{\gamma \in c^{\triangleright 1}} \left(\max_{\eta \in \gamma^{\triangleright(\delta-1)}} H(\eta) \right)$$

175 The possible configurations at time t of the oritatami system \mathcal{O} are the elongations of the
 176 seed configuration σ by t beads in the set $\mathcal{D}^t(\{\sigma\})$.

177 We say that the Oritatami system is *deterministic* if at each time t , $\mathcal{D}^t(\{\sigma\})$ is either a
 178 singleton or the empty set. In this case, we denote by c^t the configuration at time t , such
 179 that: $c^0 = \sigma$ and $\mathcal{D}^t(\{\sigma\}) = \{c^t\}$ for all $t > 0$; we say that the partial configuration c^t *folds*
 180 *(co-transcriptionally)* into the partial configuration c^{t+1} deterministically. In this case, at
 181 time t , the $(t + 1)$ -th bead of p is placed at c^{t+1} , that is at the position that maximises the
 182 number of bonds that can be made in a δ -elongation of c^t . Figure 1 illustrates the folding
 183 steps of a delay-3 oritatami glider.

184 **Turedos: Self-avoiding Turing Machines.** A *turedo* is a Turing machine working on
 185 the plane with a lookup neighborhood (like in [1]), that can only move in a self-avoiding
 186 way. Turedos are embedded in the hexagonal lattice $\mathbb{H} = (\mathbb{Z}^2, \sim)$ whose 6 unit vectors
 187 are $N_H = \{\vec{N} = (1, 1), \vec{NE} = (1, 0), \vec{SE} = (0, -1), \vec{S} = (-1, -1), \vec{SW} = (-1, 0), \vec{NW} = (0, 1)\}$.
 188 Note that \mathbb{H} 's underlying grid is rotated by 30° with respect to \mathbb{T} 's. This choice is motivated
 189 by the main simulation result of the paper where macrocells in oritatami in our figures
 190 appear in the same orientation as the hexagonal cells in turedos. We denote by $B(r)$ the
 191 hexagonal ball of radius r centered on $(0, 0)$, i.e. the set of positions in \mathbb{Z}^2 that can be
 192 written as a sum of at most r vectors from N_H . We also denote by $b(r)$ the size of $B(r)$,
 193 and $c_z(r) = (u \in B(r) \mapsto c(z + u))$ the restriction of a configuration c to the ball of radius r
 194 centered on z . Finally, we fix a universal blank symbol \perp representing unoccupied positions.

195 **► Definition 2.1.** A turedo is defined by $\mathcal{T} = (A, Q, q_0, r, \delta)$ where A is the tape alpha-
 196 bet, $\perp \in A$, Q is the set of head states with initial state $q_0 \in Q$, r is the lookup radius,
 197 $\delta : Q \times A^{B(r)} \rightarrow Q \times N_H \times A \setminus \{\perp\}$ is the local transition map.

198 A tape configuration is an element of $A^{\mathbb{Z}^2}$. A global state is an element of
 199 $\mathcal{S}_{\mathcal{T}} = A^{\mathbb{Z}^2} \times \mathbb{Z}^2 \times Q$ (tape configuration, position and state of the head). The turedo \mathcal{T}

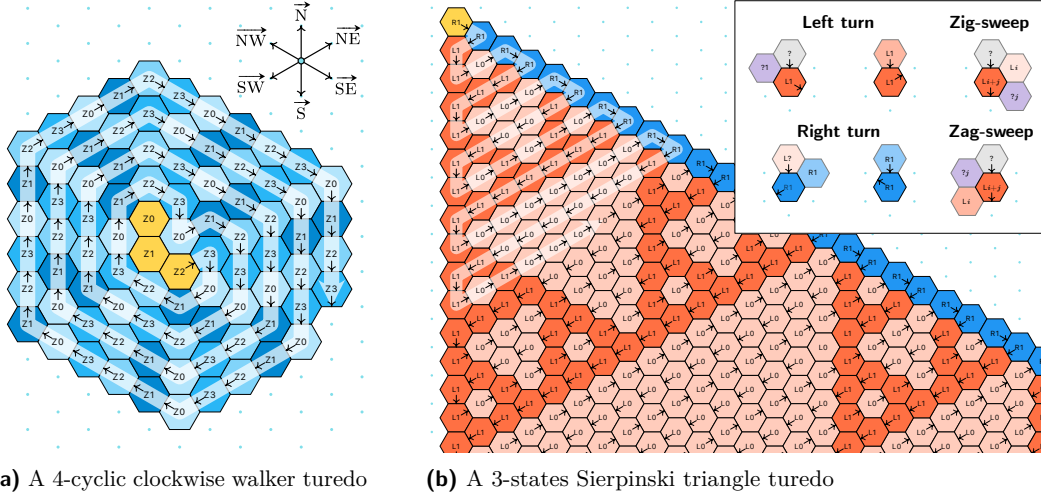


Figure 2 Two examples of radius-1 turedos. The seed configurations are displayed in yellow. The path followed by the turedo is highlighted in white. Empty (blank) positions are marked with a blue dot. **(a)** The turedo exits to the counterclockwise-most empty cell starting from its entry side. The states are just cycling in Z_0, \dots, Z_3 . **(b)** The turedo uses three states L_0, L_1, R_1 to perform a zigzag sweeping drawing the Sierpinski triangles pattern using the local rule given above.

induces a global map $F_{\mathcal{T}} : \mathcal{S}_{\mathcal{T}} \rightarrow \mathcal{S}_{\mathcal{T}}$ defined as follows:

$$F_{\mathcal{T}}(c, z, q) = \begin{cases} (c, z, q) & \text{if } c(z) \neq \perp \text{ or } c(z+d) \neq \perp, \\ (c', z+d, q') & \text{otherwise,} \end{cases}$$

where $(q', d, a) = \delta(q, c_z(r))$ and image configuration c' is defined by: $c'(z) = a$ and $c'(u) = c(u)$ for $u \neq z$. When the first case occurs, we say that the machine is blocked.

The key point of the above definition (which justifies the qualification of 'self-avoiding') is that the only way tape configurations can be altered is by turning a blank symbol into a non-blank symbol, and therefore the head cannot go back to a previously visited position (except when the machine is blocked in which case the global state is a fixed point). Positions holding a blank symbol are therefore seen as empty positions where the head can possibly move to. Two examples of radius-1 turedos are given in Figure 2.

Limit configuration and freezing time. Given an initial global state $s \in \mathcal{S}_{\mathcal{T}}$ for a turedo of global map $F_{\mathcal{T}}$, let us consider the sequence $(c^t, z_t, q_t) = F_{\mathcal{T}}^t(s)$ for $t \in \mathbb{N}$. By the self-avoiding property, it holds that for any $z \in \mathbb{Z}^2$ the sequence of symbols $(c^t(z))_{n \in \mathbb{N}}$ is ultimately constant, and, denoting its limit $c_s^\infty(z)$, we then have defined a tape configuration $c_s^\infty \in A^{\mathbb{Z}^2}$ which is called the *limit configuration* reached by F starting from s . Said differently, using the standard Cantor topology for tape configurations [14], we have that the sequence of configurations $(c^t)_t$ converges to c_s^∞ . Moreover, we can associate to the system and the initial global state s , the *freezing time* map $\tau_s : \mathbb{Z}^2 \rightarrow \mathbb{N}$ such that $\tau_s(z)$ is the minimal t for which the tape content of cell z at time t is $c_s^\infty(z)$ (in particular, $\tau_s(z) = 0$ if $c_s^\infty(z) = \perp$).

Programming turedos. Thanks to the freedom allowed in their local maps, turedos are in general much easier to design than oritatami systems. The basic building block to design

221 complex turedos is the zigzag sweeping movement which allows us to embed any 1D Turing
 222 machine/cellular automaton computation (see Fig. 2b). They can also be used as thick wires
 223 to transport information from one region to another.

224 **Simulations.** Any oritatami with delay δ can be seen as a particular turedo of radius $\delta + 1$:
 225 indeed, an oritatami transition is completely determined by the position in the sequence of
 226 beads, coded as a state of the turedo, and the local configuration in a ball of radius $\delta + 1$.

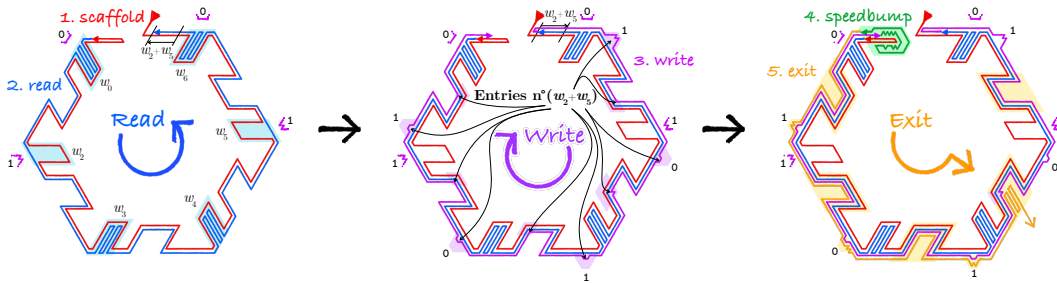
227 Our main result proven in the next section is a converse to this observation: any turedo
 228 of radius 1 can be simulated by an oritatami system of delay 3. The general idea is to
 229 reproduce the dynamics up to a linear spatio-temporal scale factor like in similar notions
 230 already considered for cellular automata or self-assembly tilings [3, 6, 2]. More precisely,
 231 each cell of the simulated system is represented by a macrocell in the simulator system, the
 232 macrocells form a linearly distorted hexagonal lattice, and a constant number of time steps
 233 is allowed for the simulator to reproduce one step of the simulated system. This notion of
 234 simulation is very strict and allows to relate properties of the limit configurations in the
 235 simulated system to the corresponding limit configuration in the simulator. This can be done
 236 without further hypothesis for computability of limit configurations, but can also be done for
 237 the density of non-blank states as soon as the simulation uses macrocells that are filled with
 238 the same high enough density.

239 **3 Delay-3 oritatami systems simulate radius-1 Turedos**

240 This section provides an overview of the design implying main Theorem 1.1. As for the 1D
 241 cellular automaton simulation in [20], our simulation proceeds in three phases: 1) reading
 242 the neighboring letters, 2) preparing for writing the new letter on the boundaries of the
 243 macrocell and 3) exiting to the computed next location. However, we must solve a significant
 244 number of new challenges to adapt to turedos. Turedos are free to move in every direction:
 245 the shape of the macrocells must then be isotropic. Furthermore, as the exit direction has
 246 to be deduced from the symbols read, the reading process must be non-blocking. Thus we
 247 cannot use the reading mechanism in [20], nor the writing flip-flap mechanism which would
 248 block any further return to a previously visited border; we cannot use its hardcoded exit
 249 mechanism either. Moreover, as we need to return to a random side after reading and writing
 250 on all sides, our oritatami system must be able to absorb up to 4 times the side length
 251 before exiting to the new macrocell and starting the next period of the transcript. It follows
 252 that we cannot park unused information on the boundary of the macrocell as in [20], but
 253 need to store information *inside* the macrocell to avoid increasing the macrocell side length
 254 uncontrollably. Similarly the speedbump module introduced in [20] must be adapted to fit
 255 inside a compact space.

256 To solve all those issues, we have developed new tools that we believe to be simple, powerful
 257 and generic enough to have their own interest. We also believe that some of them could serve as
 258 a guideline for a first biochemical implementation of computation using RNA co-transcription.
 259 Our current implementation `turedo2oritatami` uses 1735 bead types. Examples of radius-1
 260 turedos compiled as oritatami as well as a fully functional `python` compiler can be downloaded
 261 from [23]: <https://hub.darcs.net/turedo2oritatami/turedo2oritatami/python>. The
 262 resulting `.os` files are to be run with the oritatami simulator by [22].

263 **Bit-weight encoding of a Turedo.** Consider a radius-1 turedo. First, we get rid of its
 264 internal state and orientation by encoding them in the symbols of the tape configuration.



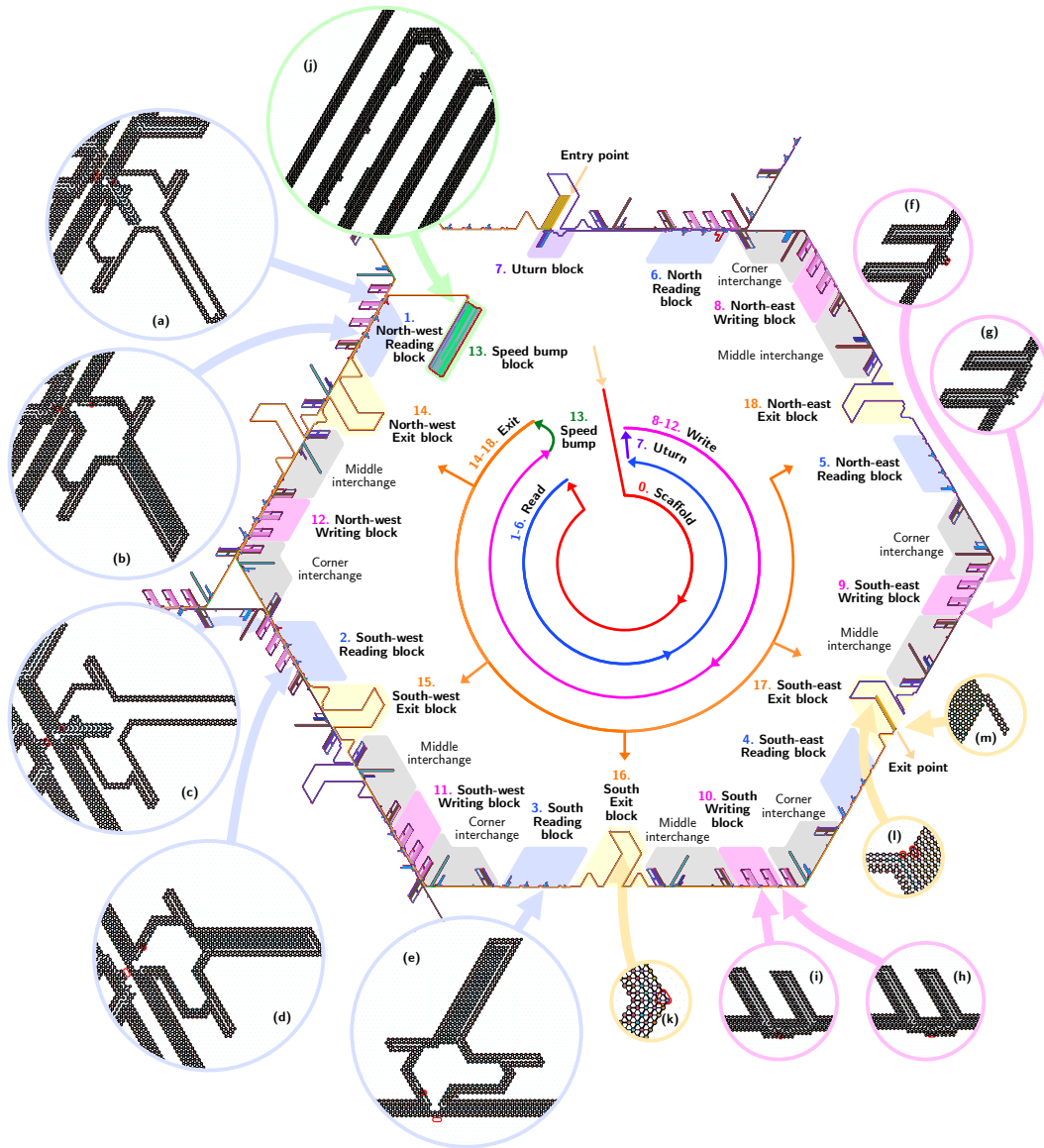
■ **Figure 3** Principle of the macrocell operation. The shift of the reading layer at the end of its folding (and thus of the writing layer) is $\sum_{i:\text{bit read}_i=1} w_i = w_2 + w_5$.

265 We then encode each symbol of the resulting tape alphabet \mathcal{A} as a string of q bits where
 266 $q = \lceil \log_2 \#\mathcal{A} \rceil$. The blank symbol \perp is encoded by the reserved word 0^q . Let $Q = 2^q$.
 267 In the following we assume that the neighboring cells of the current position are numbered
 268 in counterclockwise (CCW) order from 0 to 5 where 5 denotes the cell previously visited
 269 by the turedo. Our simulation assumes that the turedo transition function is a function
 270 $F : (2^q)^6 \rightarrow 2^q \times \{0, \dots, 4\}$, that reads the q bits $b_{i,0}, \dots, b_{i,q-1}$ encoding the symbol in the
 271 i th CCW neighboring cell for $i = 0..5$, and outputs the q bits of the symbol to be written
 272 and the CCW index of the next cell to go to.² Furthermore, we assume that F is encoded
 273 as a tuple $((w_{ij}), \Phi)$ such that $F((b_{ij})) = \Phi(\sum_{i,j} w_{ij} b_{ij})$ where the $6q$ bit-weights (w_{ij}) are
 274 non-negative integers. All transition function F can be encoded this way using the weights
 275 $w_{ij} = 2^{q_i+j}$. We denote by $\mathcal{W} = \sum_{i,j} w_{ij}$ the sum of the weights of the bits. Encoded this
 276 way, the size of the transition table of F is exactly $\mathcal{W} + 1$ for every bit and the exit direction.

277 **Principle of the macrocell operation.** Fig. 3 presents a schematic overview of the key
 278 operations in the macrocell. The transcript consists of five parts:

- 279 1. the *scaffold* of the macrocell folds, on each side of the macrocell, in front of the position
 280 of each bit to be read, “read pockets” (in blue) of size equal to the weight given by the
 281 transition function to that bit; it also builds one “exit pocket” (in orange) per side;
- 282 2. the *read layer* folds counterclockwise and fills the read pockets (outlined in blue) when
 283 it senses a 0, and jumps over it when it senses a 1, pushing the transcript forward by a
 284 shift corresponding to the sum Δ of the sizes of the pockets sensing a 1 ($\Delta = w_2 + w_5$ in
 285 the figure);
- 286 3. the *write layer* contains all the transition tables of the simulated turedo, one for each bit
 287 to write on each side, and one for each exit-or-not decision on each side; this layer folds
 288 clockwise, and as it is translated forward by Δ , it folds the Δ th entry of each transition
 289 table at the writing spots (in purple) that trigger the folding of the selected transition
 290 table entries. The shift Δ accumulated by the read layer allows then to write the output
 291 pattern on each side. It also places a “kicking bead” (in purple) in the exit pocket on the
 292 computed exit side and no-kicking beads in the other using the same shift-principle;
- 293 4. the *speedbump module* (outlined in green) absorbs the shift so that the next layer starts
 294 without any shift regardless of the values read by the read layer;
- 295 5. the *exit layer* folds counterclockwise, following the border until it hits the kick (outlined
 296 in yellow) and folds upon itself to the next macrocell.

² The case of attempting to exit towards the CCW neighboring cell $n^\circ 5$ from which the turedo came, is purposely ignored as it would unnecessarily complicate the construction.



■ **Figure 4** A macrocell for a turedo with $q = 3$ bits ($Q = 8$ tape symbols) together with the order in which layers and modules are used along its boundary as well as snapshots of important modules: (a)-(e) the read pocket in all possible situations: reading a 0/ \perp (fig. b, d and e) or a 1 (fig. a and c) from a neighboring cell (fig. a-d) (or not (fig. e)) and through its exit layer (fig. a and b) or directly from its write layer (fig. c and d) – (f)-(h) all possible situations for the write module: writing a 0 (fig. g and i) or a 1 (fig. f and h), through the exit layer (fig. h and i) or directly (fig. f and g) – (j) the shift-absorbing speedbump – (k) the exit layer folds along the exit pocket – (l)-(m) the write layer has placed the kicking bead ⊗76 in the corner that detaches the exit layer from the pocket and concludes the folding by exiting to the SW. Zoom in for details

297 Observe that the reading layer needs to "read" the bit from neighboring cells while still
 298 making room for the two next layers to fold between the reading layer and the neighboring
 299 cells. This explains why our oritatami systems has delay 3: it has to read through 3 layers.

300 This presentation was just an overview of the macrocell. The complete description of the
 301 macrocell is given in Fig. 4. We will now present some of the key tools used in our design.

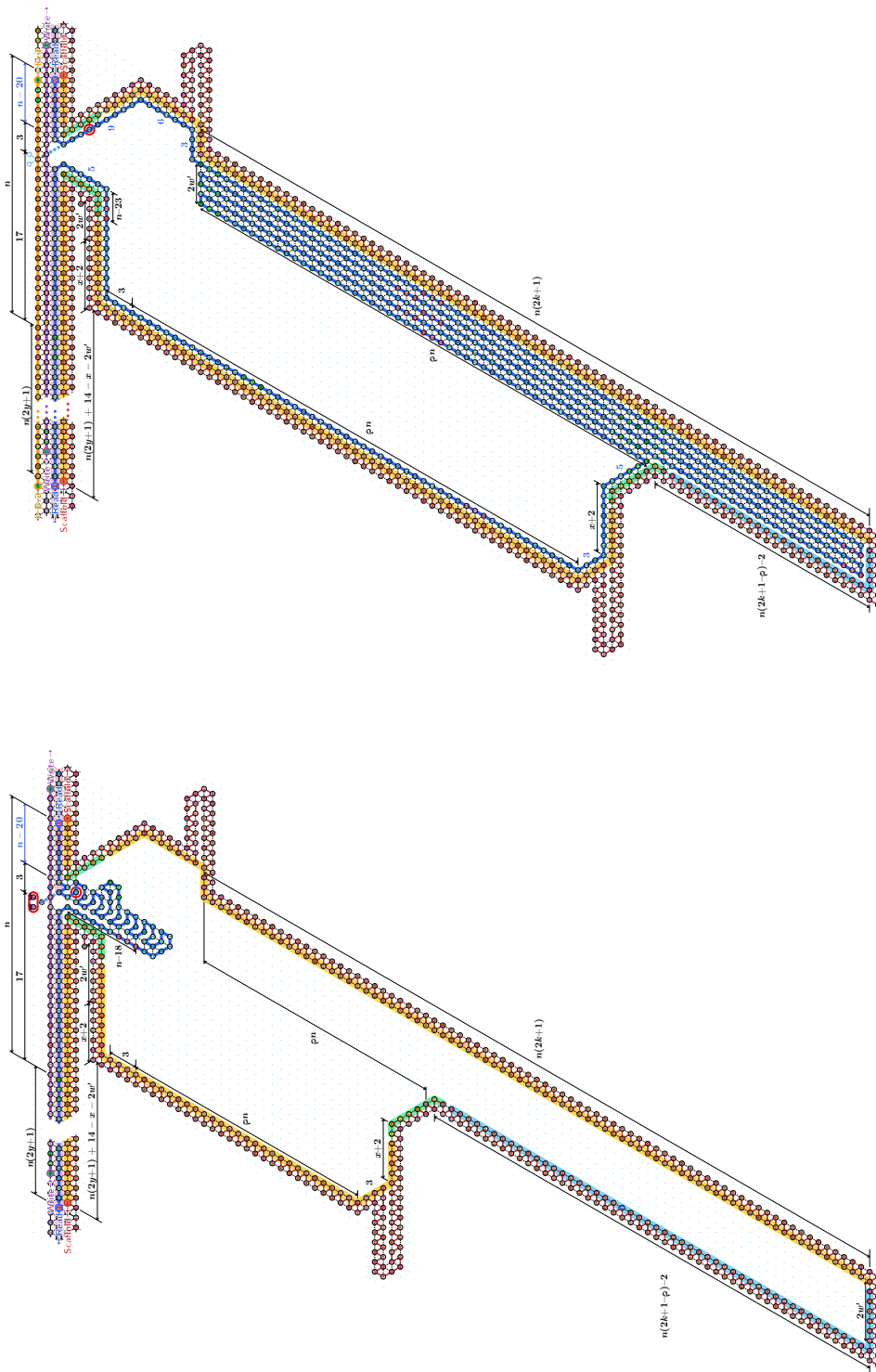
302 **Folding meter and pockets.** Our construction relies on two new simple and powerful tools:

- 303 ■ a *folding meter* is a $4n$ -periodic transcript whose period has 4 equally spaced articulation
 304 points, so that it can either: 1) follow a border if it is strongly attracted to it; 2) fold
 305 upon itself in a compact zig-zag form if the attraction to the border is weak; 3) reveal an
 306 hardcoded structure if the attraction to its surrounding is mild (see Appendix A).
- 307 ■ a *pocket* is a box which triggers the compact folding of a folding meter and which allows to
 308 hide a portion of it in a compact space. The entrance to such a pocket can be conditioned
 309 by the surrounding. For instance, the read folding meter enters a read pocket if and
 310 only if its reading head `rq88` or `rq36` is not attracted by two beads encoding a 1 in its
 311 neighborhood (see Fig. 4(a–e)), otherwise it folds into an hardcoded glider and exits the
 312 pocket rightaway.

313 Furthermore, several folding meters can be layered on top of each other in opposite directions
 314 as long as their periods match. Synchronizing and desynchronizing the two layers allow to
 315 trigger the various behaviors as well, by varying the strength of their bonds. For instance, the
 316 write layer folds into spikes encoding 0 or 1 when it passes over the read layer in Fig. 4(f–i)
 317 because its bonds are weaker with the read layer when the latter is desynchronized after
 318 having been sucked into the pitfalls that surround this area. The length n of the each segment
 319 is a priori arbitrary as long as $n \geq 6$. In the present design, n was set to 26 in order to
 320 accommodate all the desired configurations — the most demanding being the glider at the
 321 entrance of a read pocket when reading a 0, and the writing of 0 or 1 over the write module.

322 **Read layer and pocket.** Let us illustrate the folding meter/pocket mechanism with the
 323 read pocket used to induce a shift of $2n\kappa$ in the transcript every time it reads a 1 with
 324 bit-weight κ . The primary purpose of the read pocket is namely to read a bit (0/1) and
 325 to push forward the read layer by the amount equivalent to its capacity if the read layer
 326 reads a 1. The read layer folds from right to left. When the read layer reaches the entrance
 327 (see Fig. 5a), its "reading head", the bead `rq88`, "senses" whether there is a 1 written on
 328 the adjacent macrocell at this location. If there is a 1, encoded by the presence of the pair
 329 of beads `lp62` and `lp64` in Fig. 5a, then the reading head `rq88` is momentarily attracted
 330 upwards (making two temporary bonds), which results in placing bead `rq86` away from the
 331 border of the read pocket; the read layer gets then too far from the read pocket border
 332 to get attracted to it anymore, and folds into its natural hidden shape: a glider that will
 333 immediately escape from the read pocket (Fig. 5a). Otherwise, if there is a 0, encoded by the
 334 absence of these bead types at the expected location in Fig. 5b, the reading head `rq88` gets
 335 attracted downwards inside the pocket by making one bond; the read layer pursues its course
 336 downwards along the border until it reaches the bottom of the pocket which does not attract
 337 it anymore; the read layer prefers then to fold upon itself into a switchback pattern, filling
 338 the pocket completely, until it reaches the other side of the pocket which attracts it again;
 339 the read layer follows this border until it exits the pocket. This results in a shift forward of
 340 the read layer by an amount corresponding to the pocket capacity $2n\kappa$ if and only if the bit
 341 written on the adjacent macrocell is 1.

342 Remark that this novel bit reading method, using a reading head, does not obstruct the



(a) Read pocket reading 1.

(b) Read pocket reading 0.

Figure 5 Read pocket. The *capacity* of a read pocket is defined to be the difference in length between the paths taken by the transcript upon reading 1 and upon reading 0, and it is determined by the parameters k, w, ρ with $\rho < 2k + 1$ as $\text{capacity} = 2n \left(w(2k + 1) + \rho - 1 + \frac{(w+1)\rho \cdot \text{nextMultiple}(\text{of: } n)}{n} \right)$.

343 way between adjacent cells unlike the method used in [20]; this allows the write and exit
 344 layers to pass and reach the exit at an arbitrary side. Note that this is the reason why our
 345 simulation uses delay 3.

346 Note finally that the interactions between the scaffold and the read layer are extremely
 347 simple: the only places where these interactions are carefully designed are at the entrance
 348 and at the end of the pocket (the three areas highlighted in green in Fig. 5), all the other
 349 interactions are either “attract-them-all” (the areas highlighted in yellow) or “attract-none-
 350 of-them” (the areas highlighted in blue). This demonstrates the simplicity of the folding
 351 meter/pocket concept.

352 **The read and write blocks.** Fig. 6 shows in details the actual oritatami implementation of
 353 the read and write blocks and how write pockets of size equal to the size of the transition
 354 tables are used as interconnected vessels to place the correct entry of the table over each write
 355 module. The write module (responsible for writing 0 or 1 on the sides of the macrocell) and
 356 write pocket (responsible for hiding the unused entries of the transition table) are presented
 357 in details in Appendix B.

358 **Layer interchange.** Each layer is heavily interacting with its neighboring layers inside a
 359 macrocell. It follows that unwanted interferences may occur between facing layers from
 360 neighboring macrocells. For this purpose, we use three different variants of bead types in
 361 each layer: one for each half of each side (e.g. Read1 and Read2 for the read layer), plus one
 362 in the middle and the corners which interconnects one to the other and cancel the need for
 363 interactions between them (e.g., Read12 to switch between Read1 and Read2).

364 **Setting up the exit block and computing the macrocell size.** As the exit pocket needs to
 365 accommodate the remaining of the exit layer before it exits, it must have room to fold in
 366 a compact shape a folding meter of length up to four macrocell-sides long. Since the exit
 367 pocket belongs to the macrocell side, extending the exit pocket extends the macrocell side as
 368 well: we have thus to solve a fix point problem. Moreover, since a different amount of the
 369 exit layer will fold into each exit pocket depending on its location in the macrocell, we need
 370 a mechanism to make sure that, in all cases, the transcript will exit at the same position
 371 on every macrocell side, without interfering with the fix point resolution above. The latter
 372 problem is solved by using a pair of “loose ropes” of equal length, one on each side, “pulling”
 373 on the exit pocket to adapt its position to the macrocell side (see the two triangles of varying
 374 depth surrounding each of the five exit pockets in Fig. 4). Making the exit pocket deep
 375 enough allows to solve this fix point issue, which concludes the proof overview of Thm. 1.1.

376 4 Uncomputable Limit Configurations and Freezing Time

377 A configuration $c \in A^{\mathbb{Z}^2}$ is *computable* if there is a Turing machine which on input $z \in \mathbb{Z}^2$
 378 computes $c(z)$. We are interested in the computability of limit configurations obtained from
 379 finite initial configurations (*i.e.* everywhere \perp except on a finite region).

380 As said in the introduction, constructions of Turing universal oritatami systems known so
 381 far [20, 11] *do not* produce uncomputable limit configurations. The key reason is that they
 382 have a computable *escape direction*: a direction $u \in \mathbb{Z}^2$ and a computable non-decreasing
 383 function μ such that $\mu(t) \rightarrow \infty$ and for any $t \in \mathbb{N}$, the position z_t of the head after t steps
 384 verifies $u \cdot z_t \geq \mu(t)$ where ‘ \cdot ’ denotes the scalar product (*i.e.* the head globally moves away
 385 along the direction u). Such a computable escape direction appears naturally in these

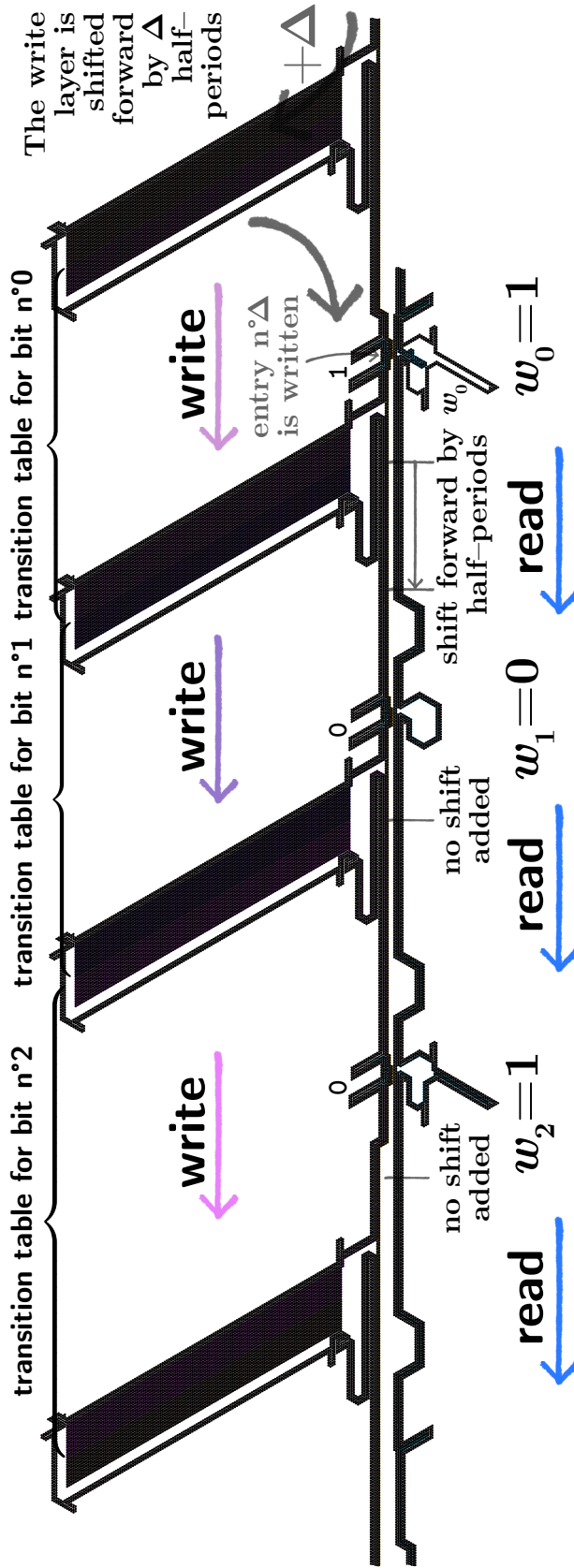
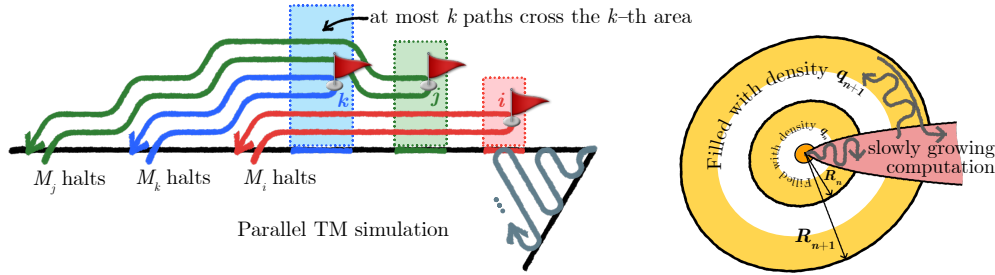


Figure 6 The read and write blocks: (bottom) when the read layer folds, it fills every read pocket facing a 0 on the neighboring macrocell, but skips the read pocket if it faces a 1, which increases the shift forward of the read layer by the weight of the corresponding bit; this yields a total additional shift of w_0 half-periods in the figure – (top) the transition tables are stored in the write layer transcript: one per half-period; the size of the write pockets is set to \mathcal{W} half-periods to accommodate all the unused entries in the $(\mathcal{W} + 1)$ -long transition tables and the transition table is located in the write layer so that its first entry is aligned with the write module when the shift is 0; when the write layer starts to fold, it is shifted forward by $\Delta = \sum_{\text{bit read}(i,j)=1} w_{ij}$ half-periods, the total shift accumulated by its preceding read layer (each transition table is highlighted in a different color in the figure); this implies that the part of the write layer folding over each write module (highlighted in blue) is the one encoding the Δ -th entry of the transition table for each bit to write on the macrocell side as expected; this part will fold into a prescribed shape which will be read as 0 or a 1 by the read layer of its upcoming neighboring macrocell. This is an actual oritatami simulation. Zoom in for details



(a) Sketch of the turedo building an uncomputable limit configuration. (b) Sketch of the turedo building a limit configuration with an arbitrary density $d \in \Pi_2$.

■ **Figure 7** Sketch of the two turedo constructions in sections 4 and 5.

386 simulations because they are fundamentally simulations of space-time of one-dimensional
 387 systems: they work by growing successive 1D finite configurations and stacking them along a
 388 direction u that corresponds to the time of the simulated system. The simulation never goes
 389 back to previously stacked layers simply because computing one step of the 1D system is
 390 performed using the last stacked 1D configuration only. More generally:

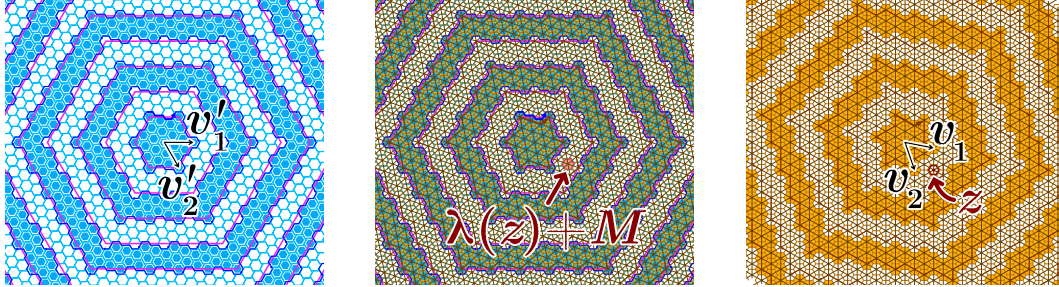
391 ► **Fact 4.1.** *For any turedo reaching limit configuration c_s^∞ from a finite global state s , the*
 392 *maps $z \mapsto c_s^\infty(z)$ and $z \mapsto \tau_s(z)$ are Turing-equivalent. Moreover, they are both computable*
 393 *if the dynamics admits a computable escape direction.*

394 In the next result, we construct a turedo that goes back uncomputably close to the origin
 395 uncomputably often in spite of following a self-avoiding trajectory. Precisely, we prove that
 396 turedos of radius 1 and therefore oritatami are powerful enough to embed any recursively
 397 enumerable set into their limit configurations reached from a finite initial configuration. As
 398 a consequence, both models produce uncomputable limit configurations.

399 ► **Theorem 4.2.** *There exists a fixed turedo of radius 1 which, when started from a fixed*
 400 *global state s with a blank tape configuration, reaches an uncomputable limit configuration*
 401 *and therefore has an uncomputable freezing time map τ_s .*

402 **Proof sketch.** The basic idea, illustrated in Fig. 7a, is to build a turedo which runs a Turing
 403 machine simulation to test all Turing machines for halt in parallel and that, when it finds
 404 that some machine i has halted, interrupts momentarily its computation and goes to write a
 405 flag in a prefabricated area $p(i)$ located at a computable in i position (initially all areas $p(i)$
 406 are empty). Areas of type $p(i)$ are progressively filled in some uncomputable and unknown
 407 order, but, at the limit, it holds that $p(i)$ contains a flag if and only if the machine i halts.
 408 Therefore the limit configuration is uncomputable because it can solve the halting problem
 409 when used as an oracle.

410 The key to implementing this idea is the layout of the paths to reach the areas $p(i)$:
 411 when we proceed as shown in Fig. 7a, no more than i paths will go across the area $p(i)$,
 412 i.e. the ones that correspond to the halting Turing machines j with $j < i$. As a zigzag of
 413 thickness $O(j)$ is enough for the turedo to reach area j , place a flag, and go back, then the
 414 flag in area $p(i)$ (if any) will never be placed higher than $O(i^2)$. It follows that these areas
 415 have quadratic size and their ground basis can be set up in advance by the turedo as it
 416 simulates the Turing machines in parallel (in particular, the turedo will start the simulation
 417 of machine i only after the ground basis of area $p(i)$ is set up). Of course, Figure 7a is a
 418 simplification and does not represent all movements of the turedo's head: in particular, when



■ **Figure 8** The linear map λ between the turedo world and the oritatami world induces a tilt between the concentric balls in the both worlds. This simulation tilt must be compensated by providing to the simulated turedo a pair of vectors as an input, so that it fills a proper discretization of the oritatami world balls when simulated: (left) the shortest radius vectors v'_1, v'_2 of a ball in the oritatami world that can be mapped exactly in the turedo world – (right) when the two corresponding vectors v_1, v_2 in the turedo world are supplied to the turedo as an input, the turedo can use them to build a proper discretization of large enough balls in the oritatami world – (middle) both turedo and oritatami worlds superposed (the target balls are drawn in purple and the discretized turedo ones in blue).

419 moving towards area $p(i)$, the turedo needs to carry on the information i and to bubble up
 420 the ground basis of each area crossed over along the way, and it cannot carry those in its
 421 state set. Using our simulation framework, Theorem 1.2 follows from Theorems 1.1 and 4.2.

5 Characterization of Possible Densities of Limit Configurations

422 We can define the (upper) density $\bar{d}(c)$ of non-blank cells in configuration c as follows:

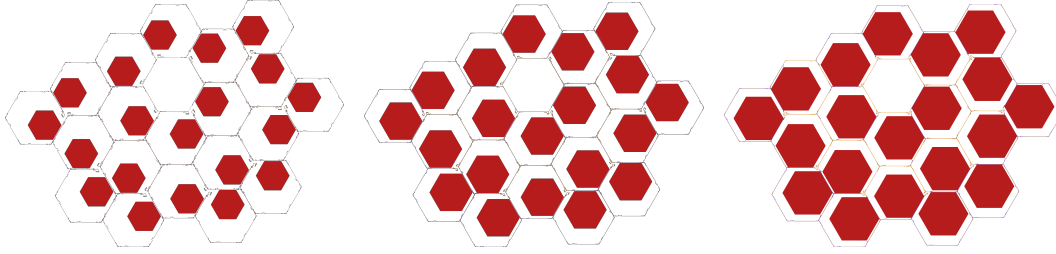
$$423 \quad \bar{d}(c) = \limsup_n \frac{\#\{z \in B(n) : c(z) \neq \perp\}}{b(n)}.$$

424 This choice is natural and gives a translation-invariant notion, but it is not unique (we could
 425 replace the sequence $(B(n))_n$ by another Følner sequence [7]). The problem is that, in a
 426 simulation, the lattice of cells is distorted into a macro-lattice of macro-cells in such a way
 427 that the macro-balls do not have the same shape as genuine balls, as shown in Fig. 8. Said
 428 differently, the reference Følner sequence is distorted into another one and this can change
 429 the density. To circumvent this problem and produce more robust results, we will consider
 430 all possible linearly distorted balls from the start: for any pair $v_1, v_2 \in \mathbb{Z}^2$ of non-colinear
 431 vectors, we consider the (upper) density \bar{d}_{v_1, v_2} of non-blank state after distortion of the
 432 lattice by the pair v_1 and v_2 .

433 We first prove that the computational complexity of $\bar{d}_{v_1, v_2}(c)$ is Π_2 -bounded as soon as c
 434 is produced as the limit of a computable process on finite configurations such that the set of
 435 non-blank positions is monotonically increasing and with diameter growing in a computable
 436 way. This bound applies to turedos but also all systems cited in section 1.

437 ▶ **Lemma 5.1** (Densities of any self-assembling systems are Π_2). *Let c^∞ be the limit configuration reached from some finite seed by some system among oritatami, turedos, freezing cellular automata or directed aTAM. Then for any pair of non-colinear vectors v_1, v_2 , the upper density $\bar{d}_{v_1, v_2}(c^\infty)$ is a Π_2 -computable number.*

438 For non-deterministic systems (both turedos and aTAM), we can state a similar lemma saying that, starting from any finite seed, there is always one orbit converging to a configuration with Π_2 density.



■ **Figure 9** Increasing the density by folding a filled hexagon inside the macrocell expanded by 50, 100 and 200 extra $2n$ -periods on each side. Actual oritatami simulation [Zoom in for details](#)

445 **Arbitrarily dense simulation.** The next theorem is a stronger version of Theorem 1.1,
 446 enforcing a *constant* and arbitrarily large density inside each macrocell of the oritatami
 447 simulation of a given turedo. Precisely, if we consider the *cell partition* of the oritatami
 448 plane into disjoint identical copies of a *macrocell tile* M induced by the map λ from the
 449 turedo world to the oritatami world, where each copy $\lambda(z) + M$ covers exactly the macrocell
 450 corresponding to the turedo position z (see Fig. 8), then:

451 ► **Theorem 5.2.** *For any turedo \mathcal{T} of radius 1, and for any $\epsilon > 0$, there exists an oritatami*
 452 *system of delay 3 that simulates \mathcal{T} and such that the number of occupied positions in each*
 453 *macrocell tile $\lambda(z) + M$ in the oritatami limit configuration is exactly k for all non- \perp*
 454 *position z of the turedo limit configuration (and 0 for \perp position), with $k \geq (1 - \epsilon) \cdot \#M$.*

455 This result is obtained by 1) expanding of the macrocell with a straight line of length L
 456 in the middle of each side so that the empty triangles between the macrocells become negligible
 457 and 2) inserting a sequence in the scaffold that folds into a filled hexagon of radius $L(1 + \alpha)$
 458 inside the space freed inside the macrocell by the expansion. The factor $\alpha > 0$ is necessary to
 459 account for the increase of the exit pocket induced by the increase of the side length (more
 460 transcript needs to fit into the pocket) (see Fig. 9). Picking L large enough concludes the
 461 proof. The case of density 1 is treated separately by an ad hoc solution.

462 **Arbitrary Π_2 -density.** We conclude with the construction of a turedo of radius 1 that
 463 is able to produce limit configurations with any possible density when starting from the
 464 appropriate finite configuration. By possible density we mean any real number $d \in [0, 1]$
 465 which is Π_2 -computable [25], *i.e.* such that there exists a computable sequence of rational
 466 numbers (q_n) with $d = \limsup_n q_n$. The construction is rather technical but the overall idea
 467 is simple (see Fig. 7b): at step n , leave a large annulus empty then densely fill another large
 468 annulus in such a way that the surface ratio between these annuli is q_n and that their sizes
 469 are large enough to dominate all the previously constructed annulus in anterior steps. The
 470 exact sequence of annuli is computed by the turedo in a sublinearly growing (hence negligible)
 471 corridor.

472 ► **Theorem 5.3.** *There exists a turedo of radius 1 such that for any Π_2 -computable number*
 473 *$d \in [0, 1]$ and any pair of non-colinear vectors v_1, v_2 , there is a finite initial global state such*
 474 *that the limit tape configuration c^∞ reached from it verifies: $\bar{d}_{v_1, v_2}(c^\infty) = d$.*

475 The Π_2 -computability limitation is unavoidable as shown in Lemma 5.1, hence our
 476 result is optimal and actually gives a characterization of densities of limit configurations
 477 of continuous sequential self-avoiding systems (resp. turedo, resp. oritatami) started from
 478 finite configurations. Using our simulation framework and Theorem 5.2 we directly deduce
 479 Theorem 1.3.

480 ——— **References** ———

- 481 1 Sebastián Barbieri, Jarkko Kari, and Ville Salo. The group of reversible Turing machines.
482 In *Cellular Automata and Discrete Complex Systems*, pages 49–62. Springer International
483 Publishing, 2016.
- 484 2 Florent Becker, Diego Maldonado, Nicolas Ollinger, and Guillaume Theyssier. Universality in
485 freezing cellular automata. In *Sailing Routes in the World of Computation - 14th Conference*
486 *on Computability in Europe, CiE 2018, Kiel, Germany, July 30 - August 3, 2018, Proceedings*,
487 pages 50–59, 2018.
- 488 3 Marianne Delorme, Jacques Mazoyer, Nicolas Ollinger, and Guillaume Theyssier. Bulking ii:
489 Classifications of cellular automata. *Theor. Comput. Sci.*, 412(30):3881–3905, 2011.
- 490 4 Erik D. Demaine, Jacob Hendricks, Meagan Olsen, Matthew J. Patitz, Trent A. Rogers, Nicolas
491 Schabanel, Shinnosuke Seki, and Hadley Thomas. Know when to fold 'em: Self-assembly
492 of shapes by folding in oritatami. In *DNA Computing and Molecular Programming - 24th*
493 *International Conference, DNA 24, Jinan, China, October 8-12, 2018, Proceedings*, volume
494 LNCS 11145, pages 19–36, 2018.
- 495 5 D. Doty, J. H. Lutz, M. J. Patitz, R. T. Schweller, and S. M. Summer. The tile assembly
496 model is intrinsically universal. In *Proceedings of the 53rd Annual Foundations of Computer*
497 *Science (FOCS)*, 2012.
- 498 6 David Doty, Jack H. Lutz, Matthew J. Patitz, Robert T. Schweller, Scott M. Summers, and
499 Damien Woods. The tile assembly model is intrinsically universal. In *FOCS2012: Proceedings*
500 *of the 53rd Annual IEEE Symposium on Foundations of Computer Science*, pages 302–310,
501 2012.
- 502 7 Erling Følner. On groups with full banach mean value. *MATHEMATICA SCANDINAVICA*,
503 3:243, dec 1955.
- 504 8 Cody Geary, Guido Grossi, Ewan K. S. McRae, Paul W. K. Rothmund, and Ebbe S. Andersen.
505 RNA origami design tools enable cotranscriptional folding of kilobase-sized nanoscaffolds.
506 *Nature Chemistry*, 13:549–558, 2021.
- 507 9 Cody Geary, Pierre-Étienne Meunier, Nicolas Schabanel, and Shinnosuke Seki. Programming
508 biomolecules that fold greedily during transcription. In *MFCS2016: Proceedings of the 41st*
509 *International Symposium on Mathematical Foundations of Computer Science*, volume 58 of
510 *LIPICs*, pages 43:1–43:14, 2016.
- 511 10 Cody Geary, Pierre-Étienne Meunier, Nicolas Schabanel, and Shinnosuke Seki. Oritatami:
512 A computational model for molecular co-transcriptional folding. *International Journal of*
513 *Molecular Sciences*, 9(2259), 2019.
- 514 11 Cody Geary, Pierre-Étienne Meunier, Nicolas Schabanel, and Shinonsuke Seki. Proving the
515 Turing universality of oritatami cotranscriptional folding. In *ISAAC 2018: Proceedings of the*
516 *29th International Symposium on Algorithms and Computation*, volume 123 of *LIPICs*, pages
517 23:1–23:13, 2018.
- 518 12 Cody Geary, Paul W. K. Rothmund, and Ebbe S. Andersen. A single-stranded architecture
519 for cotranscriptional folding of RNA nanostructures. *Science*, 345:799–804, 2014.
- 520 13 E. Goles, N. Ollinger, and G. Theyssier. Introducing freezing cellular automata. In J. Kari,
521 I. Törmä, and M. Szabados, editors, *Exploratory Papers of Cellular Automata and Discrete*
522 *Complex Systems (AUTOMATA 2015)*, pages 65–73, 2015.
- 523 14 Petr Kůrka. *Topological and symbolic dynamics*. Société Mathématique de France, 2003.
- 524 15 James I. Lathrop, Jack H. Lutz, Matthew J. Patitz, and Scott M. Summers. Computability
525 and complexity in self-assembly. *Theory Comput. Syst.*, 48(3):617–647, 2011.
- 526 16 Diego Maldonado, Anahí Gajardo, Benjamin Hellouin de Menibus, and Andrés Moreira.
527 Nontrivial turmites are Turing-universal. *J. Cell. Autom.*, 13(5-6):373–392, 2018.
- 528 17 Yusei Masuda, Shinnosuke Seki, and Yuki Ubukata. Towards the algorithmic molecular
529 self-assembly of fractals by cotranscriptional folding. In *CIAA2018: the 23rd International*
530 *Conference on Implementation and Application of Automata*, volume 10977 of *LNCS*, pages
531 261–273. Springer, 2018.

- 532 18 Nicolas Ollinger and Guillaume Theyssier. Freezing, bounded-change and convergent cellular
533 automata. *CoRR*, abs/1908.06751, 2019.
- 534 19 Matthew J. Patitz. An introduction to tile-based self-assembly and a survey of recent results.
535 *Natural Computing*, 13(2):195–224, 2014.
- 536 20 Daria Pchelina, Nicolas Schabanel, Shinnosuke Seki, and Yuki Ubukata. Simple intrinsic
537 simulation of cellular automata in oritatami molecular folding model. In Yoshiharu Kohayakawa
538 and Flávio Keidi Miyazawa, editors, *LATIN 2020: Theoretical Informatics - 14th Latin*
539 *American Symposium, São Paulo, Brazil, January 5-8, 2021, Proceedings*, volume 12118 of
540 *Lecture Notes in Computer Science*, pages 425–436. Springer, 2020.
- 541 21 Paul W. K. Rothmund, Nick Papadakis, and Erik Winfree. Algorithmic self-assembly of DNA
542 Sierpinski triangles. *PLoS Biology*, 2:2041–2053, 2004.
- 543 22 Nicolas Schabanel. Simple OS simulator, 2021. <http://perso.ens-lyon.fr/nicolas.schabanel/OSsimulator/>.
- 544 23 Nicolas Schabanel and Shinnosuke Seki. Turedo to oritatami compiler, 2022. <https://hub.darcs.net/turedo2oritatami/turedo2oritatami>.
- 545 24 Erik Winfree. *Algorithmic Self-Assembly of DNA*. PhD thesis, California Institute of Technology,
546 1998.
- 547 25 Xizhong Zheng and Klaus Weihrauch. The arithmetical hierarchy of real numbers. In Mirosław
548 Kutylowski, Leszek Pacholski, and Tomasz Wierzbicki, editors, *Mathematical Foundations of*
549 *Computer Science 1999*, pages 23–33, Berlin, Heidelberg, 1999. Springer Berlin Heidelberg.
550
551

552 **A** Transcript, Folding meter and Pocket

553 In our design, the oritatami transcript is periodic, and one period folds into one macrocell.
554 The period is divided semantically in five parts:

555 Transcript = Scaffold · Read · Write · Speedbump · Exit

556 Scaffold hardcodes the “skeleton” of the macrocell and folds clockwise. Read folds around
557 the scaffold counterclockwise. It reads the states of the adjacent macrocells, which induces a
558 shift of the transcript equal to the total weight of the bits read with value 1. Write folds on
559 top of the Read layer clockwise, and, according to the shift read, writes the bits to be output
560 on each side and marks the exit side. Speedbump annihilates the shift using a process similar
561 to [20]. Finally, Exit folds on top of the Write layer counterclockwise until it reaches the “exit
562 mark” that has been placed by the Write layer.

563 Each of the Read, Write, Exit layers have the same periodic structure that we call a
564 folding meter. A n -folding meter is a $4n$ -periodic transcript with a period R of the form:

$$565 \quad R = Rt_0, Rt_1, Rt_2, Rp_3, \dots, Rp_{n-1}, Rb_n, Rb_{n+1}, Rb_{n+2}, Rq_{n+3}, \dots, Rq_{2n-1},$$

$$566 \quad Rt_{2n}, Rt_{2n+1}, Rt_{2n+2}, Rp_{2n+3}, \dots, Rp_{3n-1}, Rb_{3n}, Rb_{3n+1}, Rb_{3n+2}, Rq_{3n+3}, \dots, Rq_{4n-1}$$

566 where the letters t and b stand for *top* and *bottom*. The internal interactions are:

$$567 \quad \underbrace{R_i \heartsuit R_{-i-1}, R_i \heartsuit R_{-i-2}}_{\text{Going down}}, \quad \underbrace{R_{n+i} \heartsuit R_{n-i}, R_{n+i} \heartsuit R_{n-i-1}}_{\text{Going up}} \quad \text{for all } i. \quad (1)$$

568 They ensure that the folding meter will either:

- 569 ■ *follow a border* if all of its beads bind to every bead on the border. Two examples are
570 the Read layer along the right border in Fig. 5, and the Write layer along the left border
571 in Fig. 11. This process allows as well to stack several n -folding meters on top of each
572 other in opposite direction and that are in-sync, i.e. such that one’s p -parts face other’s
573 q -parts. As an example, observe the folding of the three layers Read, Write and Exit at
574 the top right of the write pocket in Fig. 11;
- 575 ■ *or fold upon itself in the manner of the “folding meter” tool*, when reaching the bottom of
576 a pocket that no longer attract the layer. Two examples are the Read layer in Fig. 5b or
577 the Write layer in Fig. 11. Indeed, in Fig. 11, the beads $rp_{15..12}$ do not attract the beads
578 $wb_{26..28}$ which thus fold upwards thanks to the interactions listed in Eq. (1) yielding to
579 a switchback pattern that continues until the Write layer reaches the right side, to which
580 the Write layer is attracted and thus resumes following the border.

581 As the binding between the switchbacks of a n -folding meter are strong, they can flatten
582 any custom interactions encoded internally in either the p - or the q -beads of an n -folding
583 meter as long as these interactions are weak, i.e. do not involve more than 3 bonds per bead.
584 This allows us to hide or expose on-demand specific behaviors when the binding with the
585 lower layer is weak enough: for instance, in Fig. 10, the binding between the p -parts of the
586 Write and Read layers is weak enough to let the p -part of the Write layer fold into various
587 two-spikes patterns encoding 0 or 1 that flatten anywhere else in the macrocell.

588 In this article, $n = 26$. Note that each folding meter is essentially $2n$ -periodic with period
589 (t, p, b, q) . This $2n$ -period is repeated twice only to prevent unwanted interactions when
590 in switchback form. This is why everywhere in the paper the true unit of length is $2n$,
591 half-period of the folding meter, and not a full period. Furthermore every bead type R_i in a
592 folding meter R behaves the same as the beadtype $R(i + 2n) = R(i + 52)$. For this reason,
593 we will adopt the following notation: given a folding meter R , $R\bar{i}$ will refer to either bead
594 types R_i or $R(i + 2n)$; for instance $R\bar{12}$ refers to both bead types R_{12} and R_{64} .

595 **Notations.** For any pair of integers $x \geq 0$ and $y \geq 1$:

596 ■ $x.\text{nextMultiple}(\text{of: } y) = y\lceil x/y \rceil$ is the least multiple of y larger or equal to x

597 ■ $x.\text{complement}(\text{to: } y) = y\lceil x/y \rceil - x$ so that $x + x.\text{complement}(\text{to: } y) = x.\text{nextMultiple}(\text{of: } y)$

598 **B The write block**

599 As seen in Fig. 6, the *write block* on each side of a macrocell consists in an alternation of
600 $q + 1$ write pockets of capacity $2n\mathcal{W}$ beads, and q write modules, one for each of the q bits
601 to be written. Each write pocket hides the \mathcal{W} entries of the $(\mathcal{W} + 1)$ -long transition tables
602 that are unused, while the write module writes the selected entry. Let us start by describing
603 the write module.

604 **B.1 Write module**

605 Write modules are the places where the oritatami writes the bits output on each side. Every
606 side of a macrocell is provided with q write modules, each of which is responsible for one of
607 the q bits to be output. Precisely, this module places two beads of special type (circled in
608 red in figures) at a designated readable site to write a 1 (Figs. 10a and 10b), or deliberately
609 out of the site so that they cannot attract the reading head no matter what types they are
610 to write a 0 (Figs. 10c and 10d).

611 Depending on whether the module is located before of after the side by which the oritatami
612 will exit the macrocell, the module will be covered or not by the **Exit** layer, leading to the
613 4 possible configurations in total presented in Fig. 10. As the side by which the oritatami
614 system will exit is known as soon as the states of the neighboring macrocell are read, we can
615 use, in the **Write** layer description, the appropriate variant of the encoding for each bit on
616 each bit according of the relative position of the module with the exit direction to be taken:

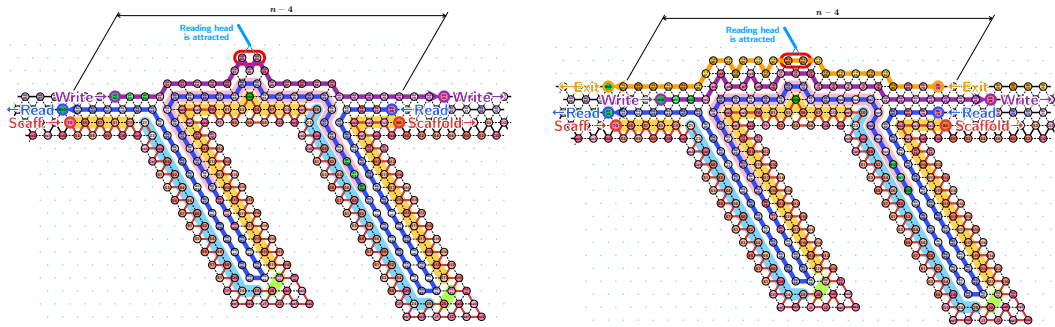
617 ■ In the case where the **Write** layer is to be covered, the **Write** layer folds into two spikes
618 either to the left (Fig. 10b) or the right (Fig. 10d) of the blue hill at the center of the
619 module; then, when the **Exit** layer folds from right to left, the two special bead types $\star\bar{3}\bar{2}$
620 and $\star\bar{3}\bar{3}$ in the **Exit** layer will either be placed at the top of the hill (where they will
621 attract the reading head) or hidden in the right side of the hill (where they cannot
622 attract the reading head), which will be interpreted by the facing macrocell as a 1 or a 0
623 respectively.

624 ■ Otherwise, the bit 0/1 is encoded directly either by two big spikes bearing the special
625 attracting bead types $\sqcup\bar{1}\bar{0}$ and $\sqcup\bar{1}\bar{2}$ at the top of the hill, which will be read as a 1
626 (Fig. 10a), or by two spikes to the left of the hill leaving the designated site empty, which
627 will be read as a 0 (Fig. 10c).

628 In order for the **Write** layer to adopt these peculiar configurations, we need it to take its
629 independence from the underlying **Read** layer. This is accomplished thanks to the two pitfalls
630 surrounding the write module. Each of them have a capacity of n beads exactly, which
631 induces a phase difference of n between the **Read** and the **Write** layers, resulting in the **p**-part
632 of the **Write** layer to fold on top of the **p**-part of **Read** layer over the write modules. These
633 two **p**-parts have specific interactions between them allowing the specific configurations to be
634 folded there and only there.

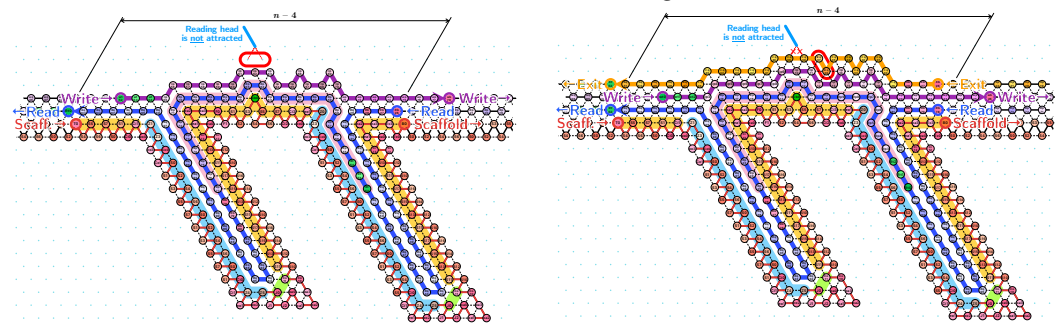
635 **B.2 Write pocket**

636 **Write pocket operation.** Its primary purpose is to hide the \mathcal{W} unused entries of the
637 $(\mathcal{W} + 1)$ -long transition tables as can be observed in Fig. 6. These pockets are placed between



(a) Write module – Top variant: the Write layer writes a 1 by forming two spikes on the top of the module, with two active beads aligned with the reading head of the adjacent macrocell.

(b) Write module – Left variant: the Write layer prepares for writing a 1 by forming two spikes to the left of the module so that the two active beads of the Exit layer get aligned with the reading head of the facing macrocell.



(c) Write module – Right variant 1: on a side located before the exit that will be taken later, the Write layer writes a 0 by forming two spikes to the right of the module; as the Exit layer will exit before reaching this position, the reading positions will stay empty, which will be interpreted as a 0 by the reading head of the facing macrocell.

(d) Write module – Right variant 2: the Write layer prepares for writing a 0 by forming two spikes to the right of the module, so that the two active beads of the Exit layer get misaligned with the reading head of the facing macrocell.

■ Figure 10 The four variants of the Write module: (a,b) writing a 1 and (c,d) writing a 0.

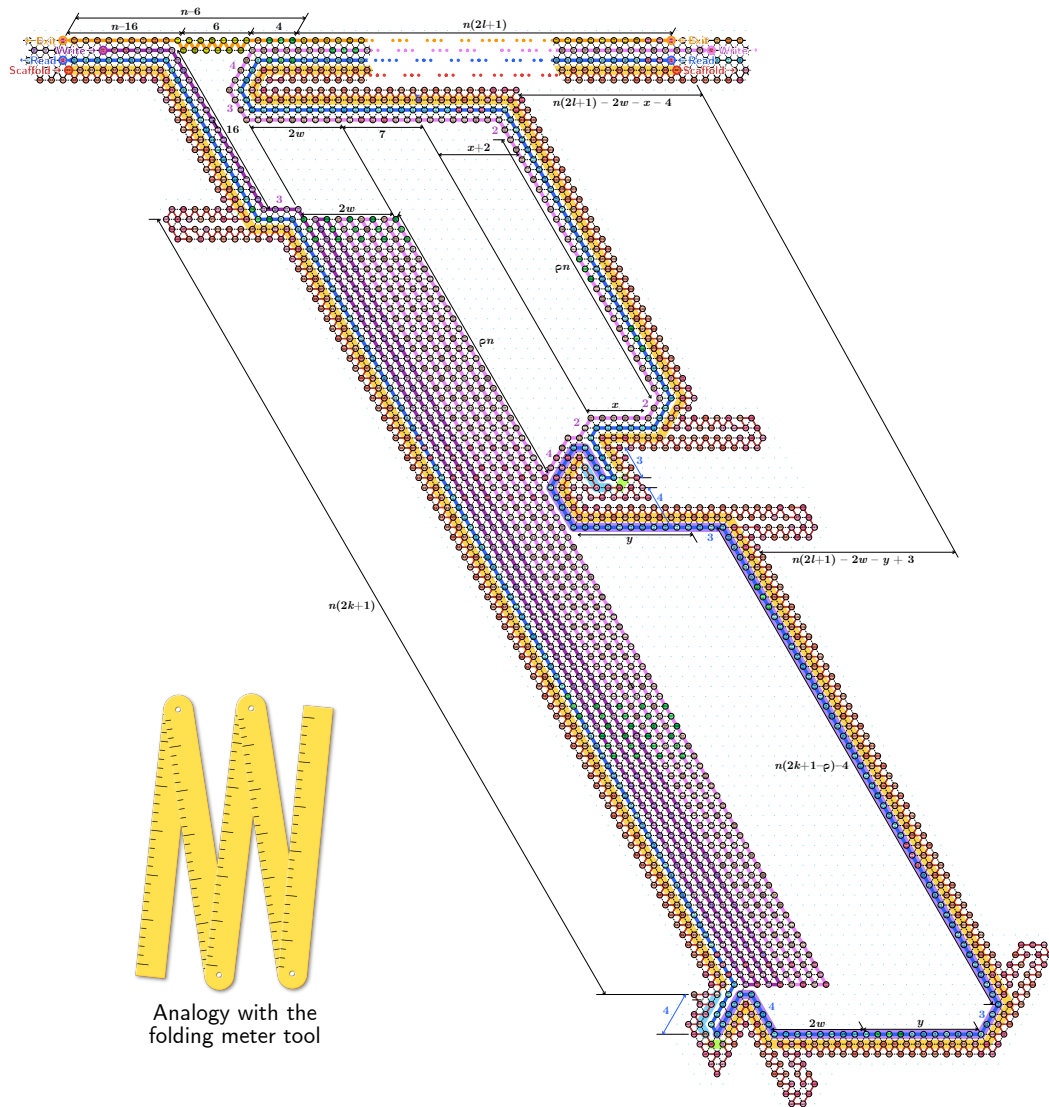
638 the write modules so that only the entries to be written are exposed on the border, at the
 639 locations of the write modules; all the others are hidden in the write pockets. The Read layer
 640 does not fill the write pocket but simply “coats” its border. The Write layer, however, enters
 641 it and folds inside into its compact switchback form, hiding away the \mathcal{W} unused entries of
 642 each transition table, encoded each in one p -part of the Write layer. Finally, the Exit layer
 643 simply jumps over it by folding into a hardcoded bridge to get across its entrance.

644 As for the read pocket, write pockets are also used in the interchange blocks to switch
 645 between the three variants of the write layer $Write1 \leftrightarrow Write12 \leftrightarrow Write2$ (see Fig. 4).

646 **Write pocket design.** This pocket differs from the read pocket in three ways: 1) as opposed
 647 to the read pocket, both layers Read and Write will enter the pocket unconditionally; 2) the
 648 Read and Write layers will enter the pocket from opposite directions; 3) only the Write layer
 649 fills the pocket: the Read layer must not fill the pocket. It follows that:

- 650 ■ Been “coated” by the Read layer, both sides of the write pocket entrance will attract the
 651 Write layer. To avoid unwanted interactions, we need thus to make the entrance wider. It
 652 follows that the Exit layer will not be able to jump over the entrance as the Write layer

653 does over the read pocket. Fortunately, as the **Exit** layer is never shifted, we can hardcode
 654 a bridge $G4..16$ in this layer at this precise location to solve this issue (see top of Fig. 11).
 655 ■ Furthermore, as the **Read** layer will enter the pocket with an arbitrary shift, the interactions
 656 of the pocket with the **Write** layer cannot be directly hardcoded. As illustrated in Fig. 5,
 657 a pocket has essentially three kinds of interactions with the layer that fills it: 1) full
 658 attraction (highlighted in yellow), 2) no attraction (in blue), and 3) localised specific
 659 attractions (in green). We create similar interactions using two mechanisms: A) two
 660 pitfalls are located at the bottom left and at the middle right of the pocket (see Fig. 11)
 661 that introduce and then cancel a phase difference between the **Read** and **Write** layers;
 662 B) the bottom and bottom right borders of the pocket are moved out of reach of the
 663 **Write** layer switchbacks. A) and B) ensure at the bottom right that the **Write** layer is not



■ **Figure 11** Write pocket. Given w , parameters x , y and ℓ must be adjusted so as to match the period of the folding meter. The part of the **Read** layer which is desynchronized with the **Write** layer is highlighted in purple.

664 attracted by the bottom nor opposite border while it folds in switchbacks. Specifically
 665 programmed interactions between the in-sync write bead types $\text{Cb}\overline{78..80}\text{q}\overline{81..83}$ and
 666 the out-of-sync read bead types $\text{rp}\overline{24..25}\text{rb}\overline{26..28}\text{rq}\overline{29}$ that appear just before the second
 667 pitfall in the middle of the right border, ensure that the **Write** layer glues back to the
 668 border once it has ended its switchback pattern and not earlier.

669 Fitting all these constraints together contributes to the choice of $n = 26$ for the period of our
 670 folding meters.

671 **Geometry.** The pocket geometry is determined by four integer parameters: its width w ,
 672 height k , remainder $\rho < 2k + 1$, and extension ℓ . The two "bubbles" to the right of the
 673 pocket are used to keep synchronised the three layers **Read**, **Write** and **Exit**. Their sizes are
 674 determined by two extra integer parameters x and y which are adjusted as follows:

- 675 (a) The length of the **Write** layer path from $\text{Wt}\overline{0}/\text{Wb}\overline{26}$ to $\text{Wt}\overline{0}$ inside the upper bubble
 676 should be equal to 0 modulo $2n$ if ρ is even or equal to n modulo $2n$ if ρ is odd. The total
 677 length is $2(x + w + 17) + \rho n$; thus, x should be set to: $\mathbf{x} = (\mathbf{w} + \mathbf{17}).\text{complement}(\text{to: } n)$.
- 678 (b) The length of the **Read** layer path from $\text{Rt}\overline{0}/\text{Rb}\overline{26}$ to $\text{Rb}\overline{26}$ inside the lower bubble should
 679 be equal to 0 modulo $2n$ if ρ is odd, or equal to n modulo $2n$ if ρ is even. The total
 680 length is $2(w + y + 12) + (2k + 1 - \rho)n$; thus, we set: $\mathbf{y} = (\mathbf{w} + \mathbf{12}).\text{complement}(\text{to: } n)$.
- 681 (c) Lastly, in order to avoid collision with other modules, we set the extension
 682 ℓ so that the pocket module ends to the right of the two bubbles, that is:
 683 $\ell = (\mathbf{2w} + \max(\mathbf{x} + \mathbf{8}, \mathbf{y} - \mathbf{3})).\text{nextMultiple}(\text{of: } 2n)/\mathbf{2n}$.

Capacity. The *capacity* of a write pocket is defined as the length of the path taken by the
Write layer from the leftmost $\text{Wt}\overline{0}$ to the rightmost $\text{Wt}\overline{0}$, and it is determined by the three
independent parameters k, w, ρ with $\rho < 2k + 1$, and one dependent parameter ℓ as:

$$\text{capacity}(w, k, \rho, \ell) = 2n((2k + 1)w + \rho) + 2(w + 17).\text{nextMultiple}(\text{of: } n) + 2n\ell.$$

684 **Building a write pocket with a given capacity.** Various parameters can yield the same
 685 capacity, thus we aim for the parameters that yield the shortest transcript. The length of the
 686 transcript is given by the **Read** layer, whose asymptotic length is $\sim 4nk + 6w$. Minimizing
 687 this value subject to a fixed asymptotic capacity of $2w(2k + 1)n$ yields to the ideal ratio of
 688 $w \sim 2nk/3$. Now, to obtain a write pocket of target capacity $2n\mathcal{W}$ we proceed as follows:

- solving $\text{capacity}(w = 2nk/3, k, \rho = 0, \ell = 0) = 2n\mathcal{W}$ yields a suggested value for k of:

$$k := \max\left(0, \left\lfloor \frac{\sqrt{12n\mathcal{W} + n^2 + 4n - 224}}{4n} - \frac{1}{2n} - \frac{1}{4} \right\rfloor\right)$$

- we then set w by solving $2n\mathcal{W} = \text{capacity}(w, k, \rho = 0, \ell \sim w/n)$ which yields:

$$w := \max\left(1, \left\lfloor \frac{n\mathcal{W} - 19}{n(2k + 1) + 2} \right\rfloor\right)$$

- 689 ■ $x, y,$ and ℓ are then computed according to the formulas (a), (b), and (c).
- we conclude by setting:

$$\rho := \max\left(0, 2n\mathcal{W} - \overbrace{\left(2n((2k + 1)w + 1) + 2(w + 17).\text{nextMultiple}(\text{of: } n) + 2n\ell\right)}^{\text{capacity}(w, k, \rho=0, \ell)}\right) / n,$$

690 if $\rho > 2k$, then rerun the two last steps with $w := w + 1$.

691 This ensures that: $2n\mathcal{W} \leq \text{capacity}(w, k, \rho, \ell) \leq 2n(\mathcal{W} + 2)$ and that $k, w,$ and ℓ are $O(\sqrt{\mathcal{W}})$.