On the power of oritatami cotranscriptional folding with unary bead sequence*

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Abstract. We investigate simple oritatami systems in an attempt to establish lower bounds on the size and complexity of computationally universal systems. In particular, we look at oritatami systems, where the folding sequence consists of a number of beads of the same type and show that under reasonable assumptions, these systems are not universal.

1 Introduction

Transcription is the first essential step of gene expression, when a DNA template sequence is copied into a complementary single stranded RNA sequence by a 'copy machine' called RNA polymerase. The RNA strand is synthesized letter by letter according to the complimentarity relation $A \to U$, $G \to C$, $C \to G$, and $C \to A$ and folds up during a process called co-transcriptional folding.

In a recent breakthrough in molecular engineering by Geary, Rothemund and Andersen [?] the co-transcriptional folding of RNA is controlled by careful design of the DNA template. As demonstrated in laboratory, this method, called RNA Origami, makes it possible to build rectangles out of RNA strands. Geary et al. [?] proposed a mathematical model for this process, called oritatami system. It has been shown [?] that the model is computationally universal by simulating cyclic tag systems introduced by Cook [?]. The simulation involves a very large and complex oritatami system. One future direction of research is to find smaller universal oritatami system.

Closely related is the question of where not to look for universal systems, i.e., what are the limitations of simple oritatami system. In search for simple oritatami systems, there are a number of restrictions one can pose on them:

- bounds on the delay, the number of bead types or the arity;

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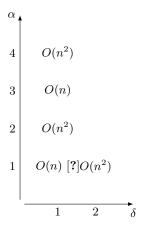


Fig. 1. Summary of the results.

- bounds on the size of the primary structure or of the attraction rule set;
- structural conditions on the primary structure or the attraction rule set.

In this paper we start a new line of study which concerns the alphabet size for the primary structure, i.e., the number of bead types.

2 Preliminaries

Let Σ be a set of types of abstract molecules, or beads. A bead of type $a \in \Sigma$ is called an a-bead. By Σ^* and Σ^ω , we denote the set of finite sequences of beads and that of one-way infinite sequences of beads, respectively. The empty sequence is denoted by λ . Let $w = b_1 b_2 \cdots b_n \in \Sigma^*$ be a sequence of length n for some integer n and bead types $b_1, \ldots, b_n \in \Sigma$. The length of w is denoted by |w|, that is, |w| = n. For two indices i, j with $1 \le i \le j \le n$, we let w[i..j] refer to the subsequence $b_i b_{i+1} \cdots b_{j-1} b_j$; if i = j, then w[i..i] is simplified as w[i]. For $k \ge 1$, w[1..k] is called a prefix of w.

Oritatami systems fold their transcript, which is a sequence of beads, over the triangular grid graph $\mathbb{T}=(V,E)$ cotranscriptionally. We designate one point in V as the origin O of \mathbb{T} . For a point $p\in V$, let N_p^d denote the set of points which lie in the regular hexagon of radius d centered at the point p. A directed path $P=p_1p_2\cdots p_n$ in \mathbb{T} is a sequence of pairwise-distinct points $p_1,p_2,\ldots,p_n\in V$ such that $\{p_i,p_{i+1}\}\in E$ for all $1\leq i< n$. Its i-th point is referred to as P[i]. Now we are ready to abstract RNA single-stranded structures in the name of conformation. A conformation C (over Σ) is a triple (P,w,H) of a directed path P in \mathbb{T} , $w\in \Sigma^*$ of the same length as P, and a set of h-interactions $H\subseteq \big\{\{i,j\}\ \big|\ 1\leq i,i+2\leq j,\{P[i],P[j]\}\in E\big\}$. This is to be interpreted as the sequence w being folded along the path P in such a manner that its i-th bead w[i] is placed at the i-th point P[i] and the i-th and j-th beads are bound

(by a hydrogen-bond-based interaction) if and only if $\{i,j\} \in H$. The condition $i+2 \leq j$ represents the topological restriction that two consecutive beads along the path cannot be bound. A rule set $R \subseteq \Sigma \times \Sigma$ is a symmetric relation over Σ , that is, for all bead types $a,b \in \Sigma$, $(a,b) \in R$ implies $(b,a) \in R$. A bond $\{i,j\} \in H$ is valid with respect to R, or simply R-valid, if $(w[i], w[j]) \in R$. This conformation C is R-valid if all of its bonds are R-valid. For an integer $\alpha \geq 1$, C is of arity α if it contains a bead that forms α bonds but none of its bead forms more. By $\mathcal{C}_{\leq \alpha}(\Sigma)$, we denote the set of all conformations over Σ whose arity is at most α ; its argument Σ is omitted whenever Σ is clear from the context.

The oritatami system grows conformations by an operation called elongation. Given a rule set R and an R-valid conformation $C_1 = (P, w, H)$, we say that another conformation C_2 is an elongation of C_1 by a bead $b \in \Sigma$, written as $C_1 \xrightarrow{R}_b C_2$, if $C_2 = (Pp, wb, H \cup H')$ for some point $p \in V$ not along the path P and set $H' \subseteq \{\{i, |w|+1\} \mid 1 \le i < |w|, \{P[i], p\} \in E, (w[i], b) \in R\}$ of bonds formed by the b-bead; this set H' can be empty. Note that C_2 is also R-valid. This operation is recursively extended to the elongation by a finite sequence of beads as: for any conformation C, $C \xrightarrow{R}_{\lambda}^* C$; and for a finite sequence of beads $w \in \Sigma^*$ and a bead $b \in \Sigma$, a conformation C_1 is elongated to a conformation C_2 by wb, written as $C_1 \xrightarrow{R}_{wb}^* C_2$, if there is a conformation C' that satisfies $C_1 \xrightarrow{R}_w^* C'$ and $C' \xrightarrow{R}_b C_2$.

An oritatami system (OS) $\Xi = (\Sigma, R, \delta, \alpha, \sigma, w)$ is composed of

- a set Σ of bead types,
- a rule set $R \subseteq \Sigma \times \Sigma$,
- a positive integer δ called the *delay*,
- a positive integer α called the arity,
- an initial R-valid conformation $\sigma \in C_{\leq \alpha}(\Sigma)$ called the *seed*, upon which
- its (possibly infinite) transcript $w \in \Sigma^* \cup \Sigma^\omega$ is to be folded by stabilizing beads of w one at a time so as to minimize energy collaboratively with the succeeding $\delta-1$ nascent beads.

The energy of a conformation C = (P, w, H), denoted by $\Delta G(C)$, is defined to be -|H|; the more bonds a conformation has, the more stable it gets. The set $\mathcal{F}(\Xi)$ of conformations foldable by the system Ξ is recursively defined as: the seed σ is in $\mathcal{F}(\Xi)$; and provided that an elongation C_i of σ by the prefix w[1..i] be foldable (i.e., $C_0 = \sigma$), its further elongation C_{i+1} by the next bead w[i+1] is foldable if

$$C_{i+1} \in \underset{C \in \mathcal{C}_{\leq \alpha} s.t.}{\operatorname{arg \, min}} \min \left\{ \Delta G(C') \mid C \xrightarrow{R}^{*}_{w[i+2...i+k]} C', k \leq \delta, C' \in \mathcal{C}_{\leq \alpha} \right\}. \quad (1)$$

$$C_{i} \xrightarrow{R}_{w[i+1]} C$$

Then we say that the bead w[i+1] and the bonds it forms are *stabilized* according to C_{i+1} . Note that an arity- α oritatami system cannot fold any conformation of arity larger than α . A conformation foldable by Ξ is *terminal* if none of its elongations is foldable by Ξ . The oritatami system Ξ is *deterministic* if for all $i \geq 0$,

there exists at most one C_{i+1} that satisfies (1). A deterministic oritatami system folds into a unique terminal conformation. An oritatami system with the empty rule set just folds into an arbitrary elongation of its seed nondeterministically. Thus, the rule set is always assumed to be non-empty.

In the second half of this paper, we consider the unary oritatami system. An oritatami system is unary if its bead type set Σ is of size 1. Its sole bead type is denoted by a, that is, $\Sigma = \{a\}$. Its only possible rule is (a, a) so that the non-empty rule set assumption implies that its rule set is $R = \{(a, a)\}$. Its transcript is a sequence of a-beads. That is to say, the behavior of a unary oritatami system is fully determined by the delay, arity, and seed.

Proposition 1. For any rule set R, arity α and conformation C = (P, w, H) it is possible to check whether C is R-valid and whether $C \in \mathcal{C}_{\leq \alpha}$ in time $\mathcal{O}(|H| \cdot |w| \cdot |R|)$.

Proof. To check whether C is R-valid:

- 1. FOR each $(i, j) \in H$:
- 2. IF $(w[i], w[j]) \notin R$ THEN answer NO and HALT
- 3. answer YES and HALT

Checking the condition in 2. can be done in $\mathcal{O}(|w|\cdot|R|)$ time for any reasonable representation of w and R, hence the whole process takes $\mathcal{O}(|H|\cdot|w|\cdot|R|)$ time. To check the arity constraint $C \in \mathcal{C}_{\leq \alpha}$:

- 1. FOR each $i \in \{1, ..., |w|\}$:
- 2. IF degree $(i) = |\{j|(i,j) \in H\}| > \alpha$ THEN answer NO and HALT
- 3. answer YES and HALT

Checking the condition in 2. can be done in $\mathcal{O}(|H|)$ time for any reasonable representation of H, hence the whole process takes $\mathcal{O}(|w| \cdot |H|)$ time.

Theorem 1. There is an algorithm that simulates any oritatami system $\Xi = (\Sigma, R, \delta, \alpha, \sigma, w)$ in time $2^{\mathcal{O}(\delta)} \cdot |R| \cdot |w|$.

Proof. Take any step in the computation, up to which some $i \geq 0$ first beads of w have been stabilized, with the last bead at a point p. The number of all possible elongations of the current conformation by the next δ-beads is $(6 \times 5^{\delta-1}) \times ((2^4)^{\delta-1} \times 2^5) \in 2^{O(\delta)}$. By Proposition 1, we can check for each of these elongations whether its arity is at most α or not and whether it is R-valid or not in time $\mathcal{O}((2^4)^{\delta-1} \cdot 2^5 \cdot \delta \cdot |R|) = 2^{\mathcal{O}(\delta)} \cdot |R|$. Therefore, the total running time is $2^{\mathcal{O}(\delta)} \cdot |R| \cdot |w|$.

Corollary 1. For fixed δ and α , the class of problems solvable by oritatami systems $(\Sigma, R, \delta, \alpha, \sigma, w)$ is included in DTIME (n^3) .

Proof. The claim follows from Theorem 1 and the fact that |R| is implicitly bounded by $|w|^2$.

Because of the time hierarchy theorems, we know that $P \subseteq EXP$, so we can conclude that OS which cannot deterministically fold transcripts of length exponential in the length of the seed are not computationally universal.

3 Problem description

In [?], Demaine et al. proved that at delay 1 and arity 1, an oritatami system can fold upon the seed of size n only the first 9n beads of a transcript deterministically. We consider this finiteness problem under various settings of the values of delay and arity. Results on this problem are summarized in Fig. 1.

4 Case of $\delta = 1$

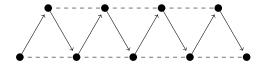


Fig. 2. zig-zag conformation

4.1 Introduction

In this section, we consider on finiteness of structures produced deterministically at delay 1. In result, cases of arity 1 and 3 can only yield finite structures which is size of $\mathcal{O}(n)$, and cases of arity 4 and more can only yield finite structures which is size of $mathcalO(n^2)$, and a case of arity 2 can yield infinite structures but they are only the zig-zag conformation shown in Fig.2.

Let Ξ be a deterministic oritatami system of delay 1 and arity 2. Assume its seed σ consists of n beads. For $i \geq 0$ let C_i be the unique elongation of σ by w[1..i] that is foldable by Ξ . Hence $C_0 = \sigma$.

Let us consider the stabilization of the *i*-th bead a_i upon C_{i-1} . The bead cannot collaborate with any succeeding bead $w[i+1], w[i+2], \cdots$ at delay 1. There are just two ways to get stabilized at delay 1. One way is to be bound to another bead. The other way is through a tunnel section. A tunnel section consists of four beads that occupy four neighbors of a point (Fig.3). Accordingly, how they are stabilized can be described by a binary sequence S of b's (bound) and t's (tunnel section); priority is given to t, that is, S[i] = t if the i-th bead w_i is stabilized not only by being bound but also through a tunnel section.

Assume that four of the six neighbors of a point p are occupied by beads $a_{j_1}, a_{j_2}, a_{j_3}, a_{j_4}$ while the other two are free. We call such a bead p as inside of a tunnel and such beads p' as entrance of a tunnel without a case that p' is inside of a tunnel. If the beads w[i-2] and w[i-1] are stabilized respectively at one of the two free neighbors and at p one after another, then the next bead w[i] cannot help but be stabilized at the other free neighbor. In this way, w[i] can get stabilized without being bound.

If a bead is stabilized through a tunnel section, then it can provide some bonds. Let us consider bonds of C_i . C_i is represented C=(W,P,H) where |W|=i+n C_i contains $\alpha\cdot(i+n)-2|H|$ bonds because C_i consists of i+n beads and a bead has just α bonds and then 2|H| of the those bonds are already used. However, even if a bead has an available bond, w[j] (j>i) might not be able to use this bond because the bond has possibility that it is blocked by transcripts w[i+1..j-1]. Number of binding capabilities does not contain that case so that it is at most $\alpha\cdot(i+n)-2|H|$.

Definition 1 (binding capability). We say that p is reachable if there exists directed path P' from q to p such that the path of P' does not cross the path of P where q_0 is a bead which is latest stabilized, and q_m is a reachable point. Let B_i be $(\{(h,i)|\forall h < i\} \cup \{(i,j)|\forall j > i\}) \cap H$. Moreover, let R_i be a set of inside $\bigcirc_{w_i}^1$ that are free and reachable. We represent the number of binding capabilities of a conformation C_j as $\#bc(C_j)$. $\#bc(C_j)$ is defined by $\sum_{k=-n+1}^j \min\{|B_k|, |R_k|\}$.

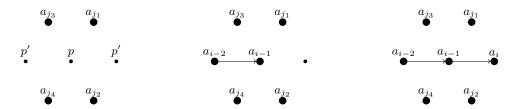


Fig. 3. Through a tunnel section

Theorem 2 (Tunnel Troll Theorem). Let Ξ be a unary oritatami system of $\delta = 1, \alpha \geq 2$. If there is a part of transcript w[i...j + 1] = bttt...tb, then $\#bc(C_i)$ is more than $\#bc(C_j)$.

Proof. Assume Ξ is deterministic. Each of beads in transcript are bound either inside of tunnel or outside (Fig. 5). If a bead is stabilized at inside of tunnel, then its successor is already decided the position either inside of a tunnel or outside. Moreover, If a bead is stabilized at outside of a tunnel, then its position is an entrance of a tunnel or otherwise.

Tunnel sections have three possible shape with considering symmetry such as straight, acute turn and obtuse turn (Fig. 4).

Let us consider each of cases of tunnel A, B, and C. Accordingly, We use lemma 1

Lemma 1. If a bead does not have any bond, then neighbors of it must be occupied by $\alpha + 2$ beads at $\delta = 1$ and unary except first and last beads.

Proof (lemma 1). A bead in transcript needs predecessor and successor except first and last beads. If the bead does not have ant bond, then it use hand with α neighbors. Thus, lemma 1 is clearly true.

Let us consider tunnel sections only tunnel A and B. See Fig. 5. From appendix (Entrance of Tunnel A, B and Exit of Tunnel), we add maximum number which each transitions provide binding capabilities for each edge where a is number of consuming binding capabilities when the bead is stabilized at position of successor in outside.

Next, we consider on tunnel C section. If w[i] is stabilized by tunnel C and S[i+1] is t, then w[i+1] is stabilized by tunnel A or B because if w[i+1] is stabilized by tunnel C, then C_{i+1} is a terminal. Hence, tunnel C section is divided cases such as Figure 6. Cases of $S[i...] = bt^l (l \ge 2)$ are already considered (Upper). According to appendix (Tunnel C), cases of S[i...i+2] = btb also consume some binding capabilities (Lower).

Thus, if a bead is stabilized through a tunnel section, then it consume some binging capabilities.

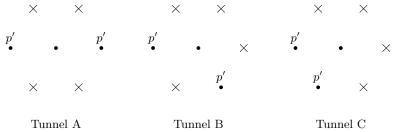


Fig. 4. All possible tunnel sections: straight, acute turn, and obtuse turn

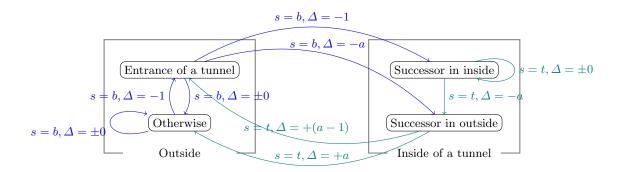


Fig. 5. Increment on Tunnel A,B

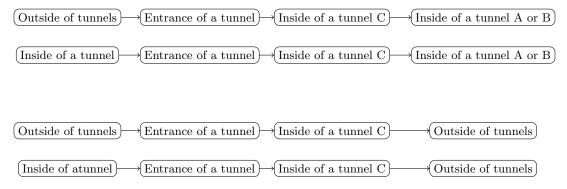


Fig. 6. Case of Tunnel C

4.2 Appendix of Tunnel Troll Theorem

Entrance of TunnelA,B Fig.7 exhibits all the three kinds of entrance of tunnel A, B. Any cases in $\delta = 1, \alpha = 2$ consume some binding capabilities into the follows.

- Case of t_0

Let us consider points of n_3 , n_4 either occupied or not. A point n_3 or n_4 is free because if both of them are occupied, p' is inside of tunnel. If n_3 is free, then p' has to be bound to a bead except n_1 due to deterministically stabilize. In this situation, at least three neighbors of n_1 are free that is n_1 leave at least one bond from lemma 1. Hence, p' must be bound to n_1 . Thus, a case of t_0 consumes two binding capabilities and it does not supply any binding capabilities.

- Case of $t_{\pm 60}$

In this case, a point n_4 or n_5 is free, too. If n_5 is free, p' has to be bound to n_1 or n_2 . If n_5 is occupied, then n_4 is free. This time, n_2 has some binding capabilities so p' has to be bound to n_2 .

In this situation, p' is able to supply a binding capability, if this capability is active, p' bind a bead into n_4 or n_5 . However, n_2 and n_3 are exist in back bone. According to Jordan curve theorem, any successors of p' cannot reach a point n_4 or n_5 so this capability is inactive. Thus, a case of $t_{\pm 60}$ consumes some binding capabilities.

- Case of $t_{\pm 120}$

Binding capabilities that p' supply are inactive according to Jordan curve theorem on n_1 and n_2 . Moreover, p' has to be bound to one of n_3 , n_4 , n_5 in order to deterministically stabilize. Thus, a case of $t_{\pm 120}$ consumes some binging capabilities.

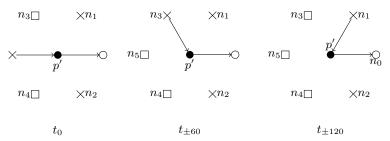


Fig. 7. Direction into a entrance

Exit of Tunnel Fig.8 exhibits all the two kings of exit of tunnel. At least one of point n_1 or n_2 is free because if both of them are occupied, p' is inside of tunnel.

 $\delta = 1, \alpha = 2$ Any cases of $\delta = 1, \alpha = 2$ supply at most a binding capabilities into follows where a is number of predecessor of p' consumes binging capabilities.

- Case of n_1 and n_2 are free
 - This case can be regarded same situation as entrance. See Fig.8 (Left). Predecessor n_5 has to be bound n_4 and n_5 because each of n_3 and n_4 leave binding capabilities. Hence, at least a=2. This time, $\alpha=2$ that is this case supply at most only a binding capabilities.
- Case of c is occupied See Fig.8 (Right). If n_1 is occupied, then n_2 is free so that n_5 has to be bound n_4 . Hence, at least a=1. This case can supply two binding capabilities but p' can bind to only one of n_0 or n_2 because n_0 or n_2 will be occupied a successor of p'. Therefore, this case supply at most only a=1 active binding capability.

 $\delta=1, \alpha\geq 3$ Any cases of $\delta=1, \alpha\geq 3$ consume some binding capabilities into follows.

- Case of n_1 and n_2 are free
 - In $\alpha \geq 3$, if three neighbors of a bead leave, then it can supply two binding capabilities. Therefore Predecessor n_5 has to be bound n_3 and n_4 , and p', too. In this case, at least consumes four bindings and supplies at most two bindings. Thus, it consumes some binding capabilities, totally.
- Case of c is occupied
- In this case, n_4 leave at least two bindings and n_3 , n_1 also leave at least one binding. Therefore n_5 has to be bound n_3 and n_4 , and p' also has to be bound n_1 and n_4 . In this case, at least consumes four bindings and supplies at most one binding. Thus, it consumes some binding capabilities, totally.

Point n_1 and n_2 are free

Point n_1 is occupied

Fig. 8. Exit of Tunnel

Tunnel C Assume w[i] is a bead which stabilized by tunnel C. Let us consider kinds of stabilization S[i-2..i] = tbt or S[i-2..i] = bbt except cases of w[i] is inside of tunnel A, B.

Case of S[i-2..i] = tbt Fig.9 exhibits all the two kinds of stabilization depending on structures of tunnel C.

- Left of Fig.9

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In this figure, Bead n_4 has at least one binding so that w[i-2] has to bound n_4 . Moreover, w[i-1] has to bound one of n_1, n_2, n_3 in order to stabilize deterministically. On the other hand, w[i] can supply two bindings but free neighbors of w[i] are two points. One of them is occupied by a successor. Therefore w[i] can only bind one of n_5, n_6 that is w[i] supplies at most one binding. Thus, this case consumes some binding capabilities.

Right of Fig.9

This cases are divided on number of capabilities that w[i-2] consumes.

- w[i] does not consume any bindings w[i-1] has to bound one of n_1, n_2, n_3 in order to stabilize deterministically. w[i] has to be bound to w[i-2] because w[i-2] has bindings. This time, let us consider either n_4 is occupied or not. If n_4 is occupied, then w[i-2] has no active bindings that is this situation consumes some binding capabilities. If n_4 is free and w[i+1] is stabilized in n_4 , then w[i-2] has to bind w[i+1]. Therefore, In this case, stabilization of w[i-2...i+1] consumes some bindings. If n_4 is free and w[i+1] is stabilized except n_4 , then this oritatami system has to use two binding capabilities in order to bind w[i+1]. Therefore, in this case consumes some bindings. Thus in this cases consume some binding capabilities.
- w_i consumes one binding In this case, w_{i-1} has to be bound one of n_1, n_2, n_3 . In addition, w[i-2] and w[i] are not supply any bindings. Thus, in this cases consume some binding capabilities.
- w_i consumes two bindings In this case, w[i-2] already consumes two binding. w[i-1] has to be bound. w[i] supplies two bindings. Thus, in this cases consume some binding capabilities.

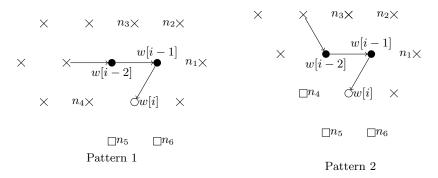


Fig. 9. Case of S[i - 2..i] = tbt

Case of S[i-2..i] = bbt Let us consider number of consumed by w[i-2] (Fig.10).

- -w[i-2] consumes one binding In this situation, w[i-2] supplies one active binding whereas w[i] consumes this binding. In addition, w[i-1] has to bound to one of n_1, n_2, n_3 . Thus, in this cases consume some binding capabilities.
- -w[i-2] consumes two bindings In this case, w[i-2] already consumes two binding. w[i-1] has to be bound. w[i] supplies at most two bindings. Thus, in this cases consume some binding capabilities.

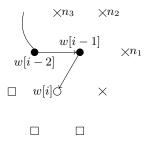


Fig. 10. Case of S[i-2..i] = bbt

By Tunnel Troll Theorem, any tunnel sections which represented in bbt^+ or bt^+bt^+ consume binding capabilities. If the sequence S is free from any subsequence of the form bt^+bt^+ , then it can factorize as $S=u_1u_2u_3\cdots$ for some $u_1,u_2,u_3,\dots\in\{b\}\cup bbt^+$. Assume the length of σ is n, seed supplies at most 2n binding capabilities. Therefore formula 2 hold.

$$\exists i \in \mathbb{N} \quad s.t. \quad u_{i-1}, u_i, u_{i+1}, u_{i+2}, \dots \in \{b\}$$
 (2)

Let us represent S as $S[i.i+1...] = v_i v_{i+1} v_{i+2} \cdots$ for some $v_i, v_{i+1}, v_{i+2}, \cdots \in \{a, o\}$ where if v_k is a, then v_{k+1} is bound to v_{k-1} , if v_k is a, then a is NOT bound to a both the a bound to a bound to a both the a bound to a both the a

Let us consider the case of v_k is o. See Fig.11. w[i-1] consumes some binding capabilities because v_{i-1} is b. If the number of w[i-1]'s bindings is one binding, then w[i+1] has to be bound except n_1 or n_2 so that w[i+1] must consumes two bindings except the case of n_1 and n_2 are occupied and w[i] consumes at least one binding. If n_1 and n_2 are occupied, then w[i-1]'s binginds are inactive that is w[i-1] consumes two binding capabilities. Therefore, this case consumes binding capabilities. If w[i-1] dose Not have any bindings, then w[i-1] already consumes two bindings. In addition, w[i] and w[i+1] consume at least one binding. Therefore this case consumes binding capabilities. Thus, the formula 3 hold and according to the formula 2 and the formula 3, the formula 4 is hold.

$$\exists j \in \mathbb{N} \quad s.t. \quad u_j, u_{j+1}, u_{j+2}, \dots \in \{a\}$$
 (3)

$$|S| > \forall m \in \mathbb{N} \quad \to \quad \exists n \in \mathbb{N} \quad s.t. \quad S[n], S[n+1], \dots \in \{a\}$$
 (4)

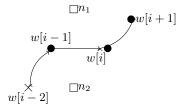


Fig. 11. Case of S[i]

5 Case of $\delta \geq 2$

5.1 $\alpha = 1$, unary

Let the point where the first transcript bead was fixed be p and let n = |seed| + 1. We will argue about the situation when the first bead is stabilized outside \bigcirc_p^n (a hexagon of radius n). Let this be the ith bead of the transcript. Without loss of generality, we can translate the origin (0,0) to the coordinates of bead i-1

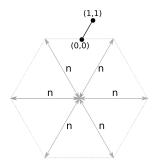
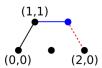


Fig. 12. N_p^n and the position (1,1) of the first bead fixed outside of it.

(which is still in \bigcirc_p^n), and we can assume that the bead outside the hexagon is fixed at (1,1) (see Fig. 12).

In the elongation that places bead i at (1,1) there are two possibilities:

- -i forms a bond with a bead at (1,0).
- -i does not bond to anything and i+1 is at (2,1) bonding with a bead at (2,0). If there is no bead at (1,0), then placing i at (1,0) instead of (1,1) results in the same number of bonds, leading to nondeterminism. Therefore, there has to be a bead at (1,0) and it is inert, otherwise it would bond to i. This is analogous to case 1. below.



The next bead, i+1, can be fixed at (2,1) or at (0,1) as all other possibilities result in nondeterministic behavior immediately, so we have two cases.

1. bead i+1 is fixed at (2,1) and can bond with a bead at (2,0). Now consider bead i+2. For i+1 to be fixed at (2,1), i+2 needs to form a bond somewhere, otherwise i+2 could go to (2,1) forming the bond with the bead at (2,0) and there would be two conformations with the maximal 1 bond. The only possibility is that there is a bead at (3,0) and i+2 can bond with it when placed at (3,1). We can apply the same argument inductively: there is some $m \geq 0$ such that grid points $(\ell,0)$ are occupied by active beads, for all $\ell \in \{2,\ldots,2+m\}$, and there is no bead at (3+m,0). Such an m exists, and it is not greater than n. Then, bead $i+\ell$ is fixed at $(\ell+1,1)$ and bonds with $(\ell+1,0)$. However, bead i+2+m cannot be fixed anywhere, because i+2+m and i+3+m can only add one bond to the conformation, and that is possible either with $i+2+m \to (2+m,1)$, $i+3+m \to (3+m,1)$ or with $i+2+m \to (2+m,2)$, $i+3+m \to (2+m,1)$.

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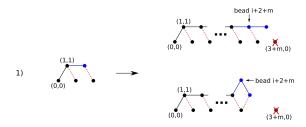


Fig. 13.

- 2. bead i + 1 is fixed at (0, 1). This is only possible if
 - (a) there is an inactive bead at (-1,0) and an active one at (-2,0). This case is symmetrical to (1).
 - (b) there is no bead at (-1,0), bead i+1 can bond with bead i-1 at (0,0) and the bead i+2 can be placed at (-1,0) where it can bond with (-2,0), (-2,-1) or (-1,-1). This leads to nondeterminism, because bead i at (-1,0) and bead i+1 at (0,1) has two bonds, just as the original conformation.
 - (c) there is a bead at (-1,0) and bead i+1 can bond with that or with bead i-1 at (0,0). However, this means that placing bead i at (0,1) at bead i+1 at (1,1) creates the same number of hydrogen bonds, thus resulting in bead i not being placed deterministically.

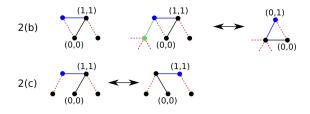


Fig. 14.

5.2 In the case of $\delta = 3$

We discuss when the length of seed is n. If the length of seed is n, the seed will be in an equilateral hexagon with a radius of n when it is given. We will begin our disscussion by considering the moment when a first bead is fixed on a side of an equilateral hexagon with a radius of n+1. Let us assume that a fixed bead on a side of an equilateral hexagon with a radius of n is able to make a hydrogen bond. If it is able to make only two hydrogen bonds, it will be nondeterministic

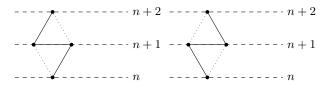


Fig. 15.

because there are two possibilities that it makes two structures in Fig.5. Therefore, it needs to make three hydrogen bonds to determine the only structure.

There are the two cases in which a first bead is fixed on a side of an equilateral hexagon with a radius of n+1 and it makes three hydrogen bonds in Fig.6. If it makes three hydrogen bonds once such as Fig6., it will make three hyrogen bonds forever to be deterministic. This becomes nondeterministic finally, such as in Fig.7., when it reaches a corner of an equilateral hexagon. Thus, the fixed bead on a side of an equilateral hexagon with a radius of n has to make a hydrogen bond with a bead in an equilateral hexagon.

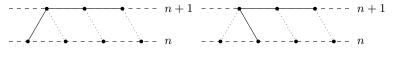


Fig. 16.

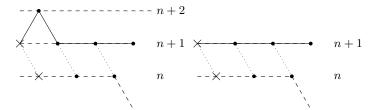


Fig. 17.

Then, we will discuss the moment after the first the is fixed on a side of an equilateral hexagon with a radius of n + 1. Let us assume that the bead is able to make a hydrogen bond. If it is able to make only two hydrogen bonds, it will be nondeterministic because there are possibilities that it makes two structures

in Fig.8. Hence, it needs to make three hydrogen bonds to determine the only structure. It needs to be a right or left figure in Fig.9. to make three hydrogen bonds.

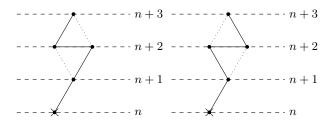


Fig. 18.

If it makes three hydrogen bonds once such as Fig.9., it will make three hyrogen bonds forever to be deterministic. This becomes nondeterministic finally, such as in Fig.7., when it reaches a corner of an equilateral hexagon. Therefore, the first fixed bead on a side of an equilateral hexagon with a radius of n+1 has to make a hydrogen bond with a molecule on a side of an equilateral hexagon with a radius of n.

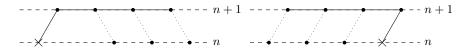


Fig. 19.

Accordingly, it is Fig.10. when a first bead is fixed on a side of an equilateral hexagon with a radius of n + 1.



Fig. 20.

It has to be a right or left figure in Fig.6. when the first bead fixed on a side of an equilateral hexagon with a radius of n + 1 make a hydrogen bond with

a bead on a side of an equilateral hexagon with a radius of n. Fig.6. becomes nondeterministic finally, such as in Fig.7., when it reaches a corner of an equilateral hexagon. It is clear that all beads on a side of an equilateral hexagon with a radius of n+1 has to make a hydrogen bond with a bead on a side of an equilateral hexagon with a radius of n. It is impossible to fix molecules on a side of an equilateral hexagon with a radius of n+2 and determine the only structure when there are only beads which are able to make a hydrogen bond in an equilateral hexagon with a radius of n.