Ciliate Gene Unscrambling with Fewer Templates

Lila Kari*  Afroza Rahman
Department of Computer Science, University of Western Ontario,
London, Ontario N6A 5B7, Canada
lila@csd.uwo.ca arahman@cs.queensu.ca

One of the theoretical models proposed for the mechanism of gene unscrambling in some species of ciliates is the template-guided recombination (TGR) system by Prescott, Ehrenfeucht and Rozenberg which has been generalized by Daley and McQuillan from a formal language theory perspective. In this paper, we propose a refinement of this model that generates regular languages using the iterated TGR system with a finite initial language and a finite set of templates, using fewer templates and a smaller alphabet compared to that of the Daley-McQuillan model. To achieve Turing completeness using only finite components, i.e., a finite initial language and a finite set of templates, we also propose an extension of the contextual template-guided recombination system (CTGR system) by Daley and McQuillan, by adding an extra control called permitting contexts on the usage of templates.

1 Introduction

This paper proposes improvements in the descriptional complexity of two theoretical models of gene unscrambling in ciliates: template-guided recombination (TGR) systems, and contextual template-guided recombination (CTGR) systems. Ciliates are a group of unicellular eukaryotic protozoans, some of which have the distinctive characteristic of nuclear dualism, i.e., they have two types of nuclei: a functionally inert micronucleus and an active macronucleus. Genes in the active macronucleus provide RNA transcripts for the maintenance of the structure and function of the cell. Genes within the micronucleus are usually inactive and assist only in the conjugation process. The process of “decryption” the micronuclear genes after conjugation, to obtain the functional macronuclear genes, is called gene unscrambling or gene assembly.

The genes within micronuclear chromosomes are composed of protein-coding DNA segments (also known as macronuclear destined sequences (MDSs)) interspersed by numerous, short, non-protein-coding DNA segments (also called internally eliminated sequences (IESs)). Furthermore, in some species of ciliates such as Oxytricha or Stylonychia, the micronuclear gene has been found to have a highly complex structure in which MDSs are stored in a permuted order. During the course of macronuclear development, these IESs are eliminated from the micronucleus by means of homologous recombination, and the permuted MDSs are sorted, resulting in a functionally complete macronucleus with MDSs present in the correct order. In the micronuclear sequence, each MDS is flanked by guiding short sequences, 3 to 20 nucleotides long, which act as pointers in a linked list. For instance, the $n$th MDS is flanked on the left by the same short sequence which flanks the $(n-1)$th MDS on the right. In the process of gene unscrambling, homologous recombination takes place between two DNA molecules that contain the identical guiding short sequences at the correct MDS-IES junctions.

Various theoretical models have been proposed in order to model the genetic unscrambling processes in ciliate organisms: the reversible guided recombination model, [10] [9], based on binary inter- and

---

*This research was supported by the Natural Sciences and Engineering Research Council of Canada Grant R2824A01, and Canada Research Chair Award to L.K.
intra-molecular DNA recombination operations; the ld, hi, dlad model, [8] 7 14, based on three unary intra-molecular DNA recombination operations; the template-guided recombination (TGR) model, [13], where a DNA molecule from the old macronucleus conducts inter-molecular DNA recombination process serving as a template; the RNA-guided DNA assembly model, [1], experimentally confirmed in [11], where either double-stranded RNA or single-stranded RNA act as templates.

This paper proposes two improvements of the descriptional complexity (size of template language, size of alphabet) of the template-guided recombination model as studied by Daley and McQuillan [2]. In [2], it has been showed that the Daley-McQuillan TGR system can generate all regular languages using iterated template-guided recombination with finite initial and template languages. For this model, the considered gene rearrangement processes take place in a stochastic style in vivo environment. In such a biological setting, it is significant to consider the size of the template language [6] 5, because sufficient copies of each template must be available throughout the recombination process. This is essential to confirm the accessibility of a template in the right place at the proper time according to the demand. Hence, the number of the unique templates should be as low as possible. The first aim of this paper is a reduction of the size of the template language by introducing a new approach to generate regular languages applying iterated TGR system with a finite initial language and a small finite set of templates.

The second aim of this paper is the reduction of the size of the template language (from regular to finite), in the extension of the template-guided recombination model called the contextual template-guided recombination system (CTGR system), [3]. Recall that a CTGR is a TGR enhanced with “deletion contexts”, the introduction of which made it possible to enhance the TGR computational power to that of Turing machines. Our reason for wanting to achieve a reduction of the template set size is the obvious one, namely that handling an infinite regular set of templates in a biological setting is impossible. To achieve our goal, we employ an additional control over the templates in the form of “permitting contexts”. We namely introduce the contextual template-guided recombination system (CTGR system) using permitting contexts as an extension of the CTGR system, and prove that an iterated version of this system has the computational power of a Turing machine, but only uses a finite initial language and a finite set of templates.

The paper is organized as follows. Section [2] introduces our new approach for generating the family of regular languages by using iterated TGR systems with \( n^2 \) templates, compared to \( n^3 \) templates in [2]. This reduction in descriptional complexity is achieved at the expense of using a filtering set to discard unintended results. Section [3] describes our proposed CTGR system using permitting contexts that, unlike CTGR systems, are able to characterize the recursively enumerable languages by using only a finite base language and a finite set of templates. This reduction in the size of the template language is achieved by introducing an additional control mechanism, the permitting context, to CTGR.

We end this introduction by some formal definitions and notations. An alphabet is a finite and nonempty set of symbols. A word or a string is a finite sequence of symbols. Let \( \Sigma \) be an alphabet. By \( \Sigma^* \) we denote the set of all words over \( \Sigma \) that includes the empty one denoted by \( \lambda \). The set of nonempty words over \( \Sigma \), i.e., \( \Sigma^* \setminus \{\lambda\} \), is denoted by \( \Sigma^+ \). The length of a word \( x \in \Sigma^* \) is denoted by \( |x| \). For \( k \in \mathbb{N} \), let \( \Sigma^{\geq k} = \{w \mid w \in \Sigma^*, |w| \geq k\} \).

For two alphabets \( \Sigma, \Delta \), a morphism is a function \( h : \Sigma^* \to \Delta^* \) satisfying \( h(xy) = h(x)h(y) \) for all \( x, y \in \Sigma^* \). A morphism \( h : \Sigma^* \to \Delta^* \) is called a coding if \( h(a) \in \Delta \) for all \( a \in \Sigma \) and a weak coding if \( h(a) \in \Delta \cup \{\lambda\} \). We denote by \( RE, CS, CF, LIN, \) and \( REG \) the families of languages generated by arbitrary, context-sensitive, context-free, linear, and regular grammars, respectively. By \( FIN \) we denote the family of finite languages. For additional formal language theory definitions and notations the reader is referred to [15].
2 TGR systems with fewer templates

This section proposes a refinement of the template-guided recombination (TGR) model as studied in the model by Daley and McQuillan [2], that is able to generate the family of regular languages by using a reduced number of templates and a smaller alphabet.

Definition 1. [2] A template-guided recombination system (or TGR system) is a four tuple $\varrho = (T, \Sigma, n_1, n_2)$ where $\Sigma$ is a finite alphabet, $T \subseteq \Sigma^*$ is the template language, $n_1$ is the minimum MDS length and $n_2$ is the minimum pointer length.

For a TGR system $\varrho = (T, \Sigma, n_1, n_2)$ and a language $L \subseteq \Sigma^*$, $\varrho(L) = \{ w \in \Sigma^* \mid (x, y) \vdash_t w \text{ for some } x, y \in L, t \in T \}$ where $(x, y) \vdash_t w$ if $x = u_1 \alpha \beta d, y = e \beta \gamma v, t = \alpha \beta \gamma v, u, v, d, e \in \Sigma^*, \alpha, \gamma \in \Sigma, \beta \in \Sigma^{\geq n_1}, \beta \in \Sigma^{\geq n_2}$. $L$ is sometimes called the base, or initial language.

Note that, if $x$ is a segment of the micronuclear DNA sequence that contains the $n$th MDS $\alpha$, and $y$ is a segment of the micronuclear DNA sequence that contains the $(n+1)$st MDS $\gamma$, then the recombination between $x$ and $y$ guided by the template $t$ will result in bringing the MDSs $n$ and $(n+1)$ in the correct order in the intermediate DNA sequence $w$, regardless of their original position in the micronuclear sequence. A sequence of such template-guided recombinations is thought to accomplish the gene unscrambling, and the transformation of the micronuclear DNA sequence in the macronuclear DNA sequence in ciliates.

For a TGR system $\varrho = (T, \Sigma, n_1, n_2)$ and a language $L \subseteq \Sigma^*$, $\varrho^*(L)$ is defined as follows:

$$\varrho^0(L) = L, \quad \varrho^{n+1}(L) = \varrho^n(L) \cup \varrho(\varrho^n(L)), \quad n \geq 0, \quad \varrho^*(L) = \bigcup_{n=0}^{\infty} \varrho^n(L).$$

If $L_1, L_2$ are two language families, then $h^*(L_1, L_2, n_1, n_2) = \{ \varrho^*(L) \mid L \in L_1, \varrho = (T, \Sigma, n_1, n_2), T \in L_2 \}$ and $h^*(L_1, L_2) = \{ h^*(L_1, L_2, n_1, n_2) \mid n_1, n_2 \in \mathbb{N} \}$.

In [2] Prop. 15, Daley and McQuillan prove that all regular languages can be generated using iterated template-guided recombination systems with finite initial and template languages, i.e., every regular language is a coding of a language in the family $h^*(\text{FIN}, \text{FIN})$. The limitation of the Daley-McQuillan model [2] is that the size of the template language and the alphabet was not meant to be optimized. Since the size of the template language will have a great impact on this type of model during in vivo computation, this is an important factor. Our aim is to reduce this number of templates. We namely introduce a new approach to generate regular languages using iterated template-guided recombination, using a finite initial language, a finite set of templates, and a weak coding. We provide a simpler construction than that of [2] Prop. 15], with fewer templates and a smaller alphabet.

Proposition 1. Each regular language $L \subseteq \Sigma^*$ can be written in the form $L = h(\varrho^*(L_0) \cap R)$, where $R$ is a regular language, $h$ is a weak coding homomorphism, $\varrho = (T, \Sigma', 1, 1)$ is a TGR system, $T$ is a finite set of templates and $L_0 \subseteq \Sigma^*$ is a finite language.

Proof. Let $L \in \text{REG}$ be generated by a regular grammar $G = (N, \Sigma, S, P)$ with the rules in $P$ of the form $X \rightarrow aY, X \rightarrow a | \lambda$, for $X, Y \in N, a \in \Sigma$. We construct a TGR system $\varrho = (T, \Sigma', n_1, n_2)$, where $n_1 = n_2 = 1$ and the alphabet $\Sigma' = N \cup \Sigma \cup \{ \# \}$. Here, $\#$ is a new symbol which assists to complete the recombination process acting as an end marker. Then we construct a finite base language $L_0 \subseteq \Sigma'$ and a template language $T \subseteq \Sigma'$ as follows.

We define the finite base language by:

$$L_1 = \{ Sa\# \mid \exists S \rightarrow a \in P, a \in \Sigma \}, \quad L_2 = \{ SaX \mid \exists S \rightarrow aX \in P, a \in \Sigma \},$$
The finite template language is defined by:

\[ T_1 = \{ aXb \mid a, b, X, Y, Z \in \Sigma, aX \rightarrow bY \in P, a \in \Sigma, b \in \Sigma \}, \]

\[ T_2 = \{ aXa \mid a, X \in \Sigma, aX \rightarrow aX \in P \}, \]

\[ T_3 = \{ aX# \mid a, X, Y, \# \in \Sigma, \exists Y \rightarrow aX \in P, \exists X \rightarrow \lambda \in P \}, \quad T = T_1 \cup T_2 \cup T_3. \]

Note that for example, \( L_4 \subseteq L_3, L_2 \subseteq L_3, \) and \( T_2 \subseteq T_1. \) However, we made these separations for the purpose of the clarity of the proof.

In order to eliminate all non terminals and the new symbol, we consider the weak coding \( h \) defined by \( h(X) = \lambda, \) for any \( X \in N, h(a) = a, \) for any \( a \in \Sigma, \) \( h(\#) = \lambda. \) Moreover, we consider the language \( R = \{ S \}(\Sigma N)^*\{\#, \#\}, \) whose purpose is to ensure that only strings of the correct form will be accepted, by removing other unintended strings.

We claim that \( L = h(g^*(L_0) \cap R). \)

For the "\( \subseteq \)" inclusion, in order to obtain a valid derivation in \( G \) and to continue recombinations, we consider the string \( SaX \) from group \( L_2 \) as the first string in the recombinations. At this stage, through recombination, the application of the rules of the form \( X \rightarrow bY \in P \) can be achieved as follows. During the recombination, a string \( XbY \) from group \( L_3 \) as the second string can be recombined with the string \( SaX, \) and an appropriate template \( aXb \) from group \( T_1, \) can be used to produce the string \( SaXbY \) which is of the form \( \{ S \}(\Sigma N)^* \), with \( X, Y \in N \) and \( a, b \in \Sigma. \) By using only the templates from group \( T_1, \) the simulation of the rules of the form \( X \rightarrow bY \in P \) is possible, because no other template from group \( T_2, \) \( T_3 \) can be used. The simulation is as follows:

\[ (SaX, XbY) \vdash_{aXb} SaXbY. \]

The above mentioned simulation process can be repeated an arbitrary number of times according to the templates in group \( T_1. \) Likewise, rules of the form \( X \rightarrow aX \in P \) can be simulated that produce the string \( SaXaX \) using a template from \( T_2 \) and considering the second string from group \( L_4 \) as follows:

\[ (SaX, XaX) \vdash_{aXa} SaXaX. \]

The application of the rules of the form \( X \rightarrow aX \in P \) can also be simulated repeatedly. In general, for representing an intermediate recombination, if \( u, v \in g^*(L_0) \) illustrates derivations of \( G \) and \( u = u' aY, \) \( v = Yb'v', \) where \( u' \in \{ S \}(\Sigma N)^*, \quad v' \in (N \Sigma)^*N, \) \( a \in \Sigma, \) \( b \in \Sigma, \) then the resulting recombined string \( (u, v) \vdash_{aYb} u' aYb'v' \) that is a string of the form \( \{ S \}(\Sigma N)^* \) can be generated which corresponds to an intermediate computation of the form \( S \Rightarrow^* \delta N \) in \( G \) where \( \delta \in \Sigma^* \).

If a template finds more than one matching point in the first string, then the template can attach to any of those points and a matching second string from \( L_3 \) as guided by the template can be recombined with the first string. For example, such a recombination can happen to a string of the form

\[ Sa_1 X_i a_2 X_2 \ldots a_i X_i a_{i+1} X_{i+1} \ldots a_{k-1} X_{k-1} a_k X_k. \]

Along this string if \( a_i X_i = a_k X_k \) for some \( 1 \leq i \leq k, \) then the recombination guided by a template \( a_i X_i b \) can take place either at the matching position \( a_i X_i \) or at the matching position \( a_k X_k \) between the above first string and the second string of the form \( X_i bY. \) This recombination will produce the resulting string either of the form \( Sa_1 X_1 a_2 X_2 \ldots a_i X_i bY \) or of the form \( Sa_1 X_1 a_2 X_2 \ldots a_i X_i a_{i+1} X_{i+1} \ldots a_{k-1} X_{k-1} a_k X_k bY, \) respectively. Note however that any recombination that does not happen at the rightmost end of the sentential form has only the effect of "resetting" the derivation a few steps backward.
Thus, without loss of generality, we will hereafter assume that any derivation that results in a terminal word has an equivalent rightmost derivation. We will only discuss these rightmost derivations.

Note also that recombinations can proceed in parallel, for example, a recombination can take place between a string of the form $Sa_1X_1a_2X_2\ldots a_{n-1}X_{n-1}a_n#$ and the corresponding template from the group $L_5$ and applying the matching template $a_{n-1}X_{n-1}a_n#$ from group $T_3$. After recombination, the produced string is $Sa_1X_1a_2X_2\ldots a_{n-1}X_{n-1}a_n# = w'$ which is of the form $\{S\}(\Sigma N)^{\ast}\{\#\}$ and corresponds to our intended terminal word. At this point, any further recombination at the right most end of this produced terminal string stops because no matching template can be found in the finite set of templates $T$ to guide recombination with this string:

$$(Sa_1X_1a_2X_2\ldots a_{n-1}X_{n-1}a_n#) \vdash a_{n-1}X_{n-1}a_n Sa_1X_1a_2X_2\ldots a_{n-1}X_{n-1}a_n#.$$

Moreover, for simulating a rule of the form $X_{n-1} \rightarrow \lambda$, the required second string is from group $L_6$ and the corresponding template from the group $T_3$. Recombination yields a string $Sa_1X_1a_2X_2\ldots a_{n-1}X_{n-1}#\#$ of the form $\{S\}(\Sigma N)^{\ast}\{\#\}$ that is the terminal string and further recombination cannot take place:

$$(Sa_1X_1a_2X_2\ldots a_{n-1}X_{n-1}#\#) \vdash a_{n-1}X_{n-1}# Sa_1X_1a_2X_2\ldots a_{n-1}X_{n-1}#.$$

By construction, it is clear that each string in $\rho^\ast(L_0)$ corresponds to a derivation in $G$, and the simulation of a derivation is possible only by using recombinations according to the corresponding template from the finite template language $T$. Accordingly, each derivation in $G$ of the form

$$S \Rightarrow a_1X_1 \Rightarrow^* \ldots \Rightarrow a_1a_2\ldots a_kX_k \Rightarrow a_1a_2\ldots a_ka_{k+1}X_{k+1} \Rightarrow^* \ldots$$

$$a_1a_2\ldots a_{n-1}X_{n-1} \Rightarrow a_1a_2\ldots a_n = w,$$

where $1 \leq k \leq n$, $X_k \rightarrow a_{k+1}X_{k+1} \in P$, $X_{n-1} \rightarrow a_n \in P$, corresponds to a computation in $\varrho$ of the form

$$(Sa_1X_1,a_1X_1a_2X_2) \vdash a_1X_1a_2X_2 \Rightarrow^* \ldots Sa_1X_1a_2X_2\ldots a_kX_ka_{k+1}X_{k+1} \Rightarrow^* \ldots Sa_1X_1a_2X_2\ldots a_kX_ka_{k+1}X_{k+1}a_{n-1}X_{n-1}a_n# = w',$$

or

$$Sa_1X_1a_2X_2\ldots a_kX_ka_{k+1}X_{k+1}a_{n-1}X_{n-1}# = w'.$$

Therefore, we can say from the above description that a terminal string according to the grammar $G$ is achievable only by starting the recombination with a string that begins with the start symbol $S$ (that means considering as the first string a string containing the start symbol $S$ at the beginning) and then proceeding by a series of recombination processes according to the appropriate templates from the finite template language $T$ for an arbitrary number of times, which end up with the end marker $#$ and simulate a derivation according to $G$. Afterwards, intersecting the language $R = \{S\}(\Sigma N)^{\ast}\{\#,\#\}$ with the set of generated strings, we obtain our intended terminal strings. In this way, we are able to find a string...
$w' \in \varrho^*(L_0) \cap R$ and then the application of the weak coding $h(w') = w \in \Sigma^*$ allows us to obtain the exact string generated by a derivation in $G$. Thus, every derivation in $G$ can be simulated.

Hence, we obtain $L \subseteq h(\varrho^*(L_0) \cap R)$. The other inclusion follows because the only recombinations that can happen according to $\varrho$ lead either to words that are eliminated by the filter, or to words in $L$ after applying the weak coding.

Let us now compare the size of the template language we obtained with that of the Daley-McQuillan model \cite{2}. The Daley-McQuillan model \cite{2} requires three production rules to construct a template based on their definition of the template language in the following. $T = \{[X,a,Y][Y,b,Z][Z,c,W]\}$ where $[X,a,Y],[Y,b,Z],[Z,c,W] \in V$, $V$ is an alphabet and $X \rightarrow aY, Y \rightarrow bZ, Z \rightarrow cW \in P$. If the number of production rules in the grammar is $|P| = n$, then based on this definition the template language has a cardinality of $n^3$.

Our construction requires two production rules to construct a template. In the worst case we can have $n^2$ templates where $n$ is the number of production rules in the simulated grammar. In addition to the size of templates, the size of the TGR alphabet $\Sigma'$ in our construction is small: one plus the number of terminals and nonterminals in the simulated grammar. In the Daley-McQuillan model as described above, the alphabet $V$ can be much larger, and it also depends on the number of productions of the grammar. Although we require fewer templates and alphabet, our model has one limitation, i.e., it requires a filter to discard unintended results, while the Daley-McQuillan model requires only the correct recombination to occur according to the constructed matching templates.

### 3 CTGR systems with permitting contexts

As shown in \cite{3,4,2}, the finiteness of the initial language and the set of templates restricts the computational power of a TGR system. In fact, even with a regular initial language and a regular set of templates, iterated TGR systems can generate at most regular languages \cite{2}.

Daley and McQuillan \cite{3} have added a new feature called “deletion context” to enhance the computational power of template-guided recombination. Their extension of the TGR system is called the contextual template-guided recombination system (CTGR system). In \cite{3}, it was shown that arbitrary recursively enumerable languages can be generated by iterated CTGR with a regular set of templates and a finite initial language, with the help of taking intersection with the Kleene star of the terminal alphabet, and a coding. From a practical viewpoint, dealing with a regular set of templates is not realistic in the sense that we cannot manage an infinite “computer”.

To achieve the finiteness of the employed component sets while preserving the computational power of CTGR, we impose an additional control on the templates in order to restrict their usage. More precisely, we associate each template with a set of “permitting contexts”: strings that must appear as subwords within the two participating words if this particular template is to be used for their recombination. The idea of permitting contexts has been previously used in the context of splicing systems, a formal model of DNA recombination that uses restriction enzymes and ligases \cite{12}.

**Definition 2.** A contextual template-guided recombination system (CTGR system) using permitting contexts is a quadruple $\varrho_p = (T, \Sigma, n_1, n_2)$, where $\Sigma$ is a finite alphabet, $n_1 \in \mathbb{N}$ is the minimum MDS length and $n_2 \in \mathbb{N}$ is the minimum pointer length, $T$ is a set of triples (templates using permitting contexts) of the form $t_p = (t; C_1, C_2)$ with $t = e_1#\alpha\beta\gamma#d_1$ being a template over $\Sigma$ and $C_1, C_2$ being finite subsets of $\Sigma^*$. To such a triple $t_p$ we associate the word

$$\tau(t_p) = e_1#\alpha\beta\gamma#d_1$a_1$&$a_2$&$b_1$&$b_2$&$b_m$$,
where $C_1 = \{a_1, \ldots, a_k\}$, $C_2 = \{b_1, \ldots, b_m\}$, $k, m \geq 0$ and $\$, \&, \# are new special symbols not included in $\Sigma$. We define $\tau(T) = \{\tau(t_p) \mid t_p \in T\}$.

For a CTGR system using permitting contexts $g_p = (T, \Sigma, n_1, n_2)$ and a language $L \subseteq \Sigma^*$, we define $g_p(L) = \{w \in \Sigma^* \mid (x, y) \vdash c \ w \text{ for some } x, y \in L \text{ and } t_p \in T\}$, where $(x, y) \vdash c \ w$ if and only if $x = u\alpha\beta d, y = e\gamma v, t_p = (e_1 \# \alpha \gamma \# d_1, a_1, \ldots, a_k, b_1, \ldots, b_m), w = u\alpha\beta\gamma v, u, v, d, e \in \Sigma^*, \alpha, \gamma \in \Sigma^{\geq n_1}, \beta \in \Sigma^{\geq n_2}$. Every element that belongs to $C_1$ appears as a substring in $x$ and every element that belongs to $C_2$ appears as a substring in $y$, i.e., $a_i \in \text{sub}(x)$ for $1 \leq i \leq k$, $b_j \in \text{sub}(y)$ for $1 \leq j \leq m$; moreover, if $C_1 = \{\lambda\}$ or $C_2 = \{\lambda\}$, then we assume that no constrain is imposed on $x$ and $y$ respectively.

For a CTGR system using permitting contexts $g_p = (T, \Sigma, n_1, n_2)$ and a language $L \subseteq \Sigma^*$, a template language $T$, we can define an iterated version of $g_p^*(L)$ similarly as for TGR systems.

The following proposition shows that iterated CTGR system using permitting contexts can generate arbitrary recursively enumerable languages using a finite initial language and a finite set of templates with the help of intersection with a filter language and, at last, applying a weak coding homomorphism.

**Proposition 2.** Every recursively enumerable language $L \subseteq \Sigma^*$ can be written in the form $L = h(L' \cap L_1)$, where $h$ is a weak coding homomorphism, $L_1$ is a regular language and $L' = g_p^*(L_0)$ with $L_0$ a finite language.

**Proof.** Consider a Chomsky type-0 grammar $G = (N, \Sigma, S, P)$ in Kuroda normal form, where $L(G) = L$ and the production rules in $P$ are of the forms $A \rightarrow EC, AE \rightarrow CD, A \rightarrow a \mid \lambda$ for $A, C, D, E \in N, a \in \Sigma$. Let us denote $U = N \cup \Sigma \cup \{B, B_1, B_2\}$, where $B, B_1, B_2$ are new symbols.

We then construct a CTGR system using permitting contexts $g_p = (T, V, 1, 1)$ where $V = N \cup \Sigma \cup \{B, B_1, B_2, X, X', Y, Z, Z'\} \cup \{Y_b \mid b \in U\}$ and $T$ contains the following templates using permitting contexts:

1. Simulate : $Z\#cavY\#uY; \{X\}, \{\lambda\}, \text{for } a, c \in U, Z, Y \in V, u \to v \in P$.
2. Rotate : $Z\#caY_b\#bY; \{X\}, \{\lambda\}, \text{for } a, b, c \in U, Z, Y \in V$.
3. $X\#X'\#d\#e\#Z; \{\lambda\}, \{Y_b\}, \text{for } b, d, e \in U, Z, X \in V$.
4. $Z\#caY\#Y_b; \{X'\}, \{\lambda\}, \text{for } Z, Y, Y_b \in V$.
5. $X\#Xac\#Z; \{\lambda\}, \{Y\}, \text{for } X', X, Z \in V$.
6. Terminate : $XXB_1B_2\#abc\#Z'; \{\lambda\}, \{Y\}, \text{for } X, Z' \in V, B, B_1, B_2 \in U$.

We define the following languages which are included in the initial finite language $L_0 \subseteq V^*$:

$L_1 = \{XBB_1B_2SY\}, L_2 = \{ZavY \mid a \in U, u \to v \in P\}$,
$L_3 = \{ZaY_b \mid a \in U\}, L_4 = \{X'baZ \mid b, a \in U\}$,
$L_5 = \{ZaY \mid a \in U\}, L_6 = \{XaZ \mid a\}, L_7 = \{abZ' \mid a, b \in U\}$.

We denote $L_0 = L_1 \cup L_2 \cup L_3 \cup L_4 \cup L_5 \cup L_6 \cup L_7$, and $L_0$ acts as the initial language.

For the construction of this system, the idea we use is the well-known proof technique, “rotate-and-simulate procedure”, which was effectively used in other contexts [12] in order to allow the simulation of a rule that applies to a symbol in the middle of the word by first moving that symbol to the right hand end of the word, simulating the rule, and returning the result to its original place.

Throughout this construction we assume $x$ and $y$ to be, respectively, the first word and the second word of the recombination as defined in Definition 2.

The starting of the simulation based on the derivation steps in $G$ requires to consider the word $XBB_1B_2SY$ as the first word. Indeed, any other choice of start word leads to derivations of words of illegal form (not in $XBB_1B_2\Sigma^*Y$). Throughout the derivation steps this word is bordered by $X$ or its variant $X'$ at the left end, as well as by $Y$ or its variant $Y_b, b \in U$ at the right end, $X, X'$ and $Y, Y_b$ make...
the left respectively right extremity of the word. Likewise, the symbol $B$ always signals the beginning of
the word, i.e., the sentential forms of $G$, which facilitates the permutation of the word and $B_1, B_2$ are
included to provide the contexts for recombination.

Note that all the templates with permitting contexts in $T$ include symbols $Z$ or $Z'$ that have thus
to be present in one of the two words taking part in the recombination. Furthermore, the words contain-
ing symbols $Z$ and $Z'$ are from the initial language $L_0$ but will not appear in the resulting word of
recombination. This guarantees that each recombination has to happen between the current word which
is produced in the previous recombination and at least one word from $L_0$. The simulation of a derivation
in $G$ initiates with the application of template 1 to $XBB_1B_2SY \in L_1$ and $ZavY \in L_2$, where initially
$w = BB_1B_2S$, $S \rightarrow v \in P$ and $a \in U$. The word obtained through the recombination is $XBB_1B_2vY$:

$$(XBB_1B_2SY, ZBB_2vY) \vdash_{t_p} XBB_1B_2vY$$

for $t_p = (Z#cavY #uY; \{X\}, \{\lambda\}) = (ZB_1B_2vY #SY; \{X\}, \{\lambda\})$, where $S \rightarrow v \in P$, $c, a \in U$ and
$w = BB_1B_2S$.

Generally, considering a word $X_1BB_1B_2xvY$ and $u \rightarrow v \in P$, the resulting word will be
$X_1BB_1B_2xvY$ applying the associated templates from group 1. Here, $w = w_1cavu = x_1BB_1B_2xu$. This
simulates a derivation step $x_2ux_1 \Rightarrow x_2vx_1$ in $G$. The derivation is as follows:

$$(Xw_1cavY, ZavY) \vdash_{t_p} Xw_1cavY$$

for $t_p = (Z#cavY #uY; \{X\}, \{\lambda\})$, where $u \rightarrow v \in P$, $c, a \in U$.

In this simulation step, no other templates from groups 2 - 6 can be applied except the templates from
group 1 because of imposed restriction as deletion contexts and permitting contexts on the usage of the
templates. Afterwards, we come to the rotation process that is necessary so as to move symbols from the
right hand end of the current word to the left hand end. This rotation process can be explained by the
following steps:

Step 1: We can start the rotation process using the corresponding template from group 2 with a
word $XwbY$, where $b \in (N \cup \Sigma)^*$, $w \in (N \cup \Sigma)^*(BB_1B_2)(N \cup \Sigma)^*$ (respectively $XwbY$, where $b \in
\{B, B_1, B_2\}, w \in (N \cup \Sigma)^*$). In this step, $x = XwbY = Xw_1cabY$, $y = ZyaY \in L_3$:

$$(Xw_1cabY, ZaYb) \vdash_{t_p} Xw_1caYb$$

for $t_p = (Z#caYb #bY; \{X\}, \{\lambda\})$, where $wb \in (N \cup \Sigma)^*(BB_1B_2)(N \cup \Sigma)^*, b \in N \cup \Sigma \cup \{B, B_1, B_2\}$.

Step 2: After applying the template from group 2 in Step 1, we obtained the word $Xw_1caYb$, which
we rewrite as $Xdew_2Yb$ where $w = w_1ca = dew_2$. Then we continue the rotation process using the
matching template from group 3 with $x = X'bdZ \in L_4$, $y = Xdeuw_2Yb$:

$$(X'bdZ, Xdeuw_2Yb) \vdash_{t_p} X'bdew_2Yb$$

for $t_p = (X'X'bd#Z; \{\lambda\}, \{Y_0\})$, where $b \in N \cup \Sigma \cup \{B, B_1, B_2\}$.

Step 3: The resulting word from the previous step is of the form $X'bdew_2Yb$, which can be written
of the form $X'wYb = Xw_3caYb$. In this step we will apply the matching template from group 4 where
$x = X'w_3caYb$, $y = ZaY \in L_5$:

$$(X'w_3caYb, ZaY) \vdash_{t_p} X'w_3caY$$

for $t_p = (Z#caY #yb; \{X'\}, \{\lambda\})$ where $c, a \in N \cup \Sigma \cup \{B, B_1, B_2\}$.
Step 4: The recombined word from Step 3 is $X^t w_3 ca Y$, in general, the outcome of step 3 is a word of the form $X^t acw_4 Y$. Lastly, we complete the rotation process by using a template from group 5 where $x = XaZ \in L_6$, $y = X^t acw_4 Y$:

$$(XaZ, X^t acw_4 Y) \vdash_{t_p} Xacw_4 Y$$

for $t_p = (X^t \# X ac \# Z; \{\lambda\}, \{Y\})$ where $a, c \in N \cup \Sigma \cup \{B, B_1, B_2\}$.

The above mentioned rotation-steps produced the word $Xbacw_4 Y = XbwY$ which implies that starting from the word $XwY$ and applying steps 1 - 4, we achieve the word $XbwY$ having the same end markers. In this way, we are able to move the symbol $b$ from the right-hand end to the left-hand end of the word that accomplishes the rotation of the underlying sentential form. These rotation-steps can be repeated an arbitrary number of times and thus provide every circular permutation of the word flanked by $X$ and $Y$.

Using a template from group 1 to each word of the form $XwY$ when $w$ ended by the left hand part of a rule in $P$, it is possible to simulate the application of all rules of $P$ at a desired position corresponding to the sentential form of $G$, by means of the four rotation-steps.

It is observed that from the initial word $XBB_1B_2SY$, each produced word in every step does not include the symbols $Z, Z'$, that is, the word is of the form $\alpha_1x_1BB_1B_2x_2\alpha_2$ in which the pair $(\alpha_1, \alpha_2)$ is one of the four pairs $(X, Y), (X, Y_b), (X', Y_b), (X', Y), b \in U$. In fact, these symbols being present in the templates of $T$ serve as permitting contexts that restrict the regulation of the recombination process of this system $\varrho_p$.

Now we come to the termination process. Applying the terminating template from group 6, we can remove $XBB_1B_2$ only when $Y$ is present and the symbols $B, B_1, B_2$ together as a word $BB_1B_2$ is adjacent to $X$. Here, $x = abZ' \in L_7$, $y = Xacw_4 Y = XBB_1B_2bcw_5 Y$:

$$(abZ', XBB_1B_2bcw_5 Y) \vdash_{t_p} abcw_5 Y$$

for $t_p = (XBB_1B_2\# abc\# Z'; \{\lambda\}, \{Y\})$, where $w \in (N \cup \Sigma)^* \{BB_1B_2\}(N \cup \Sigma)^*$, $b, a, c \in N \cup \Sigma \cup \{B, B_1, B_2\}$.

Now our achieved word is of the form $abcw_5 Y = wY = w' \in \varrho_p^*(L_0)$, i.e., $L' = \varrho_p^*(L_0)$. The intersection operation with the language $L_1 = \Sigma^* Y$ will filter out the words that are not in proper form. Furthermore, we define a weak coding homomorphism which eliminate the right end marker $Y$ leaving other letters unchanged. Let us now define a weak coding homomorphism by $h(a) = a$, for any $a \in \Sigma$, $h(Y) = \lambda$.

Thus, we obtain a word in $\Sigma^*$ by applying the weak coding homomorphism where $w \in h(L' \cap L_1)$. Finally, from the above construction we can produce each word in $L(G)$ and we say that $L(G) \subseteq h(\varrho_p^*(L_0) \cap \Sigma^* Y)$. Conversely, the opposite inclusion is held by this system. Therefore, $h(\varrho_p^*(L_0) \cap \Sigma^* Y) \subseteq L(G)$.

\[\Box\]

4 Conclusions

This paper improves on the descriptional complexity (size of the template language) from $n^3$ to $n^2$ in the case of template-guided recombination (TGR) systems, and from regular to finite in the case of contextual template-guided recombination (CTGR) systems. These reductions are obtained at the expense of using a filtering language in the case of TGR, and of an additional control (permitting contexts) in the case of CTGR.
References

[1] A. Angeleska, N. Jonoska, M. Saito & L. Landweber (2007): RNA-guided DNA assembly. Journal of Theoretical Biology 248(4), pp. 706–720.

[2] M. Daley & I. McQuillan (2005): Template guided DNA recombination. Theoretical Computer Science 330(2), pp. 237–250.

[3] M. Daley & I. McQuillan (2006): On Computational Properties of Template-Guided DNA Recombination. Lecture Notes in Computer Science 3892, pp. 27–37.

[4] M. Daley & I. McQuillan (2006): Useful Templates and Iterated Template-Guided DNA Recombination in Ciliates. Theory of Computing Systems 39(5), pp. 619–633.

[5] Michael Domaratzki (2008): Equivalence in template-guided recombination. Natural Computing 7(3), pp. 439–449.

[6] Michael Domaratzki (2009): Minimality in template-guided recombination. Information and Computation 207(11), pp. 1209–1220.

[7] T. Harju, I. Petre & G. Rozenberg (2003): Gene Assembly in Ciliates: Molecular operations. Technical Report 557, Turku Centre for Computer Science.

[8] T. Harju & G. Rozenberg (2003): Computational Processes in Living Cells: Gene Assembly in Ciliates. Lecture Notes in Computer Science 2450, pp. 1–20.

[9] L. Kari, J. Kari & L. Landweber (1999): Reversible Molecular Computation in Ciliates. In: Jewels are Forever, Contributions on Theoretical Computer Science in Honor of Arto Salomaa, Springer-Verlag, London, UK, pp. 353–363.

[10] L. Kari & L. Landweber (1999): Computational Power of Gene Rearrangement. 5th DIMACS Workshop on DNA Based Computers, pp. 207–216.

[11] M. Nowacki, V. Vijayan, Y. Zhou, K. Schotanus, T. Doak & L. Landweber (2007): RNA-mediated epigenetic programming of a genome-rearrangement pathway. Nature 451(7175), pp. 153–158.

[12] Gheorghe Paun, Grzegorz Rozenberg & Arto Salomaa (1998): DNA Computing: New Computing Paradigms. Springer.

[13] D. Prescott, A. Ehrenfeucht & G. Rozenberg (2003): Template-guided recombination for IES elimination and unscrambling of genes in stichotrichous ciliates. Journal of Theoretical Biology 222(3), pp. 323–330.

[14] D. M. Prescott, A. Ehrenfeucht & G. Rozenberg (2001): Molecular operations for DNA processing in hypotrichous ciliates. Europ. J. Protistol. 37, pp. 241–260.

[15] Arto Salomaa (1973): Formal Languages. Academic Press, New York.