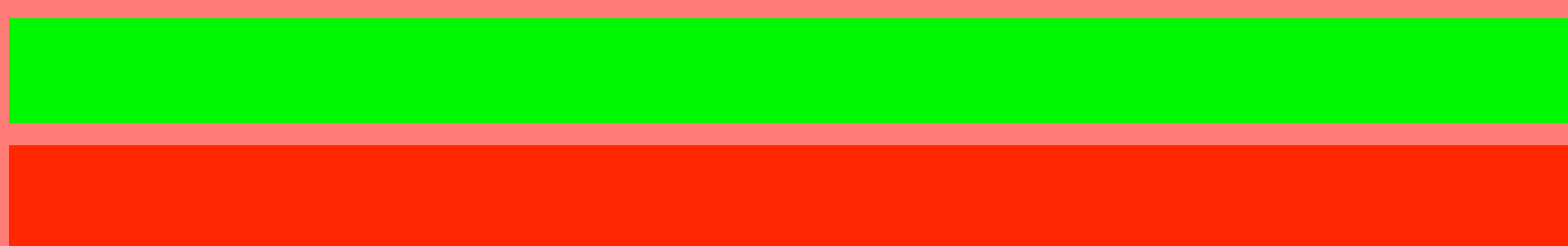
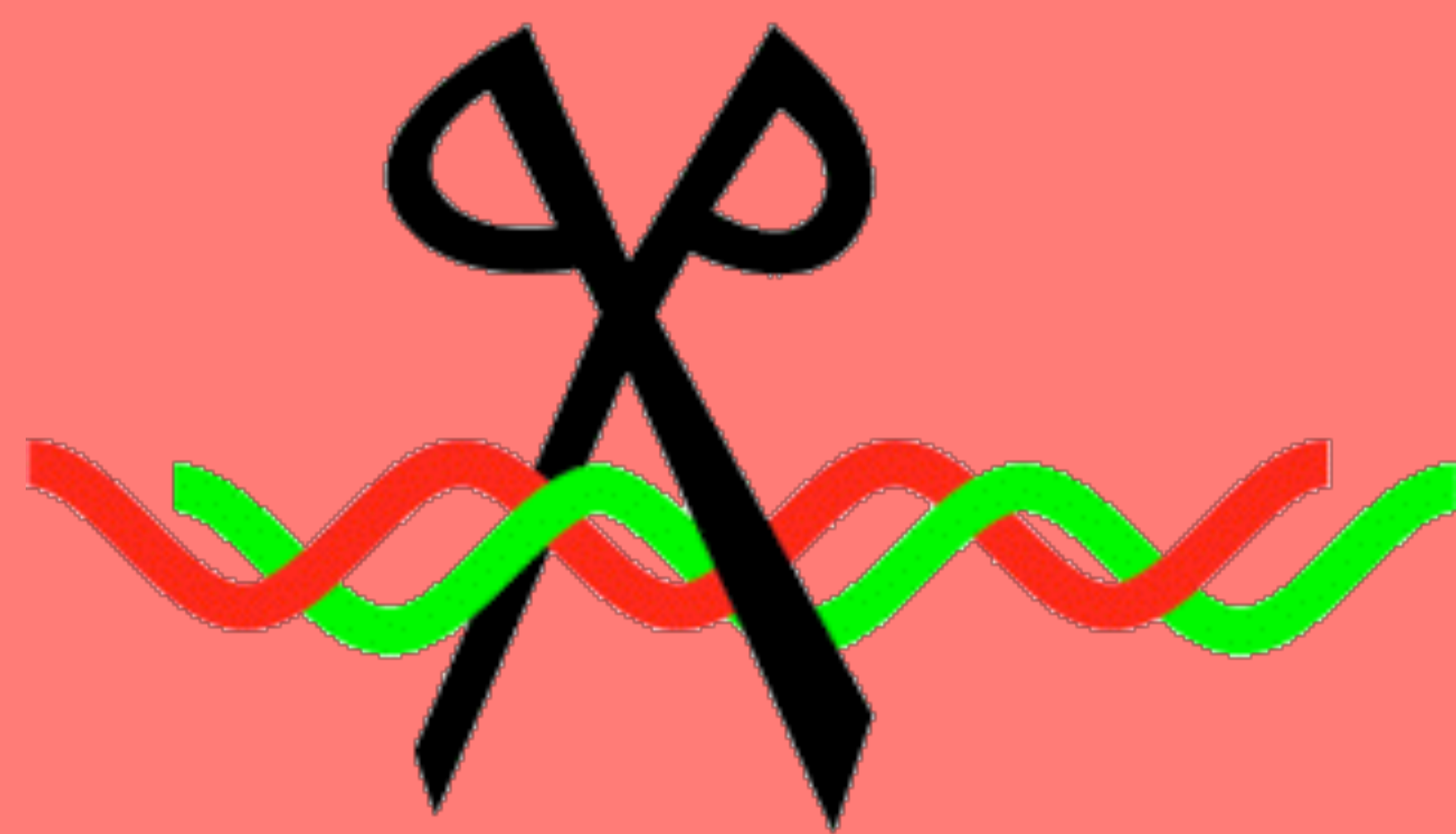


The Asian Mathematical Conference 2013


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IDEA

 double-stranded DNA (dsDNA)



The cutting process of dsDNA by restriction enzyme

 The pasting process of dsDNA by the presence of ligase

EXAMPLES OF SECOND ORDER LIMIT LANGUAGE

Example 1

Let $S = (A, I, R)$ be a Y-G splicing system consisting of two restriction enzymes namely *FauI* and *AciI*, where $A = \{a, c, g, t\}$, $I = \{\alpha cccgcttaacg\beta\}$ such that $\alpha, \beta \in A^*$ and $R = \{r_1 : r_2\}$ where $r_1 = (cccgcttaa; cg, 1)$ and $r_2 = (c; cg, c)$.

The following are the splicing languages after the first splicing has taken place,

$$I \cup \left\{ \begin{array}{l} \alpha cccgcttaacgcttaagcggg\alpha', \alpha cccgcttaacgcttaagcggg\alpha', \beta'cg\beta, \\ \alpha cccggg\alpha', \beta'cgtaagcgttaacg\beta, \alpha cccgcttaacgcttaacg\beta, \\ \alpha cccg\beta, \alpha cccgcttaacg\alpha', \beta'cgtaagcgc\beta \end{array} \right\}.$$

The second order limit language are

$$\{\alpha cccgcttaacgcttaacgcttaacg\beta, \alpha cccgcttaacgcttaacg\alpha'\}.$$

Example 2

Let $S = (A, I, R)$ be a Y-G splicing system consisting of a restriction enzyme namely *MboI*, where $A = \{a, c, g, t\}$, $I = \{aagatcgccgatcttct\}$ which consists of two recognition sites of the restriction enzyme and $R = \{(1; gatc, 1 : 1; gatc, 1)\}$.

The following are the splicing languages after the first splicing has taken place,

$$I \cup \left\{ \begin{array}{l} aagatctt, aagatcttct, aagatcgccgatctt, \\ aagatcgccgatcgccgatcttct, aagatcgccgatctt, \\ aagatcgccgatcgccgatctt, aagatcgccgatcgccgatcttct, \\ aagatcgccgatcgccgatcttct \end{array} \right\}.$$

The second order limit language are

$$\left\{ \begin{array}{l} aagatcgccgatcgccgatcgccgatcttct, \\ aagatcgccgatcgccgatcgccgatcttct, \\ aagatcgccgatcgccgatcgccgatcttct, \\ aagatcgccgatcgccgatcgccgatcgccgatcttct. \end{array} \right\}.$$

INTRODUCTION

DNA molecules are known of its functions which are coding for proteins synthesis and also self-replication that ensure an exact copy is passed on to the offspring cell [1]. These molecules are made up of thousands of complementary nucleotides commonly referred to as Adenine (A), Guanine (G), Cytosine (C) and Thymine (T). By Watson-Crick complementarity [2], A is paired with T and C is paired with G and vice versa. Then, those pairs are presented as a , g , c and t . There is an enzyme that works beautifully with DNA molecules known as the restriction enzyme.

In this research, the definition of second order limit language is given and an example is discussed to show the existence of second order limit language. Once the formation and existence of second order limit language is shown, its characteristics are illustrated by some theorems.

MAIN RESULTS

Theorem 1

If the rule of a splicing system is itself palindromic, then there will be no second order limit language.

Theorem 2

An initial string that contains two recognition sites of two rules with identical crossing sites produces second order limit language.

Corollary 1

If only an initial string and a rule is involved in a splicing system, then the second order limit language does not exist.

PRELIMINARIES

Definition 1 [3]: Y-G Splicing System

If $r \in R$, where $r = (u, x, v; y, x, v)$ and $s_1 = \alpha u x v \beta$ and $s_2 = \gamma y x z \delta$ are elements of I , then splicing s_1 and s_2 using r produces the initial string I together with $\alpha u x z \beta$ and $\gamma y x v \beta$, presented in either order where $\alpha, \beta, \gamma, \delta, u, x, v, y$ and $z \in A^*$ are the free monoid generated by A with the concatenation operation and 1 as the identity element.

Definition 2 [4]: Second Order Limit Language

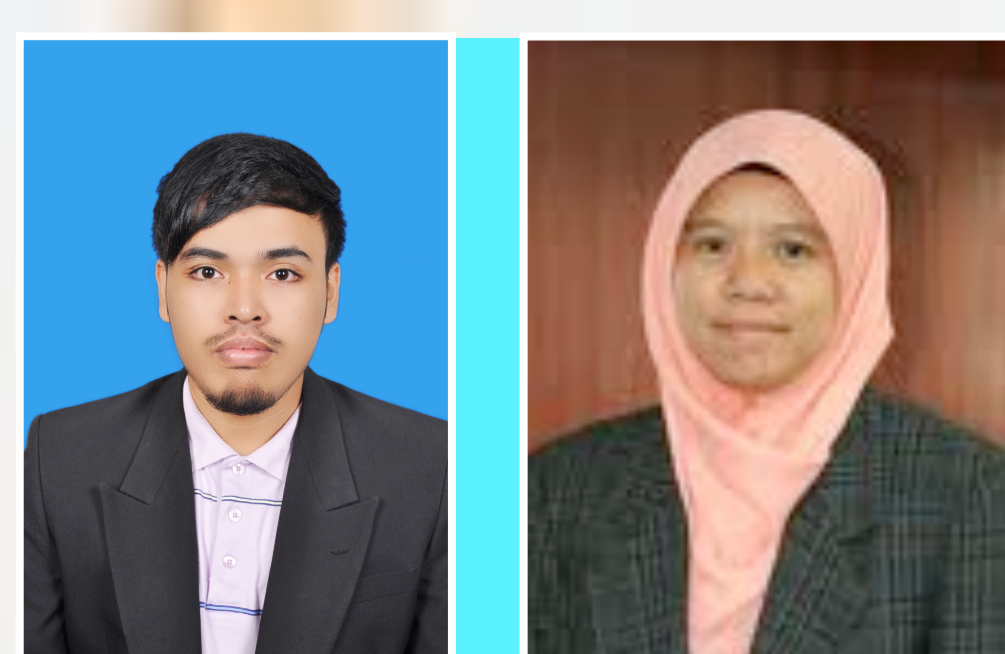
Let L_1 be the set of second order limit words of L , the set L_2 of second order limit words of L to be the set of first order limits of L_1 . We obtain L_2 from L_1 by deleting the words that are transient in L_1 .

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Muhammad Azrin Ahmad¹, Nor Haniza Sarmin², Fong Wan Heng³ and Yuhani Yusof⁴
^{1,2}Department of Mathematical Sciences, Faculty of Science,
³Ibnu Sina Institute for Fundamental Science Studies,
^{1,2,3,4}Applied Algebra Analysis Group (A³G), Nanotechnology Research Alliance Universiti Teknologi Malaysia, 81310 UTM Johor Bahru, Johor.
⁴Faculty of Industrial Science and Technology, Universiti Malaysia Pahang, 26300 UMP Gambang, Pahang.
 azrinahmad1@gmail.com¹, nhs@utm.my², fwh@ibnusina.utm.my³, yuhani@ump.edu.my⁴

