# On the Comparative Study of Some Mathematical Tools for Specific Sequences Design 

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#### Abstract

In modern communication system, cryptography and automatic test patterns, some specific sequences with strictly defined properties are required to meet the application demands. $[1,2,3,4 \ldots]$ These requirements are:


- Good pseudorandom (PN) properties (large period length, uniform distrbution...).
- Low periodic correlation property.
- Low aperiodic correlation property.
- Large linear complexity.
- Large cardinality (number of sequences in the set).

Unfortunately, there is not any set of sequence satisfying all these requirements, despite the fact that a lot of efforts have been given for design such sequences. For this purpose, different mathematical tools have been widely used such as: matrix, d-transform and trace function representations. However, to the best of our knowledge, there has been no any comparative study on these mathematicals tools carried out so far!
In this contribution, we try to fill some parts of this gap by considering some typical applications of these tools. The paper is constructed as below:

- In the introduction, the basic concepts and definitions of matrix, d-transform and trace are given (briefly).
- In section II, some typical applications (for demonstration) will be shown. In this regard, we will give some discussions and suggestions for choosing the appropriate mathematical tools for each application.
- Especially, in section III, we will show the relationship (interchanging or equivalencies) between them. Keywords: specific sequences, mathematical tool, matrix, d-transform and trace.


## 1. Introduction

### 1.1 Matrix and State-machine representation of LFSR

In this section, two typical configurations of linear feedback shift register for the same output i.e., the Galois and Fibonacci, are considered and showed as Fig 1.1a and Fig 1.1b, which can be used to generate the m-sequence for mobile communication and cryptography application [5,6]. The state at the time instant n is the contents of the shift register represented by a column vector $\mathrm{S}_{\mathrm{n}}$ :

$$
\begin{equation*}
S_{n}=\left[S_{0, \mathrm{n}} \mathrm{~S}_{1, \mathrm{n}} \ldots \mathrm{~S}_{\mathrm{m}-2, \mathrm{n}} \mathrm{~S}_{\mathrm{m}-1, \mathrm{n}}\right]^{\mathrm{T}} \tag{1}
\end{equation*}
$$

Let consider the Fig 1.1a first. Given the contents of the LFSR at the time instant $n$, then the content of the shift register at instant ( $\mathrm{n}+1$ ) can be obtained:

$$
\begin{align*}
& \mathrm{S}_{0, \mathrm{n}+1}=\mathrm{S}_{1, \mathrm{n}}+\mathrm{g}_{1} \mathrm{~S}_{0, \mathrm{n}} \\
& \mathrm{~S}_{1, \mathrm{n}+1}=\mathrm{S}_{2, \mathrm{n}}+\mathrm{g}_{2} \mathrm{~S}_{0, \mathrm{n}} \\
& \quad \vdots  \tag{2}\\
& \mathrm{~S}_{\mathrm{m}-2, \mathrm{n}+1}=\mathrm{S}_{\mathrm{m}-1, \mathrm{n}}+\mathrm{g}_{\mathrm{m}-1} \mathrm{~S}_{0, \mathrm{n}} \\
& \mathrm{~S}_{\mathrm{m}-1, \mathrm{n}+1}=\mathrm{g}_{\mathrm{r}} \mathrm{~S}_{0, \mathrm{n}}
\end{align*}
$$

Where addition and multiplication are $\bmod 2$ and $\mathrm{g}_{\mathrm{i}} \in\{0,1\}$ for $\mathrm{i}=\{1,2, \ldots, \mathrm{~m}\}$. These equations can be written in matrix form as:

$$
\mathrm{S}_{\mathrm{n}+1}=\left[\begin{array}{c}
\mathrm{S}_{0, \mathrm{n}+1}  \tag{3}\\
\mathrm{~S}_{1, \mathrm{n}+1} \\
\mathrm{~S}_{2, \mathrm{n}+1} \\
\vdots \\
\mathrm{~S}_{\mathrm{m}-1, \mathrm{n}+1}
\end{array}\right]=\left[\begin{array}{ccccc}
\mathrm{g}_{1} & 1 & 0 & \ldots & 0 \\
\mathrm{~g}_{2} & 0 & 1 & \ldots & 0 \\
\mathrm{~g}_{3} & 0 & 0 & \ldots & 0 \\
\vdots & \vdots & \vdots & \vdots & \vdots \\
\mathrm{~g}_{\mathrm{m}} & 0 & 0 & \ldots & 0
\end{array}\right] \times\left[\begin{array}{c}
\mathrm{S}_{0, \mathrm{n}} \\
\mathrm{~S}_{1, \mathrm{n}} \\
\mathrm{~S}_{2, \mathrm{n}} \\
\vdots \\
\mathrm{~S}_{\mathrm{m}-1, \mathrm{n}}
\end{array}\right]
$$

which can be written for short:

$$
\begin{equation*}
\mathrm{S}_{\mathrm{i}}=\mathrm{G}^{\mathrm{i}} \mathrm{~S}_{0} \text { for } \mathrm{i}=\{1,2, \ldots, \mathrm{n}\} \tag{4}
\end{equation*}
$$

Where G is a square Galois state transition matrix of order $\mathrm{m}, \mathrm{S}_{0}$ is the vector of initial state.

The output $b_{n}$ of the shift register at the given instant $n$ (the content of the right most shift register) can be determined as:

$$
\mathrm{b}_{\mathrm{n}}=\left[\begin{array}{lllll}
1 & 0 & 0 & \ldots & 0 \tag{5}
\end{array}\right] \times \mathrm{G}^{\mathrm{n}} \times \mathrm{S}_{0}
$$

Which presents the relationship between output sequence and the state transition matrix and the initial state $\mathrm{S}_{0}$ of the LFSR.
For the shift register configuration in Fig 1.1b, a similar result may be obtained in the same way as:

$$
\begin{equation*}
S_{n+1}=F \times S_{n} \tag{6}
\end{equation*}
$$

Where:

$$
\mathrm{F}=\left[\begin{array}{cccccc}
0 & 1 & 0 & 0 & \cdots & 0  \tag{7}\\
0 & 0 & 1 & 0 & \cdots & 0 \\
\vdots & & & & & \vdots \\
0 & 0 & 0 & 0 & \cdots & 1 \\
g_{m} & g_{m-1} & g_{m-2} & g_{m-3} & \cdots & g_{1}
\end{array}\right]
$$

And the output $b_{n}$ at the instant of time $n$ is:

$$
\mathrm{b}_{\mathrm{n}}=\left[\begin{array}{llll}
\mathrm{g}_{\mathrm{m}} & \mathrm{~g}_{\mathrm{m}-1} & \cdots & \mathrm{~g}_{1} \tag{8}
\end{array}\right] \times \mathrm{F}^{\mathrm{n}} \times \mathrm{S}_{0}
$$



Figure 1.1a. Galois feedback generator.


Figure 1.1b. Fibonaci feedback generator.

### 1.2 D-transform

The D-transform of a sequence $\left\{\mathrm{b}_{\mathrm{n}}\right\}$ over $\operatorname{GF}(\mathrm{p})$ (Galois filed (mod p ), a primitive integer) is denoted by $\mathrm{D}\left[\mathrm{b}_{\mathrm{n}}\right]$ [4,7,8,9]:

$$
\begin{equation*}
\mathrm{D}\left[\mathrm{~b}_{\mathrm{n}}\right]=\sum_{\mathrm{i}=0}^{\mathrm{m}} \mathrm{~b}_{\mathrm{i}} \mathrm{D}^{\mathrm{i}}, \mathrm{~b}_{\mathrm{i}} \in\{\mathrm{GF}(\mathrm{p})\} \tag{9}
\end{equation*}
$$

Thus, the D-transform of the sequence will have the form of a polynomial in d over $\mathrm{GF}(\mathrm{p})$ and has been conveniently used in signal and system analysis in data transmission and CDMA [8,9].
The D-transform of the generator sequence $\left\{b_{n}\right\}$ of a linear feedback shift register (LFSR) is then given by:

$$
\begin{equation*}
\mathrm{b}(\mathrm{D})=\frac{\mathrm{S}(\mathrm{D})}{\mathrm{g}(\mathrm{D})} \tag{10}
\end{equation*}
$$

Where $g(D)$ of degree $n$ is the generating polynomial of a LFSR and $S(D)$ of degree $\leq n-1$ specifies the initial condition corresponding to a particular shifted version of $\left\{b_{n}\right\}$.
1.3 Representation the m-sequence in term of the power of primitive elements (Trace function)

Let $\alpha$ be the root of the primitive polynomial $g(D)$ of the degree $m$. The trace function of $\alpha$ is mapping from $\mathrm{GF}\left(\mathrm{q}^{\mathrm{m}}\right)$ to $\mathrm{GF}(\mathrm{q})$ and is defined as:

$$
\begin{equation*}
\mathrm{T}_{\mathrm{r}} \mathrm{q}_{\mathrm{q}}^{\mathrm{m}}(\alpha)=\sum_{\mathrm{i}=0}^{\mathrm{m}-1} \alpha^{\mathrm{q}} \tag{11}
\end{equation*}
$$

In other words, the trace of $\alpha$ over $\mathrm{GF}(\mathrm{q})$ is the sum of conjugate of $\alpha$ with respect to $\mathrm{GF}(\mathrm{q})$.

## 2. Application cases

2.1 Matrix representation

### 2.1.1 State and configure description

As clearly seen in 1.1 previous section, the matrix presentation may be best suitable to describle FSR in term of Finite State Machine (FSM). It shows the close relations in the set of four element $\left\{\mathrm{S}_{\mathrm{n}}, \mathrm{S}_{\mathrm{n}+1}, \mathrm{~F}, \mathrm{~b}_{\mathrm{n}}\right\}$ which defines the FSM.
Observing the fig 1.1a, 1.1b one can easily get elements of the feedback tap $g_{n}$, state vector $S_{n}$ and vice versa. In word: matrix presentation is no doubt the most intuitive.
2.1.2 Interleaved structure representation

It is well known that the specific sequences must have very large length. Except for few Mersenne primitive integers, the natutally posses interleaved structure which draws a lot of attentions in the literature $[1,2,3,4,10,11 \ldots]$. For this case, the matrix representation can offer a very intuitive approach in represent the interleaved structure $[4,10,11 \ldots]$ but it may look complicated due to the great length of those sequences.
Decomposition of m-sequence $\left\{b_{n}\right\}$ and rearrange it Matrix form [3,4,12]. It is well known that if we decimate $\left\{b_{n}\right\}$ by $T$, we obtain:

$$
\begin{equation*}
\left\{\mathrm{a}_{\mathrm{n}}\right\}=\left\{\mathrm{b}_{\mathrm{nT}}\right\}=\operatorname{Tr}_{1}^{\mathrm{n}}\left(\alpha^{\operatorname{Tn}}\right)=\operatorname{Tr}_{0}^{\mathrm{n}}\left(\beta^{\mathrm{n}}\right) \tag{12}
\end{equation*}
$$

With $\beta=\alpha^{\mathrm{T}}$. Since $\alpha^{\mathrm{T}}$ is primitive element in $\operatorname{GF}\left(2^{\mathrm{m}}\right)$, $\left\{\mathrm{a}_{n}\right\}$ is also an m -sequence of length $\mathrm{N}=\left(2^{\mathrm{n}}-1\right) / \mathrm{T}$. When the decimation start at the first bit, we will obtain the subsequence: $\left\{\mathrm{a}_{0}, \mathrm{a}_{\mathrm{T}}, \ldots, \mathrm{a}_{\left(2^{\mathrm{m}}-2\right) \mathrm{T}}\right\}$
Similarly, we'll obtain the subsequence $\left\{\mathrm{a}_{\mathrm{t}-1}, \mathrm{a}_{\mathrm{T}+\mathrm{t}-1}, \ldots, \mathrm{a}_{\left(2^{\mathrm{m}}-2\right) \mathrm{T}+\mathrm{t}-1}\right\}$ when the decimation start at the $\mathrm{t}^{\text {th }}$-bit. Thus, on the time-domain, these subsequence (arrange in columns) can be considered as T time-multiplexed sequences $\left\{\mathrm{a}_{\mathrm{nT}}\right\}\left\{\mathrm{a}_{\mathrm{nT}+1}\right\} \ldots\left\{\mathrm{a}_{(\mathrm{n}+1) \mathrm{T}-1}\right\}$ onto T time slots:

$$
\underline{\underline{M}}=\left[\begin{array}{cccc}
a_{0} & a_{1} & \cdots & a_{T-1}  \tag{13}\\
a_{T} & a_{\mathrm{T}+1} & \cdots & a_{2 \mathrm{~T}-1} \\
\vdots & \vdots & & \vdots \\
a_{\left(2^{\mathrm{m}}-2\right) \mathrm{T}} & a_{\left(2^{\mathrm{m}}-2\right) \mathrm{T}+1} & \cdots & a_{\left(2^{\mathrm{m}}-1\right) \mathrm{T}-1}
\end{array}\right]=\left\{a_{\mathrm{nT}}\right\}\left\{a_{\mathrm{nT}+1}\right\} \ldots\left\{a_{(\mathrm{n}+1) \mathrm{T}-1}\right\}
$$

This representation is closely related to the time multiplex technique, which well understood by engineers.

Example 1: Decomposition of the sequence of 1023 bits long into 33 subsequences, each of length 31: 100100100110100110101111100110001 111100100011101111110000111000000 011111111110001110001001110110010 101110111101010001111010010101000 001011111111010101010111101000011 101001000110010110101100111101011 000110011111100101010100110011001 010011111010011100001000110110010 001010011011110111010101110011001 110111011100111010100111010000011 110110111000011000100101001011001 101000100010110100101110100110001 011000000101001001011111011110001 100011011101100001111001001110010 110001000011011111110011100011010 100101000010000100101101111101011 100010111001000011111011010101000 101111011001110011111000001110010 010101100101111001011100000101011 011001100001101011011101000101011 111101000111001101110010100011010 000001100100100010000010011011010 011110011010101100001011101101000 110000100111111101110001111000000 111011011000101000100110010000011 010010011110111110001010101101000 010100000001011011011110011110001 000111111011000111010110101000011 001101100000110000000011011011010 111010111100001010100100001011001 001100000100010010000001000000000
Based on the phase shifts between the subsequences represented by the columns and define the Null sequence by $\infty$ we can easily get the interleaving order $[4,9,12]$
$\mathrm{I}_{\mathrm{p}}^{\mathrm{T}}=\{12,21,1,29,6,17,0,6,6,2,4,16,21,7,22,4,29,16,13,16,15,8,24,12,4,5,25,15,10,18, \infty, 17,8\}$
Note:

- This sequences is still too short for practical application.
- One can get the interleaving order $\mathrm{I}_{\mathrm{p}}^{\mathrm{T}}$ (shift sequence e) only after knowing the sequences.


### 2.2 D-transform representaion

2.2.1 State and configure description

Refer to (10) we see that in D-transform the information about configuration $g(D)$, state $S(D)$ and signal $b(D)$ are given. It is short and effective to handle the relation between these three parameters. If $S(D)$ and $g(D)$ are known, we can find out the corresponding output sequence by the long division algorithm [5,13] as in example 2.
Example 2: Apply the long division algorithm to determinate shift register output when $\mathrm{S}(\mathrm{D})=1, \mathrm{~g}(\mathrm{D})=1+\mathrm{D}+\mathrm{D}^{4}$

## Solution:

The shift register output is:
$b(D)=\frac{1}{1+D+D^{4}}$
$b(D)=1+D+D^{2}+D^{3}+D^{5}+D^{7}+D^{8}+\ldots$ and $b_{n}=1,1,1,1,0,1,0,1,1, \ldots$
2.2.2 Interleaved structure representation

Since interleaving process and D-transform are both sort of time multiplexing [1, 3, 4, 9..] one can easily derive the interleaving order $\mathrm{I}_{\mathrm{p}}^{\mathrm{T}}$ straightforwardly from D-transform of the sequences. In fact, there are two methods for derive $\mathrm{I}_{\mathrm{p}}^{\mathrm{T}}$, namely: expanding and decomposition. For the sake of simplicity we just show the result as in example 3.

Example 3: Let $\mathrm{m}=3, \mathrm{n}=6$ and let $\alpha$ be a primitive element of $\operatorname{GF}\left(2^{6}\right)$ with primitive polynomial $\mathrm{b}(\mathrm{D})=\mathrm{D}^{6}+\mathrm{D}^{5}+1$ over $\operatorname{GF}(2)$. Let $\left\{b_{n}\right\}$ denote the $m$-sequence generated by $b(D)$
$\left\{b_{n}\right\}=\{011111101.010110011 .011101101 .001001110 .001011110 .010100011 .0$ $00010000\}$.
Decimation of $\left\{b_{n}\right\}$ by $T=9$, we obtain $\left\{a_{n}\right\}=\left\{b_{n 9}\right\}$ and rearrange is as:
$\mathrm{M}=\left[\begin{array}{lllllllll}0 & 1 & 1 & 1 & 1 & 1 & 1 & 0 & 1 \\ 0 & 1 & 0 & 1 & 1 & 0 & 0 & 1 & 1 \\ 0 & 1 & 1 & 1 & 0 & 1 & 1 & 0 & 1 \\ 0 & 0 & 1 & 0 & 0 & 1 & 1 & 1 & 0 \\ 0 & 0 & 1 & 0 & 1 & 1 & 1 & 1 & 0 \\ 0 & 1 & 0 & 1 & 0 & 0 & 0 & 0 & 1 \\ 0 & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 0\end{array}\right]$
We can see that the columns are shift equivalent and $\mathrm{I}_{\mathrm{p}}^{\mathrm{T}}=\{\infty, 5,3,5,6,3,3,2,5\}$, where $\infty$ represents Null sequence.
2.2.3 Hardware implementation of interleaving

D-transform can offer this representation, since it is time multiplexing from nature! It is clear that all msubsequences are shift versions generated by same LFSR. Therefore, the software controlled switch S just picks up these shift versions and deliver them in the order determined by $\mathrm{I}_{\mathrm{p}}^{\mathrm{T}} \ldots$. This approach have been successfully have been successfully simulated and implemented in [14,15,16].


Figure 2. Shift versions for interleaving

### 2.3 Trace function representation

### 2.3.1 Field structure and sequences

As seen in the definition, this is also very short and compact representation: the information about the extension Field and subfield is given (mapping between them). One can derive the sequence from it trace function (see example 4).
Example 4: The binary m-sequence $\left\{\mathrm{a}_{\mathrm{n}}\right\}$ with period $2^{\mathrm{m}}-1$ can be represented through the trace function as:
$\left\{\mathrm{a}_{\mathrm{n}}\right\}=\left\{\mathrm{a}_{0}, \mathrm{a}_{1}, \mathrm{a}_{2}, \ldots, \mathrm{a}_{2 \mathrm{~m}-1}\right\}=\operatorname{Tr}_{1}^{\mathrm{m}}\left(\alpha^{0}\right), \operatorname{Tr}_{1}^{\mathrm{m}}\left(\alpha^{1}\right), \ldots, \operatorname{Tr}_{1}^{\mathrm{m}}\left(\alpha^{2^{\mathrm{m}}-1}\right)=\operatorname{Tr}_{1}^{\mathrm{m}}\left(\alpha^{\mathrm{n}}\right)$
It is well known that, the shift version of sequence $\left\{a_{n}\right\}$ can be created bu its decimation.
The sequence $\left\{\mathrm{a}_{\mathrm{kn}}\right\}$ obtained by sampling every $\mathrm{k}^{\text {th }}$ bit of $\left\{\mathrm{a}_{\mathrm{n}}\right\}$, starting from the first bit is referred to as decimation of $\left\{a_{n}\right\}$. A decimation by $k$ of $\left\{a_{n}\right\}$ results in the sequence: $\left\{a_{n k}\right\}=\operatorname{Tr}_{1}^{m}\left(\alpha^{n}\right)$.
Let $\left\{a_{n}\right\}$ be $m$-sequence generator by $g(D)=D^{5}+D^{2}+1$ over $G F(2)$.
The corresponding binary from is: 10010.11001 .11110 .00110 .11101 .010000
Below are few sequences result from decimation of $\left\{a_{n}\right\}$ by difference $k$ [17]

| k | Sequence | Characteristic Polynomial (in octal) |
| :--- | :--- | :--- |
| 1 | 1001011001111100011011101010000 | 45476156500 |
| 3 | 111101110001010110100001100100 | 76705320620 |
| 5 | 1110100010010101100001110011011 | 72112607154 |
| 7 | 1001001100001011010100011101111 | 44605521674 |
| 11 | 1110110011100001101010010001011 | 73160651054 |
| 15 | 100001010111011000111100110100 | 41273076320 |

2.3.2 Interleaved structure representation

By nature, trace function shows the "interleaving" of subfileds inside the extension field. Therefore it is convenient to describle the interleaved structure.
Base on trace function one can easily derive the interleaving order directly as in example 5.
Example 5: Let trace function from $\mathrm{GF}\left(3^{4}\right)$ onto $\mathrm{GF}\left(3^{2}\right)$ with primitive polynomial 10021: $\mathrm{f}(\mathrm{x})=\mathrm{x}^{4}+2 \mathrm{x}+1$ with $\mathrm{n}=4$, $\mathrm{L}=3^{4}-1=80, \mathrm{~m}=2, \mathrm{~N}=3^{2}-1=8, \mathrm{~S}=\mathrm{L} / \mathrm{N}=10$.
Calcualting the trace function of x from $\mathrm{GF}\left(3^{4}\right)$ into $\mathrm{GF}\left(3^{2}\right)$ we obtained:
$\operatorname{Tr}_{\mathrm{m}}^{\mathrm{n}}(\alpha)=\sum_{\mathrm{k}=0}^{\mathrm{n} / \mathrm{m}-1} \alpha^{2 \mathrm{k}}=\alpha+\alpha^{9}$
$\alpha^{\text {Si }}$ table:
$\mathrm{i}=0 \Rightarrow \alpha^{0}=1$
$\mathrm{i}=1 \Rightarrow \alpha^{10}=1+2 \alpha+\alpha^{2}+\alpha^{3}$
$\mathrm{i}=2 \Rightarrow \alpha^{20}=\alpha+2 \alpha^{2}+2 \alpha^{3}$
$\mathrm{i}=3 \Rightarrow \alpha^{30}=1+\alpha+2 \alpha^{2}+2 \alpha^{3}$
$\mathrm{i}=4 \Rightarrow \alpha^{40}=2$
$\mathrm{i}=5 \Rightarrow \alpha^{50}=2+\alpha+2 \alpha^{2}+2 \alpha^{3}$
$\mathrm{i}=6 \Rightarrow \alpha^{60}=2 \alpha+\alpha^{2}+\alpha^{3}$
$\mathrm{i}=7 \Rightarrow \alpha^{70}=2+2 \alpha+\alpha^{2}+\alpha^{3}$
With j run from 0 to $\mathrm{S}-1$ we get:

$$
\begin{aligned}
& \mathrm{j}=0 \Rightarrow \operatorname{Tr}\left(\alpha^{0}\right)=\operatorname{Tr}(1)=1+1=2=\alpha^{40} \Rightarrow \mathrm{I}_{\mathrm{P}}^{0}=4 \\
& \mathrm{j}=1 \Rightarrow \operatorname{Tr}\left(\alpha^{1}\right)=\alpha+\alpha^{9}=2 \alpha+\alpha^{2}+\alpha^{3}=\alpha^{60} \Rightarrow \mathrm{I}_{\mathrm{P}}^{1}=6 \\
& \mathrm{j}=2 \Rightarrow \operatorname{Tr}\left(\alpha^{2}\right)=\alpha^{2}+\alpha^{18}=2 \alpha+\alpha^{2}+\alpha^{3}=\alpha^{60} \Rightarrow \mathrm{I}_{\mathrm{P}}^{2}=6 \\
& \mathrm{j}=3 \Rightarrow \operatorname{Tr}\left(\alpha^{3}\right)=\alpha^{3}+\alpha^{27}=\alpha+2 \alpha^{2}+2 \alpha^{3}=\alpha^{20} \Rightarrow \mathrm{I}_{\mathrm{P}}^{3}=2 \\
& \mathrm{j}=4 \Rightarrow \operatorname{Tr}\left(\alpha^{4}\right)=\alpha^{4}+\alpha^{36}=2+\alpha+2 \alpha^{2}+2 \alpha^{3}=\alpha^{50} \Rightarrow \mathrm{I}_{\mathrm{P}}^{4}=5 \\
& \mathrm{j}=5 \Rightarrow \operatorname{Tr}\left(\alpha^{5}\right)=\alpha^{5}+\alpha^{45}=0 \Rightarrow \mathrm{I}_{\mathrm{P}}^{5}=\infty \\
& \mathrm{j}=6 \Rightarrow \operatorname{Tr}\left(\alpha^{6}\right)=\alpha^{6}+\alpha^{54}=\alpha+2 \alpha^{2}+2 \alpha^{3}=\alpha^{20} \Rightarrow \mathrm{I}_{\mathrm{P}}^{6}=2 \\
& \mathrm{j}=7 \Rightarrow \operatorname{Tr}\left(\alpha^{7}\right)=\alpha^{7}+\alpha^{63}=1=\alpha^{0} \Rightarrow \mathrm{I}_{\mathrm{P}}^{7}=0 \\
& \mathrm{j}=8 \Rightarrow \operatorname{Tr}\left(\alpha^{8}\right)=\alpha^{8}+\alpha^{72}=2+\alpha+2 \alpha^{2}+2 \alpha^{3}=\alpha^{50} \Rightarrow I_{P}^{4}=5 \\
& \mathrm{j}=9 \Rightarrow \operatorname{Tr}\left(\alpha^{9}\right)=\alpha^{9}+\alpha^{81}=2 \alpha+\alpha^{2}+\alpha^{3}=\alpha^{60} \Rightarrow I_{P}^{1}=6
\end{aligned}
$$

The interleaving order is:
$I_{p}^{T}=\{4,6,6,2,5, \infty, 2,0,5,6\}$
The subsequences can be interleaved like this [3]
$M=\left[\begin{array}{llllllllll}1 & 0 & 0 & 0 & 1 & 0 & 0 & 2 & 1 & 0 \\ 1 & 1 & 1 & 2 & 0 & 0 & 2 & 2 & 0 & 1 \\ 0 & 2 & 2 & 1 & 1 & 0 & 1 & 0 & 1 & 2 \\ 1 & 2 & 2 & 1 & 2 & 0 & 1 & 2 & 2 & 2 \\ 2 & 0 & 0 & 0 & 2 & 0 & 0 & 1 & 2 & 0 \\ 2 & 2 & 2 & 1 & 0 & 0 & 1 & 1 & 0 & 2 \\ 0 & 1 & 1 & 2 & 2 & 0 & 2 & 0 & 2 & 1 \\ 2 & 1 & 1 & 2 & 1 & 0 & 2 & 1 & 1 & 1\end{array}\right]$
Even though trace function is very effective, it is abstract (pure mathematic) and one cannot expect it to be intuitive. Note: It has shown that the interleaving order (or shift sequence) calculated by D-transform and by trace representation are identical for the same composite sequence [4,9,12].

## 3. The links between above mentioned tools

### 3.1 Between Matrix and D-transform

In order to determinate $\mathrm{S}(\mathrm{D})$ (in D-transform) which specifies the corresponding memory content of LFSR we have use the matrix equation system. This fact can be best demonstrated by an example. Unfortunately the matrix representation for specific sequences which have very large length would take a lot of spaces, let alone the manipulation with these matrices. Therefore we consider here only the subsequence (shorter length) $\mathrm{P}_{\mathrm{n}}$ generated by:

$$
g(D)=1+D^{5}+D^{6}
$$

Example 6: The D-transform of $\mathrm{P}_{\mathrm{n}}$ is $\mathrm{W}_{\mathrm{N}}(\mathrm{D})$ with $\mathrm{N}=2^{\mathrm{m}}-1=63$, so, the length of $\mathrm{P}_{\mathrm{n}}$ is then=63 and $\mathrm{W}_{\mathrm{i}}\left(\mathrm{d}^{\mathrm{T}}\right)=\frac{\mathrm{S}_{\mathrm{i}}\left(\mathrm{d}^{\mathrm{T}}\right)}{\mathrm{g}_{\mathrm{s}}\left(\mathrm{d}^{\mathrm{T}}\right)}$
With T denotes the number of subsequences in the specific sequence.
The corresponding $\mathrm{S}_{\mathrm{i}}\left(\mathrm{d}^{\mathrm{T}}\right)$ of memory state: $\mathrm{S}_{\mathrm{i}}$ can be calculated as:
$\left(\begin{array}{c}\mathrm{p}_{0} \\ \mathrm{p}_{1} \\ \vdots \\ \mathrm{p}_{\mathrm{N}-1}\end{array}\right)=\left(\begin{array}{cccc}1 & 0 & \ldots & 0 \\ \mathrm{w} 1 & 1 & \ldots & 0 \\ \vdots & \vdots & \ddots & \vdots \\ { }^{\mathrm{w}} & \mathrm{N}-1 & { }^{w_{N}}-2 & \cdots \\ { }^{2}\end{array}\right)\left(\begin{array}{c}\mathrm{g}_{0} \\ \mathrm{~g}_{1} \\ \vdots \\ \mathrm{~g}_{\mathrm{N}-1}\end{array}\right)$
And $\left[g_{i}\right]$ denote the feedback taps of the LFSR characterized by $g(D)$.

Table $i$ the corresponding phase shifts of sequences specified by $\mathrm{S}_{\mathrm{i}}\left(\mathrm{d}^{T}\right)$

| phase shifts | $\mathrm{S}_{\mathrm{i}}\left(\mathrm{d}^{\mathrm{T}}\right)$ | phase shifts | $\mathrm{S}_{\mathrm{i}}\left(\mathrm{d}^{\text {T }}\right.$ ) |
| :---: | :---: | :---: | :---: |
| 0 | $1+\mathrm{D}^{5}$ | 32 | $1+\mathrm{D}^{3}+\mathrm{D}^{5}$ |
| 1 | $\mathrm{D}^{5}$ | 33 | $\mathrm{D}^{3}+\mathrm{D}^{5}$ |
| 2 | $\mathrm{D}^{4}$ | 34 | $\mathrm{D}+\mathrm{D}^{4}$ |
| 3 | $\mathrm{D}^{3}$ | 35 | $1+\mathrm{D}^{3}$ |
| 4 | $\mathrm{D}^{2}$ | 36 | $\mathrm{D}^{2}+\mathrm{D}^{4}+\mathrm{D}^{5}$ |
| 5 | D | 37 | $\mathrm{D}+\mathrm{D}^{3}+\mathrm{D}^{5}$ |
| 6 | 1 | 38 | $1+\mathrm{D}^{2}+\mathrm{D}^{3}$ |
| 7 | $\mathrm{D}^{4}+\mathrm{D}^{5}$ | 39 | $\mathrm{D}+\mathrm{D}^{2}+\mathrm{D}^{4}+\mathrm{D}^{5}$ |
| 8 | $\mathrm{D}^{3}+\mathrm{D}^{4}$ | 40 | $1+\mathrm{D}+\mathrm{D}^{3}+\mathrm{D}^{4}$ |
| 9 | $\mathrm{D}^{2}+\mathrm{D}^{3}$ | 41 | $1+D^{2}+D^{3}+D^{4}+D^{5}$ |
| 10 | $\mathrm{D}+\mathrm{D}^{2}$ | 42 | $\mathrm{D}+\mathrm{D}^{2}+\mathrm{D}^{3}+\mathrm{D}^{5}$ |
| 11 | 1+D | 43 | $1+\mathrm{D}+\mathrm{D}^{2}+\mathrm{D}^{4}$ |
| 12 | $1+\mathrm{D}^{4}+\mathrm{D}^{5}$ | 44 | $1+\mathrm{D}+\mathrm{D}^{3}+\mathrm{D}^{4}+\mathrm{D}^{5}$ |
| 13 | $\mathrm{D}^{3}+\mathrm{D}^{5}$ | 45 | $1+\mathrm{D}^{2}+\mathrm{D}^{3}+\mathrm{D}^{5}$ |
| 14 | $\mathrm{D}^{2}+\mathrm{D}^{4}$ | 46 | $\mathrm{D}+\mathrm{D}^{2}+\mathrm{D}^{5}$ |
| 15 | $\mathrm{D}^{1}+\mathrm{D}^{3}$ | 47 | $1+\mathrm{D}+\mathrm{D}^{4}$ |
| 16 | $1+\mathrm{D}^{2}$ | 48 | $1+D^{3}+D^{4}+D^{5}$ |
| 17 | $\mathrm{D}+\mathrm{D}^{4}+\mathrm{D}^{5}$ | 49 | $\mathrm{D}^{2}+\mathrm{D}^{3}+\mathrm{D}^{5}$ |
| 18 | $1+\mathrm{D}^{3}+\mathrm{D}^{4}$ | 50 | $\mathrm{D}+\mathrm{D}^{2}+\mathrm{D}^{4}$ |
| 19 | $\mathrm{D}^{2}+\mathrm{D}^{3}+\mathrm{D}^{4}+\mathrm{D}^{5}$ | 51 | $1+\mathrm{D}+\mathrm{D}^{3}$ |
| 20 | $\mathrm{D}+\mathrm{D}^{2}+\mathrm{D}^{3}+\mathrm{D}^{4}$ | 52 | $1+D^{2}+D^{4}+D^{5}$ |
| 21 | $1+\mathrm{D}+\mathrm{D}^{2}+\mathrm{D}^{3}$ | 53 | $\mathrm{D}+\mathrm{D}^{3}+\mathrm{D}^{5}$ |
| 22 | $1+\mathrm{D}+\mathrm{D}^{2}+\mathrm{D}^{4}+\mathrm{D}^{5}$ | 54 | $1+D^{2}+D^{4}$ |
| 23 | $1+\mathrm{D}+\mathrm{D}^{3}+\mathrm{D}^{5}$ | 55 | $\mathrm{D}+\mathrm{D}^{3}+\mathrm{D}^{4}+\mathrm{D}^{5}$ |
| 24 | $1+D^{2}+D^{5}$ | 56 | $1+\mathrm{D}^{2}+\mathrm{D}^{3}+\mathrm{D}^{4}$ |
| 25 | $\mathrm{D}+\mathrm{D}^{5}$ | 57 | $\mathrm{D}+\mathrm{D}^{2}+\mathrm{D}^{3}+\mathrm{D}^{4}+\mathrm{D}^{5}$ |
| 26 | $1+\mathrm{D}^{4}$ | 58 | $1+\mathrm{D}+\mathrm{D}^{2}+\mathrm{D}^{3}+\mathrm{D}^{4}$ |
| 27 | $\mathrm{D}^{3}+\mathrm{D}^{4}+\mathrm{D}^{5}$ | 59 | $1+\mathrm{D}+\mathrm{D}^{2}+\mathrm{D}^{3}+\mathrm{D}^{4}+\mathrm{D}^{5}$ |
| 28 | $\mathrm{D}^{2}+\mathrm{D}^{3}+\mathrm{D}^{4}$ | 60 | $1+\mathrm{D}+\mathrm{D}^{2}+\mathrm{D}^{3}+\mathrm{D}^{5}$ |
| 29 | $\mathrm{D}+\mathrm{D}^{2}+\mathrm{D}^{3}$ | 61 | $1+\mathrm{D}+\mathrm{D}^{2}+\mathrm{D}^{5}$ |
| 30 | $1+\mathrm{D}+\mathrm{D}^{2}$ | 62 | $1+\mathrm{D}+\mathrm{D}^{5}$ |
| 31 | $1+\mathrm{D}+\mathrm{D}^{4}+\mathrm{D}^{5}$ |  |  |

Note that T may be few thousands or more to create the specific sequences used in the practice!
On the other hand from $g(D)$ we can easily get the feedback tap of fig $1.1 \mathrm{a}, 1.1 \mathrm{~b}$ and therefore $\mathrm{G}_{\mathrm{i}}$ of (4), the state transition matrix $F$ of (7). If $S(D)$ is known, then we can get the output $b_{n}$.

### 3.2 Between D-transform and trace

The equivalence can be proven as follows:
Let consider the Decomposition of $m$, sequences $\left\{b_{n}\right\}$
Let $\left\{b_{n}\right\}$ be an $m$-sequence generated by $g(D)$ of degree $n=\operatorname{lm}$, and denote $T=\left(2^{n}-1\right) /\left(2^{m}-1\right)$
From $\left\{b_{n}\right\}$ sequences, we try to find out $\left\{a_{n}\right\}$ and $I_{p}^{T}$
Decimate $\left\{b_{n}\right\}$ by T, we will obtain: $\left\{a_{n}\right\}=\operatorname{Tr}_{1}^{\mathrm{n}}\left(\alpha^{\mathrm{Tn}}\right)=\operatorname{Tr}_{0}^{\mathrm{n}}\left(\beta^{\mathrm{n}}\right)$
Since $\left(\alpha^{\mathrm{Tn}}\right)$ and $\left(\beta^{\mathrm{n}}\right)$ are primitive element in $\operatorname{GF}\left(2^{\mathrm{n}}\right),\left\{\mathrm{a}_{\mathrm{n}}\right\}$ is also an m-sequence of length N with $\mathrm{N}=\left(2^{\mathrm{n}}-1\right) / \mathrm{T}$.
Let arrange $\left\{a_{n}\right\}$ in the column of the decimation matrix
$M=\left|\begin{array}{cccc}a_{0} & a_{1} & \ldots & a_{\mathrm{T}-1} \\ a_{\mathrm{T}} & a_{\mathrm{T}+1} & \ldots & a_{2 \mathrm{~T}-1} \\ a_{2 \mathrm{~T}} & a_{2 \mathrm{~T}+1} & \ldots & \\ a_{\left(2^{\mathrm{m}}-2\right) \mathrm{T}} & a_{\left(2^{\mathrm{m}}-2\right) \mathrm{T}+1} & \ldots & a_{\left(2^{\mathrm{m}}-1\right) \mathrm{T}-1}\end{array}\right|$
Each column is a decimation of $\left\{b_{n}\right\}$ by T, an m-subsequence [17].
On other hand, from d-transform property: $S\left(d^{T}\right) / g\left(d^{T}\right)$ represents a decimations of $b_{n}$ by $T=\left(2^{\mathrm{n}}-1\right) /\left(2^{\mathrm{m}}-1\right)$, which turn out to be $\left\{\mathrm{a}_{\mathrm{n}}\right\}$, an m-sequence also because: $\mathrm{S}\left(\mathrm{d}^{\mathrm{T}}\right)$ and $\mathrm{g}\left(\mathrm{d}^{\mathrm{T}}\right)$ are relative prime.
So, the subsequences in the interleaving are identical. Now, we need only to check $I_{p}^{T}$. It can be seen that $I_{p}^{T}$
calculated by [3] or lookup table [12] (similar to table 1, example 1) are identical also (they must be the same because they are derived from matrix M). That implies D-transform and trace function are equivalent in creating the sequences with interleaved structure.

## 4. Conclusion remarks and future works

In this contribution we try to compare some mathmatical Tools widely used in representation of specific sequences applied in such areas like: advanced communications, cryptography and automatic testing.

The matrix representation is very intuitive and easily understandable in describling the configuration, state transition. It can also be used for determination the particular memory content when the D-transform of either the sequences or the corresponding initial state are given. However, for the sequences of great length, the dimension of the Matrix is unbearable great, which makes the manipulation difficult.

The trace function is compact and very comfortable for investigation of the $m$-sequences in general and interleaved $m$-sequences in particular. Since the length of sequences defined in trace function is $L=p^{m}-1$ only, it cannot be used for an arbitrary interleaved sequences ( $L \neq p^{m}-1$ ) without some complicated modification as in Dtransform case. Furthermore, one cannot get the information about states and configuration of LFSR in trace representation. However, from shift sequence ( $e=I_{p}^{\mathrm{T}}$ ) one can also figure out the hardware implementation (interleaving order).

The D-transform representation is also short and easy to handle. Furthermore, it contains information about the state $(\mathrm{S}(\mathrm{D})$ ) and configuration, $\mathrm{g}(\mathrm{D})$ as well. Like Trace function representation, it also indicates the hardware implementation ( $\mathrm{I}_{\mathrm{p}}^{\mathrm{T}}$ ). We notice once more that D-transform can be applied for any periodic sequence without any new concept. We hope to prove this statement in coming papers!

The link between those tools are also pointed out, for the first time.
As mentioned in our introduction and $[1,2,3,4, \ldots]$ all the above requirements need to be carefully evaluated before application. The static properties of the sequences (correlation functions, symbol distribution...) have to satisfy some random criteria so that the processed signal can be seen as pseudo noise and the linear complexities have to be large enough to make the sequence estimation difficult [4,11,12,18,19,20]. The evaluation process is obviously more complicated so that the trace function and D-transform need to be modified correspondingly. However, due to the limited scope of this paper we concentrate only on the comparison of represenration methods and leave other issues such as analysis and sequence selection for the future contributions.

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