

necessary and sufficient conditions in (24) derived in the previous sections also hold when carrier phase delays are considered—that is, $|\mathbf{R}|$ is maximized when

$$\sum_{\ell=1}^L g_{\ell i} g_{\ell j} \boldsymbol{\Theta}_{\ell i} \mathbf{X}_{\ell} \boldsymbol{\Theta}_{\ell j}^{\top} + \mathbf{W}_{ij} = \frac{E_{ij}}{N} \mathbf{I}_N, \quad 1 \leq i, j \leq B. \quad (51)$$

And as before, it is possible that no such set of \mathbf{X}_{ℓ} exists, in which case, our results provide an upper/lower bound on sum capacity/TSC.

VI. CONCLUSION

The overall structure of collaborative but geographically dispersed bases is interesting in light of the proliferation of consumer wireless systems like 802.11 and the amount of dark fiber available from past fiber (over)deployments. In this correspondence, we considered an abstraction of such systems as multiple collaborating base stations and uniform channels between users and bases and derived bounds on sum capacity and TSC via structural properties of the received covariance matrix. We also showed that as compared to single-base systems, where maximizing sum capacity and minimizing TSC are equivalent problems, in multibase systems TSC and sum capacity optimization can lead to different results.

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Frequency Hopping Sequences With Optimal Partial Autocorrelation Properties

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Abstract—We classify some p^k -ary (p prime, k integer) generalized m -sequences and generalized Gordon–Mills–Welch (GMW) sequences of period $p^{2k} - 1$ over a residue class ring $\mathbf{R} = \text{GF}(p)[\xi]/(\xi^k)$ having optimal partial Hamming autocorrelation properties. In frequency hopping (FH) spread-spectrum systems, these sequences are useful for synchronizing process. Suppose, for example, that a transmitting p^k -ary FH patterns of period $p^{2k} - 1$ are correlated at a receiver. Usually, the length of a correlation window, denoted by L , is shorter than the pattern's overall period. In that case, the maximum value of the out-of-phase Hamming autocorrelation is lower-bounded by $\lceil \frac{L}{p^k+1} \rceil$ but the classified sequences achieve this bound with equality for any positive integer L .

Index Terms—Finite rings, frequency hopping, generalized Gordon–Mills–Welch (GGMW) sequences, Hamming correlation, partial autocorrelation.

I. INTRODUCTION

In frequency hopping multiple-access (FHMA) spread-spectrum systems employing orthogonal modulation, we have to use a set of frequency hopping patterns to minimize the maximum of Hamming out-of-phase autocorrelation and cross correlation to effectively discriminate between their own signals and reduce multiple-access interference (MAI). Specific methods to generate such sets originate from the properties of m -sequences, Reed–Solomon codes, or combinatorial methods used in the ring of integers mod p for appropriate prime p [1], [2]. For example, an optimal family of frequency hopping (FH) sequences having p^k (p is a prime and k is a positive integer) symbols can be easily constructed from m -sequence over a Galois field $\text{GF}(p)$ [3] or from a generalized m -sequence (GM) or a generalized Gordon–Mills–Welch (GGMW) sequence over a polynomial residue class ring [4], [5]. Such sequences have optimal periodic autocorrelation functions. However, usually the length of a correlation window is shorter than the period of the chosen FH sequence due to the limited synchronization time or hardware complexity. Moreover, the window length may vary from time to time depending on the channel conditions. In that case, the partial Hamming autocorrelation,

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rather than the full-period Hamming autocorrelation, will play a major role in determining the synchronization performance.

The partial Hamming correlation function between two sequences $X = \{x(j)\}$ and $Y = \{y(j)\}$, for a period N and the correlation window length L starting at t , is defined as follows:

$$H_{XY}(\tau; t | L) = \sum_{j=t}^{t+L-1} h[x(j), y(j + \tau)], \quad 0 \leq \tau < N \quad (1)$$

where $h[x, y]$ is a binary Hamming function determined as $h[x, y] = 1$ if $x = y$ and $h[x, y] = 0$ if $x \neq y$. If $t = 0$ and $L = N$, (1) represents the conventional periodic Hamming correlation function $H_{XY}(\tau)$ as defined in [3]. Then the maximum of the partial Hamming autocorrelation function (HAF) along with window length L is defined as

$$H(X | L) = \max_{0 < \tau < N, 0 \leq t < N} \{H_{XX}(\tau; t | L)\}. \quad (2)$$

In this correspondence, we classify GM and GGMW sequences having optimal partial Hamming autocorrelation properties irrespective of the length of the correlation window. The optimality of the partial Hamming autocorrelation property can be extended from the optimal criteria as presented in [3].

Definition 1: Let S be the set of all sequences of length N over a given alphabet A . We say that a sequence $X \in S$ is *strictly optimal* if $H(X | L) \leq H(X' | L)$ for all $L \leq N$ and all $X' \in S$.

II. GENERALIZED MAXIMAL LENGTH AND GMW SEQUENCES

Let R be a polynomial residue class ring defined by $R = \text{GF}(p)[\xi]/(w(\xi)^k)$, where $w(\xi)$ is an irreducible polynomial of degree m over $\text{GF}(p)$, $m \geq 1$. From this point, we will consider only the case where $m = 1$ or $R = \text{GF}(p)[\xi]/(\xi^k)$. In that case, any element $b \in R$ can be expressed via the *ideal basis representation*

$$b = b_0 + b_1\xi + \cdots + b_{k-1}\xi^{k-1} \quad (3)$$

where $b_i \in \text{GF}(p)$. Thus, R can be written as

$$R = \text{GF}(p) + \xi \text{GF}(p) + \cdots + \xi^{k-1} \text{GF}(p). \quad (4)$$

The Galois extension ring of R denoted as $\text{GR}(R, r)$ is defined as $R[x]/(f(x))$ where $f(x)$ is a basic monic irreducible polynomial of degree r over R . This $f(x)$ can be selected from the monic irreducible polynomials over $\text{GF}(p)$ since $\text{GF}(p)$ is a subring of R and any irreducible polynomial defined over the subring $\text{GF}(p)$ is obviously irreducible over R [5]. Similarly to (3) and (4), any element $\beta \in \text{GR}(R, r)$ and $\text{GR}(R, r)$ can be expressed as

$$\beta = \beta_0 + \beta_1\xi + \cdots + \beta_{k-1}\xi^{k-1}$$

$$\text{GR}(R, r) = \text{GF}(p^r) + \xi \text{GF}(p^r) + \cdots + \xi^{k-1} \text{GF}(p^r)$$

where $\beta_i \in \text{GF}(p^r)$.

For

$$\beta = \sum_{i=0}^{k-1} \beta_i \xi^i \in \text{GR}(R, r)$$

let the mapping $\sigma^s: \beta \mapsto \sum_{i=0}^{k-1} \beta_i p^{si} \xi^i$ denote a Frobenius automorphism of $\text{GR}(R, r)$. If $s|r$, the trace function $\text{Tr}_s^r(\cdot)$ from $\text{GR}(R, r)$ into its subring $\text{GR}(R, s)$ is calculated as

$$\begin{aligned} \text{Tr}_s^r(\beta) &= \sum_{i=0}^{(r/s)-1} \sigma^{si}(\beta) \\ &= \sum_{i=0}^{(r/s)-1} \sum_{j=0}^{k-1} \beta_j p^{sij} \xi^j = \sum_{j=0}^{k-1} \text{tr}_s^r(\beta_j) \xi^j \end{aligned} \quad (5)$$

where

$$\text{tr}_s^r(v) = \sum_{i=0}^{(r/s)-1} v^{p^{si}}$$

is the field trace function from $\text{GF}(p^r)$ to $\text{GF}(p^s)$. The trace function defined in (5) has the following properties:

- 1) $\text{Tr}_s^r(\beta) = \text{Tr}_s^r(\sigma^{si}(\beta))$, $\forall i$ and $\forall \beta \in \text{GR}(R, r)$;
- 2) $\text{Tr}_s^r(b\beta + c\gamma) = b\text{Tr}_s^r(\beta) + c\text{Tr}_s^r(\gamma)$, $\forall b, c \in \text{GR}(R, s)$ and $\forall \beta, \gamma \in \text{GR}(R, r)$;
- 3) for any fixed $b \in \text{GR}(R, s)$, the equation $\text{Tr}_s^r(\beta) = b$ has exactly $p^{k(r-s)}$ solutions.

When α is a root of a primitive basic irreducible polynomial $f(x)$ over $R = \text{GF}(p)[\xi]/(\xi^k)$ and the Galois extension ring is defined as $\text{GR}(R, r) = R[x]/(f(x))$, every GM sequence $S^\nu = \{s^\nu(i)\}$ over R has the following unique trace representation [5]:

$$s^\nu(i) = \text{Tr}_1^r(\nu \alpha^i), \quad \nu \in \text{GR}(R, r).$$

For $a = \sum_{i=0}^{k-1} a_i \xi^i \in \text{GR}(R, s)$, let us define a permutation monomial

$$\Psi^d: a \mapsto \sum_{i=0}^{k-1} a_i^d \xi^i.$$

Then, every GGMW sequence [5], extended from a GMW sequence over a finite field [6], [7], can be represented as

$$s^\nu(i) = \text{Tr}_1^s(\Psi^d[\text{Tr}_s^r(\nu \alpha^i)]), \quad \nu \in \text{GR}(R, r)$$

where $s|r$ and $\text{gcd}(d, p^s - 1) = 1$.

III. GM AND GGMW SEQUENCES WITH OPTIMAL PARTIAL AUTOCORRELATION PROPERTY

For a frequency hopping sequence X of period N and a given correlation window of length L ($\leq N$), we derive a lower bound on the maximum out-of-phase autocorrelation value $H(X | L)$, defined in (2). We use the special case $H(X) = H(X | N)$. We start from the minimum bound on $H(X)$ presented in [3].

Lemma 1: For every sequence $X = \{x(j)\}$ of length N over an alphabet A of size $|A| = m$

$$H(X) \geq \overline{H}(X) \geq \frac{(N-b)(N+b-m)}{m(N-1)} \quad (6)$$

where b is the least nonnegative residue of N modulo m and $\overline{H}(X)$ is the average out-of-phase value of $H_{XX}(\tau)$.

Using the preceding lemma, we can derive the following lower bound on the partial HAF maximum.

Corollary 1:

$$H(X | L) \geq \frac{L}{N} \frac{(N-b)(N+b-m)}{m(N-1)}. \quad (7)$$

Proof: Let us derive the average out-of-phase value of $H_{XX}(\tau; t | L)$ as defined in (1)

$$\begin{aligned} \overline{H}(X | L) &= \frac{\sum_{\tau=1}^{N-1} \sum_{t=0}^{N-1} H_{XX}(\tau; t | L)}{(N-1)N} \\ &= \frac{\sum_{\tau=1}^{N-1} L H_{XX}(\tau; 0 | N)}{(N-1)N} \\ &= \frac{L}{N} \frac{\sum_{\tau=1}^{N-1} H_{XX}(\tau)}{(N-1)} \\ &= \frac{L}{N} \overline{H}(X). \end{aligned}$$

Then $H(X | L) \geq \overline{H}(X | L)$ and (6) yields the result. \square

Now, we classify some *strictly optimal* GM or GGMW sequences which achieve the lower bound given in (7). First, let us choose a degree $2k$ primitive irreducible polynomial $f(x)$ over $\text{GF}(p)$ as a primitive basic irreducible polynomial over $R = \text{GF}(p)[\xi]/(\xi^k)$. Assume $f(\alpha) = 0$ and

$$\nu = \alpha^{\epsilon_0} + \alpha^{\epsilon_1}\xi + \cdots + \alpha^{\epsilon_{k-1}}\xi^{k-1} \in \text{GR}(R, 2k).$$

Then

$$s^\nu(i) = \text{Tr}_1^k\left(\Psi^d[\text{Tr}_k^{2k}(\nu\alpha^i)]\right)$$

will be a p^k -ary sequence, of period $p^{2k} - 1$, if all the α^{ϵ_j} 's are linearly independent over $\text{GF}(p)$ and $\text{gcd}(d, p^k - 1) = 1$. In that case, the maximum out-of-phase value calculated for a given L is bounded using (7), as

$$H(S^\nu | L) \geq \left\lceil \frac{L}{p^k + 1} \right\rceil. \quad (8)$$

Subsequently, the following theorem can be used to classify the *strictly optimal* GGMW sequences satisfying the equality in (8) for any positive integer L . Here L is at most $p^{2k} - 1$, the period of S^ν .

Theorem 1: Let $f(x)$ be a degree $2k$ primitive polynomial over $\text{GF}(p)$, $f(\alpha) = 0$, and $\text{gcd}(d, p^k - 1) = 1$. A GGMW sequence $\{s^\nu(i)\}$

$$s^\nu(i) = \text{Tr}_1^k\left(\Psi^d[\text{Tr}_k^{2k}(\nu\alpha^i)]\right), \\ \nu = \alpha^{\epsilon_0} + \alpha^{\epsilon_1}\xi + \alpha^{\epsilon_2}\xi^2 + \cdots + \alpha^{\epsilon_{k-1}}\xi^{k-1}$$

is *strictly optimal* if and only if $\alpha^{\epsilon_0 d}, \alpha^{\epsilon_1 d}, \alpha^{\epsilon_2 d}, \dots, \alpha^{\epsilon_{k-1} d}$ are linearly independent over $\text{GF}(p)$ and $e_i \equiv e_j \pmod{p^k + 1}$ for all $i, j, 0 \leq i, j \leq k - 1$.

Proof: Since $\alpha^{(\epsilon_j - \epsilon_0)d} \in \text{GF}(p^k)$ for all j if $e_j \equiv e_0 \pmod{p^k + 1}$

$$s^\nu(i) - s^\nu(i + \tau) = \sum_{j=0}^{k-1} \text{tr}_1^k\left(\alpha^{(e_j - e_0)d} \left[(\text{tr}_k^{2k}(\alpha^{i+\epsilon_0}))^d - (\text{tr}_k^{2k}(\alpha^{i+\tau+\epsilon_0}))^d \right] \right) \xi^j \quad (9)$$

for a GGMW sequence S^ν and a fixed nonzero τ . Then (9) will be equal to zero exactly when all ξ^j 's coefficients are zero, or equivalently

$$\text{tr}_1^k\left(\alpha^{(e_j - e_0)d} \left[(\text{tr}_k^{2k}(\alpha^{i+\epsilon_0}))^d - (\text{tr}_k^{2k}(\alpha^{i+\tau+\epsilon_0}))^d \right] \right) = 0 \quad (10)$$

for all $j, 0 \leq j \leq k - 1$. Since all the $\alpha^{(e_j - e_0)d}$'s are linearly independent over $\text{GF}(p)$ they form a basis for $\text{GF}(p^k)$. Therefore, (10) occurs only if

$$(\text{tr}_k^{2k}(\alpha^{i+\epsilon_0}))^d - (\text{tr}_k^{2k}(\alpha^{i+\tau+\epsilon_0}))^d = 0 \quad (11)$$

as proven in [8]. However, since the given monomial is a simple permutation, the number of solutions for $0 \leq i \leq p^{2k} - 2$ in (11) is the same for any value of d , namely, $d = 1$. Therefore,

$$H_{S^\nu S^\nu}(\tau; t | L) \\ = \left| \{i | s^\nu(i) - s^\nu(i + \tau) = 0, t \leq i \leq t + L - 1\} \right| \\ = \left| \{i | \text{tr}_k^{2k}(\alpha^{i+\epsilon_0}(1 - \alpha^\tau)) = 0, t \leq i \leq t + L - 1\} \right|. \quad (12)$$

To evaluate (12) with a window length L and a nonzero τ given, we need the following lemma as explained in [1].

Lemma 2: Let m -sequence $b(i)$ over $\text{GF}(p^s)$, p prime, be defined as

$$b(i) = \text{tr}_s^{rs}(\alpha^i)$$

where α is a primitive element of $\text{GF}(p^{rs})$, and let $T = (p^{rs} - 1)/(p^s - 1)$. Then every segment of T consecutive symbols from $b(i)$ contains exactly $(p^{(r-1)s} - 1)/(p^s - 1)$ zeros.

Applying Lemma 2, we observe that an m -sequence represented by $\text{tr}_k^{2k}(\alpha^i)$ produces only one zero symbol in every segment of $p^k + 1$ consecutive indices i . Since $\text{tr}_k^{2k}(\alpha^{i+\epsilon_0}(1 - \alpha^\tau))$ is a cyclic-shifted version of $\text{tr}_k^{2k}(\alpha^i)$ when $\tau \neq 0$, (12) becomes

$$H_{S^\nu S^\nu}(\tau; t | L) = \begin{cases} \frac{L}{p^k + 1}, & \text{if } (p^k + 1) | L \\ \left\lceil \frac{L}{p^k + 1} \right\rceil - 1, & \text{otherwise.} \end{cases}$$

This proves that

$$H(S^\nu | L) = \lceil L/(p^k + 1) \rceil. \quad (13)$$

Conversely, assume that S^ν is *strictly optimal*. For $L = 1$ and an arbitrary but fixed τ , it is obvious that t^* exists such that

$$H_{S^\nu S^\nu}(\tau; t^* | 1) = 1$$

where $s^\nu(t^*) - s^\nu(t^* + \tau) = 0$. Since S^ν satisfies the bound (13) for any L , it yields

$$H_{S^\nu S^\nu}(\tau; t^* | L) = w$$

for $(w - 1)(p^k + 1) < L \leq w(p^k + 1)$. This produces

$$s^\nu(t^* + i) - s^\nu(t^* + i + \tau) = \begin{cases} 0, & \text{if } i \equiv 0 \pmod{p^k + 1} \\ \text{nonzero,} & \text{otherwise.} \end{cases} \quad (14a) \quad (14b)$$

Since (14a) indicates that

$$s^\nu(t^* + j(p^k + 1)) - s^\nu(t^* + j(p^k + 1) + \tau) = 0$$

for all $j, 0 \leq j < p^k - 1$, each ξ^l 's coefficient of the equation must be zero as

$$\text{tr}_1^k\left(\alpha^{j(p^k + 1)d} \left[(\text{tr}_k^{2k}(\alpha^{t^* + \epsilon_l}))^d - (\text{tr}_k^{2k}(\alpha^{t^* + \epsilon_l + \tau}))^d \right] \right) = 0. \quad (15)$$

As j varies from 0 to $p^k - 2$, $\alpha^{j(p^k + 1)d}$ passes through all the elements in $\text{GF}(p^k)$. This indicates that the difference of two inner traces in (15) will be zero for the same reason as was applied to (10) to produce (11). Because of the permutation property of the given monomial, applying $d = 1$ to this equation will not change the solutions of e_l with t^* and τ fixed. This yields the following equation involving e_l :

$$\text{tr}_k^{2k}\left(\alpha^{t^* + \epsilon_l}(1 - \alpha^\tau)\right) = 0. \quad (16)$$

Since an m -sequence $\text{tr}_k^{2k}(\alpha^e)$ produces only one zero symbol in every segment of $p^k + 1$ consecutive indices e , we have $e_i \equiv e_j \pmod{p^k + 1}$ for all $i, j, 0 \leq i, j \leq k - 1$ in (16).

Next, to show that $\alpha^{\epsilon_0 d}, \alpha^{\epsilon_1 d}, \alpha^{\epsilon_2 d}, \dots, \alpha^{\epsilon_{k-1} d}$ are linearly independent, we assume the contrary. Then there exist c_0, c_1, \dots, c_{k-1} which are not all-zero in $\text{GF}(p)$ satisfying

$$\sum_{l=0}^{k-1} c_l \alpha^{\epsilon_l d} = 0.$$

Assuming that $c_l \neq 0$ and $\gamma \in \text{GF}(p^k)$, the following equation is obviously true:

$$\text{tr}_1^k(\alpha^{(\epsilon_l - \epsilon_0)d} \gamma) = - \sum_{j=0, j \neq l}^{k-1} \frac{c_j}{c_l} \text{tr}_1^k(\alpha^{(e_j - \epsilon_0)d} \gamma). \quad (17)$$

TABLE I
THREE 27-ARY GM SEQUENCES HAVING A PERIOD OF $3^6 - 1$

(e_0, e_1, e_2)	GM Sequences (Frequency Hopping Patterns)
(0, 1, 2)	0 0 0 9 3 1 0 0 18 15 5 1 0 9 12 13 4 1 18 6 2 9 3 10 21 7 20 15 23 25 26 ...
(0, 17, 100)	21 0 9 15 18 19 21 18 3 24 2 7 3 24 15 25 4 4 15 9 20 21 0 25 6 19 8 21 14 19 17 ...
(0, 28, 56)	24 3 6 15 24 22 21 6 18 18 5 1 24 15 0 25 4 13 9 15 14 21 18 4 3 4 20 3 26 1 2 ...

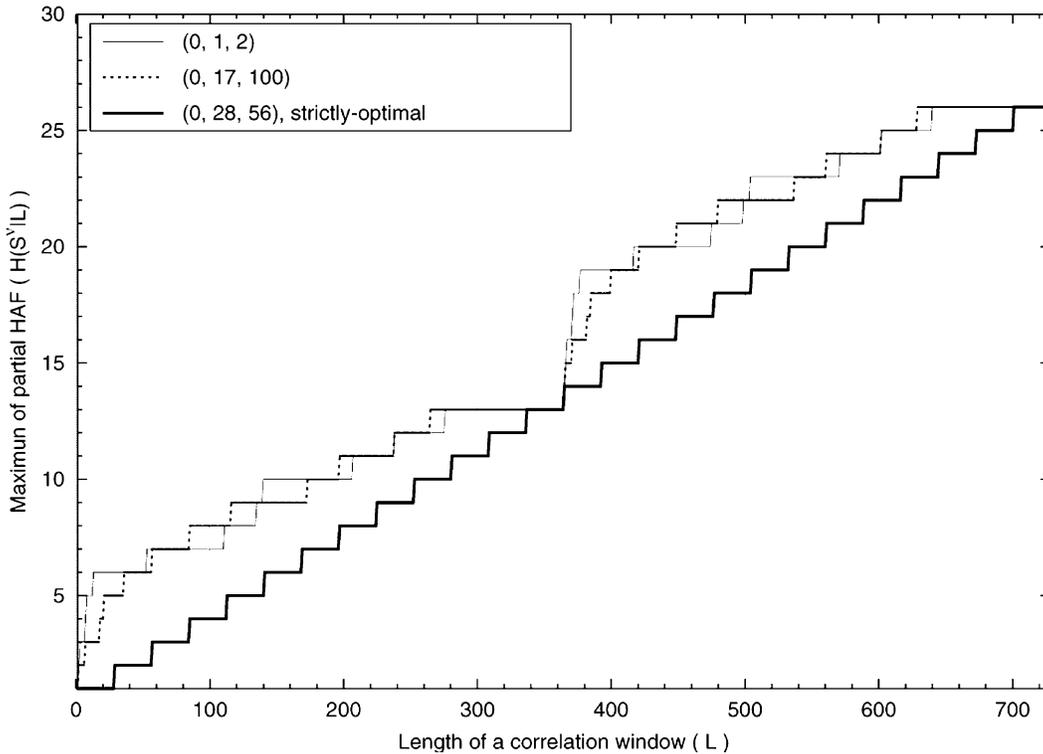


Fig. 1. $H(S^\nu|L)$ of three GM sequences represented by (e_0, e_1, e_2) .

Applying (17) to (9), the trace term of ξ^l 's coordinate can be described by a linear combination of the remaining trace terms. Since (9) is, in this case, related to the binary Hamming function of a GGMW sequence derived over $GR(R', r)$ where $R' = GF(p)[\xi]/\xi^{k-1}$, zero occurs in (9) at least $p^{k+1} - 1$ times during one period. This demonstrates that S^ν is not optimal even in the case of full-period autocorrelation, which is a contradiction. \square

For GM sequences, we can obtain a similar result.

Corollary 2: Let $f(x)$ be a degree $2k$ primitive polynomial over $GF(p)$ and $f(\alpha) = 0$. A GM sequence $\{s^\nu(i)\}$,

$$s^\nu(i) = \text{Tr}_1^{2k}(\nu\alpha^i), \quad \nu = \alpha^{e_0} + \alpha^{e_1}\xi + \alpha^{e_2}\xi^2 + \dots + \alpha^{e_{k-1}}\xi^{k-1}$$

is strictly optimal if and only if $\alpha^{e_0}, \alpha^{e_1}, \alpha^{e_2}, \dots, \alpha^{e_{k-1}}$ are linearly independent over $GF(p)$ and $e_i \equiv e_j \pmod{p^k + 1}$ for all $i, j, 0 \leq i, j \leq k - 1$.

Proof: Applying $d = 1$ in Theorem 1 yields this corollary. \square

Example 1: In Table I, we represent three GM sequences over $R = GF(3)[\xi]/\xi^3$ where

$$s^\nu(i) = \text{Tr}_1^6(\nu\alpha^i), \quad \nu = \alpha^{e_0} + \alpha^{e_1}\xi + \alpha^{e_2}\xi^2 \in GR(R, 6)$$

and α is a root of a primitive polynomial $x^6 + x + 2$ over $GF(3)$. Although both sequences $(e_0, e_1, e_2) = (0, 1, 2)$ and $(e_0, e_1, e_2) = (0, 17, 100)$ have the optimal periodic Hamming autocorrelation properties they are *not strictly optimal* as shown in Fig. 1. However, any

sequence satisfying Theorem 1 must be *strictly optimal*, for example, $(e_0, e_1, e_2) = (0, 28, 56)$ where any $e_a - e_b$ for $a \neq b$ is divisible by $28 = 3^3 + 1$, and $1, \alpha^{28}, \alpha^{2 \cdot 28}$ are linearly independent over $GF(3)$.

IV. CONCLUDING REMARKS

Optimal families of FH sequences can be constructed from these classified GM and GGMW sequences by using the same method as presented in [5]. Then all the sequences in such a family have the same optimal partial Hamming autocorrelation properties. In this correspondence, we have only considered the case in which $w(\xi) = \xi$ for $R = GF(p)[\xi]/(w(\xi)^k)$. Therefore, further study should focus on a more general case when the degree of $w(\xi)$ is greater than one.

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Compression Mappings on Primitive Sequences Over $Z/(p^e)$

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Abstract—Let $Z/(p^e)$ be the integer residue ring with odd prime $p \geq 5$ and integer $e \geq 2$. For a sequence \underline{a} over $Z/(p^e)$, there is a unique p -adic expansion $\underline{a} = \underline{a}_0 + \underline{a}_1 \cdot p + \dots + \underline{a}_{e-1} \cdot p^{e-1}$, where each \underline{a}_i is a sequence over $\{0, 1, \dots, p-1\}$, and can be regarded as a sequence over the finite field GF(p) naturally. Let $f(x)$ be a primitive polynomial over $Z/(p^e)$, and $G'(f(x), p^e)$ the set of all primitive sequences generated by $f(x)$ over $Z/(p^e)$. Set

$$\begin{aligned}\varphi_{e-1}(x_0, \dots, x_{e-1}) &= x_{e-1}^k + \eta_{e-2,1}(x_0, x_1, \dots, x_{e-2}) \\ \psi_{e-1}(x_0, \dots, x_{e-1}) &= x_{e-1}^k + \eta_{e-2,2}(x_0, x_1, \dots, x_{e-2})\end{aligned}$$

where $\eta_{e-2,1}$ and $\eta_{e-2,2}$ are arbitrary functions of $e-1$ variables over GF(p) and $2 \leq k \leq p-1$. Then the compression mapping

$$\varphi_{e-1} : \begin{cases} G'(f(x), p^e) \rightarrow \text{GF}(p)^\infty \\ \underline{a} \rightarrow \varphi_{e-1}(\underline{a}_0, \dots, \underline{a}_{e-1}) \end{cases}$$

is injective, that is, $\underline{a} = \underline{b}$ if and only if

$$\varphi_{e-1}(\underline{a}_0, \dots, \underline{a}_{e-1}) = \varphi_{e-1}(\underline{b}_0, \dots, \underline{b}_{e-1})$$

for $\underline{a}, \underline{b} \in G'(f(x), p^e)$. Furthermore, if $f(x)$ is a strongly primitive polynomial over $Z/(p^e)$, then

$$\varphi_{e-1}(\underline{a}_0, \dots, \underline{a}_{e-1}) = \psi_{e-1}(\underline{b}_0, \dots, \underline{b}_{e-1})$$

if and only if

$$\underline{a} = \underline{b} \text{ and } \varphi_{e-1}(x_0, \dots, x_{e-1}) = \psi_{e-1}(x_0, \dots, x_{e-1})$$

for $\underline{a}, \underline{b} \in G'(f(x), p^e)$.

Index Terms—Compressing mapping, integer residue ring, linear recurring sequence, primitive sequence.

I. INTRODUCTION

Suppose p is a prime and $R_e = Z/(p^e)$ is the integer residue ring modulo p^e , which can be also represented as $\{0, 1, \dots, p^e - 1\}$. In this correspondence, given positive integer $m \geq 2$, we always consider $a \pmod{m}$ as an element in $\{0, 1, \dots, m-1\}$.

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Let $f(x) = x^n + c_{n-1}x^{n-1} + \dots + c_0$ be a monic polynomial with degree $n \geq 1$ over R_e . A sequence $\underline{a} = (a(t))_{t \geq 0}$ over R_e satisfying the recursion

$$a(i+n) = -[c_0 a(i) + c_1 a(i+1) + \dots + c_{n-1} a(i+n-1)] \pmod{p^e}$$

for $i = 0, 1, 2, \dots$, is called a linear recurring sequence of degree n over R_e , generated by $f(x)$. $G(f(x), p^e)$ denotes the set of all sequences over R_e generated by $f(x)$. Reference [8] is a good introduction on linear recurring sequences over R_e .

Let $\underline{a} = (a(t))_{t \geq 0}$ and $\underline{b} = (b(t))_{t \geq 0}$ be sequences over R_e and $c \in R_e$. Define $\underline{a} + \underline{b} = (a(t) + b(t))_{t \geq 0}$, $c\underline{a} = (c \cdot a(t))_{t \geq 0}$, $\underline{a} \cdot \underline{b} = (a(t) \cdot b(t))_{t \geq 0}$, and the shift operator of sequence $x^k \underline{a} = (a(t+k))_{t \geq 0}$ for $k = 0, 1, 2, \dots$. So we have

$$G(f(x), p^e) = \{\underline{a} \in R_e^\infty \mid f(x)\underline{a} = \underline{0}\}.$$

Especially, we set

$$G'(f(x), p^e) = \{\underline{a} \in G(f(x), p^e) \mid \underline{a} \not\equiv \underline{0} \pmod{p}\}.$$

If $f(0) \not\equiv 0 \pmod{p}$, then there always exists a positive integer P such that $f(x)$ divides $x^P - 1$ over $Z/(p^e)$. The least such P is called the period of $f(x)$ over $Z/(p^e)$ and denoted by $\text{per}(f(x), p^e)$, which is upper-bounded by $p^{e-1}(p^n - 1)$, where $n = \deg f(x)$.

Definition 1: Let $f(x)$ be a monic polynomial of degree n over $Z/(p^e)$, then $f(x)$ is called a primitive polynomial if $\text{per}(f(x), p^e) = p^{e-1}(p^n - 1)$ (see [3], [7], and [19]).

Let $f(x)$ be a primitive polynomial of degree n over $Z/(p^e)$, then $f(x) \pmod{p^i}$ is also a primitive polynomial over $Z/(p^i)$, whose period is

$$\text{per}(f(x), p^i) = p^{i-1}(p^n - 1), \quad i = 1, 2, \dots, e-1.$$

In particular, $f(x) \pmod{p}$ is a primitive polynomial over the prime field GF(p), see [11]. Thus, we have

$$x^{p^i-1T} \equiv 1 + p^i h_i(x) \pmod{f(x)} \quad (1)$$

for $i = 1, 2, \dots, e-1$, where $T = p^n - 1$ and $h_i(x)$ is a polynomial over $Z/(p^e)$ of degree less than n satisfying $h_i(x) \not\equiv 0 \pmod{p}$. Clearly, $h_i(x)$ is coprime with $f(x) \pmod{p}$ over $Z/(p)$. Furthermore, we have [1], [7]

- 1) if $p = 2$, then $h_2(x) = \dots = h_{e-1}(x) \not\equiv 0 \pmod{2}$ and $h_2(x) = h_1(x) + h_1(x)^2 \pmod{f(x)}$;
- 2) if $p \geq 3$, then $h_1(x) = h_2(x) = \dots = h_{e-1}(x) \not\equiv 0 \pmod{p}$.

For the primitive polynomial $f(x)$ over $Z/(p^e)$, we always set

$$h(x) = h_1(x) \pmod{p} \quad (2)$$

where $h_1(x)$ is defined by (1). We call $f(x)$ a strongly primitive polynomial over $Z/(p^e)$, if $\deg h(x) \geq 1$ for the case of $p \geq 3$ or $p = e = 2$, and $\deg h_2(x) \geq 1$ for the case of $p = 2$ and $e \geq 3$. Obviously, we have $\deg f(x) \geq 2$ for the strongly primitive polynomial $f(x)$ over $Z/(p^e)$ with $e \geq 2$ (see [15]).