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## A Sharpened Dynamic Range of a Generalized Chinese Remainder Theorem for Multiple Integers

Huiyong Liao and Xiang-Gen Xia, Senior Member, IEEE


#### Abstract

A generalized Chinese remainder theorem (CRT) for multiple integers from residue sets has been studied recently, where the remainders in a residue set are not ordered. In this correspondence, we first propose a majority method and then based on the proposed majority method we present a sharpened dynamic range of multiple integers that can be uniquely determined from their residue sets.


Index Terms-Chinese remainder theorem (CRT), frequency determination from multiple undersampled waveforms, phase unwrapping, residue sets, sensor networks.

## I. INTRODUCTION

Chinese remainder theorem (CRT) has applications in many areas, such as computing, coding and cryptography, such as RSA-CRT and secret sharing, [8] and digital signal processing [7]. CRT gives a reconstruction of an integer from its remainders modulo several

[^0]smaller integers. The uniqueness of the reconstruction is possible if and only if the integer is smaller than the least common multiple (lcm) of the moduli that is the product of the moduli when all the moduli are co-prime. There are several generalizations of CRT, see for example [8]. Recently, a different generalization of CRT has been presented in [1]-[3]. In this generalized CRT, multiple integers are determined from their residue sets modulo several smaller integers, where the remainders in a residue set are known as the remainders of the multiple integers modulo a smaller integer but the correspondence of the remainders and the multiple integers is not known, i.e., the correct order of the remainders in a residue set is not known. As an example, consider two integers 60 and 64 and four moduli $5,7,11,13$. In this case, there are four residue sets from the two integers and the four moduli and they are $\{0,4\},\{1,4\},\{5,9\},\{8,12\}$ corresponding to the four moduli $5,7,11,13$, respectively. The problem is to uniquely determine the two integers from these four residue sets and four moduli, where the correspondence between the two integers and their remainders in a residue set is not specified, for example, in the second residue set $\{1,4\}$, it is not known whether 1 is the remainder of the first unknown integer or the second unknown integer modulo 7. Clearly, if the two integers are too large, the solution may not be unique similar to the conventional CRT. The problem we are interested in is how large the two integers can be so that they can be uniquely determined from their four residue sets (nonordered), which is called dynamic range in this correspondence. In the conventional CRT for a single integer, it is the product of the four prime moduli, i.e., $5 \cdot 7 \cdot 11 \cdot 13=5005$.

Based on the table look-up method, a dynamic range for the unique determination of the multiple integers has been presented in [1], where dynamic range means a range of integers within which multiple integers can be uniquely determined from the residue sets and the moduli. The dynamic range presented in [1] is sharpened and maximized in [2] when an additional condition on the multiple integers is imposed. More detailed descriptions of the problem and these results are stated in Section II. The motivation of the study of the above problem, i.e., the generalized CRT in [1]-[3] is the determination of multiple frequencies from multiple undersampled waveforms that may occur in, for example, phase unwrapping in synthetic aperture radar (SAR) imaging of moving targets [4], polynomial phase signal detection [5], and sensor networks where sensors have low power and low functionality [6].

In this correspondence, we propose a majority method for multiple integer determination from their residue sets. We present a sharpened dynamic range over the one presented in [1] of the unique determination of multiple integers from their residue sets when no additional condition on these integers is required. We also show an example that the sharpened dynamic range is not the maximal one, which means that further improvement is still possible.

This correspondence is organized as follows. In Section II, we describe the mathematical problem and some necessary notations. In Section III, we present a majority method for the determination and a sharpened dynamic range for multiple integers. In Section IV, we conclude this correspondence.

## II. Mathematical Problem Description

Suppose we have a set of distinct positive integers $S=\left\{N_{1}, N_{2}, \ldots, N_{\rho}\right\}$ and a set of positive integers, $P=\left\{p_{1}, p_{2}, \ldots, p_{\gamma}\right\}$, which, without loss of generality, are assumed relative co-prime, i.e., any two of $p_{r}, 1 \leq r \leq \gamma$, are co-prime, and $0<p_{1}<p_{2}<\cdots<p_{\gamma}$. The remainder (or residue) of $N_{l}$ modulo $p_{r}$ is

$$
\begin{equation*}
t_{l, r} \equiv N_{l} \bmod p_{r} \quad \text { for } 1 \leq l \leq \rho, \quad 1 \leq r \leq \gamma \tag{1}
\end{equation*}
$$

For $1 \leq r \leq \gamma$, define the residue set of $S$ modulo $p_{r}$

$$
\begin{equation*}
S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right) \triangleq \bigcup_{l=1}^{\rho}\left\{t_{l, r}\right\} \tag{2}
\end{equation*}
$$

Thus, we have $\gamma$ residue sets $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right), 1 \leq r \leq \gamma$. For each residue set $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right), 1 \leq r \leq \gamma$, there may be multiple integers in $S$ which share same residue, i.e., for each $r$, residues $t_{l, r}, l=1,2, \ldots, \rho$, may not be necessarily distinct. While all the distinct residues in each residue set $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ are known, the number of repeatings of any residue $t_{l, r}$ is not known. For each $r, 1 \leq r \leq \gamma$, we arrange the distinct elements in $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ in the following increasing order:

$$
\begin{equation*}
S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)=\left\{k_{l, r}: l=1,2, \ldots, \mu_{r}\right\} \tag{3}
\end{equation*}
$$

where $k_{l, r}<k_{m, r}$ for $1 \leq l<m \leq \mu_{r}$ and $\mu_{r}$ is the number of distinct elements of $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$. We define an onto mapping $\tau_{r}$ from the index set $I=\{1,2, \ldots, \rho\}$ of $S$ to the index set $J_{r}=$ $\left\{1,2, \ldots, \mu_{r}\right\}$ of $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ such that

$$
\begin{equation*}
t_{l, r}=k_{\tau_{r}(l), r} \quad \text { for } l=1,2, \ldots, \rho \tag{4}
\end{equation*}
$$

The mapping $\tau_{r}$ specifies the correspondence between integers in $S$ and residues in $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ for each $r$. Suppose the correspondence between residue set $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ and $p_{r} \in P$ for $1 \leq r \leq \gamma$ is specified, but the correspondence between $N_{l}$ and its remainder $k_{l, r}$ (or equivalently, the mapping $\tau_{r}$ ) is not known, i.e., the correct order of the remainders in a residue set is not known. Although $\tau_{r}$ is not known but exists.

The problem is to determine set $S$ of multiple integers $N_{1}, N_{2}, \ldots, N_{\rho}$ from the $\gamma$ residue sets $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ and their corresponding moduli $p_{r}$, where the correct order of the remainders in each residue set $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ is not known, $1 \leq r \leq \gamma$.
It is clear that, when $\rho=1$, the above problem is back to the conventional CRT and CRT provides a complete answer to the problem. As pointed out in [1], the difficulty of the above problem when $\rho>1$ comes from the fact that the correspondence between integers $N_{l}$ and their residues $k_{\tau_{r}(l), r}$ is not known, i.e., for any fixed $r$, it is not known with which integer $N_{i}$ a remainder $k_{l, r}$ satisfies $k_{l, r}=N_{i} \bmod p_{r}$, while we only know the residue set $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ that comes from a set of integers modulo $p_{r}, 1 \leq r \leq \gamma$.

The above problem has been studied in [1]-[3] motivated from multiple frequency determination using multiple undersampled waveforms as mentioned in Introduction. It can be briefly described as follows.

Consider $\gamma$ sensors with sampling rates $p_{r} \mathrm{~Hz}, 1 \leq r \leq \gamma$. Consider $\rho$ multiple frequencies $f_{1}=N_{1} \mathrm{~Hz}, \ldots, f_{\rho}=\bar{N}_{\rho} \mathrm{Hz}$ in a superpositioned waveform and these frequencies may include information interested and need to be accurately determined. At the $r$ th sensor, the received analog signal is of the following form:

$$
\begin{equation*}
x_{r}(t)=\sum_{l=1}^{\rho} A_{l, r} e^{2 \pi j f_{l} t}+w_{r}(t) \tag{5}
\end{equation*}
$$

where $A_{l, r}, 1 \leq l \leq \rho$, are nonzero complex coefficients and $w_{r}(t)$ is the additive white noise. The sampled signal at the $r$ th sensor with sampling rate $p_{r}$ Hertz is

$$
\begin{equation*}
x_{r}[n]=x_{r}\left(\frac{n}{p_{r}}\right)=\sum_{l=1}^{\rho} A_{l, r} e^{2 \pi j f_{l} n / p_{r}}+w_{r}\left(\frac{n}{p_{r}}\right) . \tag{6}
\end{equation*}
$$

The problem is to determine the multiple frequencies $f_{l}=N_{l}, 1 \leq$ $l \leq \rho$, from the above sampled data $x_{r}[n], 1 \leq r \leq \gamma$, where the sampling rates $p_{r}$ may be much lower than the signal frequencies $N_{l}$.

Based on the sampled data $x_{r}[n]$ at the $r$ th sensor, we take $p_{r}$-point DFT and obtain

$$
\begin{align*}
X_{r}[k] & =\mathrm{DFT}_{p_{r}}\left(x_{r}[n]\right) \\
& =\sum_{l=1}^{\rho} \sqrt{p_{r}} A_{l, r} \delta\left(k-t_{l, r}\right)+W_{r}[k] \tag{7}
\end{align*}
$$

for $0 \leq k \leq p_{r}-1$, where $t_{l, r}$ is the remainder of $N_{l}$ modulo $p_{r}$ and can be detected without the order information in terms of the index $l$. Thus, at the $r$ th sensor, what can be detected from the sampled waveform with sampling rate $p_{r}$ Hertz is the residue set $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ defined above and the frequency determination problem of $f_{l}, 1 \leq l \leq \rho$, precisely becomes the problem we described above. The case when there are errors in the detected residues $t_{l, r}$ has been considered in [6] with a lower dynamic range and this correspondence only considers the residue error free case.

Regarding the above problem, there are two questions. 1) When can the multiple integers in $S$ be uniquely determined from the residue sets $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ and $p_{r}$ for $\left.1 \leq r \leq \gamma ? 2\right)$ If the uniqueness is satisfied in 1), how can these multiple integers be determined? In [1], a dynamic range for the uniqueness of the determination of the multiple integers is given: If

$$
\begin{equation*}
\max \left\{N_{1}, N_{2}, \ldots, N_{\rho}\right\}<\max \left\{p, p_{1}, p_{2}, \ldots, p_{\gamma}\right\} \tag{8}
\end{equation*}
$$

where

$$
\begin{align*}
p & =\min _{1 \leq r_{1}<r_{2}<\cdots<r_{\eta} \leq \gamma} \operatorname{lcm}\left\{p_{r_{1}}, p_{r_{2}}, \ldots, p_{r_{\eta}}\right\} \\
& =p_{1} p_{2} \cdots p_{\eta} \tag{9}
\end{align*}
$$

where

$$
\begin{equation*}
\eta=\left\lfloor\frac{\gamma}{\rho}\right\rfloor \quad \text { or } \quad \gamma=\eta \rho+\theta \tag{10}
\end{equation*}
$$

for some $0 \leq \theta<\rho$. In [1], the determination method is basically look-up table method. In [3], an efficient (but may be still complicated) determination algorithm is proposed, which can be thought of as a generalization of CRT. As a special case, when $\max \left\{p, p_{1}, p_{2}, \ldots, p_{\gamma}\right\}=$ $p_{\gamma}$ in (8), all integers $N_{1}, N_{2}, \ldots, N_{\rho}$ can be uniquely determined directly from the residue set $S_{\gamma}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ alone because in this case, all the integers $N_{1}, N_{2}, \ldots, N_{\rho}$ are the same as the remainders $k_{1, \gamma}, k_{2, \gamma}, \ldots, k_{\rho, \gamma}$ themselves, i.e., $S=S_{\gamma}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$.

The dynamic range in (8) is maximized in [2] with an efficient determination algorithm when an additional condition on the multiple integers is satisfied: if

$$
\begin{equation*}
\max _{1 \leq l_{1}<l_{2} \leq \rho}\left|N_{l_{1}}-N_{l_{2}}\right|<\frac{1}{2} \min \left\{p_{1}, p_{2}, \ldots, p_{\gamma}\right\} \tag{11}
\end{equation*}
$$

and

$$
\begin{align*}
& \max \left\{N_{1}, N_{2}, \ldots, N_{\rho}\right\}<l c m\left\{p_{1}, p_{2}, \ldots, p_{\gamma}\right\} \\
& =p_{1} p_{2} \cdots p_{\gamma} \tag{12}
\end{align*}
$$

then $N_{1}, N_{2}, \ldots, N_{\rho}$ can be uniquely determined. Clearly, the dynamic range in (12) is already the maximal possible one since it is even the maximal possible one for the conventional CRT, i.e., when $\rho=1$. However, the right hand side of (12) is not the dynamic range in general. One example is: $N_{1}=5, N_{2}=4$ and $p_{1}=2, p_{2}=3$. In this case, the right hand side of (12) is $p_{1} p_{2}=6$ and the two integers $N_{1}$ and $N_{2}$ are within the range in (12). The two residue sets are $\{0,1\}$ and $\{1,2\}$, respectively. It is not hard to see there is another solution for these two sets of residues: $\hat{N}_{1}=1, \hat{N}_{2}=2$. Another example is $N_{1}=208, N_{2}=209$ and $p_{1}=2, p_{2}=3, p_{3}=5, p_{4}=7$. In this case, $\rho=2, \gamma=4$, and the right hand side of (12) is $p_{1} p_{2} p_{3} p_{4}=210$,
and the two integers $N_{1}$ and $N_{2}$ are also within the range in (12). The four residue sets are $\{0,1\},\{1,2\},\{3,4\}$, and $\{5,6\}$, respectively. One can see that $\hat{N}_{1}=13$ and $\hat{N}_{2}=194$ form another solution for the two integers $N_{1}$ and $N_{2}$, i.e., they share the same residue sets as $N_{1}$ and $N_{2}$ do.

The problems of interest of this correspondence are whether we can improve the dynamic range (8) for general multiple integers without any additional condition and how we can determine these multiple integers.

## III. Majority Method and an Improved Dynamic Range

To introduce a majority method, we first introduce some notations. An $m$-partition $\pi$ of $P=\left\{p_{1}, p_{2}, \ldots, p_{\gamma}\right\}$ is defined as a decomposition of $P$ into a union of its $m$ disjoint subsets $\left\{P_{1}^{\pi}, P_{2}^{\pi}, \ldots, P_{m}^{\pi}\right\}$ where a subset $P_{i}^{\pi}$ of $P$ can be empty, i.e.,

$$
\pi: P \rightarrow\left\{P_{1}^{\pi}, P_{2}^{\pi}, \ldots, P_{m}^{\pi}\right\}
$$

where $P=P_{1}^{\pi} \cup P_{2}^{\pi} \cup \cdots \cup P_{m}^{\pi}$ and $P_{i}^{\pi} \cap P_{j}^{\pi}=\emptyset$ for $1 \leq i \neq j \leq m$, and $P_{i}^{\pi}$ can be the empty set. For $1 \leq i \leq m$, we define $b_{i}^{\pi}$ as the product of integers $p_{r} \in P_{i}^{\pi}$ if $P_{i}^{\pi}$ is not empty and 1 if $P_{i}^{\pi}=\emptyset$, i.e.,

$$
b_{i}^{\pi} \triangleq \begin{cases}\prod_{p_{r} \in P_{i}^{\pi}} p_{r}, & \text { if } P_{i}^{\pi} \neq \emptyset  \tag{13}\\ 1, & \text { if } P_{i}^{\pi}=\emptyset\end{cases}
$$

We define $b^{\pi}$ as the minimum of $b_{i}^{\pi}$ for $1 \leq i \leq m$ and $c^{\pi}$ as the maximum of $b_{i}^{\pi}$ for $1 \leq i \leq m$, i.e.,

$$
\begin{equation*}
b^{\pi} \triangleq \min _{1 \leq i \leq m} b_{i}^{\pi} \quad \text { and } \quad c^{\pi} \triangleq \max _{1 \leq i \leq m} b_{i}^{\pi} \tag{14}
\end{equation*}
$$

Clearly, $c^{\pi} \geq b^{\pi}$ for any $m$-partition $\pi$. Let $b(m)$ be the maximal $b^{\pi}$ and $c(m)$ be the minimum $c^{\pi}$ among all the $m$-partitions $\pi$ of $P$, i.e.,

$$
b(m) \triangleq \max _{m-\text { partition } \pi \text { of } P} b^{\pi}
$$

and

$$
\begin{equation*}
c(m) \triangleq \min _{m-\text { partition } \pi \text { of } P} c^{\pi} \tag{15}
\end{equation*}
$$

which can be calculated as long as a modulus set $P$ is given while it may be complicated to do when the size of $P$ is large. Let $\pi_{1}$ and $\pi_{2}$ be the two $m$-partitions of $P$ with which the maximum and the minimum in (15) are reached, respectively. Then

$$
\begin{align*}
(b(m))^{m} & =\left(b^{\pi_{1}}\right)^{m} \leq \prod_{i=1}^{m} b_{i}^{\pi_{1}}=\prod_{i=1}^{\gamma} p_{i}=\prod_{i=1}^{m} b_{i}^{\pi_{2}} \\
& \leq\left(c^{\pi_{2}}\right)^{m}=(c(m))^{m} . \tag{16}
\end{align*}
$$

Thus

$$
\begin{equation*}
b(m) \leq c(m) \tag{17}
\end{equation*}
$$

We now introduce a majority method. Let $\pi$ be the $m$-partition with that the maximal $b(m)$ in (15) is reached. Assume all the integers $N_{l}$ in $S$ satisfy

$$
\begin{equation*}
\max \left\{N_{1}, N_{2}, \ldots, N_{\rho}\right\}<\min \{c(\rho), b(m)\} . \tag{18}
\end{equation*}
$$

For each $r, 1 \leq r \leq \gamma$, let $\sigma_{r}$ be an arbitrarily chosen onto-mapping from the index set $I=\{1, \ldots, \rho\}$ to the index set $J_{r}$ of the elements in $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$

$$
S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)=\bigcup_{j=1}^{\rho}\left\{k_{\sigma_{r}(j), r}\right\}
$$

TABLE I
Remainder Table

| integer | $\bmod p_{1}$ | $\bmod p_{2}$ | $\cdots$ | $\bmod p_{\gamma}$ |
| :---: | :---: | :---: | :--- | :---: |
| $\hat{N}_{1}$ | $k_{\sigma_{1}(1), 1}$ | $k_{\sigma_{2}(1), 2}$ | $\cdots$ | $k_{\sigma_{\gamma}(1), \gamma}$ |
| $\hat{N}_{2}$ | $k_{\sigma_{1}(2), 1}$ | $k_{\sigma_{2}(2), 2}$ | $\cdots$ | $k_{\sigma_{\gamma}(2), \gamma}$ |
| $\cdots$ | $\cdots$ | $\cdots$ | $\cdots$ | $\cdots$ |
| $\hat{N}_{\rho}$ | $k_{\sigma_{1}(\rho), 1}$ | $k_{\sigma_{2}(\rho), 2}$ | $\cdots$ | $k_{\sigma_{\gamma}(\rho), \gamma}$ |

For each subset $P_{i}^{\pi}, 1 \leq i \leq m$, we calculate the positive integers $N_{j}^{i}$ with

$$
\begin{equation*}
0 \leq N_{j}^{i}<\min \{c(\rho), b(m)\} \leq b(m) \leq b_{i}^{\pi} \tag{19}
\end{equation*}
$$

for $1 \leq j \leq \rho$ by using the conventional CRT such that

$$
\begin{equation*}
N_{j}^{i} \equiv k_{\sigma_{r}(j), r} \bmod p_{r}, \quad \forall p_{r} \in P_{i}^{\pi} \tag{20}
\end{equation*}
$$

where $k_{\sigma_{r}(j), r} \in S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ as we see from the above definitions. Note that, for an arbitrary mapping $\sigma_{r}$ above, the integers $N_{j}^{i}$ in (19) may not exist. However, if the mapping $\sigma_{r}$ is the correct mapping (or the correspondence between the integers $N_{j}$ and their remainders $\left.k_{\tau_{r}(j), r}\right)$, i.e., $\sigma_{r}=\tau_{r}$, although it may not have to be, integers $N_{j}^{i}$ in (19) do exist because the remainders are the true remainders from the set of integers $\left\{N_{1}, N_{2}, \ldots, N_{\rho}\right\}$ modulo $\left\{p_{1}, p_{2}, \ldots, p_{\gamma}\right\}$, and the assumption (18) for integers $N_{j}$ that are within the dynamic ranges of the conventional CRT for single integers. If an integer $N_{j}^{i}$ in (19) does not exist, we know that the mapping $\sigma_{r}$ is not a correct mapping, i.e., $\sigma_{r} \neq \tau_{r}$, and we then arbitrarily choose another onto-mapping from the index set $I=\{1, \ldots, \rho\}$ to the index set $J_{r}$ of the elements in $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ until integers $N_{j}^{i}$ in (19) do exist. Also note that although the searching process of the above mappings $\sigma_{r}$ so that the integers $N_{j}^{i}$ in (19) can be found may have a high complexity, we are interested more in the uniqueness of the determination than in the complexity of the determination in this correspondence. Therefore, as long as there exist mappings $\sigma_{r}$ such that (19) and (20) hold, it is sufficient for the results obtained later in this correspondence to hold.

Due to (19), for each valid mapping $\sigma_{r}$ described above, a reconstruction of integer $N_{j}^{i}$ in (20) is unique. Thus, we obtain an integer set $\hat{S}_{i}=\left\{N_{1}^{i}, N_{2}^{i}, \ldots, N_{\rho}^{i}\right\}$ for each subset $P_{i}^{\pi}, 1 \leq i \leq m$, and for each valid mapping $\sigma_{r}$. We then compare these $m$ integer sets $\hat{S}_{i}, 1 \leq i \leq m$. Clearly, these integer sets depend on the choice of the arbitrarily chosen onto-mappings $\sigma_{r}$ from $I$ to $J_{r}$ for $1 \leq r \leq \gamma$. We are interested in the case when all these integer sets are the same and have $\rho$ distinct elements, i.e., $\hat{S}_{1}=\cdots=\hat{S}_{m}=\hat{S}=\left\{\hat{N}_{1}, \ldots, \hat{N}_{\rho}\right\}$ with $\hat{N}_{i} \neq \hat{N}_{j}$ for $i \neq j$. Let
$\Omega \triangleq\left\{\left(\sigma_{1}, \ldots, \sigma_{\gamma}\right): \hat{S}_{1}=\cdots\right.$

$$
\begin{equation*}
\left.=\hat{S}_{m} \text { contains } \rho \text { distinct elements }\right\} \tag{21}
\end{equation*}
$$

It is not hard to see that the set $\Omega \neq \emptyset$ since $\left(\tau_{1}, \ldots, \tau_{\gamma}\right) \in \Omega$, where $\tau_{r}$ is the mapping from $I$ to $J_{r}$ defined before, due to the facts (14) and (18)-(20).

Suppose we have already found $\gamma$ onto-mappings $\sigma_{1}, \ldots, \sigma_{\gamma}$ for all the residue sets such that $\hat{S}_{1}=\hat{S}_{2}=\cdots=\hat{S}_{m}=\left\{\hat{N}_{1}, \ldots, \hat{N}_{\rho}\right\}$ with $\hat{N}_{i} \neq \hat{N}_{j}$ for $i \neq j$. Let us look at Table I. Note that the existence of such maps $\sigma_{r}$ is ensured by the fact that the set $\Omega$ is not empty.

For $1 \leq s \leq \rho$ and $1 \leq l \leq \rho$, we define set

$$
\begin{equation*}
\hat{Q}_{s}^{l} \triangleq\left\{p_{r}: \sigma_{r}(l)=\tau_{r}(s), 1 \leq r \leq \gamma\right\} . \tag{22}
\end{equation*}
$$

Then, for each $l, 1 \leq l \leq \rho, P=\bigcup_{s=1}^{\rho} \hat{Q}_{s}^{l}$. In order to form a $\rho$-partition of $P$, we define

$$
\begin{equation*}
Q_{s}^{l} \triangleq\left(\bigcup_{t=1}^{s} \hat{Q}_{t}^{l}\right)-\left(\bigcup_{t=1}^{s-1} \hat{Q}_{t}^{l}\right) \tag{23}
\end{equation*}
$$

for $s=1, \ldots, \rho$, such that $Q_{i}^{l} \cap Q_{j}^{l}=\emptyset$ for $1 \leq i \neq j \leq \rho$ and $\bigcup_{t=1}^{\rho} Q_{t}^{l}=P$. We denote the corresponding $\rho$-partition as $\mu^{l}$ for simplicity. For each fixed $s, 1 \leq s \leq \rho$, each fixed $l, 1 \leq l \leq \rho$, and any $p_{r} \in Q_{s}^{l}$, we have

$$
p_{r} \mid\left(\hat{N}_{l}-N_{s}\right)
$$

which is because when $p_{r} \in Q_{s}^{l}$, we have $\sigma_{r}(l)=\tau_{r}(s)$ and $k_{\sigma_{r}(l), r}=k_{\tau_{r}(s), r}=t_{s, r}$ and then it follows from (1) by replacing $l$ with $s$ in (1) and (20) by replacing $j$ with $l$ in (20). Thus, we have $\prod_{p_{r} \in Q_{s}^{l}} p_{r} \mid\left(\hat{N}_{l}-N_{s}\right)$, i.e., $b_{s}^{\mu^{l}} \mid\left(\hat{N}_{l}-N_{s}\right)$, and if $0 \leq \hat{N}_{l}, N_{s}<b_{s}^{\mu^{l}}$ then $\hat{N}_{l}=N_{s}$. Therefore, we have the following system of equations:

$$
\begin{align*}
& \hat{N}_{1}=N_{1}-a_{11} b_{1}^{\mu^{1}}=\cdots=N_{\rho}-a_{1 \rho} b_{\rho}^{\mu^{1}} \\
& \hat{N}_{2}=N_{1}-a_{21} b_{1}^{\mu^{2}}=\cdots=N_{\rho}-a_{2 \rho} b_{\rho}^{\mu^{2}}  \tag{24}\\
& \vdots \\
& \hat{N}_{\rho}=N_{1}-a_{\rho 1} b_{1}^{\mu^{\rho}}=\cdots=N_{\rho}-a_{\rho \rho} b_{\rho}^{\mu^{\rho}}
\end{align*}
$$

where the coefficient $a_{i j} \in \mathbb{Z}, 1 \leq i, j \leq \rho$, and $b_{i}^{\mu^{l}}$ is similarly defined as in (13) for an $m$-partition $\pi$ of $P$ when $m=\rho$.

We next find a dynamic range (a sufficient upper bound) for integers in $S$ such that once $\hat{S}_{1}=\hat{S}_{2}=\cdots=\hat{S}_{m}$ containing $\rho$ distinct elements occurs, or the above system of equations occur, we have $\hat{S}=$ $S$.

Theorem 1: An integer set $S$ above can be uniquely determined from its $\gamma$ residue sets $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ and moduli $p_{r}, 1 \leq r \leq \gamma$, by the above majority method if

$$
\begin{align*}
\max \left\{N_{1}, N_{2}, \ldots, N_{\rho}\right\}<\min \{c, b\} & \\
& = \begin{cases}\min \{c, b\}, & \text { when } \rho>2 \\
b, & \text { when } \rho=2\end{cases} \tag{25}
\end{align*}
$$

where $c$ and $b$ are defined similar to before:

$$
\begin{equation*}
c \triangleq \min _{\rho-\text { partition } \pi \text { of } P} c^{\pi} \quad \text { and } \quad b \triangleq \max _{2-\text { partition } \pi \text { of } P} b^{\pi} \tag{26}
\end{equation*}
$$

where $c^{\pi}$ and $b^{\pi}$ are defined in (14) for a $\rho$-partition $\pi$ and a 2-partition $\pi$ of $P=\left\{p_{1}, p_{2}, \ldots, p_{\gamma}\right\}$, respectively.

Proof: Let the 2-partition of $P$ achieving $b=$ $\max _{2-\text { partition } \pi \text { of } P b^{\pi} \quad \text { in }(25) \text { be } \pi_{0} \text {, i.e., } P=}^{=}$ $P_{1}^{\pi_{0}} \cup P_{2}^{\pi_{0}}, P_{1}^{\pi_{0}} \cap P_{2}^{\pi_{0}}=\emptyset$, and $b^{\pi_{0}}=b=\max _{2-\text { partition } \pi \text { of } P} b^{\pi}$. Using the majority method described above with $m=2$, we arrive at $\hat{S}_{1}=\hat{S}_{2}=\hat{S}=\left\{\hat{N}_{1}, \ldots, \hat{N}_{\rho}\right\}$ with $0 \leq \hat{N}_{i} \neq \hat{N}_{j}<\min \{c, b\}$ for $i \neq j$ which can be seen from (19) with $c=c(\rho)$ and $b=b(2)$. We next want to show $\hat{S}=S$.

For $1 \leq l \leq \rho$, we define set $\hat{Q}_{s}^{l}$ and $Q_{s}^{l}$ like (22) and (23), respectively, for $1 \leq s \leq \rho$. For each $l, 1 \leq l \leq \rho$, sets $Q_{s}^{l} \subset P, s=$ $1, \ldots, \rho$, also form a $\rho$-partition of $P$ and we denote it as $\mu^{l}$. We define $c^{\mu^{l}}=\max \left\{b_{1}^{\mu^{l}}, b_{2}^{\mu^{l}}, \ldots, b_{\rho}^{\mu^{l}}\right\}$. Clearly

$$
c^{\mu^{l}} \geq \min _{\rho-\text { partition } \pi \text { of } P} c^{\pi}=c .
$$

Therefore, $c^{\mu^{l}} \geq \min \{c, b\}$. Without loss of generality, we assume $c^{\mu^{l}}=b_{i_{l}}^{\mu_{l}^{l}}$. Thus, $b_{i_{l}}^{\mu^{l}} \geq \min \{c, b\}$. From (24), we obtain $b_{i_{l}}^{\mu^{l}} \mid\left(\hat{N}_{l}-\right.$ $\left.N_{i_{l}}\right)$. Combining this property with $0 \leq \hat{N}_{l}, N_{i_{l}}<\min \{c, b\} \leq b_{i_{l}}^{\mu^{l}}$,
we know $\hat{N}_{l}=N_{i_{l}} \in S$. This proves $\hat{S} \subset S$. Since both $\hat{S}$ and $S$ have $\rho$ elements, we conclude $\hat{S}=S$. The last equality in (25) is because $c=c(2) \geq b(2)=b$ from (17) when $\rho=2$.

It is not hard to see that the sequence $b(m)$ defined in (15) decreases in terms of $m$. Thus, $b=b(2)>b(m)$ for $m>2$. Therefore, although in the proof of Theorem 1, 2-partitions of $P$ are used in the majority method proposed above, the majority method for $m$-partitions also works, i.e., when $\hat{S}_{1}=\hat{S}_{2}=\cdots=\hat{S}_{m}=\hat{S}=\left\{\hat{N}_{1}, \hat{N}_{2}, \ldots, \hat{N}_{\rho}\right\}$ with $\hat{N}_{i} \neq \hat{N}_{j}$ for $i \neq j$, we then have $\hat{S}=S$. In other words, the following corollary holds.

Corollary 1: The above majority method using $m$-partitions with $m \geq 2$ provides a solution of $S$, i.e., $\hat{S}=S$.

In the following, we use the pigeon hole principle and 2-partitions of $P$ to obtain another dynamic range.

Theorem 2: When $\rho>2$, an integer set $S$ above can be uniquely determined from its $\gamma$ residue sets $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ and moduli $p_{r}, 1 \leq r \leq \gamma$, by the above majority method if

$$
\max \left\{N_{1}, N_{2}, \ldots, N_{\rho}\right\}<\prod_{i=1}^{\left\lceil\frac{\gamma}{\rho}\right\rceil} p_{i}
$$

Proof: Similar to the proof of Theorem 1, we now use a 2-partition of $P: P=P_{1} \bigcup P_{2}$, where $P_{1}=\left\{p_{1}, p_{2}, \ldots, p_{\left\lceil\frac{\gamma}{\rho}\right\rceil}\right\}$ and $P_{2}=$ $\left\{p_{\left\lceil\frac{\gamma}{\rho}\right\rceil+1}, \ldots, p_{\gamma}\right\}$. When $\rho>2$ and $\gamma \geq 2$, we have $\gamma-\left\lceil\frac{\rho}{\rho}\right\rceil \geq\left\lceil\frac{\gamma}{\rho}\right\rceil$. Thus

$$
\prod_{i=1}^{\left\lceil\frac{\gamma}{\rho}\right\rceil} p_{i}<\prod_{i=\left\lceil\frac{\gamma}{\rho}\right\rceil+1}^{\gamma} p_{i}
$$

due to the earlier assumption $p_{1}<p_{2}<\cdots<p_{\gamma}$. Following the steps of the proof of Theorem 1 until we obtain the partition sets $Q_{s}^{l}, s=$ $1, \ldots, \rho$, for $l=1, \ldots, \rho$. For each $1 \leq l \leq \rho$, all of the $\gamma$ integers in $P$ are put into $\rho$ subsets $Q_{s}^{l}, s=1, \ldots, \rho$. The pigeon hole principle states that there is one subset has at least $\left\lceil\frac{\gamma}{\rho}\right\rceil$ integers $p_{i} \in P$. Thus, for the $\rho$-partition, $Q_{s}^{l}, 1 \leq s \leq \rho$, of $P$, we have

$$
c^{\mu^{l}}=\max \left\{b_{1}^{\mu^{l}}, b_{2}^{\mu^{l}}, \ldots, b_{\rho}^{\mu^{l}}\right\} \geq \prod_{i=1}^{\left\lceil\frac{\gamma}{\rho}\right\rceil} p_{i}
$$

for $l=1, \ldots, \rho$. Without loss of generality, we assume $c^{\mu^{l}}=b_{i_{l}}^{\mu^{l}}$. Thus, $b_{i_{l}}^{\mu^{l}} \geq \prod_{i=1}^{\left\lceil\frac{\gamma}{\rho}\right\rceil} p_{i}$. From (24), we have

$$
\hat{N}_{l}=N_{i_{l}}-a_{l_{i}} b_{i_{l}}^{\mu_{l}^{l}}
$$

Since the condition $0 \leq \hat{N}_{l}, N_{i_{l}}<\prod_{i=1}^{\left\lceil\frac{\gamma}{\rho}\right\rceil} p_{i}$, we know that the above equality holds if and only if $a_{l i_{l}}=0$, equivalently, $\hat{N}_{l}=N_{i_{l}} \in S$. This proves $\hat{S} \subset S$. Since both $\hat{S}$ and $S$ have $\rho$ elements, we conclude $\hat{S}=S$.

Combining the above two results, we obtain the following improved dynamic range of the generalized CRT.

Corollary 2: Let $b$ and $c$ be defined in (26) in Theorem 1. An integer set $S$ of $\rho$ distinct integers can be uniquely determined from its $\gamma$ residue sets $S_{r}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)$ and moduli $p_{r}, 1 \leq r \leq \gamma$, by the above majority method if

$$
\max \left\{N_{1}, N_{2}, \ldots, N_{\rho}\right\}<\max \left\{\min \{c, b\}, \prod_{i=1}^{\left\lceil\frac{\gamma}{\rho}\right\rceil} p_{i}, p_{\gamma}\right\}
$$

## Dynamic range comparison



Fig. 1. Dynamic range comparison.
when $\rho>2$, and

$$
\max \left\{N_{1}, N_{2}, \ldots, N_{\rho}\right\}<\max \left\{b, p_{\gamma}\right\}
$$

when $\rho=2$.
Proof: If $\max \left\{N_{1}, N_{2}, \ldots, N_{\rho}\right\}<p_{\gamma}$, then $S_{\gamma}\left(N_{1}, N_{2}, \ldots, N_{\rho}\right)=S=\left\{N_{1}, N_{2}, \ldots, N_{\rho}\right\}$ as mentioned in Section II. The rest follows from Theorems 1 and 2 directly.

It is not hard to see that the new dynamic range presented in Corollary 2 is greater than the one (8) in [1] when there are more than two moduli, i.e., $\gamma>2$ :

$$
\begin{equation*}
\min \{c, b\}>p_{1} p_{2} \cdots p_{\eta} \tag{27}
\end{equation*}
$$

which is because of the following argument. By the definition of $c$ in (26), (14), and (15) where $c=c(\rho)$, we immediately have $c>$ $p_{1} p_{2} \cdots p_{\eta}$. When $\gamma>2$, we specify a 2 -partition $\pi$ of $P$ such that $P=\left\{p_{1}, p_{3}, \ldots, p_{2\lceil\gamma / 2\rceil-1}\right\} \cup\left\{p_{2}, p_{4}, \ldots, p_{2\lfloor\gamma / 2\rfloor}\right\}$. Thus, from (26), (14), and (15), we know $b=b(2) \geq b^{\pi}>p_{1} p_{2} \cdots p_{\lfloor\gamma / 2\rfloor} \geq$ $p_{1} p_{2} \cdots p_{\eta}$ since $\lfloor\gamma / 2\rfloor \geq \eta=\lfloor\gamma / \rho\rfloor$ when $\rho \geq 2$. When $\gamma=2$, both dynamic ranges in Corollary 2 and (8) in [1] become trivial, i.e., $p_{1}$. When $\rho=1$, it reduces to the conventional CRT.

As example, let us consider the case of $p_{1}=5, p_{2}=7, p_{3}=$ $11, p_{4}=13$, and $\rho=2$. We want to determine two integers $N_{1}, N_{2}$ from their residue sets. From Corollary 2, we conclude that they can be uniquely determined if $\max \left\{N_{1}, N_{2}\right\}<65$. Comparing with the one in (8) in [1], $\max \left\{N_{1}, N_{2}\right\}<35$, one can see that the new dynamic range, 65 , obtained in this correspondence almost doubles 35 previously obtained in [1]. The improvement becomes more significant when the size of the modulus set $P$ becomes larger, which can be seen from Fig. 1. Regarding to the application of multiple frequency determination proposed in [1], the maximal frequency in this example
is 65 Hz in a superposition of two harmonic signals so that these two frequencies can be uniquely determined from four sensors with sampling rates $5,7,11$, and 13 Hz , respectively, based on the proposed majority method above, while the maximal frequency is 35 Hz based on the method proposed in [1]. Note that, the above dynamic range 65 is a sufficient range of two integers for their unique determination and it does not mean that two integers above 65 can not be uniquely determined, i.e., the above dynamic range may not be necessary as we shall see later in another example.

For a general $\rho$, the calculation of $c$ in (26) in the dynamic range in Corollary 2 may not be easy. Due to (16), it can be easily lower bounded by

$$
\begin{equation*}
c \geq\left\lceil\left(\prod_{i=1}^{\gamma} p_{i}\right)^{\frac{1}{\rho}}\right\rceil . \tag{28}
\end{equation*}
$$

Clearly, the lower bound of $c$ in (28) is greater than the dynamic range (8) obtained in [1]:

$$
\begin{equation*}
\left\lceil\left(\prod_{i=1}^{\gamma} p_{i}\right)^{\frac{1}{\rho}}\right\rceil>p_{1} p_{2} \cdots p_{\eta} \tag{29}
\end{equation*}
$$

where $\eta$ is defined in (10).
The lower bound of $c$ in (28) provides the following lower bound for the new dynamic range in Corollary 2.

Corollary 3: The dynamic range in Corollary 2 is lower bounded by

$$
\begin{equation*}
\max \left\{\min \left\{\left\lceil\left(\prod_{i=1}^{\gamma} p_{i}\right)^{\frac{1}{\rho}}\right\rceil, b\right\}, \prod_{i=1}^{\left\lceil\frac{\gamma}{\rho}\right\rceil} p_{i}, p_{\gamma}\right\} \tag{30}
\end{equation*}
$$

when $\rho>2$, and $\max \left\{b, p_{\gamma}\right\}$ when $\rho=2$.

In Fig. 1, we compare the existing dynamic range (8) in [1] with the new dynamic range $\max \left\{b, p_{\gamma}\right\}$ when $\rho=2$ and the lower bound (30) of the new dynamic range in Corollary 2 when $\rho>2$. In Fig. 1, the moduli $p_{1}=2, p_{2}, \ldots, p_{\gamma}$ are the $\gamma$ smallest primes, and $\rho=2,3,4,5$ and $\gamma=5,10,15,20,25$ are considered. The existing dynamic ranges are plotted with dashed lines and the new dynamic ranges or their lower bounds are plotted with solid lines. One can see that the improvement of the newly obtained dynamic ranges are significantly better than the existing ones.
On the other hand, the new dynamic range presented in Corollary 2 is still not necessary. As a counter example, let us consider the case of $p_{1}=5, p_{2}=7, p_{3}=11, p_{4}=13$, and $\rho=3$. In this case, the new dynamic range from Corollary 2 is 35 , i.e., if $\max \left\{N_{1}, N_{2}, N_{3}\right\}<35$, then these three nonnegative integers can be uniquely determined. This is not necessary. In fact, it is not hard to check that if $\max \left\{N_{1}, N_{2}, N_{3}\right\}<65$, we can uniquely reconstruct $N_{1}, N_{2}, N_{3}$ from their four residue sets as follows. Arrange $p_{1}, p_{4}$ as a group and $p_{2}, p_{3}$ as another group. Then, these three integers can be determined by using the majority method described before. We omit its details here.

## IV. CONCLUSION

In this correspondence, we further studied a generalized CRT for multiple integer determination from their residue sets and moduli. We first presented a majority method for the determination and then obtained an improved dynamic range over the existing one for the unique determination of multiple integers based on the proposed majority method. Besides the mentioned application in multiple frequency determination from multiple undersampled waveforms, such as, from low functionality sensors, the above generalized CRT can be applied to cryptography, for example, for secret sharing similar to the conventional CRT [8]. As a remark, the majority method for the multiple integer determination has a high complexity. Any simplified determination algorithm for the generalized CRT with the newly proposed dynamic range would be interesting.

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## A Note on the Optimal Quadriphase Sequences Families

Xiaohu H. Tang, Member, IEEE, and<br>Parampalli Udaya, Member, IEEE


#### Abstract

In this note, by using a modification of the families $\mathcal{B}$ and $\mathcal{C}$, we obtain a larger family of optimal quadriphase sequences, $\mathcal{D}$ over $Z_{4}$. In contrast to the families $\mathcal{B}$ and $\mathcal{C}$, the family $\mathcal{D}$ has the same length and the same maximal nontrival correlation value, but with double the size.


Index Terms—Galois ring, optimal sequences, quadriphase sequences.

## I. INTRODUCTION

In code-division multiple-access (CDMA) communication systems, nonbinary signature sequences are preferred over binary sequences as they offer 3-dB improvement in signal to interference ratio [2]. This is because the lower bound on smallest possible nontrivial correlation parameter $C_{\max }$ for non binary sequences is $\sqrt{2}$ times better than that for binary sequences [5]. Among the non binary alphabets, quadriphase sequences are preferred for signature sequences because of easy implementation of modulators and availability of optimal sequences.

In the early 1990 s, the theory of $\mathbf{Z}_{4}$ maximal length sequences was established, leading to the discovery of optimal quadriphase sequences meeting the Welch and Sidelnikov bounds [2], [6], [7]. Unlike in field case, the possible periods for sequences over $\mathbf{Z}_{4}$ are $2^{n}-1$ and $2\left(2^{n}-\right.$ $1)$, where $n$ is a positive integer. There are three optimal families derived as a sequences satisfying a linear recursion over $\mathbf{Z}_{4}$. The first basic optimal family is known as family $\mathcal{A}$ which comprises of $2^{n}+$ $1 \mathbf{Z}_{4}$ maximal length sequences [2]. The second optimal family known as family $\mathcal{B}$ [2] which can be seen as interleaved version of sequences in family $\mathcal{A}$. This family consists of $2^{n-1}$ sequences of period $2\left(2^{n}-1\right)$. A third optimal family not discussed in [2] exists with the same parameters as family $\mathcal{B}$ with $n$ odd integer [7]. We refer to this family as family $\mathcal{C}$. A complete treatment of all such families of trace sequences over $\mathbf{Z}_{4}$ is given in [7] which includes three more suboptimal families.

Quadriphase sequences based on a generalization of the above $\mathbf{Z}_{4}$ families have been adopted as spectrum spreading sequences in 3G wideband CDMA standards [4]. It is expected that fourth generation CDMA systems need to handle higher data rates of up to 1 Gbytes/s.

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X. H. Tang is with the Institute of Mobile Communications, Southwest Jiaotong University, Chengdu, China (e-mail: xhutang@ieee.org).
P. Udaya is with the Department of Computer Science and Software Engineering, University of Melbourne, VIC 3010, Australia (e-mail: udaya@cs.mu.oz.au).

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    The authors are with the Department of Electrical and Computer Engineering, University of Delaware, Newark, DE 19716 USA (e-mail: liao@ee.udel.edu; xxia@ee.udel.edu).

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